

TOTAL SYNTHESIS OF CYCLOOROIDIN AND STUDIES TOWARDS SOME
OROIDIN DIMERS

by

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ABSTRACT

TOTAL SYNTHESIS OF CYCLOOROIDIN AND STUDIES TOWARDS SOME OROIDIN DIMERS

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This dissertation consists of two parts. The first part describes the synthesis of northern fragment of nagelamide R and the enantioselective total synthesis of cyclooroidin. The oroidin family of marine natural products is a growing group of marine sponge-derived alkaloids which contain 2-aminoimidazole and pyrrolicarboxamide fragments as their signature features. The structural complexity, and in many cases interesting biological profiles, of these compounds has rendered several of these natural products as targets of interest. In the context of nagelamide R, the key transformation involves the intramolecular cyclization of a pyrrolicarboxamide via the carbonyl oxygen leading to the formation of oxazoline, whose subsequent elaboration by pyrrole

bromination and imidazole azidation provides the fully functionalized northern fragment of the natural product. This chemistry suggests a possible biosynthetic pathway to the formation of nagelamide R through nagelamide B via activation and substitution of the amide carbonyl.

The key steps en route to the enantioselective total synthesis of cyclooroidin involve a coupling reaction between an imidazole chlorohydrin and pyrrole 2-carboxylic acid followed by intramolecular cyclization to form an oxazine ring. Conversion into the corresponding pyrazine, followed by bromination of the pyrrole and introduction of the azide group at the C-2 position of the imidazole provided the complete natural product framework. Removal of the protecting group and reduction of the azide to amine provide synthetic enantioselective cyclooroidin.

The second part of this dissertation describes studies towards some oroidin dimers including palau'amine and ageliferin. These studies have resulted in a concise entry into the all *trans*-substituted spiro cyclopentyl imidazolone system found in palau'amine and related natural products. These structures are accessed through an intramolecular Diels-Alder reaction of an enyne followed by an oxidative rearrangement. The final section describes a preliminary investigation of methods for the stereoselective incorporation of the chloro moiety, which is present in several oroidin dimers, including palau'amine.

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PART I TOTAL SYNTHESIS OF CYCLOOROIDIN

CHAPTER 1

INTRODUCTION

1.1 Pyrrole-Imidazole Alkaloids

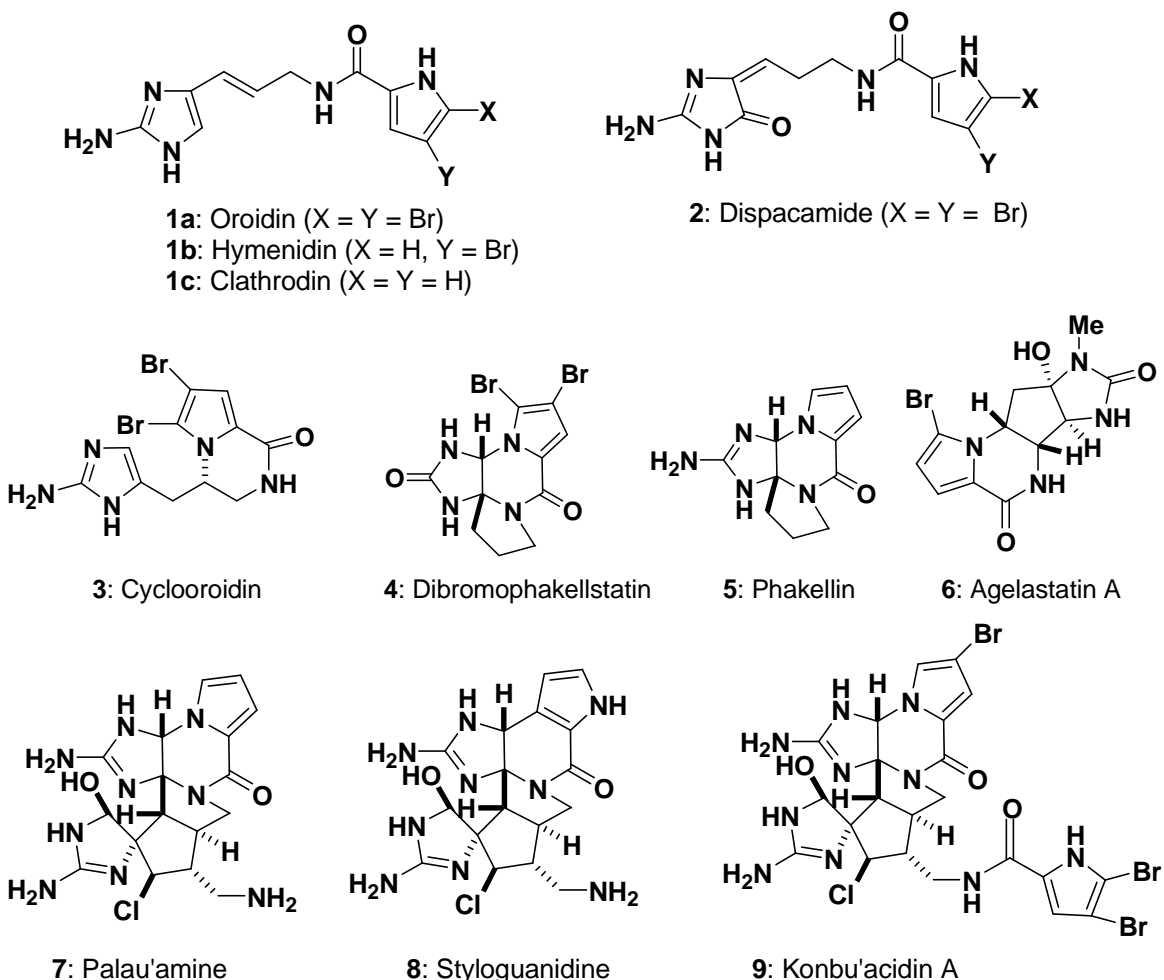


Figure 1.1 Pyrrole-imidazole marine alkaloids

Marine sponges have been, and continue to be, a rich source of structurally diverse pyrrole-imidazole alkaloids.¹⁻³ Over the past thirty years, many pyrrole-imidazole alkaloids with unprecedented skeletal diversity and a broad range of biological activities have been isolated and characterized.¹⁻³ The first member of

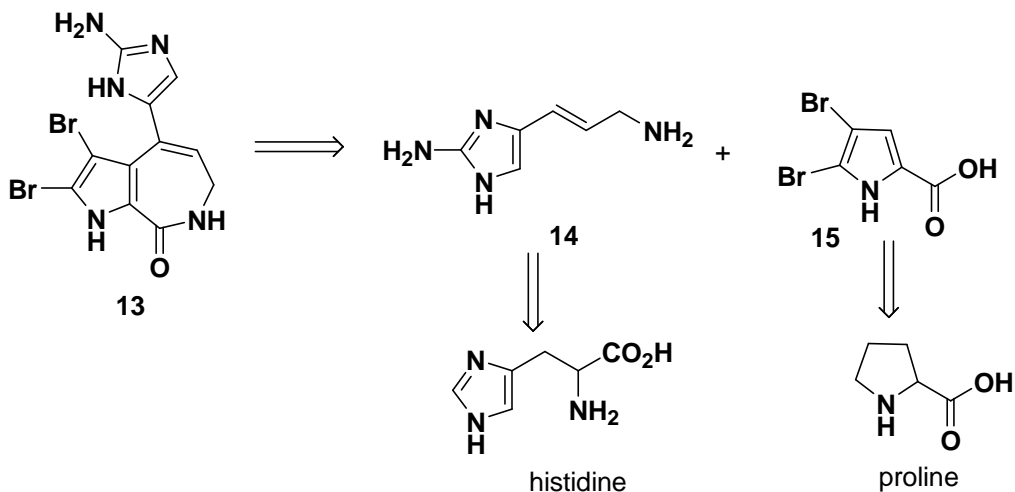
this family of alkaloids to be isolated was oroidin (**1a**), initially from *Agelas oroides* in 1971, and subsequently from a variety of other sponges.¹⁻³ Over the years, more elaborate and complex derivatives have been isolated and the whole family is frequently referred to either as the oroidin alkaloids or the pyrrole-imidazole alkaloids. Furthermore, these natural products are frequently characterized by the number of oroidin subunits and thus are described as oroidin monomers, dimers or tetramers. The more elaborate members of this family of natural products arise from various modes of functionalization, cyclization, and in some cases dimerization or tetramerization of the parent heterocycles **1a-c**, leading to the production of **2-9** (Figure 1.1).⁴

Inspired in large part by this family of natural products, our research group has been interested in the development of new and efficient synthetic methods for the elaboration of simple imidazole derivatives into highly functionalized polycyclic systems as key intermediates *en route* to the total synthesis of a number of the imidazole-containing natural products, including several illustrated in Figure 1.1. These approaches include: the inter- and intramolecular Diels-Alder reactions of 4-vinylimidazoles;^{5,6} the intramolecular radical cyclization reactions of imidazoles;⁷ ring-closing metathesis of dienylimidazoles and enynes;⁸ and oxidative chemistry.^{9,10a,10b} The work described in this dissertation represents a continuation of this effort and the development of a number of new reactions.

1.2 Biosynthetic pathways

Several groups have proposed biogenetic pathways for the assembly of pyrrole-imidazole alkaloids. Baran et al. recently proposed that several of the dimeric

members of the family come from a single precursor.¹¹ This proposal bears some resemblance to the more comprehensive hypothesis of Al Mourabit and Potier.³ However, despite numerous proposals in the literature there is very little described in terms of experimental work to support these hypotheses. Kerr et al. reported the first, and to date only biosynthetic study using cell culture of the sponge *Teichaxinella morchella*.¹² Using ¹⁴C-labeled proline and histidine they demonstrated that these amino acids are precursors of the monomer stevensine (**13**) through 2-aminoimidazolylprop-1-ene (**14**) and 4,5-dibromopyrrole carboxylic acid (**15**) (scheme 1.1). Unfortunately, this study did not reveal any details of the transformations involved.



Scheme 1.1

1.3 Cyclooridin

In 2000, Fattorusso et al. isolated a new alkaloid belonging to the pyrrole-imidazole alkaloid family from the Mediterranean sponge *Agelas oroides*, collected in the Bay of Naples, which they termed (-)-cyclooridin (**3**).¹⁴ It exhibited a negative optical rotation $[\alpha]_D -12$ (c 0.02, MeOH) and the absolute stereochemistry was established to be *S* by comparison of its CD spectrum with

the one reported for dibromophakellin. This alkaloid was also detected in an extract obtained from the Okinawan marine sponge *Agelas sp* (Figure 1.2).¹⁵ In 2006 three more natural products structurally related to cyclooroidin (**3**) were isolated. Agesamide A (**10**) and B (**11**) were obtained from the Okinawan marine sponge *Agelas sp*.¹⁶ Oxocyclostylidol **12** was extracted from the Caribbean derived sponge *Stylissa caribica*.¹⁷

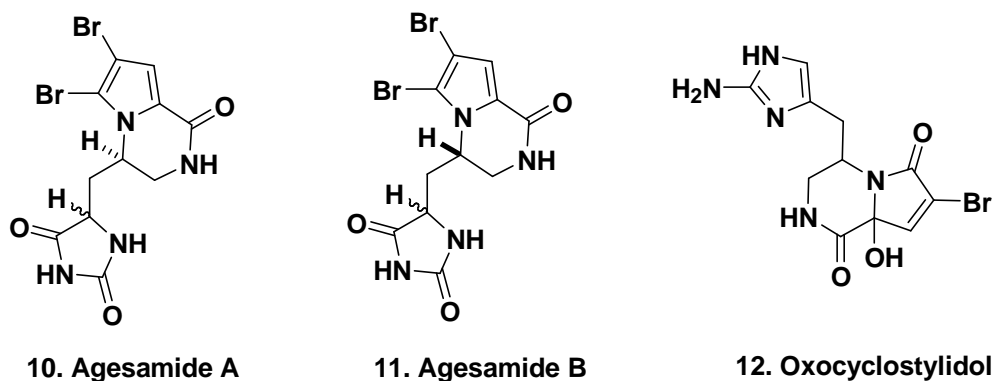
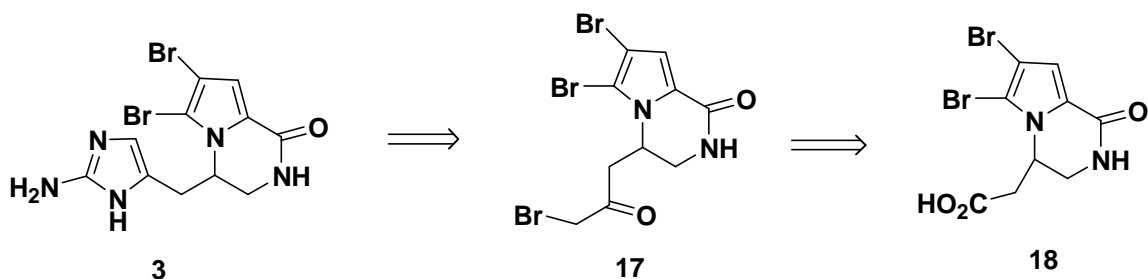


Figure 1.2 Pyrrole-imidazole marine alkaloids

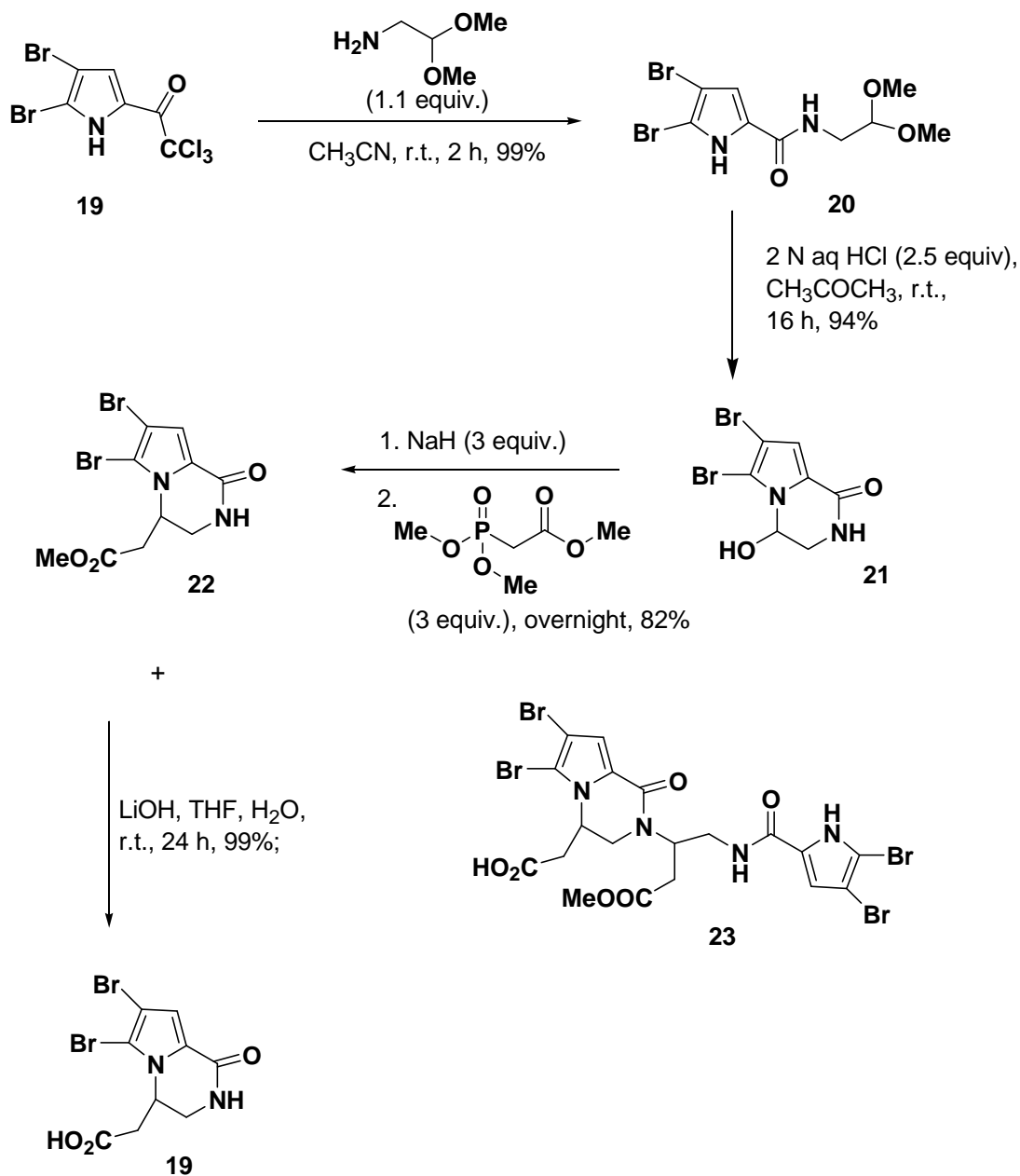
1.4 (±)-cyclooroidin by Papeo

As might be expected cyclooroidin has attracted attention from the synthetic community and very recently, two racemic syntheses have been described. The first one, published by Papeo et al.¹⁸ was achieved in nine steps with an overall yield of 10%. As depicted in (Scheme 1.2) the 2-aminoimidazole ring **16** was constructed from the corresponding α -bromoketone **17**, which was in turn derived from previously prepared from racemic longamide B (**18**)¹⁹ through a Wolff bromoketone synthesis.²⁰



Scheme 1.2

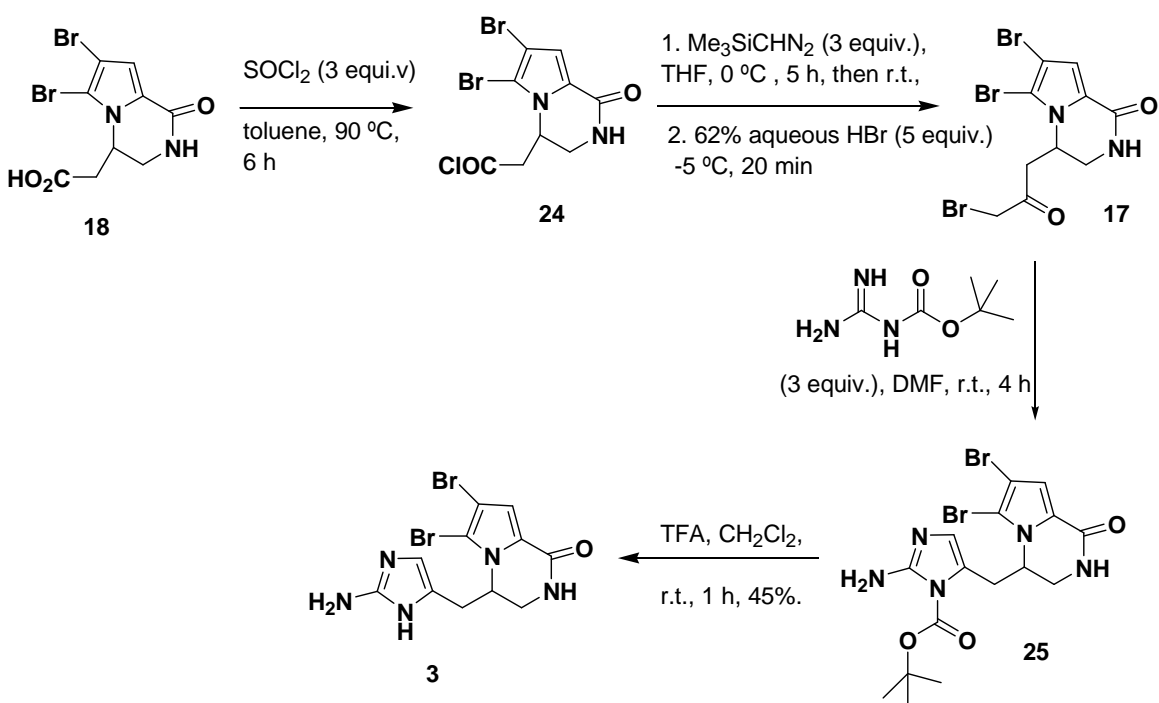
Their synthetic efforts started (Scheme 1.3) by following the previously disclosed protocol¹⁹ for the synthesis of racemic longamide B (**18**). 4,5-Dibromo-trichloroacetylpyrrole (**19**)²¹ was used as a starting material, which upon treatment with commercially available aminoacetaldehyde dimethyl acetal delivered amido-acetal **20**. Treating this amido-acetal with 2 N HCl resulted in the formation of intermediate **21** in quantitative yield.^{19,22} Then they attempted to convert the racemic carbinolamine **21** to longamide B methyl ester (**22**) by a published procedure using (MeONa/MeOH/methyl diethylphosphonoacetate).¹⁹ However, they obtained a complex reaction mixture, in which the amount of the desired product (**22**) was minor due to the presence of by-products, most of them arising from hydrolysis and competitive conjugate addition of methanol. Eventually longamide B methyl ester (**22**) was obtained by a modification of the reaction conditions and using NaH/trimethylphosphonoacetate in THF in 82% yield. Along with the expected product, they obtained a diastereomeric mixture (15/85 ratio based on HPLC analysis) of dimer **23**, arising from an intermolecular aza-Michael reaction. The yield of this diastereomeric mixture was up to 15%, depending on reaction time and reactant concentration.



Scheme 1.3

In the presence of LiOH in a mixture of THF/water at r.t. longamide B methyl ester (**22**) was then converted quantitatively into longamide B (**18**). Subsequently longamide B (**18**) was converted into the corresponding acyl chloride **24** through standard SOCl_2 treatment (Scheme 1.4). Reacting **24** with $\text{Me}_3\text{SiCHN}_2$ ²⁴ followed by exposure of the resulting α -diazoketone solution to aqueous HBr at 5 °C

delivered the α -bromoketone **17**. The crude α -bromoketone was immediately reacted with Boc protected guanidine, yielding the 15-Boc-protected cyclooroidin **25** (30% over four steps).²⁵ It was purified by extensive chromatography (AcOEt/MeOH) in order to remove unreacted Boc-guanidine. They postulated that the migration of the *tert*-butoxycarbonyl group from the exo- to the endocyclic nitrogen atom could occur during the purification step.²⁵ Removal of the protecting group, followed by column chromatography with a basic mobile phase (CH₂Cl₂/MeOH/NH₃ aq) delivered (\pm)-cyclooroidin (**16**) as the free base in 45% yield from **25**.

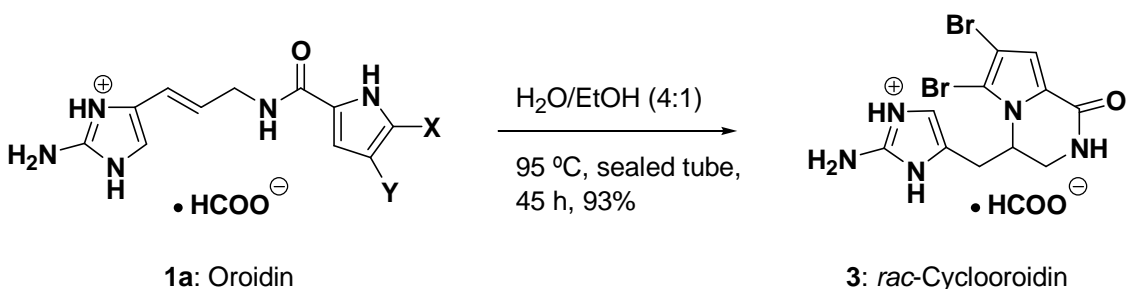


Scheme 1.4

1.5 (\pm)-cyclooroidin by Lindel

The second racemic synthesis was disclosed by Lindel and co-workers²⁶ which involved the intramolecular cyclization of oroidin formate in protic solvents (Scheme 1.5). The authors suggested that the biosynthesis of optically active

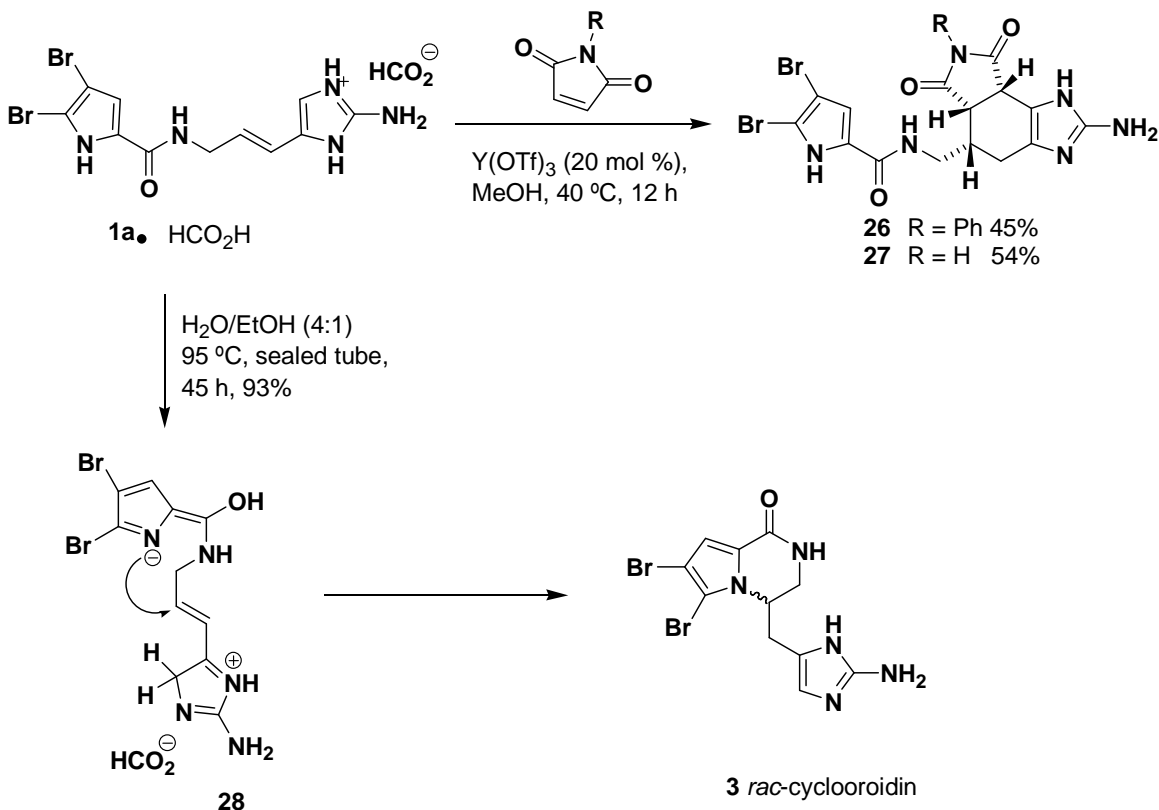
cyclooroidin from the major marine metabolite oroidin (**1a**) may be a related but enzyme-assisted process.



Scheme 1.5

It should be noted that Lindel and co-workers were actually exploring [4 + 2] cycloaddition reactions of oroidin itself and related 4(5)-alkenylimidazoles bearing an 2-amino substituent rather than attempting a total synthesis of cyclooroidin.²⁷ In this study they found natural product oroidin itself starts to react with *N*-phenylmaleimide and maleimide at room temperature. Addition of $\text{Y}(\text{OTf})_3$ (20 mol%) led to acceleration of the Diels-Alder reactions of oroidin, providing the *endo* cycloadducts **26** and **27** in good yields (Scheme 1.6).²⁸ During this study they attempted a [4+2]-dimerization of oroidin (**1a**) and they noticed a new product was forming on heating **1a**. HCO_2H (in the absence of maleimides) above 65 °C in protic solvents. They discovered that the double bond had disappeared, whereas the product had similar molecular formula as oroidin. They concluded that instead of a Diels-Alder reaction, a cyclization had taken place affording *rac*-cyclooroidin formate **3** in almost quantitative yield. They proposed an azafulvene tautomer **28** as an intermediate which reacts via the pyrrole nitrogen. It is of note, that the enolic oxygen does not engage in cyclization to give an oxazole, a pathway which has recently been described by Al-Mourabit^{43b} and by our lab.²⁹

Although we would note in the case of Al-Mourabit report they describe the use of slightly different conditions ($\text{CH}_3\text{SO}_3\text{H}$, $80\text{ }^\circ\text{C}$, 3h) and thus the products obtained may simply reflect kinetic versus thermodynamic preferences in the cyclization.

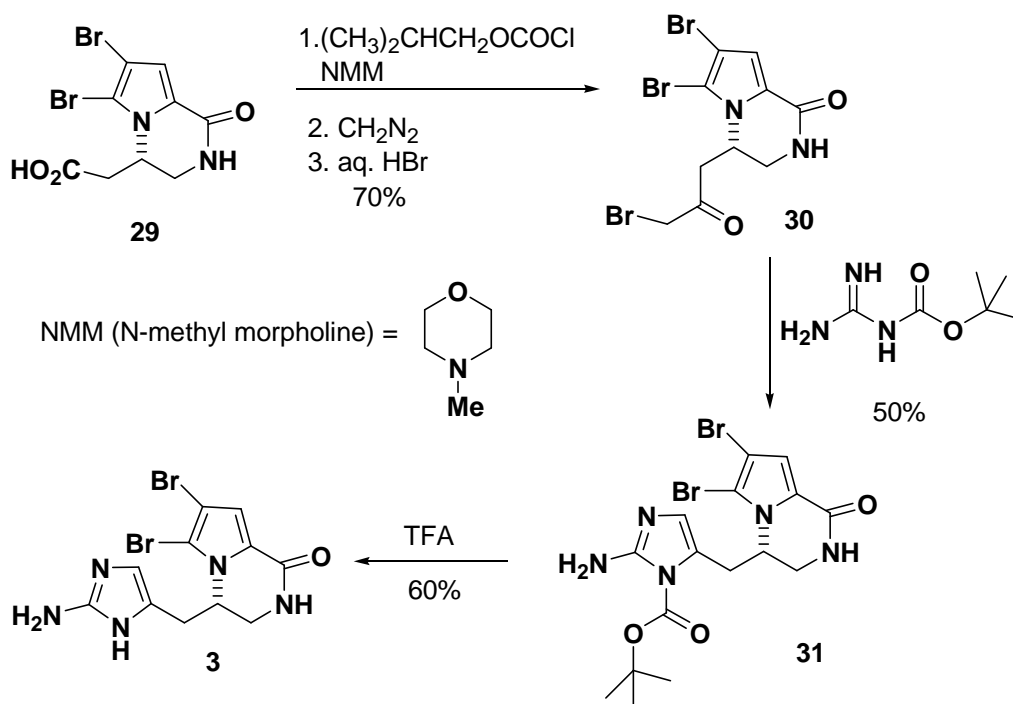


Scheme 1.6

1.6 (S)-cyclooroidin by Pelloux-Léon

In 2006 Pelloux-Léon et al.³² (Scheme 1.7) first synthesized non-racemic (S)-(-)-cyclooroidin (**3**)¹⁴ taking advantage of their earlier synthesis of (S)-(-)-longamide B.³⁰ Starting from enantiopure sample of (S)-(-)-longamide B (**29**), the carboxylic acid was transformed into a mixed anhydride with isobutyl chloroformate,³¹ which was then treated with CH_2N_2 and aqueous HBr ³¹ to furnish optically active bromoketone **30**. The α -bromoketone was converted to 15-N-Boc protected

cyclooroidin **31**. Finally enantiopure cyclooroidin was obtained by adding trifluoroacetic acid to (**3**).



Scheme 1.7

The CD spectra of synthetic cyclooroidin (**3**) and of (*S*)-longamide B methyl ester were compared and they observed no difference in the positions of the positive and negative Cotton effects. Their results therefore suggest that these two alkaloids have the same stereochemistry at C9.

Finally, they measured specific rotation. In the first attempt, they mimicked the conditions used for the natural product.¹⁴ In that regard, their synthetic sample exhibited the same value as the one described for natural cyclooroidin (-12.5 (c 0.02, MeOH)). The second measurement was performed using a more concentrated solution ($[\alpha]^{25}_{\text{D}} - 33.0$ (c 1, MeOH)). These experiments confirmed that natural cyclooroidin has an (*S*)-configuration at C-9 as it was postulated by Taglialatela-Scafati and co-workers.¹⁴

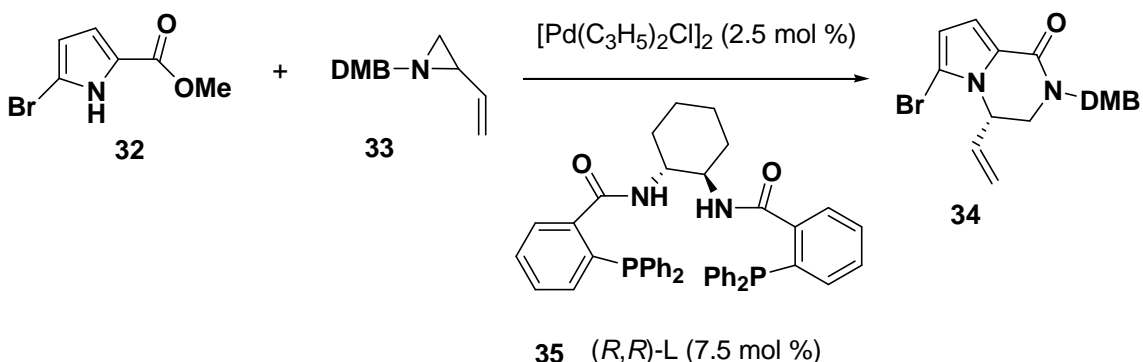
1.7 (S)-cyclooroidin by Trost

Recently Trost et al.³³ (Scheme 1.8) have developed an enantioselective, Pd-catalyzed alkylation-annulation reaction to construct a pyrrolo-piperazinone from a 5-bromopyrrole-2-carboxylate ester and a vinyl aziridine. Application of this protocol leads to the concise asymmetric synthesis of longamide B in five steps from **32** and **33**, while cyclooroidin can be synthesized in four additional steps from longamide B based on the work of Papeo et al.¹⁸ and Pelloux-Léon et al described in (scheme 1.7).³²

They envisioned that by using a 5-bromopyrrole-2-carboxylate derivative, the nitrogen on the pyrrole would behave as a good nucleophile to open the aziridine ring regioselectively during the AAA (asymmetric allylic alkylation), while the ester group on the pyrrole would act as a nitrogen acceptor to form a 6-membered lactam. Thus, the pyrrolo-piperazinone skeleton can be formed in single step.

Initially they examined the reaction between methyl 5-bromopyrrole-2-carboxylate **32** and vinyl aziridine **33**³⁴ (Table 1.1) in the presence of 2.5 mol % of $[\text{Pd}(\text{C}_3\text{H}_5)\text{Cl}]_2$ and 7.5 mol % of (*R,R*)-L. It had been shown previously that enantioselectivity in the Pd-catalyzed cycloadditions of isocyanates to vinyl aziridines was enhanced greatly by the addition of catalytic amount of HOAc.³⁵ By using 10 mol % of HOAc they obtained the desired pyrrolo-piperazinone **34** in 71% ee but very low yield due to the decomposition of starting materials. Addition of 50 mol % of Cs_2CO_3 gave much higher ee (89%) but only slightly improved the

yield (41%). However, they found that without any additives, the annulation product **34** was obtained in 72% yield and 95% ee.



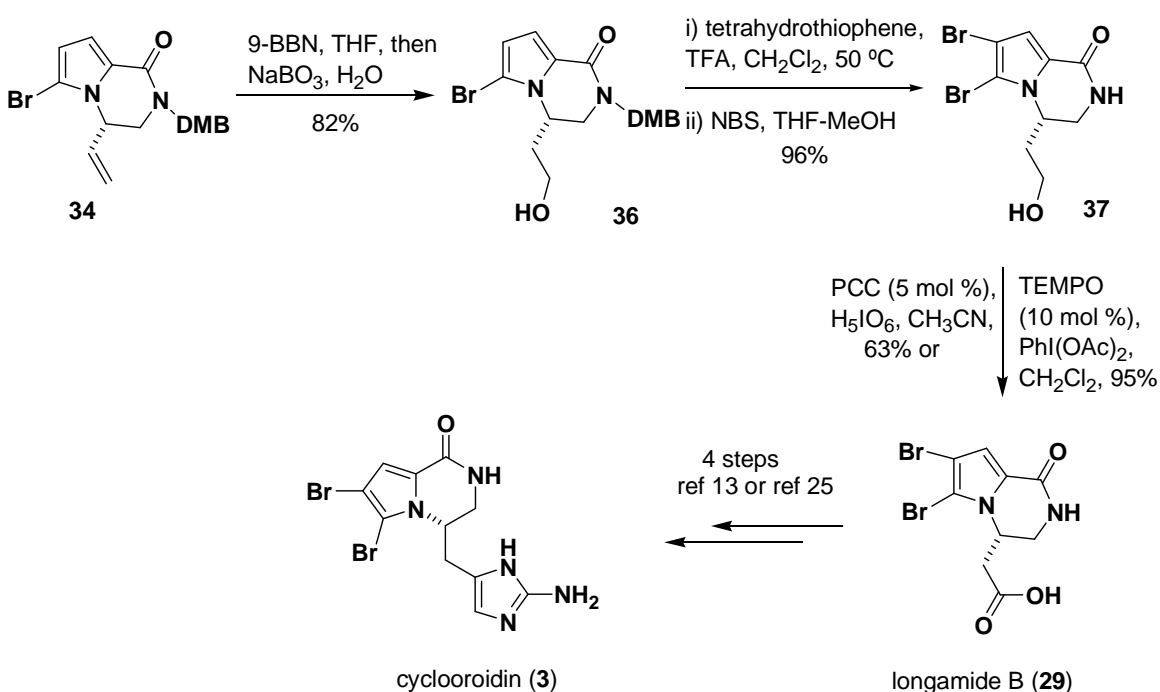
Scheme 1.8

Table 1.1 Effect of additives in Pd-catalyzed asymmetric allylic alkylation

entry	Additives	% yield	% ee
1	10 mol % HOAc	19	71
2	50 mol % Cs ₂ CO ₃	41	89
3	None	72	95

As depicted in (Scheme 1.9) primary alcohol **36** was obtained in high yield and excellent regioselectivity by hydroboration of **34** with 9-BBN followed by oxidation with sodium perborate. The DMB group was cleaved by treating **36** with 5 equiv of tetrahydrothiophene in TFA/DCM (1:1). The deprotected amide **30** was contaminated with its protodebrominated counterpart product, however they obtained dibromo alcohol **37** in 96% yield over two steps by NBS-mediated bromination of **36**. They screened conditions for oxidizing primary alcohol **30** to the carboxylic acid, and determined the PCC-catalyzed oxidation with H₅IO₆³⁶ gave (*S*)-(-)-longamide B (**29**) in 63% (brsm 90%) yield. A more effective

oxidation was found later in an attempt to oxidize **37** to the corresponding aldehyde. They found that treatment of **37** with 10 mol% of TEMPO and 2.5 equiv of $\text{PhI}(\text{OAc})_2$ in CH_2Cl_2 ³⁷ produced carboxylic acid **29** instead of the aldehyde in 95% yield. The total synthesis of (*S*)-(-)-longamide B also confirmed the absolute configuration of the Pd-catalyzed asymmetric annulation reaction. As indicated above cyclooridin (**3**) can be synthesized in four steps from longamide B based on the work of Papeo et al.¹⁸ and Pelloux-Léon et al.³²



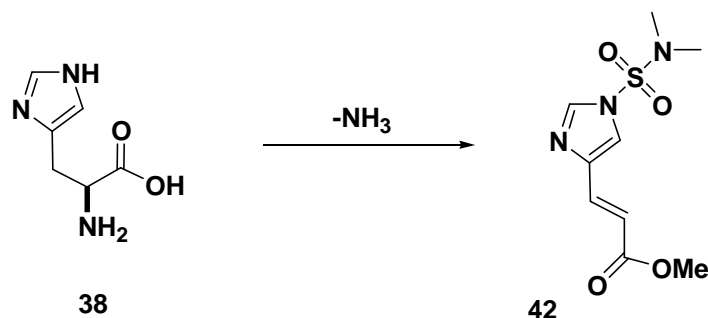
Scheme 1.9

In the next chapter, we will delineate our own asymmetric total synthesis of cyclooridin, which adopts a strategically different approach to the previously reported syntheses. In the chemistry to be outline below, we will use histidine as both our source of chirality and also to provide one of the heterocyclic rings.

CHAPTER 2
RESULTS AND DISCUSSION

2.1 Initial efforts

Our group uses large amounts of urocanic acid as a starting material, and while this is commercially available, it is relatively expensive and its supply was sometimes unreliable. Driven by these concerns, we set out to develop a more cost effective approach to urocanic acid (Scheme 2.1)³⁸ and it was in the course of this study that an approach to the cyclooroidin problem evolved.



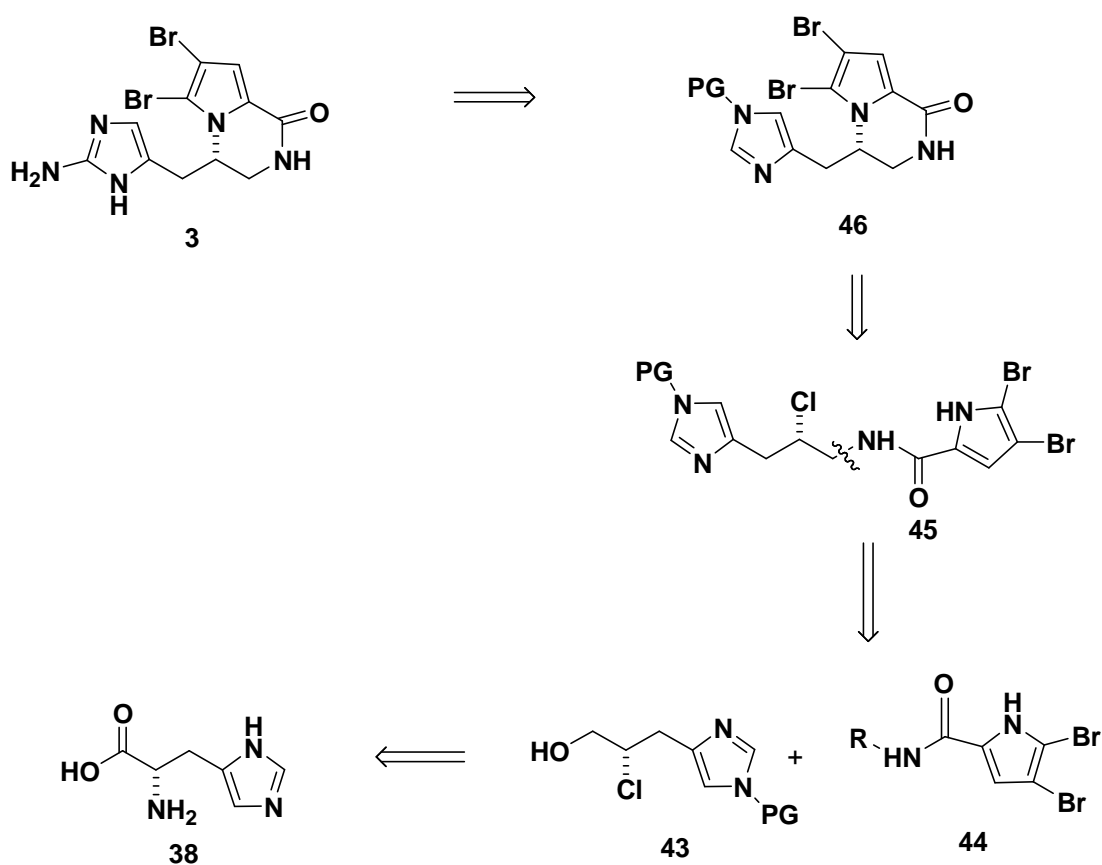
Scheme 2.1

2.2 Preparation of the methyl ester of Urocanic acid.

When the structures of histidine and urocanic acid are compared, the net loss of ammonia from **38** provides urocanic acid. In practice, such a direct conversion is not straightforward and therefore, we converted the amino group into a better leaving group through the incorporation of a chloro moiety.

Diazotization was carried out on L-histidine **38** using a mixture of concentrated hydrochloric acid and sodium nitrate at 0-5 °C according to the report of

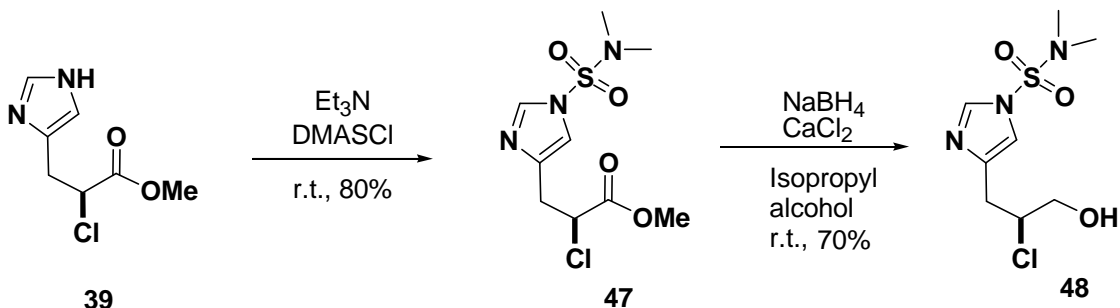
displacement of the chloride with the pyrrole nitrogen would give the basic skeleton of cyclooroidin (Scheme 2.3). In a forward sense, the key steps involve a Mitsunobu reaction between alcohol **43** and a novel cyclic pyrrole imide **49** followed by basecatalyzed cyclization to form the core skeleton of the molecule. Once the basic core of the molecule is assembled, the next goal is to introduce two bromines into the pyrrole of the molecule **46**. Finally an azide needs to be introduced at the C-2 position of the imidazole of the molecule **46**. Synthetic cyclooroidin can be obtained by reducing the azide **46** to amine first and then ultimately removing off the protecting group.



Scheme 2.3

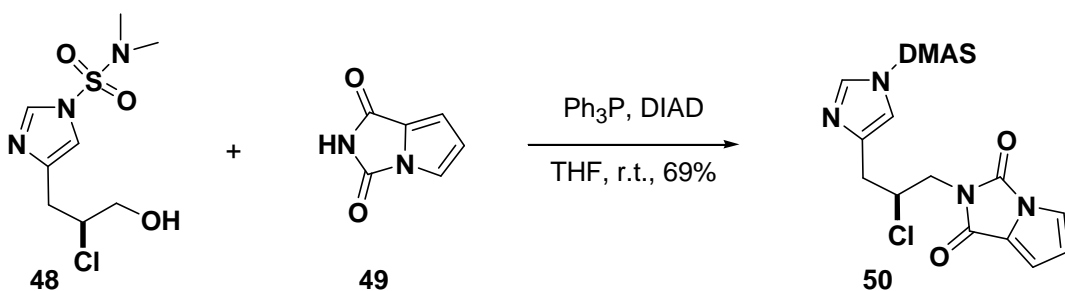
2.4 Efforts towards synthesis of cyclooroidin

The α -chloro ester **39** was protected with dimethylaminosulfonyl chloride at room temperature to obtain **47** which was then reduced to alcohol **48** (Scheme 2.4) using sodium borohydride, calcium chloride and isopropyl alcohol as solvent (at room temperature).⁴⁰

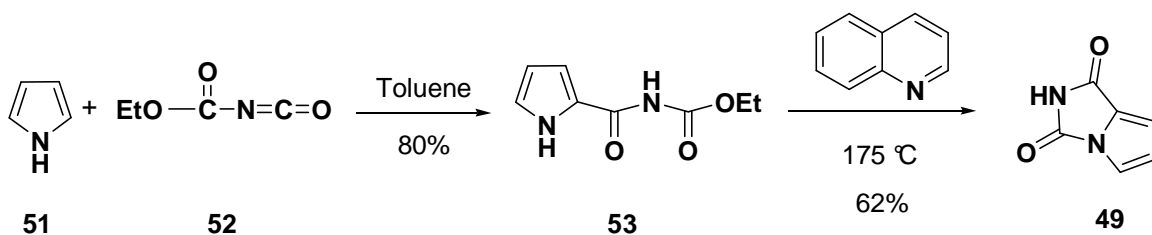


Scheme 2.4

After screening several nucleophiles or pronucleophiles including *N*-(1*H*-pyrrole-2-carbonyl)-benzenesulfonamide,^{41a} 4-methyl-*N*-(1*H*-pyrrole-2-carbonyl)benzenesulfonamide,^{41b} diphenyl phosphoryl azide,^{41b} it was found that the alcohol **48** underwent Mitsunobu reaction⁴¹ with imide **49** to obtain **50** (Scheme 2.5). Imide **49**⁴² was obtained by reacting ethoxycarbonyl isocyanate (**52**)⁴² with pyrrole (**51**) to provide *N*-ethoxycarbonylpyrrole-2-carboxamide (**53**) (Scheme 2.6).⁴² Cyclization occurred on heating a quinoline solution of **53** at 175 °C affording the known compound **49**.⁴²

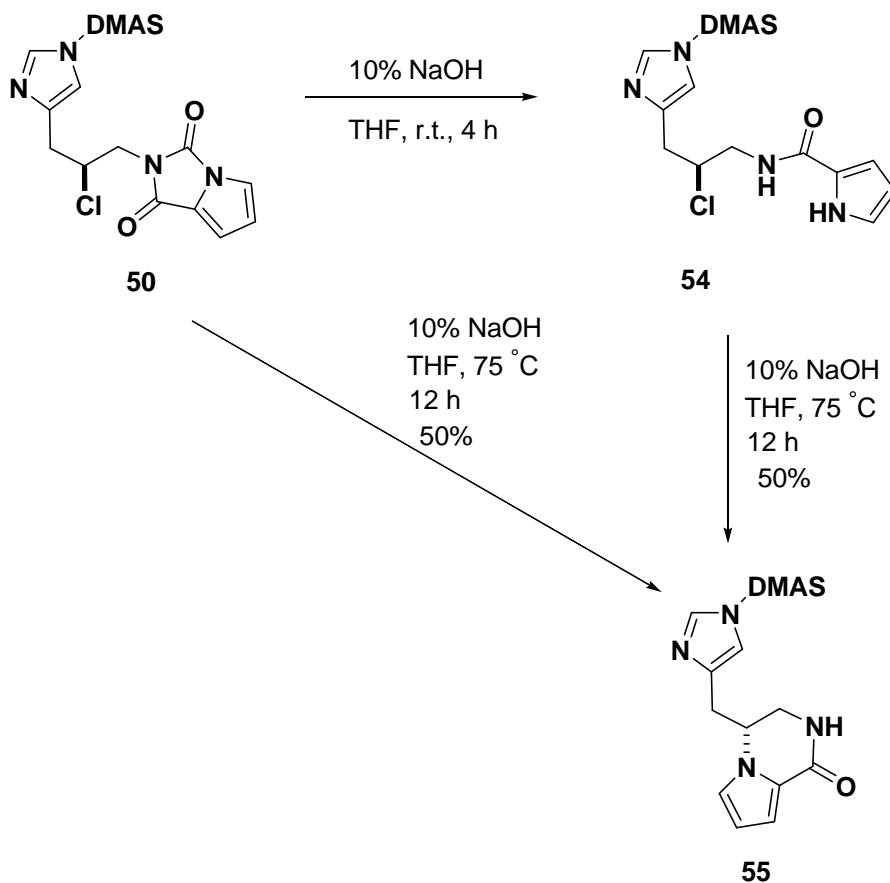


Scheme 2.5



Scheme 2.6

Stirring a solution **50** in THF and 10% NaOH solution for about 4 h, led to hydrolytic opening of hydantoin moiety and affording compound **54** (Scheme 2.7).⁴²



Scheme 2.7

When compound **54** (contaminated with triphenylphosphine oxide) was heated for an additional period time under the same reaction conditions it underwent

cyclization to form what we initially believed was compound **55**. We also found that **50** undergoes tandem hydrolysis and cyclization by simply running the reaction for 12 h at 75 °C. While we initially thought that we had constructed the basic core ring system of cyclooroidin **55**, there were some troubling discrepancies between the spectroscopic data of the isolated natural product (Table 2.1) and with our synthetically prepared compound **55**. It was found that although it was largely similar, there were inconsistencies in the ^{13}C NMR spectrum, in particular a signal due to one of the aliphatic carbons corresponding to C9 in cyclooroidin did not match well with the natural product. The particular signal in question was 19.0 ppm further downfield than anticipated, suggesting that this carbon may be bonded to oxygen rather than with nitrogen i.e., an oxazoline had been formed rather than a pyrazine. Comparison of the NMR data of similar compounds from the literature (Table 2.1) suggested that instead of forming a C-N bond, the cyclization has occurred through formation of a C-O bond.^{43a-b, 44}

Table 2.1 Comparison of ^{13}C NMR studies of C9 for cyclooroidin and several related oxazolines.

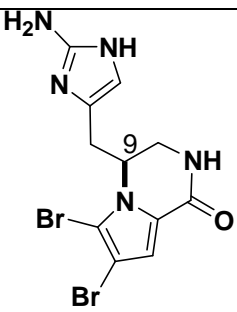
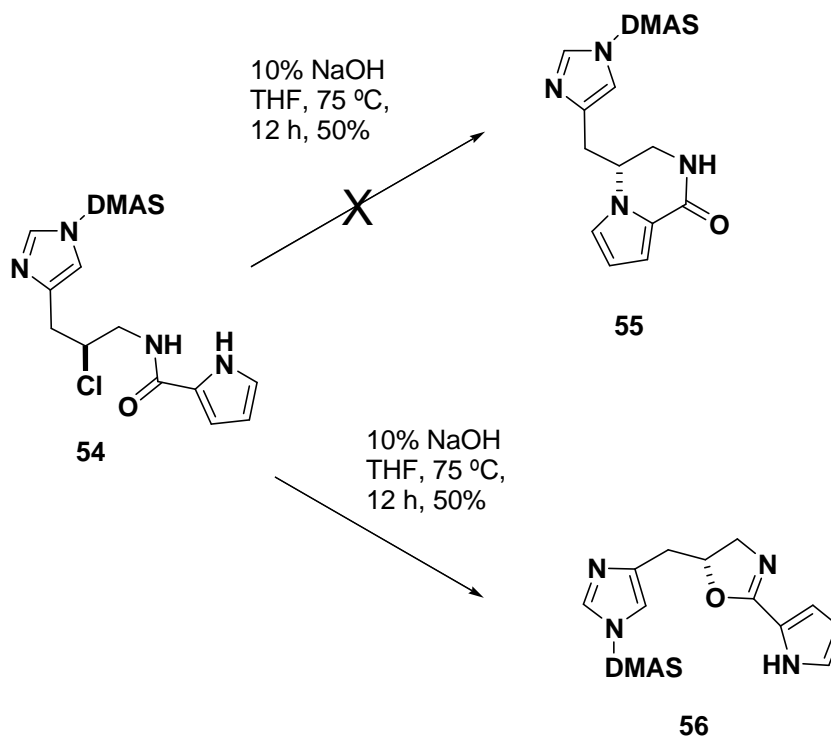
Compound	δ_{C9}
 <p style="text-align: center;">3</p>	57.9

Table 2.1-continued

<p>57</p>	89.0
<p>55</p>	77.6
<p>Nagelamide R</p>	79.5
<p>Nagelamide T</p>	80.2

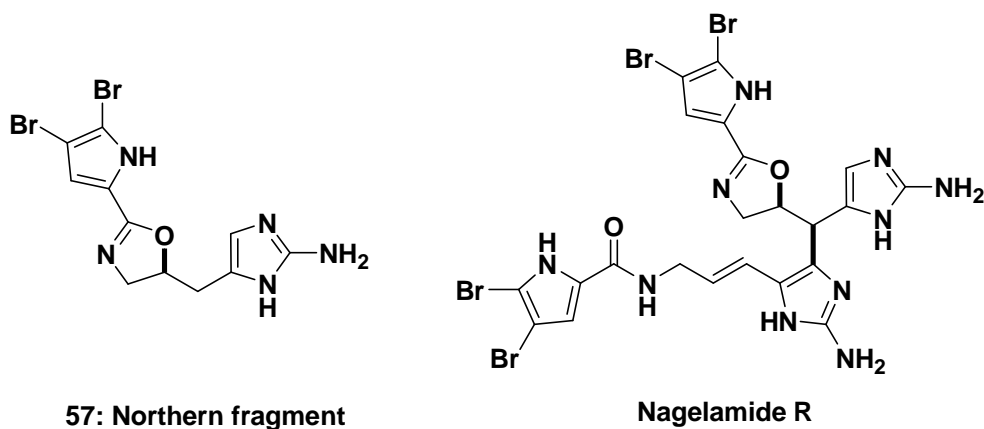
So when a THF solution of compound **54** was heated under basic condition (10% NaOH solution) it underwent cyclization through the amide oxygen instead

of pyrrole nitrogen to give an oxazole ring, thus resulting in the formation of compound **56** (Scheme 2.8).

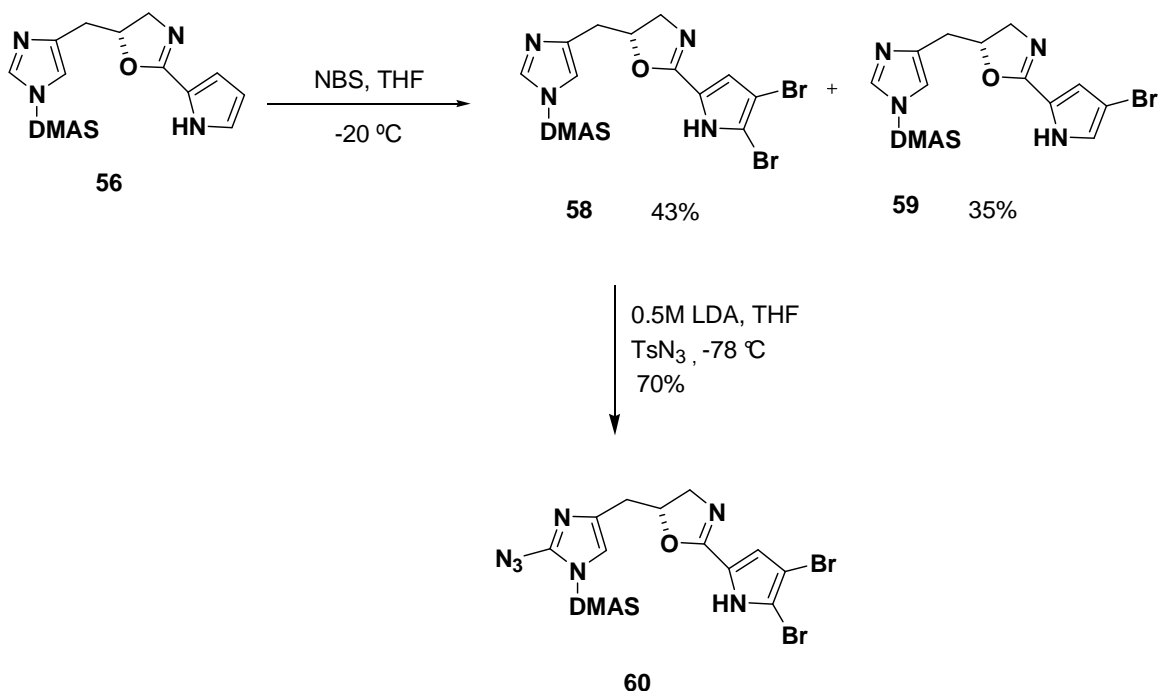


Scheme 2.8

Compound **56** has the appropriate structural fragments as the northern sector of the natural product nagelamide R.⁴⁴ So we decided to continue our investigation to establish whether **56** could be further elaborated to **57**, which would be useful in the pursuit of a total synthesis of nagelamide R.

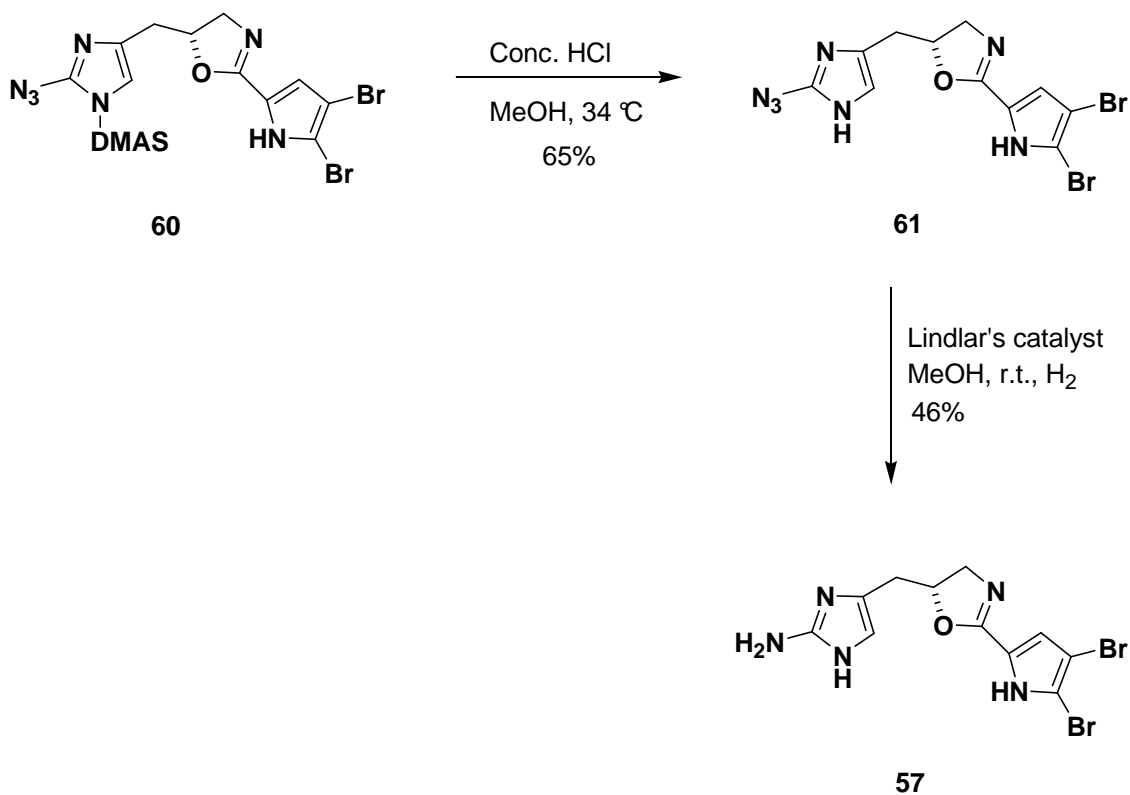


The cyclic compound **56** was subjected to bromination with NBS at $-20\text{ }^{\circ}\text{C}$ leading to the formation of compound **58** in modest yield.⁴⁵ The yield of the bromination was relatively low due to competitive formation of mono brominated compound **59** in 35%. Subsequently the azide moiety was introduced to the brominated compound **58** by generating the anion at C-2 position with 4.1 equiv. of LDA and adding TsN_3 while maintaining the reaction temperature throughout at $-78\text{ }^{\circ}\text{C}$, thus leading to formation of compound **60** (Scheme 2.9).⁴⁶



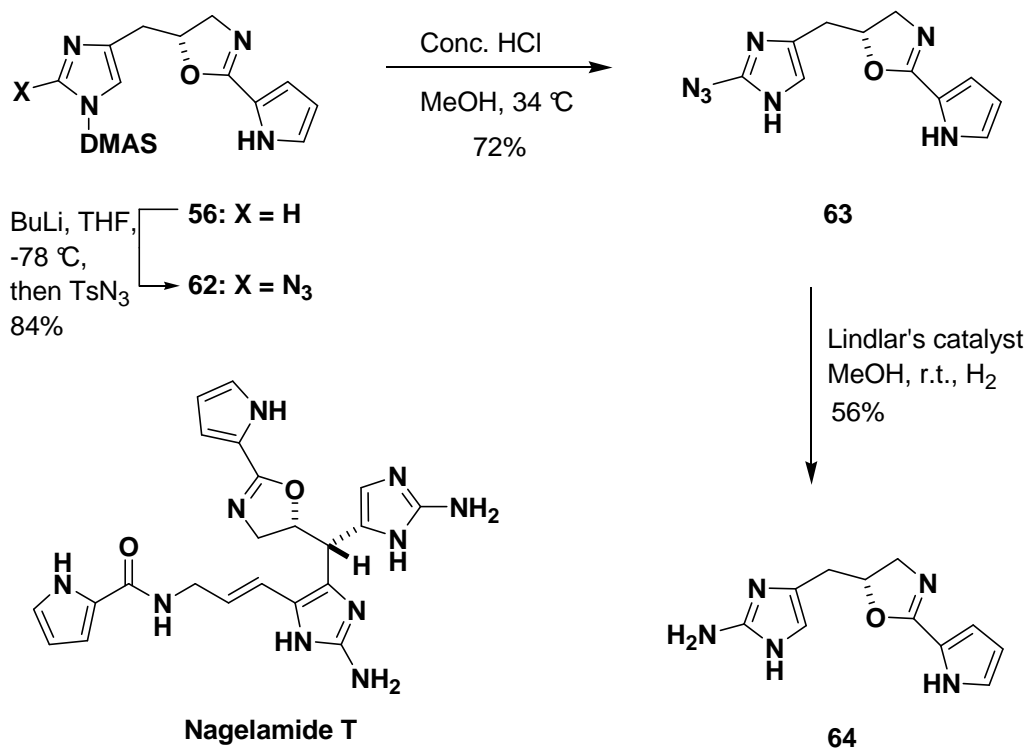
Scheme 2.9

The dimethylsulfamoyl protecting group was removed by adding conc. HCl to the compound **60** dissolved in methanol and heating the resulting mixture to $34\text{ }^{\circ}\text{C}$ to provide compound **61**.⁴⁷ Finally the azide was reduced to amine **61** in the of Lindlar's catalyst and hydrogen atmosphere to give **62** (Scheme 2.10).⁴⁷

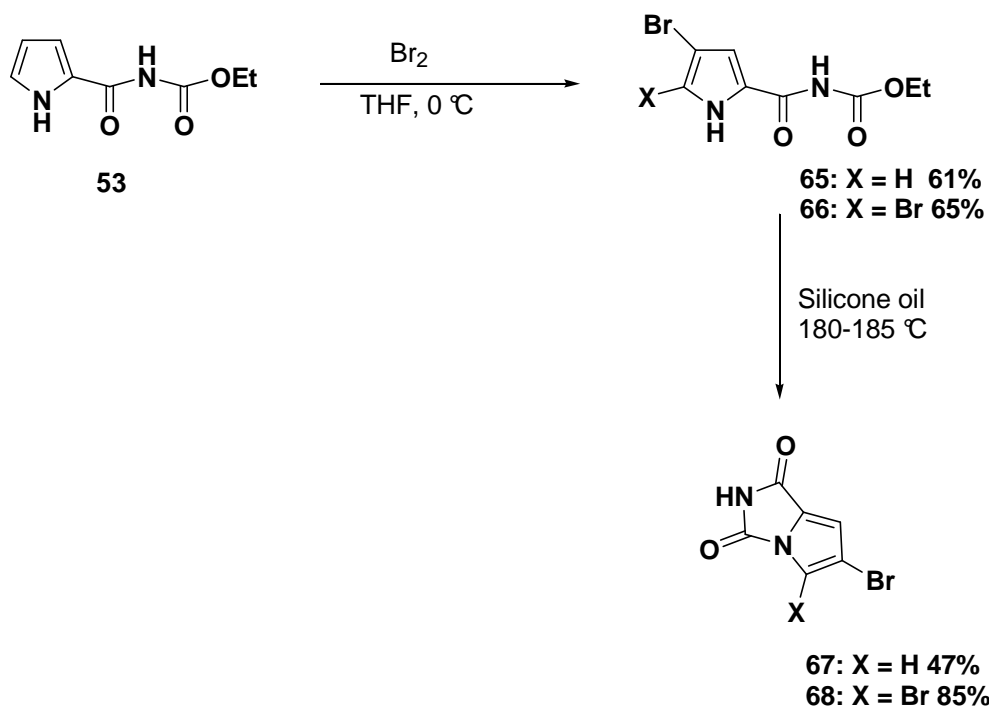


Scheme 2.10

We would note that initial studies on the final few steps were conducted on the non-brominated congener to determine viability of the deprotective and reduction steps. This sequence of reactions provides the oxazoline portion of the nagelamide T, although at the time these reactions were conducted, nagelamide T had not been described. The azidation in this case can be performed with *n*-BuLi rather than LDA, this chemistry is depicted in (Scheme 2.11).



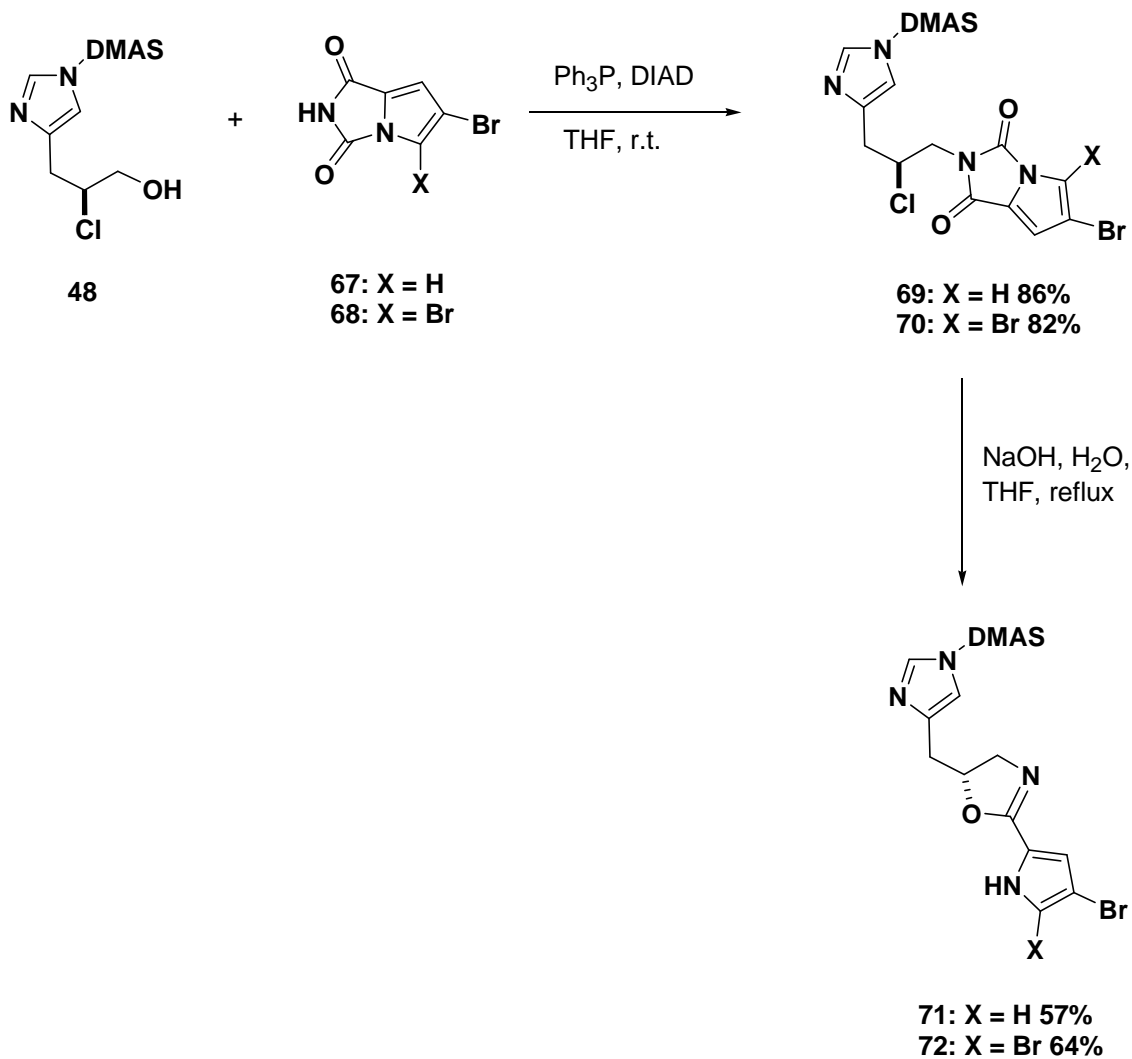
Scheme 2.11



Scheme 2.12

We were somewhat disappointed with the efficiency of the bromination step in the earlier sequence (Scheme 2.9, **56**→**58**) due to the formation of the monobromo product. We reasoned that if the bromo substituents were introduced prior to the Mitsunobu reaction, this issue would be less of a problem. Accordingly we have examined the use of other pyrrole hydantoin derivatives, specifically brominated derivatives (**67** and **68**), in the Mitsunobu sequence as mono- and dibrominated pyrrole substituents are widely distributed among the oroidin alkaloids. It is well known that the corresponding pyrrole trichloro ketones can be brominated in a controlled fashion and thus it was reasoned that imide **53** should behave similarly. Gratifyingly, treatment of **53** with 1 or 2 equiv of Br₂ provided the corresponding mono- and dibrominated derivatives. Monobromopyrrole **65** was mixed with silicone oil in a round bottom flask which was evacuated (40 mm, Hg) and heated to 180-185 °C.⁴⁸ using a procedure adapted from Spoering's dissertation.⁴⁸ After 4 h, the reaction was cooled and then the residue was crushed. Hexane was added and the solids were collected by filtration and washed repeatedly with hexane. The procedure was repeated and the resulting product was purified by chromatography, providing **67** in 47% yield. The dibrominated hydantoin **68** was also prepared in a similar fashion as described above. We have found that both the mono and the dibrominated derivatives **67** and **68** participate in the substitution reaction (Scheme 2.12).^{27,47} These alkylated hydantoin derivatives **69** and **70** were subjected to hydrolysis under basic conditions, and in a similar fashion to **54** underwent concomitant hydrolysis, cyclization to provide the corresponding oxazoles **71** and **72** (Scheme

2.13). In the case of the dibrominated derivative **72**, this provides a more efficient preparation of the oxazoline than bromination of the parent system **56**. It is challenging to unequivocally prove the location of the bromine atoms on the pyrroles, but fortunately both **69** and **72** were nice crystalline solids that provide good X-ray crystal structures.



Scheme 2.13

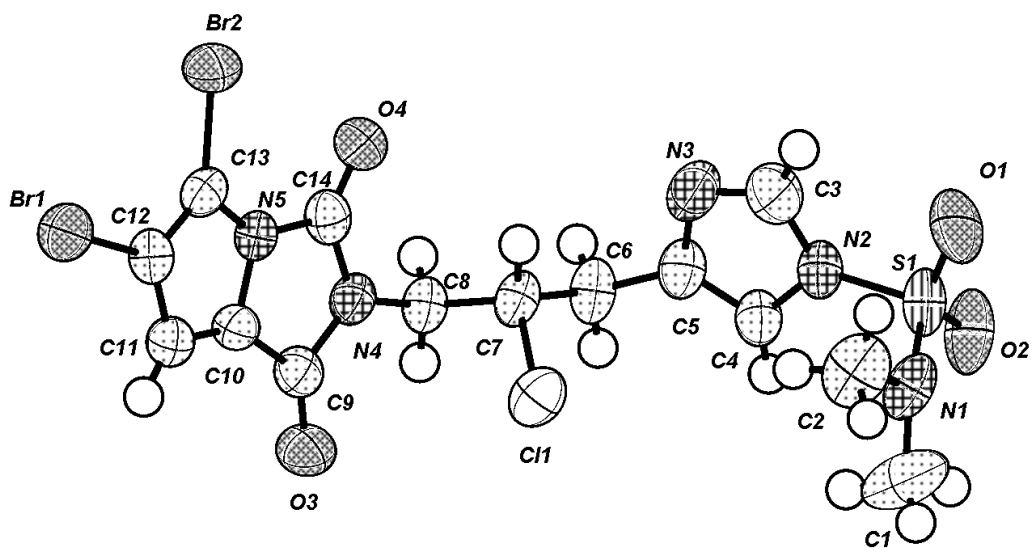


Figure 2.1 X-ray Crystal structure of compound **70**

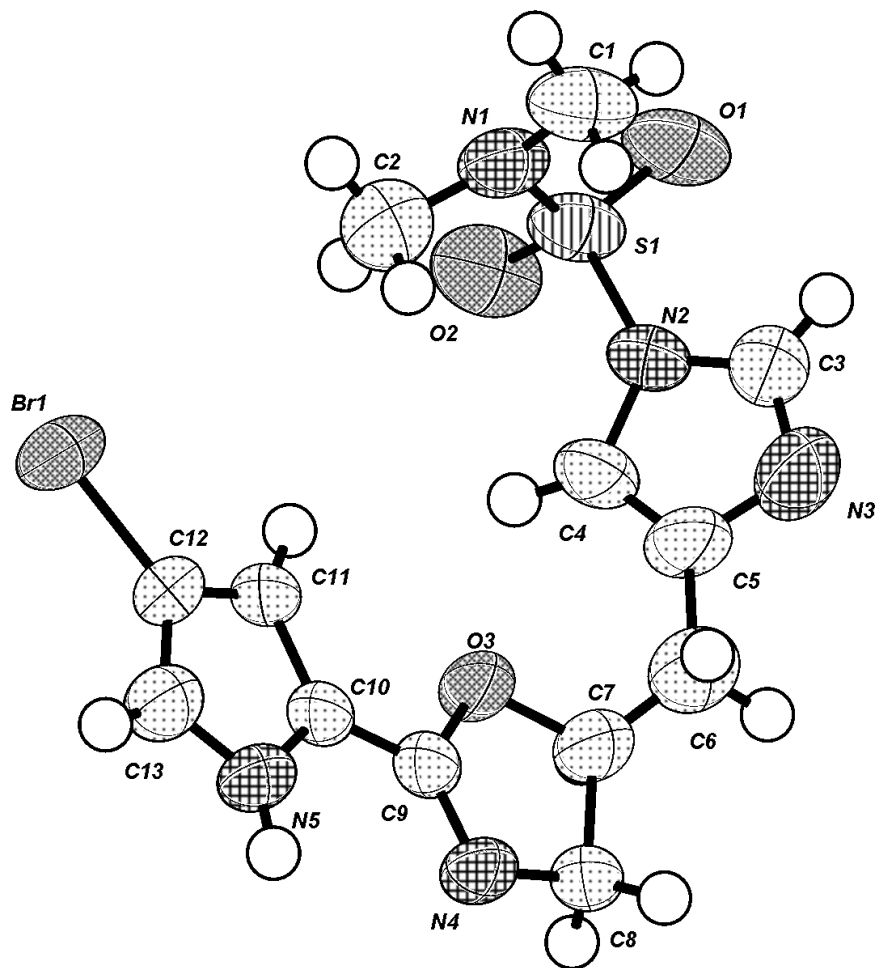
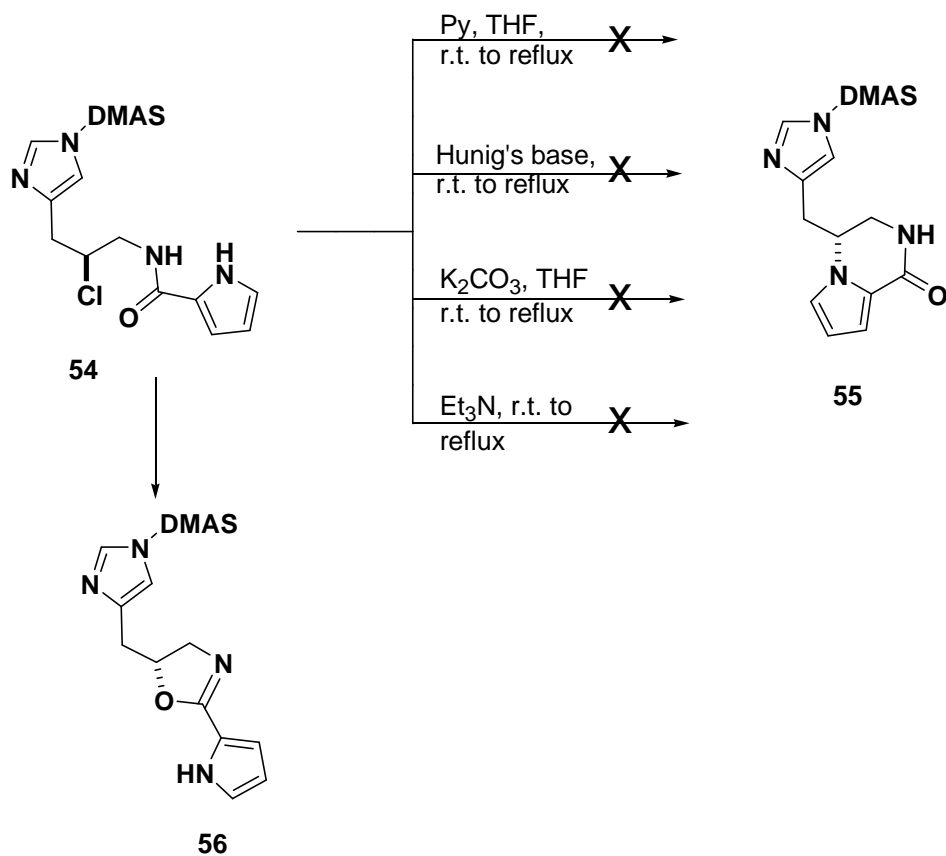


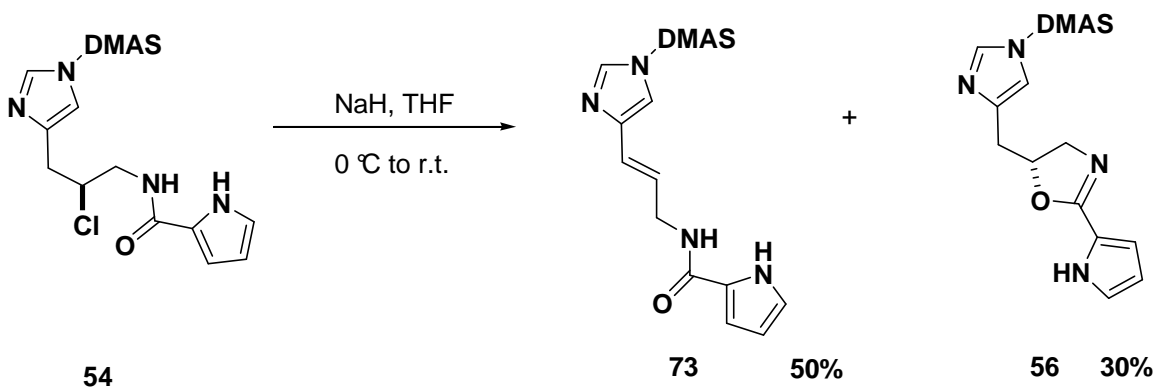
Figure 2.2 X-ray Crystal structure of compound **71**

In regards to advancing the total synthesis of cyclooroidin we attempted to determine conditions leading to the cyclization of compound **54** via the pyrrole nitrogen. A variety of different bases such as pyridine, Hunig's base, Et_3N , K_2CO_3 were evaluated however cyclization only occurred through the amide oxygen instead of pyrrole nitrogen (Scheme 2.14).



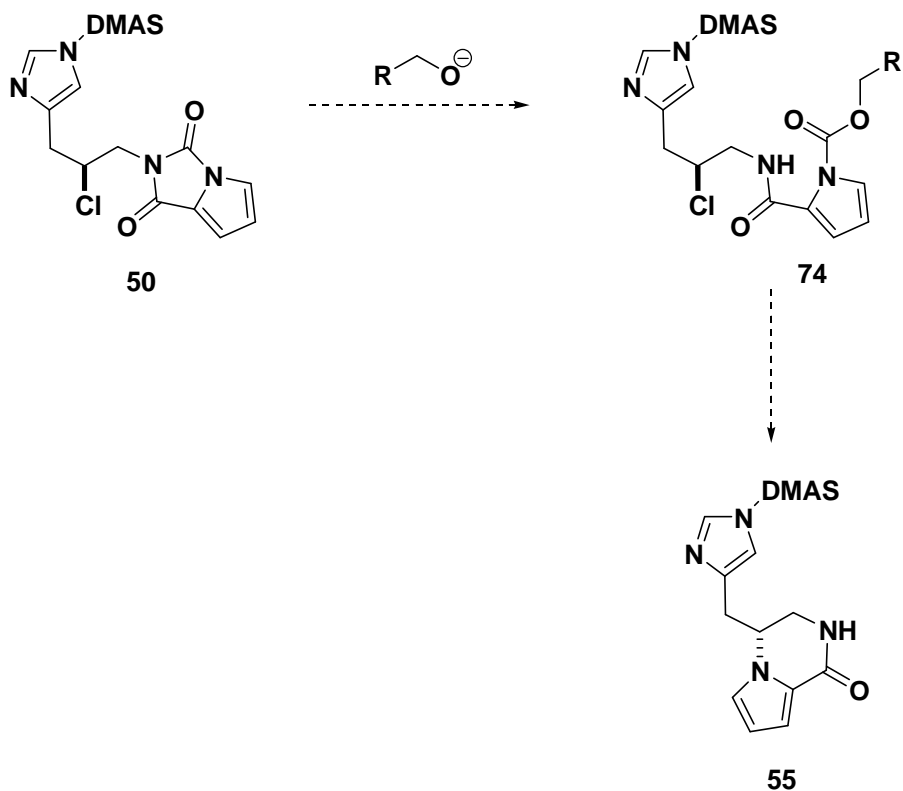
Scheme 2.14

Then we tried to cyclize compound **54** via the pyrrole nitrogen using a stronger base like NaH but a mixture of two compounds was obtained one of them being compound **56** and the other one was dehydrochlorinated compound **73** (Scheme 2.15).



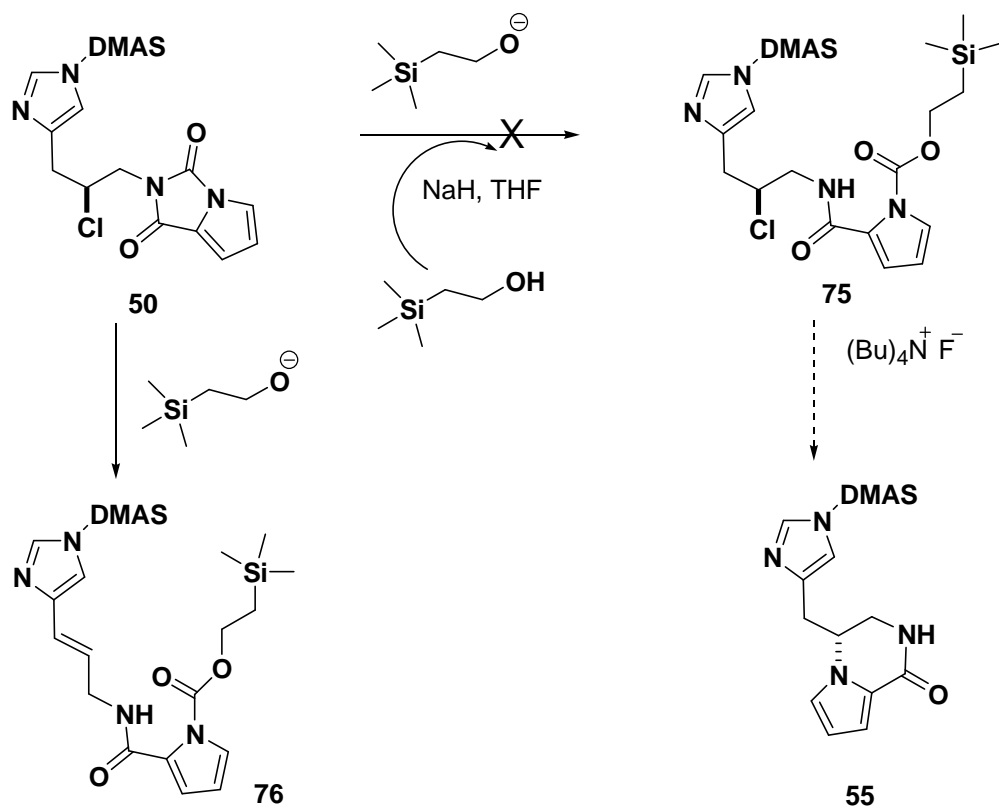
Scheme 2.15

Since our initial attempt to cyclize compound **54** directly via the pyrrole nitrogen had failed we considered using alternative nucleophiles to cleave the urea in compound **50**. This would result in the pyrrole nitrogen being protected; subsequent amide protection would then provide a differentially protected substrate. Addition of an appropriate reagent to the reaction mixture we can generate the pyrrole anion rather than the amide N-H (Scheme 2.16).⁴⁹



Scheme 2.16

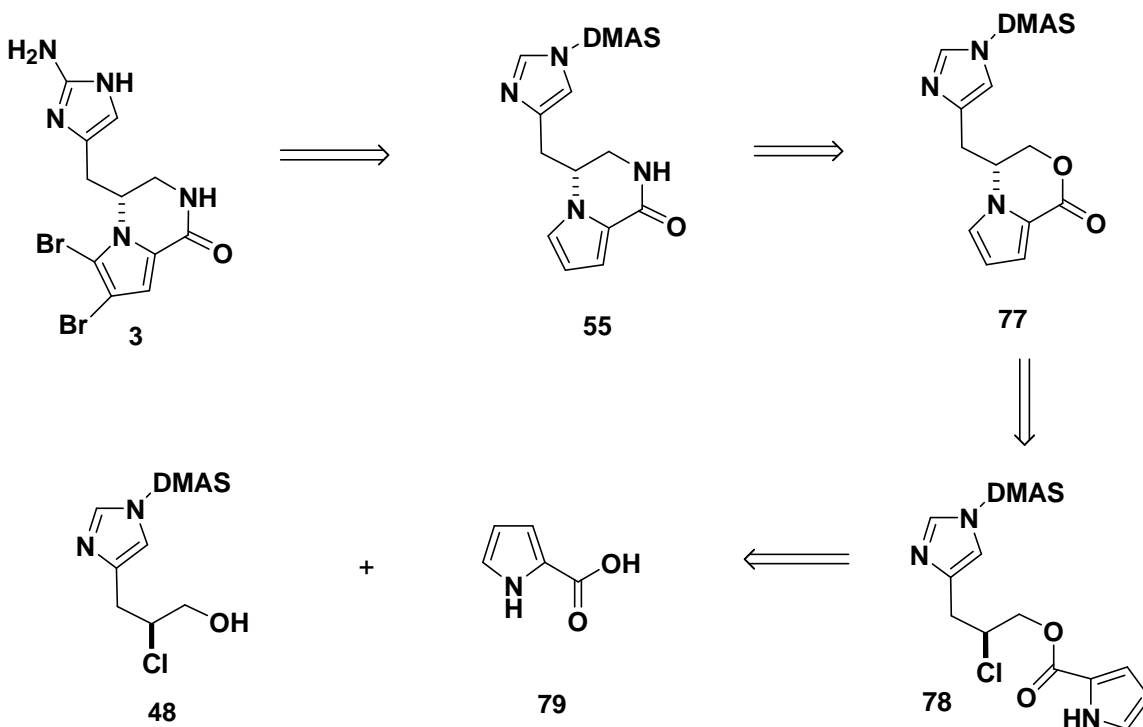
Unfortunately, attempts to execute this strategy using the anion generated from 2-trimethylsilyl alcohol were not successful. Instead of obtaining compound **75**, compound **77** was isolated which represents a dead end for synthesis of cyclooroidin (Scheme 2.17).



Scheme 2.17

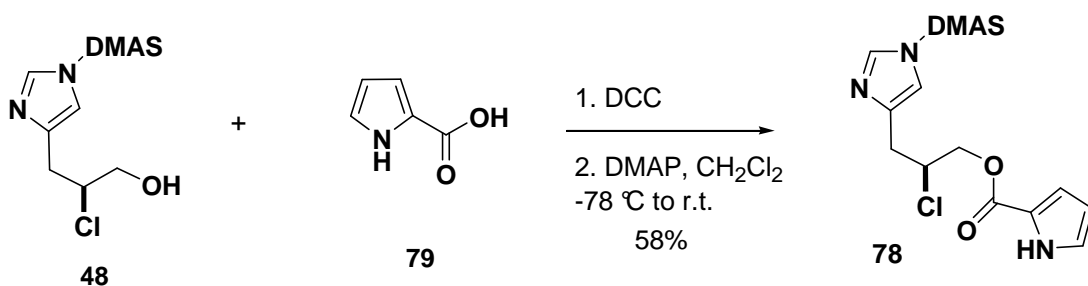
In order to address the incorrect cyclization manifold, the synthetic strategy was redesigned to avoid this pitfall and thus we developed the idea of replacing the amide with an ester. The advantage with oxygen atom as a linker is that there is no longer a problem of wrong cyclization through the amide oxygen instead of pyrrole nitrogen (Scheme 2.18). An alternative pathway involving protection of the amide nitrogen was considered, but this approach appeared to be lengthy.

2.5 Modified retrosynthetic analysis



Scheme 2.18

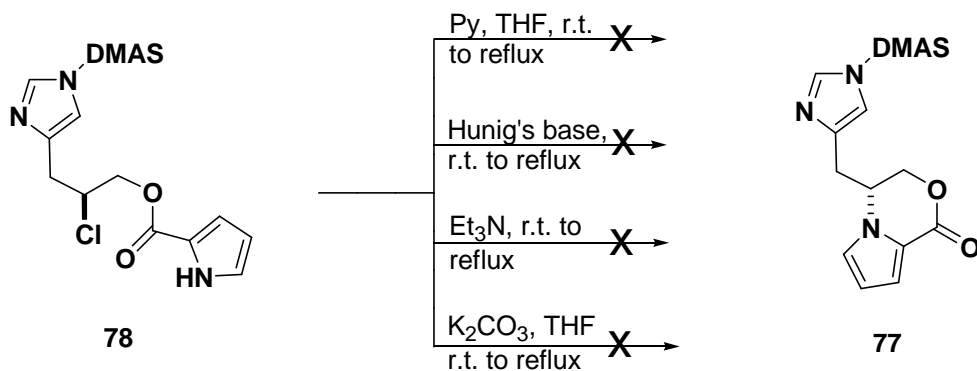
Accordingly, we esterified chloro alcohol **48** with pyrrole-2-carboxylic acid **79** in the presence of DCC,⁵⁰ obtaining ester **78** in 58% yield (Scheme 2.19).



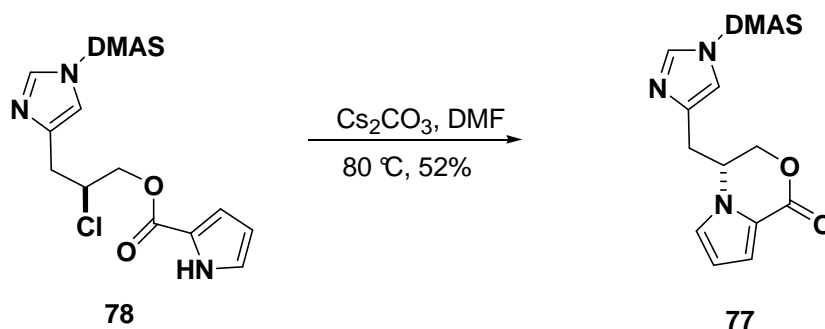
Scheme 2.19

Once we had compound **78** our next aim was to cyclize the molecule via pyrrole nitrogen. In this respect we tried mild conditions (Scheme 2.20) to remove the pyrrole N-H which in turn will cyclize intramolecularly in a S_N2 fashion. We already knew that strong bases like NaH should be avoided as they promote

dehydrochlorination. Initial studies with amine bases and K_2CO_3 were not successful. However, finally the desired cyclization was achieved by dissolving compound **78** in dry DMF and adding cesium carbonate⁵¹ to the reaction mixture and heating the reaction mixture to 80 °C for 12 h (Scheme 2.21).



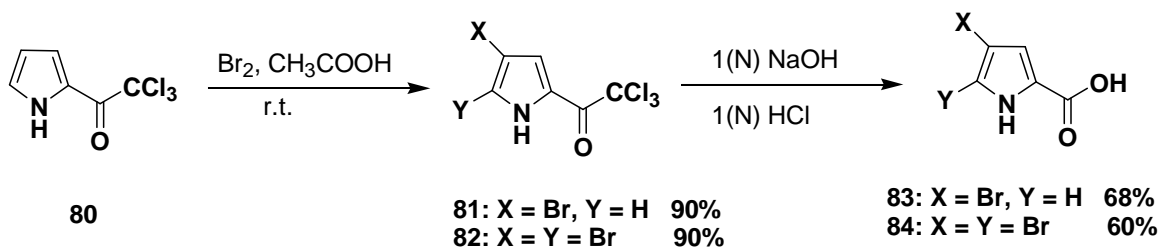
Scheme 2.20



Scheme 2.21

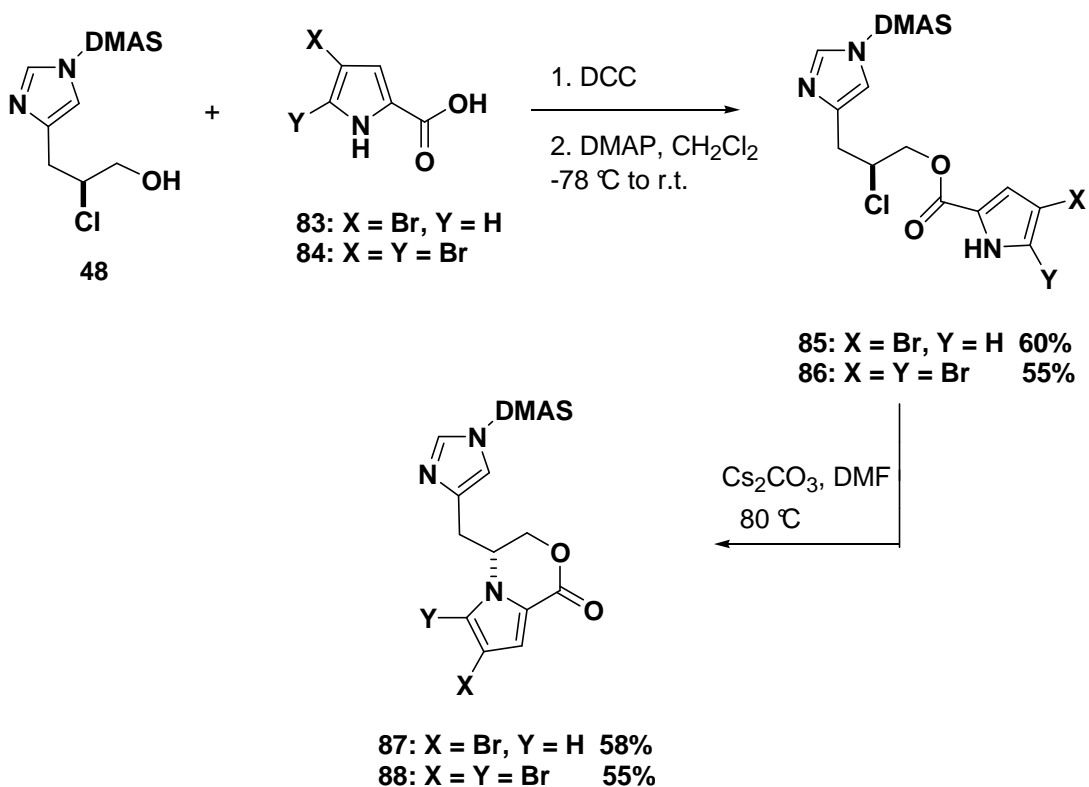
Once compound **77** was in hand our next goal was to introduce the bromine group into the pyrrole moiety as our target molecule cyclooroidin (**3**) has two bromine atom on the pyrrole moiety. In this regard we started with trichloroacetylpyrrole **80** and selectively performed either mono and dibromination on it and obtained known compounds **81** and **82**⁵² which were further transformed to mono **83** and dibrominated pyrrole carboxylic acid **84** by

hydrolysis with NaOH⁵² and finally neutralizing the solution with HCl (Scheme 2.22).



Scheme 2.22

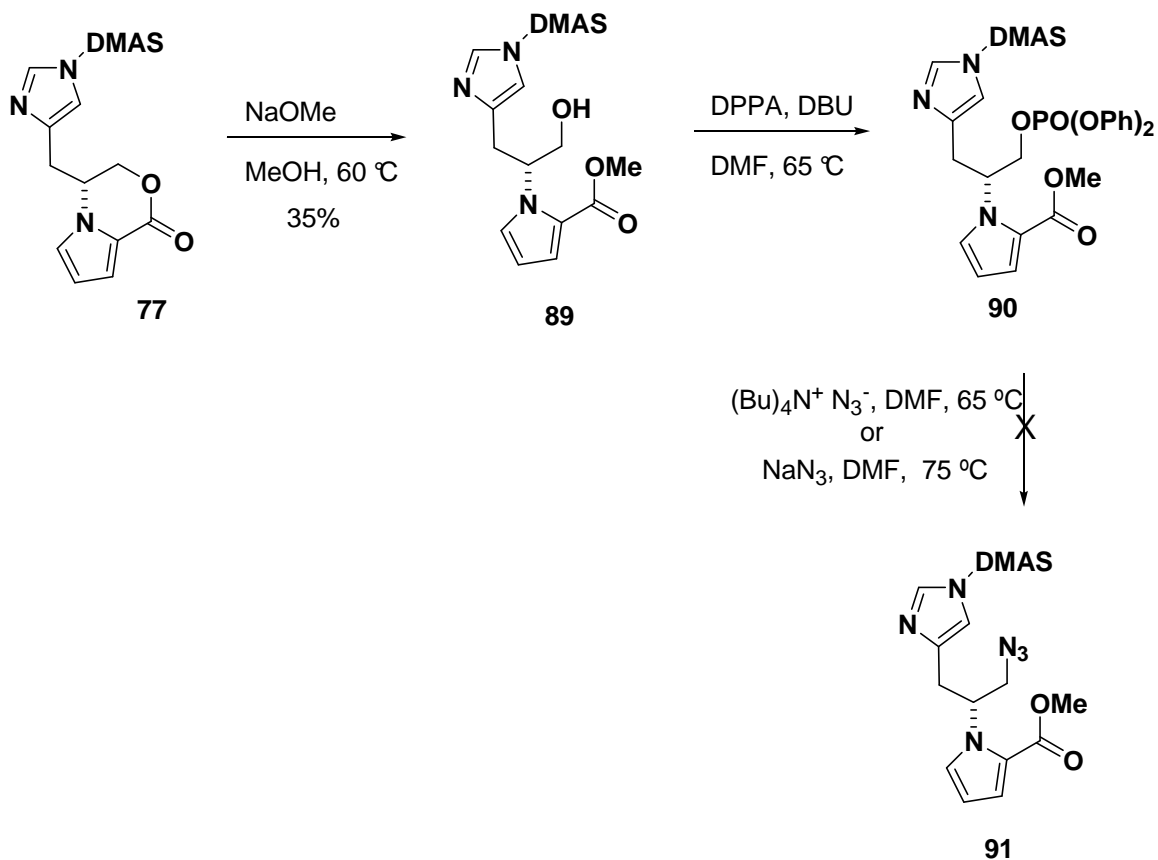
Once the mono- and dibromocarboxylic acids **83** and **84** were prepared they were coupled with chloro alcohol **48** in presence of DCC⁵⁰ at -78 °C to obtain the compound **85** and **86**. Subjection of both esters to cyclization by in DMF in the presence of cesium carbonate and then heating it to 80 °C (Scheme 2.23) led to the formation of **87** and **88**.



Scheme 2.23

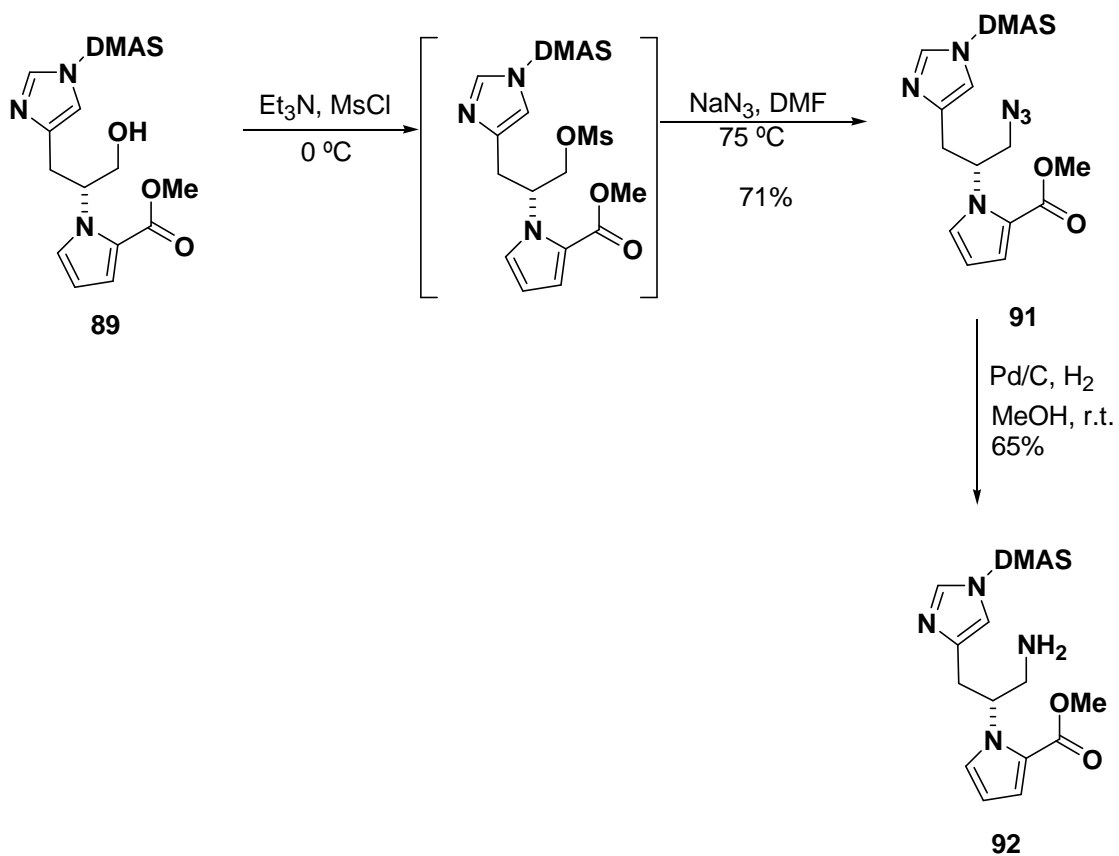
With the correct cyclized products in hand our next task was to replace the oxygen atom of the six membered lactone ring with a nitrogen atom. In order to achieve this transformation, it was decided to ring-open the lactone ring with sodium methoxide⁵³ to provide the hydroxy ester. Conversion of free alcohol to primary azide⁵⁴ and subsequent reduction of the azide to primary amine⁵⁵ should allow construction of the six membered ring.

The ring-opening of the lactone ring with sodium methoxide was achieved but the yield of the reaction was relatively low, providing the hydroxy ester **89** in 35% yield, with the remainder being recovered starting material (55%). The extent of conversion could not be improved suggesting that an equilibrium distribution has been reached between **89** and **77**. While this was not optimal, the two components can be separated and **77** can be resubjected to the ring-opening reaction. Compound **89** was subjected to DPPA and DBU in order to obtain the azide with the ultimate goal to convert this intermediate to the primary azide.⁵⁴ Interestingly the primary azide was not obtained rather only the phosphoryl intermediate **90** was isolated. Attempts to convert this phosphoryl intermediate to the primary azide with an external nucleophile like sodium azide and heating the reaction mixture to 75 °C were unsuccessful (Scheme 2.24).



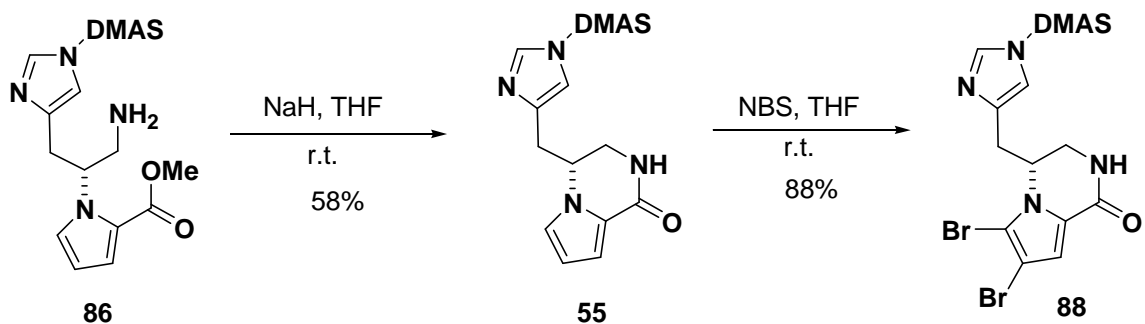
Scheme 2.24

Since this initial reaction sequence did not yield the primary azide, we decided to convert the primary alcohol to mesylate⁵⁶ and then displace the mesylate with azide ion to obtain the alkyl azide. Using this sequence the desired primary azide **91** was formed in 71% yield. The resulting azide was reduced to the primary amine⁵⁵ by using Pd/C under a hydrogen atmosphere at room temperature in 65% yield (Scheme 2.25).



Scheme 2.25

The primary amine **86** was cyclized by using NaH in THF at room temperature to obtain pyrazine **55** in 58% yield. Bromination of compound **55** using *N*-bromosuccinimide at room temperature provided dibrominated compound **93** in 88% yield (Scheme 2.26).³¹ Note here the improved efficiency compared to the oxazolines described earlier in (scheme 2.9).



Scheme 2.26

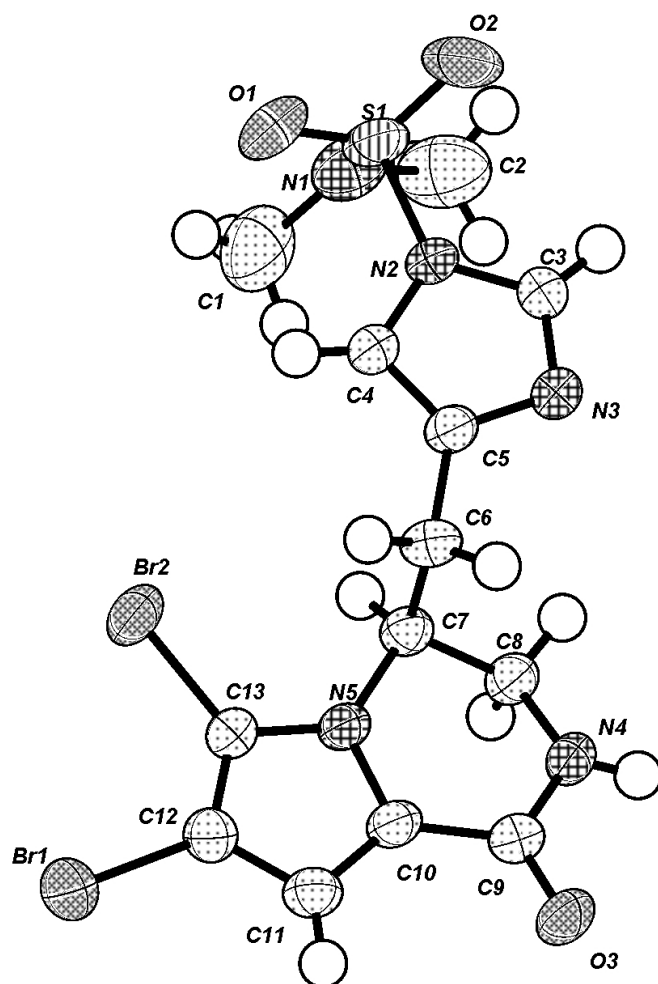
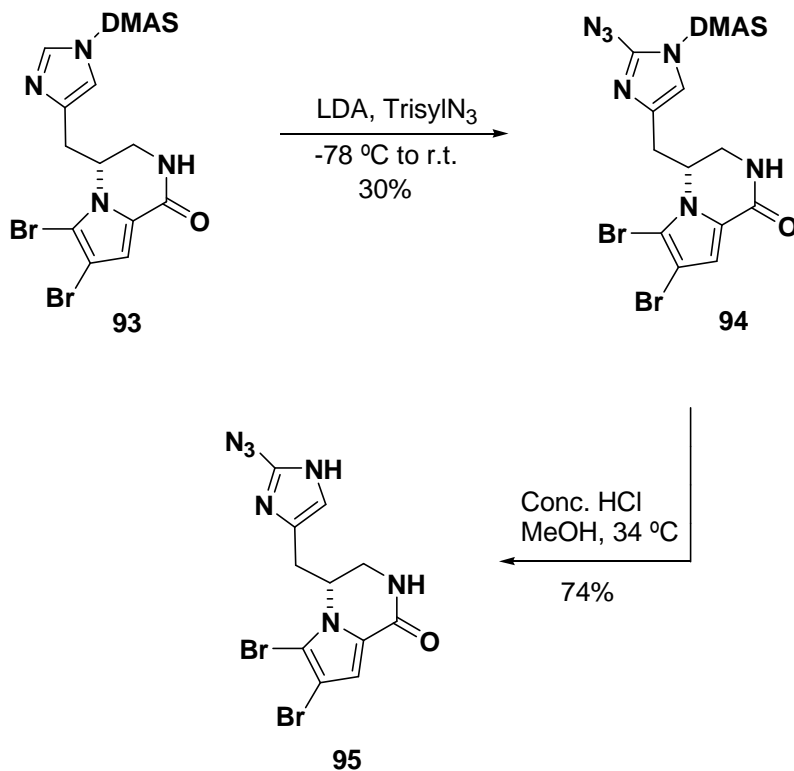


Figure 2.3 X-ray crystal structure of compound **88**

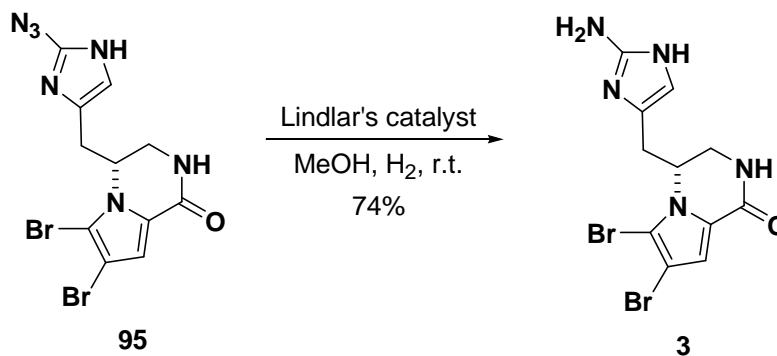
At this stage, all that remained to complete the synthesis was the incorporation of the C-2 amino group and removal of protecting group. Accordingly azide group was introduced at the C-2 position of the compound **93** by LDA at $-78\text{ }^{\circ}\text{C}$ to generate the anion at the C-2 position and then treatment with TrisylN_3 .⁴⁶ Attempts to improve the efficiency of the C-2 azidation were not successful. The

imidazole protecting group was removed by using conc. HCl in MeOH at 34 °C to produce compound **94** (Scheme 2.27).⁴⁷



Scheme 2.27

Azide reduction with Lindlar's catalyst at the C-2 position provided the primary amine, which was achieved in 74% yield (Scheme 2.28).⁴⁷ Finally the NMR data of our synthetically cyclooroidin matched with the isolated cyclooroidin and the specific rotation in our synthetic version was of correct magnitude ($[\alpha]_D = 10.6$ ($c = 0.02$, MeOH)) but of opposite direction compared to the natural product ($[\alpha]_D - 12$ ($c 0.02$, MeOH)).



Scheme 2.28

2.6 Summary

In summary, we have developed an approach to the oxazoline-containing fragment of the oroidin dimer nagelamide R and the recently isolated nagelamide T. The key transformation involves the intramolecular cyclization of a pyrrolecarboxamide via the carbonyl oxygen leading to the formation of oxazoline, whose subsequent elaboration by bromination and azidation provides the fully functionalized northern fragment of nagelamide R. This chemistry suggests a possible biosynthetic pathway to the formation of nagelamide R through nagelamide B via activation and substitution of the amide carbonyl.

We have also developed a concise enantioselective route for total synthesis of cyclooroidin. The key steps here involved a coupling reaction between alcohol **48** and pyrrole 2-carboxylic acid followed by an intramolecular cyclization to form lactone ring **77**. The lactone was then converted to pyrazine **55**. Bromine was introduced into the pyrrole ring of the molecule **55** and finally azide was introduced at the C-2 position of the imidazole of the molecule **93**. Synthetic, cyclooroidin was obtained by removing the protecting group and finally reducing the C-2 azide to the amine.

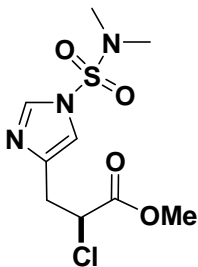
CHAPTER 3

EXPERIMENTAL DETAILS

3.1 General procedures

All reagents were purchased from commercial suppliers and were used as received unless otherwise noted. ^1H and ^{13}C NMR (δ in ppm) spectra were recorded in CDCl_3 (unless otherwise noted) at 300 and 75 MHz, respectively; using a JEOL Eclipse+ 300 spectrometer unless otherwise noted using residual CHCl_3 as reference (^1H NMR and carbon absorption of CDCl_3 for ^{13}C NMR). Infrared spectra were recorded either as neat films or as KBr pellets using a Bruker Vector 22 spectrometer. Elemental analyses were performed using a Perkin-Elmer 2400 CHN analyzer. Optical rotation was measured on a Perkin-Elmer 241MC polarimeter ($c = \text{g}/100 \text{ mL}$) and the observed value was an average of 2-3 runs. The solvent used for optical rotation was MeOH unless otherwise noted. High resolution mass spectra (HR-MS) were obtained by Dr. Powell through the mass spectrometry service at the University of Florida, Gainesville, Florida.

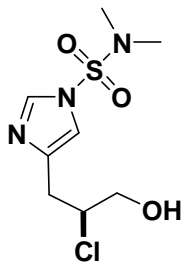
(S)-methyl 2-chloro-3-(1-(*N,N*-dimethylsulfamoyl)-1H-imidazol-4-yl)propanoate (47):



In a round bottom flask containing dry DMF (100 mL) chloroester **45**¹ (10.0 g, 0.05 mol) was added under N₂ atmosphere. The reaction mixture was stirred and to it triethylamine (11.1 mL, 0.08 mol) was added dropwise. Then the reaction mixture was cooled to 0 °C and DMASCI (8.4 mL, 0.08 mol) was added dropwise. The reaction mixture was allowed to come to r.t. and then the reaction mixture was heated to 57 °C and stirred for 9 h. Aqueous sat. NaHCO₃ (100 mL) and water (100 mL) were added to the reaction mixture and then the aqueous layer was repeatedly washed with ethyl acetate. The combined organic layer was dried using anhydrous Na₂SO₄ and concentrated to afford the crude product, which was purified by chromatography (hexane/EtOAc: 1/1) to give a colorless solid (11.1 g, 71%). [α]_D = -19.6 (c = 0.4, MeOH). m.p. 75-77 °C. ¹H NMR: δ = 7.81 (s, 1H), 7.09 (s, 1H), 4.59 (t, *J* = 7.2 Hz, 1H), 3.75 (s, 3H), 3.30 (dd, *J* = 15.1, 6.9 Hz, 1H), 3.15 (dd, *J* = 15.1, 7.2 Hz, 1H), 2.82 (s, 6H); ¹³C NMR: δ = 169.5, 138.4, 136.6, 115.7, 55.4, 53.1, 38.2, 33.7; FT-IR (neat, cm⁻¹): 3125, 2955, 1747, 1391, 1274, 1174, 1083, 963, 728, 594; HR-MS (*m/z*): calc for [M+H]⁺ C₉H₁₅ClN₃O₄S is 296.0472 found 296.0466.

(S)-4-(2-chloro-3-hydroxypropyl)-N,N-dimethyl-1H-imidazole-1-sulfonamide

(48): The chloro ester **47** (2.00 g, 6.75 mmol) was dissolved in dry isopropyl



alcohol (30 mL) under N₂ atmosphere. The reaction mixture was

stirred and to it was added anhydrous CaCl₂ (2.24 g, 2.0 mmol)

followed by the addition of anhydrous NaBH₄ (0.50 g, 1.3 mmol).

The resulting reaction mixture was stirred 24 h. After completion

of the reaction, water (40 mL) was added and then dil. HCl was

added until the solution pH becomes ~ 4 and finally NaHCO₃ was added to

reaction mixture to make the pH 8. Then the aqueous layer was repeatedly

extracted with CH₂Cl₂. The combined organic solutions were dried using

anhydrous Na₂SO₄ and concentrated. Purification of the crude product by

chromatography (EtOAc) provided the product as a colorless liquid (1.4 g, 70%).

[α]_D = -5.0 (*c* = 0.4, MeOH) ¹H NMR: δ = 7.84 (s, 1H), 7.12 (s, 1H), 4.31-4.28 (m,

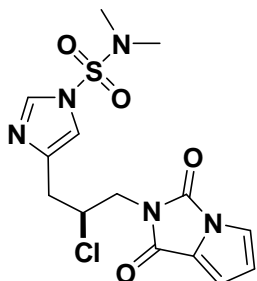
1H), 3.74-3.72 (m, 2H), 3.70 (brs, 1H), 3.10 (d, *J* = 5.8 Hz, 2H), 2.84 (s, 6H). ¹³C

NMR (CDCl₃, 125.8 MHz): δ = 139.1, 136.5, 115.8, 65.8, 61.4, 38.3, 33.2. FT-IR

(neat, cm⁻¹): 3127, 2927, 1647, 1570, 1473, 1391, 964, 837, 675. HR-MS (*m/z*):

calc for [M+H]⁺ C₈H₁₅ClN₃O₃S 268.0523 Found 268.0528.

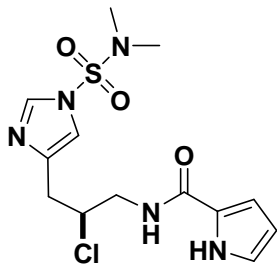
(S)-4-(2-chloro-3-(1,3-dioxo-1H-pyrrolo[1,2-c]imidazol-2(3H)-yl)propyl)-N,N-dimethyl-1H-imidazole-1-sulfonamide (50): In a round bottom flask containing



hydantoin² **49** (0.61 g, 4.5 mmol), triphenylphosphine (1.18 g, 4.5 mmol), were dissolved in dry THF (32 mL) under an N₂ atmosphere. The reaction mixture was cooled to (0 °C) and DIAD (1.06 g, 5.25 mmol) was added dropwise. After 30 min, chloroalcohol **48** (1.00 g, 3.75 mmol) dissolved in THF (5 mL) was added to the reaction mixture dropwise.

The reaction mixture was stirred for 12 h and then concentrated. The crude product was purified by chromatography (hexane/EtOAc, 3:7) to provide **50** as a yellow oil (0.98 g, 68%). ¹H NMR: δ = 7.81 (s, 1H), 7.26 (s, 1H), 7.17 (s, 1H), 6.79 (d, *J* = 3.4 Hz, 1H), 6.47 (t, *J* = 3.4 Hz, 1H), 4.59 (quint, *J* = 7.6 Hz, 1H), 3.96–3.92 (m, 2H), 3.15 (dd, *J* = 15.1, 5.2 Hz, 1H), 3.04 (dd, *J* = 15.1, 7.6 Hz, 1H), 2.84 (s, 6H); ¹³C NMR: δ = 158.1, 148.6, 139.1, 136.5, 125.1, 119.5, 117.8, 115.8, 114.2, 57.3, 44.5, 38.3, 34.9; FT-IR (neat, cm⁻¹): 2969, 2926, 1793, 1733, 1373, 1339, 1266. HR-MS (*m/z*): calc for [M+H]⁺ C₁₄H₁₇ClN₅O₄S 386.0690 Found 386.0681.

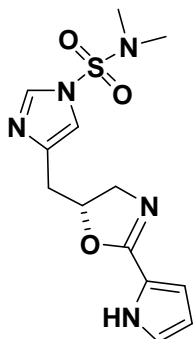
(S)-N-(2-chloro-3-(1-(*N,N*-dimethylsulfamoyl)-1H-imidazol-4-yl)propyl)-1H-pyrrole-2-carboxamide (54**):**



The Mitsunobu product **50** (0.90 g, 2.3 mmol) was dissolved in THF (10 mL) and to it was added 20% NaOH (7 mL, 1.40 g, 3.5 mmol). The reaction mixture was stirred at r.t. for 1 h. The organic solution was separated and the aqueous solution was extracted with EtOAc (3x100 mL). The combined organic solutions were dried with anhydrous Na₂SO₄ and concentrated, and the crude product was purified by chromatography (CHCl₃/MeOH, 99:1) to afford a colorless liquid (0.58 g, 70%). ¹H NMR: δ = 9.45 (bs, 1H), 7.86 (s, 1H), 7.15 (s, 1H), 6.93 (s, 1H), 6.70 (t, *J* = 6.5 Hz, 1H), 6.65 (d, *J* = 2.4 Hz, 1H), 6.25 (d, *J* = 3.1 Hz, 1H), 4.43 (m, 1H), 3.78 (m, 1H), 3.66 (m, 1H), 3.08 (d, *J* = 5.8 Hz, 2H), 2.85 (s, 6H); ¹³C NMR: δ = 161.0, 139.0, 136.5, 125.6, 121.8, 115.9, 110.1, 109.4, 59.7, 44.3, 38.3, 34.6; FT-IR (neat, cm⁻¹): 2969, 2926, 1620, 1450, 1392, 1272, 1166, 1074, 953, 732. HR-MS (*m/z*): calc for [M+H]⁺ C₁₃H₁₉ClN₅O₃S 360.0897 Found 360.0887.

(R)-4-((2-(1H-pyrrol-2-yl)-4,5-dihydrooxazol-5-yl)methyl)-N,N-dimethyl-1H-

imidazole-1-sulfonamide (56): The Mitsunobu product **50** (0.90 g, 2.0 mmol)



was dissolved in THF (10 mL) and to it was added 20% NaOH (7 mL, 1.40 g, 3.5 mmol). The reaction mixture was heated at 75 °C for 12 h. After cooling to r.t. the organic solution was separated and the aqueous solution was extracted with EtOAc (3x100 mL). The combined organic solutions were dried with anhydrous Na₂SO₄ and concentrated, and the crude product

was purified by chromatography (CHCl₃/MeOH, 98:2) to afford a white solid (0.45 g, 60%). [α]_D = -18.0 (c = 0.6, MeOH). m.p. 155-157 °C. ¹H NMR: δ = 9.71 (bs, 1H), 7.82 (s, 1H), 7.05 (s, 1H), 6.88 (s, 1H), 6.70 (d, J = 3.4 Hz, 1H), 6.23 (t, J = 3.1 Hz, 1H), 5.02 (m, 1H), 4.06 (dd, J = 14.4, 9.3 Hz, 1H), 3.85 (dd, J = 14.4, 6.9 Hz, 1H), 2.98 (d, J = 5.5 Hz, 2H), 2.69 (s, 6H); ¹³C NMR: δ = 158.9, 138.8, 136.8, 122.5, 119.9, 115.4, 112.9, 109.7, 77.6, 58.3, 37.9, 33.6; FT-IR (neat, cm⁻¹): 2969, 2926, 1793, 1733, 1373, 1339, 1266. HR-MS (m/z): calc for [M+H]⁺ C₁₃H₁₈N₅O₃S 324.1130 Found 324.1120.

(R)-4-((2-(4,5-dibromo-1H-pyrrol-2-yl)-4,5-dihydrooxazol-5-yl)methyl)-N,N-

dimethyl-1H-imidazole-1-sulfonamide (58) and (R)-4-((2-(4-bromo-1H-pyrrol-

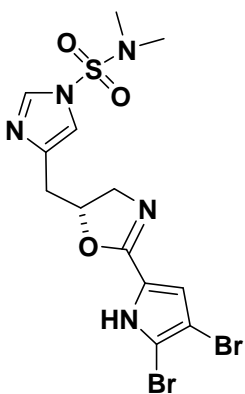
2-yl)-4,5-dihydrooxazol-5-yl)methyl)-N,N-dimethyl-1H-imidazole-1-

sulfonamide (59): The cyclized compound **56** (400 mg, 1.23 mol) was dissolved

in THF (12 mL). The reaction mixture was cooled to -20 °C, NBS (330 mg, 1.83 mmol) was added and then stirred at -20 °C for 8 h. The reaction mixture was

quenched by addition of aq. NaHCO₃ and then the organic layer was separated. The aqueous layer was extracted with EtOAc (3x50 mL), then the combined organic solutions were dried with anhydrous Na₂SO₄ and concentrated. The crude product was purified by chromatography (CHCl₃/MeOH, 98:2) providing **58** as a white solid (240 mg, 43%). [α]_D = -22.1 (c = 0.04, MeOH)

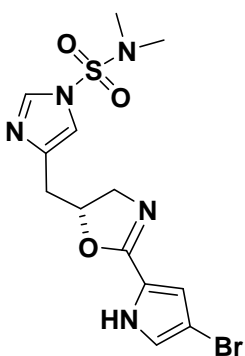
m.p. 62-64 °C. ¹H NMR: δ = 7.84 (s, 1H), 7.07 (s, 1H), 6.69 (s, 1H), 5.05 (quint, J = 5.9 Hz, 1H), 4.12 (dd, J = 14.4, 7.6 Hz, 1H), 3.92 (dd, J = 14.2, 6.9 Hz, 1H), 3.02 (dd, J = 15.1, 6.4 Hz, 1H), 2.95 (dd, J = 14.7, 5.9 Hz, 1H), 2.79 (s, 6H); ¹³C NMR: δ = 157.8, 138.7, 136.6, 121.4, 115.9, 115.3, 105.8, 99.6, 79.3, 58.1, 37.8, 33.6; FT-IR (neat, cm⁻¹): 1701, 1645, 1433, 1389, 1177, 1086, 962, 826, 738. HR-MS (m/z): calc for [M+H]⁺ C₁₃H₁₆Br₂N₅O₃S is 479.9341 found 479.9335.



Further elution provided the monobrominated product **59** as a white solid (150 mg, 34%). [α]_D = -35.2 (c = 0.04, MeOH). m.p 145-147 °C. ¹H NMR: δ = 7.81

(s, 1H), 7.05 (s, 1H), 6.85 (s, 1H), 6.66 (s, 1H), 5.04–5.02 (m, 1H), 4.06 (dd, J = 13.8, 9.6 Hz, 1H), 3.87 (dd, J = 14.7, 6.9 Hz, 1H), 2.98–2.94 (m, 2H) 2.71 (s, 6H); ¹³C NMR: δ = 157.9, 138.9, 136.6, 122.2, 120.5, 115.3, 114.7, 97.4, 78.6, 58.3, 38.0, 33.6; FT-IR (neat, cm⁻¹): 1649, 1559, 1537, 1504, 1457, 1398, 1173, 1078, 962. HR-MS (m/z): calc for

[M+H]⁺ C₁₃H₁₇BrN₅O₃S 402.0235 Found 402.0229.



(R)-2-azido-4-((2-(4,5-dibromo-1H-pyrrol-2-yl)-4,5-dihydrooxazol-5-

yl)methyl)-N,N-dimethyl-1H-imidazole-1-sulfonamide (60): Dibromopyrrole

58 (160 mg, 0.31 mmol) was dissolved in anhydrous THF (8 mL) and the reaction

mixture was cooled to -78 °C and 0.5 M LDA (2.72 mL,

4.1 eq) in THF was added dropwise to the reaction

mixture. The reaction mixture was left stirring for 30 min

at -78 °C and then TsN₃ (0.28 g, 1.07 mmol) was added.

The reaction mixture was allowed to come to r.t. and then

stirred for an additional 1 h, followed by addition of

aqueous NH₄Cl to quench the reaction mixture. The

organic layer was separated and the aqueous layer was extracted with EtOAc

(3x50 mL). The organic solutions were combined, dried with anhydrous Na₂SO₄

and concentrated. The crude product was purified by chromatography

(CHCl₃/MeOH, 98:2) giving **22** as a white solid (120 mg, 71%). [α]_D = -36.5 (c =

0.04, MeOH). m.p. 70-72 °C. ¹H NMR (CDCl₃, 500 MHz): δ = 6.93 (s, 1H), 6.69

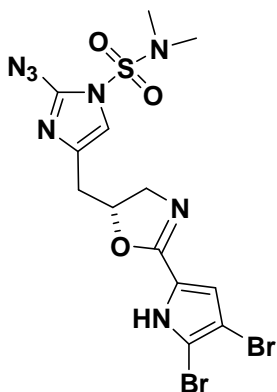
(s, 1H), 5.01 (quint, J = 6.4 Hz, 1H), 4.11 (dd, J = 14.5, 7.6 Hz, 1H), 3.87 (dd, J =

14.2, 6.8 Hz, 1H), 2.93 (s, 6H), 2.91- 2.88 (m, 2H); ¹³C NMR: δ = 143.7, 139.8,

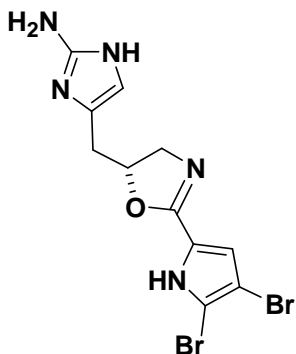
135.1, 129.4, 126.3, 121.3, 116.5, 116.3, 78.9, 57.8, 38.8, 33.3; FT-IR (neat, cm⁻¹):

2145, 1649, 1534, 1509, 1457, 1398, 1336, 1181, 1086, 990. HR-MS (m/z):

calc for [M+H]⁺ C₁₃H₁₅Br₂N₈O₃S is 520.9355 found 520.9355.

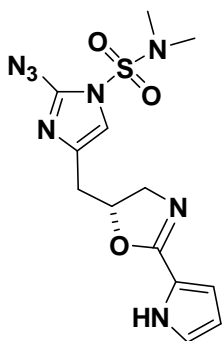


4-(((R)-2-(4,5-dibromo-1H-pyrrol-2-yl)-4,5-dihydrooxazol-5-yl)methyl)-2,3-dihydro-1H-imidazol-2-amine (57):⁴⁷



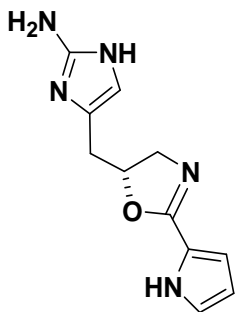
Conc. HCl (0.003 mL) was added to the solution of compound **60** (100 mg, 0.19 mmol) in methanol (3.5 mL) and then stirred at 34 °C for 12 h. Then the mixture was concentrated under vacuum and the residue was neutralized with NaHCO₃ and extracted with EtOAc (3 times). The organic extracts were combined, concentrated and purified by column chromatography (CHCl₃/methanol/NH₃, 94:5:1) to give the free imidazole (50 mg, 63%) as yellow thick liquid. This compound (30 mg, 0.05 mmol) was dissolved in methanol (2 mL) and Lindlar catalyst (20 mg) was added and the reaction mixture was stirred for 8 h under a hydrogen atmosphere (balloon). Finally the reaction mixture was filtered through a bed of Celite and then the crude reaction mixture was loaded onto a column as a solid, preadsorbed on silica gel. Then the crude product was purified by chromatography (CH₂Cl₂/MeOH/NH₄OH: 85/14/1). A colorless thick liquid was obtained (13 mg, 46%). $[\alpha]_D = -8.5$ ($c = 0.04$, MeOH) ¹H NMR (CD₃OD): $\delta = 6.73$ (s, 1H), 6.37 (s, 1H), 4.94–4.87 (m, 1H), 3.99 (dd, $J = 15.2, 10.3$ Hz, 1H), 3.69 (dd, $J = 14.4, 5.8$ Hz, 1H), 2.78–2.72 (m, 2H); ¹³C NMR (CD₃OD): $\delta = 157.7, 149.6, 128.4, 121.5, 115.0, 111.1, 104.9, 99.1, 79.6, 58.4, 32.1$. This compound has been recently been described in the literature.^{43b}

(R)-4-((2-(1H-pyrrol-2-yl)-4,5-dihydrooxazol-5-yl)methyl)-2-azido-N,N-dimethyl-1H-imidazole-1-sulfonamide (62):



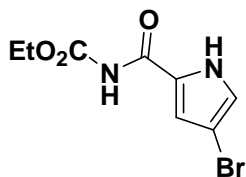
The cyclized compound **56** (160 mg, 0.49 mmol) was dissolved in anhydrous THF (8 mL) and the reaction mixture was cooled to $-78\text{ }^{\circ}\text{C}$ and 0.5 M LDA (2.06 mL, 2.1 eq) was added dropwise to the reaction mixture. The reaction mixture was left stirring for 30 min at $-78\text{ }^{\circ}\text{C}$ and then TsN_3 (0.30 g, 1.56 mmol) was added. The reaction mixture was allowed to come to r.t. and then stirred for an additional 1 h, followed by addition of aqueous NH_4Cl to quench the reaction mixture. The organic layer was separated and the aqueous layer was extracted with EtOAc (3x50 mL). The organic solutions were combined, dried with anhydrous Na_2SO_4 and concentrated. Then the crude product was purified by chromatography ($\text{CHCl}_3/\text{MeOH}$, 98:2) giving **62** as a yellow liquid (150 mg, 84%). ^1H NMR (CDCl_3 , 500 MHz): δ = 6.92 (s, 1H), 6.89 (s, 1H), 6.72 (d, J = 4.1 Hz, 1H), 6.23 (t, J = 3.7 Hz, 1H), 4.96 (quint, J = 5.9 Hz, 1H), 4.05 (dd, J = 15.1, 8.7 Hz, 1H), 3.78 (dd, J = 13.3, 7.3 Hz, 1H), 2.93 (s, 6H), 2.89- 2.82 (m, 2H); ^{13}C NMR: δ = 158.5, 134.8, 121.9, 120.6, 116.8, 112.6, 110.0, 78.7, 58.7, 38.4, 33.8; FT-IR (neat, cm^{-1}): 2971, 2145, 1651, 1534, 1509, 1457, 1394, 1342, 1170, 1067, 950. HR-MS (m/z): calc for $[\text{M}+\text{H}]^+$ $\text{C}_{13}\text{H}_{17}\text{N}_8\text{O}_3\text{S}$ is 365.1139 found 365.1143.

4-(((R)-2-(1H-pyrrol-2-yl)-4,5-dihydrooxazol-5-yl)methyl)-2,3-dihydro-1H-imidazol-2-amine (64):



Conc. HCl (0.004 mL) was added to the solution of compound **62** (150 mg, 0.41 mmol) in methanol (3.5 mL) and stirred at 34 °C for 12 h. Then the mixture was concentrated under vacuum and the residue was neutralized with NaHCO₃ and extracted with EtOAc (3 times). The organic layer were combined, concentrate and purified by column chromatography (CHCl₃/methanol/NH₄OH, 92:7:1) to give DMAS-deprotected product (76 mg, 72%) as yellow thick liquid. This compound (50 mg, 0.19 mmol) was dissolved in methanol (3 mL) and Lindlar catalyst (30 mg) was added and the reaction mixture was stirred for 8 h under a hydrogen atmosphere (balloon). Finally the reaction mixture was filtered through a bed of Celite and then the crude reaction mixture was loaded onto a column as a solid, preadsorbed on silica gel. Then the crude product was purified by chromatography (CH₂Cl₂/MeOH/NH₃: 82/17/1). A colorless thick liquid **64** was obtained (26 mg, 56%). ¹H NMR (CD₃OD): δ = 6.89 (s, 1H), 6.72 (d, *J* = 3.7 Hz, 1H), 6.37 (s, 1H), 6.12 (t, *J* = 3.4 Hz, 1H), 4.95–4.90 (m, 1H), 3.94 (dd, *J* = 13.7, 9.4 Hz, 1H), 3.69 (dd, *J* = 14.1, 6.8 Hz, 1H), 2.82 (dd, *J* = 14.8, 5.8 Hz, 1H), 2.74 (dd, *J* = 14.8, 6.5 Hz, 1H); ¹³C NMR (CD₃OD): δ = 159.5, 149.5, 128.5, 121.8, 119.2, 112.8, 111.3, 109.1, 78.9, 58.8, 32.2.; FT-IR (neat, cm⁻¹): 3050, 2887, 1601, 1407, 1364, 1260, 1164, 1054, 927. HR-MS (*m/z*): calc for [M+H]⁺ C₁₁H₁₄N₅O is 232.1198 found 232.1193.

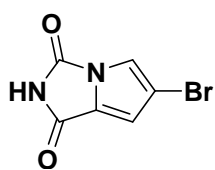
Ethyl 4-bromo-1H-pyrrole-2-carboxylate (65):



Ethyl-1-pyrrole-2-carboxylate (3.0 g, 0.02 mol, 1 eq) was dissolved in AcOH (59 mL). A solution of bromine (0.76 mL, 0.01 mol, 0.9 eq) in AcOH (6.4 mL) was added dropwise at r.t. The reaction mixture was stirred for 30 mins, after which a white precipitate appeared in the reaction flask. The resulting mixture was filtered and the collected solids were washed with cold dichloromethane (10 mL). Then dried under vacuum to give white solid (2.4 g, 56%). m.p. 193-195 °C. ¹H NMR (DMSO-d₆): δ = 12.14 (b, 1H), 10.56 (s, 1H), 7.21 (s, 1H), 7.13 (s, 1H), 4.10 (q, *J* = 7.3 Hz, 2H) 1.22 (t, *J* = 7.3 Hz, 3H); ¹³C NMR (DMSO): δ = 157.8, 151.8, 125.9, 124.8, 115.5, 96.1, 61.5, 14.8; FT-IR (neat, cm⁻¹): 3126, 2975, 2931, 1707, 1681, 1370, 1187, 1002, 979, 919, 736. HR-MS (*m/z*): calc for [M+H]⁺ C₈H₁₀BrN₂O₃ is 260.9875 found 260.9869.

6-bromo-1H-pyrrolo[1,2-c]imidazole-1,3(2H)-dione (67):

Monobromopyrrole **65** (2.0 g, 9.3 mmol) was mixed with silicone oil (2 mL) in a

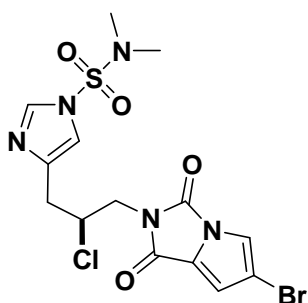


round bottom flask which was evacuated (40 mmHg) and heated to 180-185 °C. After 4 h, the reaction was cooled and then the residue was crushed. Hexane was added and the solids were collected by filtration and washed repeatedly with hexane. The procedure was repeated with (1.5 mL) silicone oil. The product was purified by chromatography (hexane/EtOAc: 8/2). A yellow solid was thus obtained (750 mg, 47%). m.p. 216-218 °C. ¹H NMR (DMSO-d₆, 500 MHz): δ = 11.60 (b, 1H), 7.81

(s, 1H), 7.01 (s, 1H); ^{13}C NMR (DMSO- d_6 , 125.8 MHz): δ = 159.2, 148.7, 126.9, 119.1, 113.9, 104.7; FT-IR (neat, cm^{-1}): 3285, 1801, 1733, 1558, 1452, 1409, 1371, 1311. Anal. Calcd. for $\text{C}_6\text{H}_3\text{BrN}_2\text{O}_2$: C, 33.52; H, 1.41; N, 13.03. Found: C, 33.62; H, 1.54; N, 12.62.

(S)-4-(3-(6-bromo-1,3-dioxo-1H-pyrrolo[1,2-c]imidazol-2(3H)-yl)-2-

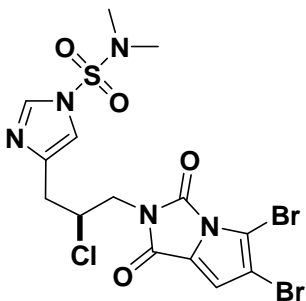
chloropropyl)-N,N-dimethyl-1H-imidazole-1-sulfonamide (69): In a round



bottom flask containing the bromohydantoin **67** (120 mg, 0.56 mmol), triphenylphosphine (150 mg, 0.56 mmol), was dissolved in dry THF (5 mL) under N_2 atmosphere. The mixture was cooled to (0 $^\circ\text{C}$) and DEAD (0.26 mL, 0.56 mmol, 40% wt in toluene) was

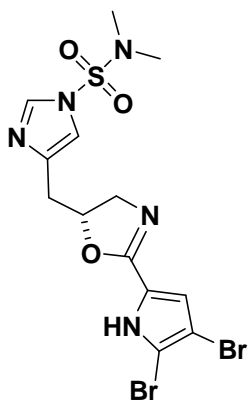
added dropwise. After 30 min, the chloroalcohol **48** (100 mg, 0.37 mmol) dissolved in THF (2 mL) was added to the reaction mixture was dropwise. The reaction mixture was stirred for 12 h, concentrated and then the crude product was purified by chromatography (hexane/EtOAc, 6:4) providing a colorless solid (0.15 g, 86%). $[\alpha]_D = -5.0$ ($c = 0.04$, MeOH). m.p. 154-156 $^\circ\text{C}$. ^1H NMR: δ = 7.82 (s, 1H), 7.29 (s, 1H), 7.16 (s, 1H), 6.79 (s, 1H), 4.60–4.56 (m, 1H), 3.96–3.93 (m, 2H), 3.10 (dd, $J = 15.1, 5.5$ Hz, 1H), 3.06 (dd, $J = 14.8, 7.6$ Hz, 1H), 2.85 (s, 6H); ^{13}C NMR: δ = 157.4, 147.6, 138.9, 136.5, 124.8, 118.7, 115.8, 115.7, 105.9, 57.2, 44.7, 38.2, 34.9; FT-IR (neat, cm^{-1}): 1797, 1733, 1562, 1392, 1272, 1166, 1074, 953, 732. HR-MS (m/z): calc for $[\text{M}+\text{H}]^+$ $\text{C}_{14}\text{H}_{16}\text{BrClN}_5\text{O}_4\text{S}$ is 463.9795 found 463.9789.

(S)-4-(2-chloro-3-(5,6-dibromo-1,3-dioxo-1H-pyrrolo[1,2-c]imidazol-2(3H)-yl)propyl)-N,N-dimethyl-1H-imidazole-1-sulfonamide (70):



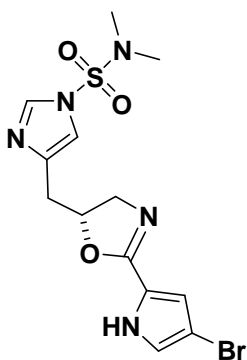
Dibromohydantoin³ **68** (100 mg, 0.34 mmol) and triphenylphosphine (90 mg, 0.33 mmol) were dissolved in dry THF (5 mL) under N₂ atmosphere and cooled to 0 °C. DEAD (0.15 mL, 0.33 mmol, 40% wt in toluene) was added dropwise then stirred for 30 min. Chloroalcohol **48** (60 mg, 0.22 mmol) dissolved in THF (3 mL) was added to the reaction mixture was dropwise. The reaction mixture was stirred for 12 h and then concentrated. The crude product was purified by chromatography (hexane/EtOAc, 7:3) affording a yellow liquid (0.10 g, 82%). $[\alpha]_D = -3.0$ ($c = 0.04$, MeOH). ¹H NMR: $\delta = 7.82$ (s, 1H), 7.16 (s, 1H), 6.85 (s, 1H), 4.61–4.56 (m, 1H), 3.96–3.94 (m, 2H), 3.10 (dd, $J = 15.1, 5.5$ Hz, 1H), 3.04 (dd, $J = 14.8, 7.6$ Hz, 1H), 2.84 (s, 6H); ¹³C NMR: $\delta = 156.1, 146.8, 139.1, 136.6, 125.3, 115.9, 115.7, 109.4, 105.0, 57.1, 44.5, 38.0, 34.8$; FT-IR (neat, cm⁻¹): 2907, 2359, 1796, 1734, 1423, 1388, 1260, 1167, 1093, 953, 736. HR-MS (m/z): calc for $[M+H]^+$ C₁₄H₁₅Br₂ClN₅O₄S is 541.8900 found 541.9506.

(R)-4-((2-(4,5-dibromo-1H-pyrrol-2-yl)-4,5-dihydrooxazol-5-yl)methyl)-N,N-dimethyl-1H-imidazole-1-sulfonamide (58):



The Mitsunobu product (80 mg, 0.15 mmol) was dissolved in THF (6 mL) and to it (0.4 mL, 80 mg, 2.2 mmol) of 20% NaOH was added. The reaction mixture was heated to 75 °C for 6 h. After cooling to r.t., the organic solution was separated and the aqueous solution was extracted with EtOAc (3x50 mL) times. The organic solutions were combined and dried with anhydrous Na₂SO₄ and concentrated. Then the crude product was purified by chromatography (CHCl₃/MeOH, 98:2) providing a white solid (40 mg, 57%) which was identical in all respects to that described above.

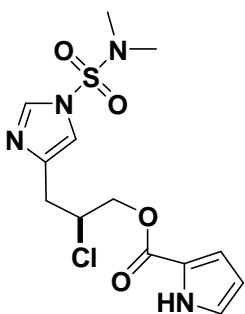
(R)-4-((2-(4-bromo-1H-pyrrol-2-yl)-4,5-dihydrooxazol-5-yl)methyl)-N,N-dimethyl-1H-imidazole-1-sulfonamide (59):



The Mitsunobu product (110 mg, 0.24 mmol) was dissolved in THF (6 mL) and 20% NaOH (0.7 mL) was added. The reaction mixture was heated to 75 °C for 8 h. After cooling to r.t., the organic solution was separated and the aqueous solution was extracted with EtOAc (3x50 mL). The organic solutions were combined together, dried with anhydrous Na₂SO₄ and concentrated.

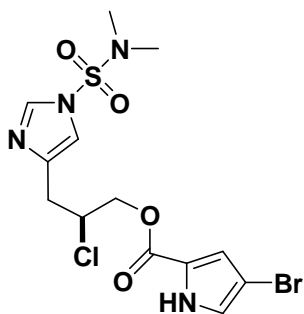
The crude product was purified by chromatography (CHCl₃/MeOH, 98:2) providing a white solid (60 mg, 64%) that was identical in all respects to the material described above.

(S)-2-chloro-3-(1-(*N,N*-dimethylsulfamoyl)-1*H*-imidazol-4-yl)propyl 1*H*-pyrrole-2-carboxylate (78**):**



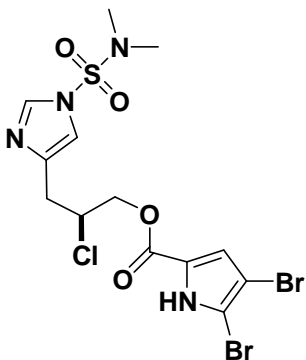
In a round bottom flask the alcohol **48** (2.60 g, 9.73 mmol), acid **79** (1.60 g, 14.4 mmol), DMAP (120 mg, 0.98 mmol) and camphorsulfonic acid (140 mg, 0.60 mmol) were dissolved in dry CH₂Cl₂ (150 mL). The mixture was cooled to (-78 °C) and DCC (2.80 g, 13.6 mmol) in dry CH₂Cl₂ (8 mL) was added dropwise. The reaction mixture was allowed to warm up to r.t. and stirred for an additional 8 h. The resulting mixture was filtered through Celite and the filter cake was washed with CH₂Cl₂. Concentration of the filtrate provided the crude product, which was purified by chromatography (hexane/EtOAc, 1:9) to provide **78** as a thick colorless liquid (2.04 g, 58%). [α]_D = -2.0 (c = 0.04, MeOH). ¹H NMR: δ = 9.36 (b, 1H), 7.85 (d, J = 1.0 Hz, 1H), 7.14 (s, 1H), 7.00-6.96 (m, 2H), 6.28 (q, J = 3.1 Hz, 1H), 4.55-4.44 (m, 3H), 3.21 (dd, J = 14.4, 4.8 Hz, 1H), 3.02 (dd, J = 14.4, 7.6 Hz, 1H), 2.84 (s, 6H); ¹³C NMR: δ = 160.4, 139.3, 136.6, 123.5, 122.0, 116.2, 115.7, 110.8, 66.5, 57.6, 38.3, 33.9; FT-IR (neat, cm⁻¹): 3150, 2928, 1695, 1422, 1396, 1173, 1074, 959, 722, 599; HR-MS (m/z): calc for [M+H]⁺ C₁₃H₁₈ClN₄O₄S 361.0737 found 361.0747.

(S)-2-chloro-3-(1-(*N,N*-dimethylsulfamoyl)-1*H*-imidazol-4-yl)propyl 4-bromo-1*H*-pyrrole-2-carboxylate (85**):**



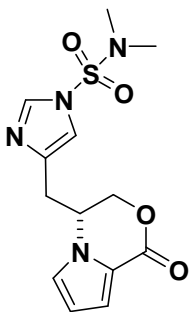
In a round bottom flask the alcohol **48** (1.00 g, 3.74 mmol), acid **83** (1.06 g, 5.61 mmol), DMAP (45 mg, 0.37 mmol) and camphorsulfonic acid (52 mg, 0.22 mmol) were dissolved in dry CH₂Cl₂ (100 mL). The mixture was cooled (-78 °C) and DCC (1.16 g, 5.61 mmol) in dry CH₂Cl₂ (4 mL) was added dropwise. The reaction mixture was allowed to warm up to r.t. and stirred for an additional 8 h. The resulting mixture was filtered through Celite and the filter cake was washed with CH₂Cl₂. Concentration of the filtrate provided the crude product, which was purified by chromatography (hexane/EtOAc, 1:9) to provide **85** as a thick colorless liquid (0.88 g, 54%). [α]_D = -1.5 (c = 0.06, MeOH). ¹H NMR (CDCl₃, 500 MHz): δ = 9.80 (b, 1H), 7.96 (s, 1H), 7.17 (s, 1H), 6.97-6.95 (m, 2H), 4.50-4.41 (m, 3H), 3.22 (dd, *J* = 15.1, 4.1 Hz, 1H), 3.06 (dd, *J* = 15.1, 7.3 Hz, 1H), 2.86 (s, 6H); ¹³C NMR (CDCl₃ 125.8 MHz): δ = 159.5, 138.6, 136.4, 123.4, 122.4, 118.7, 117.8, 116.0, 98.1, 66.9, 57.2, 38.3, 33.6; FT-IR (neat, cm⁻¹): 3095, 2915, 1712, 1540, 1393, 1311, 1173, 1079, 969, 725, 590; HR-MS (*m/z*): calc for [M+H]⁺ C₁₃H₁₇BrClN₄O₄S 438.9837 found 438.9843.

(S)-2-chloro-3-(1-(*N,N*-dimethylsulfamoyl)-1H-imidazol-4-yl)propyl 4,5-dibromo-1H-pyrrole-2-carboxylate (86**):**



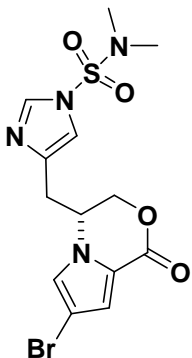
In a round bottom flask the alcohol **48** (200 mg, 0.75 mmol), acid **84** (300 mg, 1.12 mmol), DMAP (10 mg, 0.07 mmol) and camphorsulfonic acid (10 mg, 0.04 mmol) were dissolved in dry CH₂Cl₂ (20 mL). The mixture was cooled to (-78 °C) and DCC (230 mg, 1.12 mmol) in dry CH₂Cl₂ (4 mL) was added dropwise. The reaction mixture was allowed to warm up to r.t. and stirred for an additional 8 h. The resulting mixture was filtered through Celite and the filter cake was washed with CH₂Cl₂. Concentration of the filtrate provided the crude product, which was purified by chromatography (hexane/EtOAc, 1:9) to provide **86** as a thick colorless liquid (210 mg, 55%). [α]_D = -1.4 (*c* = 0.06, MeOH). ¹H NMR (CDCl₃, 500 MHz): δ = 10.68 (b, 1H), 7.90 (s, 1H), 7.16 (s, 1H), 6.94 (s, 1H), 4.50-4.41 (m, 3H), 3.21 (dd, *J* = 15.1, 4.6 Hz, 1H), 3.06 (dd, *J* = 15.1, 6.9 Hz, 1H), 2.86 (s, 6H); ¹³C NMR: δ = 158.8, 138.8, 136.6, 123.2, 118.7, 116.1, 107.7, 100.9, 66.8, 57.0, 38.3, 33.7; FT-IR (neat, cm⁻¹): 3085, 2890, 1706, 1405, 1323, 1182, 1082, 959, 907, 722, 584; HR-MS (*m/z*): calc for [M+H]⁺ C₁₃H₁₆Br₂ClN₄O₄S 516.8948 found 516.8997.

(R)-N,N-dimethyl-4-((1-oxo-3,4-dihydro-1H-pyrrolo[2,1-c][1,4]oxazin-4-yl)methyl)-1H-imidazole-1-sulfonamide (77):



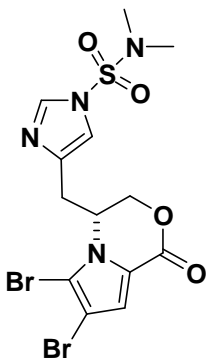
In a round bottom flask compound **78** (350 mg, 0.97 mmol) was dissolved in DMF (10 mL) and cesium carbonate (3.16 g, 9.70 mmol) was added to the reaction mixture. The reaction mixture was heated at 80 °C for 12 h. Water was added to the reaction mixture and it was extracted with EtOAc. The organic extract was dried with anhydrous Na₂SO₄, concentrated, and the residue was purified by chromatography (EtOAc) providing **77** as a thick colorless liquid (163 mg, 52%). $[\alpha]_D = -16.0$ ($c = 0.02$, MeOH). ¹H NMR: $\delta = 8.01$ (d, $J = 1.0$ Hz, 1H), 7.20 (dd, $J = 3.8, 1.4$ Hz, 1H), 6.99 (d, $J = 0.7$ Hz, 1H), 6.80 (t, $J = 1.7$ Hz, 1H), 6.33 (dd, $J = 2.4, 1.7$ Hz, 1H), 4.81-4.77 (m, 2H), 4.61 (dd, $J = 12.7, 3.1$ Hz, 1H), 3.23 (d, $J = 7.2$ Hz, 2H), 2.95 (s, 6H); ¹³C NMR: $\delta = 158.7, 138.5, 137.0, 124.3, 118.8, 117.9, 115.8, 110.7, 68.3, 52.6, 38.0, 31.8$; FT-IR (neat, cm⁻¹): 3204, 2885, 1665, 1551, 1478, 1390, 1334, 1173, 1079, 734, 596; HR-MS (m/z): calc for [M+H]⁺ C₁₃H₁₇N₄O₄S 325.0971 found 325.0986.

(R)-4-((7-bromo-1-oxo-3,4-dihydro-1H-pyrrolo[2,1-c][1,4]oxazin-4-yl)methyl)-*N,N*-dimethyl-1H-imidazole-1-sulfonamide (87**):**



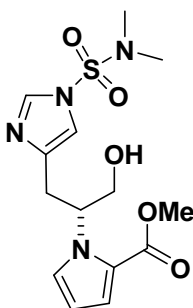
In a round bottom flask compound **85** (50 mg, 0.11 mmol) was dissolved in DMF (2 mL) and cesium carbonate (0.37 g, 1.13 mmol) was added to the reaction mixture. The reaction mixture was heated at 80 °C for 12 h. Water was added to the reaction mixture and it was extracted with EtOAc. The organic extract was dried with anhydrous Na₂SO₄, concentrated, and the residue was purified by chromatography on silica gel (EtOAc) providing **87** as a thick colorless liquid (25 mg, 55%). [α]_D = -6.0 (*c* = 0.02, MeOH). ¹H NMR: δ = 7.85 (d, *J* = 1.0 Hz, 1H), 7.08 (d, *J* = 4.1 Hz, 1H), 6.91 (s, 1H), 6.35 (d, *J* = 4.1 Hz, 1H), 4.75-4.68 (m, 1H), 4.59 (d, *J* = 1.7 Hz, 2H), 3.15-3.06 (m, 2H), 2.87 (s, 6H); ¹³C NMR: δ = 157.6, 138.1, 137.0, 119.8, 118.5, 115.8, 113.6, 107.3, 68.3, 51.8, 38.2, 30.8; FT-IR (neat, cm⁻¹): 3121, 2926, 1711, 1387, 1172, 1080, 1007, 962, 726, 590; HR-MS (*m/z*): calc for [M+H]⁺ C₁₃H₁₆BrN₄O₄S 403.0069 found 403.0069.

(R)-4-((6,7-dibromo-1-oxo-3,4-dihydro-1H-pyrrolo[2,1-c][1,4]oxazin-4-yl)methyl)-N,N-dimethyl-1H-imidazole-1-sulfonamide (88):



In a round bottom flask compound **86** (70 mg, 0.13 mmol) was dissolved in DMF (2.5 mL) and cesium carbonate (430 mg, 1.34 mmol) was added to the reaction mixture. The reaction mixture was heated at 80 °C for 12 h. Water was added to the reaction mixture and it was extracted with EtOAc. The organic extract was dried with anhydrous Na₂SO₄, concentrated, and the residue was purified by chromatography on silica gel (EtOAc) providing **86** as a thick colorless liquid (35 mg, 52%). [α]_D = -6.4 (*c* = 0.12, MeOH). ¹H NMR: δ = 7.85 (s, 1H), 7.12 (s, 1H), 6.95 (s, 1H), 4.71-4.61 (m, 1H), 4.60-4.56 (m, 2H), 3.20-2.99 (m, 2H), 2.86 (s, 6H); ¹³C NMR: δ = 156.3, 137.7, 137.1, 120.2, 119.7, 115.7, 108.9, 102.0, 68.3, 52.8, 38.3, 30.4; FT-IR (neat, cm⁻¹): 3130, 2950, 1733, 1399, 1331, 1170, 1085, 959, 910, 740, 584; HR-MS (*m/z*): calc for [M+H]⁺ C₁₃H₁₅Br₂N₄O₄S 480.9181 found 480.9178.

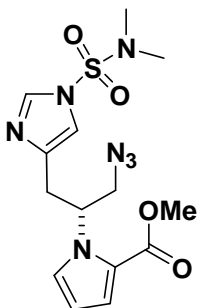
(R)-methyl 1-(1-(1-(N,N-dimethylsulfamoyl)-1H-imidazol-4-yl)-3-hydroxypropan-2-yl)-1H-pyrrole-2-carboxylate (89):



In a round bottom flask compound **77** (160 mg, 0.49 mmol) was dissolved in anhydrous methanol (10 mL) under N₂ atmosphere and to the reaction mixture freshly generated 0.23 M sodium methoxide (10.7 mL, 0.23 mmol) was added. The reaction mixture was stirred at r.t. for 8 h. Finally NH₄Cl and water were

added to the reaction mixture and it was repeatedly extracted with EtOAc. The organic layer was dried with anhydrous Na_2SO_4 concentrated and the residue was purified by chromatography on silica gel ($\text{CH}_2\text{Cl}_2/\text{MeOH}$, 98:2) providing **89** as a thick colorless liquid (61 mg, 35%). In addition recovered starting material (80 mg, 50%) was obtained. $[\alpha]_D = -14.1$ ($c = 0.04$, MeOH). $^1\text{H NMR}$: $\delta = 7.80$ (s, 1H), 7.24 (t, $J = 2.4$ Hz, 1H), 6.93 (dd, $J = 3.8, 1.7$ Hz, 1H), 6.86 (s, 1H), 6.15 (t, $J = 3.8$ Hz, 1H), 5.60-5.52 (m, 1H), 3.95 (d, $J = 5.2$ Hz, 2H), 3.79 (s, 3H), 3.16-3.13 (dd, $J = 7.4, 2.4$ Hz, 2H), 2.80 (s, 6H); $^{13}\text{C NMR}$: $\delta = 162.0, 139.9, 136.4, 126.3, 122.1, 118.3, 114.9, 108.8, 64.8, 57.8, 51.2, 38.2, 31.1$; FT-IR (neat, cm^{-1}): 3290, 3121, 1698, 1399, 1176, 1106, 1079, 959, 734, 590; HR-MS (m/z): calc for $[\text{M}+\text{H}]^+ \text{C}_{14}\text{H}_{21}\text{N}_4\text{O}_5\text{S}$ 357.1233 found 357.1242.

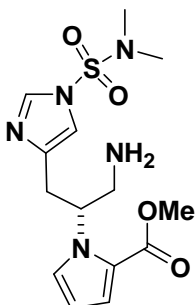
(R)-methyl 1-(1-azido-3-(1-(*N,N*-dimethylsulfamoyl)-1H-imidazol-4-yl)propan-2-yl)-1H-pyrrole-2-carboxylate (91):



In a round bottom flask compound **89** (40 mg, 0.11 mmol) was dissolved in anhydrous CH_2Cl_2 (5 mL) under N_2 atmosphere and to the reaction mixture triethylamine (0.02 mL, 0.13 mmol) was added dropwise after cooling the reaction mixture to 0°C . Then to the resulting reaction mixture $\text{CH}_3\text{SO}_2\text{Cl}$ (0.02 mL, 0.13 mmol) was added dropwise. The reaction mixture was stirred at 0°C for 2 h. Then the solvent was concentrated under vacuum and the crude reaction mixture was dissolved in DMF (2 mL) and to it was added NaN_3 (90 mg, 1.40 mmol). The reaction mixture was heated at 80°C for 8 h. After cooling to r.t. water was added

to the reaction mixture and it was extracted with EtOAc. The organic extract was dried with anhydrous Na₂SO₄, concentrated, and the residue was purified by chromatography on silica gel (CH₂Cl₂/MeOH, 99:1) providing **91** (30 mg, 71%) as a thick yellow liquid. $[\alpha]_D = -3.8$ ($c = 0.02$, MeOH). ¹H NMR: $\delta = 7.77$ (s, 1H), 7.01 (s, 1H), 6.94 (s, 1H), 6.75 (s, 1H), 6.16 (t, $J = 3.8$ Hz, 1H), 5.84-5.68 (m, 1H), 3.79 (s, 3H), 3.75-3.66 (m, 2H), 3.17 (d, $J = 7.2$ Hz, 2H), 2.73 (s, 6H); ¹³C NMR: $\delta = 161.7, 139.5, 136.5, 122.6, 118.7, 114.9, 109.1, 77.3, 54.9, 51.3, 38.1, 31.5$; FT-IR (neat, cm⁻¹): 2950, 2108, 1695, 1387, 1220, 1173, 1106, 959, 725, 593; HR-MS (m/z): calc for [M+H]⁺ C₁₄H₂₀N₇O₄S 382.1297 found 382.1303.

(R)-methyl 1-(1-amino-3-(1-(N,N-dimethylsulfamoyl)-1H-imidazol-4-yl)propan-2-yl)-1H-pyrrole-2-carboxylate (92):

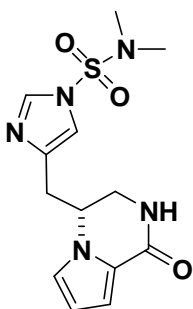


Compound **91** (50 mg, 0.13 mmol) was dissolved in dry methanol (3 mL). To the reaction mixture 10% Pd/C (50 mg) was added. The heterogeneous reaction mixture was stirred at r.t. for 8 h under a hydrogen atmosphere (balloon). The reaction mixture was filtered over Celite and the filter cake was repeatedly washed with hot methanol. The filtrate was evaporated under reduced pressure followed by purification of the residue by chromatography on silica gel (CHCl₃/MeOH, 92:8) furnishing the primary amine compound **92** (30 mg, 65%) as a thick colorless liquid. $[\alpha]_D = 29.7$ ($c = 0.2$, MeOH). ¹H NMR: $\delta = 7.75$ (s, 1H), 7.00 (t, $J = 1.7$ Hz, 1H), 6.88 (dd, $J = 1.7, 3.8$ Hz, 1H), 6.66 (s, 1H), 6.16 (t, $J = 3.4$ Hz, 1H), 5.66-5.55 (m, 1H), 3.78 (s, 3H),

3.18-2.97 (m, 4H), 2.71 (s, 6H); ^{13}C NMR: $\delta = 162.0, 140.1, 136.3, 123.0, 118.3, 118.1, 114.2, 109.1, 103.8, 77.3, 51.2, 46.8, 37.9, 32.3$; FT-IR (neat, cm^{-1}): 3250, 2366, 2334, 1720, 1502, 1449, 1207, 1144, 1010, 737, 708. HR-MS (m/z): calc for $[\text{M}+\text{H}]^+ \text{C}_{14}\text{H}_{21}\text{N}_5\text{O}_4\text{S}$ 356.1393 the expected molecular ion was not observed, but that for the cyclized product (**55**) $[\text{M}+\text{H}]^+ \text{C}_{13}\text{H}_{18}\text{N}_5\text{O}_3\text{S}$ 324.1126 was found 324.1123.

(R)-N,N-dimethyl-4-((1-oxo-1,2,3,4-tetrahydropyrrolo[1,2-a]pyrazin-4-yl)methyl)-1H-imidazole-1-sulfonamide (55**):**

In a round bottom flask **92** (50 mg, 0.14 mmol) was dissolved in dry THF (4 ml)

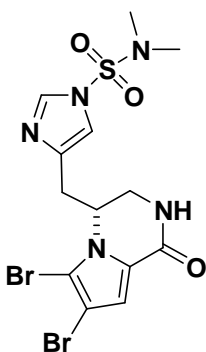


under N_2 atmosphere. The reaction mixture was cooled to $0\text{ }^\circ\text{C}$ and to it 60% sodium hydride (30 mg, 1.4 mmol) was added. The reaction mixture was stirred at r.t. for 8 h. NH_4Cl solution was added and the reaction mixture was repeatedly extracted with EtOAc. The combined extracts were dried with anhydrous

Na_2SO_4 , concentrated, and the residue was purified by chromatography ($\text{CH}_2\text{Cl}_2/\text{MeOH}$, 94:6) providing **55** as a thick colorless solid (34 mg, 58%). m.p. $192\text{-}194\text{ }^\circ\text{C}$. $[\alpha]_{\text{D}} = 17.0$ ($c = 0.4$, MeOH). ^1H NMR: $\delta = 7.86$ (d, $J = 1.4$ Hz, 1H), 6.90 (dd, $J = 4.1, 1.4$ Hz, 1H), 6.76 (s, 1H), 6.55 (t, $J = 2.4$ Hz, 1H), 6.20 (b, 1H), 6.11 (dd, $J = 2.7, 2.4$ Hz, 1H), 4.64-4.62 (m, 1H), 3.92 (dd, $J = 12.7, 4.1$ Hz, 1H), 3.53 (dd, $J = 12.7, 2.4$ Hz, 1H), 3.06 (dd, $J = 14.4, 6.9$ Hz, 2H), 2.81 (s, 6H); ^{13}C NMR: $\delta = 161.0, 139.1, 136.9, 128.4, 123.2, 115.6, 114.0, 109.4, 53.6, 44.3, 38.2, 32.3$; FT-IR (neat, cm^{-1}): 3140, 2934, 1674, 1551, 1460, 1387, 1170, 1088,

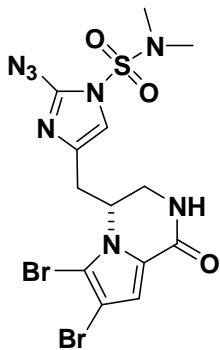
962, 722, 602; HR-MS (m/z): calc for $[M+H]^+$ $C_{13}H_{18}N_5O_3S$ 324.1126 found 324.1130.

(R)-4-((6,7-dibromo-1-oxo-1,2,3,4-tetrahydropyrrolo[1,2-a]pyrazin-4-yl)methyl)-N,N-dimethyl-1H-imidazole-1-sulfonamide (93):



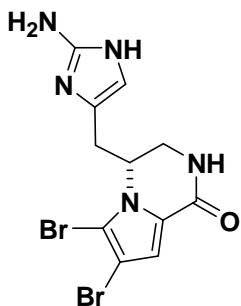
The cyclized compound **55** (90 mg, 0.27 mmol) was dissolved in THF (3.5 mL). The reaction mixture was cooled to 0 °C and to it NBS (100 mg, 0.58 mmol) was added. The reaction mixture was stirred at r.t. for 1 h. The reaction mixture was quenched by addition of aq. $NaHCO_3$ and then the organic layer was separated. The aqueous layer was extracted with EtOAc (3x50 mL), then the combined organic solutions were dried with anhydrous Na_2SO_4 and concentrated. The crude product was purified by chromatography ($CH_2Cl_2/MeOH$, 97:3) providing **93** as a brown solid (117 mg, 88%). m.p 183-185 °C. $[\alpha]_D = 14.0$ ($c = 0.004$, MeOH). 1H NMR: $\delta = 7.86$ (s, 1H), 7.00 (s, 1H), 6.85 (s, 1H), 5.90 (b, 1H), 4.68-4.63 (m, 1H), 3.91-3.85 (dd, $J = 13.1, 4.1$ Hz, 1H), 3.67-3.61 (dd, $J = 13.1, 5.2$ Hz, 1H), 3.11-3.03 (dd, $J = 14.4, 7.9$ Hz, 1H), 3.02-2.94 (dd, $J = 14.4, 5.8$ Hz, 1H), 2.87 (s, 6H); ^{13}C NMR: $\delta = 158.8, 138.6, 137.1, 124.9, 116.1, 115.5, 107.1, 100.6, 53.5, 43.4, 38.2, 31.2$; FT-IR (neat, cm^{-1}): 3180, 2960, 1680, 1545, 1480, 1350, 1210, 1070, 972, 725, 635. HR-MS (m/z): calc for $[M+H]^+$ $C_{13}H_{16}Br_2N_5O_3S$ 479.9341 found 479.9355.

(R)-2-azido-4-((6,7-dibromo-1-oxo-1,2,3,4-tetrahydropyrrolo[1,2-a]pyrazin-4-yl)methyl)-N,N-dimethyl-1H-imidazole-1-sulfonamide (94):



Dibromopyrrole **93** (60 mg, 0.12 mmol) was dissolved in anhydrous THF (3 mL) and the reaction mixture was cooled to -78 °C and 0.5 M LDA (1.22 mL, 4.2 equiv) was added dropwise to the reaction mixture. The reaction mixture was brought to 0 °C and then left stirring for 45 min at 0 °C and then again the reaction mixture was cooled to -78 °C and TrisylN₃ (0.28 g, 1.07 mmol) was added. The reaction mixture was allowed to come to r.t. and then stirred for an additional 1 h, followed by addition of aqueous NH₄Cl (3 mL) to quench the reaction mixture. The organic layer was separated and the aqueous layer was extracted with EtOAc (3x50 mL). The organic solutions were combined, dried with anhydrous Na₂SO₄ and concentrated. The crude product was purified by chromatography (CHCl₃/MeOH, 98:2) giving **94** as a yellow liquid (17 mg, 30%). In addition 30 mg, (50%) of recovered starting material was obtained. ¹H NMR: δ = 7.00 (s, 1H), 6.77 (s, 1H), 5.76 (b, 1H), 4.63-4.60 (m, 1H), 3.86 (dd, *J* = 13.8, 4.1 Hz, 1H), 3.66 (d, *J* = 13.8 Hz, 1H), 2.94 (s, 6H), 2.86 (d, *J* = 7.6 Hz, 2H); ¹³C NMR: δ = 158.5, 140.7, 134.4, 124.5, 117.1, 115.9, 107.4, 100.6, 53.8, 42.7, 38.0, 30.7; FT-IR (neat, cm⁻¹): 2985, 2145, 1649, 1534, 1509, 1457, 1398, 1336, 1181, 1086, 990; HR-MS (*m/z*): calc for [M+Na]⁺ C₁₃H₁₄Br₂N₈O₃SNa is 542.9169 found 542.9177.

(ent)-Cyclooroidin (3):



Conc. HCl (0.003 mL) was added to the solution of compound **94** (22 mg, 0.04 mmol) in methanol (1.2 mL) and stirred at 34 °C for 12 h. Then the mixture was concentrated under vacuum and the residue was neutralized with NaHCO₃ and extracted with EtOAc (3 times). The organic layer were combined, concentrate and purified by column chromatography (CHCl₃/methanol/NH₄OH, 92:5:3) to give the free imidazole (13 mg, 76%) as a thick yellow liquid. This compound (10 mg, 0.02 mmol) was dissolved in methanol (1 mL) and Lindlar catalyst (10 mg) was added and the reaction mixture was stirred for 8 h under a hydrogen atmosphere (balloon). Finally the reaction mixture was filtered through a bed of Celite and then the crude reaction mixture was loaded onto a column as a solid, preadsorbed on silica gel. Then the crude product was purified by chromatography (CH₂Cl₂/MeOH/NH₄OH: 80/15/5). A colorless thick liquid was obtained (5 mg, 74%). $[\alpha]_D = 10.6$ ($c = 0.02$, MeOH). ¹H NMR (CD₃OD) : $\delta = 6.92$ (s, 1H), 6.28 (s, 1H), 4.60-4.56 (m, 1H), 3.75 (dd, $J = 13.4, 4.1$ Hz, 1H), 3.52 (dd, $J = 13.4, 4.1$ Hz, 1H), 2.80 (d, $J = 7.2$ Hz, 2H); ¹³C NMR (CD₃OD): $\delta = 159.6, 149.6, 127.9, 124.3, 115.2, 111.3, 107.3, 100.0, 54.2, 42.1, 29.3$; HR-MS (m/z): calc for [M+H]⁺ C₁₁H₁₁Br₂N₅O 387.9403 found 387.9404.

Part II STUDIES TOWARDS SOME OROIDIN DIMERS

CHAPTER 4

INTRODUCTION

4.1 Introduction

During the last few years the oroidin class of polyheterocyclic, nitrogen-rich alkaloids has received much attention as they rank among the most challenging synthetic targets.^{4,11,56-60} In addition, several family members exhibit substantial biological activity in particular several of the dimeric congeners. Examples of these alkaloids include sceptrin (**103**), the axinellamines A (**99**) and B (**100**), palau'amine (**105**), and ageliferin (**104**). These marine natural products appear to arise from one precursor, oroidin (**1a**) which was first identified in 1971.⁶⁰ These densely functionalized, highly oxidized, polycyclic alkaloids are believed to obtain from dimerization of oroidin and then followed by consecutive functionalizations (Figure 4.1).^{3,11,62,63} However, to date there is only one reported biosynthetic study and so the precise origin of these alkaloids remains to be fully elucidated.¹² Recently the palau'amine (**105**) structure was revised and thus as a consequence, the synthetic approaches to this molecule required redesign. In the original report the stereochemistry of the junction of the two fused five-membered rings had been assigned as *cis*.⁶⁴ However, a thorough spectroscopic re-investigation^{11,65} was recently complemented by synthesis (*vide infra*),^{66,67} suggesting that the C-11/C-12 ring fusion in **105** to be in the thermodynamically less stable *trans* configuration (Figure 4.1). In addition the stereochemistry of C-

17 chloro bearing center was revised such that palau'amine now shares a common stereochemical arrangement around the carbocyclic E ring with the axinellamines and massadine families. As a result, it appears that all of these systems, **99-101** and **105** may be derived from a common, late stage biosynthetic intermediate.

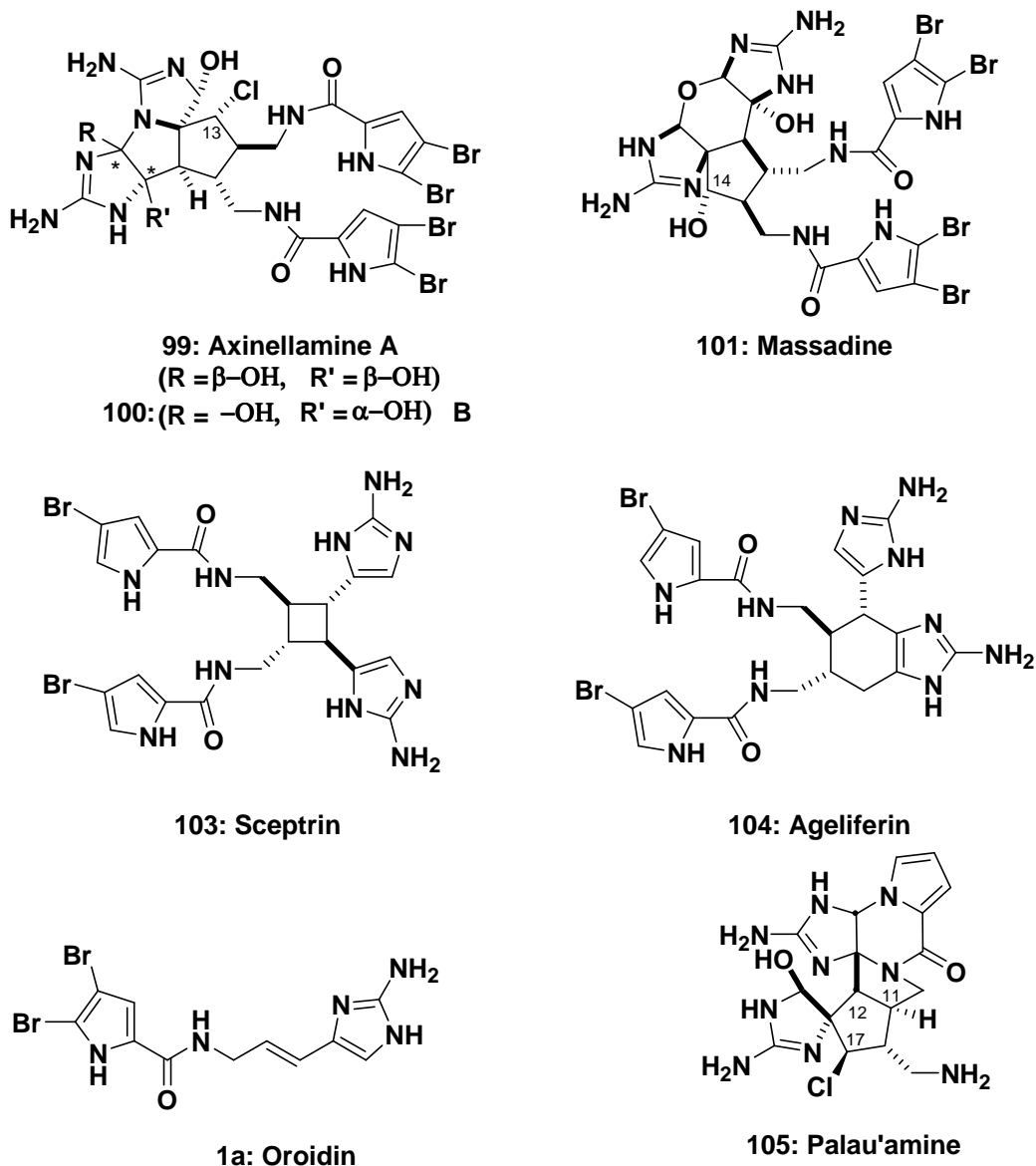
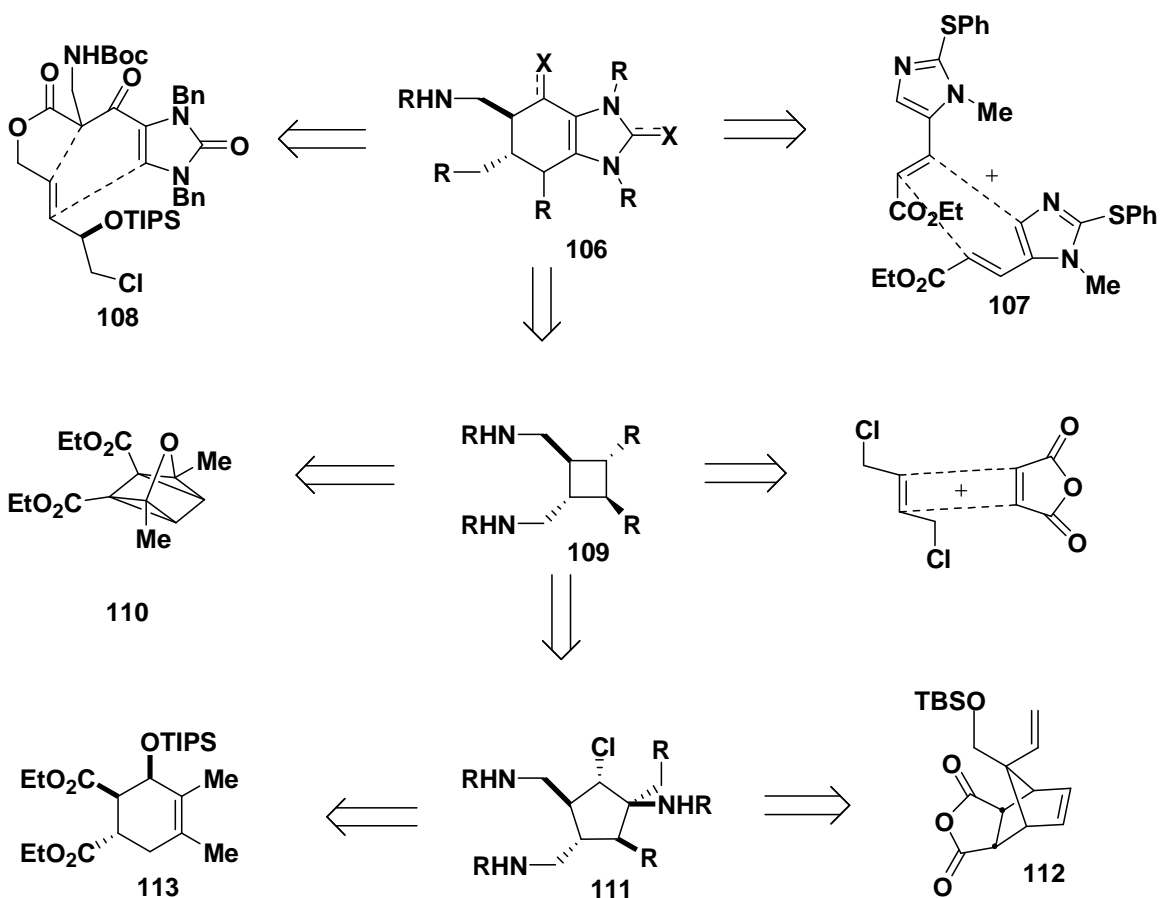


Figure 4.1 Selected Oroidin alkaloids of marine origin.

All of these pyrrole–imidazole alkaloids feature a central four-, five-, or six-membered carbocyclic ring decorated with different arrangements of the remaining heterocycles. Due to their unique structural patterns many synthetic efforts have been pursued which in turn have led to distinct solutions for assembly of each of the scaffolds (Scheme 4.1).⁵⁹ The six-membered ring of the ageliferins **106** was proposed to arise from a [4+2] cycloaddition in one early biosynthetic hypothesis.⁶² This was first implemented by Ohta et al. (**107**→**106**) although this results in the synthesis of a non-natural occurring material.⁷⁰ Our group has pursued a related, but intramolecular strategy, towards the ageliferins.^{10a} Chen and Tan developed a Mn(III)-promoted radical cascade annulation from the imidazolone **108**.⁷¹ Baran et al. successfully demonstrated that ageliferin **104** can be obtained from sceptrin (**103**) through a double ring-enlargement through a vinylcyclobutane rearrangement of sceptrin (**103**) under high temperature conditions in their total synthesis of **104**.⁷² Starting from (*E*)-1,4-dichloro-2-butene and maleic anhydride⁷³ or by fragmentation of the photochemically accessible oxaquadricyclane **110** the sceptrin scaffold **109** has been synthesised by [2+2] photocycloadditions.^{24,74} The fully substituted five-membered-ring scaffolds **111** present in the axinellamines and palau'amines is the biggest challenge as of now. For these molecules, many linear biomimetic pathways have been proposed, including ring enlargements of a four-membered sceptrin-like precursor **109**, or oxidative ring contractions of six-membered ageliferin-like precursors **106**.^{3,4,11,56-59, 62,63} Experimentally the ring-enlargement of **109**→**111** still waits to be realized whereas ring-contractions **106**→**111** have

been executed with considerable success.^{10b,76} However, “abiotic” syntheses of scaffolds **111** have proven to be the most successful to date.^{4,11,56-59} The first enantioselective synthesis of the axinellamine core **111** by desymmetrization of anhydride **112** was reported by Carreira et al. in 2000.^{4,76,77,63} The first total synthesis of the axinellamines was completed by Baran et al.⁷⁸ (**99**, **100**) in racemic form by using a ring opening-ring-contraction of the cyclohexene **113**.⁷⁸

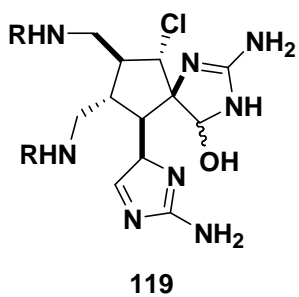
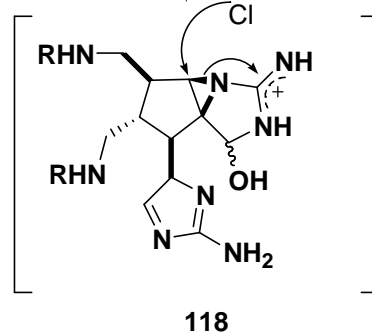
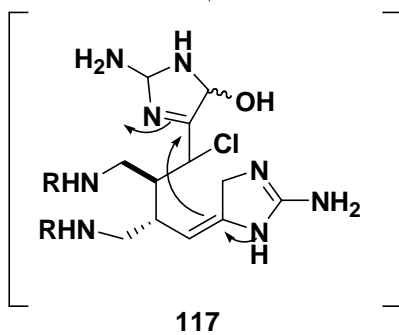
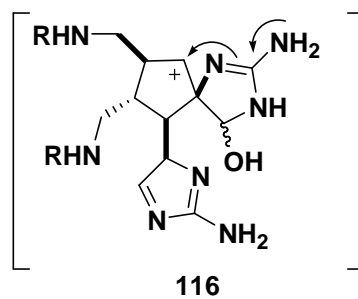
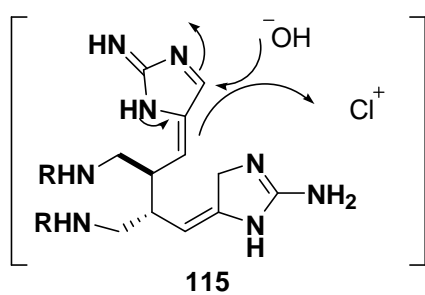
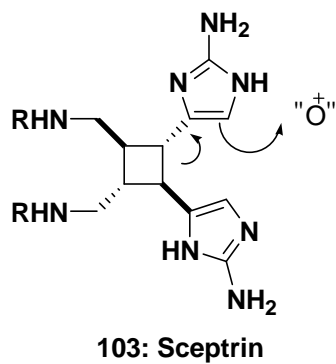
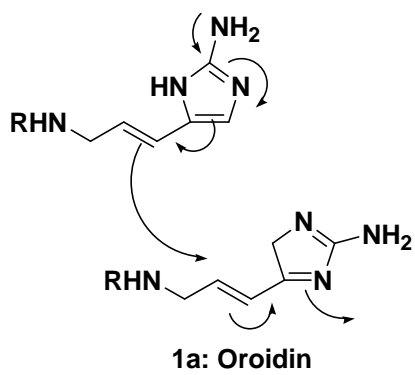


Scheme 4.1 Synthetic routes to the core structures of related oroidin alkaloids.

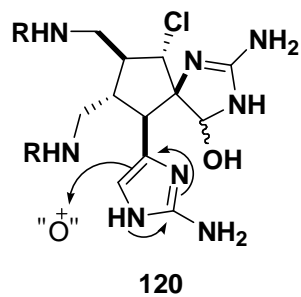
4.2 Three possible biosynthetic pathways to “preaxinellamine”

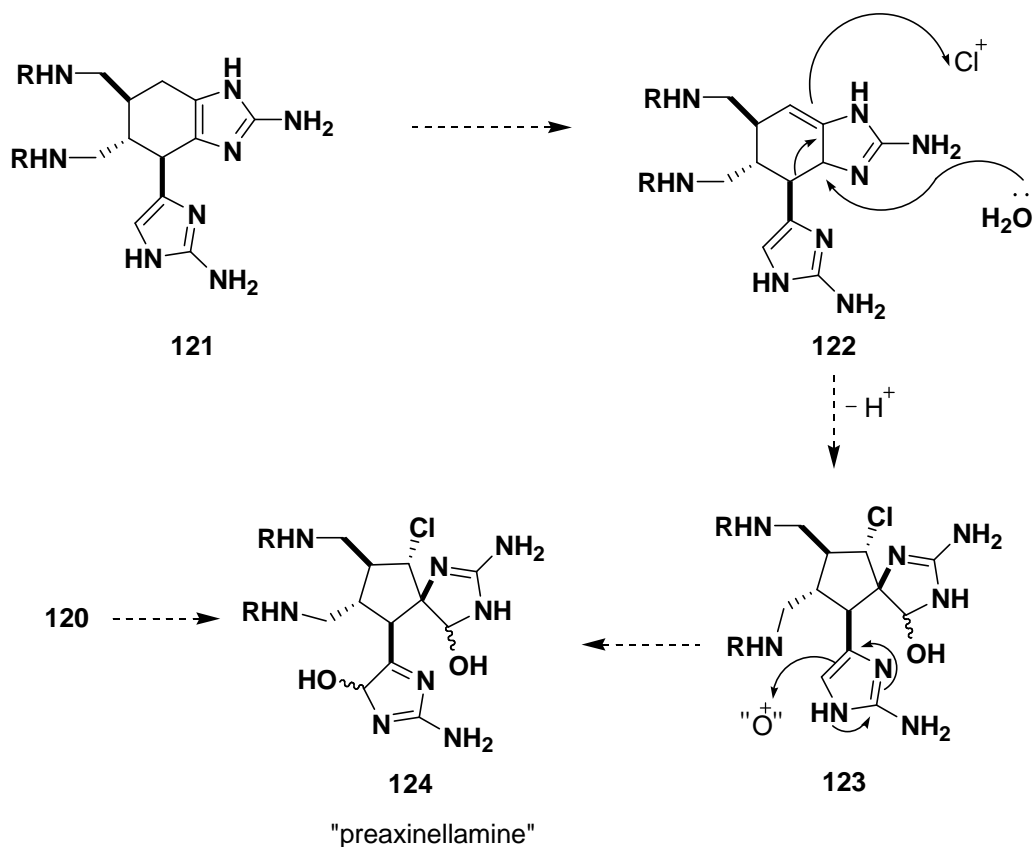
Scheme 4.2 depicts three possible biosynthetic pathways to “preaxinellamine” (**124**). In the first proposal (linear), oroidin-like molecules **1a** undergo an enantio- and diastereocontrolled dimerization to form **115**.^{3,11} **119** was formed by reacting

with an electrophilic chlorine source, a hydration reaction and then cyclization via **117**. After tautomerization and oxidation of the remaining aminoimidazole, “preaxinellamine” (**124**) is formed. An alternative proposal was recently put forth which involves a ring expansion of the cyclobutane nucleus of sceptrin (**103**).⁷² Since extremely high concentrations of **103** have been found in many of the organisms that produce ageliferin (**104**)^{80,81} (compound **103** has also been isolated from the same organism as palau’amine (**105**)⁶²) has led to the proposition that **103** is a possible biosynthetic precursor of the other more-complex members of this family. The proposal was that exposure of **103** to an oxidant leads to the formation cation **116**. Cation-initiated expansions of cyclobutanes to cyclopentanes have good precedent in the literature,⁸² and have been implicated in the biosynthesis of hirsutene terpenes from illudanes.⁸³ Compound **120** can be formed by attack of a chloride ion, possibly through the intermediacy of aziridine **118**. The third pathway (ring contraction) bears its conceptual roots in the earlier hypothesis for the genesis of palau’amine (**7**).⁶² **124** can be obtained by tautomerization of ageliferin (**104**) followed by reaction with an electrophilic chlorine source, ring contraction, and hydration. Both baran et al. and our lab have recently demonstrated that the ring-contraction pathway can lead to axinellamine-like structures from **104** but with the incorrect stereochemistry.^{75,94}



taut





Scheme 4.2 Biosynthetic pathways of marine Oroidin alkaloids

4.3 Carreira's approach towards Massadine

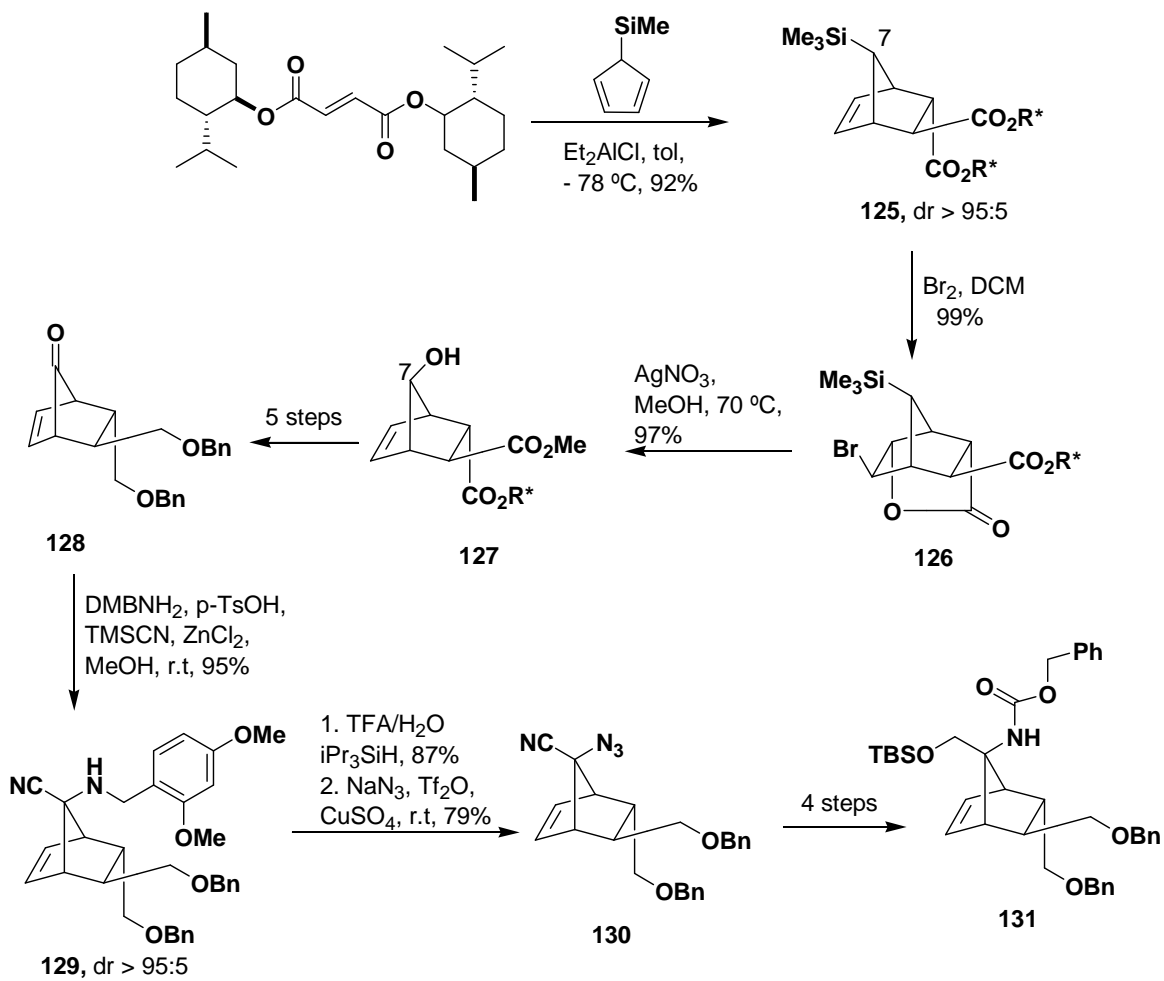
Carreira et al. have developed a synthetic route^{77,84} to access the fully substituted cyclopentane ring of the massadine-type alkaloids.⁸⁴ The goal was to produce an enantiopure norbornene intermediate as a conformationally rigid precursor to the cyclopentane and then to elaborate functionality at the C7 position (norbornene numbering) in a chemo- and diastereoselective manner followed by oxidative cleavage of the olefin to provide the polysubstituted cyclopentane. Their synthesis started (Scheme 4.3)^{77,84} with a diastereoselective cycloaddition of 5-(trimethylsilyl)cyclopentadiene with a chiral, non-racemic fumarate to obtain the bicyclo[2.2.1]heptane efficiently. The C7 position is then oxygenated through bromo-lactonization of **125**, and then followed by an Ag (I)-

mediated rearrangement of the C7-silane to generate the inverted alcohol **127** via methanolysis of the putative lactone intermediate. The alcohol was converted to ketone **128** over 5 steps from which a single amino-nitrile isomer **129** was obtained by Strecker reaction. The diastereoselectivity was achieved due to steric shielding of one diastereoface of the iminium ion by the bulky *exo*-oriented benzyloxymethylene. The silyl ether **131** was obtained by reduction of azidonitrile **130**, which was in turn generated by Cu (II)-catalyzed diazotransfer. Repeated attempts at reduction of variously protected amino-nitrile substrates led to decomposition.

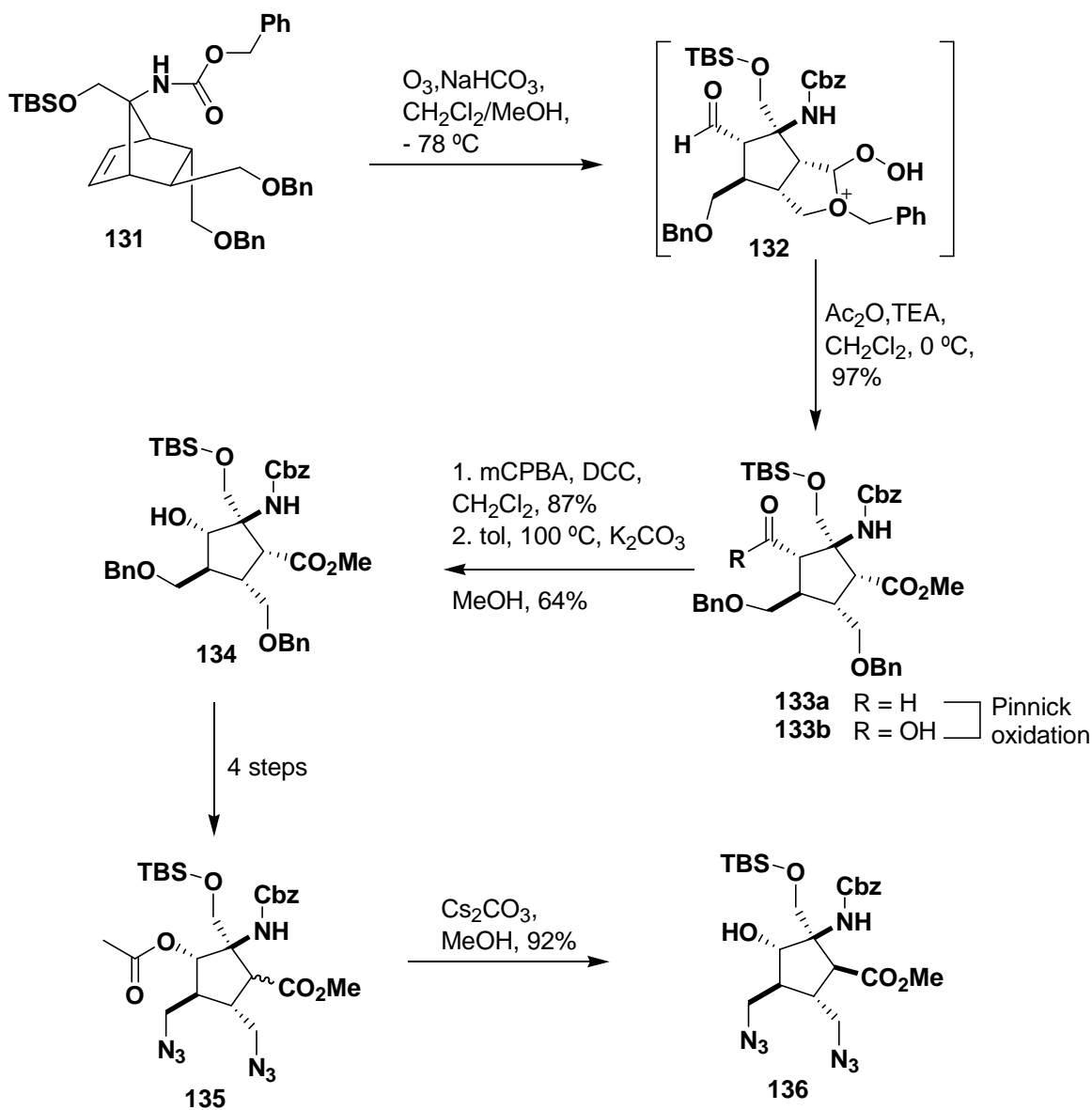
In (Scheme 4.4), ozonolysis provided regioselectively the ester/aldehyde **133a** in excellent yield. The putative oxonium species **132** was generated by anchimeric participation of one benzyloxy group in the rearrangement of the primary ozonide followed by methanolysis and dehydration to **133a**.

By generating the unsymmetrical diacyl peroxide from carboxylic acid **133b** using DCC and peracid, “carboxy-inversion” (**133b**→**134**, Scheme 4.4)⁸⁶ was realized. The alcohol **134** was obtained by heating the intermediate to promote rearrangement to the mixed carbonate and then subsequent methanolysis. A diastereomeric mixture of diazides **135** was obtained in four additional steps.

The massadine core fragment **136** was obtained in an enantioselective fashion by using tandem acetate saponification/ester epimerization with Cs₂CO₃ (the massadine intermediate was synthesized by 24 linear steps).



Scheme 4.3 Carreira's enantioselective synthesis of a massadine D ring precursor.

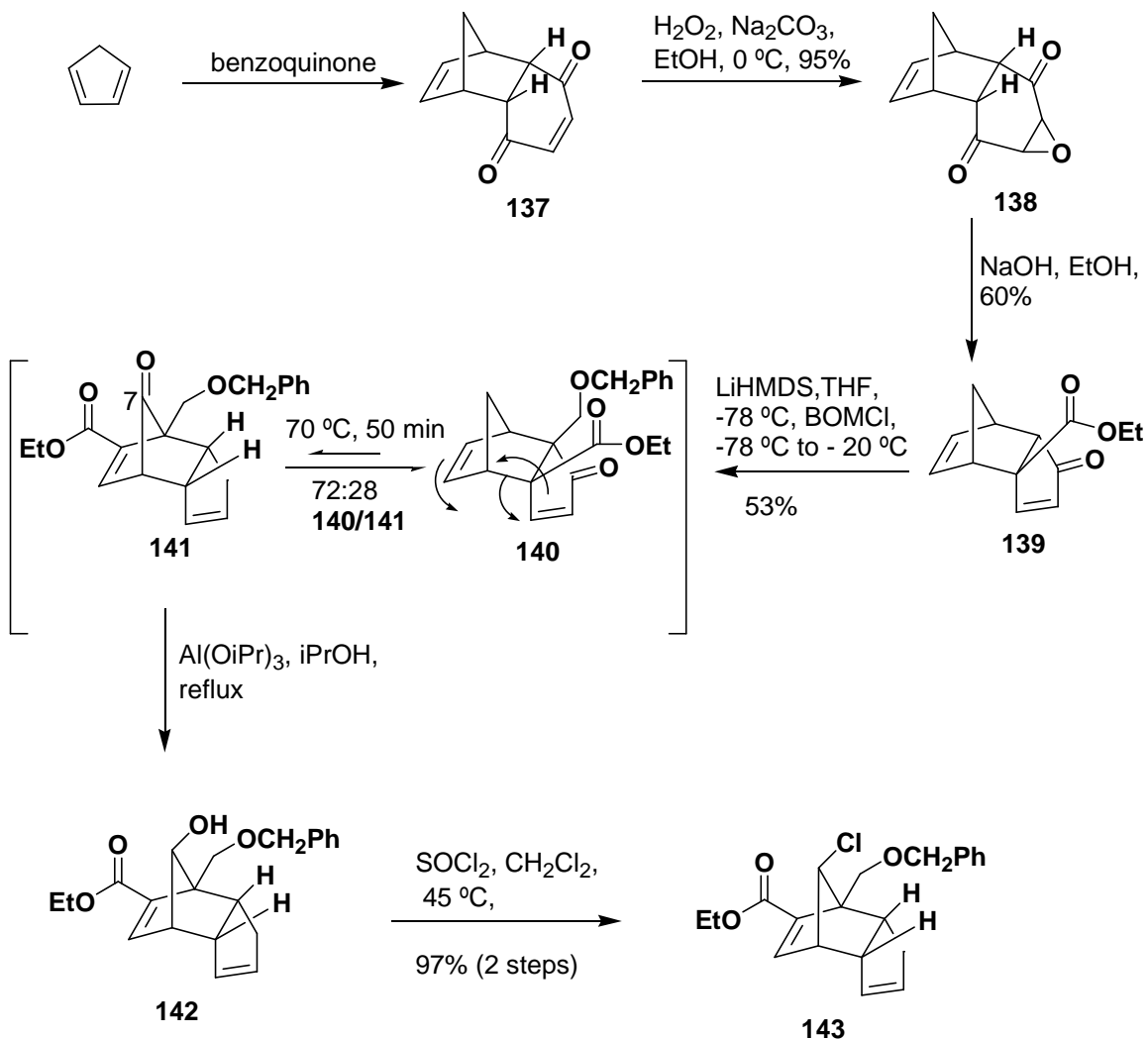


Scheme 4.4 Completion of D-ring-core fragment of massadine.

4.4 Gin's approach towards Palau'amine

Gin and co-workers synthesized the chlorocyclopentane core palau'amine by using a known [3,3]-sigmatropic rearrangement of bridged tricyclodecadienes (e.g. **140**, Scheme 4.5)⁸⁷ in order to obtain a C7-functionalized norbornene intermediate suitable for elaboration to the chlorocyclopentane core of **105**. The racemic Diels–Alder adduct **137** was epoxidized chemoselectively providing a

keto-epoxide which can be subjected to a Favorskii-type ring-contraction. The rearrangement proceeds by enolization followed by intramolecular oxirane ring-opening and finally dehydrative ring-contraction to provide the enone/ester **139** in good yield.

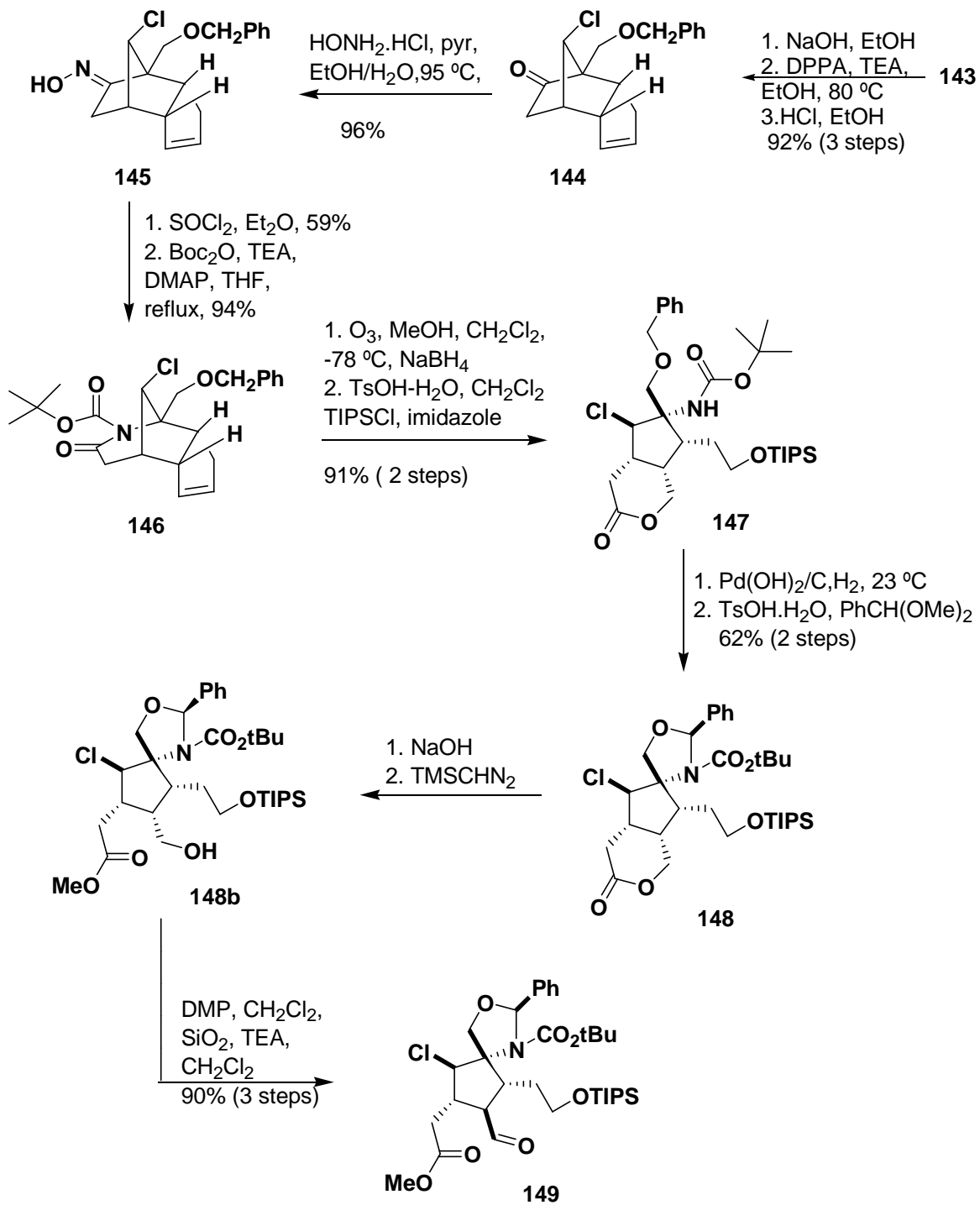


Scheme 4.5 Gin's synthesis of bridged lactam intermediate

Ketone **140**, that bears the 1,5-diene system, and can equilibrate via a [3,3]-sigmatropic rearrangement (**140**→**141**) was obtained by α -alkylation as described by Woodward.⁸⁸ This process provides a latent chloride functionality at C7 of norbornene **141** in the form of a ketone. The **140/141** mixture in a

thermodynamic ratio of 72:28 was reduced under Meerwein–Verley–Ponorf conditions, advancing selectively intermediate **141** through rearrangement and reduction. The efficiency of the above reaction depends on exploitation of the Curtin–Hammett principal of the dynamic **140/141** system. The alcohol **142** was chlorinated with net retention of configuration providing chloride **143**.

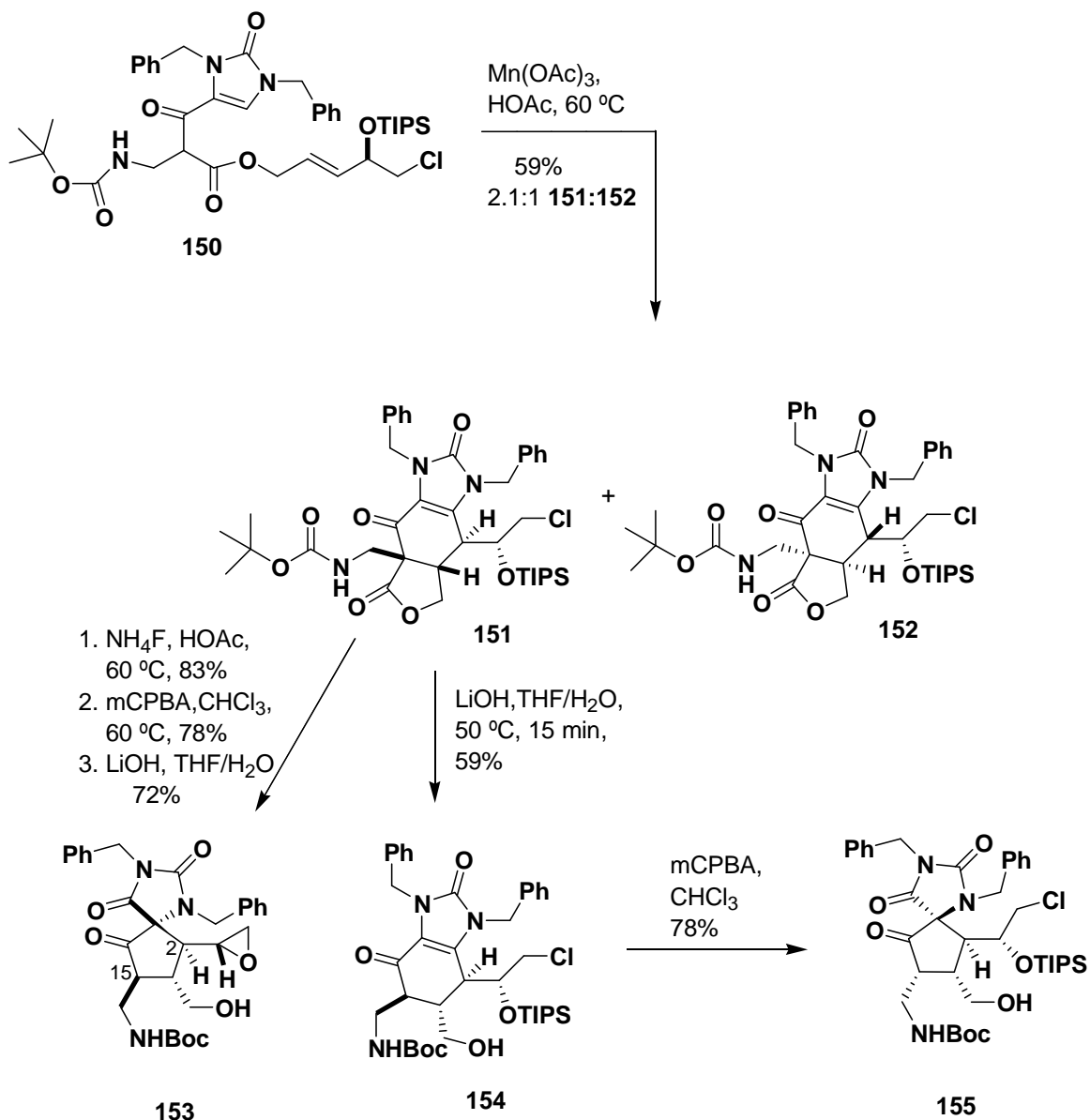
The ketone **144** was generated by an ene-carbamate Curtius rearrangement. Beckmann ring expansion of the oxime initiated with thionyl chloride was completely regioselective resulting from migration of the more substituted alkyl group to provide the lactam **146** after Boc protection. Reductive ozonolysis (Scheme 4.6) of the cyclopentene **146** and subsequent intramolecular alcoholysis of the imide gave **147** after protection of the primary alcohol as the silyl ether. In two additional steps oxazoline **148** was generated and then the lactone moiety was chemoselectively hydrolyzed and esterified with TMSCHN₂ to afford **148b**. Finally, aldehyde **149** was obtained by oxidation of the alcohol followed by a epimerization as a single diastereomer (19 steps from cyclopentadiene).



Scheme 4.6 Synthesis of diastereomeric Palau'amine chlorocyclopentane core.

4.5 Chen's Mn(III) mediated oxidative strategy towards oroidin alkaloids.

A Mn(OAc)₃ promoted oxidative radical heterobicyclization of β-keto esters tethered to unsaturated N-heterocyclic ring systems was developed by Chen and coworkers (Scheme 4.7).⁷¹ By a cascade bicyclization reaction imidazolone **150** undergoes single electron oxidation to provide an electrophilic α-radical^{91,92} that cyclizes by a 5-*exo*/6-*endo* pathway to generate **151** and **152** when it is treated with Mn(OAc)₃ in warm acetic acid. In this single step two C–C bonds and three contiguous stereocenters are established; as only two of a possible eight diastereomers are obtained. The alcohol **154** was then obtained by decarboxylation of the lactone **151**. This intermediate was converted to the spirocyclic cyclopentanone-hydantoin **155** by oxidative rearrangement with concomitant epimerization of the methylene amino-bearing stereocenter. Starting from **151** diastereomeric cyclopentanone **153** can also be prepared by deprotection of the TIPS group and then the resulting lactone was oxidized with mCPBA. The relative stereochemistry of C2 and C15 of **153** was epimeric to the massadine/palau'amine core-ring fragment.



Scheme 4.7 Chen's Mn(III) mediated strategy to access core skeleton of oroidin alkaloids.

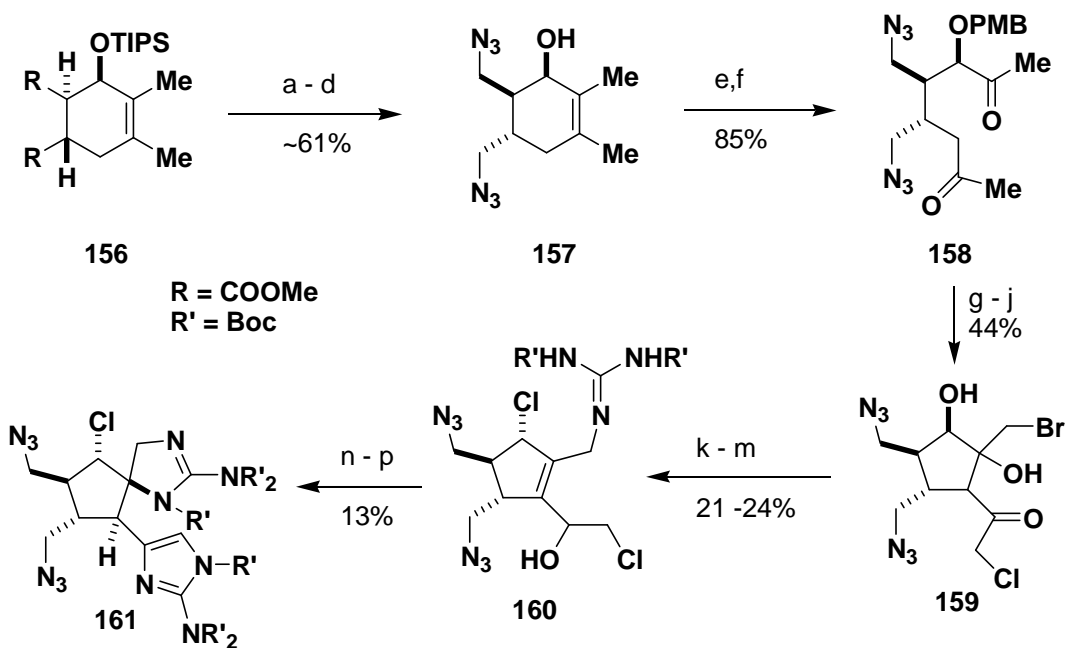
4.6 Baran's total synthesis of the axinellamines

The first total synthesis of the axinellamines was reported by Baran et al. in 2008 (Scheme 4.8).⁷⁸ First, the racemic Diels–Alder product **156** was transformed to diazide **157** in four standard conversions.⁷⁹ Protection of the secondary alcohol as the PMB ether was followed by ozonolysis provided diketone **158**, which was

α,ω -dibrominated via the bis-silylenol ether. Intramolecular aldol reaction under solvent-free conditions provided the cyclopentane derivative. Exchange of the more reactive bromide for a more stable chloride was achieved by treatment with LiCl, and deprotection of the PMB-ether gave diol **159**. The C17-chloro substituent was installed by eliminating the tertiary hydroxy group and the secondary hydroxy group was displaced by Cl⁻ using SO₂Cl₂ in a one-pot reaction. After Luche reduction of the enone, chemoselective displacement of the bromide substituents was achieved with protected guanidine (**159**→**160**). Upon reoxidation to the enone, spirocyclization occurred produced two diastereomers (1.3:1). Formation of the major (and correct) diastereomer was favored by high temperatures. The 2-aminoimidazole moiety was then introduced by displacement and in situ condensation with Boc-guanidine, providing spirocycle **161** after derivatization (Boc₂O) and purification. The protected axinellamine precursor **161** with two heterocycles already installed was achieved in 16 steps from **156** (overall yield 0.7%).

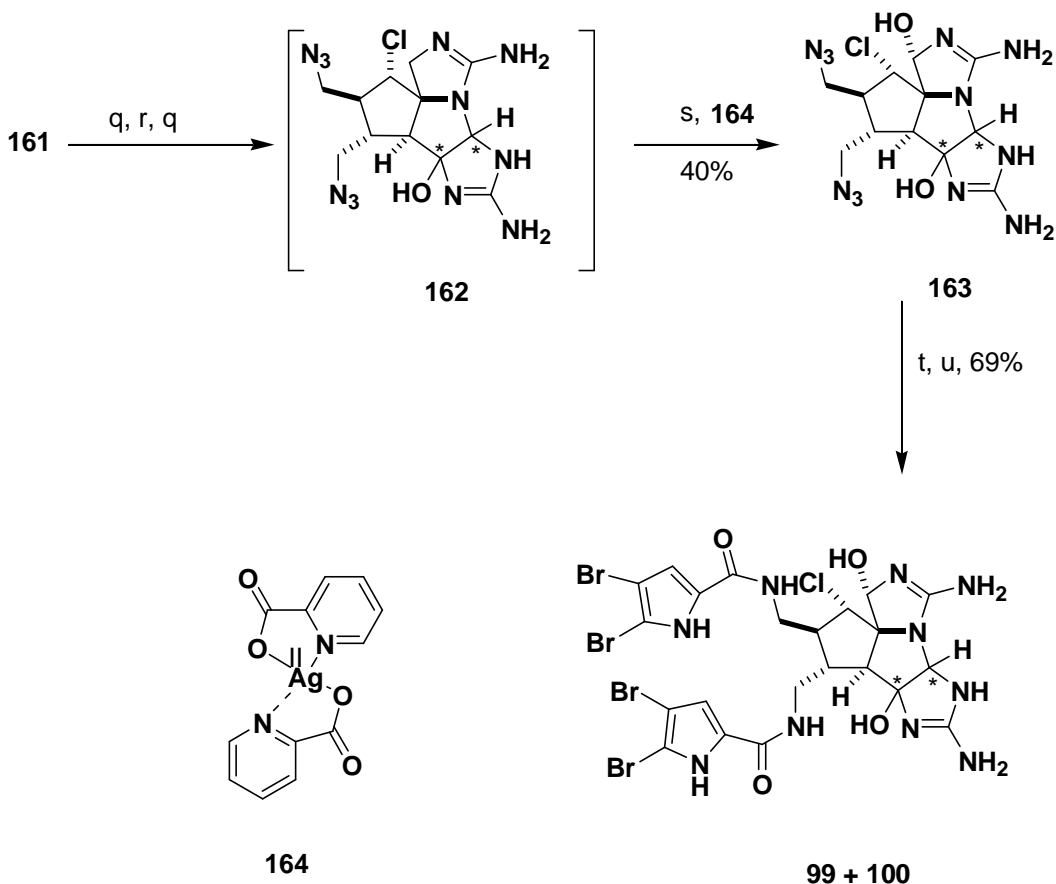
Compound **161** was separated from the axinellamine target connectivity by only two oxidations (Scheme 4.9).⁷⁸ Baran et al. realized that the aminoimidazole 4,5-double bond might be selectively oxidized,⁵⁷ and indeed found that after Boc deprotection of **161** the respective diol could be formed by DMDO oxidation, which on treatment with TFA condensed to carbinolamine **162**. The tetracycle **163** was obtained by chemoselective oxidation with the Ag (II) complex **164**, in only four steps and 40% yield from **161**. Finally the axinellamines **99** and **100** were obtained by reduction of the azide groups followed by acylation with

suitable pyrrole building blocks. The first total synthesis of **99** and **100** was thereby completed in 22 steps from **156** (overall yield 0.2%).



Scheme 4.8 Diastereoselective synthesis of the axinellamine core.

a) LiAlH_4 , THF; b) MsCl , pyridine; c) NaN_3 , DMF, 100 °C; d) TBAF; e) PMBCl , NaH, DMF; f) O_3 , MeOH; g) TMSOTf , EtNiPr_2 , then NBS; h) SiO_2 , no solvent, 47 °C; i) LiCl , DMF; j) 10% TFA; k) SO_2Cl_2 , 2,6-lutidine, CH_2Cl_2 ; l) NaBH_4 , CeCl_3 , MeOH; m) $\text{N,N}'\text{-bis-Boc-guanidine}$, DBU, DMF; n) IBX, benzene, reflux; o) Boc-guanidine , THF, reflux; p) Boc_2O , NEt_3 , cat. DMAP, CH_2Cl_2 . R = COOMe, R' = Boc, PMB = para-methoxybenzyl, MsCl = methylsulfonyl chloride, TBAF = tetrabutylammoniumfluoride, TMS = trimethylsilyl, NBS = N-bromosuccinimide, TFA = trifluoroacetic acid, DBU = 1,8-diazabicyclo-[5.4.0]undec-7-ene, IBX = o-iodoxybenzoic acid.



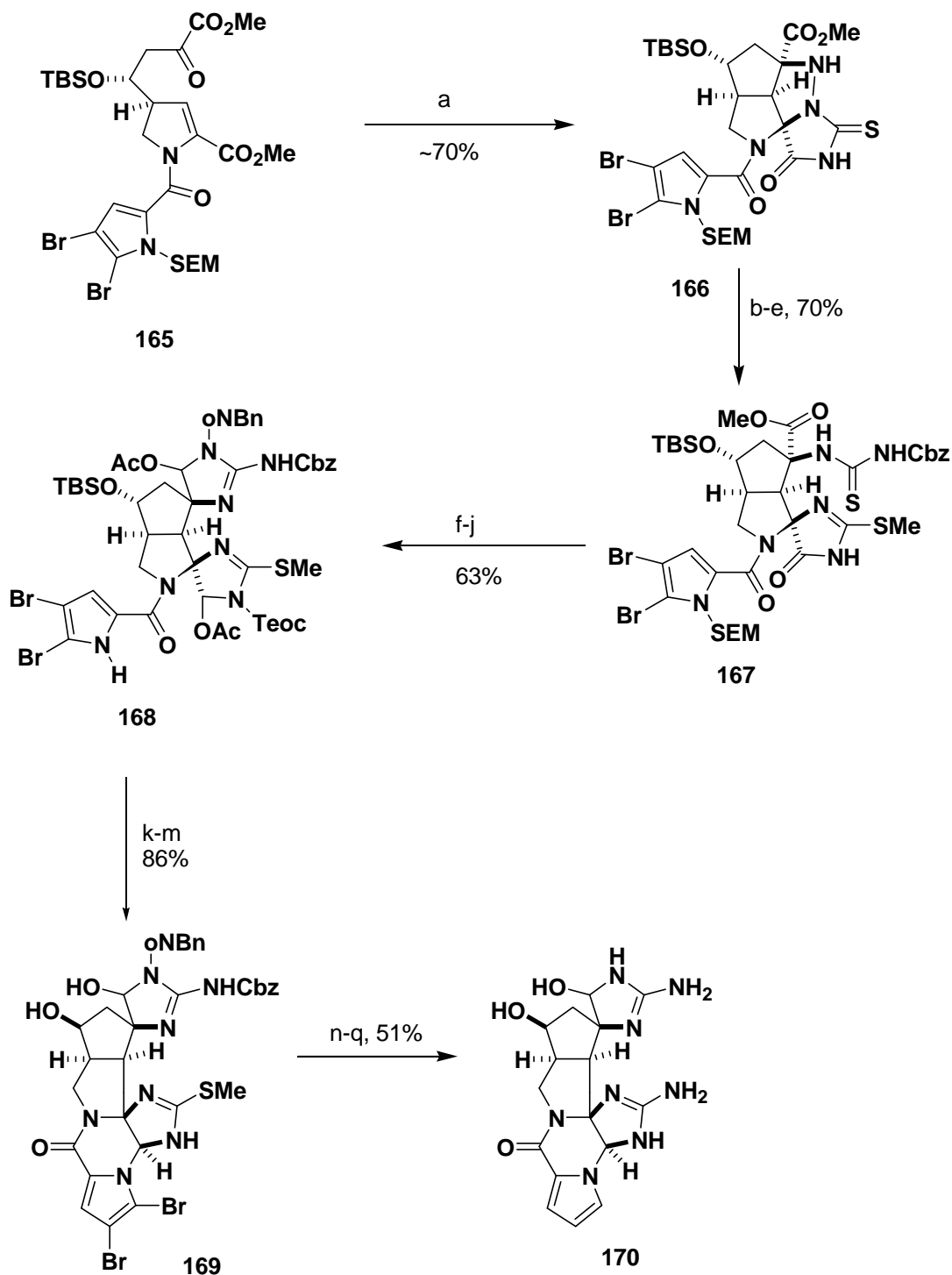
Scheme 4.9 Completion of the axinellamine total synthesis

q) 67% TFA; r) DMDO, H₂O, 0 °C; s) H₂O, 50 °C; t) 1,3-propanedithiol, NEt₃, MeOH; u) 4,5-dibromopyrrole-2-yl-trichloromethyl ketone, EtNiPr₂, DMF, 45 °C. DMDO = 2, 2-dimethyldioxirane.

4.7 Synthetic studies towards Palau'amine by Overman

Overman et al. recently provided a synthetic strategy towards the original stereochemistry of **105** (Scheme 4.10). The key steps towards the synthesis⁶⁷ of the *epi*-palau'amine scaffold **170** include a bicyclization, which was achieved by an intramolecular 1,3-dipolar cycloaddition, providing tetracycle **166** from the dihydropyrrole **165** in 70% yield.⁶⁶ Three contiguous stereocenters were fixed by this elegant transformation, two of them quaternary. The thiourea **167** was obtained by cleaving the N-N bond, thiohydantoin protection and amine acylation.

Compound **167** was then transformed into protected imidazolone, doubly reduced, and protected to give the bisaminal **168**. The Teoc and TBS protecting groups were removed by using TBAF and also closure of the ketopiperazine ring was obtained. The hemiaminal **169** was obtained by inversion of the secondary alcohol followed by an oxidation–reduction sequence. Conversion of the isothiourea into the guanidine and deprotection yielded the *cis*-configured compound **170** in 17 steps and 14% total yield from **165** (ca. 31 steps and 2.4% yield from monoprotected 2-butene-diol), which was found by NMR spectroscopy to differ significantly from natural palau'amine (**105**).



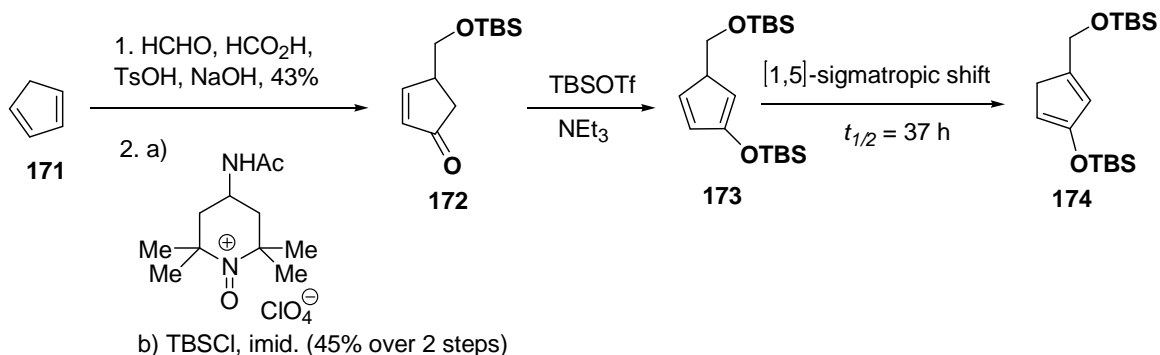
Scheme 4.10 Synthesis of the fully elaborated *epi*-palau'amine scaffold.

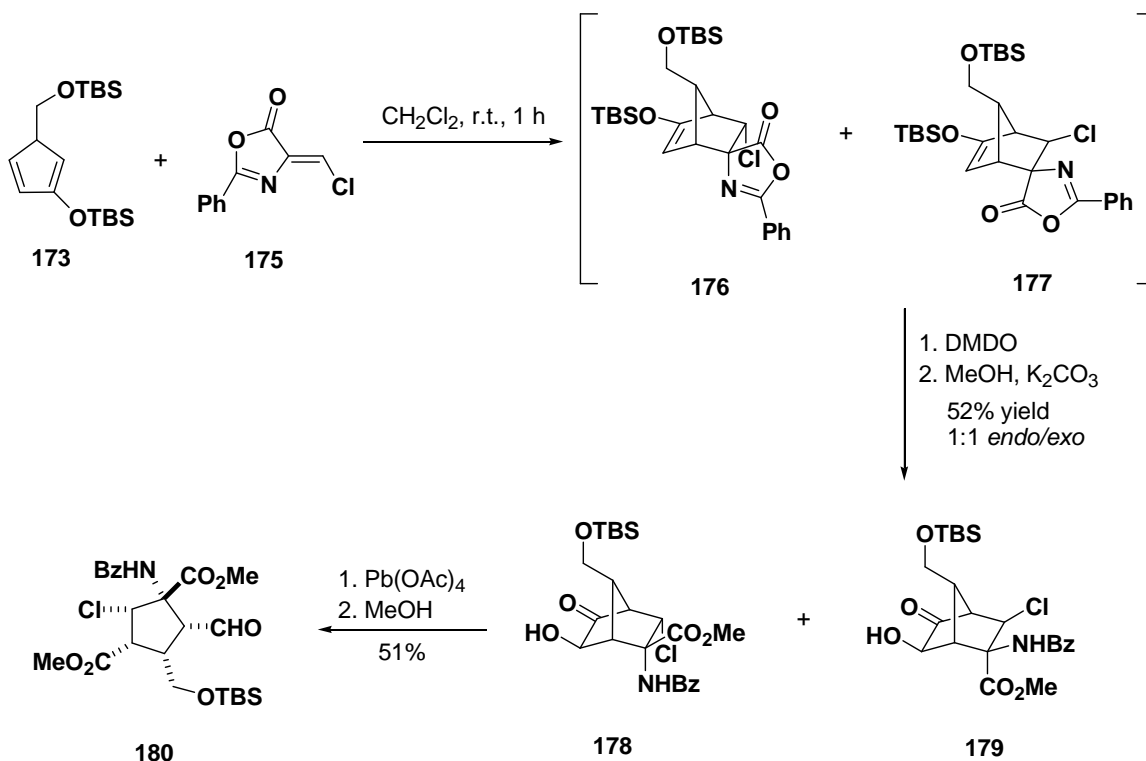
a) Thiosemicarbazide, EtOH, 110 °C; b) SmI_2 , THF/MeOH; c) MeI, EtNiPr_2 , DMAP, CH_2Cl_2 ; d) TeocCl, EtNiPr_2 , CH_2Cl_2 ; e) Cbz-NCS, CH_2Cl_2 ; f) EDC, oNBn-NH₂, EtNiPr_2 , CH_2Cl_2 ; g) 10% TFA; h) TeocCl, EtNiPr_2 , CH_2Cl_2 ; i) NaBH_4 , MeOH/THF; j) Ac_2O , pyridine, DMAP; k) TBAF, THF; l) IBX,

DMSO; m) NaBH₄, MeOH, 0 °C; n) mCPBA, CH₂Cl₂; o) NH₃, CH₂Cl₂, -78 °C; p) hν, dioxane; q) H₂, Pd/C, aq dioxane. SEM = 2-(trimethylsilyl)ethoxymethyl, Teoc = trimethylsilylethoxycarbonyl, oNBn = ortho-nitrobenzyl, NCS = N-chlorosuccinimide, MCPBA = m-chloroperoxybenzoic acid.

4.8 Synthetic studies towards Palau'amine by Gleason

Gleason and co-workers have demonstrated that 2-siloxy substitution markedly stabilizes 5-substituted cyclopentadienes almost by 30-fold at 23 °C relative to 5-methylcyclopentadienes⁶⁸ towards [1,5]-sigmatropic shifts (e.g. **173**→**174**, Scheme 4.11). This allows for productive application of similar dienes in Diels-Alder cycloaddition reactions performed at room temperature. They showed that diene **173** undergoes a variety of [4+2] reactions exhibiting different levels of *endo/exo* selectivity.^{68,69} Cycloaddition reaction between diene **173** and **175** at room temperature gave 1:1 mixture of cycloadducts (**176** and **177**) which were directly subjected to DMDO oxidation followed by methanolysis to give hydroxy ketones **178** and **179**. After separating **178** from **179**, it was oxidatively ring-opened to deliver fully substituted cyclopentane **180**.



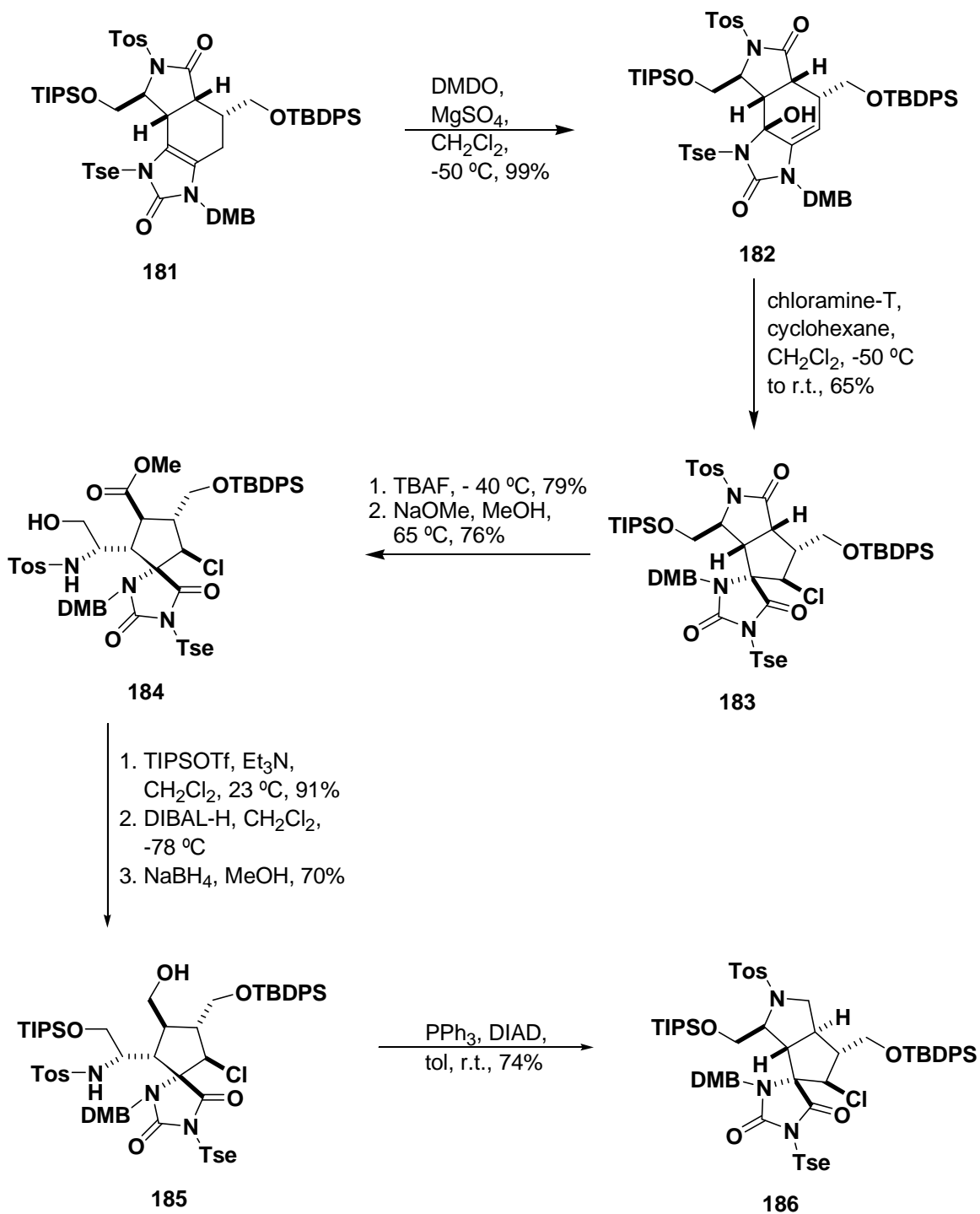


Scheme 4.11 Highly functionalized chloropentane core of palau'amine

4.9 Synthetic studies towards Palau'amine by Romo

Romo and co-workers^{52,76,93} have demonstrated that the palau'amine core skeleton can be obtained from Diels–Alder adducts such as **181** (Scheme 4.12) by a two-step oxidation/chlorination/ring-contraction sequence.^{75,96} In this sequence the imidazolone **181** (prepared via a Diels–Alder reaction) is first subjected to DMSO oxidation at low temperature providing the allylic alcohol **182** in excellent yield as a single diastereomer. Treating **182** with an electrophilic chlorinating agent resulted in the incorporation of the key chlorine atom on the convex face of the tricyclic systems. Then a 1,2-alkyl migration on this iminium species yielded the ring-contracted spirocycle **183** in good yield on a gram scale.⁹³ The use of cyclohexene as a HOCl buffer suppresses formation of aromatized over-oxidation by-products. The all *trans* relative stereochemistry

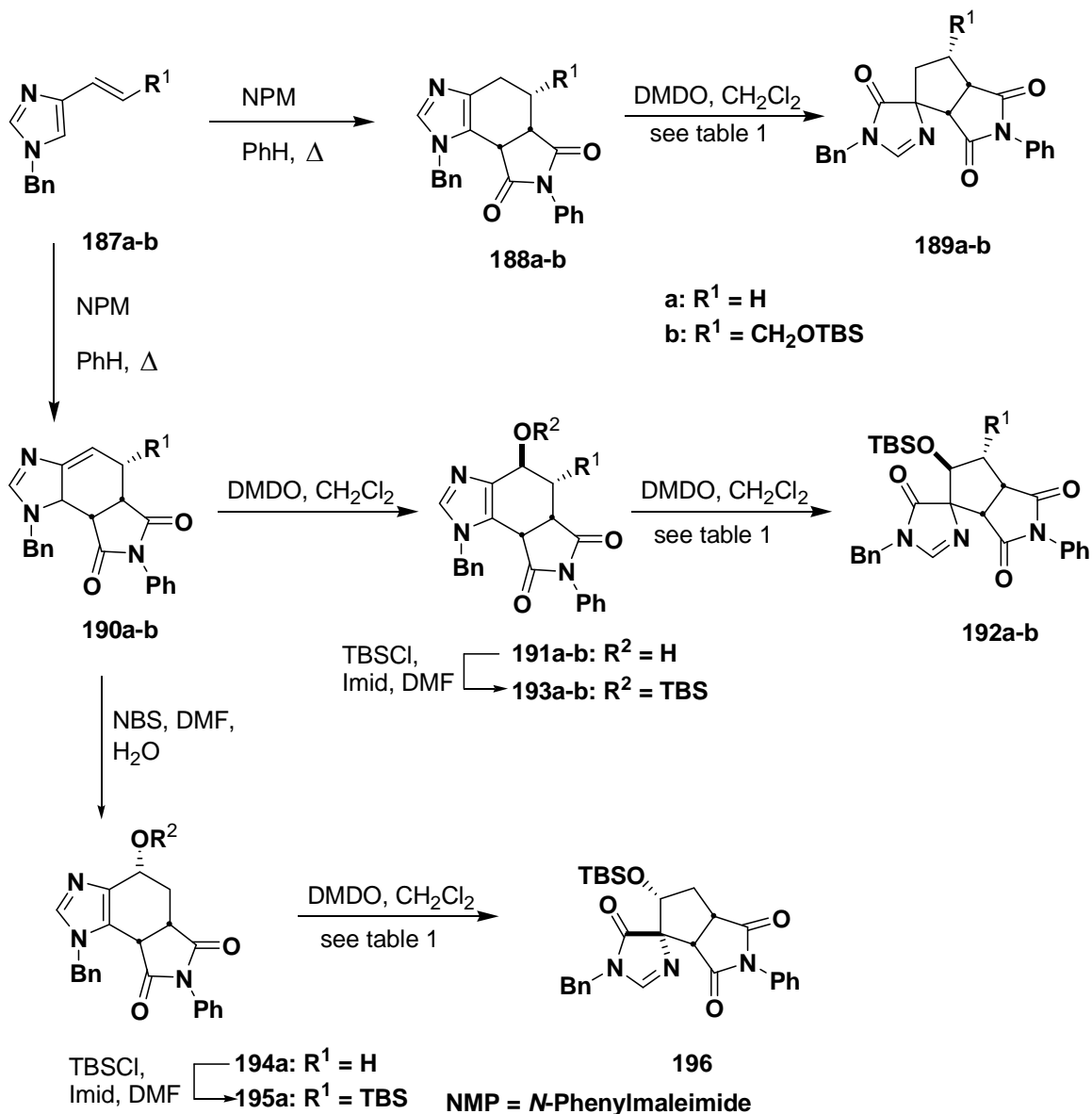
common to oroidin alkaloids **101** and **105** was achieved by cleaving the lactam with concomitant epimerization of the methoxy carbonyl-containing stereocenter. The relative configuration of this *anti*-chlorocyclopentane **184** was verified crystallographically.⁵² The azabicyclooctane **186** was obtained by employing Mitsunobu conditions at room temperature in 74% yield.⁵² The coupling constants of several key protons in this core substructure correlate well with the revised structure of palau'amine. For eventual total synthesis of **105** demonstration of the feasibility of this intramolecular cyclization to obtain the D–E *trans*-fused 5,5 ring system of the revised palau'amine structure is a critical achievement.



Scheme 4.12 Romo's Diels-Alder/oxidation/tandem chlorination-1,2-alkyl shift sequence for synthesis of core structure of Palau'amine.

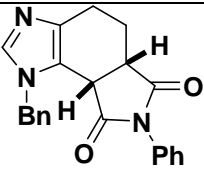
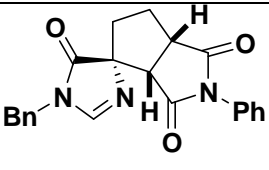
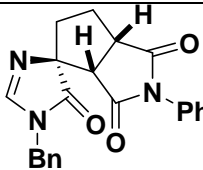
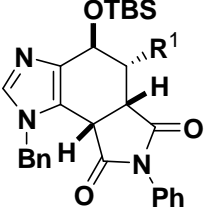
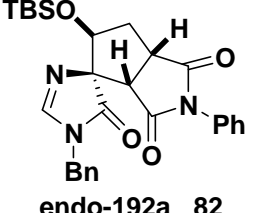
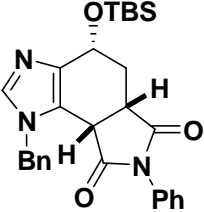
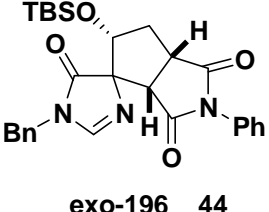
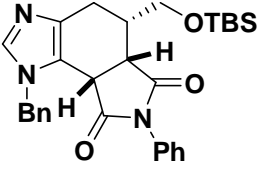
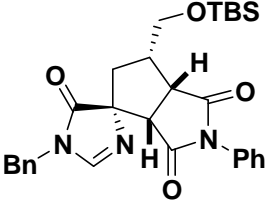
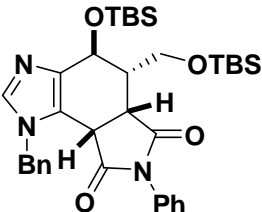
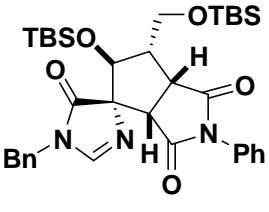
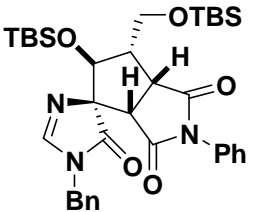
Our own group has explored approaches to palau'amine based on a similar strategy to the Romo lab. The first generation plan focused on the construction of

the DEF-rings of palau'amine through elaboration of the succinimide-containing substrates related to **192** (Scheme 4.13),^{10b} which were assembled *via* an *intermolecular* DA reaction between a vinylimidazole and subsequent oxidative rearrangement to obtain the spiro ring using DMDO or Davis' oxaziridine.

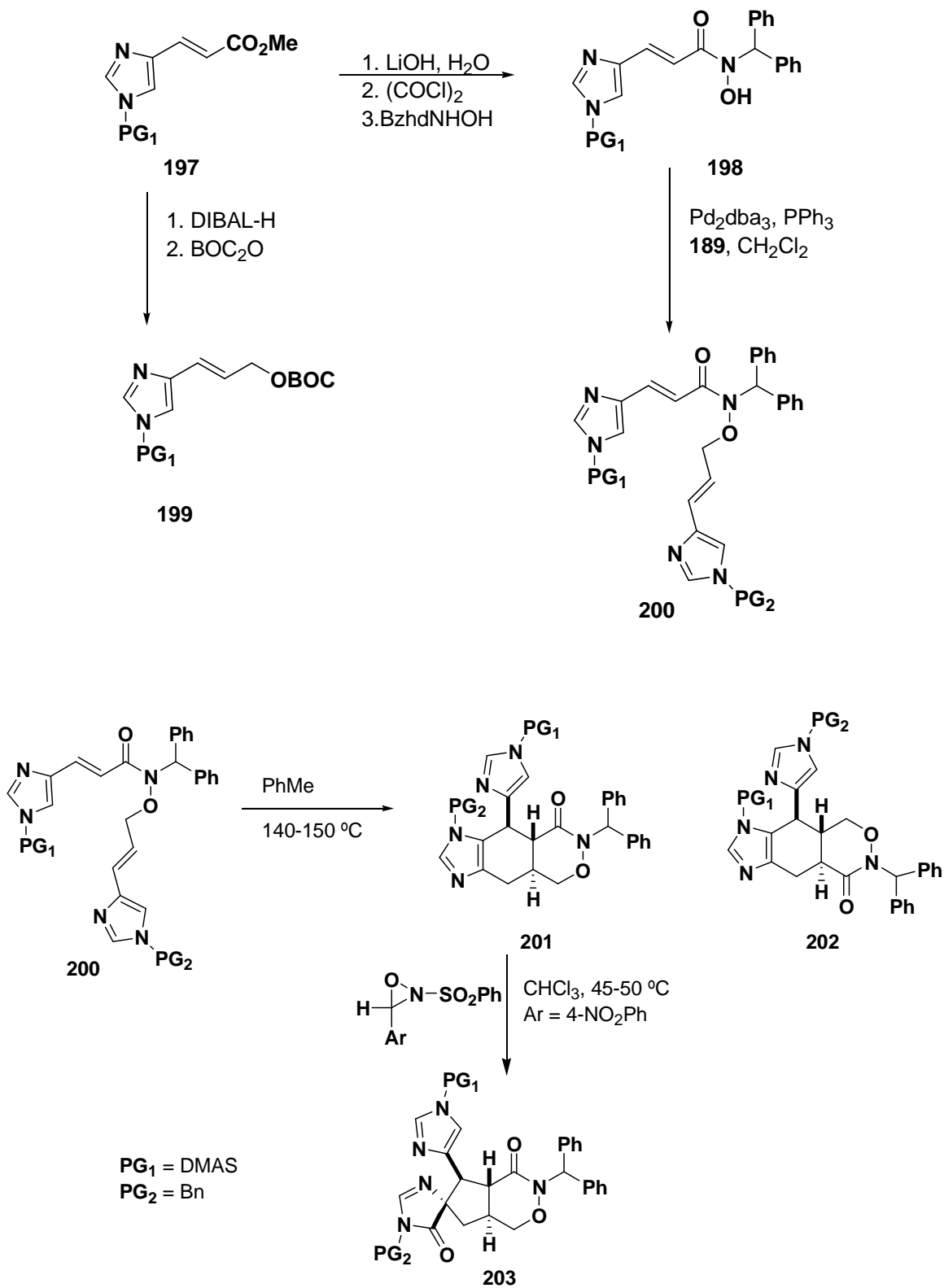


Scheme 4.13

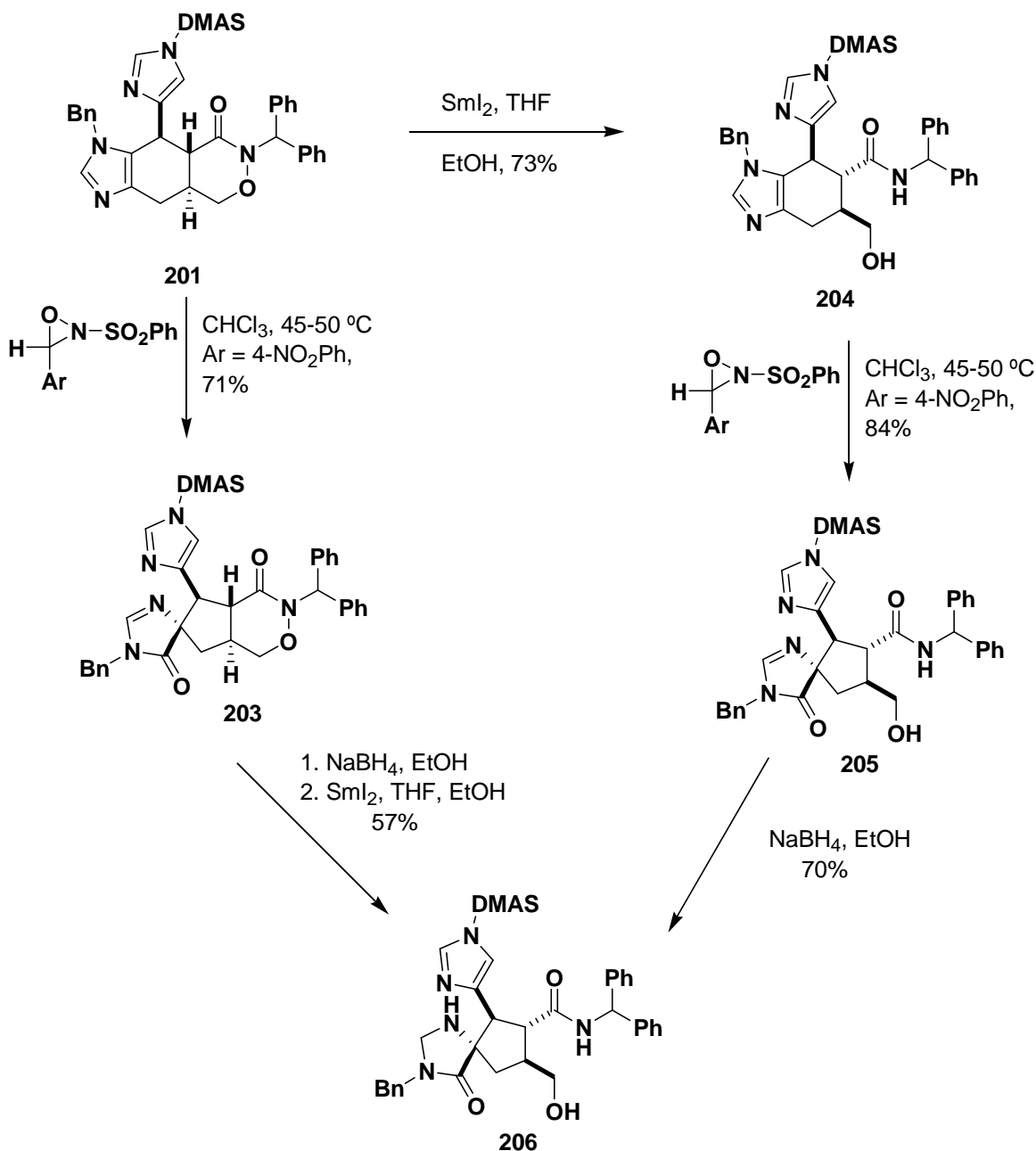
Table 4.1 Yields and products from the oxidative Rearrangement

entry	substrate	Product-yield%	Product-yield%
1	 <p>188a</p>	 <p>exo-189a 56</p>	 <p>endo-189b 27</p>
2	 <p>193</p>	 <p>endo-192a 82</p>	
3	 <p>195a</p>	 <p>exo-196 44</p>	
4	 <p>188b</p>	 <p>exo-189b 60</p>	
5	 <p>193b</p>	 <p>exo-192b 56</p>	 <p>endo-192b 14</p>

However, several aspects of this approach became less attractive in light of developments in a parallel investigation in our lab on the *intramolecular* DA (IMDA) reaction and as a result of the structural revision of palau'amine.⁹⁴ It was found in the course of these studies that fairly elaborate systems, e.g., **200** (Scheme 4.14) could be assembled extremely rapidly and these substrates would engage in cycloadditions with reasonable efficiencies. Most notable were systems that potentially allowed access into the ageliferin⁹⁴ and axinellamine⁹⁴ families through the cyclization of pseudo dimeric substrates such as **200** which afford the all *trans* substituted tetrahydrobenzimidazoles **201** on IMDA reaction (Scheme 4.15). Subsequent reductive cleavage of the hydroxamate and oxidative rearrangement with an *N*-sulfonyloxaziridine provided the spiro fused imidazolone **203**. Unfortunately, the rearranged product was epimeric at the spiro fused center preventing the further use of this particular intermediate en route to axinellamine A (**99**) or palau'amine (**105**).



Scheme 4.14 Cycloaddition and Oxidative Rearrangement

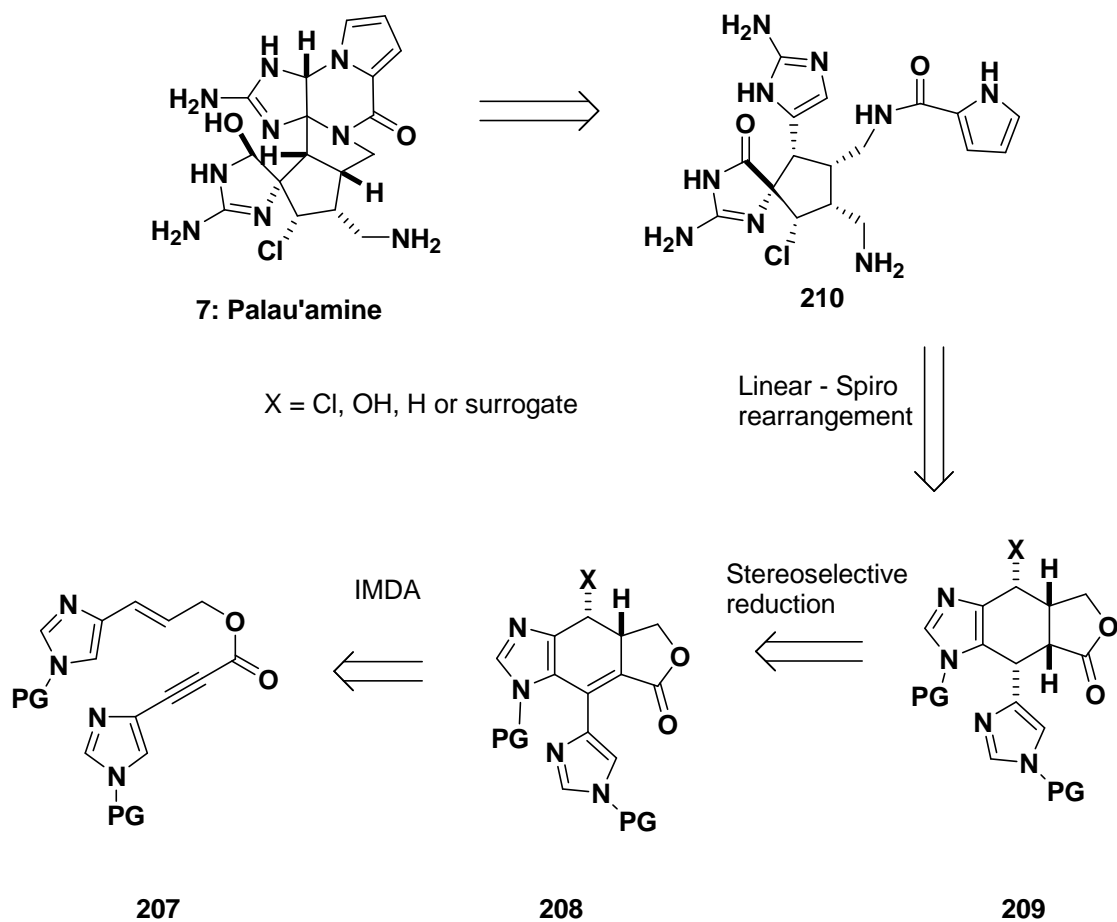


Scheme 4.15 Elaboration of the Cycloadduct

However, if the stereochemical issue of the rearrangement could be corrected, a concise approach for the construction of the key EF-ring system of palau'amine would be possible.

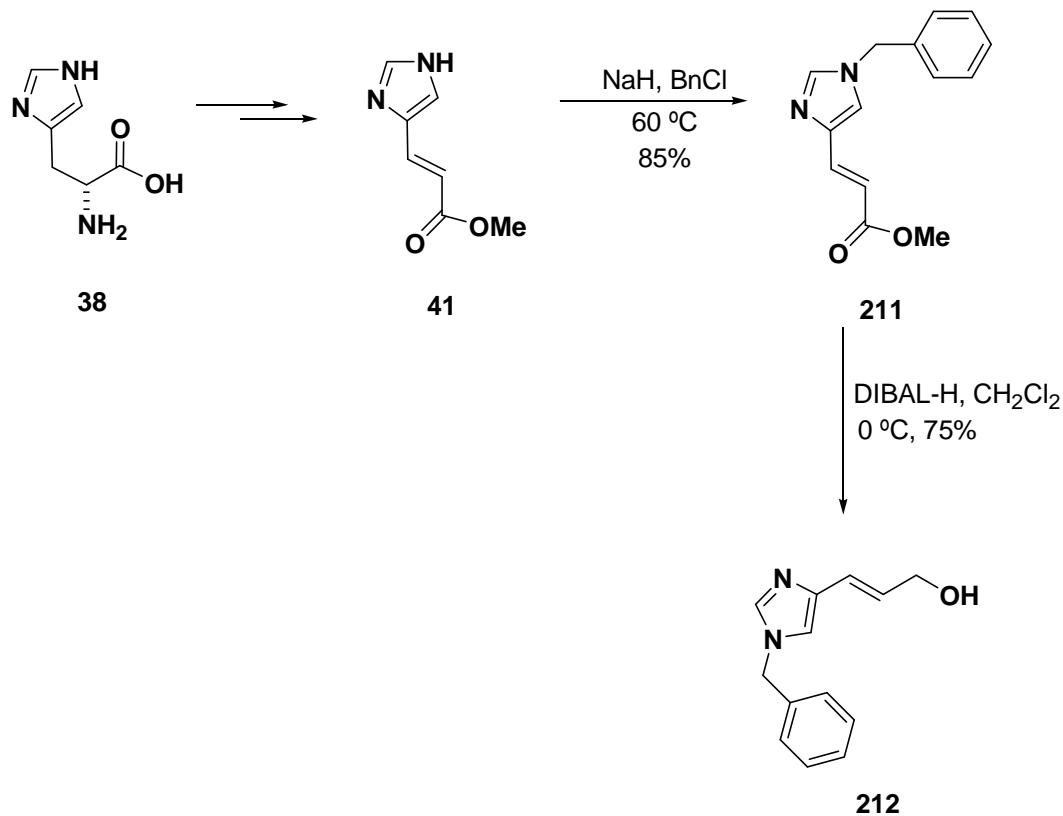
At the time these studies were initiated they were directed toward the original structure of palau'amine, *i.e.*, **7** (Fig. 1.1) but required modification of the strategy

when the structural revision appeared. As indicated above, the use of bis imidazole substrates potentially would increase convergency, and would provide substrates that could utilize chemistries related to those used for the construction of the monomeric oroidin alkaloid phakellin for the formation of the BD-rings (**210** to **7**, Scheme 4.15).⁹⁴ Thus, if an all *cis* analog of **206**, *i.e.*, **210** could be constructed, access to a palau'amine-like precursor could be envisioned. However, rather than use a *cis* substituted dienophile in our bis-imidazole approach delineated in Scheme 4.15, we chose to employ a propiolic acid derivative in the cycloaddition and perform a diastereocontrolled hydrogenation on the resulting cycloalkene to establish the relative stereochemistry (**207** to **208** to **209**). This strategy was adopted as we reasoned that this would circumvent any potential issues associated with epimerization post-cycloaddition. Furthermore, based on earlier studies from our lab with aryl propiolate derivatives we were confident that the cycloaddition strategy would be successful.⁴⁷ We were able to propose an approach to **207** and **208** depicted retrosynthetically in (Scheme 4.16), which involves an intramolecular DA reaction and its subsequent rearrangement and elaboration of lactone **208** to **209** as we have already developed key transformations like the Diels-Alder reactions of vinyl imidazoles⁵⁻¹⁰ and the oxidative rearrangement of tetrahydrobenzimidazoles.^{10b}



Scheme 4.16 Retrosynthetic analysis

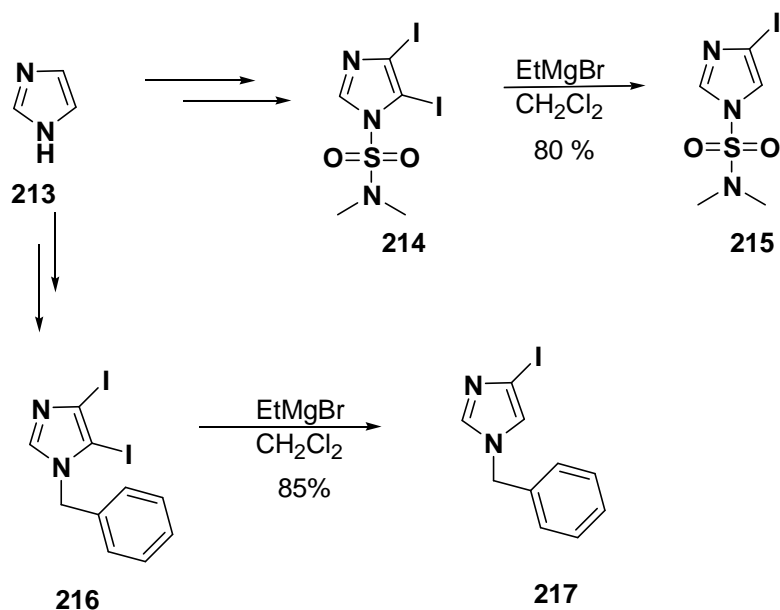
CHAPTER 5
RESULTS AND DISCUSSION



Scheme 5.1

Our studies began by converting L-histidine (**38**) to the desired urocanate ester **41** and then the free imidazole N-H was protected using NaH and benzyl chloride in dry THF to give benzyl protected urocanic ester **211**.⁹⁵ The ester was reduced to allylic alcohol **212** by using DIBAL-H (Scheme 5.1). The benzyl protecting group was used since previous work from our group has shown that an electron

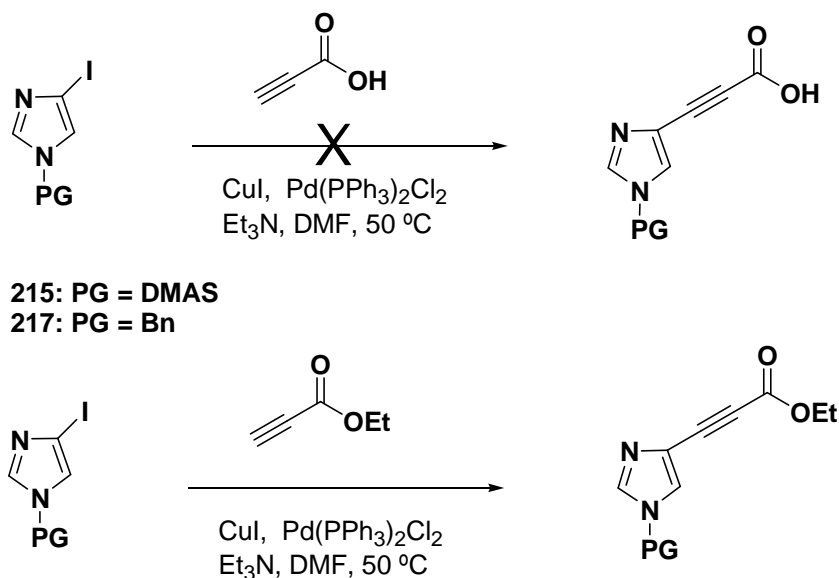
donating group at this position is critical in the formation of spiro ring in the oxidative rearrangement.⁹⁶



Scheme 5.2

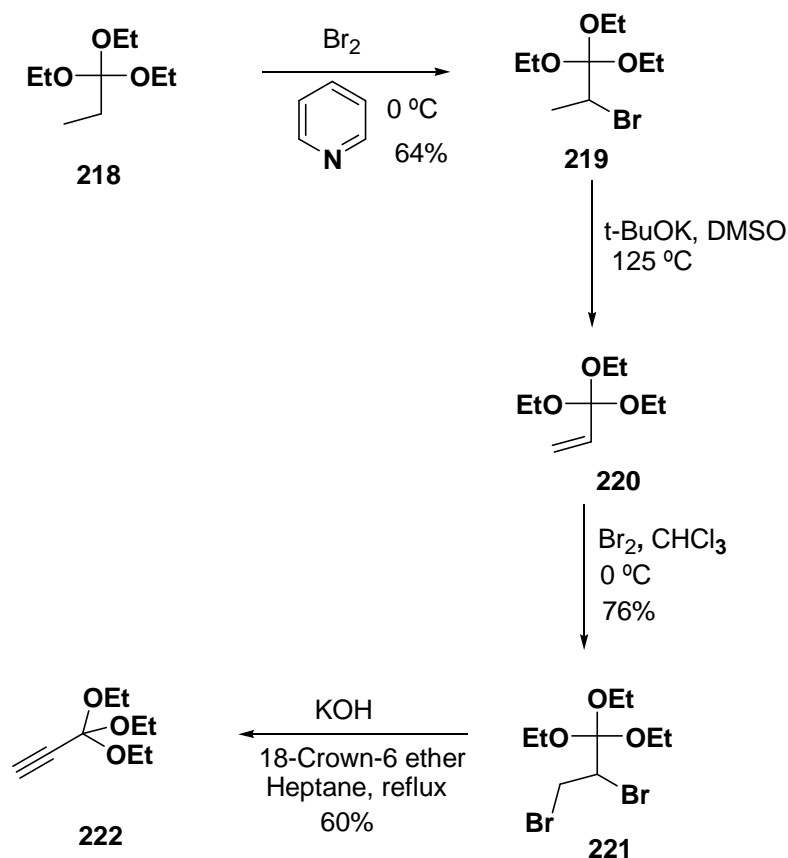
The second fragment was prepared from imidazole **213** by diiodination⁹⁷ with molecular iodine and potassium iodide in the presence of 4M NaOH (Scheme 5.2). The nitrogen atom of imidazole was protected with one the two different protecting groups so that we had flexibility at later stages of the synthesis. The iodine at the C-5 position of imidazole was removed by treatment with ethylmagnesium bromide to give **215** and **217**. With the mono iodoimidazoles in hand (**215** and **217**), our aim was to perform a Sonogashira reaction between the iodoimidazoles and a suitable alkyne coupling partner. In this regard we screened many alkynes including propiolic acid and propiolic esters (Scheme 5.3). In the case of propiolic acid we did not observe any desired Sonogashira reaction. However with propiolic ester, the desired product was obtained but the

yield was rather low (10%) as it can also act as a Michael acceptor leading to the formation many side products and scalability was also an issue with the ester.



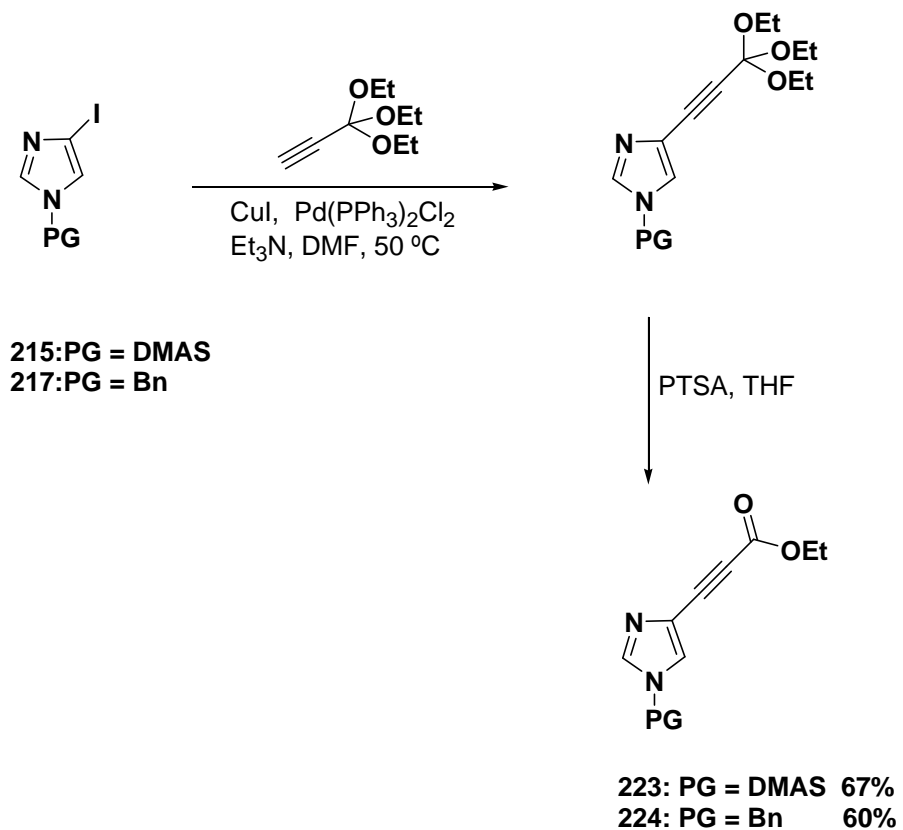
Scheme 5.3

To address this problem we decided to use 3,3,3-triethoxy-1-propyne (**222**)⁹⁸ as our other partner for the Sonogashira reaction as it is a masked propiolic ester and so it will not act as a Michael acceptor. In order to prepare this intermediate we started with the ortho ester **207**, brominated it with bromine and pyridine as a solvent (Scheme 5.4). Then compound **208** was dissolved in dry DMSO and *t*-BuOK was added, heating to 125 °C led to the formation of the alkene **209**. The resulting alkene was dibrominated by treatment with bromine in chloroform to provide the dibrominated compound **210**. To obtain compound **211**, we performed a double elimination hydrogen bromide which was achieved by using KOH in the presence of 18-crown-6 and heating the reaction mixture to 115 °C.



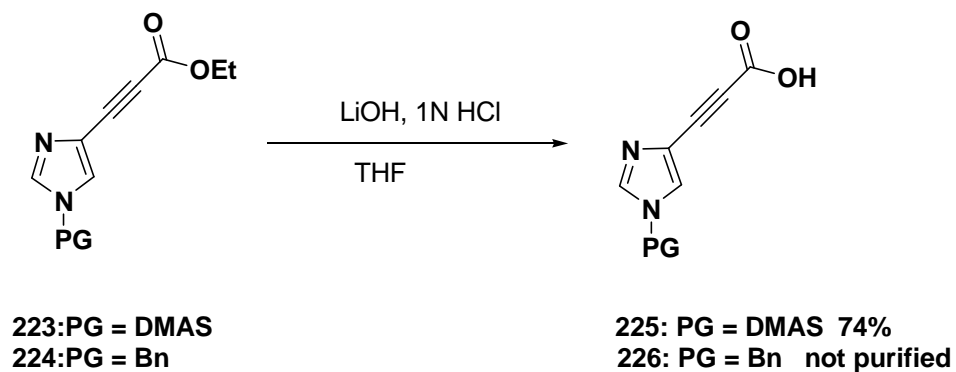
Scheme 5.4

Next a Sonogashira reaction was performed between 4-iodoimidazole (**215** and **217**) and 3,3,3-triethoxy-1-propyne (**222**) using a Pd(II) precatalyst in the presence of CuI and triethylamine and heating the reaction mixture to 50 °C (Scheme 5.5).⁹⁹ Some hydrolysis of the ortho ester occurred on purification and so the Sonogashira product was subjected directly to in situ hydrolysis using PTSA (catalytic) to give compounds **223** and **224** in 67% and 60% yield respectively.



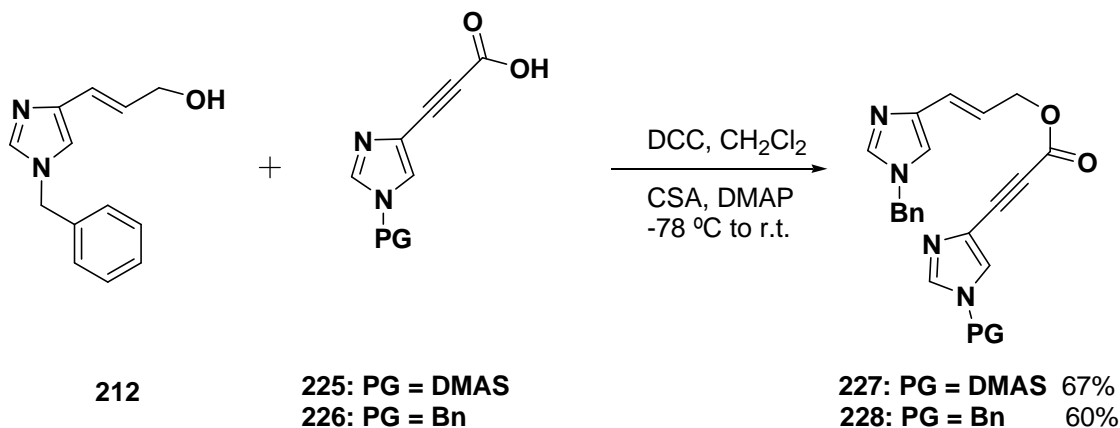
Scheme 5.5

Compounds **223** and **224** were converted into the corresponding acids **225** and **226** using aqueous lithium hydroxide and then neutralized with 1N HCl (Scheme 5.6).



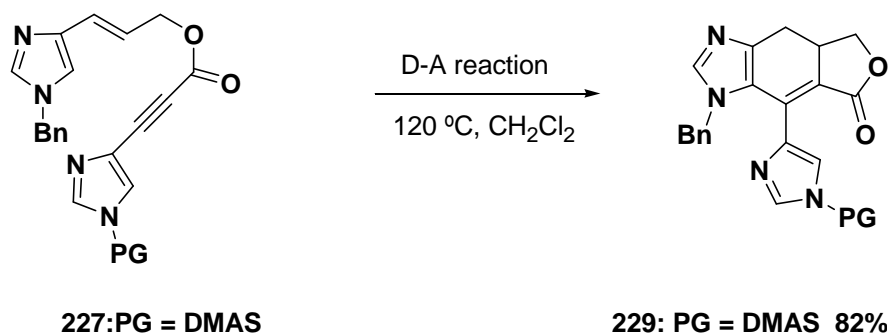
Scheme 5.6

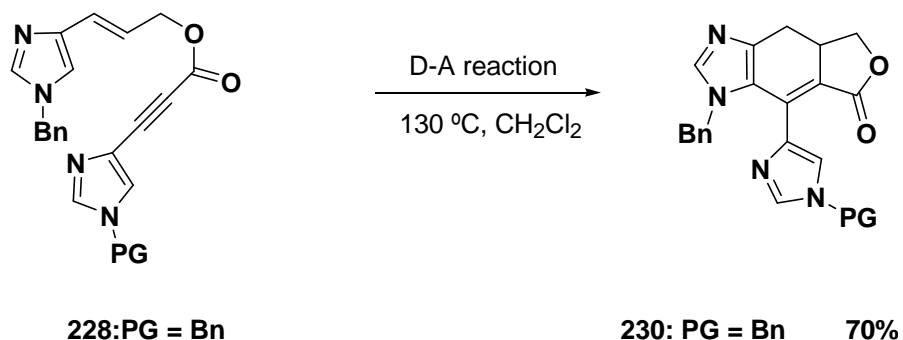
The resulting acids **225** and **226** were coupled with alcohol **212** mediated by DCC in the presence of a catalytic amount of camphorsulfonic acid (CSA) and 4-dimethylaminopyridine (DMAP) at $-78\text{ }^{\circ}\text{C}$ to obtain the enyne compounds **227** and **228** (Scheme 5.7).⁴⁹



Scheme 5.7

The ene-yne compounds **227** and **228** were dissolved in dichloromethane and heated to $\sim 120\text{ }^{\circ}\text{C}$ in a sealed tube to provide the dihydrobenzimidazole Diels-Alder products **229** and **230** (Scheme 5.8). To obtain this Diels-Alder product we screened other solvents like benzene and toluene but the desired Diels-Alder product was obtained only when we used dichloromethane as solvent.

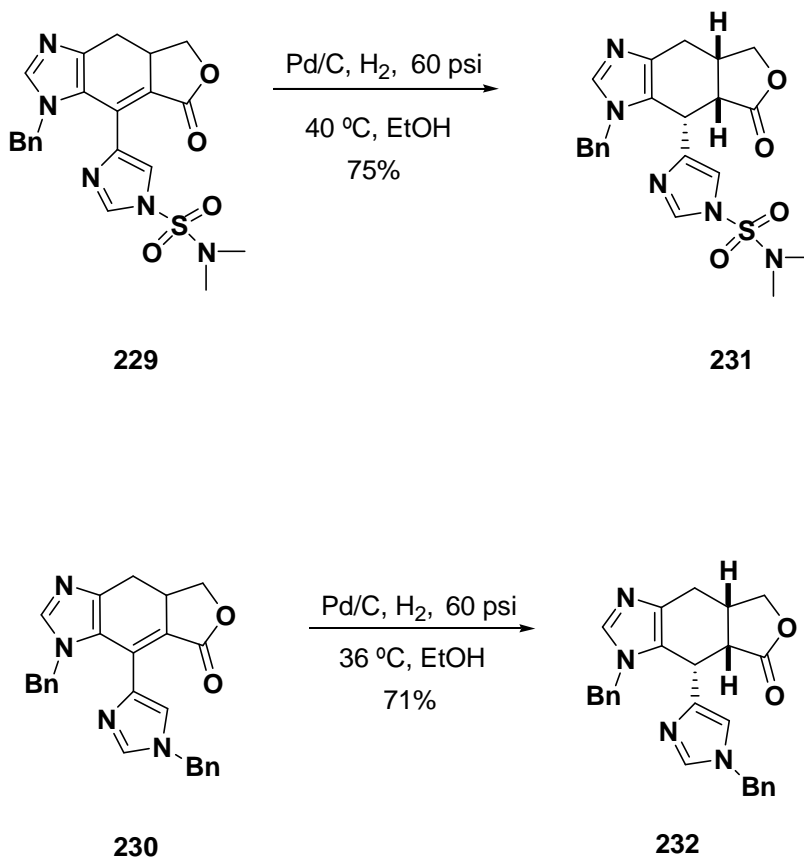




Scheme 5.8

The alkene in the six membered ring of the cycloadducts **229**, **230** was saturated using Pd/C under a hydrogen atmosphere (60 psi) with heating of the reaction mixture to 40 °C (Scheme 5.9).

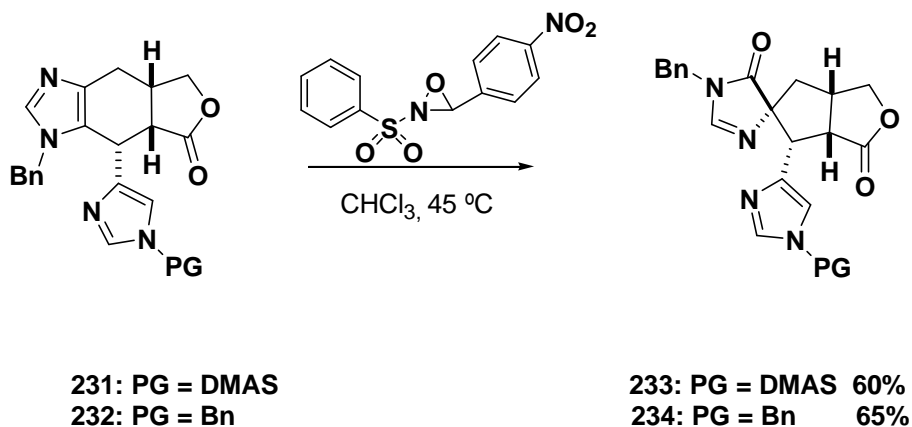
The initial assignment of the relative stereochemistry of the reduction product was obtained by examination of the relevant coupling constants of the bridgehead and benzylic protons. It has been determined that $J_{4a,7a} = 8.7$ Hz and $J_{7a,8} = 8.7$ Hz, which are completely consistent with the indicated stereochemistry. We have prepared a large number of cycloadducts related to **229** and **230** with the exception that they are all *trans* substituted, and in these cases the corresponding coupling constants are substantially larger $J_{4a,7a} = 12.8$ -13.6 Hz, and $J_{7a,8} = 10.1$ -10.8 Hz.^{10a} Further, in several cases the relative configurations of these all *trans* derivatives have been rigorously established by X-ray crystallography.⁹⁴



Scheme 5.9

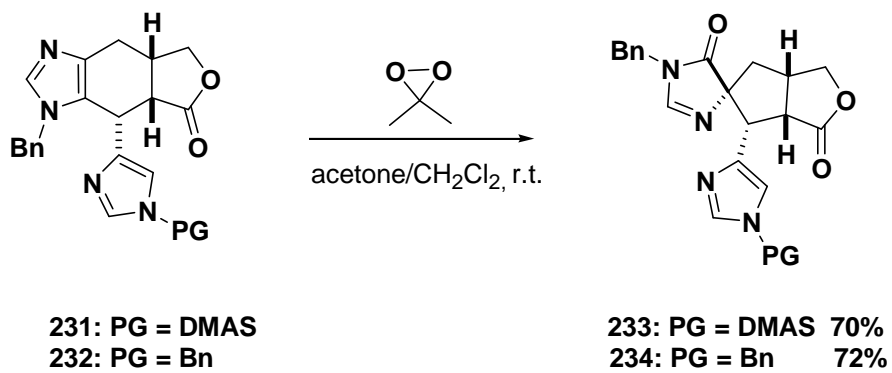
With the all-*cis* cycloadducts **231** and **232** in hand, our next task involved their elaboration, including rearrangement into the spiro fused system. We have shown previously that this can be accomplished through an oxidative rearrangement using dimethyldioxirane,^{10b} and while this works well in many cases some aspects of this chemistry were unattractive, in particular the need to prepare isolated reagent. It has been found (recently) that this same rearrangement can be performed using Davis' reagents (*N*-sulfonyloxaziridines).¹⁰⁰ Accordingly, when **231** and **232** were treated with two equivalents of Davis reagent in CHCl_3 at $40\text{ }^\circ\text{C}$, it undergoes a smooth oxidative rearrangement providing a single spiro imidazolone **233**, **234** in good yield

(Scheme 5.10). The exquisite chemoselectivity for the more substituted (and presumably more electron rich) imidazole in this reaction is noteworthy. We were also fortunate that both **233** and **234** were well-behaved crystalline solids, which gave crystals suitable for X-ray crystallography. This not only confirms the stereochemical sense of the spiro fusion, but also that the catalytic hydrogenation occurs *syn* to the hydrogen at the ring junction and leads to an all-*cis* fusion. Of particular note is the stereochemical outcome of this rearrangement. Our group previously reported studies with all *trans* substituted systems⁹⁴ and also one related substrate reported by Baran and coworkers⁷⁵ gives rise to an imidazolone with the opposite stereochemistry, and this seemingly subtle change in the relative stereochemistry of the substrate leads to a complete changeover in selectivity.



Scheme 5.10

For comparison, we also examined the rearrangement with DMDO and found that this proceeded in marginally higher yield. The other advantage with DMDO is that the purification is substantially easier as the byproducts from the oxidant are readily removed.



Scheme 5.10

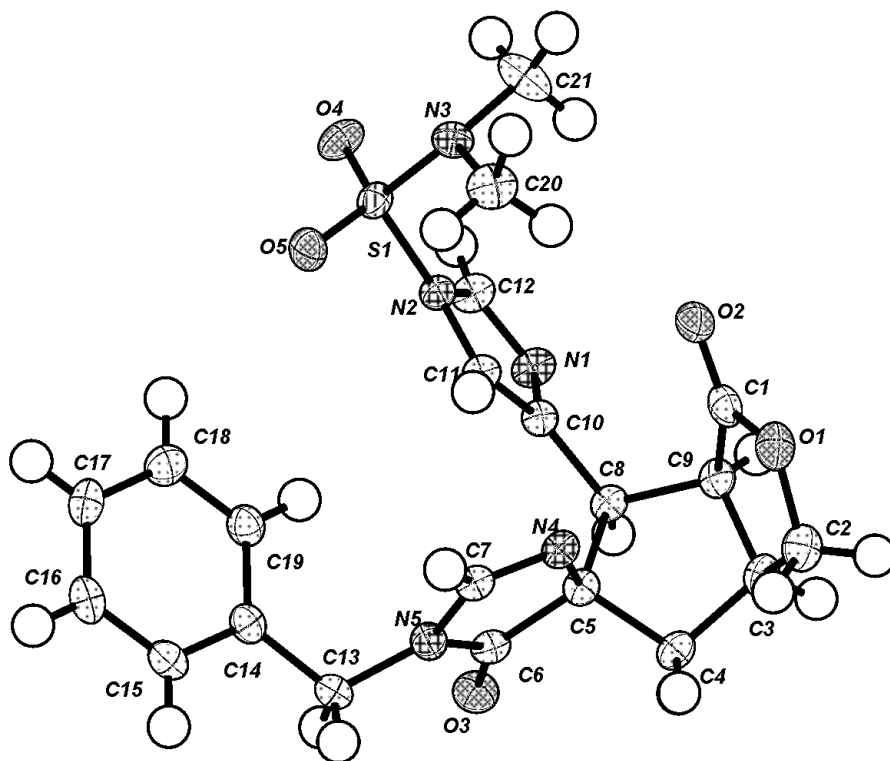
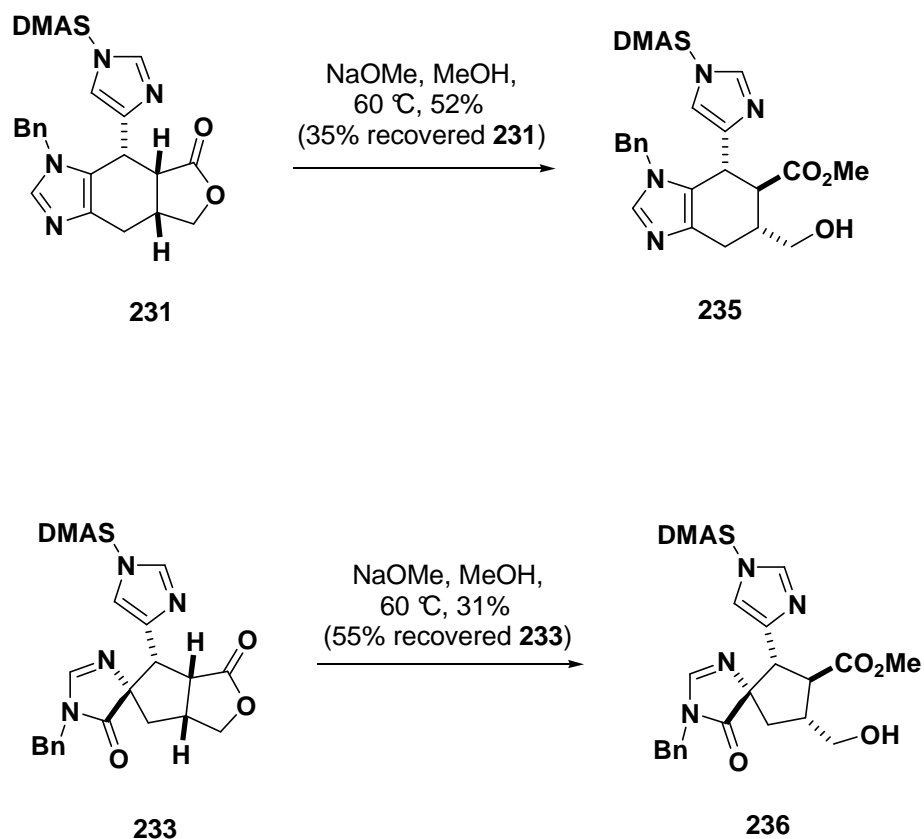


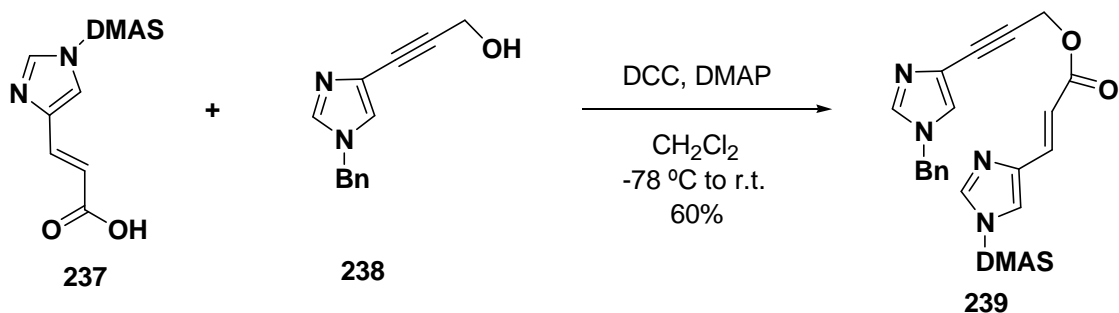
Figure 5.2 X-ray crystal structure of compound 233

After some experimentation it was found that the lactone **233** underwent ring opening-epimerization on treatment with NaOMe in MeOH at 60 °C, providing the hydroxy ester in a modest 35% yield, with 55% recovered starting material (Scheme 5.11).⁵² It was also determined that the tetrahydrobenzimidazole **231** would undergo ring-opening/epimerization, leading to an ageliferin-like stereochemical arrangement. Although both ring opening/epimerization reactions, require further optimization, access to the spiro fused core of palau'amine and related compounds has been developed. In addition, the cycloadducts **231** provide an entry to the ageliferin and potentially the related nagelamide family of natural products by suitable adjustments of the stereochemistry, and these efforts are underway in our lab.



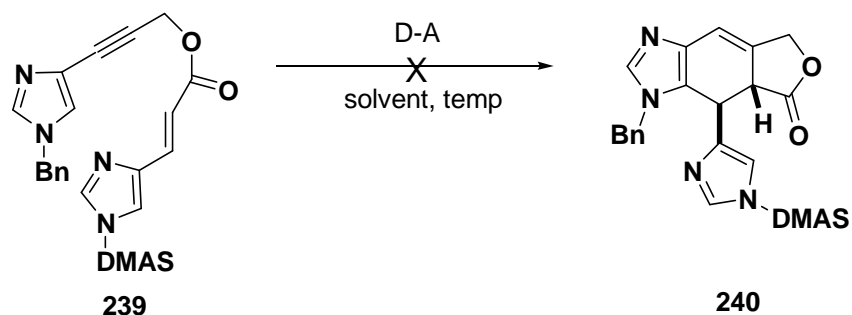
Scheme 5.11

After obtaining all *trans*-substituted spiro cyclopentyl imidazolone system found in palau'amine and related natural products, we turned our attention towards investigating methods for the stereoselective incorporation of the chloro moiety, and for the construction of the remaining rings. In order to incorporate the chloro or hydroxy moiety in the molecule, our initial strategy was to transpose the enyne part of the molecule **227** so that after the Diels-Alder reaction, the molecule will have a double bond in the six-membered ring which we can utilize to incorporate a chloro or a hydroxy group later. In order to prepare this substrate **239** compound **237** was coupled with **238** in presence of DCC (Scheme 5.12).⁴⁹



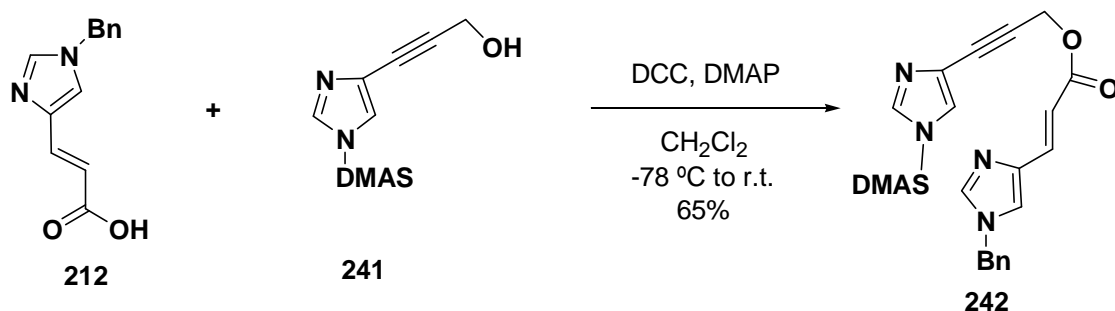
Scheme 5.12

Once we had compound **239**, we subjected it to Diels-Alder reaction under various solvent and at various reaction temperature starting from 90 to 120 °C but the desired product **228** was never isolated and starting material was recovered (Scheme 5.13).



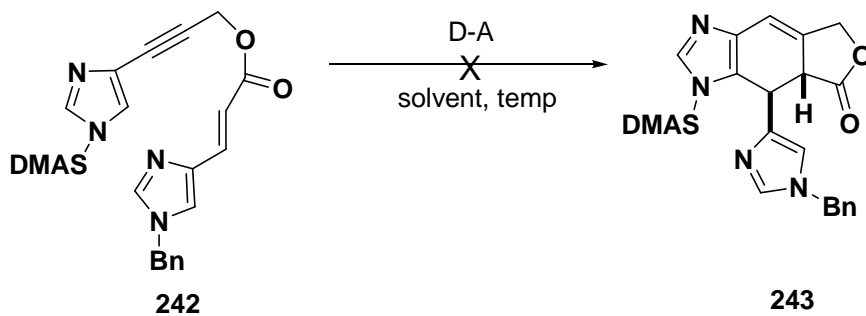
Scheme 5.13

Since compound **239** did not undergo Diels-Alder reaction, we wondered whether interchanging the protecting groups on the enyne compound **239** would lead to a productive D-A reaction (Scheme 5.14).



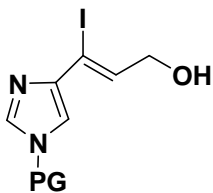
Scheme 5.14

Compound **242** was dissolved in dichloromethane in sealed tube and then it was subjected to Diels-Alder reaction at various reaction temperature starting from 90 to 120 °C but the desired product **243** was never isolated (Scheme 5.15).



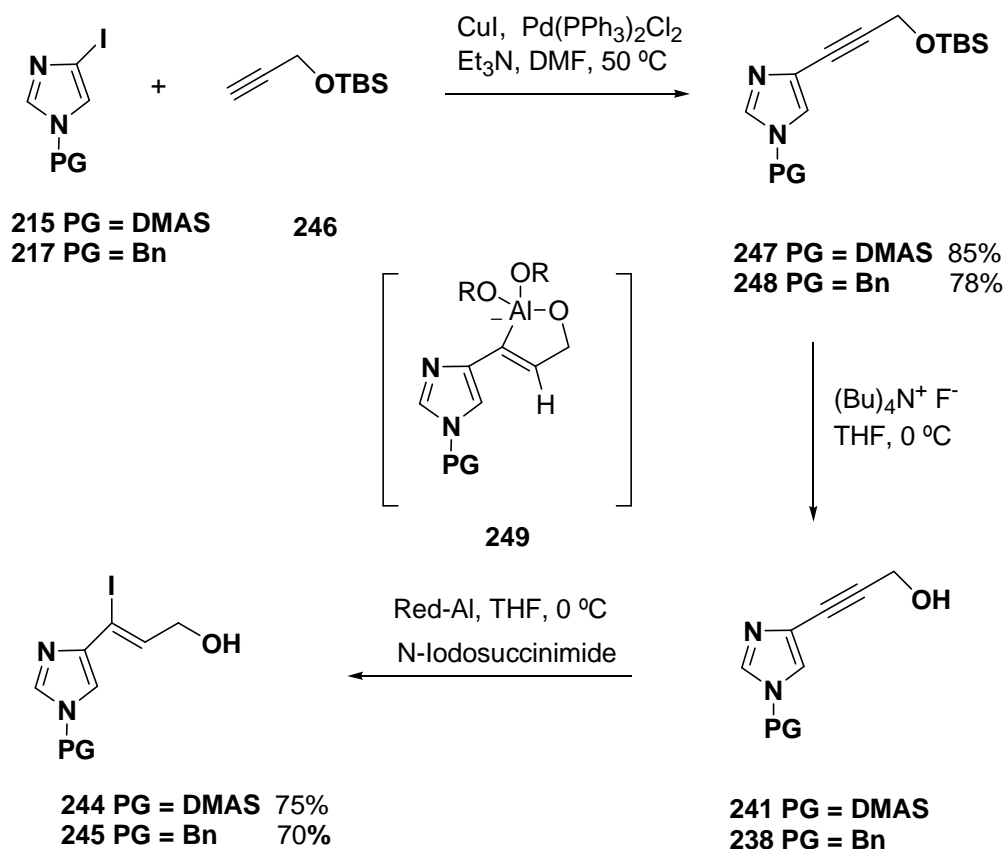
Scheme 5.15

Since compounds **239** and **242** did not undergo Diels-Alder reaction as expected, a change of strategy was necessary, specifically we chose to evaluate substrates in which the vinyl moiety was substituted. In this regard we considered using a halide, as it can be utilized either as a latent handle for incorporating other substituents at this position or in the case of chlorine as the required substituent.



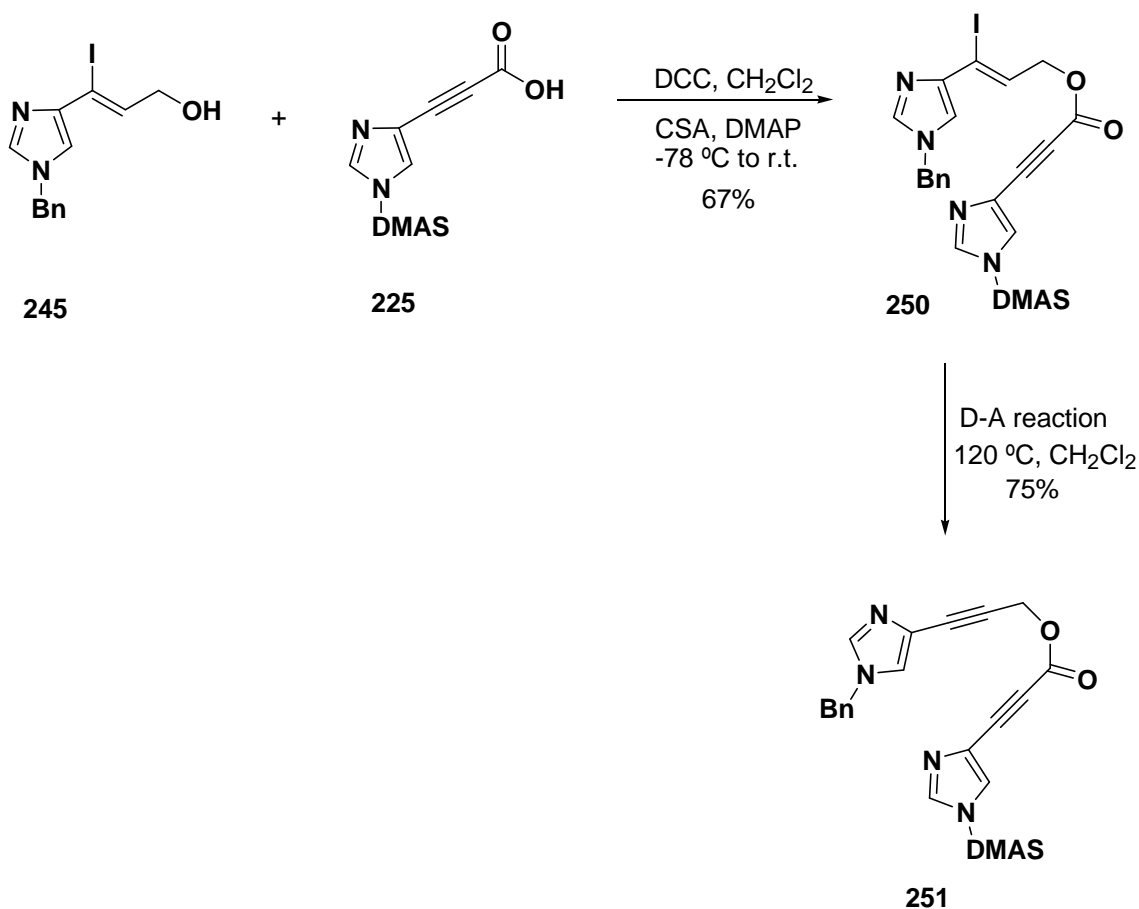
244 PG = DMAS
245 PG = Bn

First a Sonogashira reaction between iodoimidazoles **215** and **217** and the protected propargylic alcohol provided **247** and **248** (Scheme 5.16). Subsequent treatment of silyl ethers **247** and **248** with tetrabutylammonium fluoride provided alcohols **241** and **238**. The desired substituted alcohols **244** and **245** were obtained by hydroalumination with Red-Al¹⁰¹ which upon addition of N-iodosuccinimide to the reaction mixture at 0 °C provided the vinyl iodides.



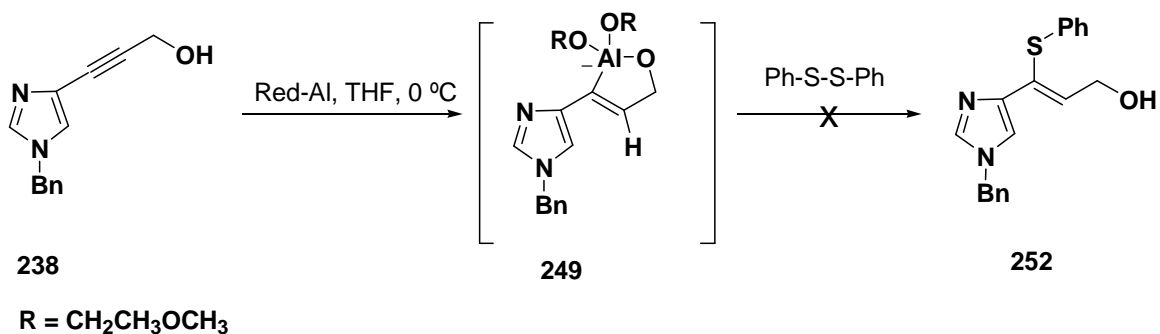
Scheme 5.16

Alcohol **245** was coupled with acid **225** mediated by DCC to obtain the enyne compound **237** (Scheme 5.17). Compound **250** was then subjected to an attempted Diels-Alder reaction at various temperatures ranging from 90 to 130 °C. We observed that at 90 °C the starting material does not react, so we increased the temperature to 100 °C still the starting material does not react. The same thing was observed as the temperature increased to 110 °C and 115 °C respectively. However, when the temperature was increased to 120 °C, a new spot was observed developing on TLC plate which upon characterization was found to be compound **251** resulting from dehydroiodination.



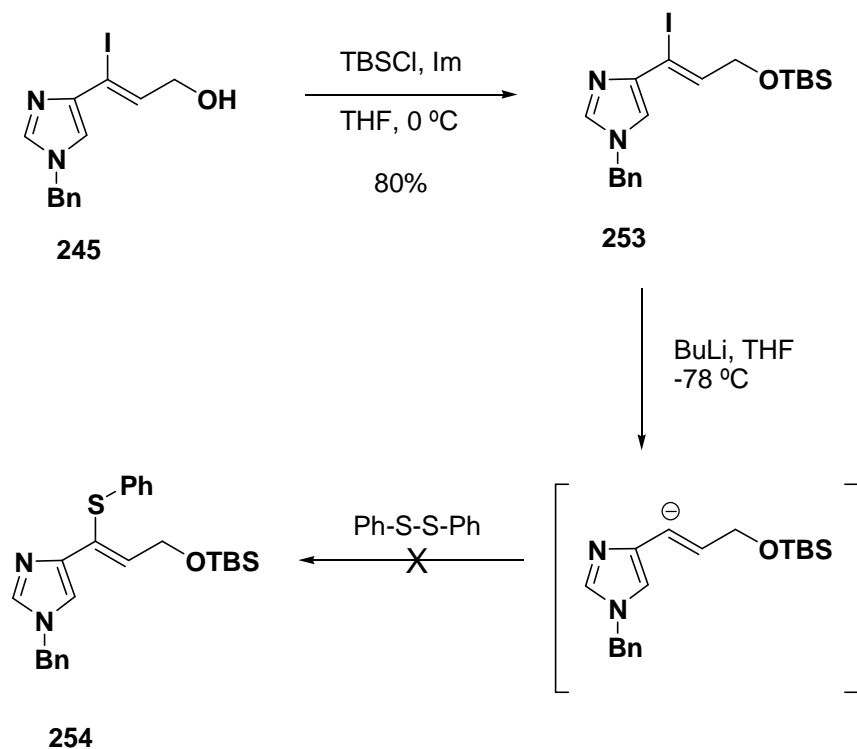
Scheme 5.17

While we were not surprised that the iodine substituent was somewhat labile under the Diels-Alder conditions, we were confident that the general strategy was worth pursuing and thus substituents tolerant of the Diels-Alder reaction were required but that would function as latent halide equivalent. In this regard we tried to incorporate a thiophenyl moiety in place of iodine. Compound **238** was dissolved in THF and Red-Al was added at 0 °C, then diphenyl disulfide was added to the mixture and was allowed to come to r.t. but we were unable to isolate compound **252**. Raising the reaction temperature to 50 °C, also failed to lead to a productive reaction (Scheme 5.18).



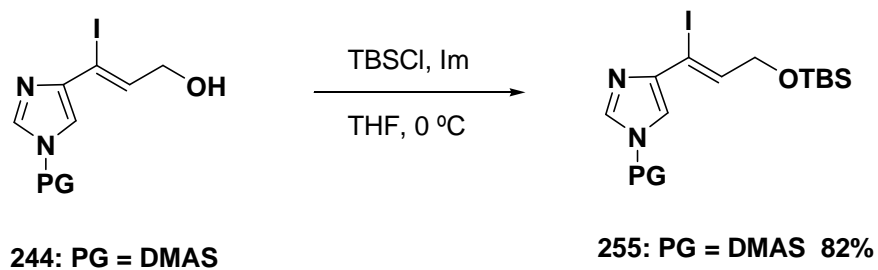
Scheme 5.18

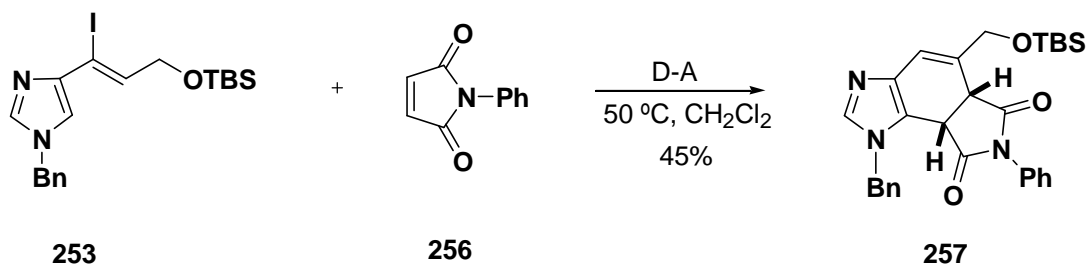
Presumably, we were not able to obtain the compound **252** because intermediate **249** is not sufficiently nucleophilic to attack the diphenyl disulfide. Therefore it was thought that generating the vinyl anion by iodine-lithium exchange and then adding diphenyl disulfide may provide a solution to the synthesis of **252**. The alcohol **245** was protected with TBSCl and then treatment of **253** in THF with BuLi at $-78\text{ }^\circ\text{C}$ resulted in iodine-lithium exchange to the desired anion. Diphenyl disulfide was added cooling the reaction mixture to $-78\text{ }^\circ\text{C}$. After allowing the reaction mixture to warm up to the r.t., it was found that the desired product has not formed. Even heating the reaction mixture to $50\text{ }^\circ\text{C}$ did not provide the desired product **254** (Scheme 5.19) instead we obtained the reduced allylic alcohol.



Scheme 5.19

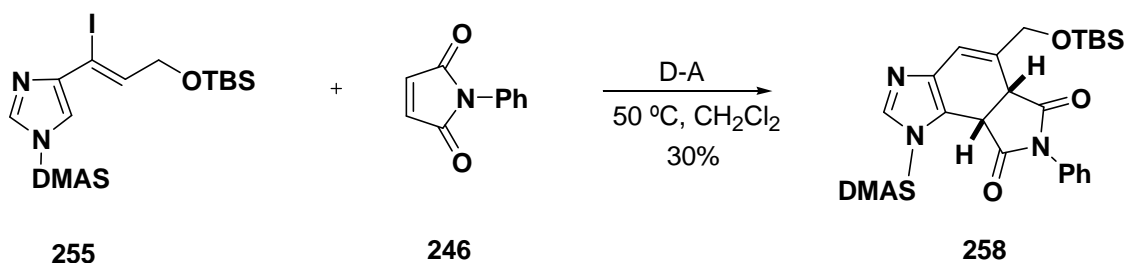
Since we were not able to introduce the sulfur group at the vinyl position, we went back and protected the alcohols **244** and **245** with TBS group and then tried the intramolecular Diels-Alder reaction with the reactive *N*-phenylmaleimide at much lower temperature such as 50 °C instead of 120 °C. Even at 50 °C we found that dehydroiodination was occurring, resulting in the formation of compound **257** (Scheme 5.20).^{5,102}





Scheme 5.20

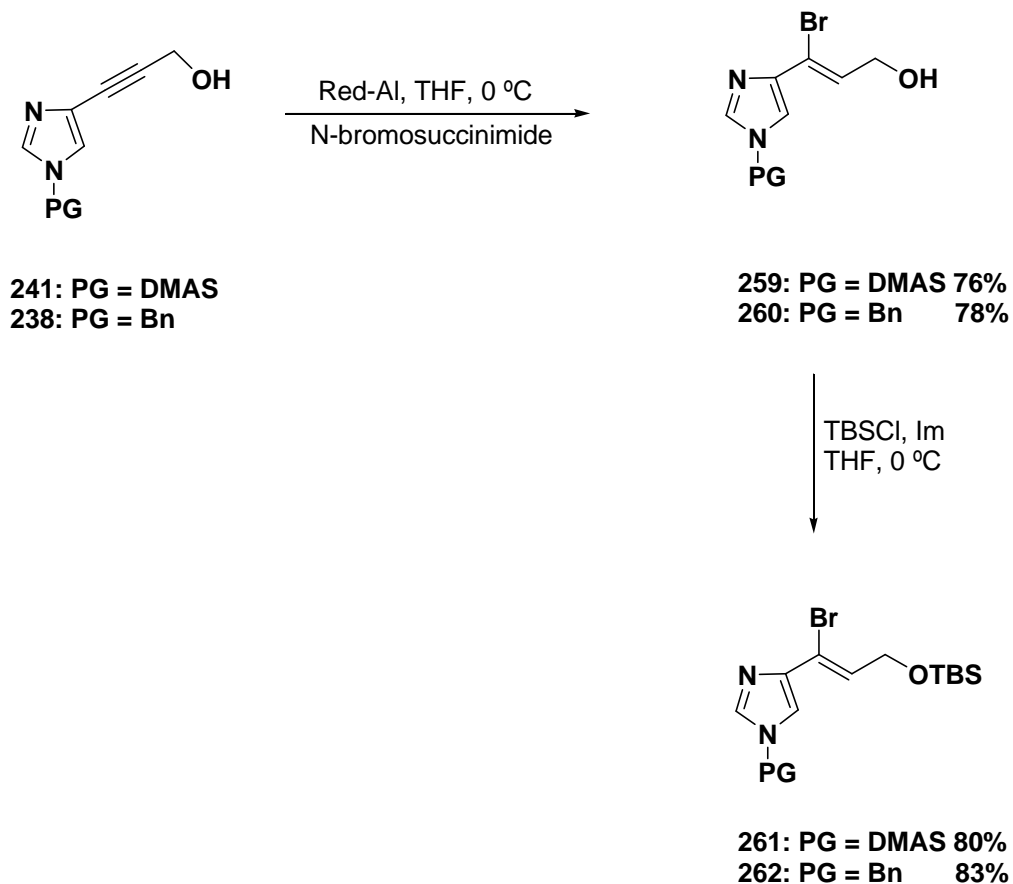
In order to stop this dehydroiodination we used $Y(OTf)_3$ as a Lewis acid since this Lewis acid has been reported to promote successful Diels-Alder reactions at much lower temperature including with vinylimidazoles.¹⁰³ So we dissolved compound **253** and **256** in dichloromethane and to the reaction mixture we added 0.01 equivalent of $Y(OTf)_3$ and then heated the reaction mixture in a sealed tube at 38 °C. Even at this temperature we obtained the same compound **257**. A similar outcome was observed with compound **255** when subjected to Diels-Alder reaction (Scheme 5.21).



Scheme 5.21

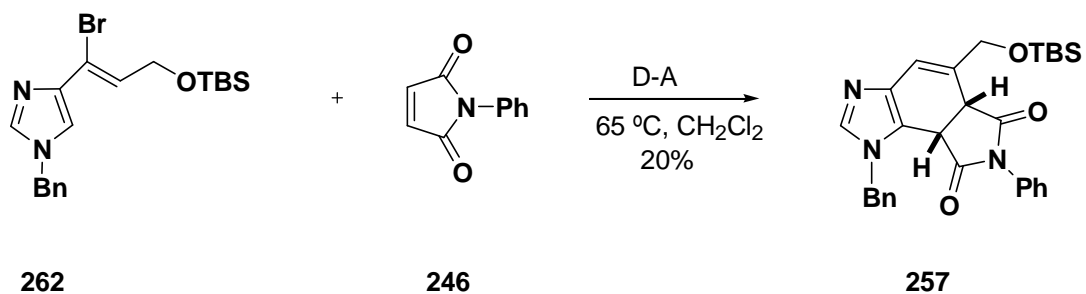
Since the iodo substituent was somewhat labile under the reaction conditions we thought that replacing iodine either with bromine or chlorine may provide more stable adducts under the Diels-Alder reaction conditions. Although the resulting 4-halo adducts are benzylic halides and thus be prone to elimination. In that

regard compound **259** and **260** were obtained by hydroalumination with Red-Al which upon treatment of *N*-bromosuccinimide to the reaction mixture at 0 °C provided the vinylbromide. Alcohols **259** and **260** were then protected with TBS group to provide compounds **261** and **262** (Scheme 5.22).



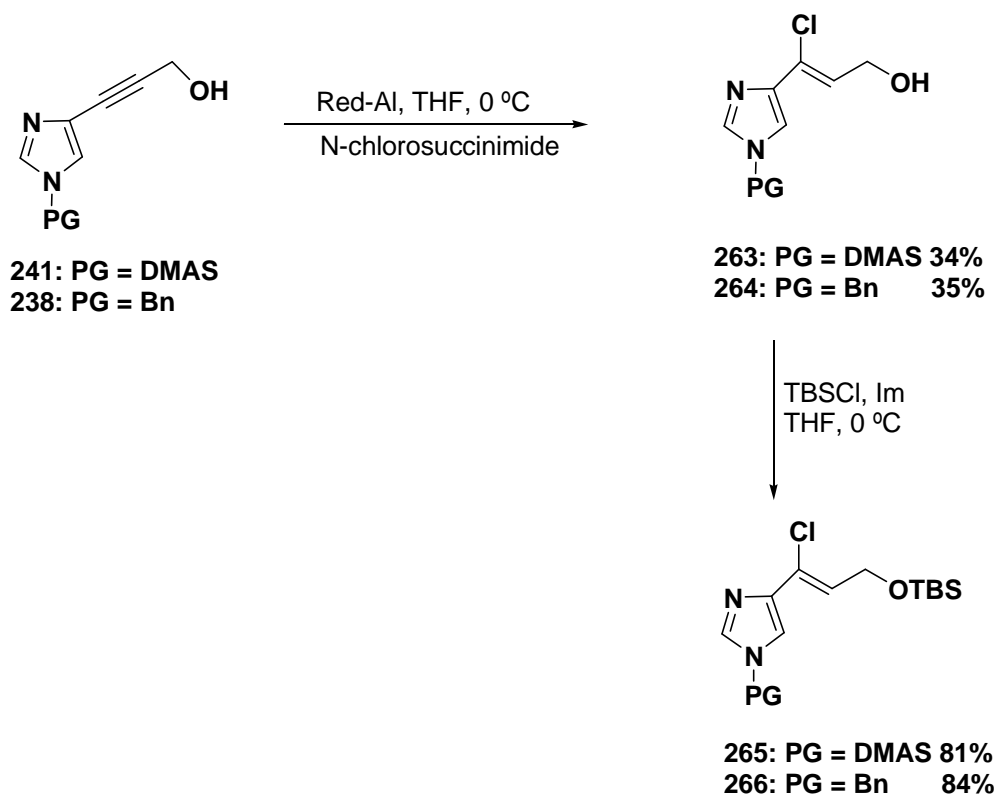
Scheme 5.22

Compound **262** was subjected to the Diels-Alder reaction at various temperatures such as from 35 °C to 65 °C and it was determined that starting material remains intact up to 60 °C. At around 65 °C a new component was observed on TLC plate which upon characterization found to be compound **257** resulting from dehydrobromination in 20% yield (Scheme 5.23).



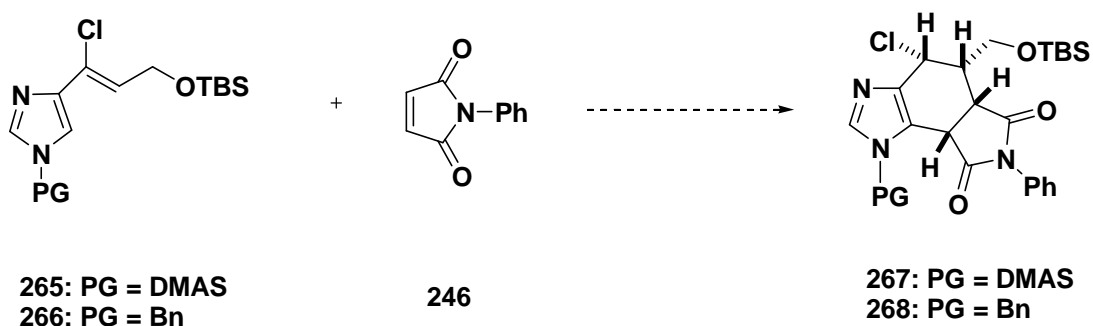
Scheme 5.23

Since the Diels-Alder product derived from compound **262** also underwent dehydrobromination under the reaction conditions we decided to make the chloro version. Compound **263** and **264** were obtained by hydroalumination with Red-Al which upon treatment with *N*-chlorosuccinimide at 0 °C provided the vinylchloride. Alcohols **263** and **264** were then protected with a TBS group to provide compound **265** and **266** (Scheme 5.24).



Scheme 5.24

Compound **265** and **266** were dissolved in dichloromethane and to each of the reaction mixture **246** was added and then the resulting reaction mixture were subjected to the Diels-Alder reaction conditions. The reactions were monitored starting from around 40 to 75 °C, but the desired product was not obtained instead starting materials were recovered (Scheme 5.25). It looks like the chloro atom is deactivating the diene and so to push this reaction towards forward direction we need to heat the reaction mixture at higher temperature. This reaction needs further investigation.



Scheme 5.25

In conclusion, we have developed a concise entry into the all *trans*-substituted spiro cyclopentyl imidazolone system found in palau'amine and related natural products *via* an IMDA reaction of an enyne followed by an oxidative rearrangement. Initial experiments aimed at elaborating this intermediate have demonstrated that precursors suitable for investigating various end-game strategies can be constructed by differentiating the two hydroxymethyl handles. Finally we have investigated a method for the stereoselective incorporation of the chloro moiety, which is present in parent molecule palau'amine.

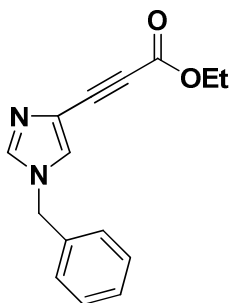
CHAPTER 6

EXPERIMENTAL DETAILS

3.1 General procedures

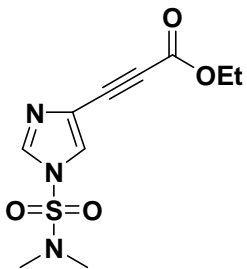
All reagents were purchased from commercial suppliers and were used as received unless otherwise noted. ^1H and ^{13}C NMR (δ in ppm) spectra were recorded in CDCl_3 (unless otherwise noted) at 500 and 125.8 MHz, respectively; using a JEOL Eclipse+ 500 spectrometer unless otherwise noted using residual CHCl_3 as reference (^1H NMR and carbon absorption of CDCl_3 for ^{13}C NMR). Infrared spectra were recorded either as neat films or as KBr pellets using a Bruker Vector 22 spectrometer. Elemental analyses were performed using a Perkin-Elmer 2400 CHN analyzer. Optical rotation was measured on a Perkin-Elmer 241MC polarimeter ($c = \text{g}/100 \text{ mL}$) and the observed value was an average of 2-3 runs. The solvent used for optical rotation was MeOH unless otherwise noted. High resolution mass spectra (HR-MS) were obtained from Dr. Powell's lab in the University of Florida, Gainesville, Florida.

Ethyl 3-(1-benzyl-1H-imidazol-4-yl)propiolate (224):



DMF (50 mL) was first purged with N₂ then Bn-protected 4-iodoimidazole **217** (5.0 g, 0.02 mole) was added followed by Pd(PPh₃)₂Cl₂ (0.37 g, 0.53 mmol), copper (I) iodide (200 mg, 1.06 mmol), orthoester^{98a} (4.54 mL, 26 mmol) and finally triethylamine (6.1 mL, 40 mmol) under a N₂ atmosphere. The reaction mixture was heated at 50 °C for 8 h. *p*-TsOH (400 mg) was added to the above reaction mixture and then stirred at r.t. overnight. The organic solvent was washed with water (200 mL), the organic layer was dried with anhydrous Na₂SO₄ concentrated and the residue was purified by chromatography on silica gel (hexane/EtOAc, 3:2) providing **224** as a thick yellow liquid (2.68 g, 60%). ¹H NMR (300 MHz): δ = 7.38-7.36 (m, 4H), 7.19-7.17 (m, 3H), 5.11 (s, 2H), 4.24 (q, *J* = 7.6 Hz, 2H), 1.31 (t, *J* = 7.6 Hz, 3H); ¹³C NMR (75 MHz): δ = 154.2, 138.4, 134.9, 129.3, 128.9, 127.6, 127.5, 121.9, 81.7, 80.8, 61.9, 51.5, 14.2; FT-IR (neat, cm⁻¹): 1749, 1716, 1699, 1497, 1489, 1473, 1457, 1436, 1419, 1396, 1374, 1362, 1339, 1318, 1094; HR-MS (*m/z*): calc for [M+H]⁺ C₁₅H₁₄N₂O₂ is 255.1128 found 255.1120.

Ethyl 3-(1-(*N,N*-dimethylsulfamoyl)-1H-imidazol-4-yl)propiolate (223): DMF

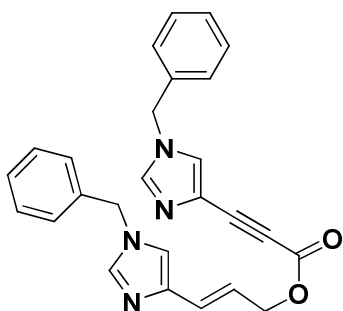


(55 mL) was purged by bubbling N₂ through it for 10 min then DMAS-protected 4-iodoimidazole **215** (5.5 g, 18 mmol) was added followed by Pd(PPh₃)₂Cl₂ (0.38 g, 0.54 mmol), copper (I) iodide (0.21 g, 1.09 mmol), ortho ester (4.7 mL,

27 mmol) and triethylamine (6.3 mL, 45 mmol) under an N₂ atmosphere. The reaction mixture was heated at 45-50 °C for 8 h. *p*-TsOH (500 mg) was added to the above reaction mixture followed by stirring at r.t. overnight. The organic solvent was washed with water (250 mL), the organic layer was dried with anhydrous Na₂SO₄ concentrated and the residue was purified by chromatography on silica gel (hexane/EtOAc, 3:2) to provided **223** as a yellow solid (3.30 g, 67%). m.p. 72-74 °C. ¹H NMR: δ = 7.85 (s, 1H), 7.59 (s, 1H), 4.25 (q, *J* = 6.9 Hz, 2H), 2.87 (s, 6H), 1.31 (t, *J* = 7.3 Hz, 3H); ¹³C NMR: δ = 153.6, 137.1, 124.6, 123.4, 82.5, 78.2, 62.3, 38.3, 14.1; FT-IR (neat, cm⁻¹): 3131, 2984, 2223, 1706, 1474, 1421, 1396, 1332, 1288, 1218, 1176, 1080, 1024, 1010, 965, 856, 728; HR-MS (*m/z*): calc for [M+H]⁺ C₁₀H₁₃N₃O₄S 272.0699, found 272.0700.

(*E*)-3-(1-benzyl-1H-imidazol-4-yl)allyl 3-(1-benzyl-1H-imidazol-4-yl)propioate

(228): Ester **224** (1.7 g, 6.6 mmol) was dissolved in a mixture of THF (17 mL)

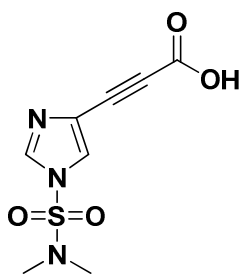


and LiOH (1N in water, 19.6 mL) and stirred at r.t. for 3 h. The pH of the solution was adjusted to pH = 4 through the addition of 1N HCl, then the resulting solution was extracted with EtOAc. The organic solvent was evaporated and a yellow solid

(1.46 g) was obtained which consists of the corresponding acid and a trace of ester. The acid was not purified any further and was used directly in the preparation of the propioate derivative. In a round bottom flask the crude acid **226** (0.90 g, 3.97 mmol), alcohol **212** (1.02 g, 4.77 mmol), DMAP (0.04 g, 0.39

mmol) and camphorsulfonic acid (500 mg, 0.21 mmol) were dissolved in dry CH₂Cl₂ (30 mL). The mixture was cooled to (-78 °C) and DCC (1.23 g, 5.96 mmol) in dry CH₂Cl₂ (8 mL) was added dropwise. The reaction mixture was allowed to warm up to r.t. and stirred for 2 h. The resulting mixture was filtered through Celite and the filter cake was washed with CH₂Cl₂. Concentration of the filtrate provided the crude product, which was purified by chromatography (hexane/EtOAc, 1:19) to provide **228** a thick colorless liquid (1.08 g, 60%) over two steps. ¹H NMR: δ = 7.50 (s, 1H), 7.46 (s, 1H), 7.37-7.31 (m, 6H), 7.28 (s, 1H), 7.17-7.12 (m, 4H), 6.84 (s, 1H), 6.56 (d, *J* = 15.6 Hz, 1H), 6.38 (td, *J* = 16.0, 6.4 Hz, 1H), 5.11 (s, 2H), 5.05 (s, 2H), 4.79 (d, *J* = 6.4 Hz, 2H); ¹³C NMR: δ = 153.9, 139.9, 138.3, 137.9, 135.9, 134.9, 129.3, 129.1, 128.9, 128.4, 127.6, 127.6, 127.4, 126.7, 121.9, 120.6, 117.9, 81.5, 81.2, 66.3, 51.4, 51.0; FT-IR (neat, cm⁻¹): 2988, 2215, 1701, 1540, 1496, 1455, 1377, 1292, 1219, 1148, 1044, 969, 940, 840, 727, 632; HR-MS (*m/z*): calc for [M+H]⁺ C₂₆H₂₂N₄O₂ 423.1816 found 423.1856.

3-(1-(*N,N*-dimethylsulfamoyl)-1H-imidazol-4-yl)propionic acid (225): Ester **223**

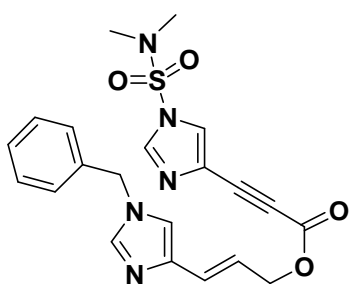


(3.10 g, 11.4 mmol) was dissolved in THF (20 mL) and LiOH (1N in water, 31 mL) and stirred at r.t. for 2.5 h. The pH of the basic solution was adjusted to pH = 4 by the addition of 1N HCl, and the resulting precipitate was collected by vacuum filtration and washed with small amount of cold water. After drying in vacuo, the corresponding acid **225** was

obtained as a colorless solid (2.00 g, 75%). m.p 138-140 °C. ¹H NMR (DMSO, 300 MHz): δ = 8.32 (s, 1H), 8.29 (s, 1H), 2.81 (s, 6H); ¹³C NMR (DMSO, 75 MHz): δ = 154.7, 138.7, 126.7, 122.1, 83.4, 78.6, 39.7; FT-IR (KBr, cm⁻¹): 3393, 3143, 2945, 1693, 1481, 1458, 1426, 1394, 1336, 1282, 1242, 1205, 1187, 1096, 999, 972, 727, 669, 618; HR-MS (*m/z*): calc for [M+H]⁺ C₈H₉N₃O₄S 244.0392 found 244.0387.

(E)-3-(1-benzyl-1H-imidazol-4-yl)allyl-3-(1-(*N,N*-dimethylsulfamoyl)-1H-

imidazol-4-yl)propiolate (227): In a round bottom flask the acid **225** (2.00 g,



8.22 mmol), alcohol **212** (2.11 g, 9.86 mmol), DMAP

(100mg, 0.82 mmol) and camphorsulfonic acid

(110mg, 0.49 mmol) was dissolved in dry CH₂Cl₂ (40

mL) under N₂ atmosphere. The mixture was cooled

to (-78 °C) and DCC (2.54 g, 12.0 mmol) dissolved

in dry CH₂Cl₂ (15 mL) was added dropwise. The reaction mixture was allowed to

warm up to r.t. and stirred for 2 h. The mixture was filtered over Celite and the

filter cake was washed with CH₂Cl₂. The filtrate was concentrated and the crude

product was purified by chromatography (hexane/EtOAc, 1:19) affording **227** as a

white solid (2.40 g, 68%). m.p 118-120 °C. ¹H NMR (300 MHz): δ = 7.85 (d, *J* =

0.9 Hz, 1H), 7.73 (s, 1H), 7.60 (d, *J* = 0.9 Hz, 1H), 7.37-7.32 (m, 3H), 7.17 (d, *J* =

6.9 Hz, 2H), 6.88 (s, 1H), 6.53 (d, *J* = 15.6 Hz, 1H), 6.47 (td, *J* = 15.8, 6.4 Hz, 1H),

5.11 (s, 2H), 4.83 (d, *J* = 6.4 Hz, 2H), 2.88 (s, 6H); ¹³C NMR (75 MHz): δ =

153.4, 138.7, 137.5, 137.1, 135.4, 129.2, 128.7, 127.6, 125.6, 124.7, 123.4,

121.6, 117.9, 82.3, 78.6, 66.5, 51.4, 38.3; FT-IR (neat, cm^{-1}): 3125, 2928, 2222, 1705, 1539, 1456, 1421, 1395, 1332, 1286, 1180, 1080, 1008, 967, 842, 728, 617; HR-MS (m/z): calc for $[\text{M}+\text{H}]^+$ $\text{C}_{21}\text{H}_{21}\text{N}_5\text{O}_4\text{S}$ 440.1387 found 440.1409.

(*R)-1-benzyl-8-(1-benzyl-1H-imidazol-4-yl)-4a,5-dihydro-1H-isobenzofuro[5,6-d]imidazol-7(4H)-one (230):**

CH_2Cl_2 (75 mL) was placed in a resealable thick-walled tube and was purged

with N_2 for 10 min, then ester **228** (0.70 g, 1.65

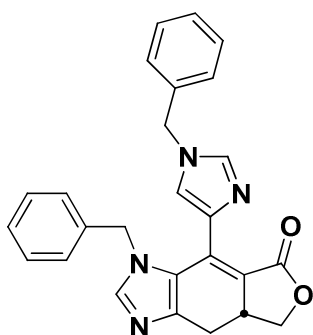
mmol) was added and again the reaction mixture

was purged with N_2 for an additional 5 min. After

sealing the tube with a Teflon screw cap, the

reaction mixture was heated at 130 °C for 12 h. The

reaction mixture was cooled to r.t. and the CH_2Cl_2



was evaporated under vacuum. The crude product was purified by

chromatography (acetone/EtOAc, 7:3) to provide **230** (490 mg, 70%) as a yellow

solid. m.p 202-204 °C. ^1H NMR: δ = 7.59 (s, 1H), 7.55 (s, 1H), 7.50 (s, 1H), 7.38-

7.32 (m, 3H), 7.22 (d, J = 6.4 Hz, 2H), 7.15 (d, J = 7.3 Hz, 1H), 7.06 (t, J = 7.3

Hz, 2H), 6.59 (d, J = 7.3 Hz, 2H), 5.15-5.09 (m, 3H), 4.96 (d, J = 15.6 Hz, 1H),

4.62 (t, J = 9.2 Hz, 1H), 3.98 (t, J = 9.2 Hz, 1H), 3.55-3.49 (m, 1H), 2.98 (dd, J =

15.6, 8.3 Hz, 1H), 2.68 (t, J = 16.0 Hz, 1H); ^{13}C NMR: δ = 168.6, 144.5, 140.9,

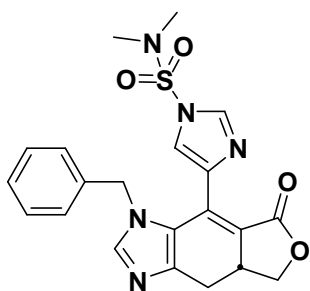
136.5, 136.3, 135.7, 133.0, 131.5, 129.2, 128.6, 127.9, 127.7, 126.4, 125.4,

116.9, 70.9, 51.3, 51.0, 38.4, 27.7; FT-IR (neat, cm^{-1}): 1732, 1604, 1520, 1497,

1455, 1372, 1252, 1223, 1164, 1097, 1075, 1016, 913, 852, 766, 727; HR-MS (m/z): calc for $[M+H]^+$ $C_{26}H_{22}N_4O_2$ 423.1816 found 423.1856.

(*R*^{*})-4-(1-benzyl-7-oxo-4,4a,5,7-tetrahydro-1H-isobenzofuro[5,6-d]imidazol-

8-yl)-*N,N*-dimethyl-1H-imidazole-1-sulfonamide (229): CH_2Cl_2 (80 mL) was



placed in a thick-walled pressure tube and purged with

N_2 for 10 min, then ester **227** (500 mg, 11.4 mmol) was

added and again the reaction mixture was purged with

N_2 for 5 min. After sealing the tube with a Teflon screw

cap, the reaction mixture was heated to 120 °C for 12

h. The reaction mixture was cooled to r.t., and then the reaction mixture was

concentrated. The crude product was purified by chromatography

(acetone/EtOAc, 7:3) to provide a yellow solid (410 mg, 82%). m.p 134-135 °C.

1H NMR: δ = 7.95 (d, J = 1.8 Hz, 1H), 7.52 (s, 1H), 7.34 (d, J = 1.4 Hz, 1H), 7.21-

7.19 (m, 3H), 6.65 (d, J = 6.6 Hz, 2H), 4.99 (d, J = 15.6 Hz, 1H), 4.94 (d, J = 15.6

Hz, 1H), 4.66 (t, J = 8.7 Hz, 1H), 3.99 (t, J = 8.7 Hz, 1H), 3.59-3.55 (quintet, J =

8.7 Hz, 1H), 3.03 (dd, J = 8.7, 8.3 Hz, 1H), 2.88 (s, 6H), 2.70 (dd, J = 16.0, 15.6

Hz, 1H); ^{13}C NMR: δ = 168.4, 144.9, 141.4, 135.9, 135.8, 131.9, 131.2, 128.9,

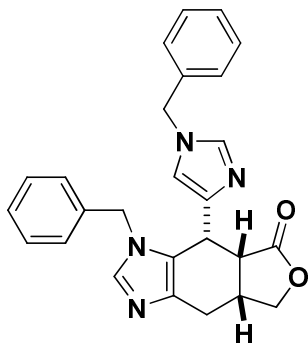
128.1, 126.9, 126.1, 121.8, 118.6, 71.1, 50.9, 38.4, 38.2, 27.7; FT-IR (neat, cm^{-1}):

3122, 2926, 1736, 1615, 1557, 1521, 1457, 1420, 1392, 1334, 1255, 1176,

1083, 1011, 963, 848, 728, 702, 648; HR-MS (m/z): calc for $[M+H]^+$ $C_{21}H_{22}N_5O_4S$

is 440.1387 found 440.1428.

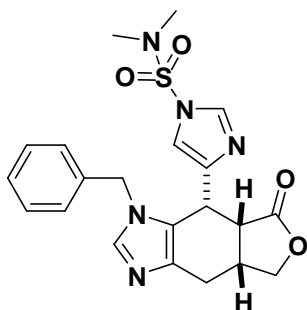
(4aR*,7aS*,8R*)-1-benzyl-8-(1-benzyl-1H-imidazol-4-yl)-4a,5,7a,8-tetrahydro-1H-isobenzofuro[5,6-d]imidazol- 7(4H)-one (232):



The Diels-Alder product **230** (160 mg, 0.37 mmol) was dissolved in dry ethanol (10 mL) and 10% Pd/C (100 mg) was added to the reaction mixture. The heterogeneous reaction mixture was stirred at 36 °C for 8 h under a hydrogen atmosphere (60 psi). The reaction mixture was filtered over Celite and the filter cake was repeatedly washed with hot ethanol. The filtrate was evaporated under reduced pressure followed by purification of the residue by chromatography on silica gel (CHCl₃/MeOH, 49:1) furnished the hydrogenated compound **232** (110 mg, 71%) as a thick colorless liquid. ¹H NMR: δ = 7.45 (s, 1H), 7.37 (s, 1H), 7.34-7.30 (m, 3H), 7.25-7.21 (m, 3H), 7.07 (d, *J* = 7.3 Hz, 2H), 6.86 (d, *J* = 7.8 Hz, 2H), 6.53 (s, 1H), 4.98 (q, *J* = 15.1 Hz, 2H), 4.85 (d, *J* = 15.6 Hz, 1H), 4.57 (d, *J* = 16.0 Hz, 1H), 4.31-4.24 (m, 3H), 3.14-3.01 (m, 3H), 2.74 (d, *J* = 16.0 Hz, 1H); ¹³C NMR (75 MHz): δ = 177.3, 139.8, 137.7, 137.4, 136.2, 136.1, 135.6, 129.1, 128.9, 128.4, 128.1, 127.2, 126.9, 124.5, 118.4, 72.5, 50.9, 48.7, 44.4, 34.9, 30.8, 22.4; FT-IR (neat, cm⁻¹): 2366, 2334, 1773, 1502, 1449, 1207, 1144, 1010, 737, 708. HR-MS (*m/z*): calc for [M+H]⁺ C₂₆H₂₅N₄O₂ 425.1972 found 425.1999.

4-((4a*R,7a*S**,8*R**)-1-benzyl-7-oxo-4,4a,5,7,7a,8-hexahydro-1H-isobenzofuro[5,6-d]imidazol-8-yl)-*N,N*-dimethyl-1H-imidazole-1-sulfonamide**

(231): The Diels-Alder product **229** (200 mg, 0.45 mmol) was dissolved in dry ethanol (15 mL) and then 10% Pd/C (120 mg) was added. The resulting heterogeneous reaction mixture was stirred at 40 °C for 12 h under hydrogen atmosphere (60 psi). On completion of the reaction, the reaction mixture was filtered through Celite and the filter cake was washed repeatedly with hot ethanol.



The filtrate was evaporated under reduced pressure followed by purification of the residue by chromatography on silica gel (CHCl₃/MeOH, 49:1) which furnished reduced compound **231** (150 mg, 75%) as yellow solid. m.p 158-160 °C. ¹H NMR: δ = 7.68 (s, 1H), 7.54 (s, 1H), 7.27-7.26 (m, 3H), 6.92 (m, 2H), 6.91-6.90 (s, 1H), 4.94 (d, *J* = 15.6 Hz, 1H), 4.68 (d, *J* = 15.6 Hz, 1H), 4.28-4.22 (m, 2H), 4.21 (t, *J* = 8.7 Hz, 1H), 3.11-3.06 (m, 3H), 2.80 (d, *J* = 2.8 Hz, 1H), 2.77 (s, 6H); ¹³C NMR: δ = 176.7, 140.6, 138.2, 136.6, 136.2, 135.6, 129.0, 128.3, 126.8, 123.1, 116.4, 72.3, 49.1, 43.8, 38.3, 34.8, 30.6, 22.7; FT-IR (neat, cm⁻¹): 2360, 1770, 1716, 1652, 1558, 1540, 1497, 1457, 1418, 1390, 1268, 1175, 1078, 1008, 962, 728; HR-MS (*m/z*): calc for [M+H]⁺ C₂₁H₂₄N₅O₄S 442.1544 found 442.1581.

(3a*R,4'*S**,6*R**,6a*S**)-1'-benzyl-6-(1-benzyl-1*H*-imidazol-4-yl)-3a,4,6,6a-tetrahydrospiro[cyclopenta[*c*]furan-5,4'-imidazole]-1,5'(1'*H*,3*H*)-dione (234):**

The hydrogenated product **232** (90 mg, 0.21 mmol) was dissolved in chloroform

(5 mL) and Davis' oxaziridine (160 mg, 0.53 mmol)

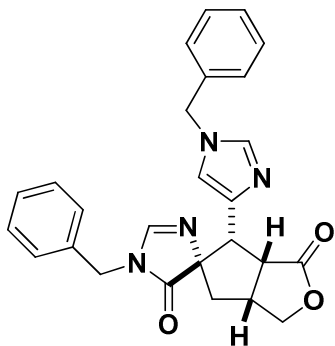
was added to the reaction mixture. The reaction

mixture was stirred at reflux for 8 h. The organic

layer was washed with 2M NaOH solution and the

organic layer was dried with anhydrous Na₂SO₄

concentrated and the residue was purified by



chromatography on silica gel (CHCl₃/MeOH, 49:1) furnished rearranged product

234 (60 mg, 64%) as a yellow solid. ¹H NMR (300 MHz): δ = 7.43 (s, 1H), 7.32-

7.22 (m, 9H), 7.05-6.99 (m, 2H), 6.85 (s, 1H), 4.93 (s, 2H), 4.59 (d, *J* = 15.1 Hz,

1H), 4.53 (d, *J* = 8.6 Hz, 1H), 4.41 (d, *J* = 15.5 Hz, 1H), 4.27 (dd, *J* = 9.3, 3.1 Hz,

1H), 4.07 (d, *J* = 8.3 Hz, 1H), 3.48-3.35 (m, 2H), 2.63 (dd, *J* = 13.4, 8.9 Hz, 1H),

1.86 (d, *J* = 13.4 Hz, 1H); ¹³C NMR (75 MHz): δ = 181.7, 177.0, 152.6, 136.2,

135.7, 135.5, 133.8, 128.9, 128.2, 127.7, 127.3, 119.4, 79.5, 73.6, 52.0, 50.9,

46.3, 44.7, 43.4, 37.4; FT-IR (neat, cm⁻¹): 2363, 2336, 1770, 1733, 1651, 1557,

1542, 1508; HR-MS (*m/z*): calc for [M+H]⁺ C₂₂H₂₅N₄O₃ 441.1927 found 441.1936.

4-((3a*R,4'*S**,6*R**,6a*S**)-1'-benzyl-1,5'-dioxo-1,1',3,3a,4,5',6,6a-octahydrospiro[cyclopenta[c]furan-5,4'-imidazole]-6-yl)-*N,N*-dimethyl-1*H*-imidazole-1-sulfonamide (**233**):**

The tetrahydrobenzimidazole **231** (90 mg, 0.20

mmol) was dissolved in chloroform (3.5 mL) and

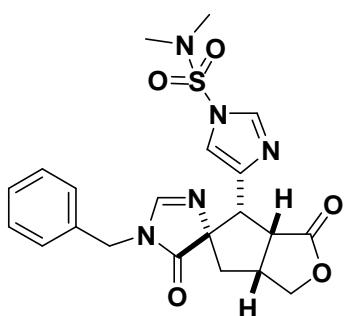
Davis' oxaziridine (150 mg, 0.5 mmol) was added to

the reaction mixture. The resulting mixture was stirred

at reflux for 8 h. The organic layer was washed with

2M NaOH solution and the organic layer was dried

with anhydrous Na₂SO₄ concentrated and the residue



was purified by chromatography on silica gel (CHCl₃/MeOH, 49:1) to provide the

spiro imidazolone **233** (50 mg, 60%) as a colorless solid. m.p 188–190 °C. ¹H

NMR: δ = 7.69 (d, *J* = 0.9 Hz, 1H), 7.60 (s, 1H), 7.29 – 7.25 (m, 3H), 7.09 (d, *J* =

1.4 Hz, 1H), 6.94 (d, *J* = 7.6 Hz, 2H), 4.72 (d, *J* = 15.6 Hz, 1H), 4.56 (t, *J* = 8.7

Hz, 1H), 4.42 (d, *J* = 15.1 Hz, 1H), 4.26 (dd, *J* = 9.2, 3.8 Hz, 1H), 4.06 (d, *J* = 8.3

Hz, 1H), 3.43-3.41 (m, 2H), 2.78 (s, 6H), 2.61 (dd, *J* = 15.3, 9.2 Hz, 1H), 1.88 (d,

J = 13.7 Hz, 1H); ¹³C NMR: δ = 181.1, 176.7, 153.4, 135.1, 134.9, 129.2, 128.4,

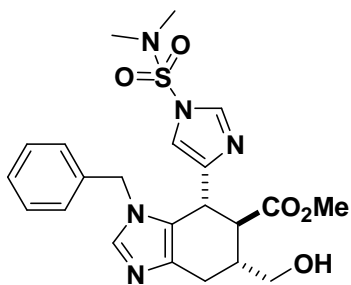
127.3, 117.2, 79.1, 73.7, 51.1, 46.1, 44.9, 43.7, 38.3, 37.2; FT-IR (neat, cm⁻¹):

2932, 1766, 1727, 1598, 1390, 1162, 1079, 954, 726; HR-MS (*m/z*): calc for

[*M*+*H*]⁺ C₂₁H₂₃N₅O₅S 458.1471 found 458.1471.

(5*R,6*R**,7*R**)-methyl-1-benzyl-7-(1-(*N,N*-dimethylsulfamoyl)-1*H*-imidazol-4-yl)-5-(hydroxymethyl)-4,5,6,7-tetrahydro-1*H*-benzo[d]imidazole-6-**

carboxylate (235): The tetrahydrobenzimidazole **231** (200 mg, 0.42 mmol) was



dissolved in MeOH (60 mL) under nitrogen atmosphere. To this reaction mixture 0.23 M sodium methoxide was added dropwise. After stirring this reaction at r.t. for 1 h it was heated to 60 °C for 2 h.

Then cooling this reaction mixture to room

temperature, (20 mL) of saturated ammonium chloride solution was added followed by addition of equal amount of water. Then this aqueous solution was

extracted repeatedly by EtOAc. The organic extracts were dried with anhydrous

Na₂SO₄ concentrated and the residue was purified by chromatography on silica

gel (CH₂Cl₂/MeOH, 4:1) furnishing the product **235** (120 mg, 52%) as a colorless

oil. In addition, unreacted lactone **231** (70 mg, 35%) was recovered. ¹H NMR: δ

= 7.74 (s, 1H), 7.40 (s, 1H), 7.31-7.25 (m, 3H), 6.85 (d, *J* = 7.3 Hz, 2H), 6.81 (s, 1H), 4.85 (d, *J* = 16.9 Hz, 1H), 4.48 (d, *J* = 18.3 Hz, 1H), 4.20 (d, *J* = 10.5 Hz,

1H), 3.64–3.59 (m, 2H), 3.57 (s, 3H), 3.00 (t, *J* = 9.2 Hz, 1H), 2.82–2.78 (m,

1H), 2.75 (s, 6H), 2.72–2.69 (m, 1H) 2.37–2.34 (m, 1H); ¹³C NMR: δ = 174.5,

142.7, 137.9, 137.2, 136.1, 129.0, 127.9, 126.3, 115.4, 65.1, 51.9, 51.6, 49.2,

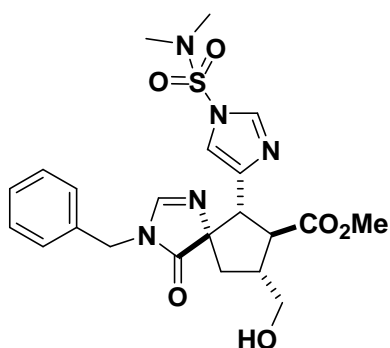
40.6, 38.1, 35.8, 26.8; FT-IR (neat, cm⁻¹): 3112, 2935, 1723, 1455, 1394, 1265,

1170, 1074, 953, 733, 603, 598; calc for [M+H]⁺ C₂₂H₂₇N₅O₅S 474.1806 found

474.1830.

(5*S,6*R**,7*R**,8*R**)-methyl-3-benzyl-6-(1-(*N,N*-dimethylsulfamoyl)-1*H*-imidazol-4-yl)-8-(hydroxymethyl)-4-oxo-1,3-diazaspiro[4.4]non-1-ene-7-**

carboxylate (236): The spiro imidazolone **233** (200 mg, 0.41 mmol) was

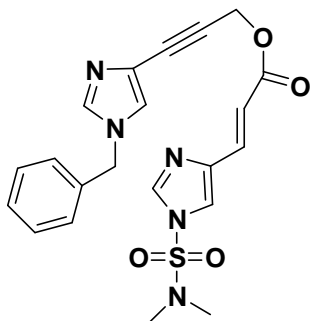


dissolved in MeOH (60 mL) under a nitrogen atmosphere. To this reaction mixture 0.23 M sodium methoxide in MeOH (49 mL) was added dropwise. After stirring this reaction at r.t. for 1 h, it was heated to 60 °C for 18 h. The reaction mixture was cooled to r.t., then saturated NH₄Cl

solution (20 mL) was added followed by addition of equal amount of water. The resulting aqueous solution was extracted repeatedly with EtOAc. The organic extracts were dried with anhydrous Na₂SO₄ and concentrated and the residue was purified by chromatography on silica gel (CH₂Cl₂/MeOH; 44:6) furnished the desired product **236** (65 mg, 31%) as a colorless liquid. In addition, unreacted lactone **233** (110 mg, 55%) was recovered. ¹H NMR: δ = 7.60 (s, 1H), 7.39 (s, 1H), 7.34-7.28 (m, 3H), 7.15 (d, *J* = 7.2 Hz, 2H), 7.00 (s, 1H), 4.71 (d, *J* = 15.1 Hz, 1H), 4.50 (d, *J* = 15.1 Hz, 1H), 4.02 (d, *J* = 11.0 Hz, 1H), 3.80-3.78 (m, 2H), 3.68 (s, 3H), 3.66-3.62 (m, 1H), 2.93-2.91 (m, 1H), 2.74 (s, 6H), 2.65-2.63 (m, 1H), 1.74 (d, *J* = 13.8 Hz, 1H); ¹³C NMR (75 MHz): δ = 180.9, 174.5, 152.6, 139.8, 135.8, 134.9, 129.1, 128.3, 127.9, 114.5, 79.0, 65.3, 52.2, 51.0, 48.6, 44.9, 43.8, 39.9, 38.2; FT-IR (neat, cm⁻¹): 2944, 1735, 1397, 1166, 1091, 962, 730; calc for [M+H]⁺ C₂₂H₂₇N₅O₆S 490.1755 found 490.1758.

(E)-3-(1-benzyl-1H-imidazol-4-yl)prop-2-ynyl-3-(1-(N,N-dimethylsulfamoyl)-

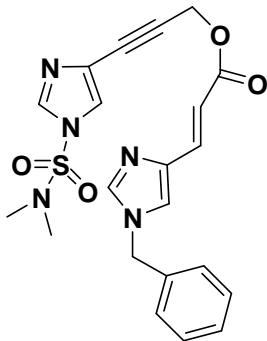
1H-imidazol-4-yl)acrylate (239): In a round bottom flask the alcohol **238** (200



mg, 0.94 mmol), acid **237** (350 mg, 1.41 mmol), DMAP (10 mg, 0.09 mmol) and camphorsulfonic acid (13 mg, 0.06 mmol) was dissolved in dry CH₂Cl₂ (20 mL) under N₂ atmosphere. The mixture was cooled to (-78 °C) and DCC (291 mg, 1.41 mmol) dissolved in dry CH₂Cl₂

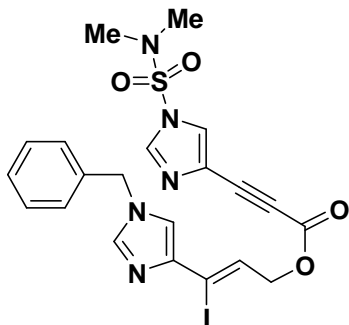
(5 mL) was added dropwise. The reaction mixture was allowed to warm up to r.t. and stirred for 2 h. The mixture was filtered over Celite and the filter cake was washed with CH₂Cl₂. The filtrate was concentrated and the crude product was purified by chromatography (hexane/EtOAc, 1:19) affording **239** as a colorless thick liquid (240 mg, 60%). ¹H NMR (300 MHz): δ = 7.87 (s, 1H), 7.54 (d, *J* = 16.0 Hz, 1H), 7.46 (d, *J* = 0.9 Hz, 1H), 7.38-7.33 (m, 4H), 7.15 (dd, *J* = 1.8, 8.0 Hz, 2H), 7.10 (d, *J* = 1.4 Hz, 1H), 6.68 (d, *J* = 15.6 Hz, 1H), 5.07 (s, 2H), 4.99 (s, 2H), 2.88 (s, 6H); ¹³C NMR (75 MHz): δ = 166.1, 139.4, 137.7, 137.5, 135.4, 135.2, 129.2, 128.7, 127.5, 124.1, 124.0, 118.9, 118.8, 83.2, 80.7, 52.7, 50.6, 38.3; FT-IR (neat, cm⁻¹): 3125, 2936, 2363, 1712, 1486, 1391, 1245, 1150, 1007, 775, 637, 592; HR-MS (*m/z*): calc for [M+H]⁺ C₂₁H₂₂N₅O₄S 440.1393 found 440.1359.

(E)-3-(1-(*N,N*-dimethylsulfamoyl)-1H-imidazol-4-yl)prop-2-ynyl-3-(1-benzyl-1H-imidazol-4-yl)acrylate (242**):**



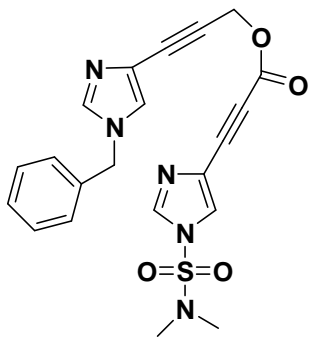
In a round bottom flask the alcohol **241** (200 mg, 0.87 mmol), acid **212** (290 mg, 1.31 mmol), DMAP (9 mg, 0.08 mmol) and camphorsulfonic acid (11 mg, 0.05 mmol) was dissolved in dry CH₂Cl₂ (20 mL) under N₂ atmosphere. The mixture was cooled to (-78 °C) and DCC (270 mg, 1.31 mmol) dissolved in dry CH₂Cl₂ (5 mL) was added dropwise. The reaction mixture was allowed to warm up to r.t. and stirred for 2 h. The mixture was filtered over Celite and the filter cake was washed with CH₂Cl₂. The filtrate was concentrated and the crude product was purified by chromatography (hexane/EtOAc, 1:19) affording **242** as a colorless thick liquid (249 mg, 65%). ¹H NMR (300 MHz): δ = 7.82 (s, 1H), 7.57 (d, *J* = 15.6 Hz, 1H), 7.55 (s, 1H), 7.39-7.36 (m, 4H), 7.18 (d, *J* = 7.8 Hz, 2H), 7.10 (s, 1H), 6.56 (d, *J* = 16.0 Hz, 1H), 5.10 (s, 2H), 4.98 (s, 2H), 2.87 (s, 6H); ¹³C NMR (75 MHz): δ = 166.7, 138.9, 138.7, 137.3, 136.6, 135.4, 129.2, 128.7, 127.5, 125.5, 122.0, 121.3, 115.2, 85.6, 78.3, 52.4, 51.2, 38.3; FT-IR (neat, cm⁻¹): 3125, 2928, 2241, 1709, 1479, 1387, 1255, 1149, 1097, 1006, 961, 831, 728, 631; HR-MS (*m/z*): calc for [M+H]⁺ C₂₁H₂₂N₅O₄S 440.1393 found 440.1358.

(Z)-3-(1-benzyl-1H-imidazol-4-yl)-3-iodoallyl-3-(1-(*N,N*-dimethylsulfamoyl)-1H-imidazol-4-yl)propiolate (250**):**



In a round bottom flask the alcohol **245** (200 mg, 0.58 mmol), acid **225** (210 mg, 0.88 mmol), DMAP (7 mg, 0.06 mmol) and camphorsulfonic acid (8 mg, 0.04 mmol) was dissolved in dry CH₂Cl₂ (15 mL) under N₂ atmosphere. The mixture was cooled to (-78 °C) and DCC (180 mg, 0.88 mmol) dissolved in dry CH₂Cl₂ (5 mL) was added dropwise. The reaction mixture was allowed to warm up to r.t. and stirred for 8 h. The mixture was filtered over Celite and the filter cake was washed with CH₂Cl₂. The filtrate was concentrated and the crude product was purified by chromatography (hexane/EtOAc, 2:18) affording **250** as a white solid (230 mg, 68%). mp 135-137 °C. ¹H NMR (300 MHz): δ = 7.86 (s, 1H), 7.61 (d, *J* = 0.9 Hz, 1H), 7.56 (s, 1H), 7.38-7.34 (m, 3H), 7.18 (d, *J* = 6.9 Hz, 2H), 7.11 (s, 1H), 6.83 (t, *J* = 15.6 Hz, 1H), 5.08 (s, 2H), 4.96 (d, *J* = 6.4 Hz, 2H), 2.90 (s, 6H); ¹³C NMR (75 MHz): δ = 153.4, 140.9, 137.6, 137.2, 134.6, 129.9, 129.5, 129.2, 127.9, 124.9, 123.4, 122.1, 82.2, 79.2, 70.1, 52.1, 38.4; FT-IR (neat, cm⁻¹): 3138, 2849, 2219, 1703, 1453, 1386, 1333, 1212, 1167, 1086, 1048, 995, 850, 738, 617; HR-MS (*m/z*): calc for [M+H]⁺ C₂₁H₂₁IN₅O₄S 566.0353 found 566.0352.

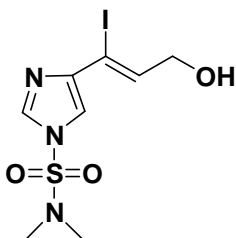
3-(1-benzyl-1H-imidazol-4-yl)prop-2-ynyl-3-(1-(*N,N*-dimethylsulfamoyl)-1H-imidazol-4-yl)propiolate (251):



CH₂Cl₂ (10 mL) was placed in a thick-walled pressure tube and purged with N₂ for 10 min, then ester **250** (120 mg, 11.4 mmol) was added and again the reaction mixture was purged with N₂ for 5 min. After sealing the tube with a Teflon screw cap, the reaction mixture was heated to 120 °C for 12 h. The reaction mixture was cooled to r.t., and then the reaction mixture was concentrated. The crude product was purified by chromatography (acetone/EtOAc, 7:3) to provide a yellow solid **251** (69 mg, 75%). m.p 209-211 °C. ¹H NMR (300 MHz): δ = 8.14 (s, 1H), 8.00 (d, *J* = 0.9 Hz, 1H), 7.87 (s, 1H), 7.19-7.16 (m, 3H), 6.88 (d, *J* = 0.9 Hz, 1H), 6.58 (d, *J* = 6.9 Hz, 2H), 5.38 (s, 2H), 5.33 (s, 2H), 2.88 (s, 6H); ¹³C NMR (75 MHz): δ = 170.0, 150.0, 149.2, 140.5, 135.8, 135.5, 133.4, 132.2, 128.9, 128.1, 125.3, 119.5, 118.5, 118.4, 113.7, 68.0, 50.7, 38.3; FT-IR (neat, cm⁻¹): 3101, 2927, 2241, 2219, 1759, 1501, 1456, 1390, 1264, 1167, 1077, 965, 734, 593; HR-MS (*m/z*): calc for [M+H]⁺ C₂₁H₂₀N₅O₄S 438.1236 found 438.1241.

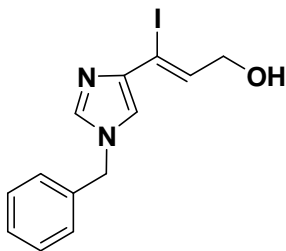
(Z)-4-(3-hydroxy-1-iodoprop-1-enyl)-N,N-dimethyl-1H-imidazole-1-

sulfonamide (244):



In a round bottom flask compound **241** (1.0 g, 4.36 mmol) was dissolved under N₂ atmosphere in anhydrous THF (50 mL). The reaction mixture was cooled to 0 °C and then to it Red-Al (65 wt.% in toluene) (1.63 mL, 5.20 mmol) was added dropwise. After stirring the reaction mixture for 30 min, N-Iodosuccinimide (1.27 g, 5.60 mmol) dissolved in anhydrous THF (4 mL) was added. Then the reaction mixture was stirred for another 1 h at r.t. Finally the reaction was quenched with NH₄Cl (10 mL) then the organic layer was separated. The aqueous layer was extracted with EtOAc (3x50 mL), then the combined organic solutions were dried with anhydrous Na₂SO₄ and concentrated. The crude product was purified by chromatography (EtOAc/Hexane, 95:5) providing **244** as a yellow solid (1.27 g, 82%). The proton and carbon contains residual acetone impurity. m.p 94-96 °C. ¹H NMR (300 MHz): δ = 7.95 (d, *J* = 1.4 Hz, 1H), 7.34 (d, *J* = 1.4 Hz, 1H), 6.99 (t, *J* = 5.8 Hz, 1H), 4.39 (d, *J* = 6.2 Hz, 2H), 2.89 (s, 6H); ¹³C NMR (75 MHz): δ = 144.2, 137.1, 136.2, 118.5, 91.1, 67.2, 38.3; FT-IR (neat, cm⁻¹): 3250, 1392, 1167, 1079, 962, 599, 508; HR-MS (*m/z*): calc for [M+H]⁺ C₈H₁₃IN₃O₃S 357.9722 found 357.9723.

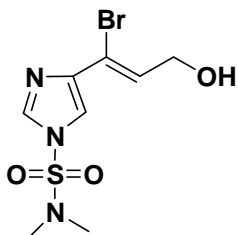
(Z)-3-(1-benzyl-1H-imidazol-4-yl)-3-iodoprop-2-en-1-ol (245):



In a round bottom flask compound **238** (1.2 g, 5.65 mmol) was dissolved under N₂ atmosphere in anhydrous THF (70 mL). The reaction mixture was cooled to 0 °C and then to it Red-Al (65 wt.% in toluene) (2.10 mL, 6.78 mmol) was added dropwise. After stirring the reaction mixture for 30 min, N-Iodosuccinimide (1.65 g, 7.35 mmol) dissolved in anhydrous THF (10 mL) was added. Then the reaction mixture was stirred for another 1 h at r.t. Finally the reaction was quenched with NH₄Cl (10 mL) then the organic layer was separated. The aqueous layer was extracted with EtOAc (3x50 mL), then the combined organic solutions were dried with anhydrous Na₂SO₄ and concentrated. The crude product was purified by chromatography (EtOAc 100%) providing **245** as a yellow solid (1.68 g, 88%). m.p 84-86 °C. ¹H NMR (300 MHz): δ = 7.56 (d, *J* = 1.4 Hz, 1H), 7.37-7.32 (m, 3H), 7.18 (dd, *J* = 2.1, 7.4 Hz, 2H), 7.05 (d, *J* = 1.4 Hz, 1H), 6.83 (t, *J* = 6.2 Hz, 1H), 5.10 (s, 2H), 4.35 (d, *J* = 6.2 Hz, 2H), 2.95 (s, 1H); ¹³C NMR (75 MHz): δ = 143.4, 137.9, 135.6, 133.4, 129.3, 128.6, 127.5, 121.2, 94.0, 67.3, 51.2; FT-IR (neat, cm⁻¹): 3192, 2850, 1633, 1493, 1457, 1355, 1156, 1041, 940, 840, 716; HR-MS (*m/z*): calc for [M+H]⁺ C₁₃H₁₄IN₂O 341.0151 found 341.0146.

(Z)-4-(1-bromo-3-hydroxyprop-1-enyl)-N,N-dimethyl-1H-imidazole-1-sulfonamide (259):

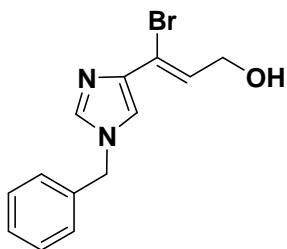
In a round bottom flask compound **241** (300 mg, 1.31 mmol) was dissolved under N₂ atmosphere in anhydrous THF (20 mL). The reaction



mixture was cooled to 0 °C and then to it Red-Al (65 wt.% in toluene) (0.5 mL, 1.57 mmol) was added dropwise. After stirring the reaction mixture for 30 min, N-bromosuccinimide

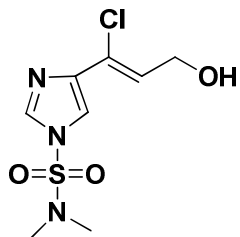
(310 mg, 1.70 mmol) dissolved in anhydrous THF (5 mL) was added. Then the reaction mixture was stirred for another 1 h at r.t. Finally the reaction was quenched with NH₄Cl (10 mL) then the organic layer was separated. The aqueous layer was extracted with EtOAc (3x50 mL), the combined organic solutions were dried with anhydrous Na₂SO₄ and concentrated. The crude product was purified by chromatography (EtOAc 100%) providing **259** as a yellow solid (315mg, 78%). m.p 101-103 °C. ¹H NMR: δ = 7.88 (d, *J* = 1.4 Hz, 1H), 7.34 (s, 1H), 7.05 (t, *J* = 5.9 Hz, 1H), 4.43 (d, *J* = 5.5 Hz, 2H), 2.87 (s, 6H); ¹³C NMR: δ = 142.1, 137.2, 129.9, 116.8, 114.9, 62.2, 38.3; FT-IR (neat, cm⁻¹): 2980, 2934, 1501, 1381, 1258, 1188, 1129, 839; HR-MS (*m/z*): calc for [M+H]⁺ C₈H₁₃BrN₃O₃S 309.9861 found 309.9870.

(Z)-3-(1-benzyl-1H-imidazol-4-yl)-3-bromoprop-2-en-1-ol (260):



In a round bottom flask compound **238** (200 mg, 0.94 mmol) was dissolved under N₂ atmosphere in anhydrous THF (20 mL). The reaction mixture was cooled to 0 °C and then to it Red-Al (65 wt.% in toluene) (0.35 mL, 1.13 mmol) was added dropwise. After stirring the reaction mixture for 30 min, N-bromosuccinimide (250 mg, 1.41 mmol) dissolved in anhydrous THF (2.5 mL) was added. Then the reaction mixture was stirred for another 1 hr at r.t. Finally the reaction was quenched with NH₄Cl (10 mL) then the organic layer was separated. The aqueous layer was extracted with EtOAc (3x50 mL), then the combined organic solutions were dried with anhydrous Na₂SO₄ and concentrated. The crude product was purified by chromatography (EtOAc 100%) providing **260** as a yellow thick liquid (220 mg, 81%). ¹H NMR (300 MHz): δ = 7.52 (d, *J* = 1.4 Hz, 1H), 7.39-7.33 (m, 3H), 7.18 (dd, *J* = 2.1, 7.4 Hz, 2H), 7.10 (t, *J* = 1.0 Hz, 1H), 6.94 (t, *J* = 6.2 Hz, 1H), 5.08 (s, 2H), 4.43 (d, *J* = 6.2 Hz, 2H), 1.94 (b, 1H); ¹³C NMR (75 MHz): δ = 141.7, 137.9, 135.6, 129.2, 128.6, 127.4, 126.2, 119.2, 117.2, 62.4, 51.2; FT-IR (neat, cm⁻¹): 3246, 2935, 1640, 1537, 1454, 1231, 1160, 1043, 830, 711; HR-MS (*m/z*): calc for [M+H]⁺ C₁₃H₁₄BrN₂O 293.0284 found 293.0281.

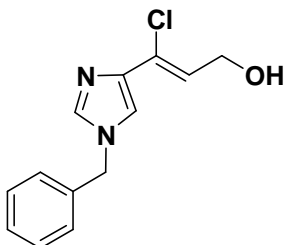
(Z)-4-(1-chloro-3-hydroxyprop-1-enyl)-N,N-dimethyl-1H-imidazole-1-sulfonamide (263):



In a round bottom flask compound **241** (300 mg, 1.31 mmol) was dissolved under N₂ atmosphere in anhydrous THF (25 mL). The reaction mixture was cooled to 0 °C and then to it Red-Al (65 wt.% in toluene) (0.50 mL, 1.57 mmol) was added dropwise. After stirring the reaction mixture for 30 min, N-chlorosuccinimide (250 mg, 1.70 mmol) dissolved in anhydrous THF (5 mL) was added. Then the reaction mixture was stirred for another 1 h at r.t. Finally the reaction was quenched with NH₄Cl (10 mL) then the organic layer was separated. The aqueous layer was extracted with EtOAc (3x50 mL), then the combined organic solutions were dried with anhydrous Na₂SO₄ and concentrated. The crude product was purified by chromatography (EtOAc 100%) providing **263** as a yellow solid (120 mg, 34%). ¹H NMR: δ = 7.85 (d, *J* = 1.4 Hz, 1H), 7.37 (d, *J* = 1.0 Hz, 1H), 6.84 (t, *J* = 6.2 Hz, 1H), 4.50 (d, *J* = 6.2 Hz, 2H), 2.90 (s, 6H), 5; ¹³C NMR (75 MHz): δ = 141.3, 137.2, 125.9, 125.1, 115.6, 59.7, 38.3; FT-IR (neat, cm⁻¹): 3356, 2987, 1635, 1396, 1173, 1082, 968, 731, 599; HR-MS (*m/z*): calc for [M+H]⁺ C₈H₁₃ClN₃O₃S 266.0361 found 266.0357.

(Z)-3-(1-benzyl-1H-imidazol-4-yl)-3-chloroprop-2-en-1-ol (264):

In a round bottom flask compound **238** (250 mg, 1.17 mmol) was dissolved under



N_2 atmosphere in anhydrous THF (20 mL). The reaction

mixture was cooled to 0 °C and then to it Red-Al (65 wt.% in toluene) (0.45 mL, 1.41 mmol) was added dropwise.

After stirring the reaction mixture for 30 min, N-

chlorosuccinimide (210 mg, 1.53 mmol) dissolved in anhydrous THF (2.5 mL)

was added. Then the reaction mixture was stirred for another 1 hr at r.t. Finally

the reaction was quenched with NH_4Cl (10 mL) then the organic layer was

separated. The aqueous layer was extracted with EtOAc (3x50 mL), then the

combined organic solutions were dried with anhydrous Na_2SO_4 and

concentrated. The crude product was purified by chromatography (EtOAc 100%)

providing **264** as a yellow thick liquid (102 mg, 35%). 1H NMR (300 MHz): δ =

7.98 (s, 1H), 7.39-7.36 (m, 3H), 7.23 (dd, J = 2.1, 7.4 Hz, 2H), 7.01 (s, 1H), 6.85

(t, J = 6.2 Hz, 1H), 5.17 (s, 2H), 4.44 (d, J = 6.2 Hz, 2H), 4.03 (b, 1H); ^{13}C

NMR (75 MHz): δ = 138.2, 137.5, 134.7, 129.3, 128.9, 127.9, 125.6, 124.1,

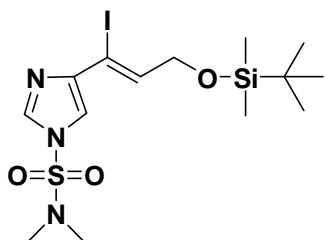
118.5, 59.6, 51.9; FT-IR (neat, cm^{-1}): 3230, 2948, 1610, 1512, 1415, 1331, 1140,

1080, 835, 712, 623; HR-MS (m/z): calc for $[M+H]^+ C_{13}H_{14}ClN_2O$ 249.0797 found

249.0790.

(Z)-4-(3-(tert-butyldimethylsilyloxy)-1-iodoprop-1-enyl)-N,N-dimethyl-1H-imidazole-1-sulfonamide (255):

In a round bottom flask compound **244** (400 mg, 1.12 mmol) was dissolved in



CH₂Cl₂ (20 mL) under nitrogen atmosphere. The

solution was cooled to 0 °C and then to the reaction

mixture imidazole (110 mg, 1.68 mmol) and TBSCl

(220 mg, 1.45 mmol) were added. The reaction

mixture was stirred for 8 h, then NH₄Cl (5 mL) was

added and the organic layer was separated. The aqueous layer was extracted

with EtOAc (3x50 mL), then the combined organic solutions were dried with

anhydrous Na₂SO₄ and concentrated. The crude product was purified by

chromatography (EtOAc 100%) providing **255** as a colorless solid (480 mg,

92%). m.p 75-77 °C ¹H NMR (300 MHz): δ = 7.88 (d, *J* = 1.0 Hz, 1H), 7.31 (d, *J* =

1.4 Hz, 1H), 6.89 (t, *J* = 5.5 Hz, 1H), 4.39 (d, *J* = 5.5 Hz, 2H), 2.86 (s, 6H), 0.89

(s, 9H), 0.08 (s, 6H); ¹³C NMR (75 MHz): δ = 144.3, 137.4, 136.9, 117.9, 89.4,

68.4, 38.3, 26.0, 18.4, -5.1 ; FT-IR (neat, cm⁻¹): 2934, 2852, 1703, 1592, 1396,

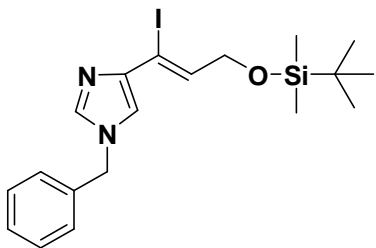
1173, 1148, 1079, 833, 731, 596; HR-MS (*m/z*): calc for [M+H]⁺ C₁₄H₂₇IN₃O₃SSi

472.0582 found 472.0596.

(Z)-1-benzyl-4-(3-(tert-butyldimethylsilyloxy)-1-iodoprop-1-enyl)-1H-imidazole (253):

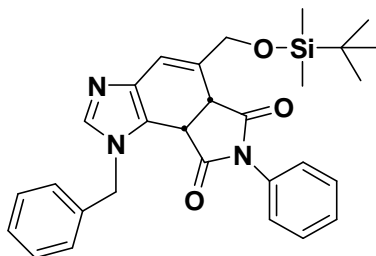
In a round bottom flask compound **245** (0.77 g, 2.25 mmol) was dissolved in

CH₂Cl₂ (25 mL) under a nitrogen atmosphere. The solution was cooled to 0 °C



and then to the reaction mixture imidazole (240 mg, 3.60 mmol) and TBSCl (440, 2.94 mmol) was added. The reaction mixture was stirred for another 8 h, then NH₄Cl (10 mL) was added and the organic layer was separated. The aqueous layer was extracted with EtOAc (3x50 mL), then the combined organic solutions were dried with anhydrous Na₂SO₄ and concentrated. The crude product was purified by chromatography (EtOAc 100%) providing **253** as a colorless thick liquid (920 mg, 90%). ¹H NMR (300 MHz): δ = 7.55 (d, *J* = 1.4 Hz, 1H), 7.40-7.33 (m, 3H), 7.16 (dd, *J* = 7.6, 1.4 Hz, 2H), 7.05 (*J* = 1.4 Hz, 1H), 6.73 (t, *J* = 5.5 Hz, 1H), 5.10 (s, 2H), 4.40 (d, *J* = 5.5 Hz, 2H), 0.92 (s, 9H), 0.10 (s, 6H); ¹³C NMR (75 MHz): δ = 143.6, 137.7, 134.0, 129.2, 128.5, 127.4, 120.3, 92.3, 77.8, 68.1, 50.9, 25.9, 18.4, 17.9, -5.1; FT-IR (neat, cm⁻¹): 2928, 2849, 1498, 1449, 1258, 1097, 1065, 842, 769, 707; HR-MS (*m/z*): calc for [M+H]⁺ C₁₉H₂₈IN₂OSi 455.1010 found 455.1014.

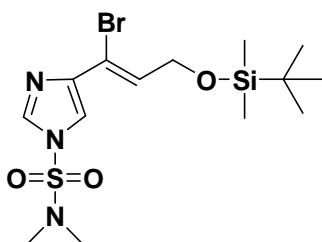
(5aR*,8aS*)-1-benzyl-5-((tert-butyldimethylsilyloxy)methyl)-7-phenyl-7,8a-dihydroimidazo[4,5-e]isoindole-6,8(1H,5aH)-dione (257):



CH₂Cl₂ (3 mL) was placed in a resealable thick-walled tube and was purged with N₂ for 5 minutes, then compound **253** (140 mg, 0.31 mmol) and compound **256** (65 mg, 0.37 mmol) was added and again the reaction mixture was purged with N₂ for an additional 5 minutes. After sealing the tube with a Teflon screw cap, the

reaction mixture was heated at 50 °C for 12 h. The reaction mixture was cooled to r.t. and the CH₂Cl₂ was evaporated under vacuum. The crude product was purified by chromatography (EtOAc/Hexane, 1:1) to provide **257** (69 mg, 45%) as a yellow solid. m.p 202-204 °C. ¹H NMR: δ = 7.52 (s, 1H), 7.49-7.46 (m, 2H), 7.42-7.32 (m, 4H), 7.24 (d, *J* = 7.8 Hz, 2H), 7.17 (d, *J* = 7.3 Hz, 2H), 6.67 (s, 1H), 5.77 (d, *J* = 15.6 Hz, 1H), 5.28 (d, *J* = 15.6 Hz, 1H), 4.53 (s, 2H), 4.20 (d, *J* = 11.4 Hz, 1H), 4.09 (d, *J* = 11.4 Hz, 1H), 0.92 (s, 9H), 0.10 (s, 6H); ¹³C NMR: δ = 175.3, 175.1, 139.6, 137.3, 135.9, 131.7, 129.3, 129.2, 129.0, 128.8, 128.3, 127.5, 126.5, 119.1, 118.9, 64.6, 50.0, 43.4, 39.8, 26.1, 18.5, -5.2; FT-IR (neat, cm⁻¹): 2934, 2858, 1721, 1498, 1375, 1194, 1123, 839; HR-MS (*m/z*): calc for [M+H]⁺ C₂₉H₃₄N₃O₃Si 500.2364 found 500.2371.

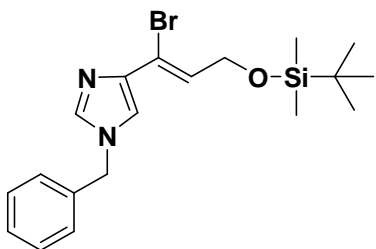
(Z)-4-(1-bromo-3-(tert-butyldimethylsilyloxy)prop-1-enyl)-N,N-dimethyl-1H-imidazole-1-sulfonamide (261):



In a round bottom flask compound **259** (280 mg, 0.90 mmol) was dissolved in CH₂Cl₂ (15 mL) under a nitrogen atmosphere. The solution was cooled to 0 °C and then to the reaction mixture imidazole (90 mg, 1.35 mmol) and TBSCl (177 mg, 1.17 mmol) was added. The reaction mixture was stirred for 8 h, then NH₄Cl (5 mL) was added and the organic layer was separated. The aqueous layer was extracted with EtOAc (3x50 mL), then the combined organic solutions were dried with anhydrous Na₂SO₄ and concentrated. The crude product was purified by chromatography (EtOAc 100%)

providing **261** as a colorless solid (336 mg, 88%). m.p 91-93 °C. ¹H NMR: δ = 7.86 (d, *J* = 1.4 Hz, 1H), 7.34 (d, *J* = 1.4 Hz, 1H), 6.97 (t, *J* = 5.5 Hz, 1H), 4.48 (d, *J* = 5.5 Hz, 2H), 2.88 (s, 6H), 0.92 (s, 9H), 0.10 (s, 6H); ¹³C NMR: δ = 142.4, 137.0, 130.9, 116.3, 113.3, 63.4, 38.3, 26.0, 18.4, -5.1; FT-IR (neat, cm⁻¹): 2938, 2845, 1592, 1397, 1165, 1150, 1085, 987, 840, 717, 590; HR-MS (*m/z*): calc for [M+H]⁺ C₁₄H₂₇BrN₃O₃SSi 424.0726 found.

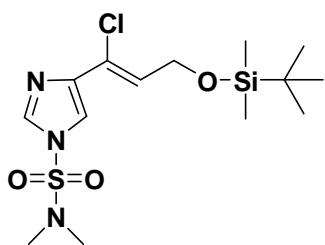
(Z)-1-benzyl-4-(1-bromo-3-(tert-butyldimethylsilyloxy)prop-1-enyl)-1H-imidazole (262):



In a round bottom flask compound **260** (170 mg, 0.58 mmol) was dissolved in CH₂Cl₂ (10 mL) under a nitrogen atmosphere. The solution was cooled to 0 °C and then to the reaction mixture imidazole (60 mg, 3.60 mmol) and TBSCl (110 mg, 0.75 mmol) was added. The reaction mixture was stirred for 8 h, then NH₄Cl (5 mL) was added and the organic layer was separated. The aqueous layer was extracted with EtOAc (3x50 mL), then the combined organic solutions were dried with anhydrous Na₂SO₄ and concentrated. The crude product was purified by chromatography (EtOAc 100%) providing **262** as a colorless thick liquid (210 mg, 91%). ¹H NMR (300 MHz): δ = 7.51 (d, *J* = 1.4 Hz, 1H), 7.37-7.34 (m, 3H), 7.16 (dd, *J* = 7.6, 1.4 Hz, 2H), 7.05 (*J* = 1.4 Hz, 1H), 6.83 (t, *J* = 5.5 Hz, 1H), 5.07 (s, 2H), 4.46 (d, *J* = 5.5 Hz, 2H), 0.90 (s, 9H), 0.09 (s, 6H); ¹³C NMR (75 MHz): δ = 141.5, 137.8, 135.7, 129.2, 128.6, 127.6, 127.4, 119.1, 115.3, 63.6, 51.2, 26.0,

18.4, -5.1; FT-IR (neat, cm^{-1}): 2943, 2840, 1604, 1495, 1457, 1255, 1115, 1065, 781, 704; HR-MS (m/z): calc for $[\text{M}+\text{H}]^+$ $\text{C}_{19}\text{H}_{28}\text{BrN}_2\text{OSi}$ 407.1149 found 407.1136 .

(Z)-4-(3-(tert-butyldimethylsilyloxy)-1-chloroprop-1-enyl)-N,N-dimethyl-1H-imidazole-1-sulfonamide (265):

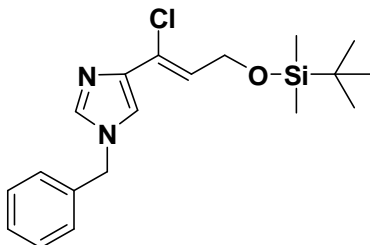


In a round bottom flask compound **263** (130 mg, 0.49 mmol) was dissolved in CH_2Cl_2 (10 mL) under a nitrogen atmosphere. The solution was cooled to 0 °C and then to the reaction mixture imidazole (50 mg, 0.73 mmol) and TBSCl (100 mg, 0.64 mmol) was added.

The reaction mixture was stirred for 8 h, then NH_4Cl (5 mL) was added and the organic layer was separated. The aqueous layer was extracted with EtOAc (3x50 mL), then the combined organic solutions were dried with anhydrous Na_2SO_4 and concentrated. The crude product was purified by chromatography (EtOAc 100%) providing **265** as a colorless solid (161 mg, 87%). m.p 88-90 °C ^1H NMR (300 MHz): δ = 7.84 (s, 1H), 7.34 (s, 1H), 6.76 (t, J = 5.9 Hz, 1H), 4.52 (d, J = 5.9 Hz, 2H), 2.89 (s, 6H), 0.91 (s, 9H), 0.10 (s, 6H); ^{13}C NMR (75 MHz): δ = 141.6, 137.1, 127.4, 123.2, 115.2, 60.7, 38.3, 26.0, 18.4, -5.1; FT-IR (neat, cm^{-1}): 2954, 2860, 1630, 1481, 1381, 1173, 1085, 827, 778, 754, 617; HR-MS (m/z): calc for $[\text{M}+\text{H}]^+$ $\text{C}_{14}\text{H}_{26}\text{ClN}_3\text{O}_3\text{SSi}$ 380.1231 found 380.1231.

(Z)-1-benzyl-4-(3-(tert-butyldimethylsilyloxy)-1-chloroprop-1-enyl)-1H-

imidazole (266):



In a round bottom flask compound **264** (100 mg, 0.40 mmol) was dissolved in CH₂Cl₂ (10 mL) under a nitrogen atmosphere. The solution was cooled to 0 °C and then to the reaction mixture imidazole (40 mg, 0.60 mmol) and TBSCl (78 mg, 0.52 mmol) was added. The reaction mixture was stirred for 8 h, then NH₄Cl (5 mL) was added and the organic layer was separated. The aqueous layer was extracted with EtOAc (3x50 mL), then the combined organic solutions were dried with anhydrous Na₂SO₄ and concentrated. The crude product was purified by chromatography (EtOAc 100%) providing **266** as a colorless solid (135 mg, 92%). m.p 76-78 °C ¹H NMR (300 MHz): δ = 7.54 (d, *J* = 1.4 Hz, 1H), 7.35-7.31 (m, 3H), 7.15 (dd, *J* = 7.4, 1.7 Hz, 2H), 7.03 (*J* = 1.0 Hz, 1H), 6.64 (t, *J* = 5.8 Hz, 1H), 5.06 (s, 2H), 4.49 (d, *J* = 5.8 Hz, 2H), 0.89 (s, 9H), 0.08 (s, 6H); ¹³C NMR (75 MHz): δ = 140.3, 137.8, 135.6, 129.2, 128.6, 127.5, 124.6, 124.4, 118.0, 60.8, 51.2, 26.2, 18.4, -5.1; FT-IR (neat, cm⁻¹): 2934, 2858, 1590, 1475, 1384, 1167, 1082, 822, 740, 596; HR-MS (*m/z*): calc for [M+H]⁺ C₁₉H₂₇ClN₂OSi 363.1659 found.

APPENDIX 1
¹H AND ¹³C NMR SPECTRUM OF
(*R*)-methyl-2-chloro-3-(1-(*N,N*-dimethylsulfamoyl)-1*H*-imidazol-4-yl)propanoate (47)



```

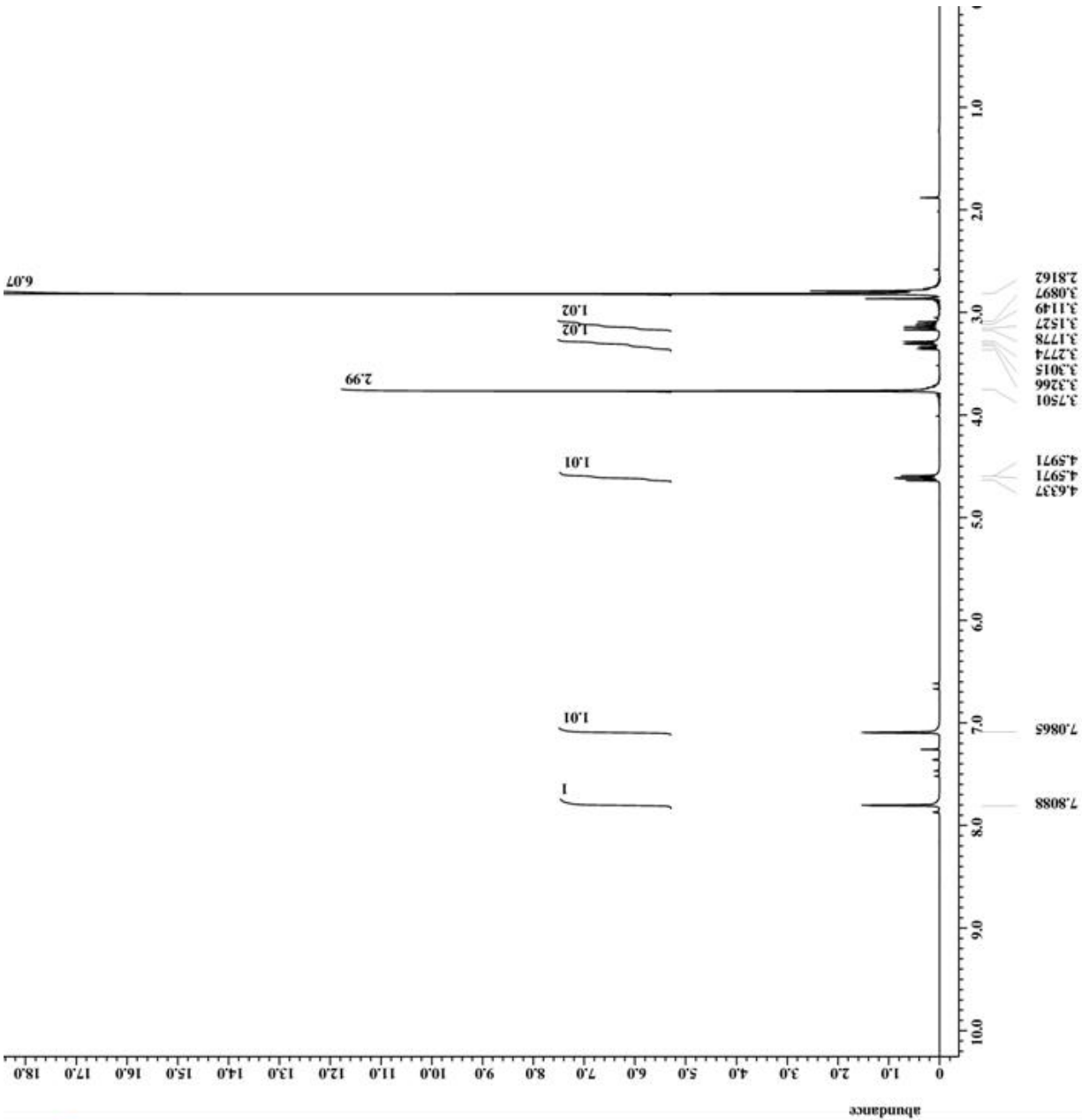
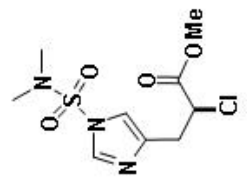
ilname      = sm_v_137_i1-4_jdf
author
xparment   = delta
sample_id  = S8659746
solvent     = CHLOROFORM-D
reaction_time = 11-JUN-2009 17:51:03
evision_time = 19-APR-2010 23:49:14
current_time = 20-APR-2010 19:36:39

comment
ate_format = single_pulse
im_size     = 1D_COMPLEX
im_title    = 13107
im_units    = [ppm]
ite         = X
ite         = ECX 300
ite         = DELTA2_NMR

ield_strength = 7.0586013[T] (300[MHz]
-acq_duration = 2.90717696[s]
-domain       = 1H
-freq        = 300.52965592[MHz]
-offset      = 5[ppm]
-points      = 16384
-prescans    = 0
-resolution  = 0.34397631[Hz]
-sweep       = 5.63570784[kHz]
rr_domain   = 1H
rr_freq     = 300.52965592[MHz]
rr_offset   = 5[ppm]
ri_domain   = 1H
ri_freq     = 300.52965592[MHz]
ri_offset   = 5[ppm]
lipped      = FALSE
bd_return   = 1
cans        = 24
stal_scans  = 24

-90_width   = 13.01[us]
-acq_time   = 2.90717696[s]
-angle      = 45[deg]
-atn        = 4[db]
-pulse      = 6.505[us]
rr_mode     = Off
ri_mode     = Off
ante_presat = FALSE
ntial_wait  = 1[e]
svr_gain    = 38
elaxation_delay = 5[s]
epitation_time = 7.90717696[s]
emp_get     = 22.9[dc]

```



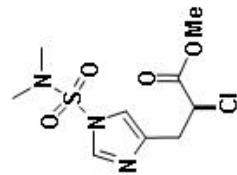
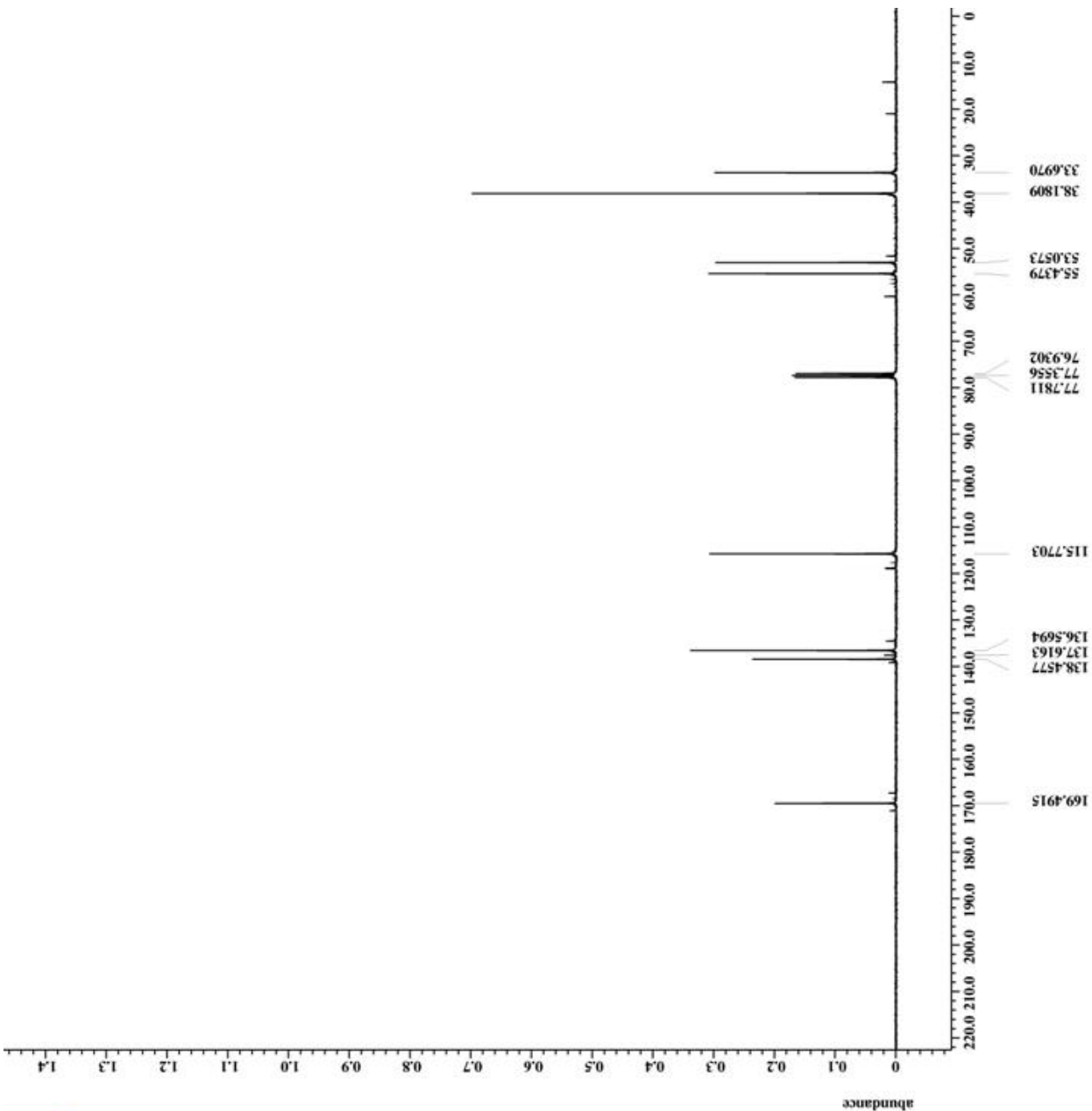


```
filename = sm_1_213_pure-4.jdf
author = delta
experiment = single_pulse_dec
sample_id = S8778317
solvent = CHLOROFORM-D
reaction_time = 10-JUN-2006 03:05:44
revision_time = 10-JUN-2009 19:09:11
current_time = 27-MAR-2010 16:21:11

comment = single pulse decouple
date_format = ID_COMPLEX
im_size = 52428
im_title = 13C
im_units = [ppm]
imensions = X
ite = ECX 300
spectrometer = DELTA2_NMR

field_strength = 7.0586013[T] (300[MHz]
-acq_duration = 2.76824064[s]
-domain = 13C
-freq = 75.56823426[MHz]
-offset = 100[ppm]
-points = 65336
-prescans = 4
-resolution = 0.36124027[Hz]
-sweep = 23.67424242[KHz]
rr_domain = 1H
rr_freq = 300.52965592[MHz]
rr_offset = 5[ppm]
lipped = FALSE
bd_return = 10
cans = 3000
stal_scans = 3000

-90_width = 9.75[us]
-acq_time = 2.76824064[s]
-angle = 30[deg]
-atn = 8[db]
-pulse = 3.25[us]
rr_atn_dec = 25[db]
rr_atn_noe = 25[db]
rr_noise = WALTZ
scoupling = TRUE
nit1al_wait = 1[s]
ce = TRUE
se_time = 3[s]
svr_gain = 50
relaxation_delay = 3[s]
acquisition_time = 5.76824064[s]
emp_get = 22.2[dC]
```

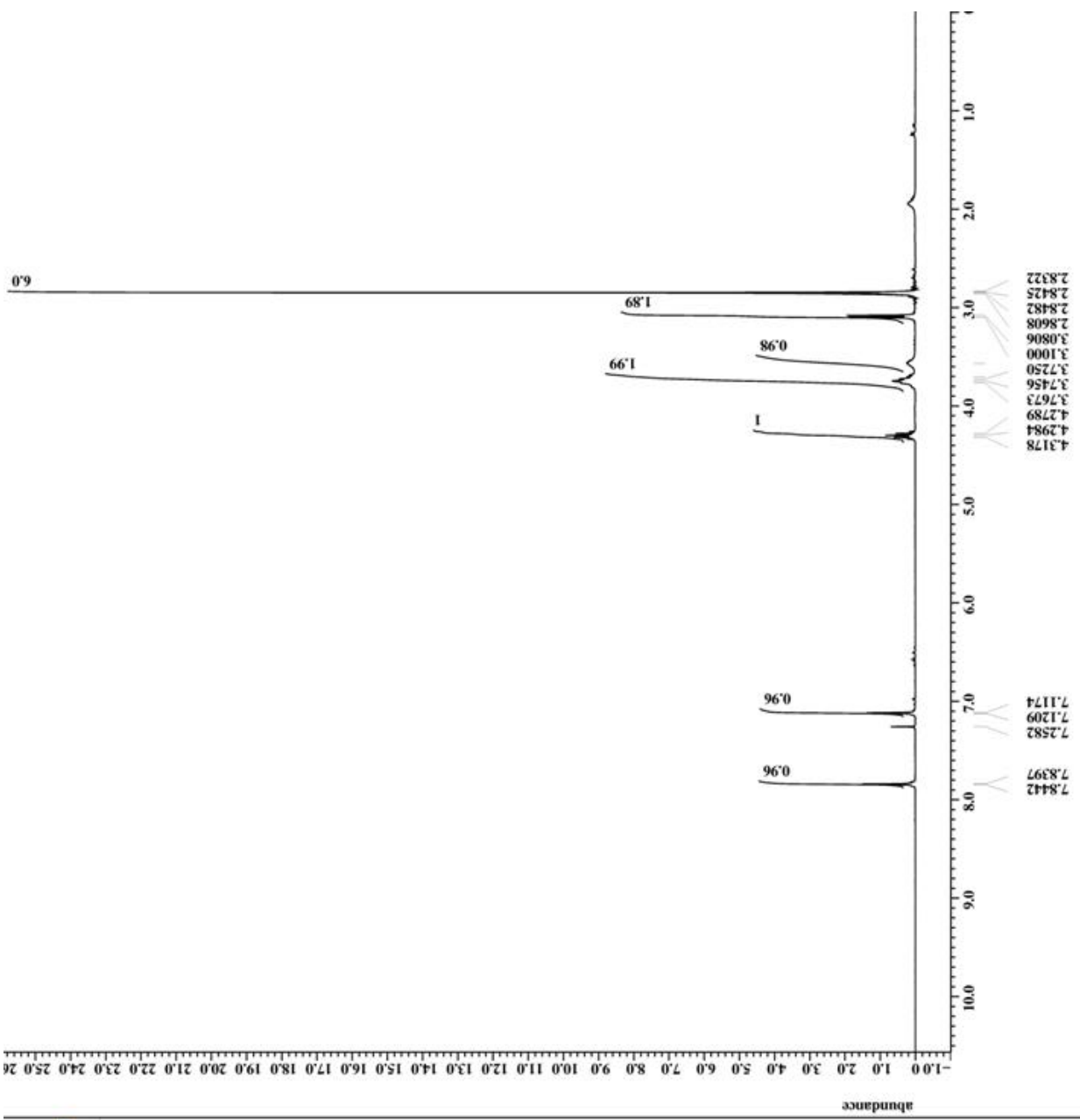
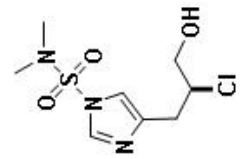


APPENDIX 2
¹H AND ¹³C NMR SPECTRUM OF
(*R*)-4-(2-chloro-3-hydroxypropyl)-*N,N*-dimethyl-1*H*-imidazole-1-sulfonamide
(48)



```

filename = sm_11_23_alcohol_pure
author = delta
experiment = single_pulse.ex2
sample_id = S#386706
solvent = CHLOROFORM-D
creation_time = 27-NOV-2006 11:33:17
revision_time = 27-MAR-2010 16:29:19
current_time = 27-MAR-2010 16:33:32
comment =
  = single_pulse
  = ID COMPLEX
in_size = 13107
in_title = 1H
in_units = [ppm]
inmensions = X
ite = ECX 300
spectrometer = DELTA2_NMR
ield_strength = 7.0586013[T] (300[MHz]
_acq_duration = 2.90717696[s]
_domain = 1H
_freq = 300.52965592[MHz]
_offset = 5[ppm]
_points = 16384
_prescans = 0
_resolution = 0.34397621[Hz]
_sweep = 5.63570784[kHz]
tr_domain = 1H
tr_freq = 300.52965592[MHz]
tr_offset = 5[ppm]
ri_domain = 1H
ri_freq = 300.52965592[MHz]
ri_offset = 5[ppm]
lipped = FALSE
od_return = 1
otal_scans = 24
_90_width = 13.01[us]
_acq_time = 2.90717696[s]
_angle = 45[deg]
_atn = 4[dB]
_pulse = 6.505[us]
tr_mode = Off
ri_mode = Off
ante_presat = FALSE
nitial_wait = 1[s]
scrvr_gain = 42
relaxation_delay = 5[s]
epitition_time = 7.90717696[s]
emp_get = 23.5[dC]
  
```



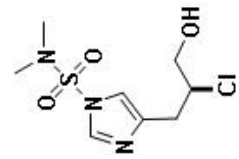
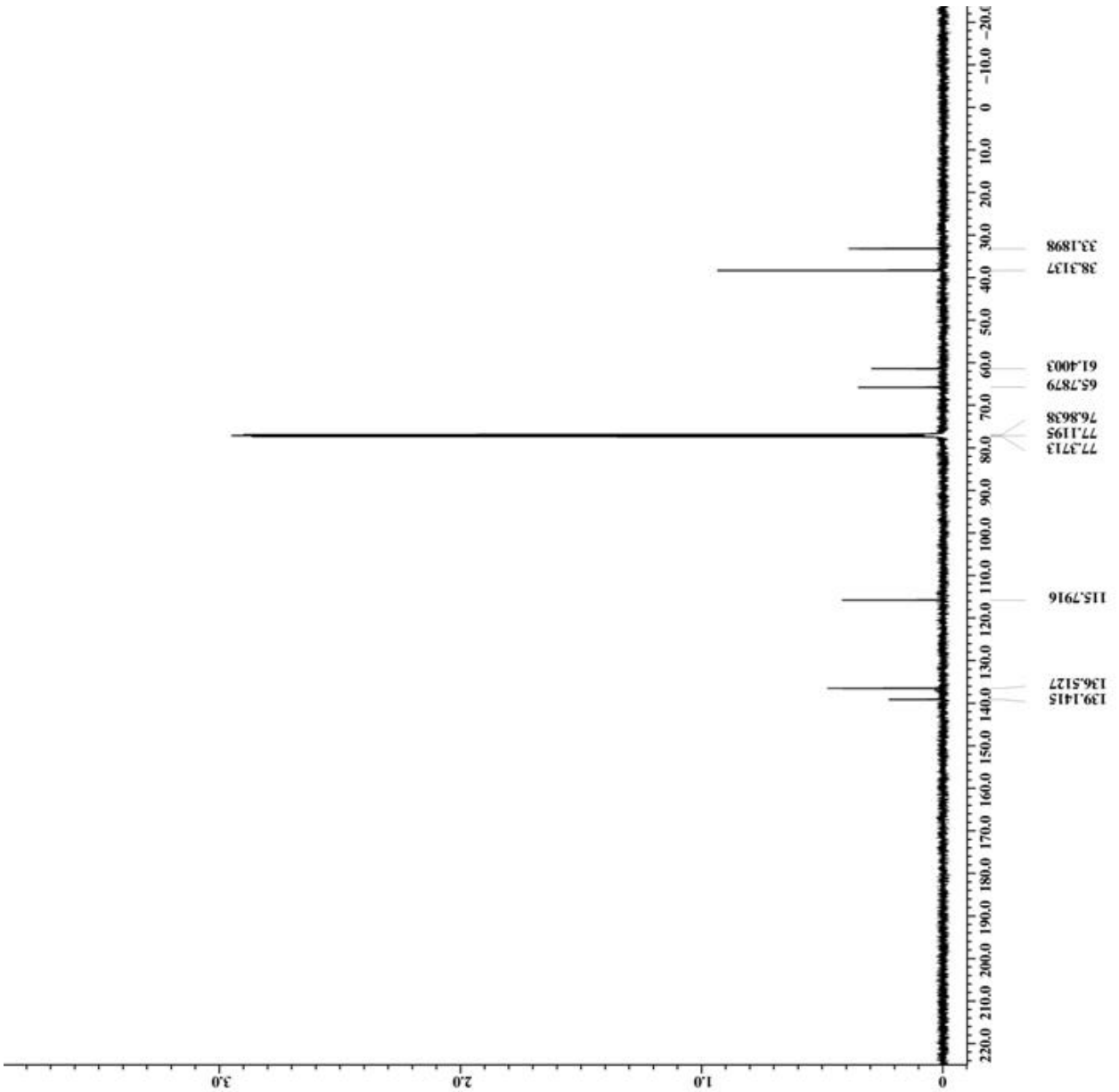


```

filename = sm_11_alcohol_pure-2.
author = delta
experiment = single_pulse_dec
sample_id = S#708634
solvent = CHLOROFORM-D
reaction_time = 10-DEC-2006 01:54:30
revision_time = 10-DEC-2006 15:31:21
current_time = 27-MAR-2010 16:38:46
comment = single pulse decouple
data_format = ID COMPLEX
im_size = 65536
im_title = 13C
im_units = [ppm]
imensions = X
ite = Eclipse+ 500
spectrometer = DELTA_NMR
field_strength = 11.7473579[T] (500[MH
acq_duration = 2.0840448[s]
domain = 13C
freq = 125.76529768[MHz]
offset = 100[ppm]
points = 65536
prescans = 4
resolution = 0.47983613[Hz]
sweep = 31.44654088[kHz]
tr_domain = 1H
tr_freq = 500.15991521[MHz]
tr_offset = 5[ppm]
lipped = TRUE
bd_return = 10
cans = 3000
otal_scans = 3000
_90_width = 14.2[us]
_acq_time = 2.0840448[s]
_angle = 30[deg]
_pulse = 4.73333333[us]
_nitai_wait = 1[s]
_pe_time = 1[s]
_base_preset = 3[us]
_scvr_gain = 29
_relaxation_delay = 2[s]
_omp_get = 26.8[dc]
_nblank_time = 2[us]

```

(Millions)



APPENDIX 3
¹H AND ¹³C NMR SPECTRUM OF
(*R*)-4-(2-chloro-3-(1,3-dioxo-1H-pyrrolo[1,2-c]imidazol-2(3H)-yl)propyl)-N,N-
dimethyl-1H-imidazole-1-sulfonamide (50)



```

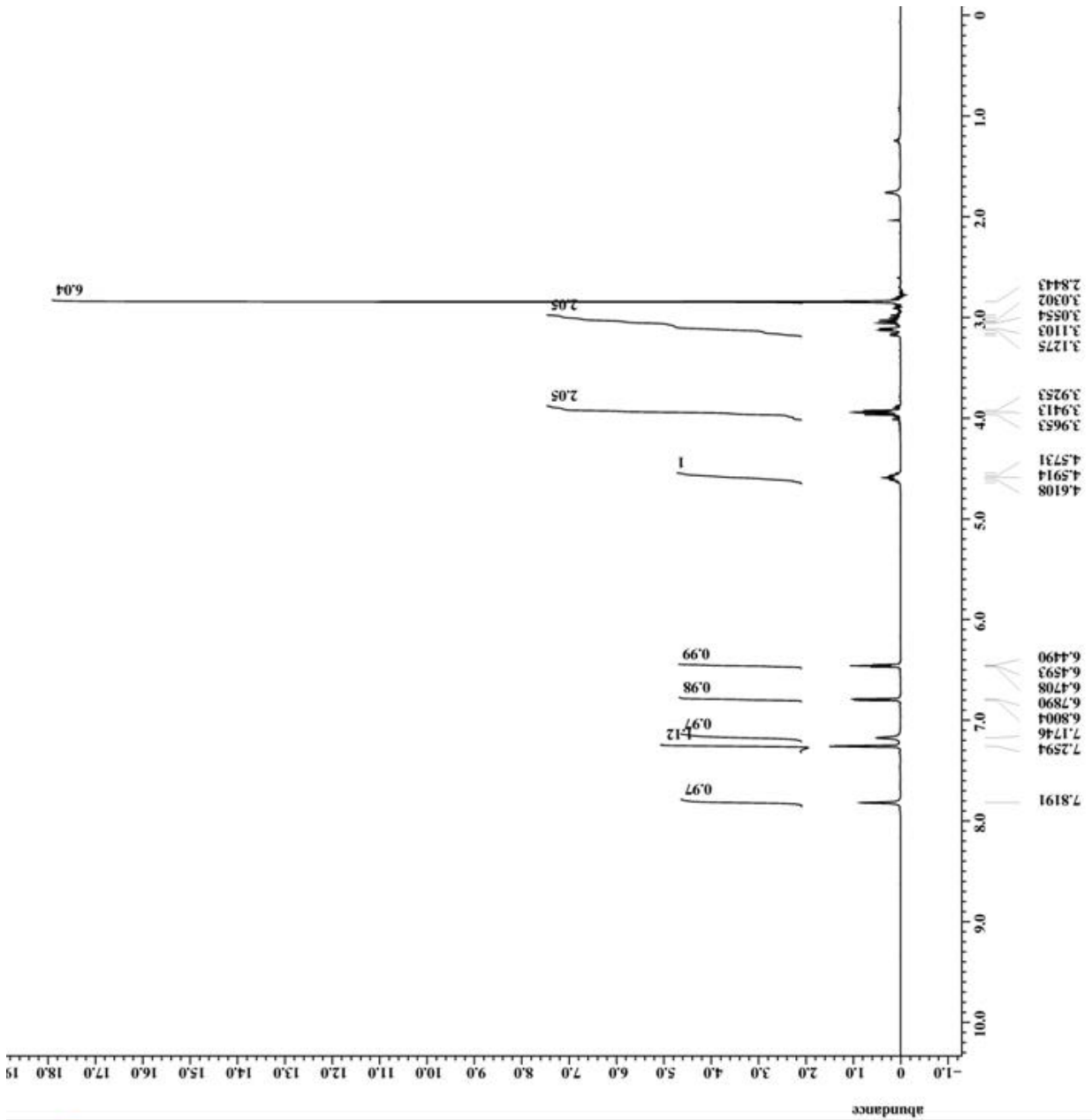
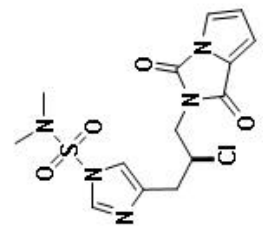
filename = sm_1183_pure ii-4_jd
author = delta
experiment = single_pulse.ex2
sample_id = S9595971
solvent = CHLOROFORM-D
reaction_time = 28-FEB-2006 17:30:58
acquisition_time = 27-MAR-2010 16:46:52
current_time = 27-MAR-2010 16:47:12

comment = single_pulse
ata_format = ID COMPLEX
in_size = 13107
in_title = 1H
in_units = [ppm]
inmensions = X
ite = ECX 300
spectrometer = DELTA2_NMR

ield_strength = 7.0586013[T] (300[MHz]
_acq_duration = 2.90717696[s]
_domain = 1H
_freq = 300.52965592[MHz]
_offset = 5[ppm]
_points = 16384
_prescans = 0
_resolution = 0.34397621[Hz]
_sweep = 5.63570784[kHz]
tr_domain = 1H
tr_freq = 300.52965592[MHz]
tr_offset = 5[ppm]
ri_domain = 1H
ri_freq = 300.52965592[MHz]
ri_offset = 5[ppm]
lipped = FALSE
od_return = 1
otal_scans = 14

_90_width = 13.01[us]
_acq_time = 2.90717696[s]
_angle = 45[deg]
_atn = 4[db]
_pulse = 6.505[us]
tr_mode = Off
ri_mode = Off
ante_presat = FALSE
nitial_wait = 1[s]
scrvr_gain = 44
relaxation_delay = 5[s]
epetition_time = 7.90717696[s]
emp_get = 21.6[dc]

```



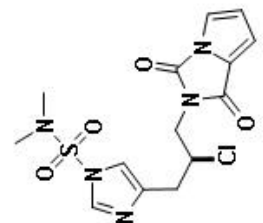
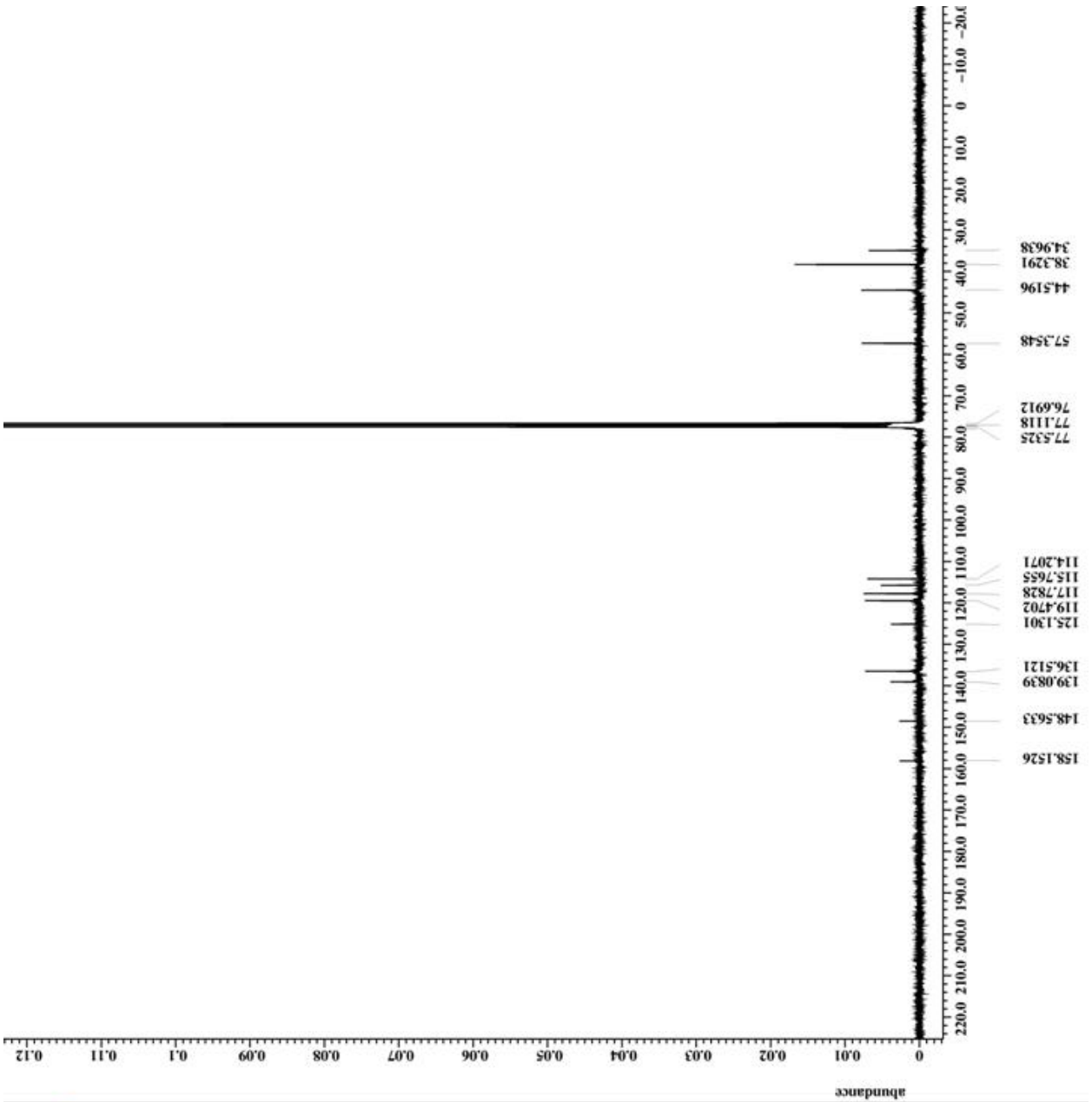


```
filename = sm_V_31_PURE-3_3.jdf
author = delta
experiment = single_pulse_dec
sample_id = S8502438
solvent = CHLOROFORM-D
reaction_time = 23-DEC-2008 20:16:49
revision_time = 23-DEC-2008 20:03:52
current_time = 27-MAR-2010 16:52:08

comment = single pulse decouple
ata_format = ID REAL
in_size = 52428
in_title = 13C
in_units = [ppm]
inmensions = X
ite = ECX 300
spectrometer = DELTA2_NMR

ield_strength = 7.0586013[T] (300[MHz]
acq_duration = 2.76824064[s]
domain = 13C
freq = 75.56823426[MHz]
offset = 100[ppm]
points = 65536
prescans = 4
resolution = 0.36124027[Hz]
sweep = 23.67424242[kHz]
tr_domain = 1H
tr_freq = 300.52965592[MHz]
tr_offset = 5[ppm]
lipped = FALSE
bd_return = 10
cans = 3775
otal_scans = 3775

_90_width = 9.75[us]
acq_time = 2.76824064[s]
angle = 30[deg]
atn = 8[dB]
pulse = 3.25[us]
tr_atn_dec = 25[dB]
tr_atn_noe = 25[dB]
tr_noise = WALTZ
scoupling = TRUE
nitial_wait = 1[s]
oe_time = TRUE
oe_time = 3[s]
ecvr_gain = 50
elaxation_delay = 3[s]
epetition_time = 5.76824064[s]
emp_get = 23.2[dc]
```



APPENDIX 4
¹H AND ¹³C NMR SPECTRUM OF
(*R*)-N-(2-chloro-3-(1-(*N,N*-dimethylsulfamoyl)-1H-imidazol-4-yl)propyl)-1H-
pyrrole-2-carboxamide (54)



```

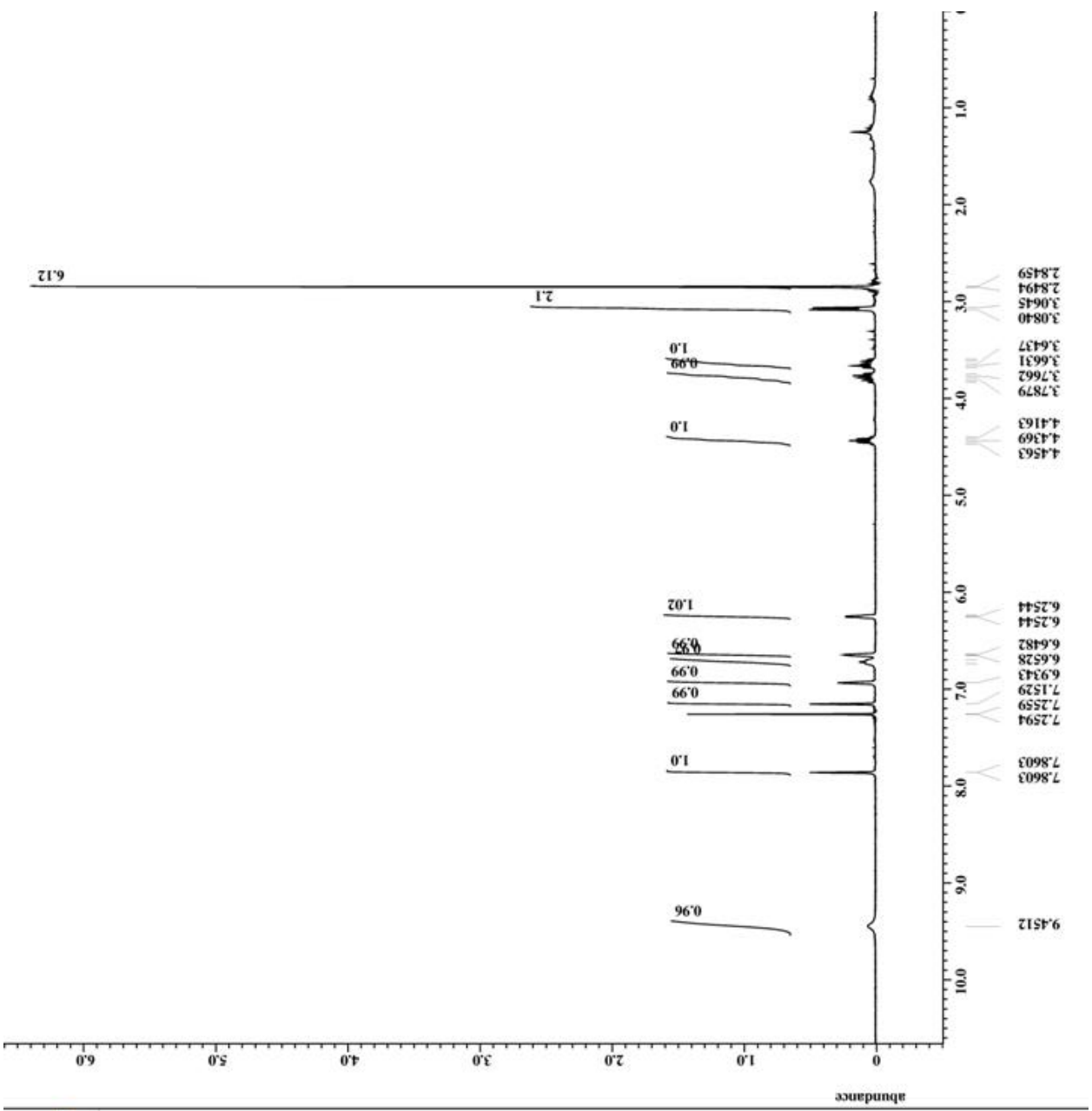
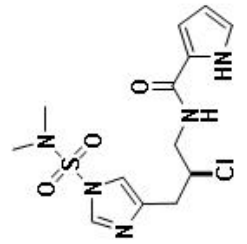
filename = sm_1137_pure-5.jdf
author = delta
experiment = single_pulse.ex2
sample_id = S#463736
solvent = CHLOROFORM-D
reaction_time = 17-NOV-2005 13:33:13
acquisition_time = 27-MAR-2010 17:01:41
current_time = 27-MAR-2010 17:03:16

comment = single_pulse
ata_format = ID COMPLEX
in_size = 13107
in_title = 1H
in_units = [ppm]
in_dimensions = X
ite = ECX 300
spectrometer = DELTA2_NMR

ield_strength = 7.0586013[T] (300 [MHz]
_acq_duration = 2.90717696[s]
_domain = 1H
_freq = 300.52965592 [MHz]
_offset = 5[ppm]
_points = 16384
_prescans = 0
_resolution = 0.34397621 [Hz]
_sweep = 5.63570784 [kHz]
tr_domain = 1H
tr_freq = 300.52965592 [MHz]
tr_offset = 5[ppm]
ri_domain = 1H
ri_freq = 300.52965592 [MHz]
ri_offset = 5[ppm]
lipped = FALSE
od_return = 1
otal_scans = 24

_90_width = 13.01[us]
_acq_time = 2.90717696[s]
_angle = 45[deg]
_atn = 4[db]
_pulse = 6.505[us]
tr_mode = Off
ri_mode = Off
ante_presat = FALSE
nitai_wait = 1[s]
scrvr_gain = 46
relaxation_delay = 5[s]
petition_time = 7.90717696[s]
emp_get = 23[dc]

```





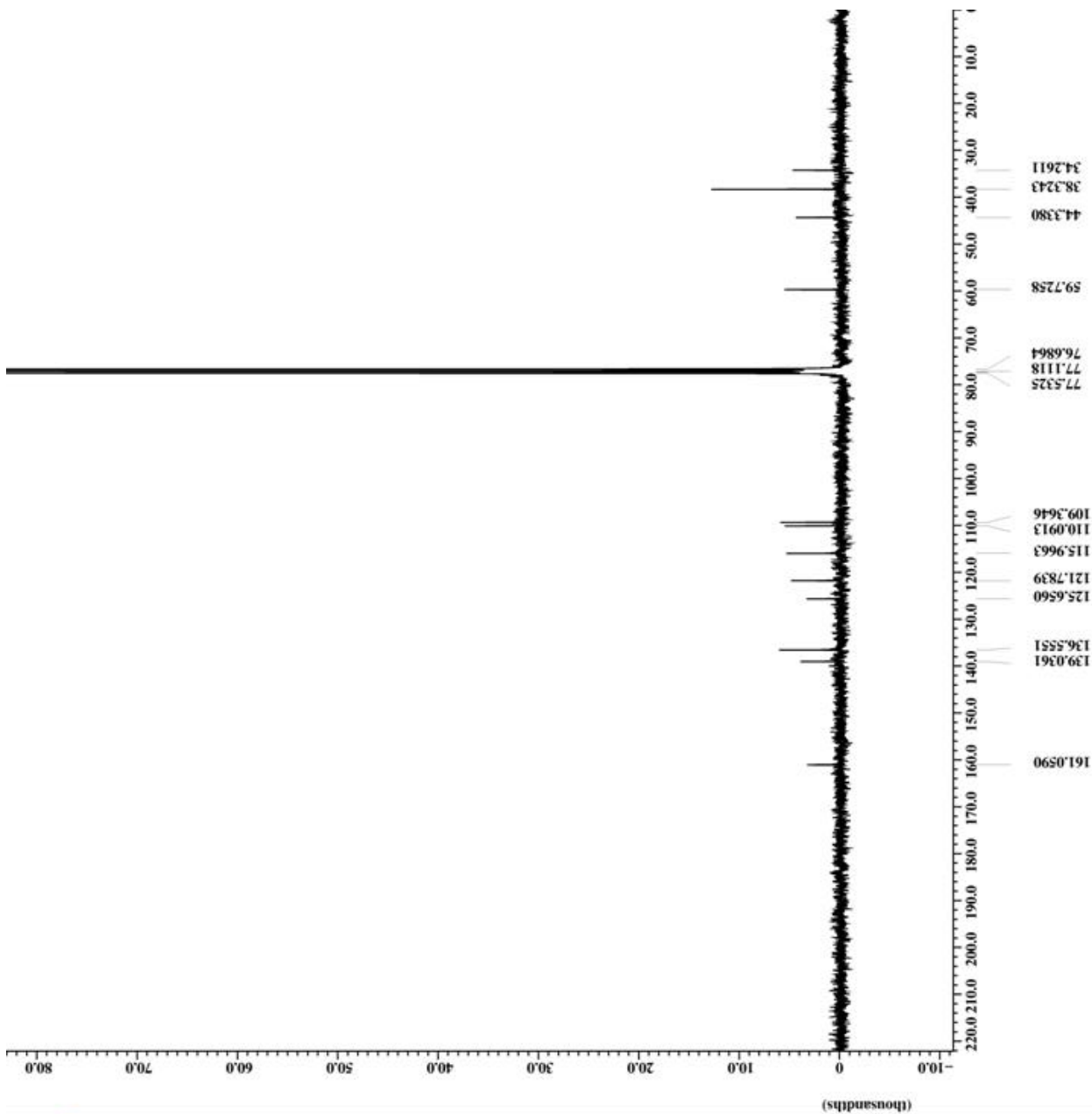
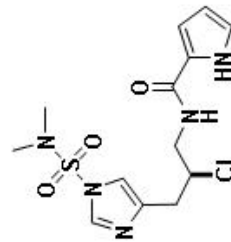
```

ilname      = sm_1_137_pure-2.jdf
author      = delta
experiment  = single_pulse_dec
sample_id   = S8617488
solvent     = CHLOROFORM-D
revision_time = 17-NOV-2005 20:42:29
revision_time = 17-NOV-2005 21:56:49
current_time = 27-MAR-2010 17:08:07

comment     = single pulse decouple
ate_format  = ID COMPLEX
im_size     = 52428
im_title    = 13C
im_units    = [ppm]
imensions  = X
ite         = ECX 300
ite         = DELTA2_NMR

ield_strength = 7.0586013[T] (300[MHz]
-acq_duration = 2.76824064[s]
-domain       = 13C
-freq         = 75.56823426[MHz]
-offset       = 100[ppm]
-points       = 65536
-prescans     = 4
-resolution   = 0.36124027[Hz]
-sweep        = 23.67424242[KHz]
-rr_domain   = 1H
-rr_freq      = 300.52965592[MHz]
-rr_offset    = 5[ppm]
-lipped       = FALSE
-bd_return    = 10
-cans         = 1627
-stal_scans   = 1627

-90_width    = 9.75[us]
-acq_time     = 2.76824064[s]
-angle        = 30[deg]
-atn          = 8[db]
-pulse        = 3.25[us]
-rr_atn_dec   = 25[db]
-rr_atn_noe   = 25[db]
-rr_noise     = WALTZ
-ecoupling    = TRUE
-nit1al_wait  = 1[s]
-be           = TRUE
-be_time      = 3[s]
-scvr_gain    = 50
-relaxation_delay = 3[s]
-epitation_time = 5.76824064[s]
-emp_get      = 23.3[dc]
  
```



APPENDIX 5
¹H AND ¹³C NMR SPECTRUM OF
(S)-4-((2-(1H-pyrrol-2-yl)-4,5-dihydrooxazol-5-yl)methyl)-N,N-dimethyl-1H-
imidazole-1-sulfonamide (56)



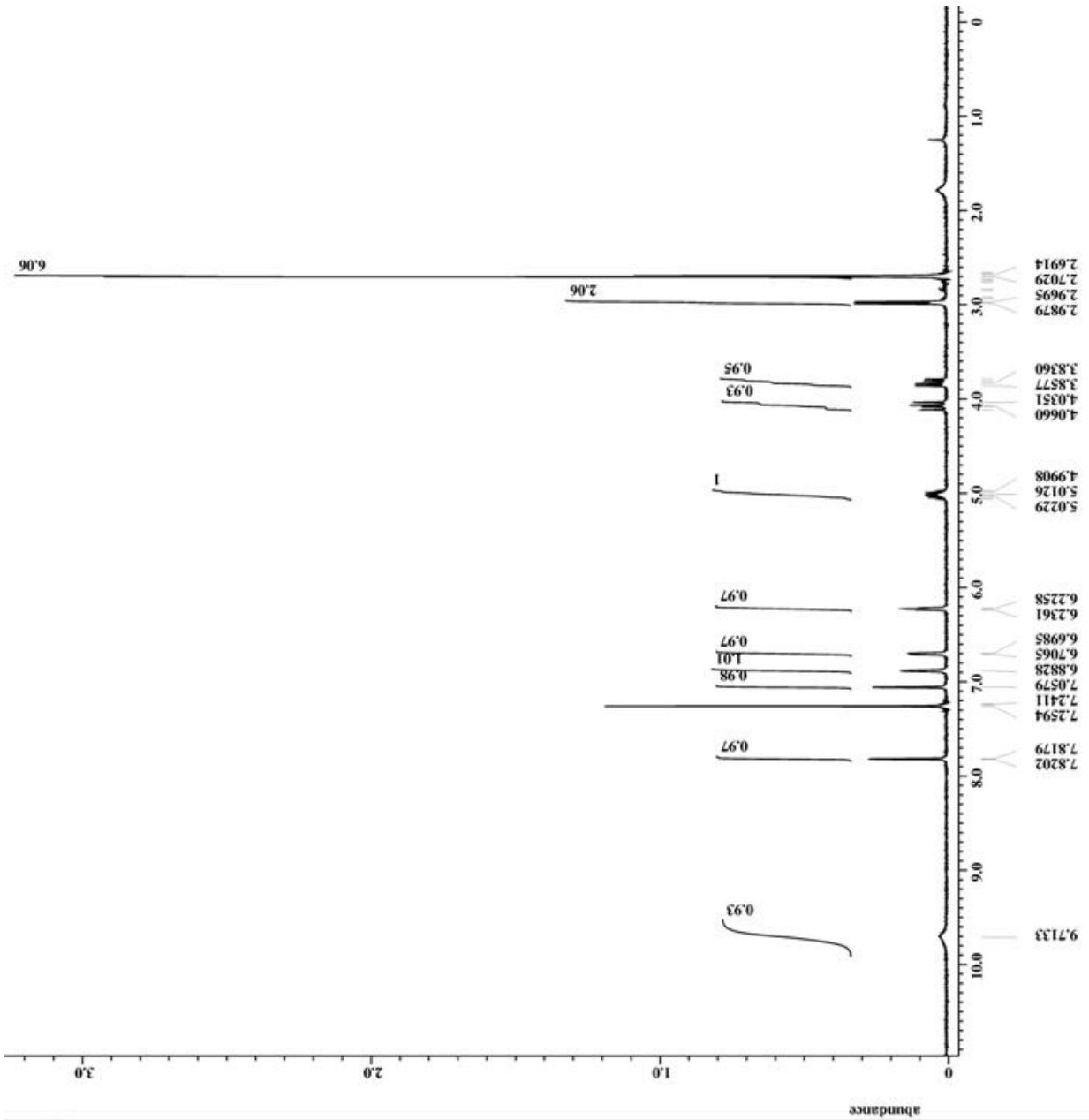
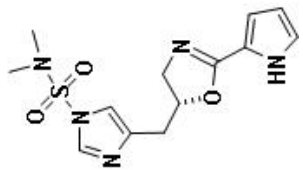
```

ilname = sm_precursor to azida
uthor = delta
xperiment = single_pulse.ex2
mple_id = S#402502
olvent = CHLOROFORM-D
reation_time = 31-AUG-2006 11:53:19
evision_time = 27-MAR-2010 17:13:30
urrent_time = 27-MAR-2010 17:13:59

omment = single_pulse
ata_format = ID COMPLEX
m_size = 13107
m_title = 1H
m_units = [ppm]
mensions = X
ite = ECX 300
ectrometer = DELTA2_NMR

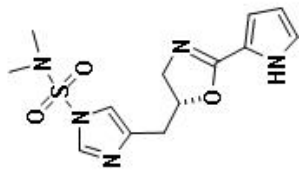
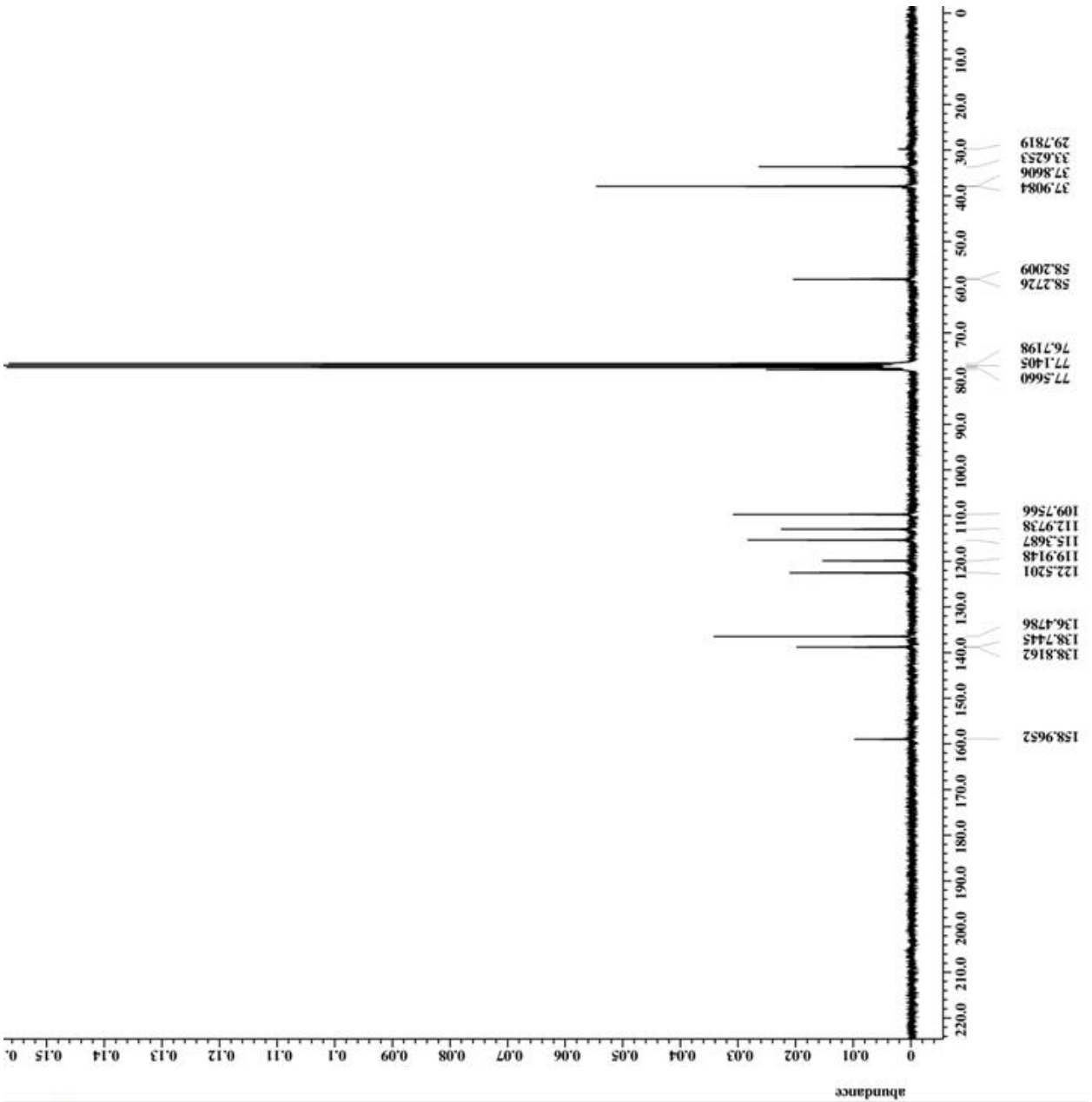
ield_strength = 7.0586013[T] (300[MHz]
cq_duration = 2.90717696[s]
omain = 1H
req = 300.52965592[MHz]
ffset = 5[ppm]
oints = 16384
rescans = 0
esolution = 0.34397621[Hz]
weep = 5.63570784[kHz]
r_domain = 1H
r_freq = 300.52965592[MHz]
r_offset = 5[ppm]
r_domain = 1H
r_offset = 300.52965592[MHz]
r_offset = 5[ppm]
lipped = FALSE
od_return = 1
cans = 19
atal_scans = 19

_90_width = 13.01[us]
cq_time = 2.90717696[s]
angle = 45[deg]
_atn = 4[db]
pulse = 6.505[us]
r_mode = Off
ri_mode = Off
ante_presat = FALSE
nitial_wait = 1[s]
ecvr_gain = 46
elaxation_delay = 5[s]
epetition_time = 7.90717696[s]
emp_get = 22.8[dc]
  
```





```
filename = sm_cyclic_int-2.jdf
author = delta
experiment = single_pulse_dec
sample_id = S#781820
solvent = CHLOROFORM-D
reaction_time = 16-JAN-2007 08:04:08
evision_time = 16-JAN-2007 10:11:11
current_time = 27-MAR-2010 17:17:35
comment = single pulse decouple
ata_format = ID COMPLEX
in_size = 52428
in_title = 13C
in_units = [ppm]
inmensions = X
ite = ECX 300
spectrometer = DELTA2_NMR
ield_strength = 7.0586013[T] (300[MHz]
_acq_duration = 2.76824064[s]
_domain = 13C
_freq = 75.56823426[MHz]
_offset = 100[ppm]
_points = 65536
_prescans = 4
_resolution = 0.36124027[Hz]
_sweep = 23.67424242[kHz]
tr_domain = 1H
tr_freq = 300.52965592[MHz]
tr_offset = 5[ppm]
lipped = FALSE
bd_return = 10
cans = 6000
atal_scans = 6000
_90_width = 9.75[us]
_acq_time = 2.76824064[s]
_angle = 30[deg]
_atn = 8[db]
_pulse = 3.25[us]
tr_atn_dec = 25[db]
tr_atn_noe = 25[db]
tr_noise = WALTZ
scoupling = TRUE
nitial_wait = 1[s]
oe_time = TRUE
oe_time = 3[s]
ecvr_gain = 50
elaxation_delay = 3[s]
epetition_time = 5.76824064[s]
emp_get = 23[dc]
```



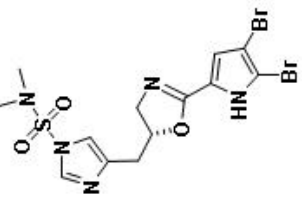
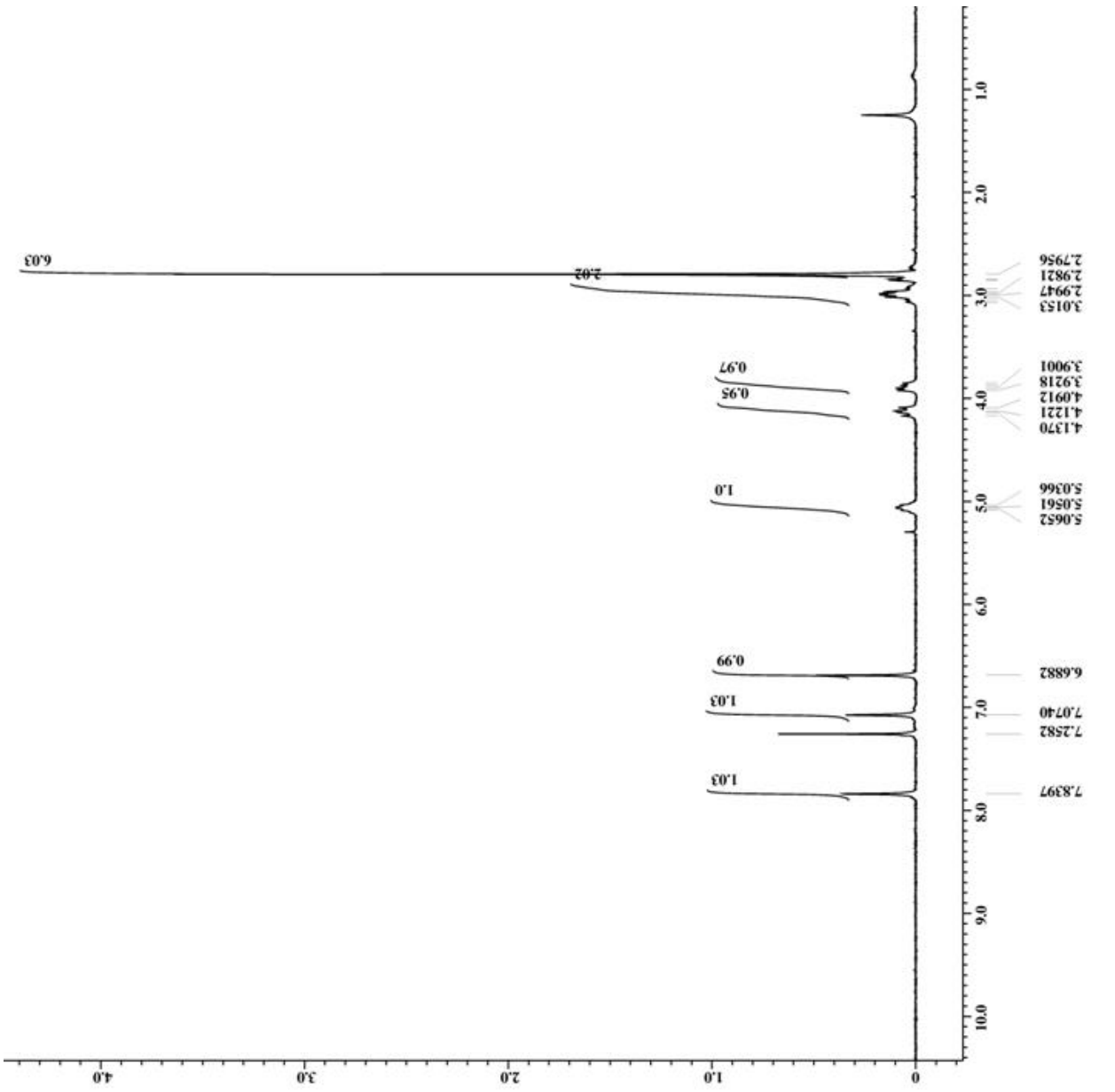
APPENDIX 6
¹H AND ¹³C NMR SPECTRUM OF
(S)-4-((2-(4,5-dibromo-1H-pyrrol-2-yl)-4,5-dihydrooxazol-5-yl)methyl)-N,N-
dimethyl-1H-imidazole-1-sulfonamide (58)



```

filename = sm_V_157_pure_iv-3.jd
author = delta
experiment = single_pulse.ex2
sample_id = S882671
solvent = CHLOROFORM-D
reaction_time = 9-JUL-2009 23:00:23
acquisition_time = 9-JUL-2009 23:12:45
current_time = 27-MAR-2010 17:21:21
comment =
  = single_pulse
  = ID REAL
  = 13107
  = 1H
  = [ppm]
  = X
  = ECX 300
  = DELTA2_NMR
  = 7.0586013[T] (300[MHz]
  = 2.90717696[s]
  = 1H
  = 300.52965592[MHz]
  = 5[ppm]
  = 16384
  = 0
  = 0.34397621[Hz]
  = 5.63570784[kHz]
  = 1H
  = 300.52965592[MHz]
  = 5[ppm]
  = 1H
  = 300.52965592[MHz]
  = 5[ppm]
  = FALSE
  = 1
  = 16
  = 16
  = 13.01[us]
  = 2.90717696[s]
  = 45[deg]
  = 4[db]
  = 6.505[us]
  = Off
  = Off
  = FALSE
  = 1[s]
  = 46
  = 5[s]
  = 7.90717696[s]
  = 23.1[dc]
  emp_get
  
```

abundance





```

filename = sm_V_157_pure_iii-2.j
author = delta
experiment = single_pulse_dec
sample_id = S8642639
solvent = CHLOROFORM-D
acquisition_time = 5-JUL-2009 04:53:39
revision_time = 5-JUL-2009 16:04:38
current_time = 27-MAR-2010 17:29:36

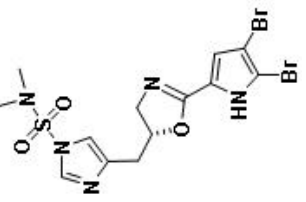
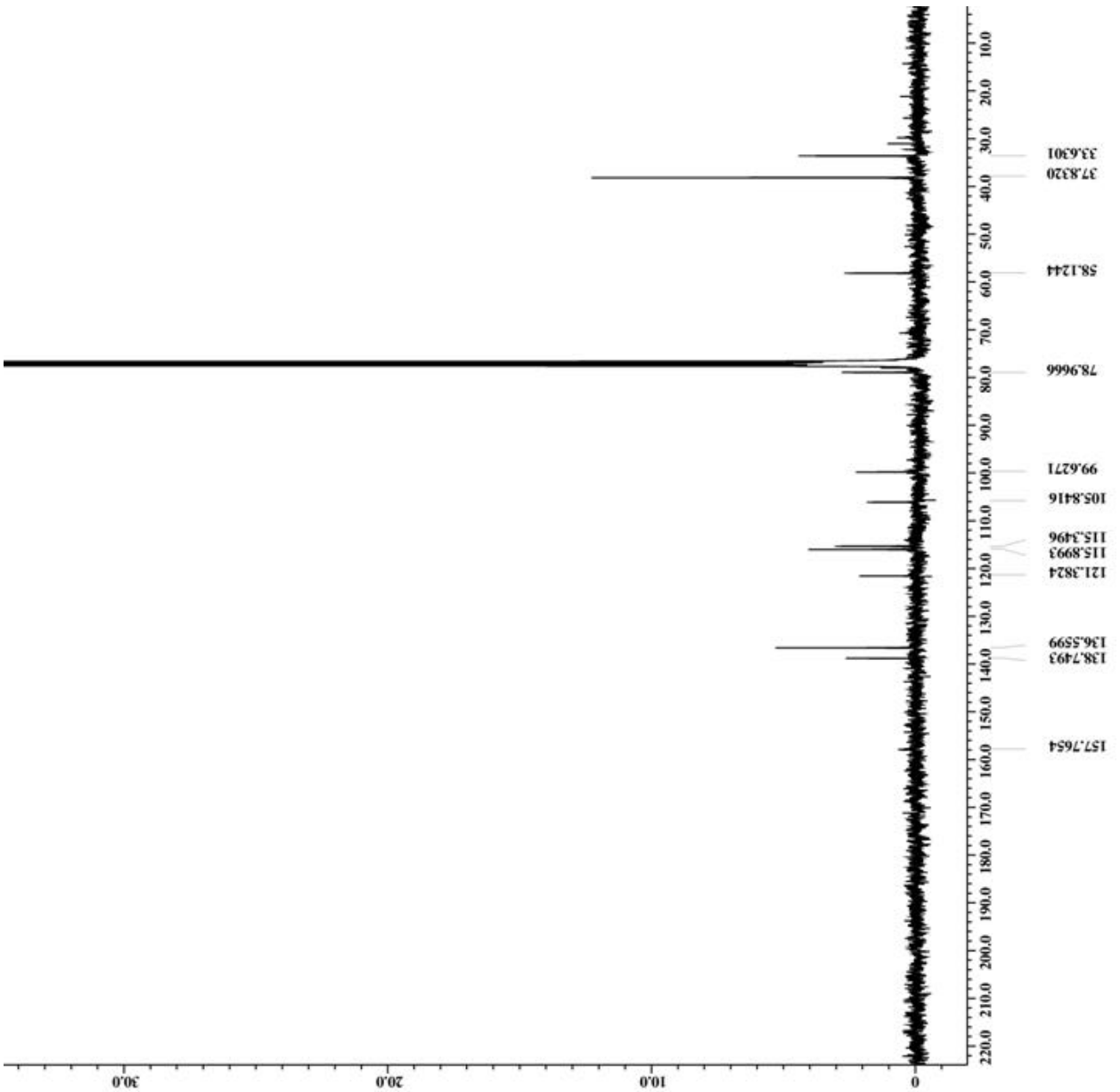
comment = single pulse decouple
ata_format = ID COMPLEX
in_size = 52428
in_title = 13C
in_units = [ppm]
inmensions = X
ite = ECX 300
spectrometer = DELTA2_NMR

ield_strength = 7.0586013[T] (300[MHz]
acq_duration = 2.76824064[s]
domain = 13C
freq = 75.56823426[MHz]
offset = 100[ppm]
points = 65536
prescans = 4
resolution = 0.36124027[Hz]
sweep = 23.67424242[kHz]
tr_domain = 1H
tr_freq = 300.52965592[MHz]
tr_offset = 5[ppm]
lipped = FALSE
bd_return = 10
cans = 7000
atal_scans = 7000

_90_width = 9.75[us]
acq_time = 2.76824064[s]
angle = 30[deg]
atn = 8[dB]
pulse = 3.25[us]
tr_atn_dec = 25[dB]
tr_atn_noe = 25[dB]
tr_noise = WALTZ
scoupling = TRUE
nitial_wait = 1[s]
oe_time = TRUE
oe_time = 3[s]
ecvr_gain = 50
relaxation_delay = 3[s]
petition_time = 5.76824064[s]
emp_get = 23.3[dc]

```

(thousands)



APPENDIX 7
¹H AND ¹³C NMR SPECTRUM OF
(S)-4-((2-(4-bromo-1H-pyrrol-2-yl)-4,5-dihydrooxazol-5-yl)methyl)-N,N-
dimethyl-1H-imidazole-1-sulfonamide (59)



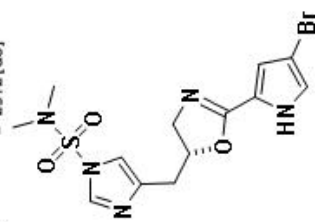
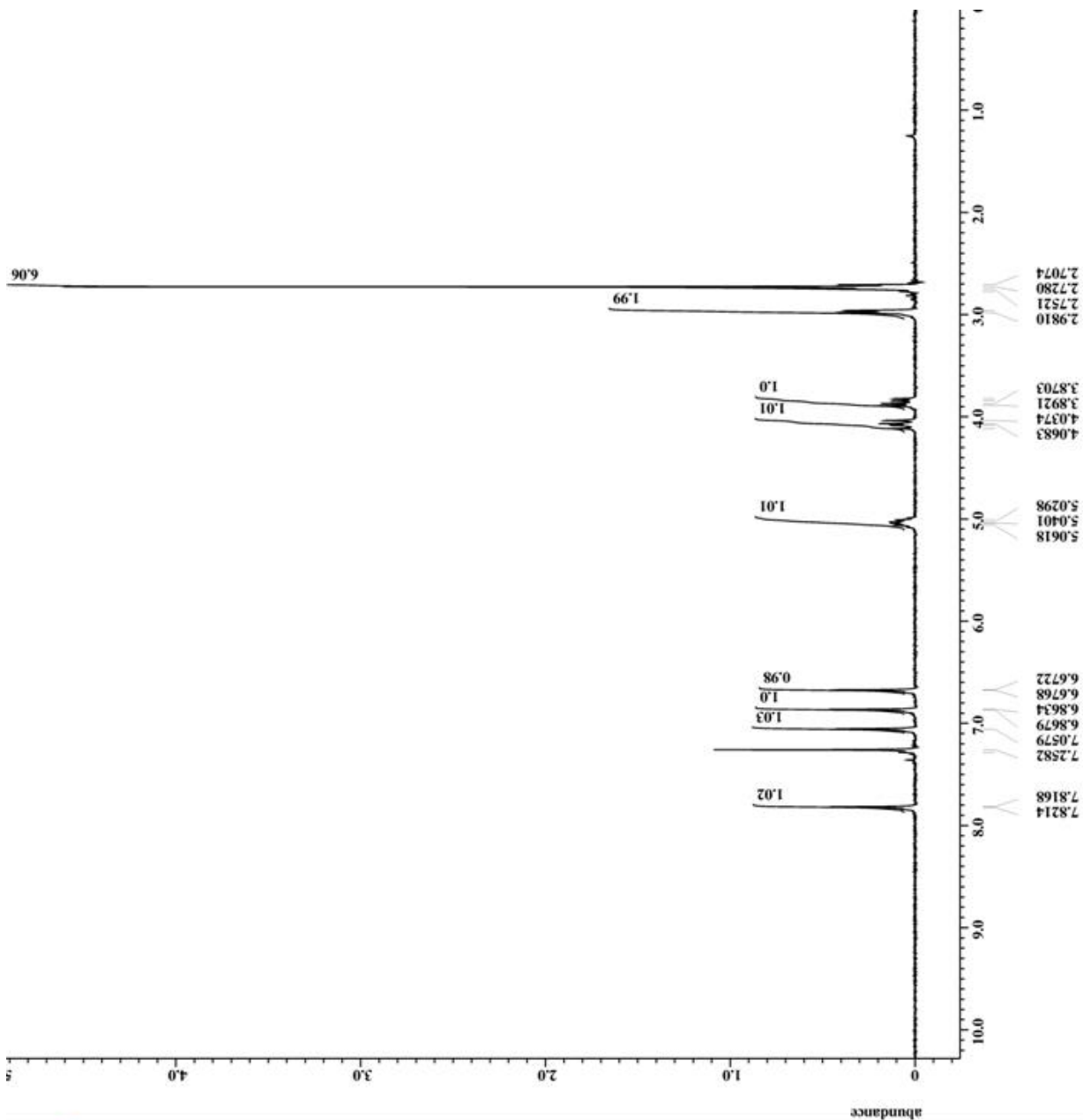
```

ilname      = sm_v_156_pure_ii-4_jd
author      = delta
experiment  = single_pulse.ex2
sample_id   = S8620333
solvent     = CHLOROFORM-D
reaction_time = 1-JUL-2009 17:05:04
revision_time = 27-MAR-2010 17:38:18
current_time = 27-MAR-2010 17:39:20

comment     = single_pulse
date_format = DD.MM.YY
im_size     = 13107
im_title    = 1H
im_units    = [ppm]
imensions  = X
ite         = ECX 300
p1          = DELTA_NMR

field_strength = 7.0586013[T] (300[MHz])
-acq_duration  = 2.90717696[s]
-domain        = 1H
-freq          = 300.52965592[MHz]
-gamma         = 5[ppm]
-points        = 16384
-prescans      = 0
-resolution    = 0.34397631[Hz]
-sweep         = 5.63570784[kHz]
-rr_domain    = 1H
-rr_freq      = 300.52965592[MHz]
-rr_offset    = 5[ppm]
-ri_domain    = 1H
-ri_freq      = 300.52965592[MHz]
-ri_offset    = 5[ppm]
-lipped       = FALSE
-bd_return    = 1
-cans         = 12
-stal_scans   = 12

-90_width     = 13.01[us]
-acq_time     = 2.90717696[s]
-angle        = 45[deg]
-atn          = 4[dB]
-pulse        = 6.505[us]
-rr_mode      = Off
-ri_mode      = Off
-ante_presat  = FALSE
-ntital_wait  = 1[s]
-ecvr_gain    = 1[e]
-relaxation_delay = 5[s]
-acquisition_time = 7.90717696[s]
-emp_get      = 23.2[dC]
  
```





```

filename = sm_V_156_pure_ii-2.jd
author = delta
experiment = single_pulse_dec
sample_id = S#621828
solvent = CHLOROFORM-D
acquisition_time = 1-JUL-2009 19:29:53
revision_time = 1-JUL-2009 22:02:41
current_time = 27-MAR-2010 17:43:06

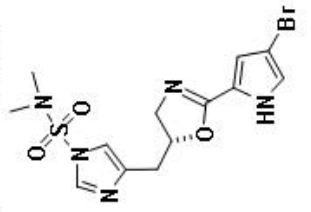
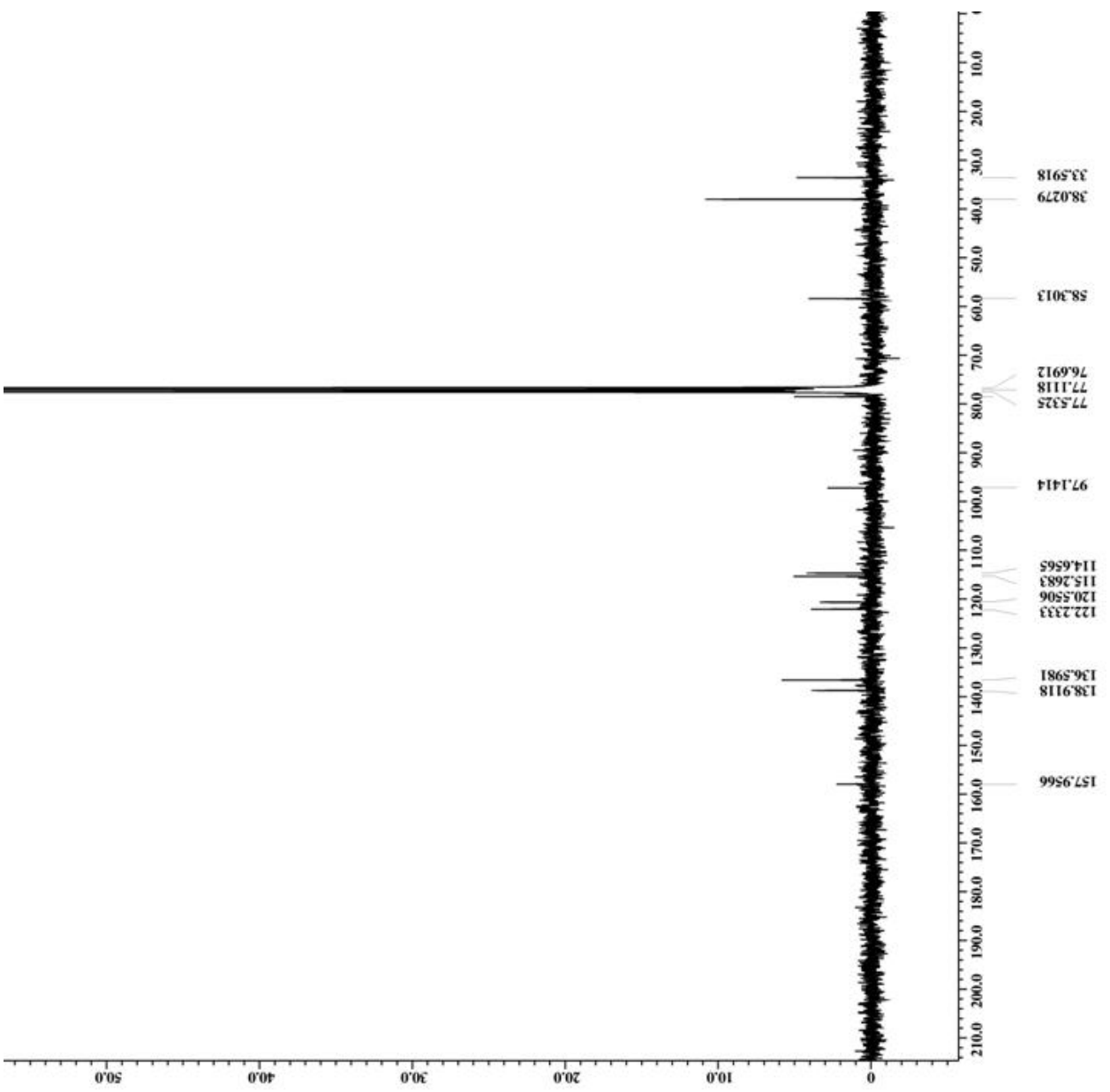
comment = single pulse decouple
ata_format = ID COMPLEX
in_size = 52428
in_title = 13C
in_units = [ppm]
inmensions = X
ite = ECX 300
spectrometer = DELTA2_NMR

ield_strength = 7.0586013[T] (300[MHz]
_acq_duration = 2.76824064[s]
_domain = 13C
_freq = 75.56823426[MHz]
_offset = 100[ppm]
_points = 65536
_prescans = 4
_resolution = 0.36124027[Hz]
_sweep = 23.67424242[kHz]
tr_domain = 1H
tr_freq = 300.52965592[MHz]
tr_offset = 5[ppm]
lipped = FALSE
bd_return = 10
cans = 1500
otal_scans = 1500

_90_width = 9.75[us]
_acq_time = 2.76824064[s]
_angle = 30[deg]
_atn = 8[db]
_pulse = 3.25[us]
tr_atn_dec = 25[db]
tr_atn_noe = 25[db]
tr_noise = WALTZ
scoupling = TRUE
nitial_wait = 1[s]
oe_time = TRUE
oe_time = 3[s]
ecvr_gain = 50
elaxation_delay = 3[s]
epetition_time = 5.76824064[s]
emp_get = 23.4[dc]

```

(thousands)



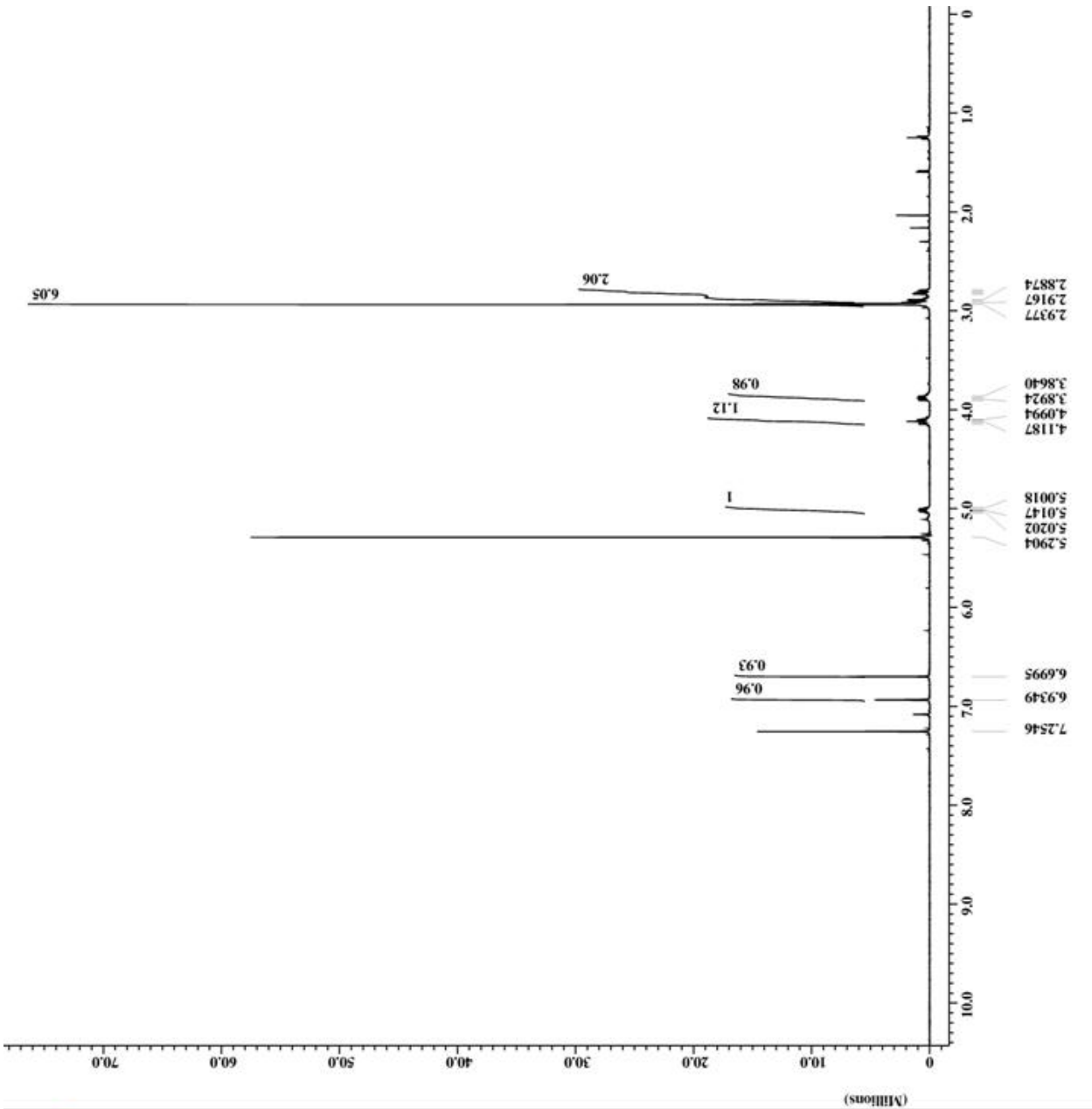
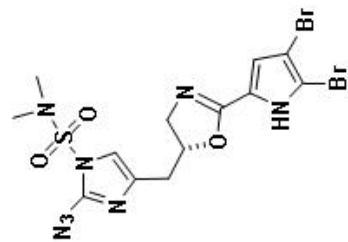
APPENDIX 8
¹H AND ¹³C NMR SPECTRUM OF
(S)-2-azido-4-((2-(4,5-dibromo-1H-pyrrol-2-yl)-4,5-dihydrooxazol-5-yl)methyl)-N,N-dimethyl-1H-imidazole-1-sulfonamide (60)



```

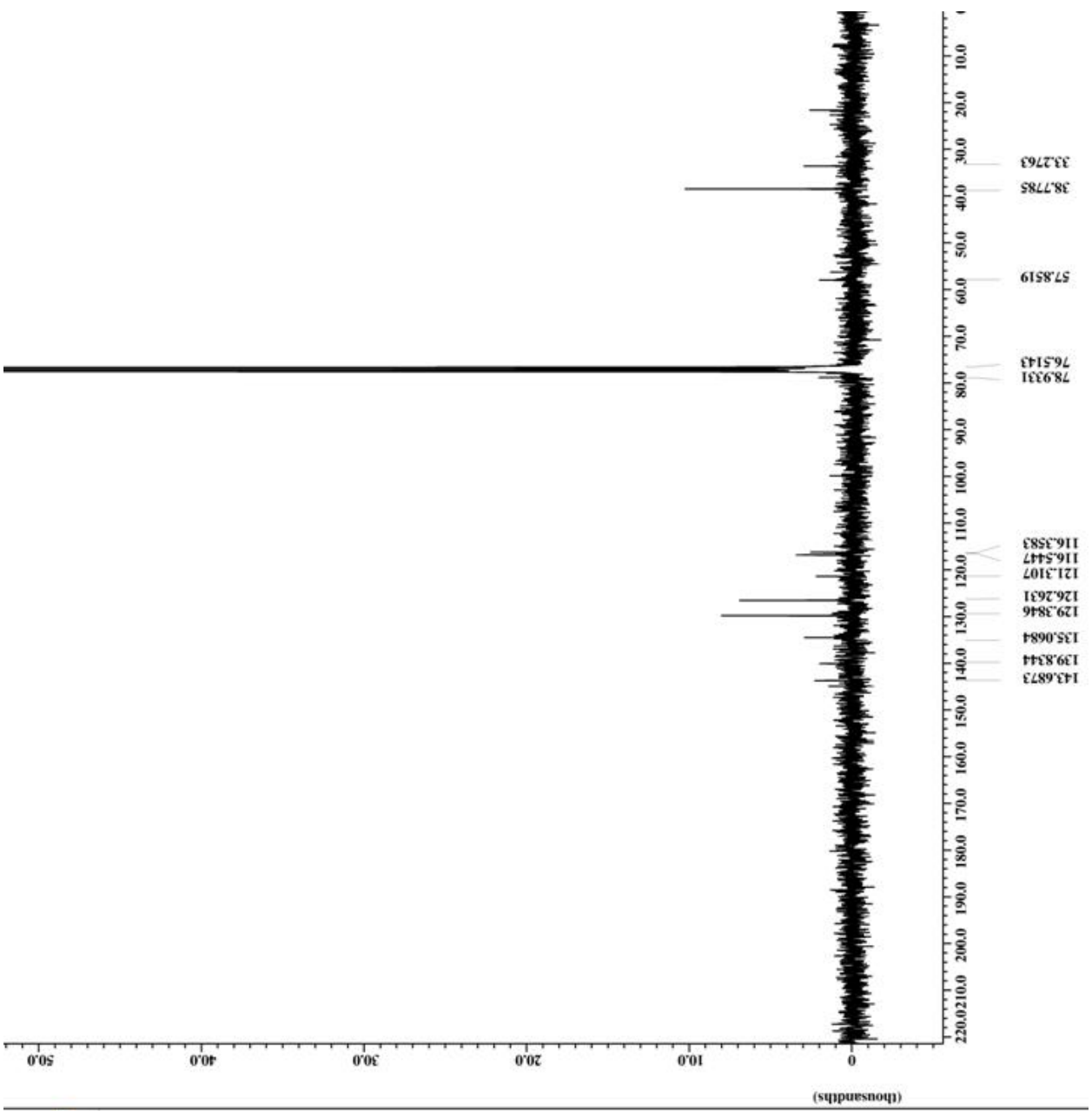
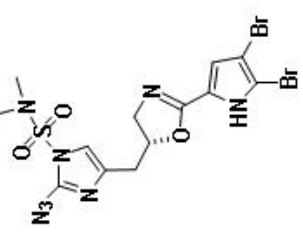
filename = sm_V_21_PURE-4_jdf
author = delta
experiment = single_pulse_exp
sample_id = S831163
solvent = CHLOROFORM-D
reaction_time = 9-DEC-2008 08:55:50
revision_time = 27-MAR-2010 18:03:03
current_time = 27-MAR-2010 18:04:01
comment = Single Pulse Experiment
data_format = ID REAL
in_size = 16384
in_title = 1H
in_units = [ppm]
instruments = X
ite = Eclipse+ 500
nucleus1 = DELTA_NMR
p1 = 11.7473579[T] (500[MH]
p2 = 2.1823486[s]
p3 = 1H
p4 = 500.15991521[MHz]
p5 = 5[ppm]
p6 = 16384
p7 = 0
p8 = 0.45822189[Hz]
p9 = 7.50750751[kHz]
p10 = FALSE
p11 = 1
p12 = 8
p13 = 8
p14 = 18.5[us]
p15 = 2.1823486[s]
p16 = 45[deg]
p17 = 7.25[us]
p18 = 1[s]
p19 = 2[us]
p20 = 20
p21 = 4[s]
p22 = 25.9[dc]
p23 = 2[us]

```





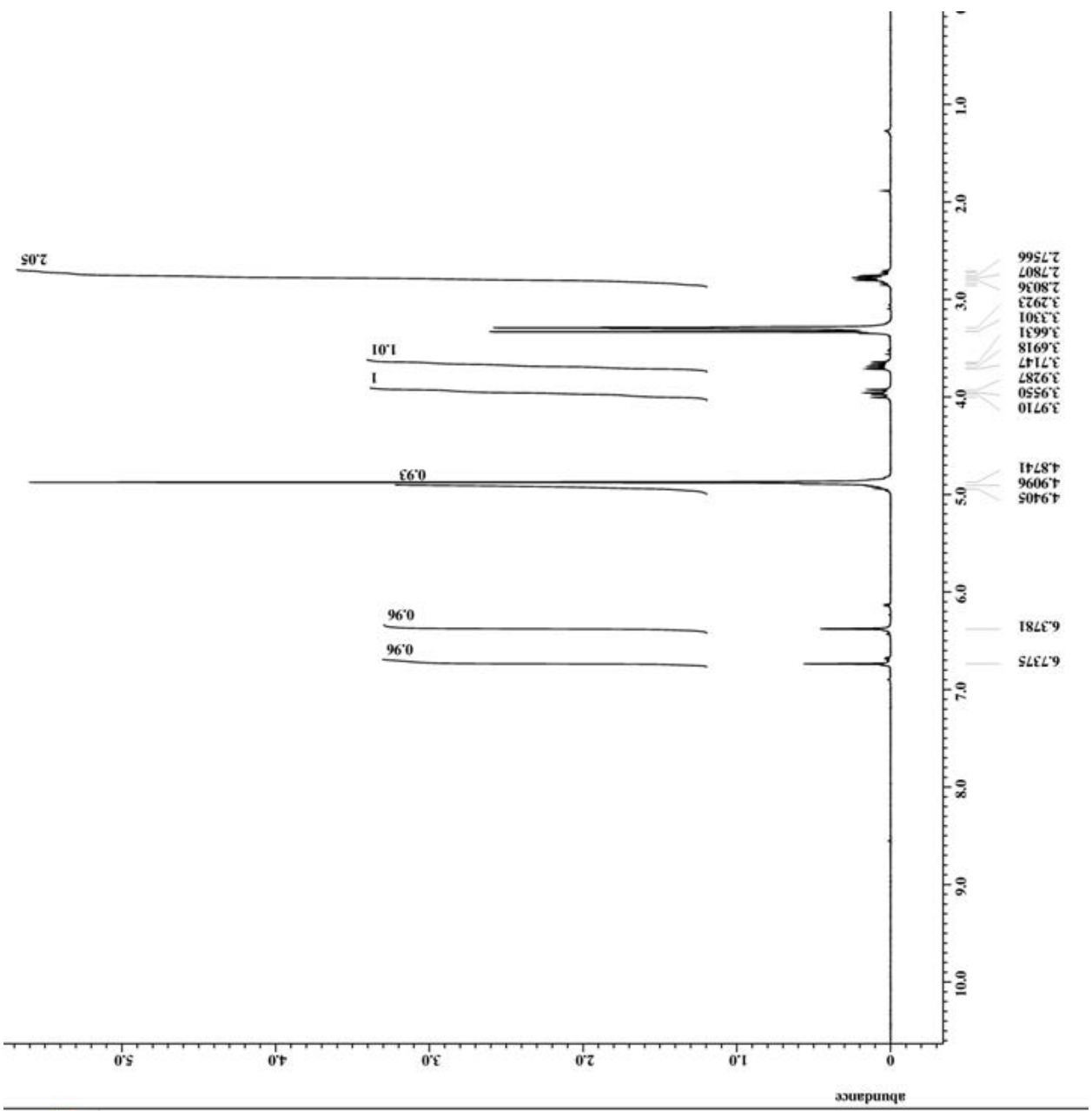
```
filename = sm_V_24_ii-5.jdf
author =
experiment = single_pulse_dec
sample_id = S#490177
solvent = CHLOROFORM-D
acquisition_time = 21-JUL-2009 15:04:31
revision_time = 27-MAR-2010 18:07:40
current_time = 27-MAR-2010 18:08:46
comment = single pulse decouple
ata_format = ID COMPLEX
in_size = 52428
in_title = 13C
in_units = [ppm]
in_dimensions = X
ite = ECX 300
spectrometer = DELTA2_NMR
kernel_strength = 7.0586013[T] (300[MHz]
acq_duration = 2.76824064[s]
domain = 13C
freq = 75.56823426[MHz]
offset = 100[ppm]
points = 65536
prescans = 4
resolution = 0.36124027[Hz]
sweep = 23.67424242[kHz]
tr_domain = 1H
tr_freq = 300.52965592[MHz]
tr_offset = 5[ppm]
lipped = FALSE
bd_return = 10
cans = 1000
atal_scans = 1000
_90_width = 9.75[us]
acq_time = 2.76824064[s]
angle = 30[deg]
atn = 8[db]
pulse = 3.25[us]
tr_atn_dec = 25[db]
tr_atn_noe = 25[db]
tr_noise = WALTZ
scoupling = TRUE
nitral_wait = 1[s]
be_time = TRUE
be_time = 3[s]
ecvr_gain = 50
relaxation_delay = 3[s]
petition_time = 5.76824064[s]
emp_get = 23.1[dc]
```



APPENDIX 9
¹H AND ¹³C NMR SPECTRUM OF
4-(((S)-2-(4,5-dibromo-1H-pyrrol-2-yl)-4,5-dihydrooxazol-5-yl)methyl)-2,3-
dihydro-1H-imidazol-2-amine (57)



```
filename = sm_V_30_pure-5.jdf
author = delta
experiment = single_pulse.ex2
sample_id = S#754445
solvent = METHANOL-D3
creation_time = 12-DEC-2008 21:16:11
revision_time = 12-DEC-2008 21:07:54
current_time = 27-MAR-2010 18:16:17
comment = single_pulse
ata_format = ID REAL
in_size = 13107
in_title = 1H
in_units = [ppm]
inmensions = X
ite = ECX 300
spectrometer = DELTA2_NMR
ield_strength = 7.0586013[T] (300[MHz]
_acq_duration = 2.90717696[s]
_domain = 1H
_freq = 300.52965592[MHz]
_offset = 5[ppm]
_points = 16384
_prescans = 0
_resolution = 0.34397621[Hz]
_sweep = 5.63570784[kHz]
rr_domain = 1H
rr_freq = 300.52965592[MHz]
rr_offset = 5[ppm]
ri_domain = 1H
ri_freq = 300.52965592[MHz]
ri_offset = 5[ppm]
lipped = FALSE
od_return = 1
otal_scans = 24
_90_width = 13.01[us]
_acq_time = 2.90717696[s]
_angle = 45[deg]
_atn = 4[dB]
_pulse = 6.505[us]
rr_mode = Off
ri_mode = Off
ante_presat = FALSE
nitial_wait = 1[s]
scrvr_gain = 44
relaxation_delay = 5[s]
petition_time = 7.90717696[s]
emp_get = 23.2[deg]
```





```

filename = sm_v_30_PURE-2.jdf
author = delta
experiment = single_pulse_dec
pulse_id = S#757410
solvent = METHANOL-D3
reaction_time = 13-DEC-2008 06:54:30
revision_time = 13-DEC-2008 08:18:14
current_time = 27-MAR-2010 18:19:24

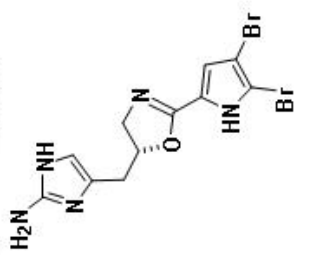
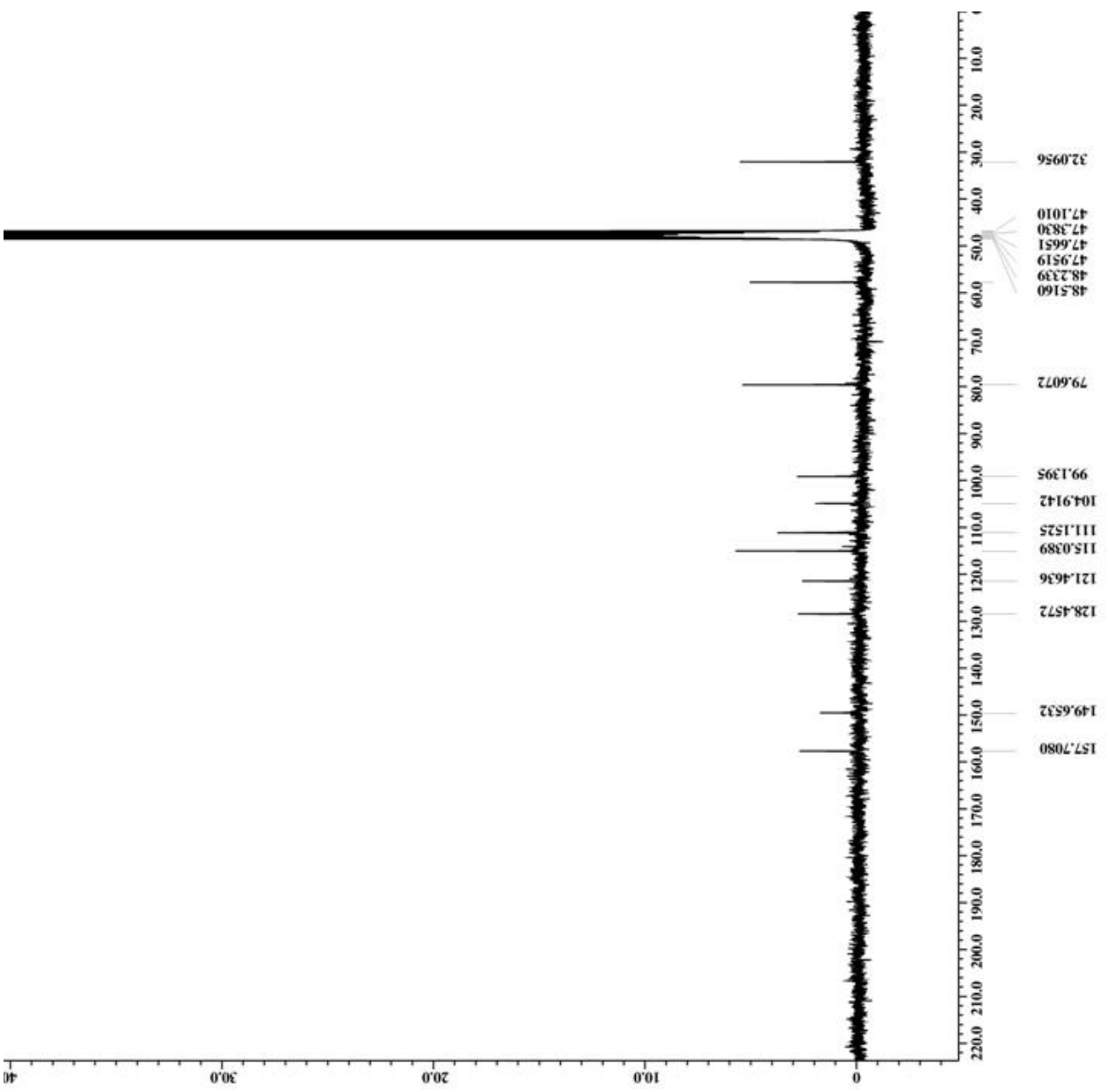
comment = single pulse decouple
ata_format = ID COMPLEX
in_size = 52428
in_title = 13C
in_units = [ppm]
in_dimensions = X
ite = ECX 300
spectrometer = DELTA2_NMR

ield_strength = 7.0586013[T] (300[MHz]
acq_duration = 2.76824064[s]
domain = 13C
freq = 75.56823426[MHz]
offset = 100[ppm]
points = 65536
prescans = 4
resolution = 0.36124027[Hz]
sweep = 23.67424242[kHz]
rr_domain = 1H
rr_freq = 300.52965592[MHz]
rr_offset = 5[ppm]
lipped = FALSE
bd_return = 10
cans = 6000
total_scans = 6000

_90_width = 9.75[us]
acq_time = 2.76824064[s]
angle = 30[deg]
atn = 8[dB]
pulse = 3.25[us]
rr_atn_dec = 25[dB]
rr_atn_noe = 25[dB]
rr_noise = WALTZ
scoupling = TRUE
initial_wait = 1[s]
oe = TRUE
oe_time = 3[s]
ecvr_gain = 50
relaxation_delay = 3[s]
petition_time = 5.76824064[s]
emp_get = 23.3[dc]

```

(thousands)



APPENDIX 10
¹H AND ¹³C NMR SPECTRUM OF
(S)-4-((2-(1H-pyrrol-2-yl)-4,5-dihydrooxazol-5-yl)methyl)-2-azido-N,N-
dimethyl-1H-imidazole-1-sulfonamide (62)



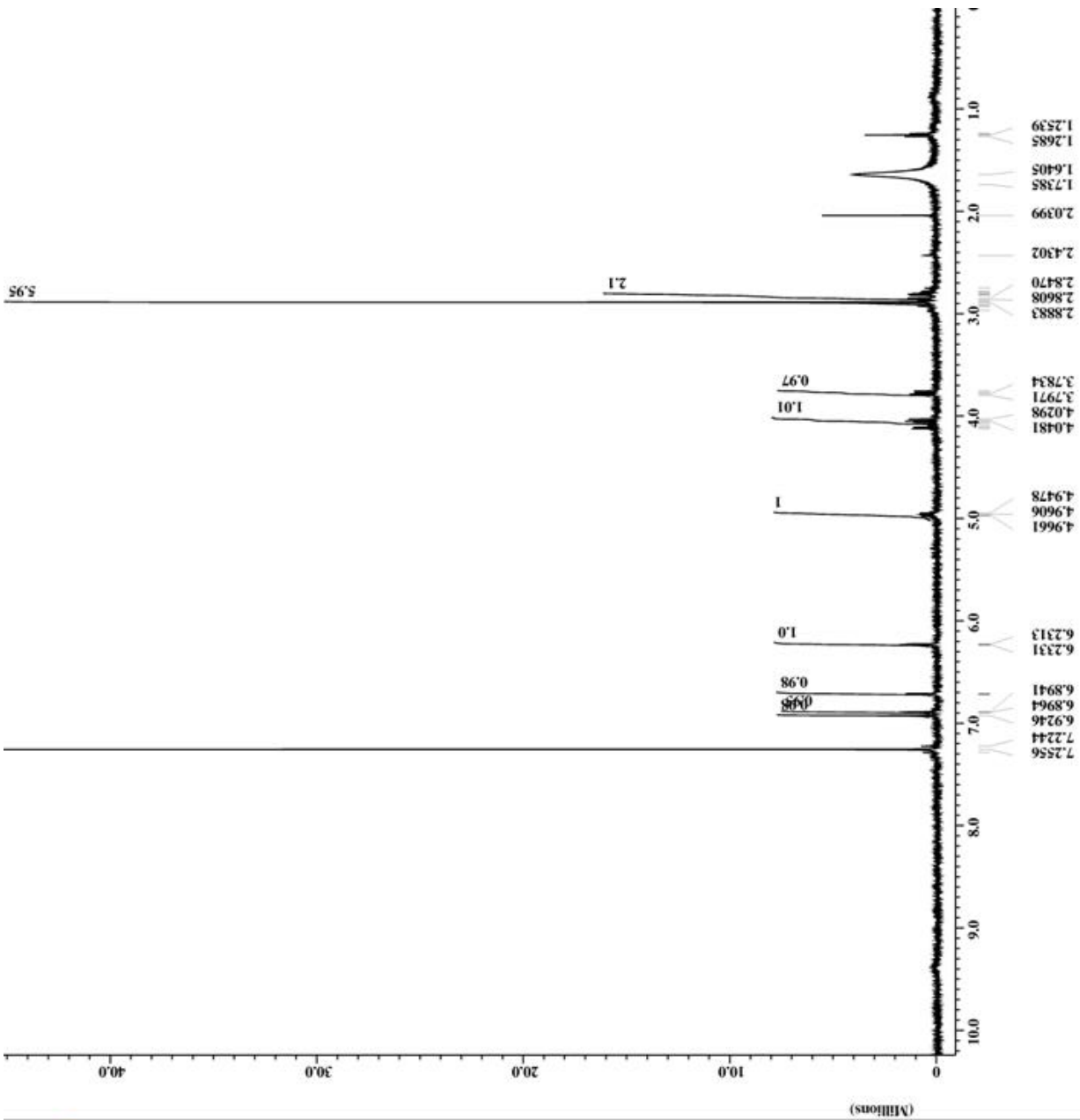
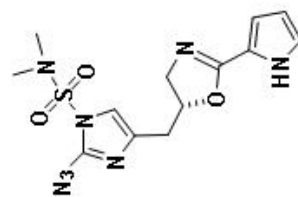
```

ilname      = sm_azide_old-3.jdf
author      = delta
experiment  = single_pulse_exp
sample_id   = S8675675
solvent     = CHLOROFORM-D
revision    = 13-JUN-2007 22:21:26
evision     = 27-MAR-2010 18:36:56
current_time = 27-MAR-2010 18:37:37

comment     = Single Pulse Experiment
sta_format  = ID COMPLEX
im_size     = 16384
im_title    = 1H
im_units    = [ppm]
imensions  = X
ite         = Eclipse+ 500
spectrometer = DELTA_NMR

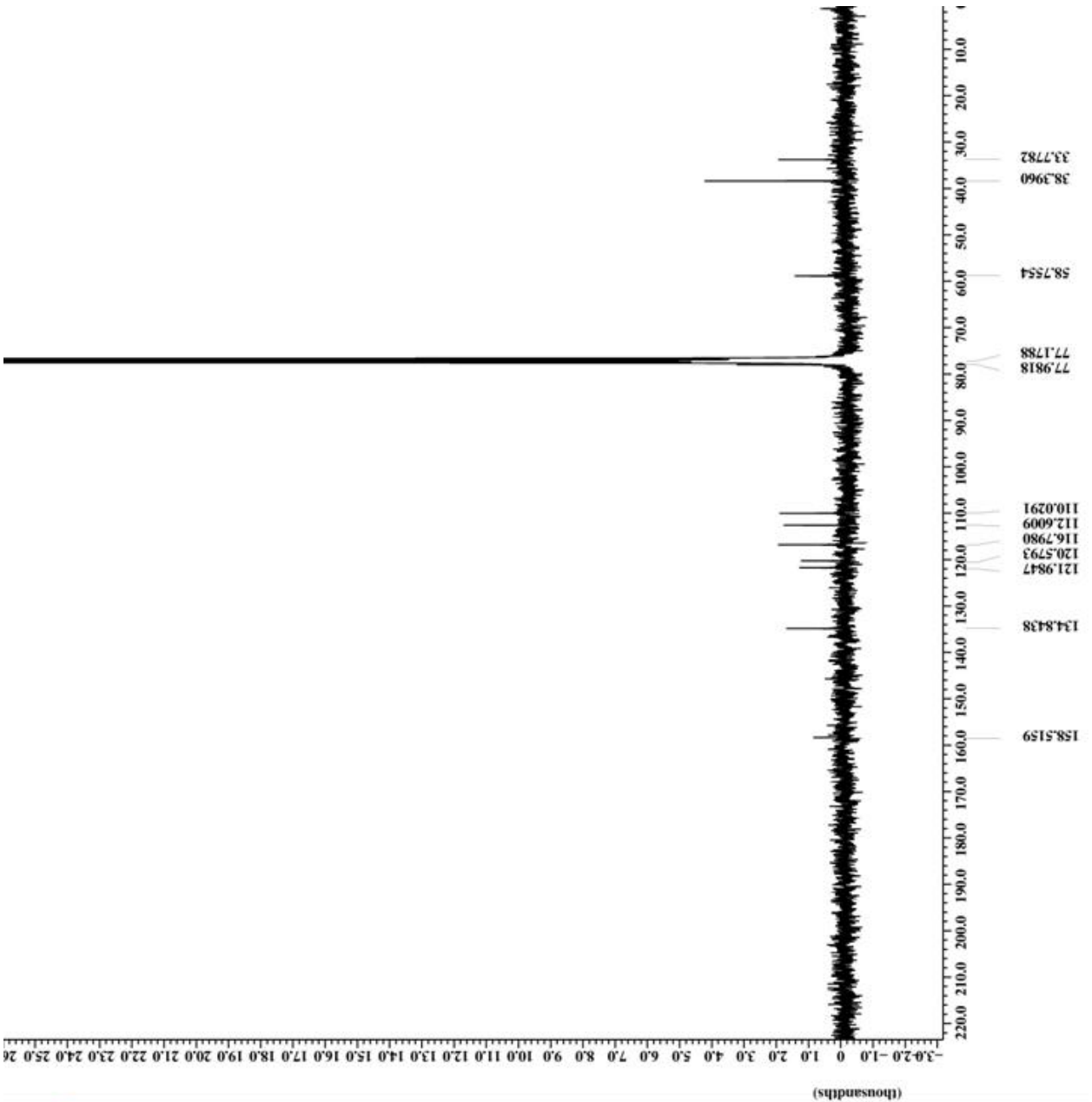
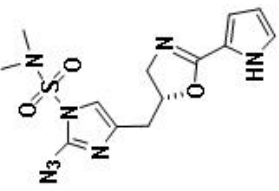
field_strength = 11.7473579[T] (500[MH
-acq_duration = 2.1823488[s]
-domain       = 1H
-freq        = 500.15991521[MHz]
-offset      = 5[ppm]
-points      = 16384
-prescans    = 0
-resolution  = 0.45822189[Hz]
-sweep       = 7.50750751[kHz]
-lipped      = FALSE
-bd_return   = 1
-cans        = 8
-stal_scans  = 8

-90_width    = 18.5[us]
-acq_time    = 2.1823488[s]
-angle       = 45[deg]
-pulse       = 9.25[us]
-pitch_wait  = 5[s]
-base_preset = 5[us]
-sevr_gain   = 27
-relaxation_delay = 4[s]
-emp_get     = 24.9[dc]
-nblank_time = 2[us]
  
```





```
ilname = sm_II_AZIDE_PURE-3_jd
uthor = delta
xperiment = single_pulse_dec
mple_id = S8709802
ilvent = CHLOROFORM-D
reation_time = 14-JAN-2007 06:02:45
evision_time = 27-MAR-2010 18:43:04
urrent_time = 27-MAR-2010 18:43:18
omment = single pulse decouple
ate_format = 1D COMPLEX
m_size = 52428
m_title = 13C
m_units = [ppm]
mensions = X
ite = ECX 300
ectrometer = DELTA2_NMR
ield_strength = 7.0586013[T] (300[MHz]
cq_duration = 2.76824064[s]
omain = 13C
req = 75.56823426[MHz]
ffset = 100[ppm]
oints = 65536
rescans = 4
esolution = 0.36124027[Hz]
sweep = 23.67424242[KHz]
r_domain = 1H
r_freq = 300.52965592[MHz]
r_offset = 5[ppm]
lipped = FALSE
d_return = 10
ans = 6000
tal_scans = 6000
_90_width = 9.75[us]
cq_time = 2.76824064[s]
ngle = 30[deg]
atn = 8[db]
pulse = 3.25[us]
r_atn_dec = 25[db]
r_atn_noe = 25[db]
r_noise = WALTZ
coupling = TRUE
nitial_wait = 1[s]
ce = TRUE
e_time = 3[s]
svr_gain = 50
elaxation_delay = 3[s]
epitation_time = 5.76824064[s]
emp_get = 23.6[dc]
```



APPENDIX 11
¹H AND ¹³C NMR SPECTRUM OF
4-(((S)-2-(1H-pyrrol-2-yl)-4,5-dihydrooxazol-5-yl)methyl)-2,3-dihydro-1H-
imidazol-2-amine (64)



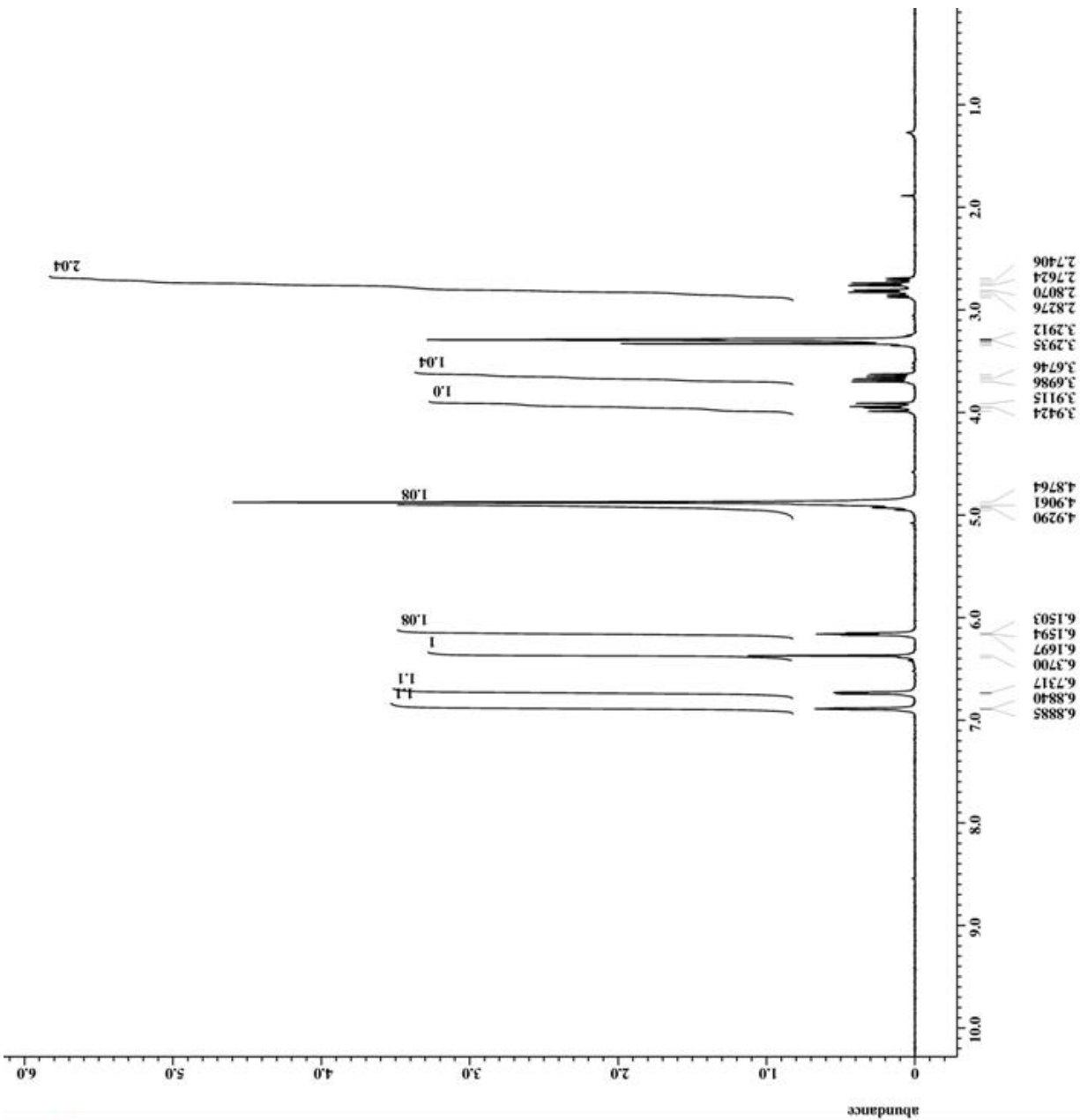
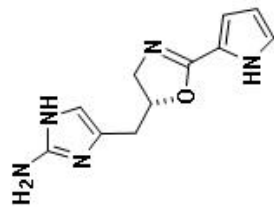
```

ilname      = sm_v_28_pure_2-3_3.jdf
author      = delta
experiment  = single_pulse.ex2
sample_id   = 2
solvent     = METHANOL-D3
reaction_time = 11-DEC-2008 22:37:18
revision_time = 11-DEC-2008 22:24:14
current_time = 27-MAR-2010 18:47:18

comment     = single_pulse
date_format = DD.MM.YY
im_size     = 13107
im_title    = 1H
im_units    = [ppm]
imensions  = X
ite         = ECX 300
spectrometer = DELTA2_NMR

field_strength = 7.0586013[T] (300[MHz])
-acq_duration  = 2.90717696[s]
-domain        = 1H
-freq          = 300.52965592[MHz]
-gamma         = 5[ppm]
-offset        = 16384
-points        = 0
-prescans      = 0
-resolution    = 0.34397631[Hz]
-sweep         = 5.63570784[kHz]
-rr_domain    = 1H
-rr_freq       = 300.52965592[MHz]
-rr_offset     = 5[ppm]
-ri_domain     = 1H
-ri_freq       = 300.52965592[MHz]
-ri_offset     = 5[ppm]
-lipped        = FALSE
-bd_return     = 1
-cans          = 11
-stal_scans    = 11

-90_width      = 13.01[us]
-acq_time      = 2.90717696[s]
-angle         = 45[deg]
-atn           = 4[dB]
-pulse         = 6.505[us]
-rr_mode       = Off
-rr_mode       = Off
-ante_presat   = FALSE
-ntial_wait    = 1[e]
-scvr_gain     = 46
-relaxation_delay = 5[s]
-acquisition_time = 7.90717696[s]
-emp_get       = 23.1[dC]
  
```





```

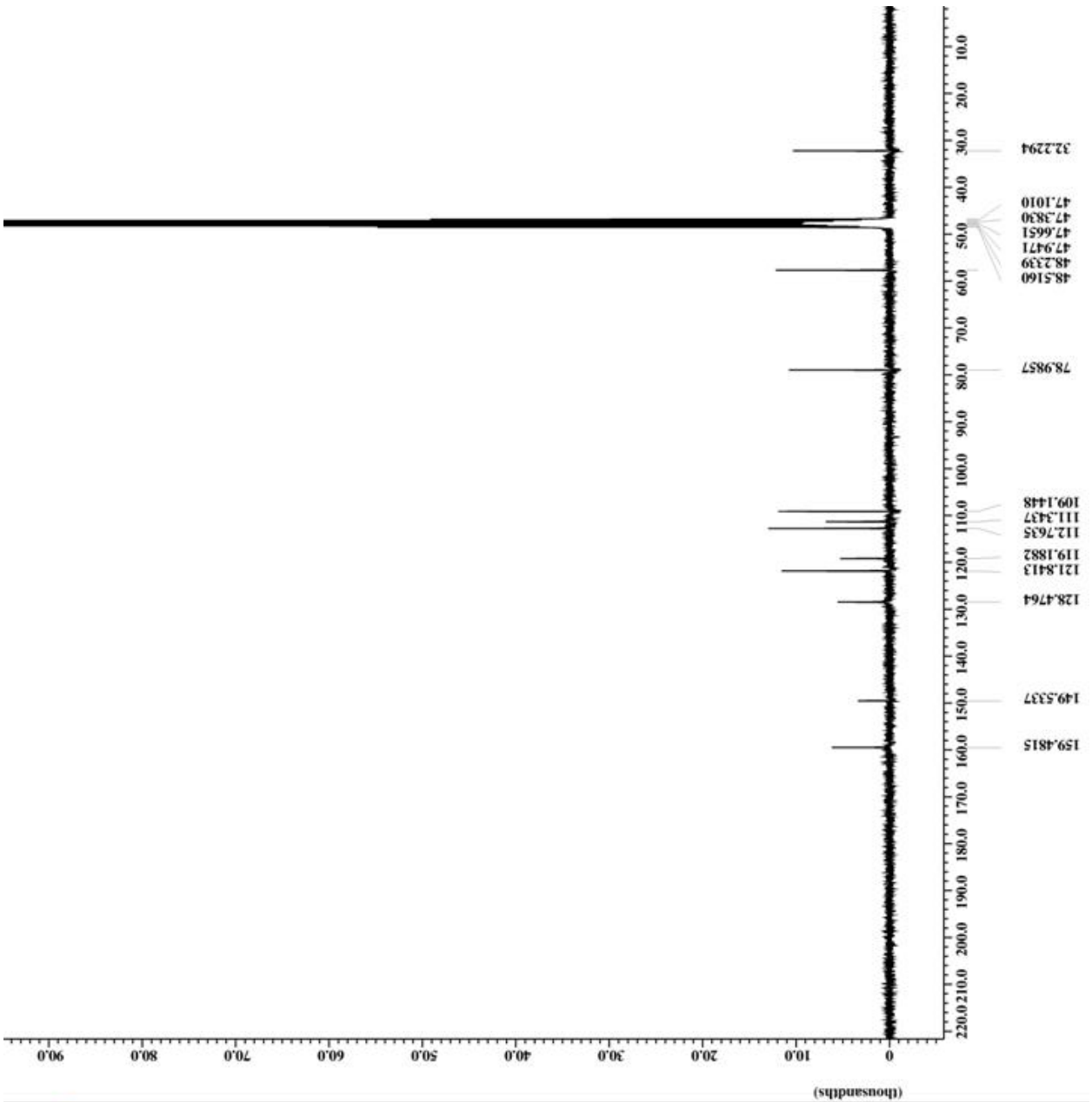
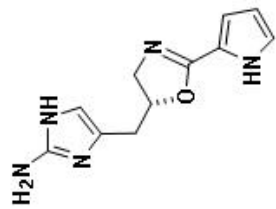
filename = sm_V_28_2-3_3.jdf
author = delta
experiment = single_pulse_dec
sample_id = S#805616
solvent = METHANOL-D3
reaction_time = 12-DEC-2008 05:02:27
revision_time = 12-DEC-2008 14:21:39
current_time = 27-MAR-2010 18:50:48

comment = single pulse decouple
ata_format = ID REAL
in_size = 52428
in_title = 13C
in_units = [ppm]
inmensions = X
ite = ECX 300
spectrometer = DELTA2_NMR

ield_strength = 7.0586013[T] (300[MHz]
acq_duration = 2.76824064[s]
domain = 13C
freq = 75.56823426[MHz]
offset = 100[ppm]
points = 65536
prescans = 4
resolution = 0.36124027[Hz]
sweep = 23.67424242[kHz]
tr_domain = 1H
tr_freq = 300.52965592[MHz]
tr_offset = 5[ppm]
lipped = FALSE
bd_return = 10
cans = 4000
otal_scans = 4000

_90_width = 9.75[us]
acq_time = 2.76824064[s]
angle = 30[deg]
atn = 8[db]
pulse = 3.25[us]
tr_atn_dec = 25[db]
tr_atn_noe = 25[db]
tr_noise = WALTZ
scoupling = TRUE
nitial_wait = 1[s]
oe_time = TRUE
oe_time = 3[s]
ecvr_gain = 50
elaxation_delay = 3[s]
epetition_time = 5.76824064[s]
emp_get = 23.5[dc]

```



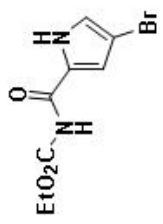
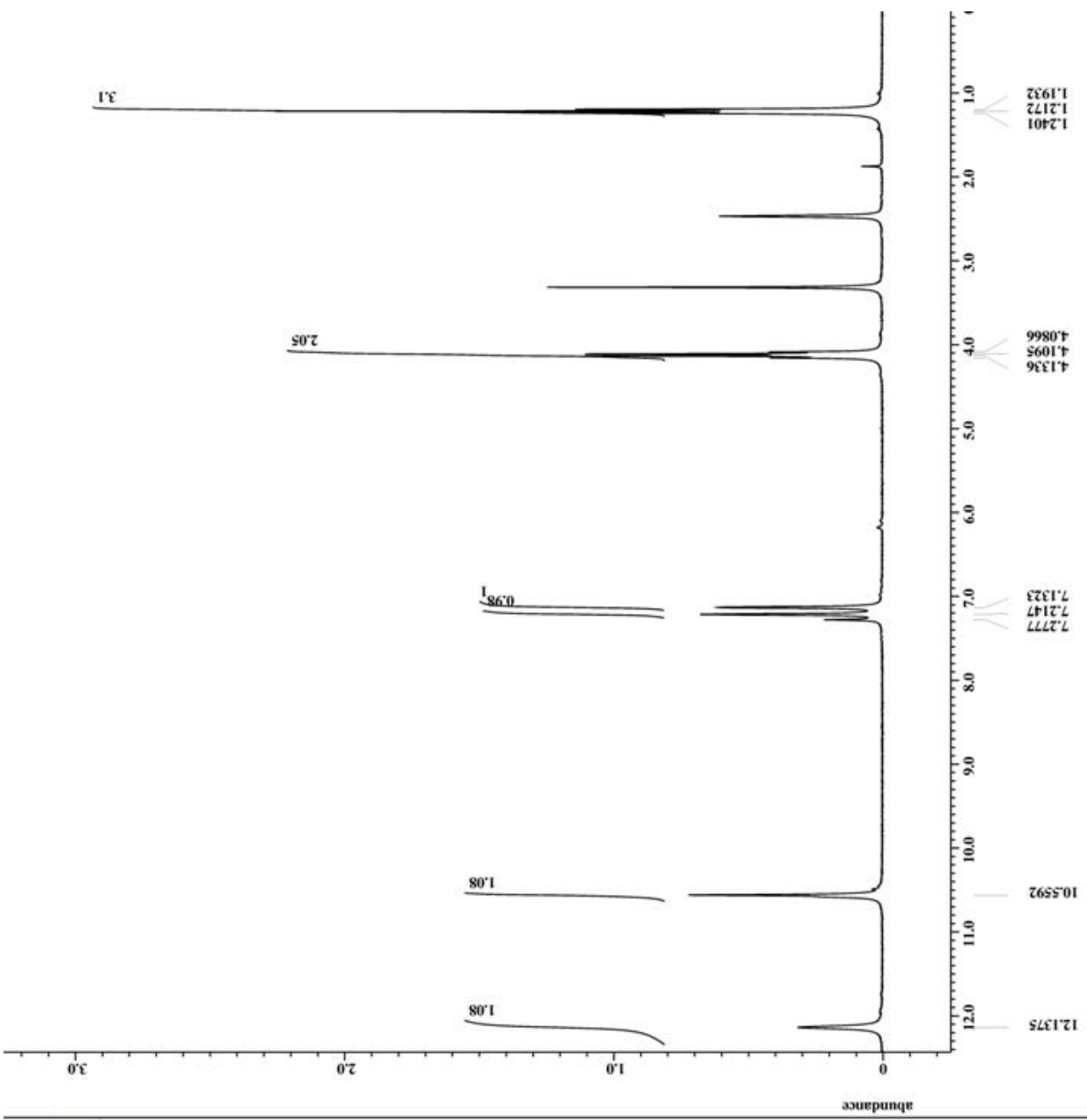
APPENDIX 12
¹H AND ¹³C NMR SPECTRUM OF
Ethyl 4-bromo-1H-pyrrole-2-carbonylcarbamate (65)



```

filename = sm_VI_88_pure-4_jdf
author = delta
experiment = single_pulse.ex2
sample_id = S821820
solvent = DMSO-D6
reaction_time = 8-JAN-2010 22:38:56
acquisition_time = 27-MAR-2010 18:57:10
current_time = 27-MAR-2010 18:57:43
comment = single_pulse
ata_format = ID COMPLEX
in_size = 13107
in_title = 1H
in_units = [ppm]
inmensions = X
ite = ECX 300
spectrometer = DELTA2_NMR
ield_strength = 7.0586013[T] (300[MHz]
_acq_duration = 2.90717696[s]
_domain = 1H
_freq = 300.52965592[MHz]
_offset = 5[ppm]
_points = 16384
_prescans = 0
_resolution = 0.34397621[Hz]
_sweep = 5.63570784[kHz]
tr_domain = 1H
tr_freq = 300.52965592[MHz]
tr_offset = 5[ppm]
ri_domain = 1H
ri_freq = 300.52965592[MHz]
ri_offset = 5[ppm]
lipped = FALSE
od_return = 1
otal_scans = 24
_90_width = 13.01[us]
_acq_time = 2.90717696[s]
_angle = 45[deg]
_atn = 4[dB]
_pulse = 6.505[us]
tr_mode = Off
ri_mode = Off
ante_presat = FALSE
ntial_wait = 1[s]
scvr_gain = 42
relaxation_delay = 5[s]
petition_time = 7.90717696[s]
emp_get = 20.8[dc]

```



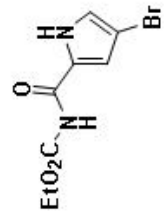
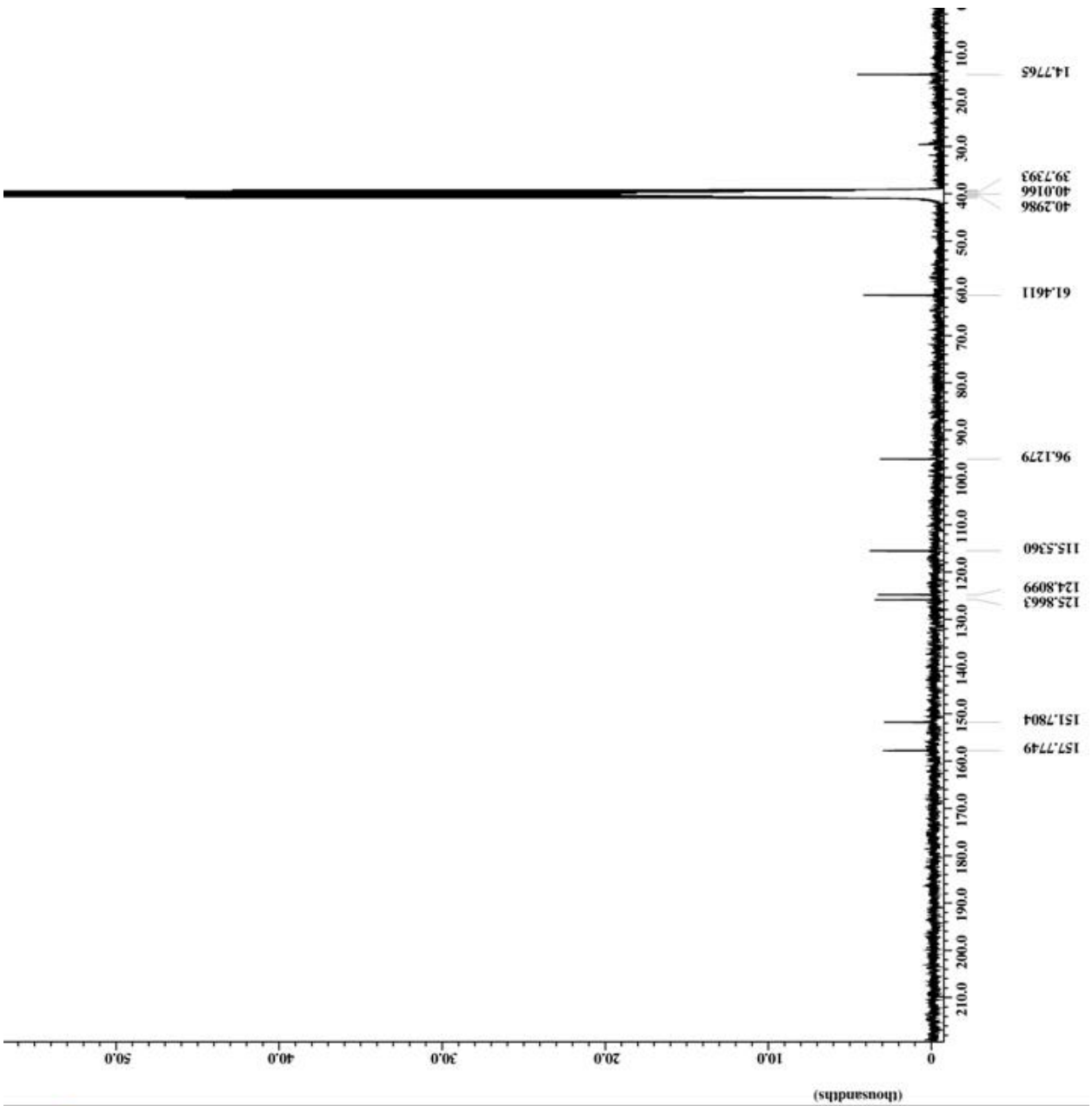


```
ilname = sm_VI_precursor_monob
uthor = delta
xperiment = single_pulse_dec
mple_id = S837678
olvent = DMSO-D6
eaction_time = 4-FEB-2010 08:42:09
evison_time = 4-FEB-2010 11:39:13
urrent_time = 19-APR-2010 19:29:02

omment = single pulse decouple
ate_format = ID COMPLEX
m_size = 52428
m_title = 13C
m_units = [ppm]
mensions = X
ite = ECX 300
ectrometer = DELTA2_NMR

ield_strength = 7.0586013[T] (300[MHz]
cq_duration = 2.76824064[s]
omain = 13C
req = 75.56823426[MHz]
ffset = 100[ppm]
oints = 65336
rescans = 4
esolution = 0.36124027[Hz]
sweep = 23.67424242[KHz]
r_domain = 1H
r_freq = 300.52965592[MHz]
r_offset = 5[ppm]
lipped = FALSE
d_return = 10
ans = 6000
tal_scans = 6000

_90_width = 9.75[us]
cq_time = 2.76824064[s]
angle = 30[deg]
atn = 8[db]
pulse = 3.25[us]
r_atn_dec = 25[db]
r_atn_noe = 25[db]
r_noise = WALTZ
scoupling = TRUE
nitital_wait = 1[s]
ce = TRUE
e_time = 3[s]
ecvr_gain = 50
elaxation_delay = 3[s]
eapitation_time = 5.76824064[s]
emp_get = 19.7[dC]
```



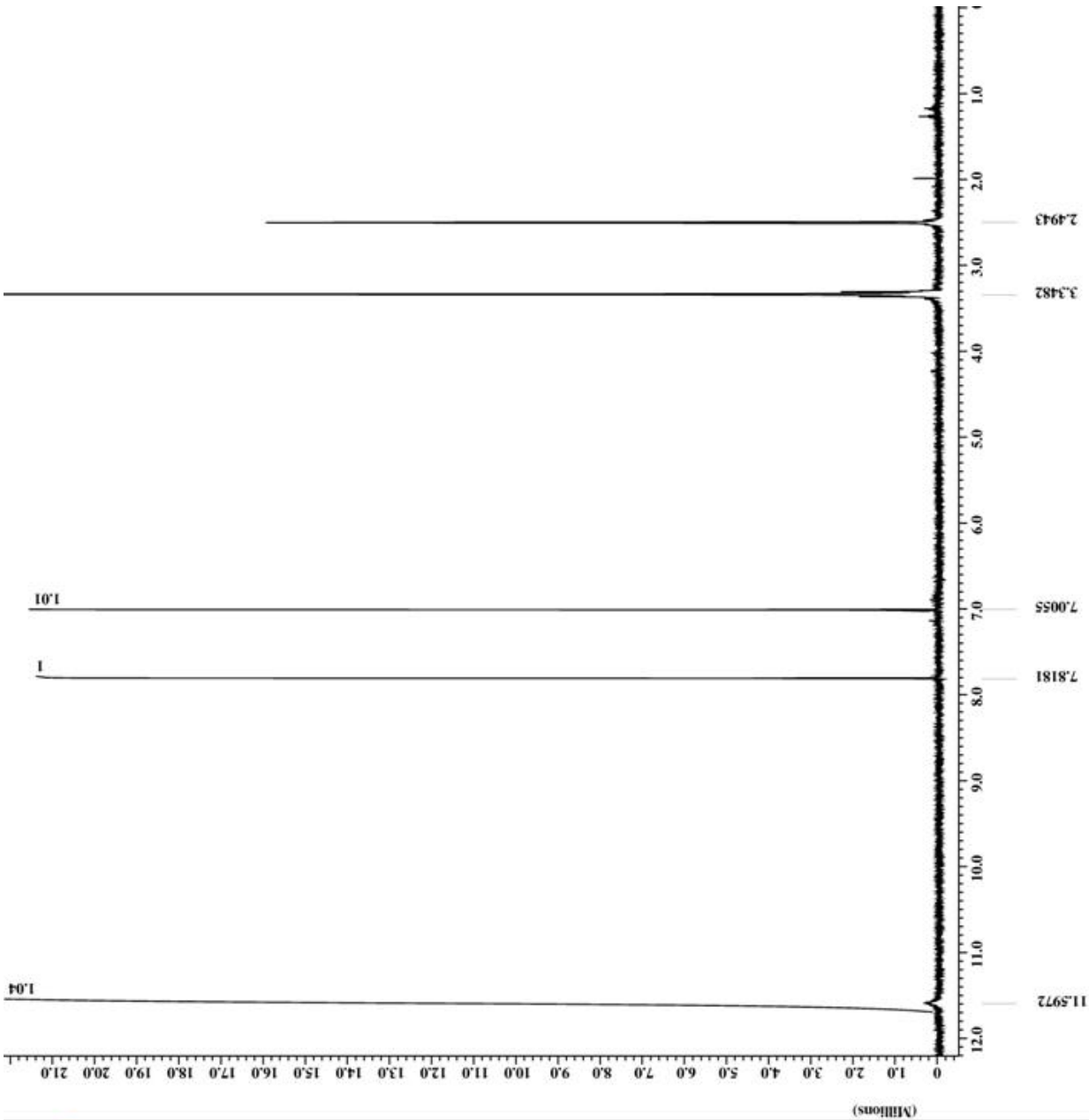
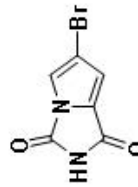
APPENDIX 13
¹H AND ¹³C NMR SPECTRUM OF
6-bromo-1H-pyrrolo[1,2-c]imidazole-1,3(2H)-dione (67)



```

filename = sm_IV_98_pure-5.jdf
author = delta
experiment = single_pulse.exp
sample_id = S9767561
solvent = DMSO-d6
reaction_time = 13-SEP-2008 04:47:28
revision_time = 27-MAR-2010 19:17:27
current_time = 27-MAR-2010 19:17:43
comment = Single Pulse Experiment
data_format = ID COMPLEX
im_size = 16384
im_title = 1H
im_units = [ppm]
imensions = X
ite = Eclipse+ 500
spectrometer = DELTA_NMR
field_strength = 11.747379[T] (500[MH
-acq_duration = 2.182348[s]
-domain = 1H
-freq = 500.15991521[MHz]
-gamma = 5[ppm]
-points = 16384
-prescans = 0
-resolution = 0.45822189[Hz]
-sweep = 7.50750751[kHz]
-lipped = FALSE
od_return = 1
cans = 3
otal_scans = 3
_90_width = 18.5[us]
-acq_time = 2.182348[s]
-angle = 45[deg]
-pulse = 9.25[us]
-nitral_wait = 1[s]
-base_preset = 2[us]
-ecvr_gain = 22
-relaxation_delay = 4[s]
-emp_get = 44.7[dc]
-nblank_time = 2[us]

```





```

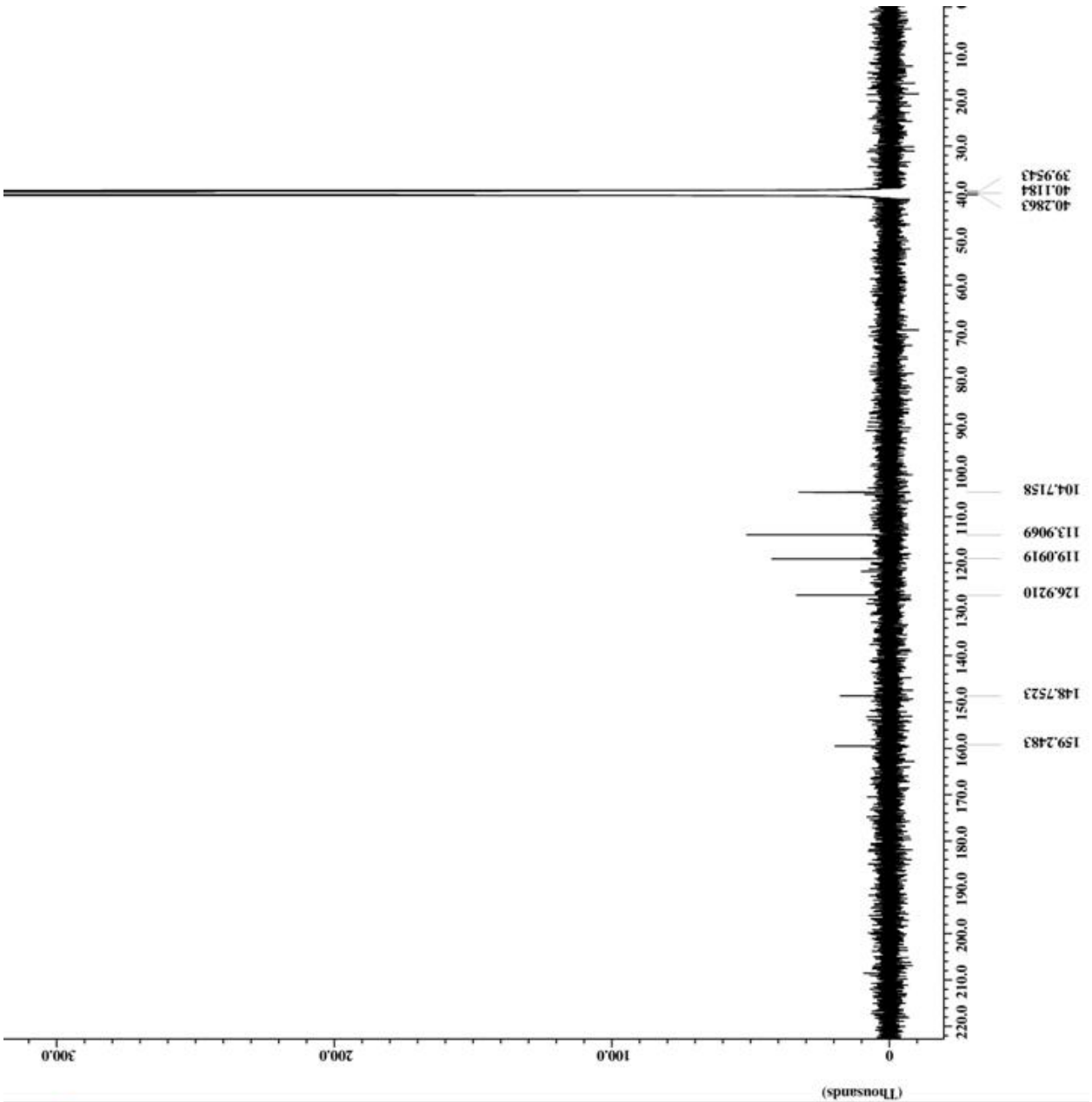
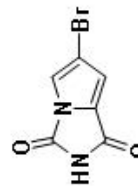
filename = sm_IV_98_pure-3.jdf
author = delta
experiment = single_pulse_dec
pulse_id = S#768816
solvent = DMSO-d6
reaction_time = 13-SEP-2008 16:08:08
revision_time = 13-SEP-2008 11:44:21
current_time = 27-MAR-2010 19:20:43

comment = single pulse decouple
ata_format = ID REAL
im_size = 65536
im_title = 13C
im_units = [ppm]
imensions = X
ite = Eclipse+ 500
spectrometer = DELTA_NMR

field_strength = 11.7473579[T] (500[MH
-acq_duration = 2.0840448[s]
-domain = 13C
-freq = 125.76529768[MHz]
-offset = 100[ppm]
-points = 65536
-prescans = 4
-resolution = 0.47983613[Hz]
-sweep = 31.44654088[kHz]
-tr_domain = 1H
-tr_freq = 500.15991521[MHz]
-tr_offset = 5[ppm]
-tipped = TRUE
-bd_return = 10
-cans = 8000
-etal_scans = 8000

_90_width = 14.2[us]
-acq_time = 2.0840448[s]
-angle = 30[deg]
-pulse = 4.73333333[us]
-nitai_wait = 1[s]
-pe_time = 1[s]
-base_preset = 3[us]
-ecvr_gain = 30
-relaxation_delay = 2[s]
-emp_get = 28.8[dC]
-nblank_time = 2[us]

```



APPENDIX 14
¹H AND ¹³C NMR SPECTRUM OF
(*R*)-4-(3-(6-bromo-1,3-dioxo-1H-pyrrolo[1,2-c]imidazol-2(3H)-yl)-2-chloropropyl)-N,N-dimethyl-1H-imidazole-1-sulfonamide (69)



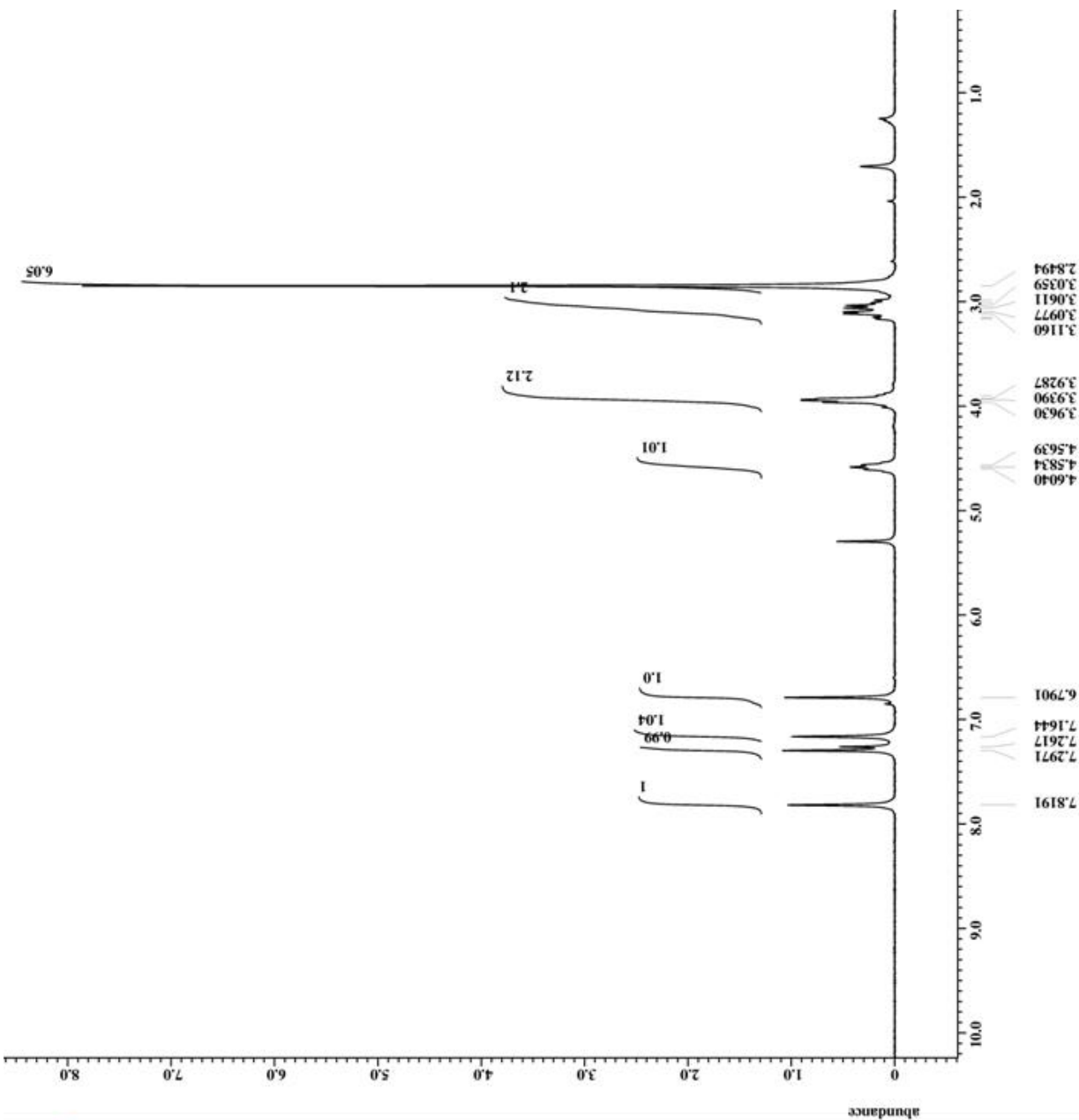
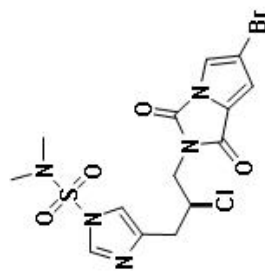
```

ilname      = sm_V_154_pure_i1-4_jd
author      = delta
experiment  = single_pulse.ex2
sample_id   = S8739199
solvent     = CHLOROFORM-D
reaction_time = 29-JUN-2009 20:23:34
acquisition_time = 27-MAR-2010 19:24:06
current_time    = 27-MAR-2010 19:24:41

comment     = single_pulse
ata_format  = ID REAL
in_size     = 13107
in_title    = 1H
in_units    = [ppm]
in_dims     = X
ite         = ECX 300
spectrometer = DELTA2_NMR

ield_strength = 7.0586013[T] (300 [MHz]
_acq_duration = 2.90717696[s]
_domain       = 1H
_freq         = 300.52965592 [MHz]
_offset       = 5[ppm]
_points       = 16384
_prescans     = 0
_resolution   = 0.34397621[Hz]
_sweep        = 5.63570784[kHz]
tr_domain    = 1H
tr_freq      = 300.52965592 [MHz]
tr_offset    = 5[ppm]
ri_domain    = 1H
ri_freq      = 300.52965592 [MHz]
ri_offset    = 5[ppm]
lipped       = FALSE
od_return    = 1
otal_scans   = 21

_90_width    = 13.01[us]
_acq_time    = 2.90717696[s]
_angle       = 45[deg]
_atn         = 4[db]
_pulse       = 6.505[us]
tr_mode      = Off
ri_mode      = Off
ante_presat  = FALSE
nitial_wait  = 1[s]
ecvr_gain    = 50
relaxation_delay = 5[s]
opetition_time = 7.90717696[s]
emp_get      = 23.2[dc]
  
```





```

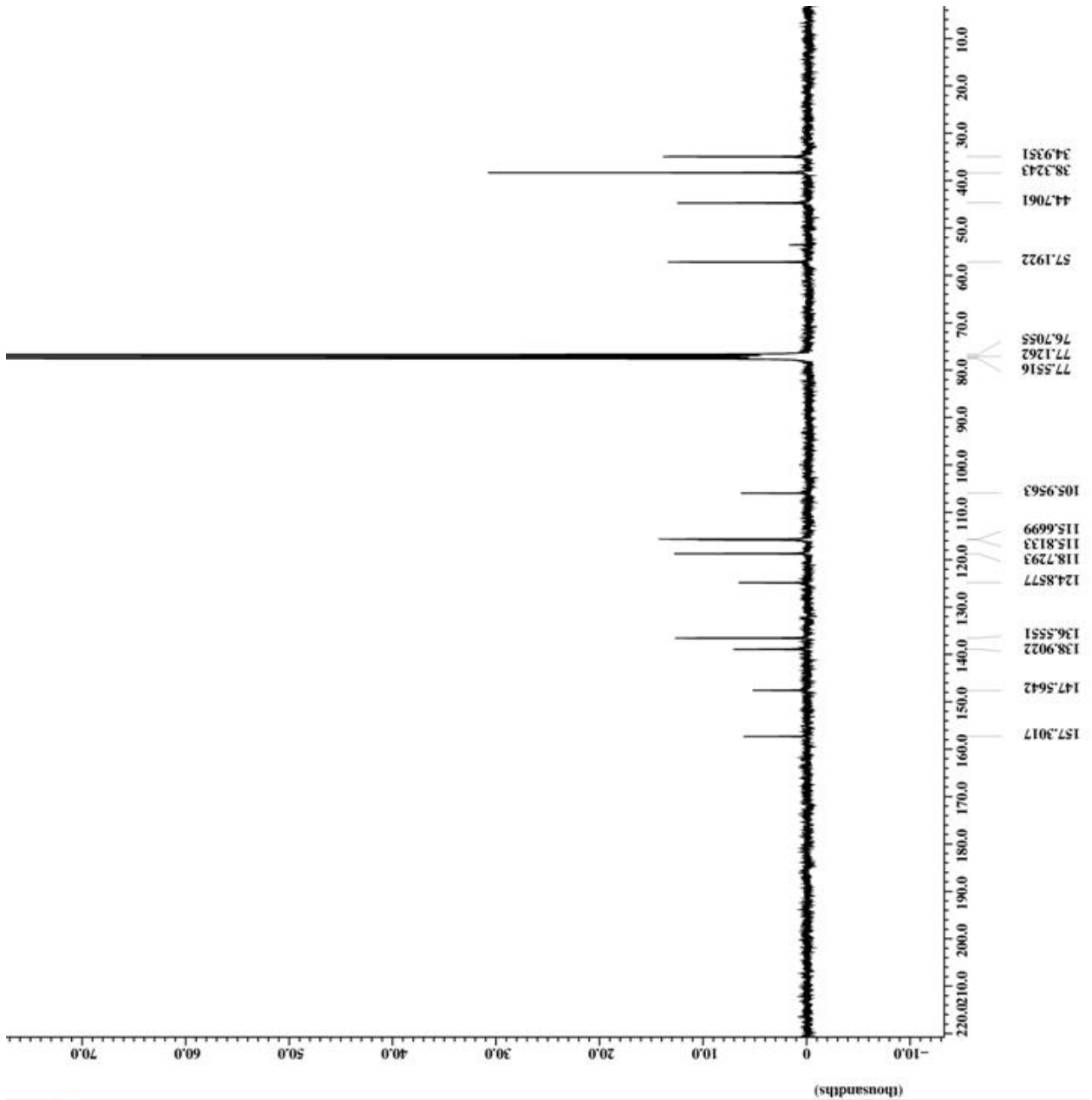
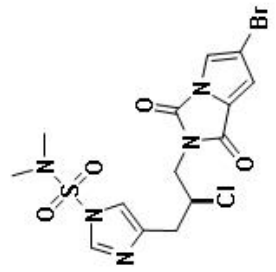
filename = sm_V_154_pure_ii-3.jd
author = delta
experiment = single_pulse_dec
sample_id = S#741095
solvent = CHLOROFORM-D
acquisition_time = 30-JUN-2009 02:48:45
revision_time = 30-JUN-2009 12:04:36
current_time = 27-MAR-2010 19:27:23

comment = single pulse decouple
ata_format = ID COMPLEX
in_size = 52428
in_title = 13C
in_units = [ppm]
in_dimensions = X
ite = ECX 300
spectrometer = DELTA2_NMR

ield_strength = 7.0586013[T] (300[MHz])
_acq_duration = 2.76824064[s]
_domain = 13C
_freq = 75.56823426[MHz]
_offset = 100[ppm]
_points = 65536
_prescans = 4
_resolution = 0.36124027[Hz]
_sweep = 23.67424242[kHz]
_tr_domain = 1H
_tr_freq = 300.52965592[MHz]
_tr_offset = 5[ppm]
_lipped = FALSE
_bd_return = 10
_cans = 4000
_otat_scans = 4000

_90_width = 9.75[us]
_acq_time = 2.76824064[s]
_angle = 30[deg]
_atn = 8[db]
_pulse = 3.25[us]
_tr_atn_dec = 25[db]
_tr_atn_noe = 25[db]
_tr_noise = WALTZ
scoupling = TRUE
initial_wait = 1[s]
oe_time = TRUE
oe_time = 3[s]
ecvr_gain = 50
relaxation_delay = 3[s]
spetition_time = 5.76824064[s]
emp_get = 23.3[dc]

```

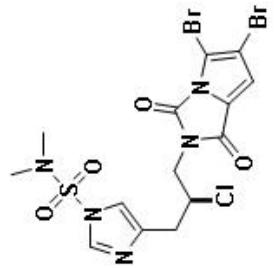
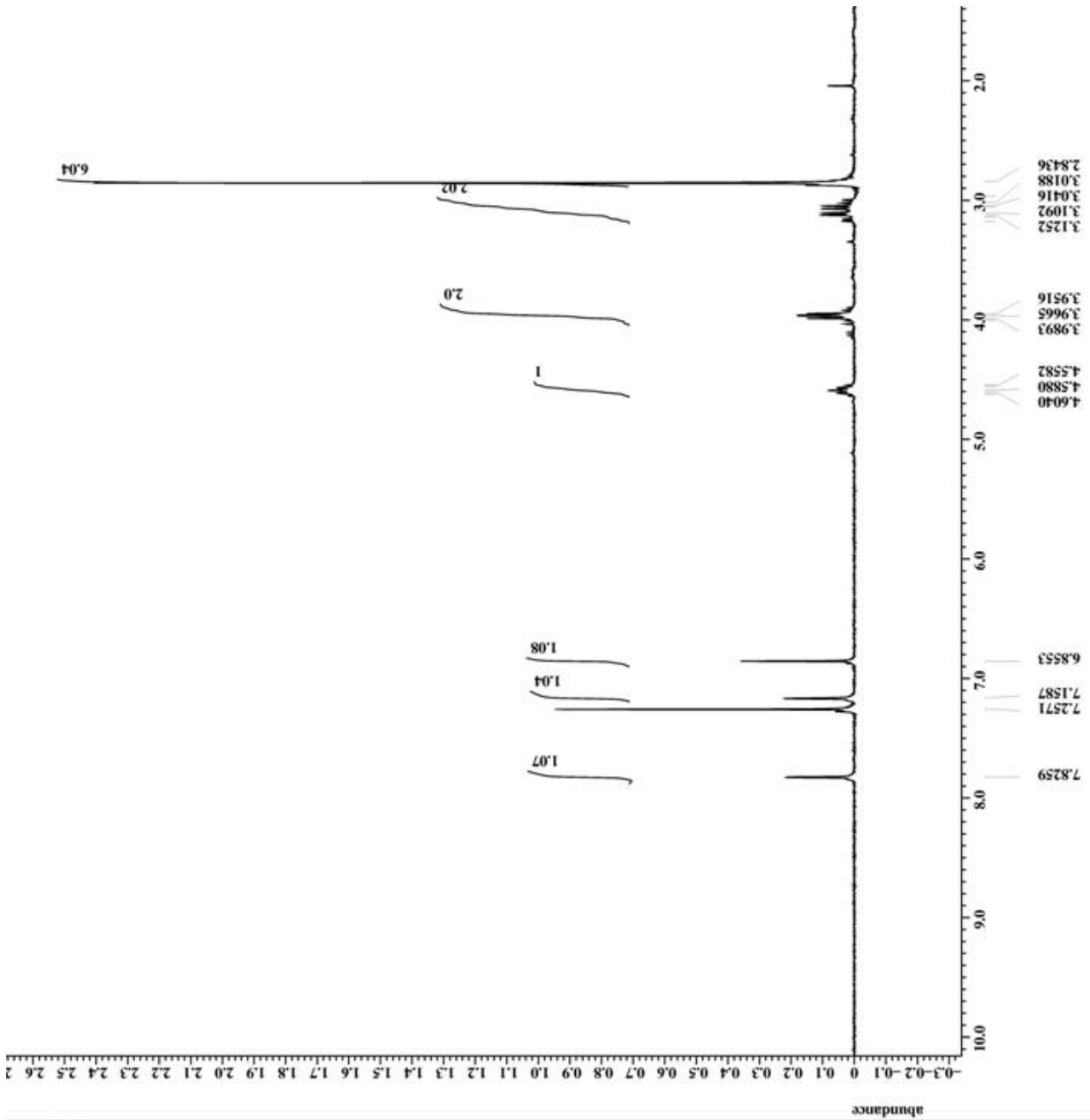


APPENDIX 15
¹H AND ¹³C NMR SPECTRUM OF
(*R*)-4-(2-chloro-3-(5,6-dibromo-1,3-dioxo-1H-pyrrolo[1,2-c]imidazol-2(3H)-
yl)propyl)-N,N-dimethyl-1H-imidazole-1-sulfonamide (70)



```

filename = sm_V_155_pure_ii-4_jd
author = delta
experiment = single_pulse.ex2
sample_id = S#596405
solvent = CHLOROFORM-D
reaction_time = 17-JUL-2009 16:28:05
acquisition_time = 27-MAR-2010 19:32:48
current_time = 27-MAR-2010 19:33:02
comment =
  = single_pulse
  = ID REAL
  = 13107
  = 1H
  = [ppm]
  = X
  = ECX 300
  = DELTA2_NMR
  = DELTA2_NMR
  = 7.0586013[T] (300[MHz]
  = 2.90717696[s]
  = 1H
  = 300.52965592[MHz]
  = 5[ppm]
  = 16384
  = 0
  = 0.34397621[Hz]
  = 5.63570784[kHz]
  = 1H
  = 300.52965592[MHz]
  = 5[ppm]
  = 1H
  = 300.52965592[MHz]
  = 5[ppm]
  = FALSE
  = 1
  = 21
  = 21
  = 13.01[us]
  = 2.90717696[s]
  = 45[deg]
  = 4[dB]
  = 6.505[us]
  = Off
  = Off
  = FALSE
  = 1[s]
  = 46
  = 5[s]
  = 7.90717696[s]
  = 23[dc]
  
```





```

filename = sm_V_32_pure-3.jdf
author = delta
experiment = single_pulse_dec
sample_id = S8725973
solvent = CHLOROFORM-D
acquisition_time = 24-DEC-2008 00:00:03
revision_time = 27-MAR-2010 19:36:10
current_time = 27-MAR-2010 19:36:26

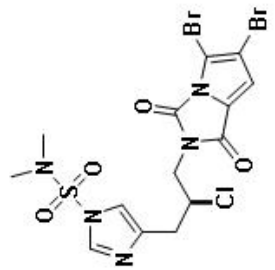
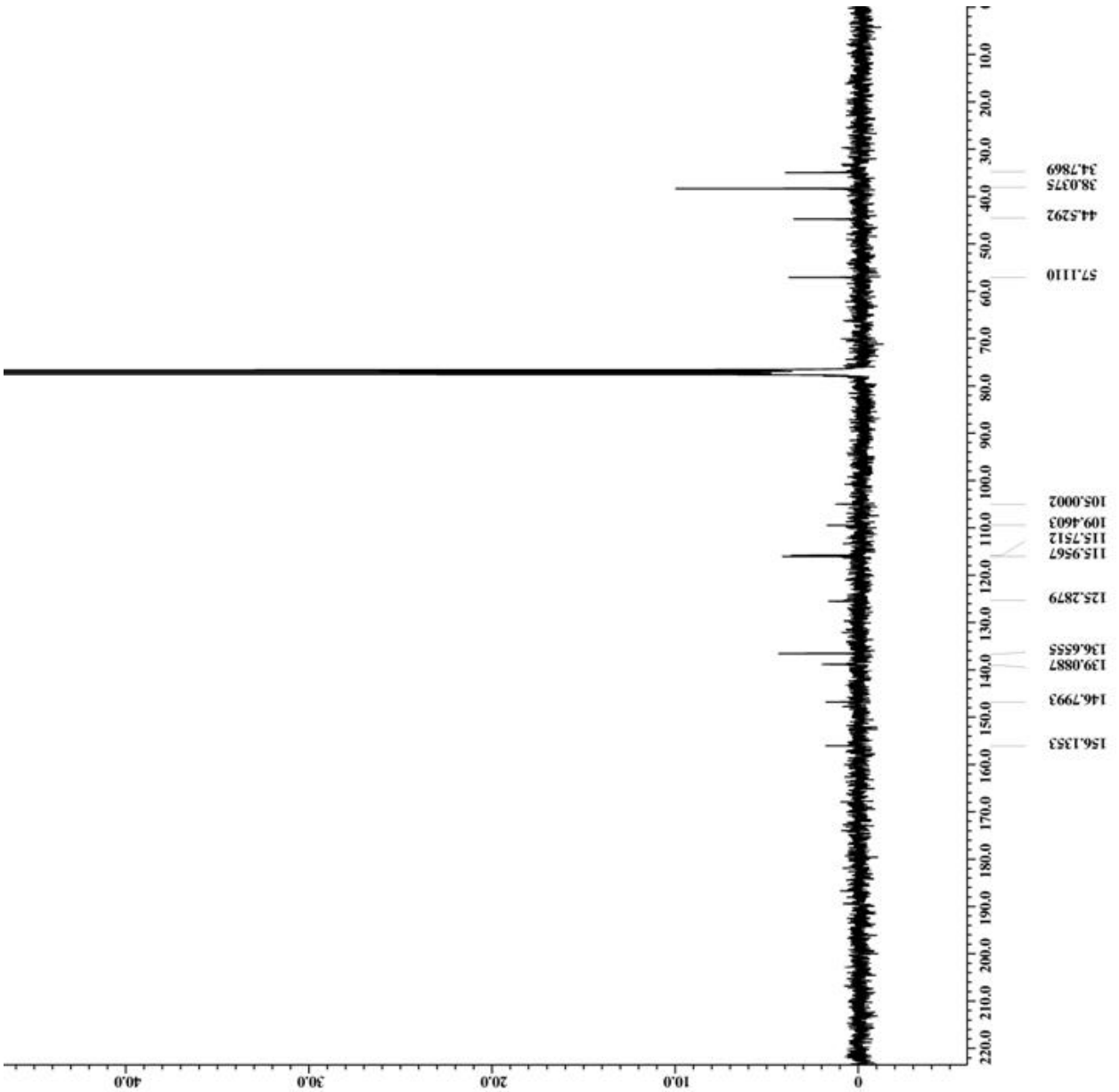
comment = single pulse decouple
ata_format = ID COMPLEX
in_size = 52428
in_title = 13C
in_units = [ppm]
inmensions = X
ite = ECX 300
spectrometer = DELTA2_NMR

ield_strength = 7.0586013[T] (300[MHz]
acq_duration = 2.76824064[s]
domain = 13C
freq = 75.56823426[MHz]
offset = 100[ppm]
points = 65536
prescans = 4
resolution = 0.36124027[Hz]
sweep = 23.67424242[kHz]
tr_domain = 1H
tr_freq = 300.52965592[MHz]
tr_offset = 5[ppm]
lipped = FALSE
bd_return = 10
cans = 2221
otal_scans = 2221

_90_width = 9.75[us]
acq_time = 2.76824064[s]
angle = 30[deg]
atn = 8[db]
pulse = 3.25[us]
tr_atn_dec = 25[db]
tr_atn_noe = 25[db]
tr_noise = WALTZ
scoupling = TRUE
nitial_wait = 1[s]
oe_time = TRUE
oe_time = 3[s]
ecvr_gain = 50
elaxation_delay = 3[s]
petition_time = 5.76824064[s]
emp_get = 23.4[dc]

```

(thousands)





```

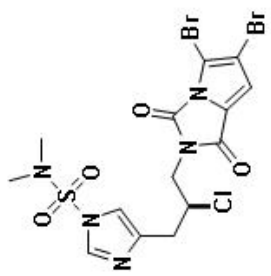
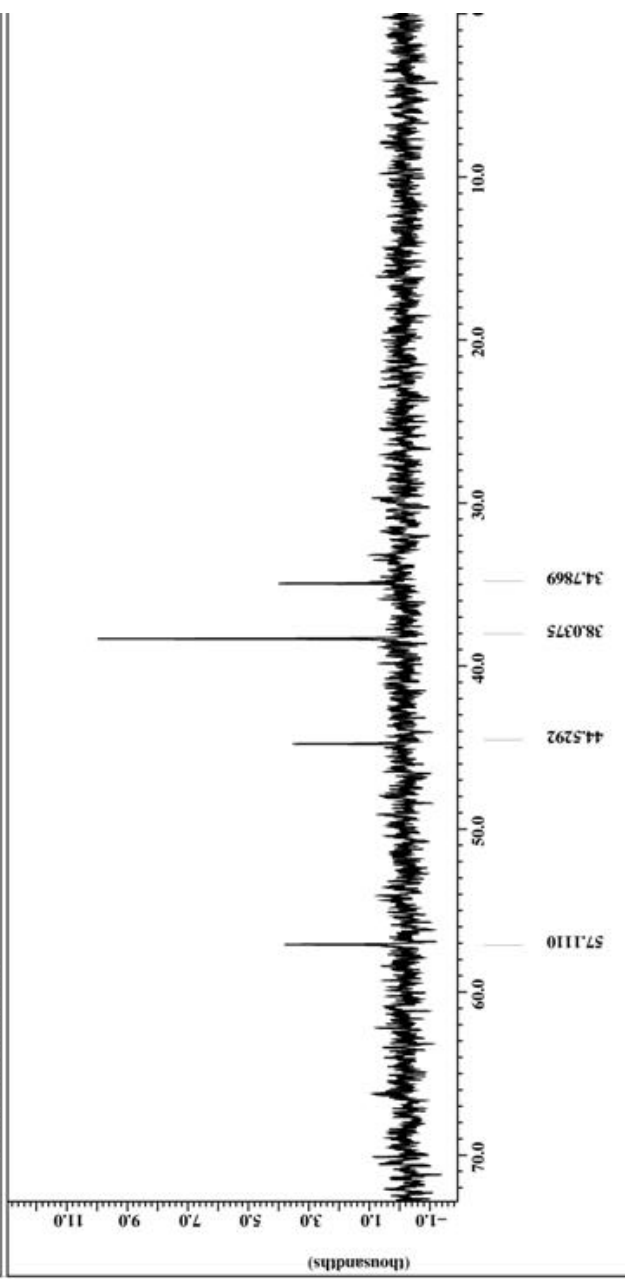
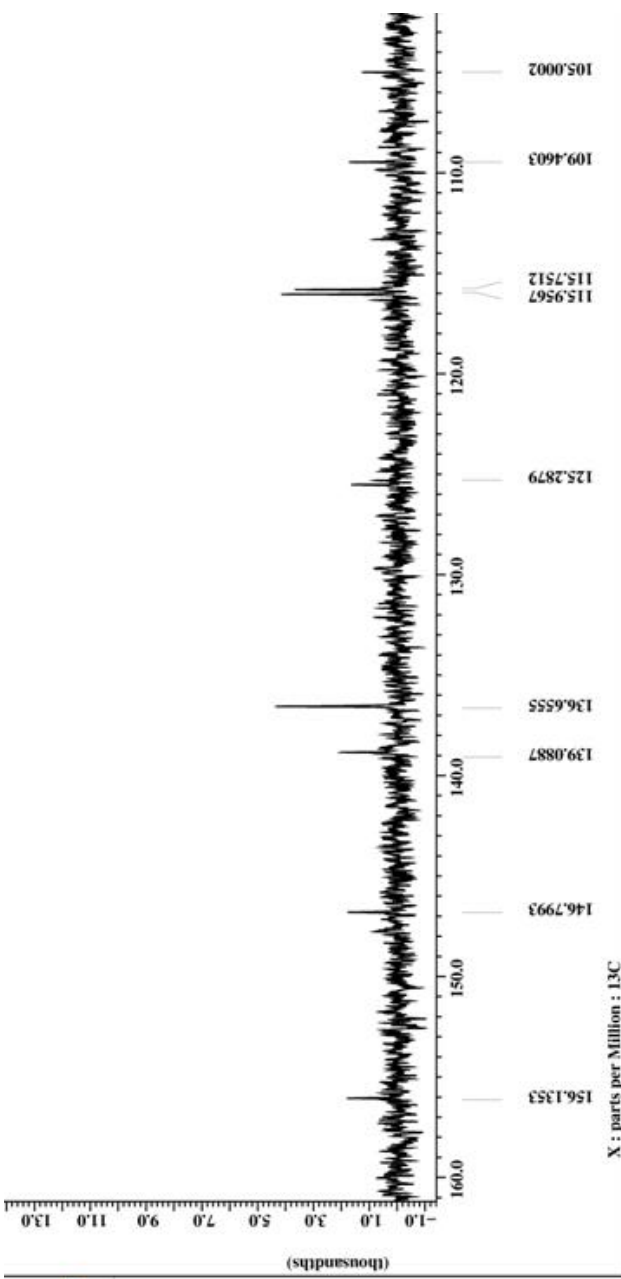
filename = sm_V_32_pure-3.jdf
author = delta
experiment = single_pulse_dec
sample_id = S8725973
solvent = CHLOROFORM-D
reaction_time = 24-DEC-2008 00:00:03
evision_time = 27-MAR-2010 19:36:10
current_time = 27-MAR-2010 19:39:27

comment = single pulse decouple
ata_format = ID COMPLEX
in_size = 52428
in_title = 13C
in_units = [ppm]
inmensions = X
ite = ECX 300
spectrometer = DELTA2_NMR

ield_strength = 7.0586013[T] (300[MHz]
-acq_duration = 2.76824064[s]
-domain = 13C
-freq = 75.56823426[MHz]
-offset = 100[ppm]
-points = 65536
-prescans = 4
-resolution = 0.36124027[Hz]
-sweep = 23.67424242[kHz]
-rr_domain = 1H
-rr_freq = 300.52965592[MHz]
-rr_offset = 5[ppm]
-lipped = FALSE
-bd_return = 10
-cans = 2221
-otal_scans = 2221

_90_width = 9.75[us]
-acq_time = 2.76824064[s]
-angle = 30[deg]
-atn = 8[db]
-pulse = 3.25[us]
-rr_atn_dec = 25[db]
-rr_atn_noe = 25[db]
-rr_noise = WALTZ
-ecoupling = TRUE
-nitial_wait = 1[s]
-be_time = TRUE
-be_time = 3[s]
-ecvr_gain = 50
-relaxation_delay = 3[s]
-periment_time = 5.76824064[s]
-emp_get = 23.4[dc]

```



APPENDIX 16
¹H AND ¹³C NMR SPECTRUM OF
(*R*)-2-chloro-3-(1-(*N,N*-dimethylsulfamoyl)-1*H*-imidazol-4-yl)propyl 1*H*-
pyrrole-2-carboxylate (78)

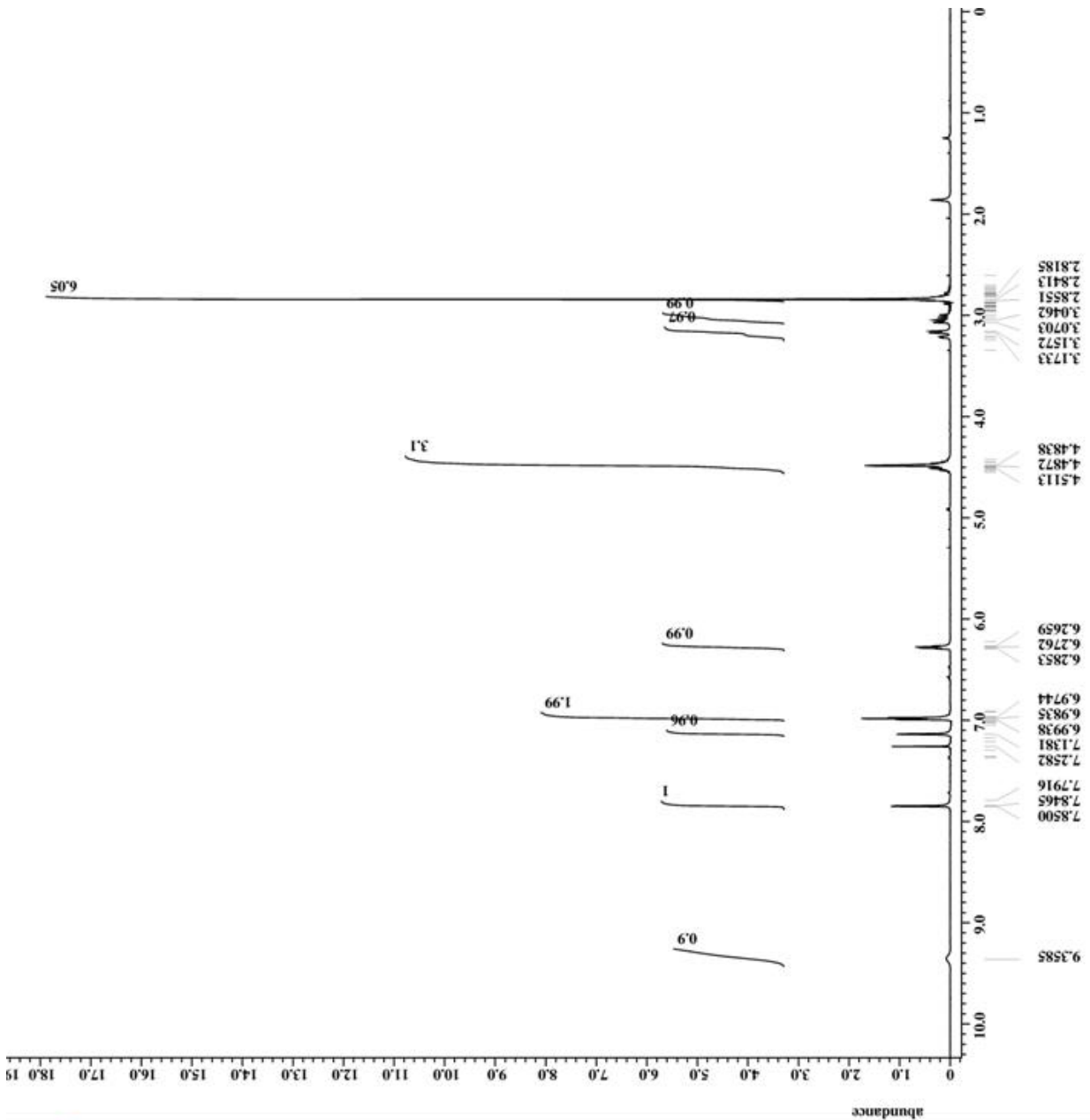
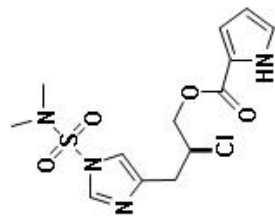


```

filename = sm_V_107_pure_iii-5.j
author = delta
experiment = single_pulse.ex2
sample_id = S835287
solvent = CHLOROFORM-D
creation_time = 27-MAR-2009 22:34:04
revision_time = 27-MAR-2010 19:45:24
current_time = 27-MAR-2010 19:47:25
comment = single_pulse
ata_format = ID REAL
in_size = 13107
in_title = 1H
in_units = [ppm]
in_dimensions = X
ite = ECX 300
spectrometer = DELTA2_NMR
ield_strength = 7.0586013[T] (300[MHz])
_acq_duration = 2.90717696[s]
_domain = 1H
_freq = 300.52965592[MHz]
_offset = 5[ppm]
_points = 16384
_prescans = 0
_resolution = 0.34397621[Hz]
_sweep = 5.63570784[kHz]
rr_domain = 1H
rr_freq = 300.52965592[MHz]
rr_offset = 5[ppm]
ri_domain = 1H
ri_freq = 300.52965592[MHz]
ri_offset = 5[ppm]
lipped = FALSE
od_return = 1
otal_scans = 18

_90_width = 13.01[us]
_acq_time = 2.90717696[s]
_angle = 45[deg]
_atn = 4[db]
_pulse = 6.505[us]
rr_mode = Off
ri_mode = Off
ante_presat = FALSE
ntial_wait = 1[s]
ecvr_gain = 46
elaxation_delay = 5[s]
epetition_time = 7.90717696[s]
emp_get = 23.3[dc]

```



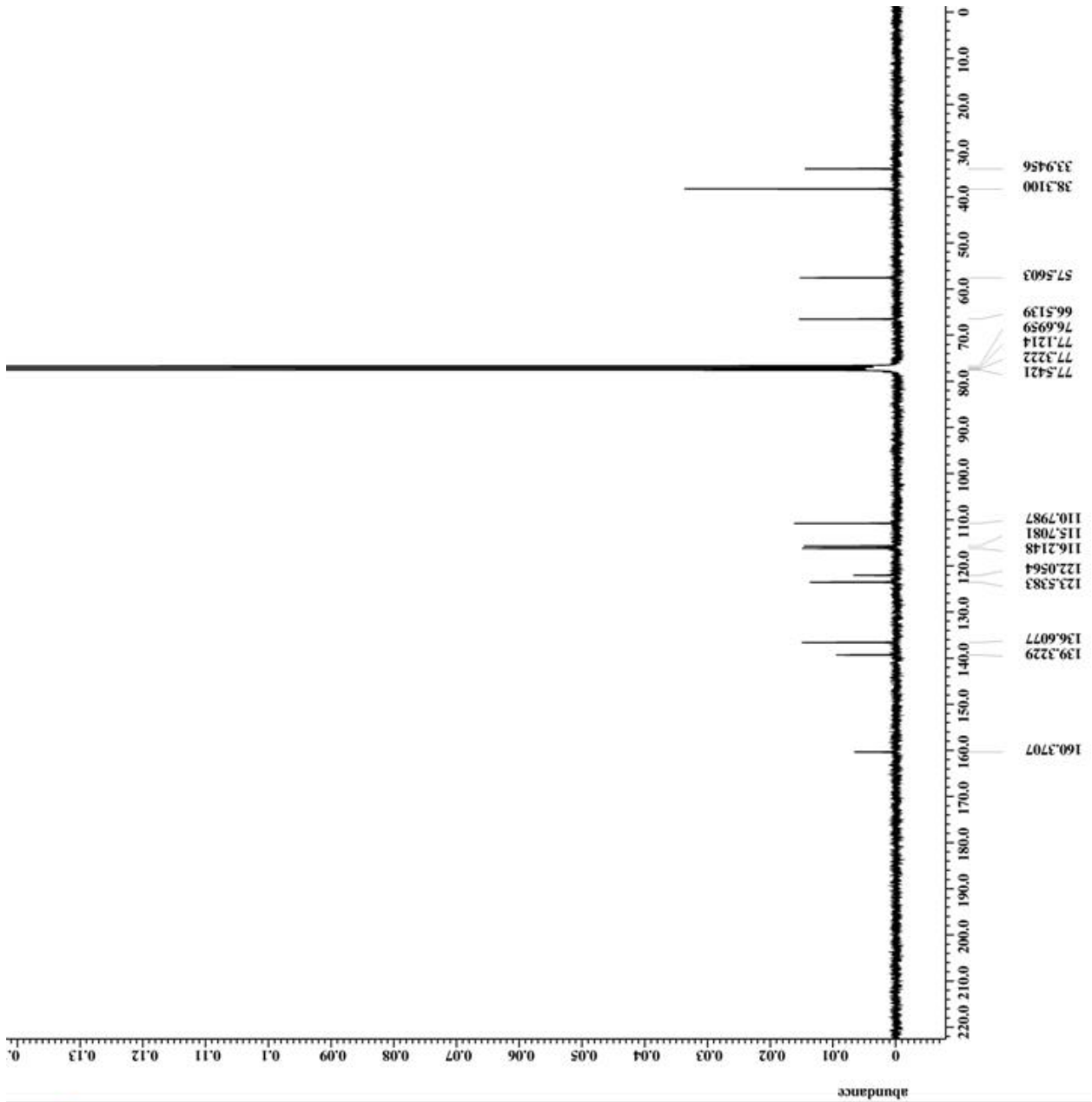
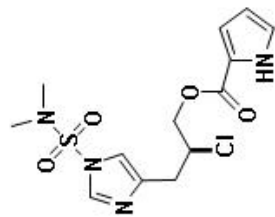


```
filename = sm_v_107_pure_iii-2.j
author = delta
experiment = single_pulse_dec
pulse_id = S837261
solvent = CHLOROFORM-D
reaction_time = 28-MAR-2009 06:35:40
acquisition_time = 28-MAR-2009 14:46:35
current_time = 27-MAR-2010 19:50:10

comment = single pulse decouple
ata_format = ID COMPLEX
in_size = 52428
in_title = 13C
in_units = [ppm]
in_dimensions = X
ite = ECX 300
spectrometer = DELTA2_NMR

field_strength = 7.0586013[T] (300[MHz])
acq_duration = 2.76824064[s]
domain = 13C
freq = 75.56823426[MHz]
offset = 100[ppm]
points = 65536
prescans = 4
resolution = 0.36124027[Hz]
sweep = 23.67424242[kHz]
tr_domain = 1H
tr_freq = 300.52965592[MHz]
tr_offset = 5[ppm]
lipped = FALSE
bd_return = 10
cans = 5000
atal_scans = 5000

_90_width = 9.75[us]
acq_time = 2.76824064[s]
angle = 30[deg]
atn = 8[dB]
pulse = 3.25[us]
tr_atn_dec = 25[dB]
tr_atn_noe = 25[dB]
tr_noise = WALTZ
scoupling = TRUE
nitral_wait = 1[s]
be_time = TRUE
be_time = 3[s]
scvr_gain = 50
relaxation_delay = 3[s]
petition_time = 5.76824064[s]
emp_get = 23.3[dc]
```





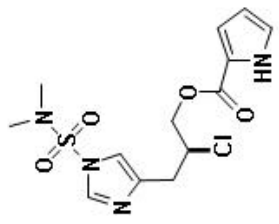
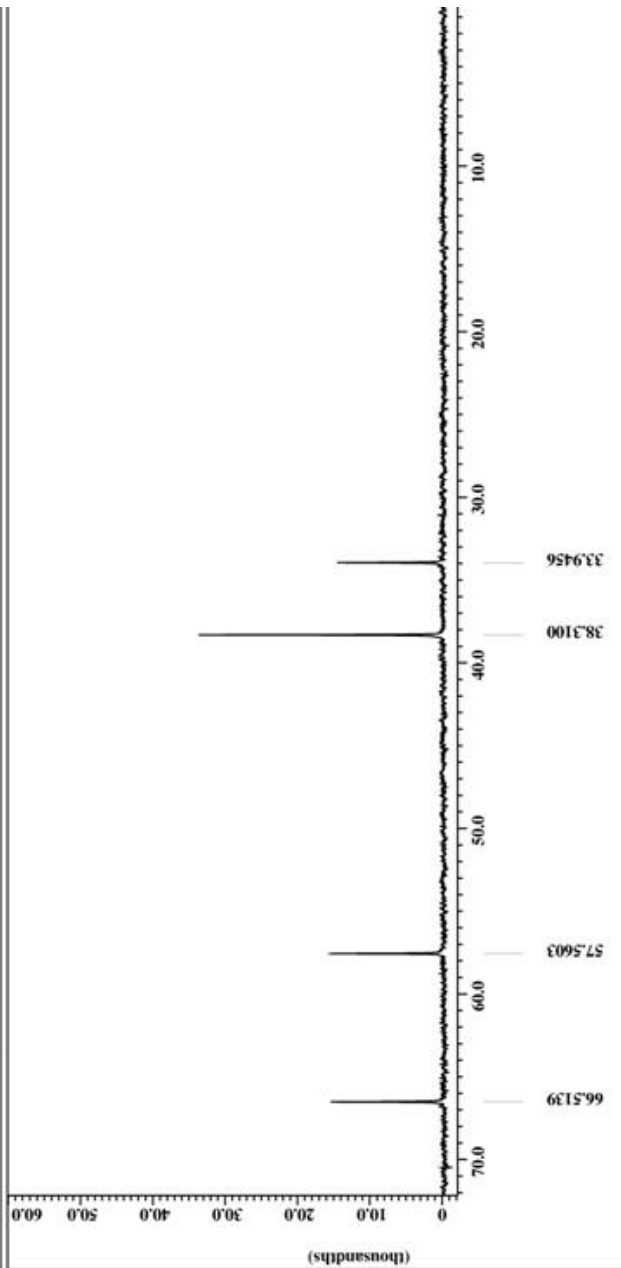
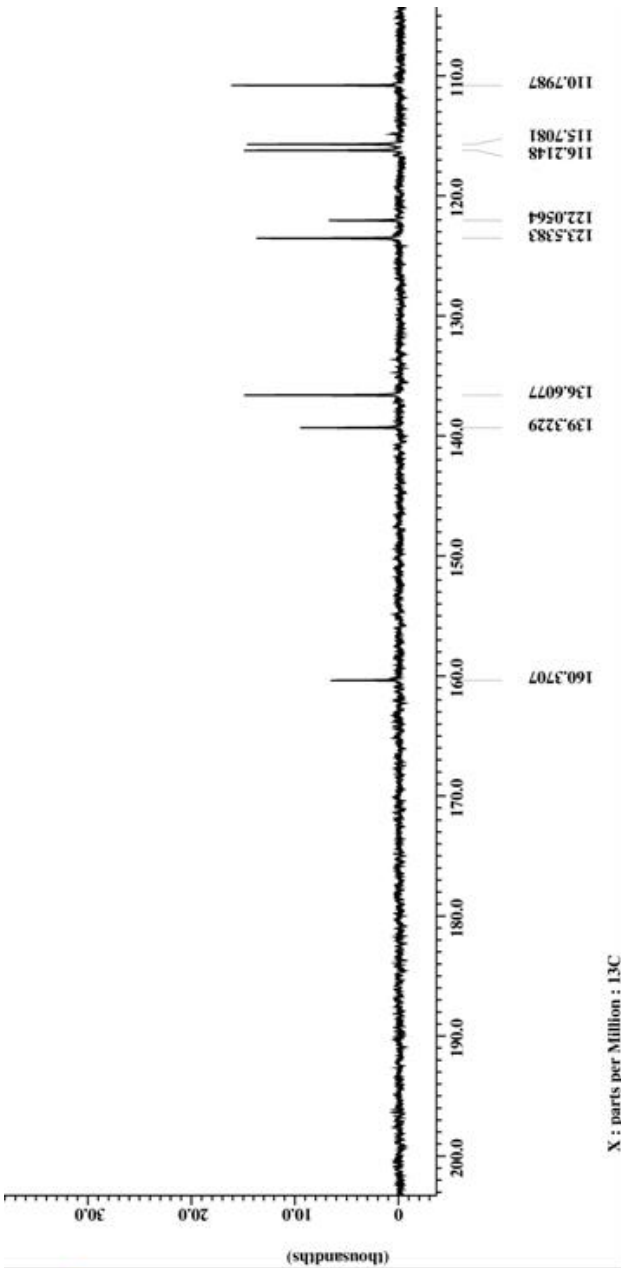
```

filename = sm_V_107_pure_iii-2.j
author = delta
experiment = single_pulse_dec
sample_id = S837261
solvent = CHLOROFORM-D
reaction_time = 28-MAR-2009 06:35:40
revision_time = 28-MAR-2009 14:46:35
current_time = 27-MAR-2010 19:52:05

comment = single pulse decouple
ata_format = ID COMPLEX
in_size = 52428
in_title = 13C
in_units = [ppm]
inmensions = X
ite = ECX 300
spectrometer = DELTA2_NMR

ield_strength = 7.0586013[T] (300 [MHz]
_acq_duration = 2.76824064[s]
_domain = 13C
_freq = 75.56823426[MHz]
_offset = 100[ppm]
_points = 65536
_prescans = 4
_resolution = 0.36124027[Hz]
_sweep = 23.67424242[kHz]
rr_domain = 1H
rr_freq = 300.52965592[MHz]
rr_offset = 5[ppm]
lipped = FALSE
od_return = 10
cans = 5000
atal_scans = 5000

_90_width = 9.75[us]
_acq_time = 2.76824064[s]
_angle = 30[deg]
_atn = 8[db]
_pulse = 3.25[us]
rr_atn_dec = 25[db]
rr_atn_noe = 25[db]
rr_noise = WALTZ
scoupling = TRUE
nitial_wait = 1[s]
oe_time = TRUE
oe_time = 3[s]
scvr_gain = 50
relaxation_delay = 3[s]
petition_time = 5.76824064[s]
emp_get = 23.3[dc]
  
```



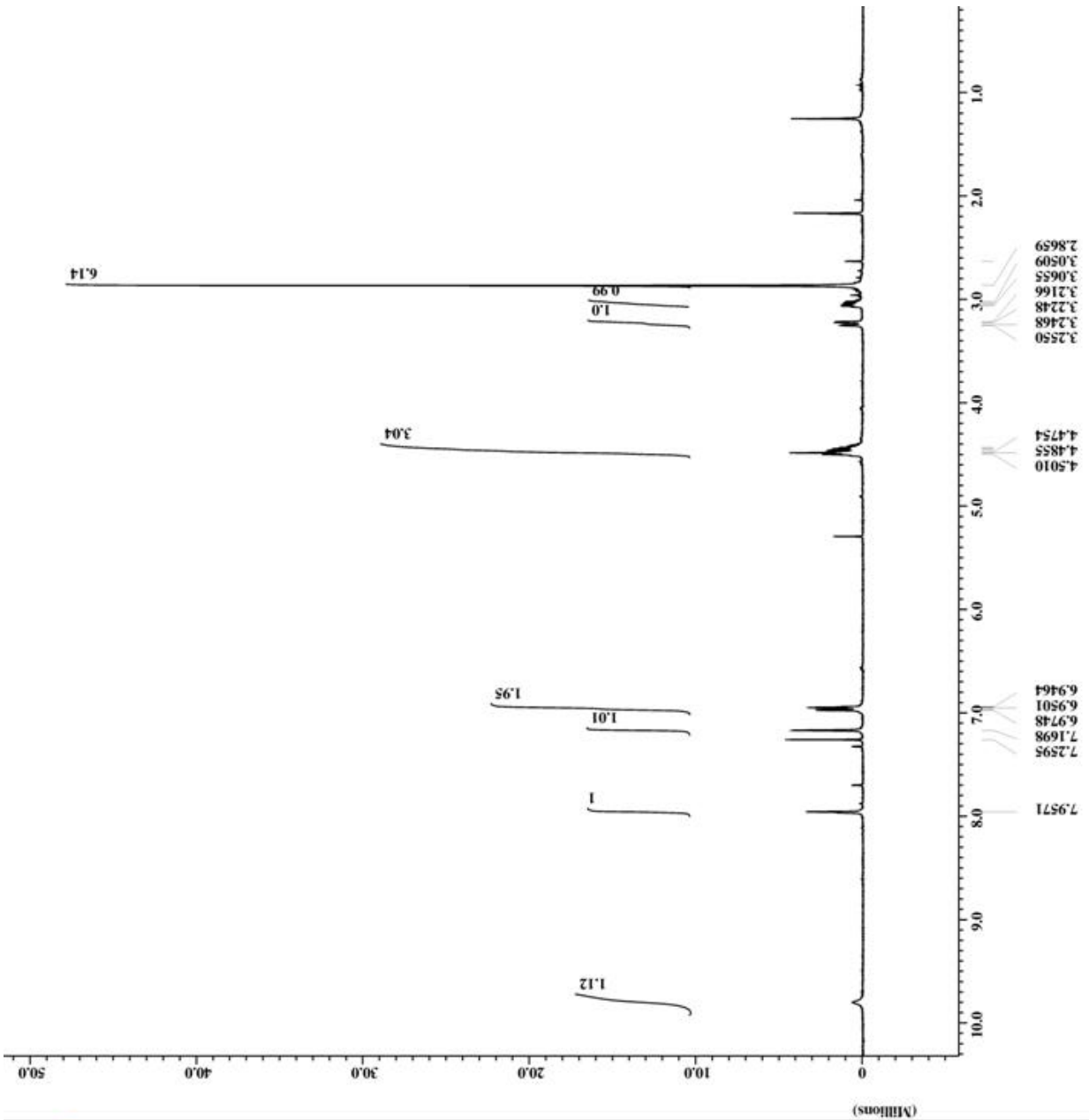
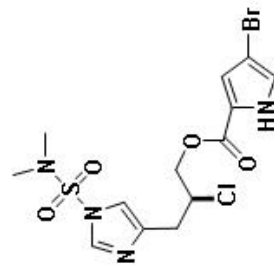
APPENDIX 17
¹H AND ¹³C NMR SPECTRUM OF
(*R*)-2-chloro-3-(1-(*N,N*-dimethylsulfamoyl)-1H-imidazol-4-yl)propyl 4-bromo-1H-pyrrole-2-carboxylate (85)



```

filename = sm_VI_49_pure-3.jdf
author = delta
experiment = single_pulse_exp
sample_id = S#855015
solvent = CHLOROFORM-D
reaction_time = 11-DEC-2009 15:17:04
revision_time = 27-MAR-2010 19:57:11
current_time = 27-MAR-2010 19:57:28
comment = Single Pulse Experiment
data_format = ID COMPLEX
in_size = 16384
in_title = 1H
in_units = [ppm]
inmensions = X
ite = Eclipse+ 500
spectrometer = DELTA_NMR
field_strength = 11.7465928[T] (500[MH
acq_duration = 2.1839872[s]
domain = 1H
freq = 500.12734003[MHz]
offset = 5[ppm]
points = 16384
prescans = 0
resolution = 0.45787814[Hz]
sweep = 7.50187547[kHz]
tipped = FALSE
od_return = 1
total_scans = 8
_90_width = 18.5[us]
acq_time = 2.1839872[s]
angle = 45[deg]
pulse = 7.25[us]
nitel_wait = 1[s]
base_preset = 2[us]
ecvr_gain = 20
relaxation_delay = 4[s]
emp_get = 25.8[dc]
nblank_time = 2[us]

```





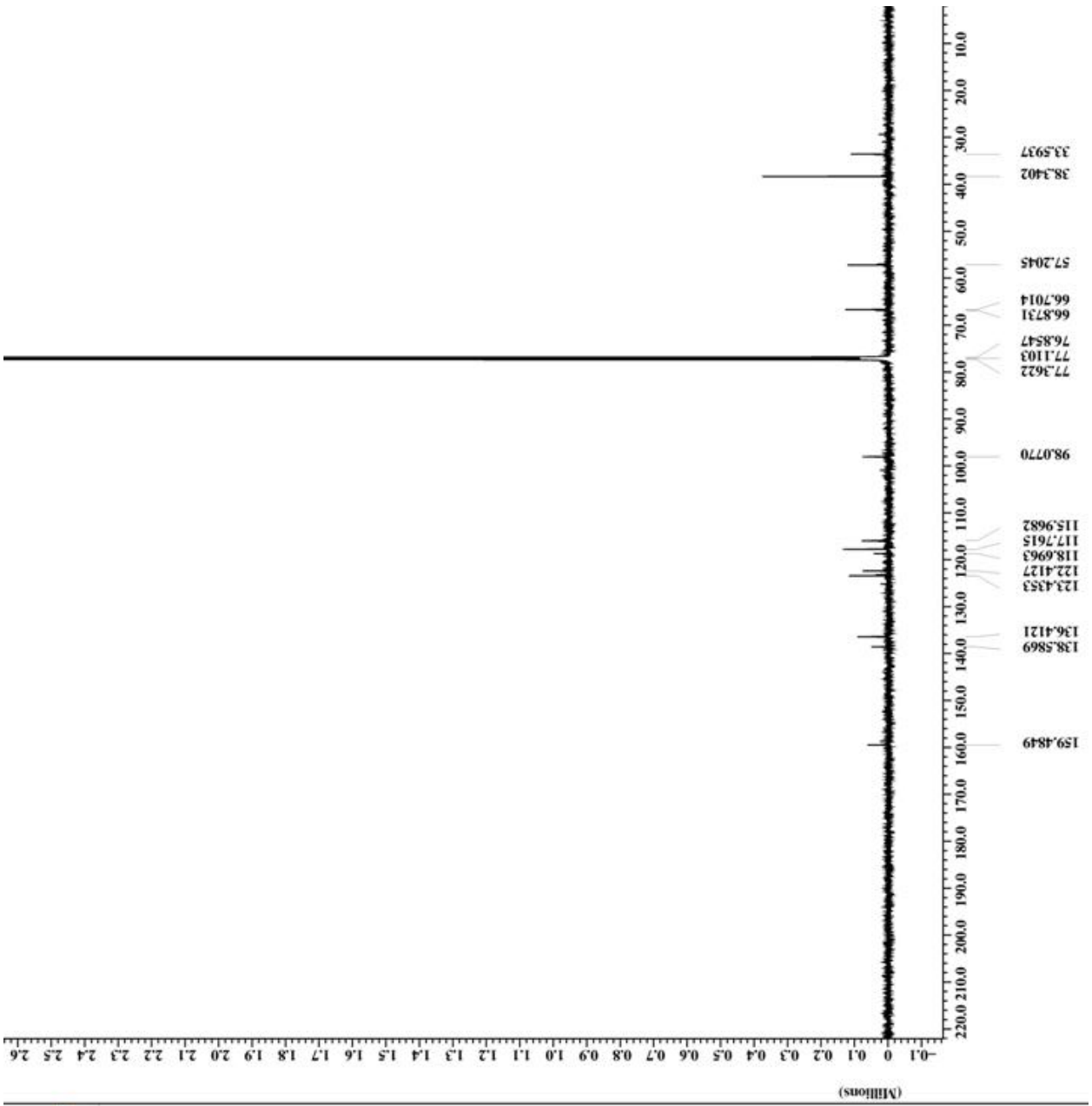
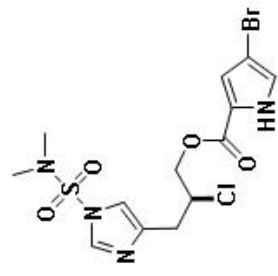
```

filename = sm_VI_49_pure-3.jdf
author = delta
experiment = single_pulse_dec
sample_id = S#856074
solvent = CHLOROFORM-D
acquisition_time = 11-DEC-2009 22:22:59
revision_time = 27-MAR-2010 20:00:11
current_time = 27-MAR-2010 20:00:24

comment = single pulse decouple
ata_format = ID COMPLEX
im_size = 65536
im_title = 13C
im_units = [ppm]
imensions = X
ite = Eclipse+ 500
spectrometer = DELTA_NMR

field_strength = 11.7465928[T] (500[MH
-acq_duration = 2.0840448[s]
-domain = 13C
-freq = 125.75710665[MHz]
-offset = 100[ppm]
-points = 65536
-prescans = 4
-resolution = 0.47983613[Hz]
-sweep = 31.44654088[kHz]
-tr_domain = 1H
-tr_freq = 500.12734003[MHz]
-tr_offset = 5[ppm]
-tipped = TRUE
-bd_return = 10
-cans = 5000
-etal_scans = 5000

_90_width = 14.2[us]
-acq_time = 2.0840448[s]
-angle = 30[deg]
-pulse = 4.73333333[us]
-nitral_wait = 1[s]
-pe_time = 1[s]
-base_preset = 3[us]
-ecvr_gain = 30
-relaxation_delay = 2[s]
-emp_get = 28.3[dC]
-nblank_time = 2[us]
  
```



APPENDIX 18
¹H AND ¹³C NMR SPECTRUM OF
(*R*)-2-chloro-3-(1-(*N,N*-dimethylsulfamoyl)-1*H*-imidazol-4-yl)propyl 4,5-
dibromo-1*H*-pyrrole-2-carboxylate (86)

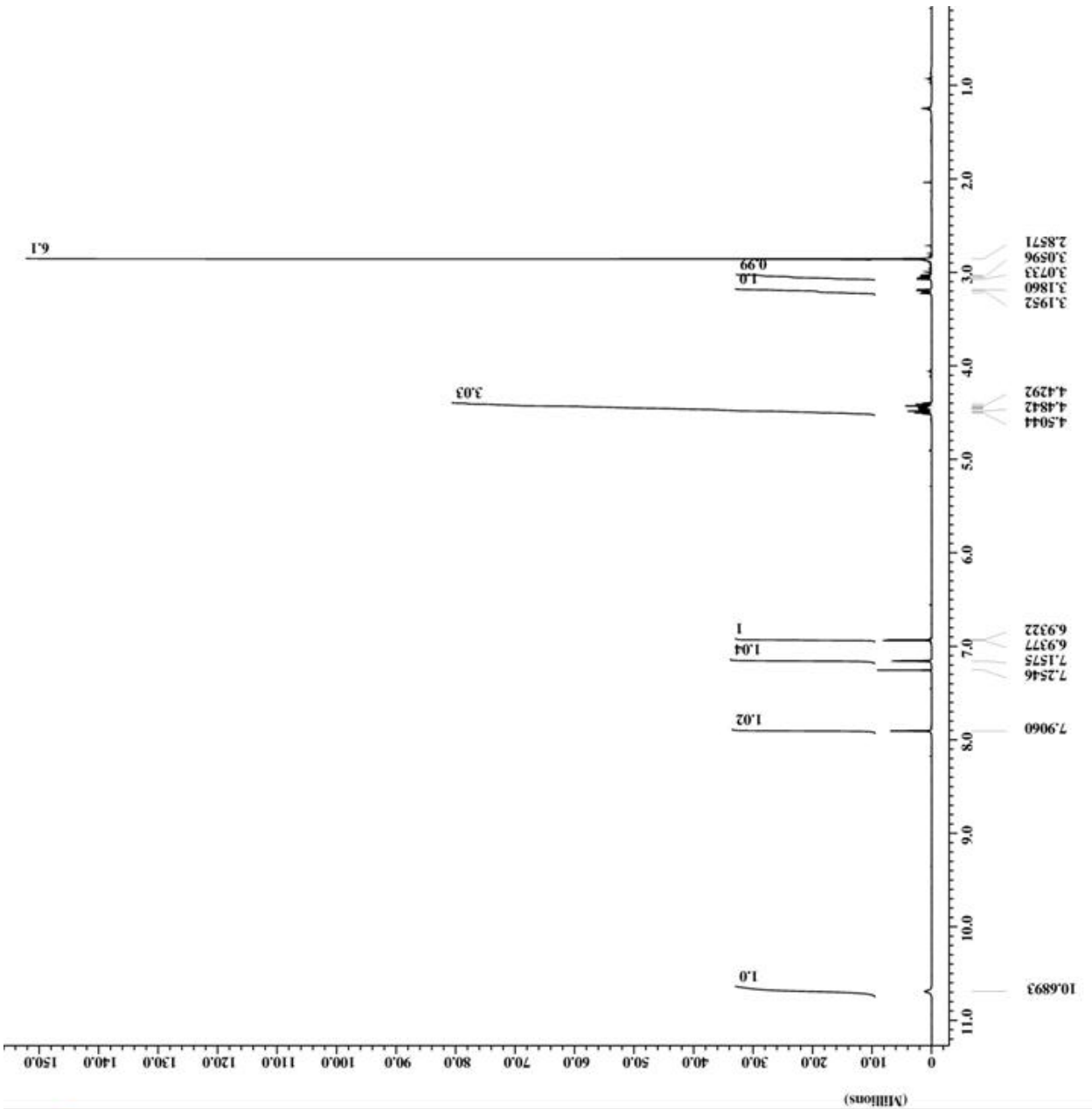
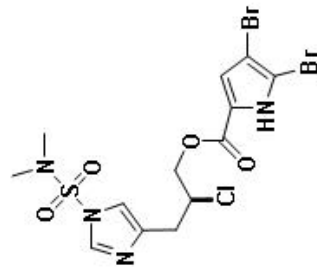


```

filename = sm_V_179_pure-4.jdf
author = delta
experiment = single_pulse_exp
sample_id = S#571802
solvent = CHLOROFORM-D
reaction_time = 8-AUG-2009 05:59:14
evision_time = 27-MAR-2010 20:12:14
current_time = 27-MAR-2010 20:13:49
comment = Single Pulse Experiment
ata_format = ID REAL
in_size = 16384
in_title = 1H
in_units = [ppm]
inmensions = X
ite = Eclipse+ 500
spectrometer = DELTA_NMR
field_strength = 11.7473579[T] (500[MH
-acq_duration = 2.1823486[s]
-domain = 1H
-freq = 500.15991521[MHz]
-offset = 5[ppm]
-points = 16384
-prescans = 0
-resolution = 0.45822189[Hz]
-sweep = 7.50750751[kHz]
lipped = FALSE
od_return = 1
otal_scans = 8

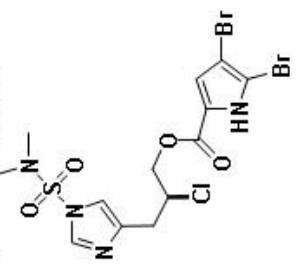
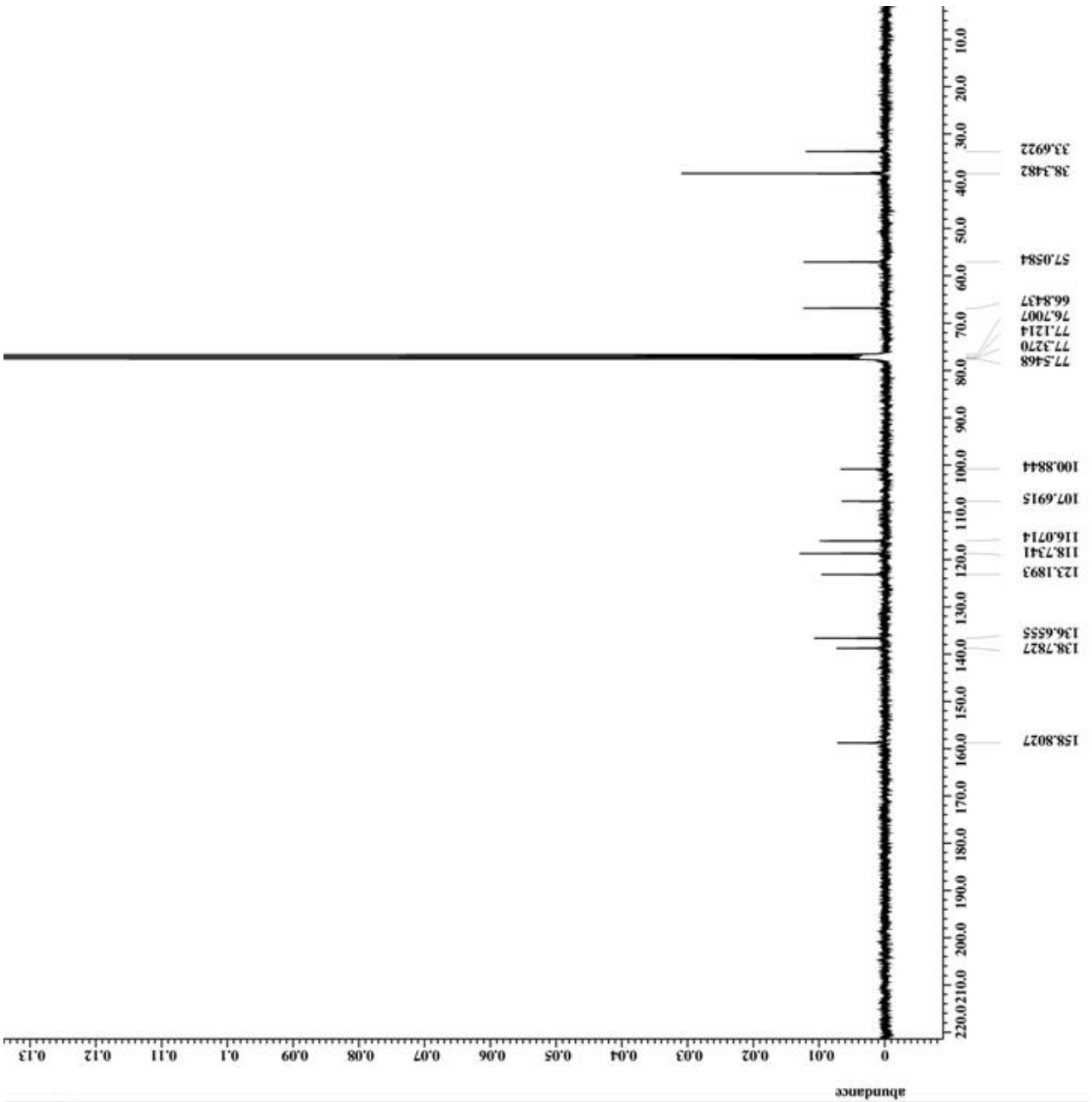
_90_width = 18.5[us]
-acq_time = 2.1823486[s]
-angle = 45[deg]
-pulse = 7.25[us]
ntial_wait = 1[s]
hase_preset = 2[us]
ecvr_gain = 21
elaxation_delay = 4[s]
emp_get = 27[dc]
nblank_time = 2[us]

```





```
filename = sm_V_179_pure_ii-2.jd
author = delta
experiment = single_pulse_dec
sample_id = S#553989
solvent = CHLOROFORM-D
acquisition_time = 9-AUG-2009 20:06:16
revision_time = 9-AUG-2009 23:00:22
current_time = 27-MAR-2010 20:16:48
comment = single pulse decouple
ata_format = ID COMPLEX
in_size = 52428
in_title = 13C
in_units = [ppm]
in_dimensions = X
ite = ECX 300
spectrometer = DELTA2_NMR
ield_strength = 7.0586013[T] (300[MHz]
_acq_duration = 2.76824064[s]
_domain = 13C
_freq = 75.56823426[MHz]
_offset = 100[ppm]
_points = 65536
_prescans = 4
_resolution = 0.36124027[Hz]
_sweep = 23.67424242[kHz]
tr_domain = 1H
tr_freq = 300.52965592[MHz]
tr_offset = 5[ppm]
lipped = FALSE
bd_return = 10
cans = 3000
atal_scans = 3000
_90_width = 9.75[us]
_acq_time = 2.76824064[s]
_angle = 30[deg]
_atn = 8[db]
_pulse = 3.25[us]
tr_atn_dec = 25[db]
tr_atn_noe = 25[db]
tr_noise = WALTZ
coupling = TRUE
initial_wait = 1[s]
be_time = TRUE
be = 3[s]
scvr_gain = 50
relaxation_delay = 3[s]
petition_time = 5.76824064[s]
emp_get = 22.9[dc]
```



APPENDIX 19
¹H AND ¹³C NMR SPECTRUM OF
(S)-N,N-dimethyl-4-((1-oxo-3,4-dihydro-1H-pyrrolo[2,1-c][1,4]oxazin-4-yl)methyl)-1H-imidazole-1-sulfonamide (77)


```

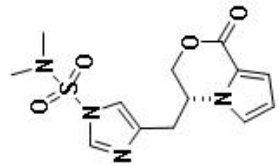
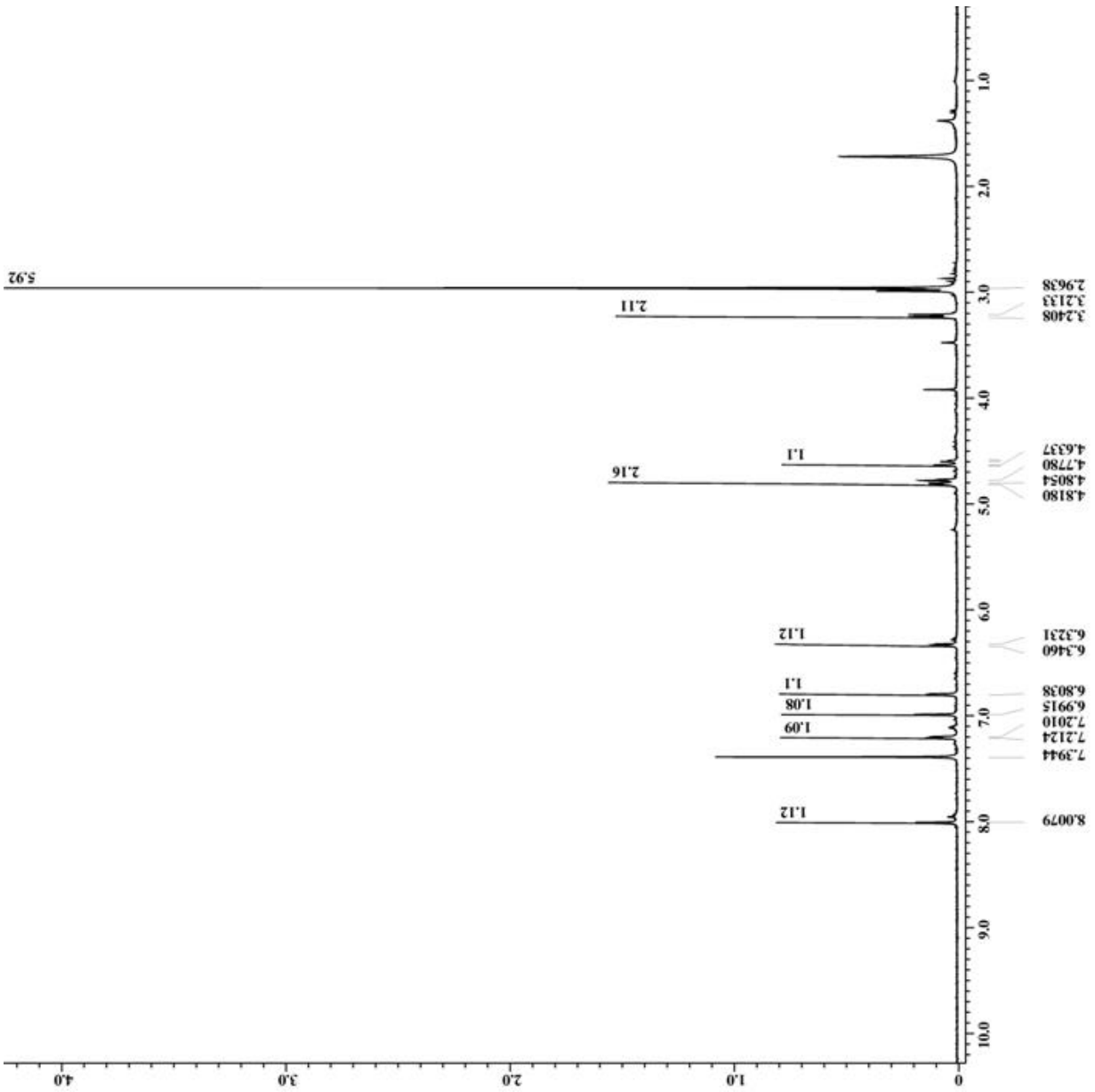
filename = sm_V_165_pure-3_jdf
author = delta
experiment = single_pulse.ex2
sample_id = S#508172
solvent = CHLOROFORM-D
reaction_time = 12-JUL-2009 14:00:37
acquisition_time = 27-MAR-2010 20:30:16
current_time = 27-MAR-2010 20:32:44

comment = single_pulse
ata_format = ID COMPLEX
in_size = 13107
in_title = 1H
in_units = [ppm]
in_dimensions = X
ite = ECX 300
spectrometer = DELTA2_NMR

field_strength = 7.0586013[T] (300[MHz])
acq_duration = 2.90717696[s]
domain = 1H
freq = 300.52965592[MHz]
offset = 5[ppm]
points = 16384
prescans = 0
resolution = 0.34397621[Hz]
sweep = 5.63570784[kHz]
tr_domain = 1H
tr_freq = 300.52965592[MHz]
tr_offset = 5[ppm]
ri_domain = 1H
ri_freq = 300.52965592[MHz]
ri_offset = 5[ppm]
lipped = FALSE
od_return = 1
otal_scans = 24

_90_width = 13.01[us]
acq_time = 2.90717696[s]
angle = 45[deg]
atn = 4[dB]
pulse = 6.505[us]
tr_mode = Off
ri_mode = Off
ante_presat = FALSE
initial_wait = 1[s]
scvr_gain = 46
relaxation_delay = 5[s]
petition_time = 7.90717696[s]
temp_get = 23[degC]
  
```

abundance



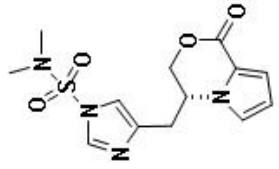
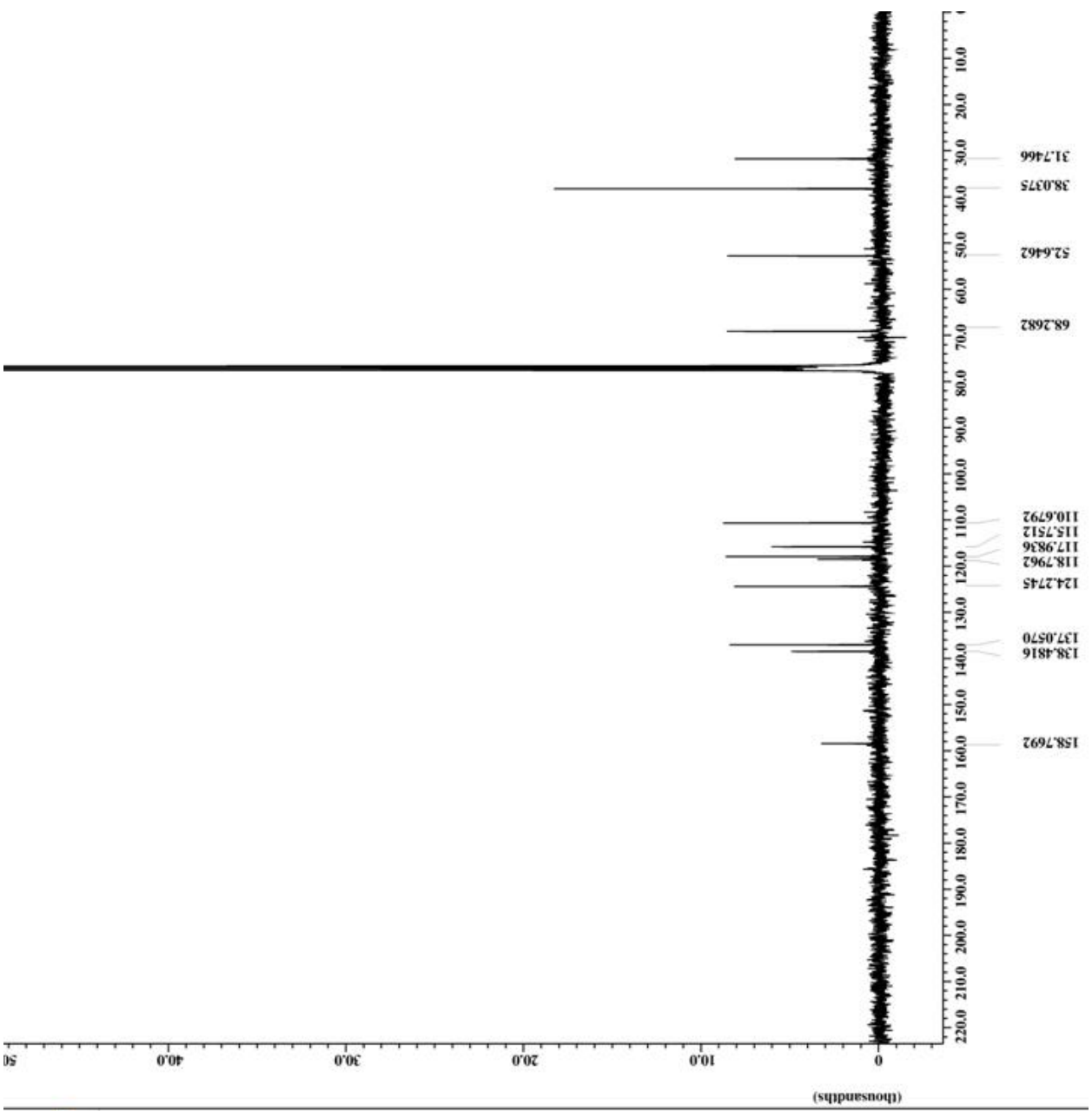


```
filename = sm_V_108_pure-2.jdf
author = delta
experiment = single_pulse_dec
sample_id = S882009
solvent = CHLOROFORM-D
acquisition_time = 3-APR-2009 03:32:08
revision_time = 27-MAR-2010 20:36:38
current_time = 27-MAR-2010 20:36:49

comment = single pulse decouple
ata_format = ID COMPLEX
in_size = 52428
in_title = 13C
in_units = [ppm]
in_dimensions = X
ite = ECX 300
spectrometer = DELTA2_NMR

ield_strength = 7.0586013[T] (300[MHz]
_acq_duration = 2.76824064[s]
_domain = 13C
_freq = 75.56823426[MHz]
_offset = 100[ppm]
_points = 65536
_prescans = 4
_resolution = 0.36124027[Hz]
_sweep = 23.67424242[kHz]
tr_domain = 1H
tr_freq = 300.52965592[MHz]
tr_offset = 5[ppm]
lipped = FALSE
bd_return = 10
cans = 3000
atal_scans = 3000

_90_width = 9.75[us]
_acq_time = 2.76824064[s]
_angle = 30[deg]
_atn = 8[db]
_pulse = 3.25[us]
tr_atn_dec = 25[db]
tr_atn_noe = 25[db]
tr_noise = WALTZ
scoupling = TRUE
nitral_wait = 1[s]
oe_time = TRUE
oe_time = 3[s]
ecvr_gain = 50
relaxation_delay = 3[s]
epetition_time = 5.76824064[s]
emp_get = 23.4[dc]
```



APPENDIX 20
¹H AND ¹³C NMR SPECTRUM OF
(S)-4-((7-bromo-1-oxo-3,4-dihydro-1H-pyrrolo[2,1-c][1,4]oxazin-4-yl)methyl)-
N,N-dimethyl-1H-imidazole-1-sulfonamide (87)



```

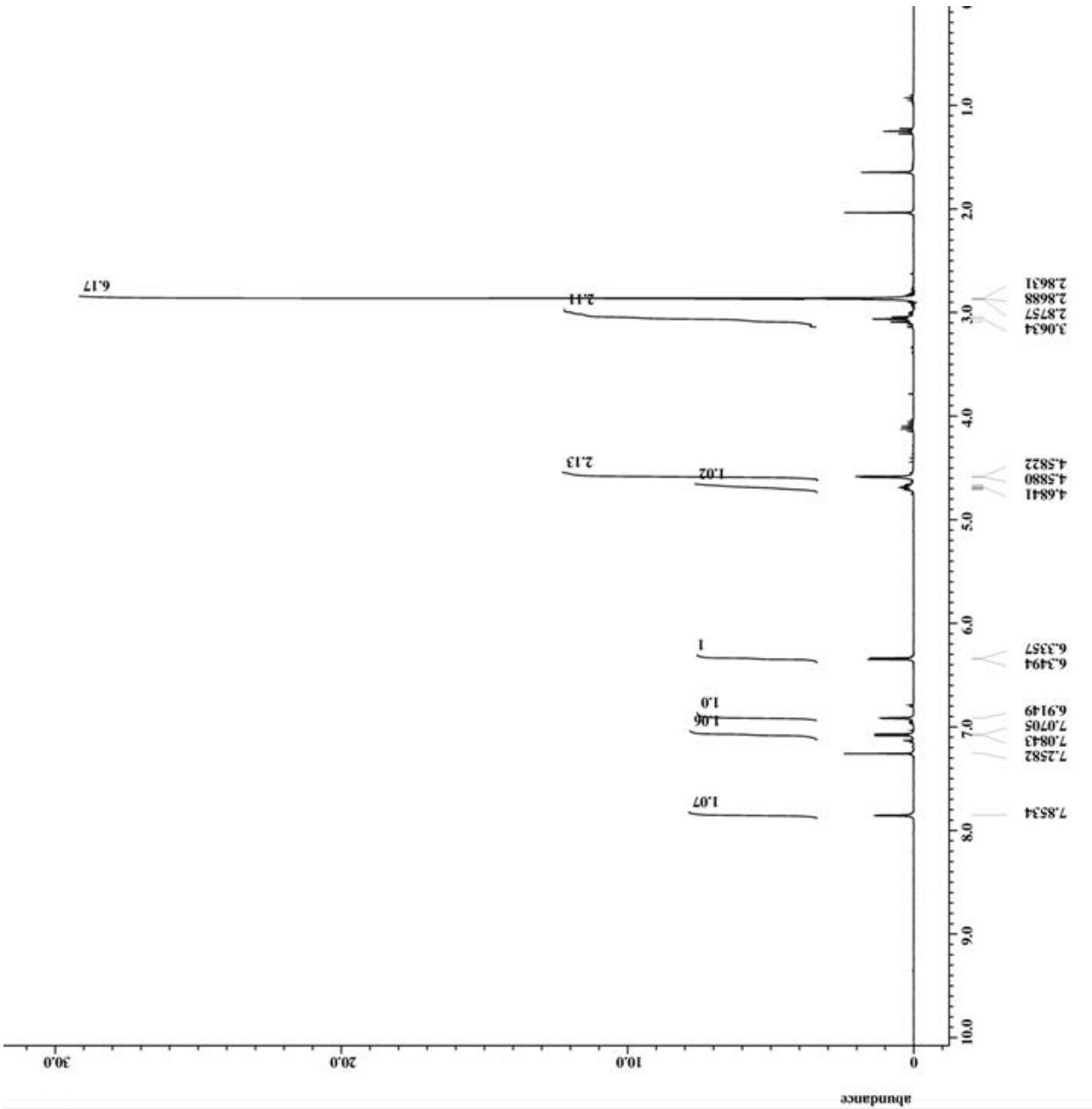
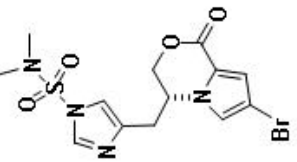
filename = sm_VI_50_pure-2.jdf
author = delta
experiment = single_pulse.ex2
sample_id = S#791896
solvent = CHLOROFORM-D
reaction_time = 2-DEC-2009 22:09:25
revision_time = 2-DEC-2009 22:06:02
current_time = 27-MAR-2010 20:41:38

comment = single_pulse
ata_format = ID COMPLEX
in_size = 13107
in_title = 1H
in_units = [ppm]
inmensions = X
ite = ECX 300
spectrometer = DELTA2_NMR

ield_strength = 7.0586013[T] (300[MHz]
_acq_duration = 2.90717696[s]
_domain = 1H
_freq = 300.52965592[MHz]
_offset = 5[ppm]
_points = 16384
_prescans = 0
_resolution = 0.34397621[Hz]
_sweep = 5.63570784[kHz]
tr_domain = 1H
tr_freq = 300.52965592[MHz]
tr_offset = 5[ppm]
ri_domain = 1H
ri_freq = 300.52965592[MHz]
ri_offset = 5[ppm]
lipped = FALSE
od_return = 1
otal_scans = 24

_90_width = 13.01[us]
_acq_time = 2.90717696[s]
_angle = 45[deg]
_atn = 4[dB]
_pulse = 6.505[us]
tr_mode = Off
ri_mode = Off
ante_presat = FALSE
initial_wait = 1[s]
scvr_gain = 50
relaxation_delay = 5[s]
petition_time = 7.90717696[s]
emp_get = 23.2[deg]

```





```

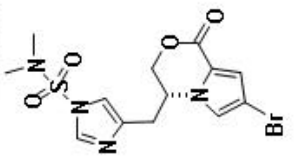
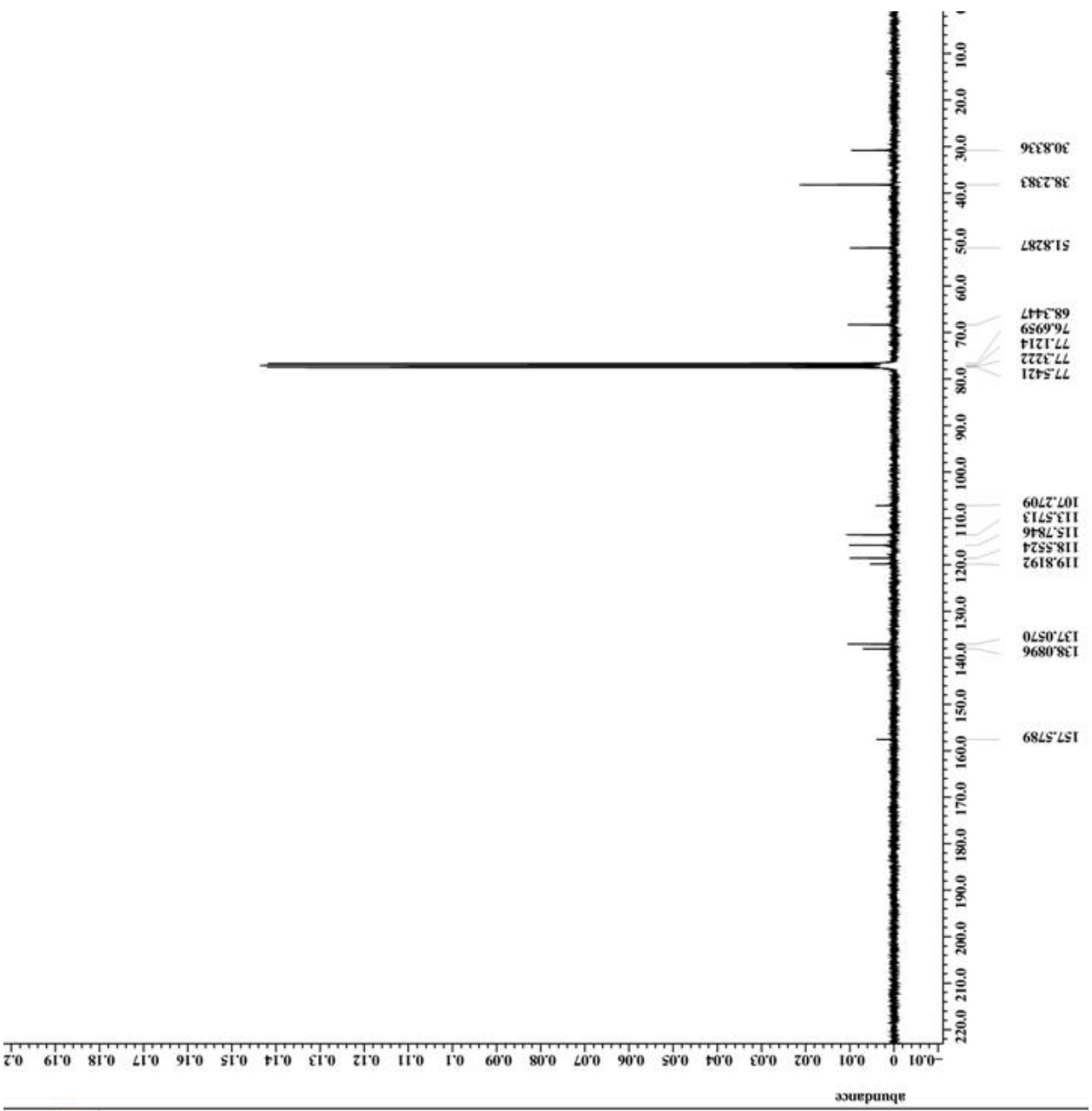
filename = sm_VI_50_pure-2.jdf
author =
experiment = single_pulse_dec
sample_id = S#796170
solvent = CHLOROFORM-D
reaction_time = 3-DEC-2009 06:14:17
revision_time = 3-DEC-2009 15:55:51
current_time = 27-MAR-2010 20:44:53

comment = single pulse decouple
ata_format = ID COMPLEX
in_size = 52428
in_title = 13C
in_units = [ppm]
inmensions = X
ite = ECX 300
spectrometer = DELTA2_NMR

ield_strength = 7.0586013[T] (300[MHz]
_acq_duration = 2.76824064[s]
_domain = 13C
_freq = 75.56823426[MHz]
_offset = 100[ppm]
_points = 65536
_prescans = 4
_resolution = 0.36124027[Hz]
_sweep = 23.67424242[kHz]
tr_domain = 1H
tr_freq = 300.52965592[MHz]
tr_offset = 5[ppm]
lipped = FALSE
bd_return = 10
cans = 5000
atal_scans = 5000

_90_width = 9.75[us]
_acq_time = 2.76824064[s]
_angle = 30[deg]
_atn = 8[db]
_pulse = 3.25[us]
tr_atn_dec = 25[db]
tr_atn_noe = 25[db]
tr_noise = WALTZ
coupling = TRUE
nitial_wait = 1[s]
oe_time = TRUE
oe = 3[s]
scvr_gain = 50
relaxation_delay = 3[s]
petition_time = 5.76824064[s]
emp_get = 23.1[dc]

```



APPENDIX 21
¹H AND ¹³C NMR SPECTRUM OF
(S)-4-((6,7-dibromo-1-oxo-3,4-dihydro-1H-pyrrolo[2,1-c][1,4]oxazin-4-yl)methyl)-N,N-dimethyl-1H-imidazole-1-sulfonamide (88)



```

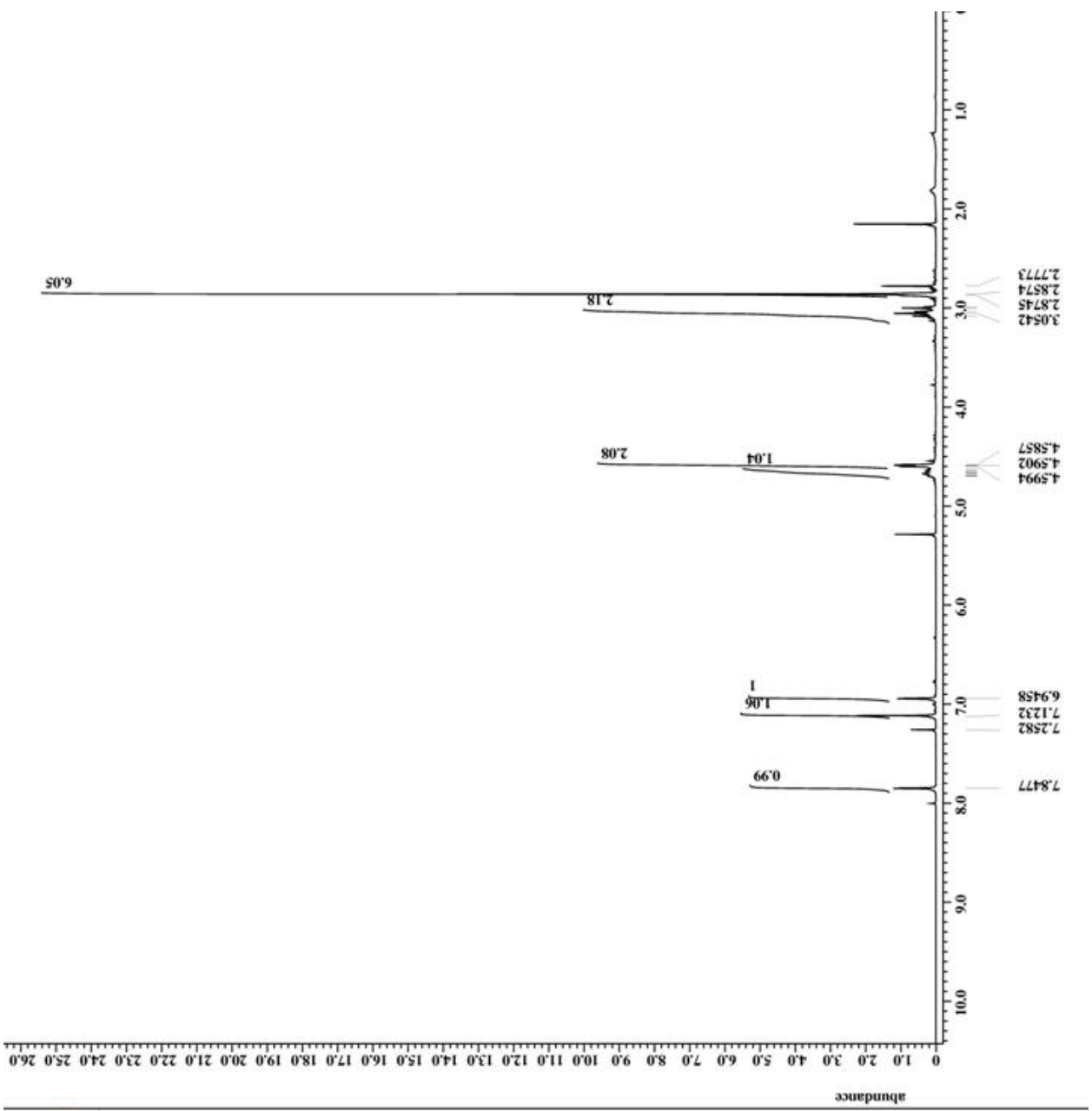
filename = sm_VI_47_pure-3_jdf
author = delta
experiment = single_pulse.ex2
sample_id = S833575
solvent = CHLOROFORM-D
reaction_time = 1-DEC-2009 23:19:18
acquisition_time = 27-MAR-2010 20:48:49
current_time = 27-MAR-2010 20:49:20

comment = single_pulse
ata_format = ID COMPLEX
in_size = 13107
in_title = 1H
in_units = [ppm]
in_dimensions = X
ite = ECX 300
spectrometer = DELTA2_NMR

ield_strength = 7.0586013[T] (300[MHz])
_acq_duration = 2.90717696[s]
_domain = 1H
_freq = 300.52965592[MHz]
_offset = 5[ppm]
_points = 16384
_prescans = 0
_resolution = 0.34397621[Hz]
_sweep = 5.63570784[kHz]
tr_domain = 1H
tr_freq = 300.52965592[MHz]
tr_offset = 5[ppm]
ri_domain = 1H
ri_freq = 300.52965592[MHz]
ri_offset = 5[ppm]
lipped = FALSE
od_return = 1
otal_scans = 24

_90_width = 13.01[us]
_acq_time = 2.90717696[s]
_angle = 45[deg]
_atn = 4[dB]
_pulse = 6.505[us]
tr_mode = Off
ri_mode = Off
ante_presat = FALSE
nitial_wait = 1[s]
ecvr_gain = 42
relaxation_delay = 5[s]
petition_time = 7.90717696[s]
emp_get = 23.2[deg]

```



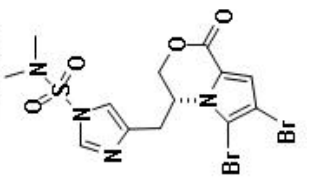
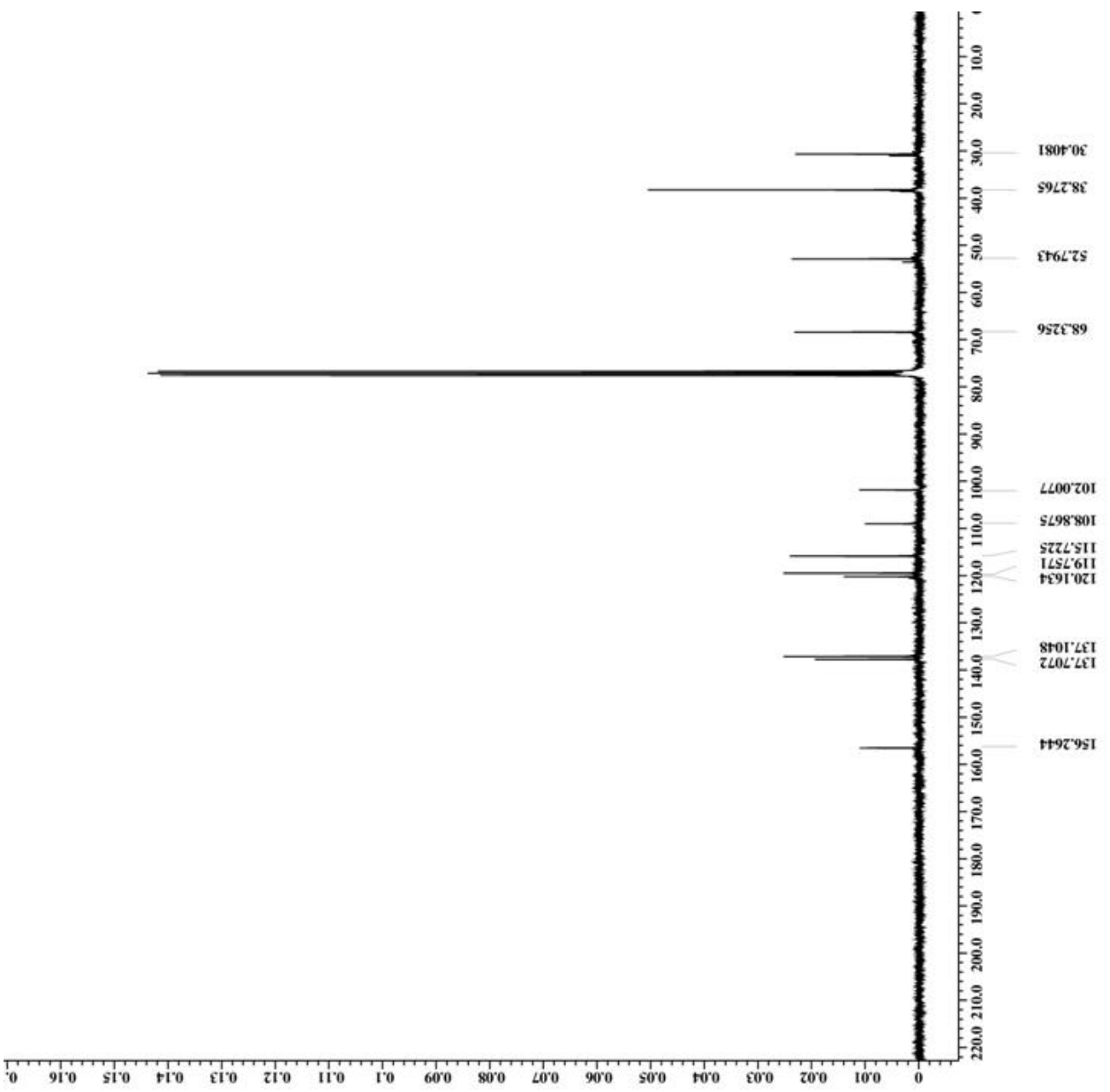


```

filename = sm_VI_47_pure-2.jdf
author = delta
experiment = single_pulse_dec
sample_id = S835989
solvent = CHLOROFORM-D
reaction_time = 2-DEC-2009 07:20:39
revision_time = 27-MAR-2010 20:52:42
current_time = 27-MAR-2010 20:52:36
comment = single pulse decouple
ata_format = ID COMPLEX
in_size = 52428
in_title = 13C
in_units = [ppm]
inmensions = X
ite = ECX 300
spectrometer = DELTA2_NMR
ield_strength = 7.0586013[T] (300[MHz]
_acq_duration = 2.76824064[s]
_domain = 13C
_freq = 75.56823426[MHz]
_offset = 100[ppm]
_points = 65536
_prescans = 5
_resolution = 0.36124027[Hz]
_sweep = 23.67424242[kHz]
tr_domain = 1H
tr_freq = 300.52965592[MHz]
tr_offset = 5[ppm]
lipped = FALSE
bd_return = 10
cans = 5000
otal_scans = 5000
_90_width = 9.75[us]
_acq_time = 2.76824064[s]
_angle = 30[deg]
_atn = 8[db]
_pulse = 3.25[us]
tr_atn_dec = 25[db]
tr_atn_noe = 25[db]
tr_noise = WALTZ
scoupling = TRUE
nitial_wait = 1[s]
oe_time = TRUE
oe_time = 3[s]
ecvr_gain = 50
elaxation_delay = 3[s]
epetition_time = 5.76824064[s]
emp_get = 23.3[dc]

```

abundance

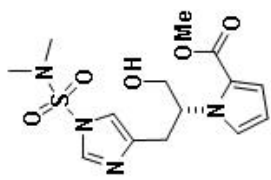
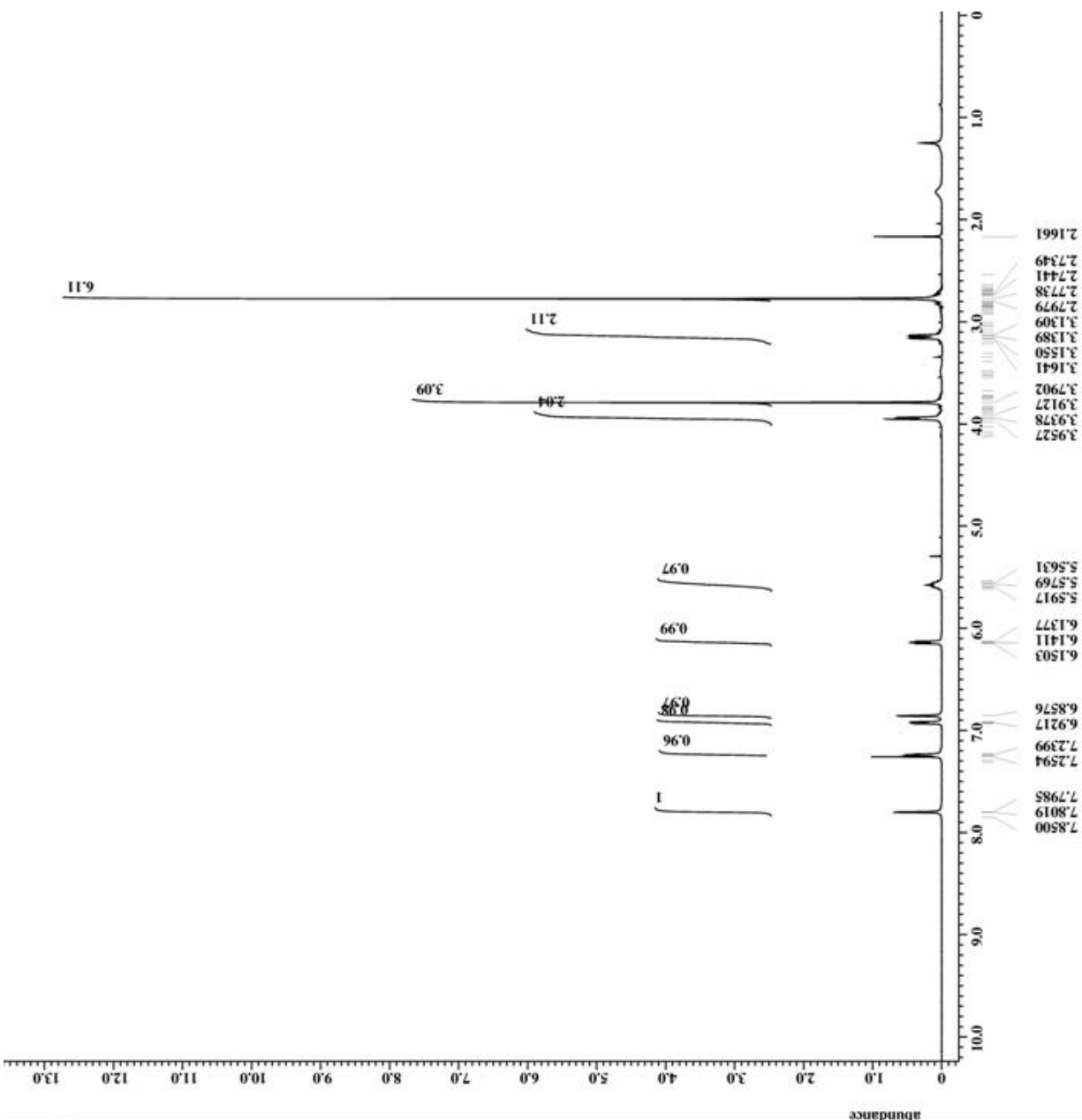


APPENDIX 22
¹H AND ¹³C NMR SPECTRUM OF
(S)-methyl 1-(1-(1-(N,N-dimethylsulfamoyl)-1H-imidazol-4-yl)-3-
hydroxypropan-2-yl)-1H-pyrrole-2-carboxylate (89)



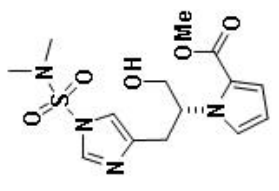
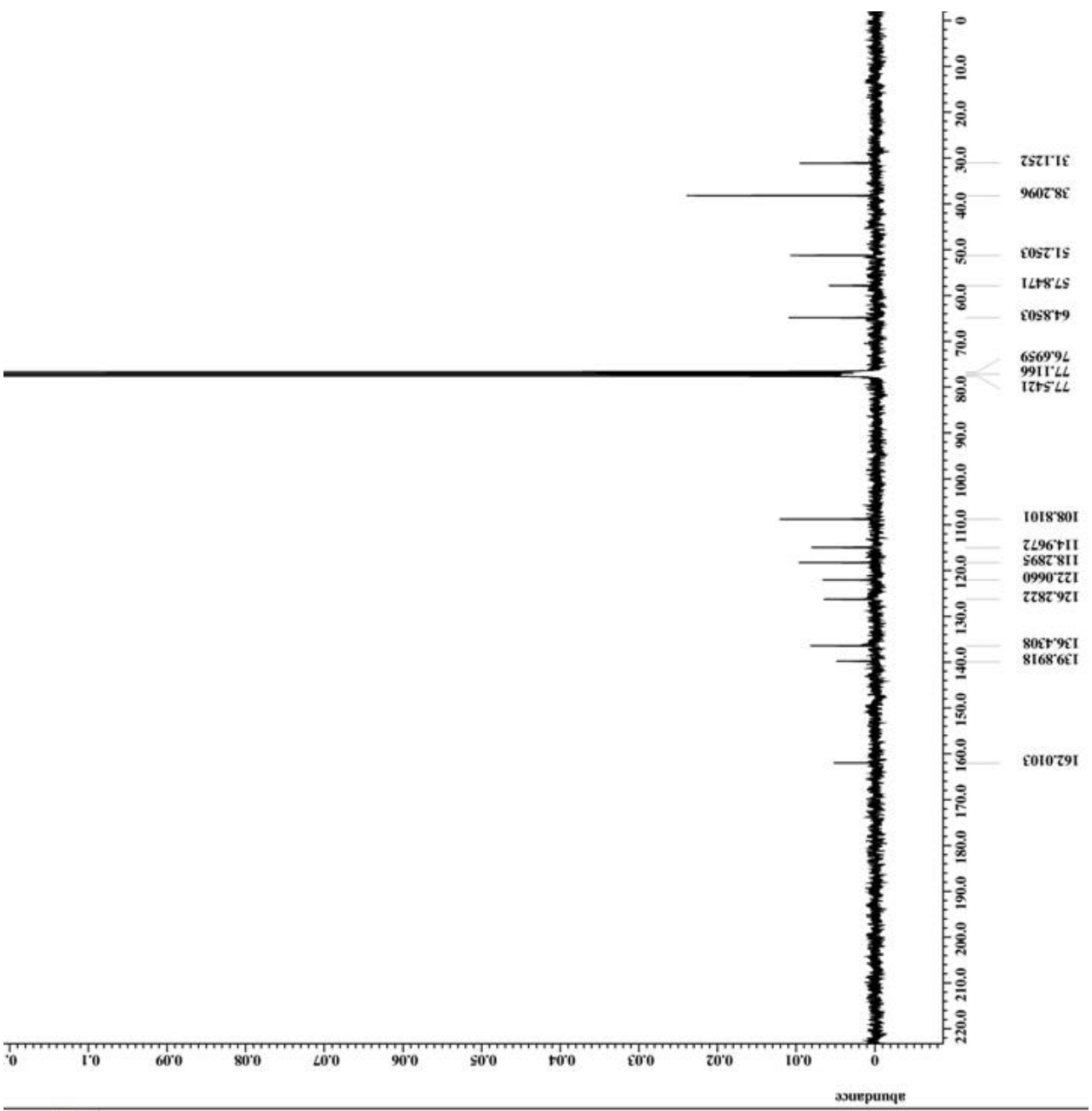
```

filename = sm_V_106_pure-4_jdf
author = delta
experiment = single_pulse.ex2
sample_id = S#503627
solvent = CHLOROFORM-D
creation_time = 26-MAR-2009 13:21:57
revision_time = 28-MAR-2010 11:11:40
current_time = 28-MAR-2010 11:12:13
comment = single_pulse
ata_format = ID REAL
in_size = 13107
in_title = 1H
in_units = [ppm]
inmensions = X
ite = ECX 300
spectrometer = DELTA2_NMR
ield_strength = 7.0586013[T] (300[MHz]
_acq_duration = 2.90717696[s]
_domain = 1H
_freq = 300.52965592[MHz]
_offset = 5[ppm]
_points = 16384
_prescans = 0
_resolution = 0.34397621[Hz]
_sweep = 5.63570784[kHz]
tr_domain = 1H
tr_freq = 300.52965592[MHz]
tr_offset = 5[ppm]
ri_domain = 1H
ri_freq = 300.52965592[MHz]
ri_offset = 5[ppm]
lipped = FALSE
od_return = 1
otal_scans = 24
_90_width = 13.01[us]
_acq_time = 2.90717696[s]
_angle = 45[deg]
_atn = 4[db]
_pulse = 6.505[us]
tr_mode = Off
ri_mode = Off
ante_presat = FALSE
ntial_wait = 1[s]
scvr_gain = 46
relaxation_delay = 5[s]
petition_time = 7.90717696[s]
emp_get = 23.1[dc]
  
```





```
filename = sm_V_106_pure-2.jdf
author = delta
experiment = single_pulse_dec
sample_id = S8516083
solvent = CHLOROFORM-D
reaction_time = 27-MAR-2009 15:15:46
revision_time = 27-MAR-2009 15:59:36
current_time = 28-MAR-2010 11:16:33
comment = single pulse decouple
ata_format = ID COMPLEX
in_size = 52428
in_title = 13C
in_units = [ppm]
in_dimensions = X
ite = ECX 300
ite_preamplifier = DELTA2_NMR
ield_strength = 7.0586013[T] (300[MHz])
_acq_duration = 2.76824064[s]
_domain = 13C
_freq = 75.56823426[MHz]
_offset = 100[ppm]
_points = 65536
_prescans = 4
_resolution = 0.36124027[Hz]
_sweep = 23.67424242[kHz]
rr_domain = 1H
rr_freq = 300.52965592[MHz]
rr_offset = 5[ppm]
lipped = FALSE
bd_return = 10
cans = 1000
atal_scans = 1000
_90_width = 9.75[us]
_acq_time = 2.76824064[s]
_angle = 30[deg]
_atn = 8[dB]
_pulse = 3.25[us]
rr_atn_dec = 25[dB]
rr_atn_noe = 25[dB]
rr_noise = WALTZ
scoupling = TRUE
initial_wait = 1[s]
be = TRUE
be_time = 3[s]
ecvr_gain = 50
relaxation_delay = 3[s]
petition_time = 5.76824064[s]
emp_get = 23.3[dc]
```



APPENDIX 23
¹H AND ¹³C NMR SPECTRUM OF
(S)-methyl 1-(1-azido-3-(1-(N,N-dimethylsulfamoyl)-1H-imidazol-4-yl)propan-2-yl)-1H-pyrrole-2-carboxylate (91)



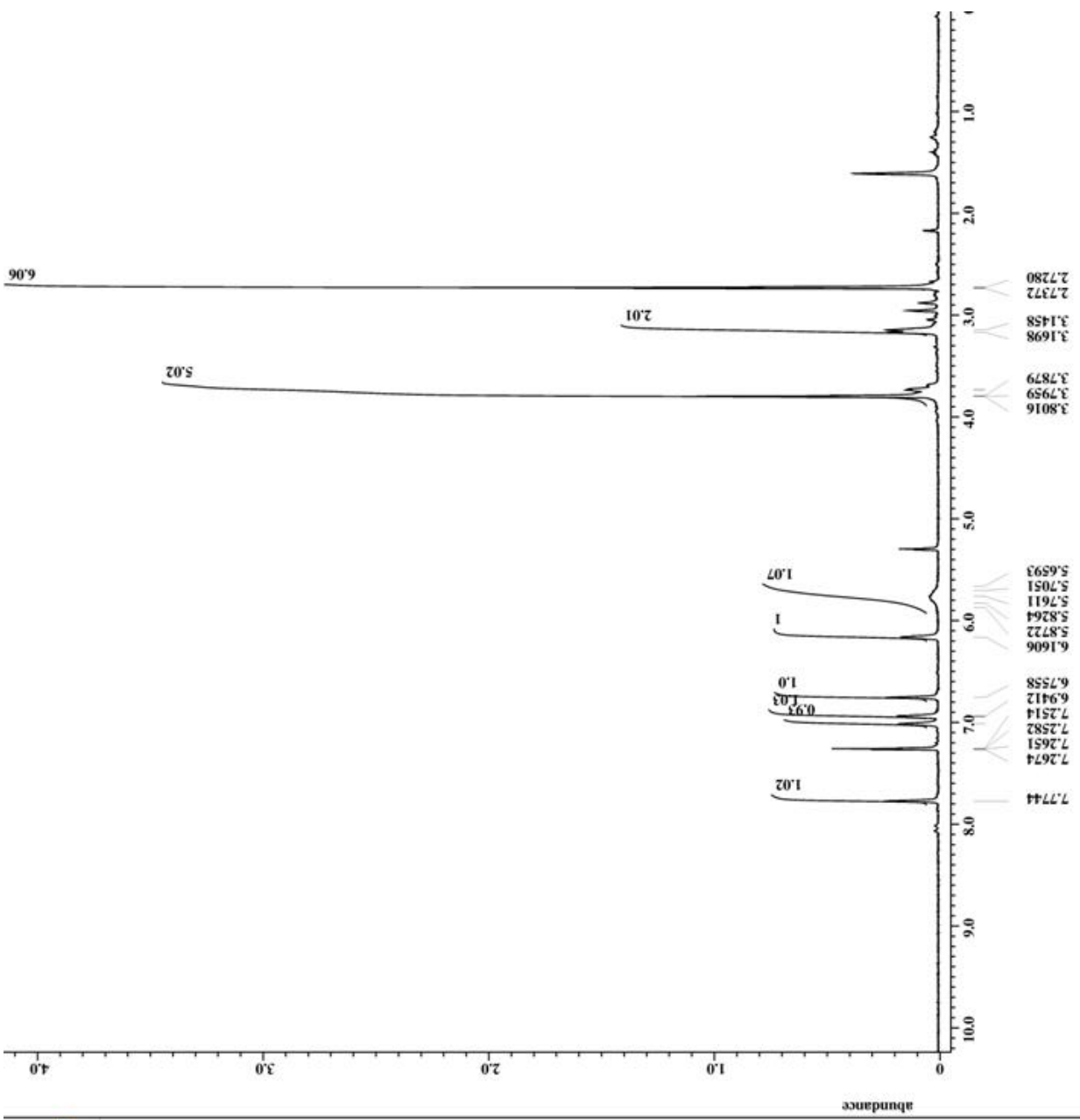
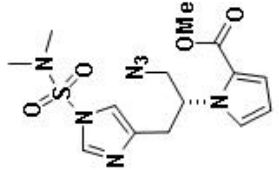
```

filename = sm_VI_39_pure-3_jdf
author = delta
experiment = single_pulse.ex2
sample_id = S8638625
solvent = CHLOROFORM-D
creation_time = 26-NOV-2009 17:53:41
revision_time = 28-MAR-2010 11:21:55
current_time = 28-MAR-2010 11:22:12

comment = single_pulse
ata_format = ID COMPLEX
in_size = 13107
in_title = 1H
in_units = [ppm]
inmensions = X
ite = ECX 300
spectrometer = DELTA2_NMR

ield_strength = 7.0586013[T] (300[MHz]
_acq_duration = 2.90717696[s]
_domain = 1H
_freq = 300.52965592[MHz]
_offset = 5[ppm]
_points = 16384
_prescans = 0
_resolution = 0.34397621[Hz]
_sweep = 5.63570784[kHz]
rr_domain = 1H
rr_freq = 300.52965592[MHz]
rr_offset = 5[ppm]
ri_domain = 1H
ri_freq = 300.52965592[MHz]
ri_offset = 5[ppm]
lipped = FALSE
od_return = 1
atal_scans = 23

_90_width = 13.01[us]
_acq_time = 2.90717696[s]
_angle = 45[deg]
_atn = 4[db]
_pulse = 6.505[us]
rr_mode = Off
ri_mode = Off
ante_presat = FALSE
nitial_wait = 1[s]
scvr_gain = 46
relaxation_delay = 5[s]
petition_time = 7.90717696[s]
emp_get = 23.4[dc]
  
```



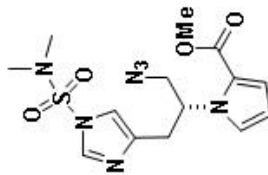
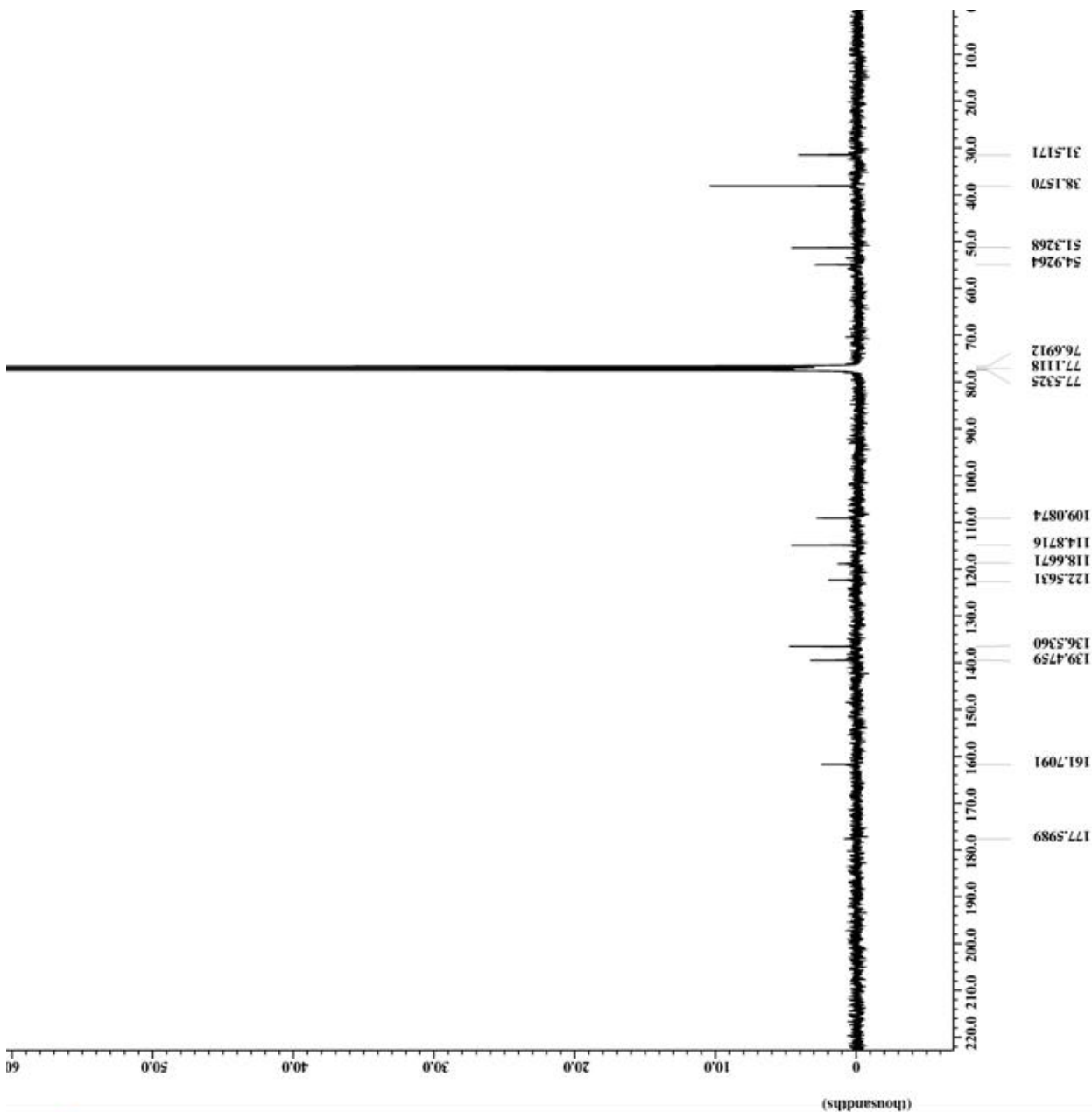


```
filename = sm_VI_39_pure-2.jdf
author =
experiment = single_pulse_dec
pulse_id = S8641182
solvent = CHLOROFORM-D
reaction_time = 27-NOV-2009 00:19:11
revision_time = 27-NOV-2009 17:46:00
current_time = 28-MAR-2010 11:30:13

comment = single pulse decouple
ata_format = ID COMPLEX
in_size = 52428
in_title = 13C
in_units = [ppm]
inmensions = X
ite = ECX 300
spectrometer = DELTA2_NMR

ield_strength = 7.0586013[T] (300[MHz]
acq_duration = 2.76824064[s]
domain = 13C
freq = 75.56823426[MHz]
offset = 100[ppm]
points = 65536
prescans = 4
resolution = 0.36124027[Hz]
sweep = 23.67424242[kHz]
tr_domain = 1H
tr_freq = 300.52965592[MHz]
tr_offset = 5[ppm]
lipped = FALSE
bd_return = 10
cans = 4000
atal_scans = 4000

_90_width = 9.75[us]
acq_time = 2.76824064[s]
angle = 30[deg]
atn = 8[db]
pulse = 3.25[us]
tr_atn_dec = 25[db]
tr_atn_noe = 25[db]
tr_noise = WALTZ
scoupling = TRUE
nitral_wait = 1[s]
oe_time = TRUE
oe_time = 3[s]
ecvr_gain = 50
relaxation_delay = 3[s]
petition_time = 5.76824064[s]
emp_get = 22.9[dc]
```



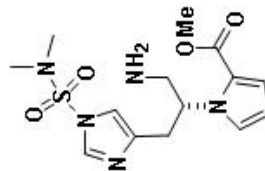
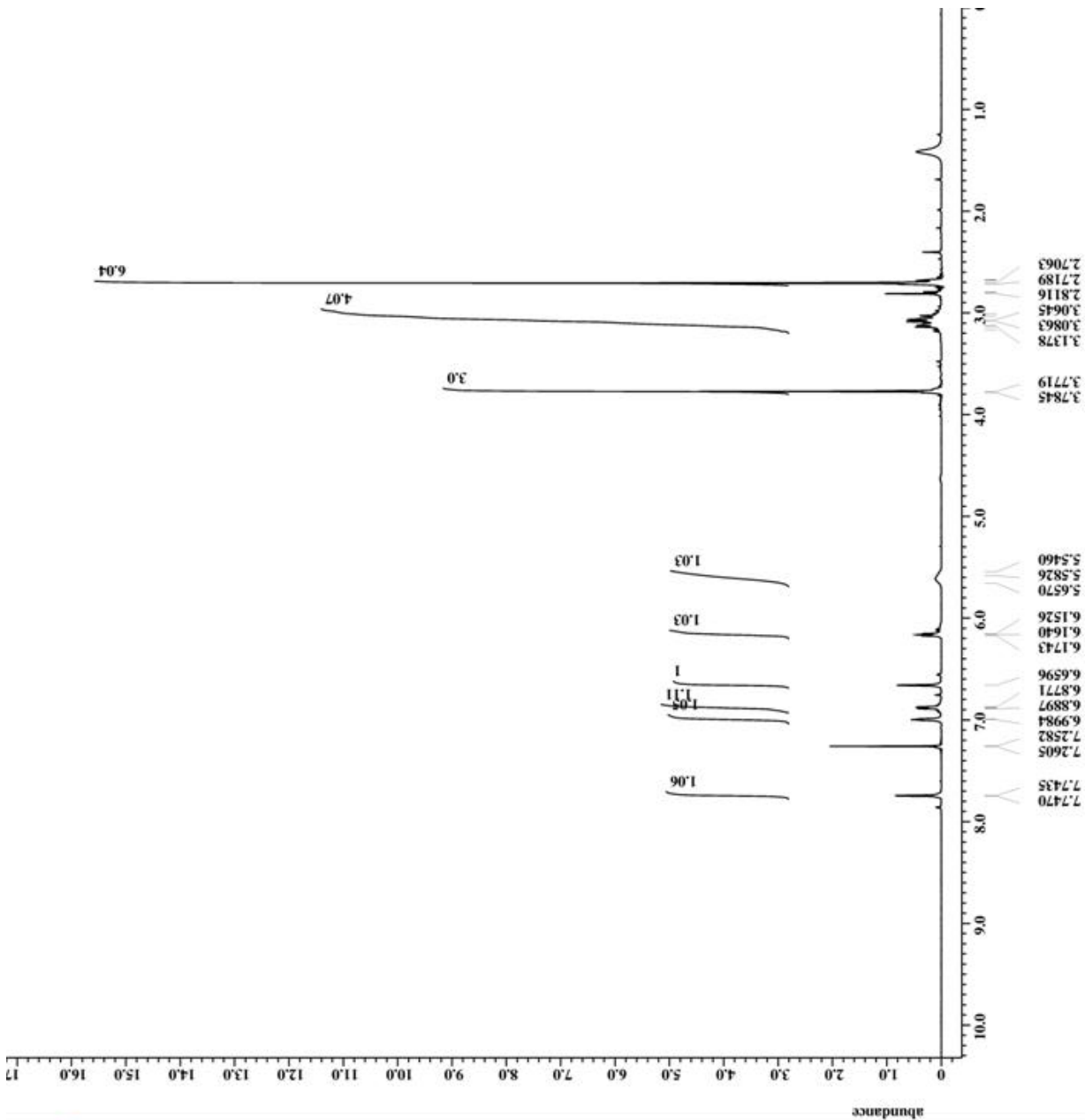
APPENDIX 24
¹H AND ¹³C NMR SPECTRUM OF
(S)-methyl 1-(1-amino-3-(1-(N,N-dimethylsulfamoyl)-1H-imidazol-4-yl)propan-2-yl)-1H-pyrrole-2-carboxylate (92)



```

filename = sm_V_181-4.jdf
author = delta
experiment = single_pulse.ex2
sample_id = S8664788
solvent = CHLOROFORM-D
reaction_time = 3-AUG-2009 18:24:07
evision_time = 28-MAR-2010 11:34:56
current_time = 28-MAR-2010 11:35:12
comment = single_pulse
ata_format = ID REAL
in_size = 13107
in_title = 1H
in_units = [ppm]
inmensions = X
ite = ECX 300
spectrometer = DELTA2_NMR
ield_strength = 7.0586013[T] (300[MHz]
_acq_duration = 2.90717696[s]
_domain = 1H
_freq = 300.52965592[MHz]
_offset = 5[ppm]
_points = 16384
_prescans = 0
_resolution = 0.34397621[Hz]
_sweep = 5.63570784[kHz]
rr_domain = 1H
rr_freq = 300.52965592[MHz]
rr_offset = 5[ppm]
ri_domain = 1H
ri_freq = 300.52965592[MHz]
ri_offset = 5[ppm]
lipped = FALSE
od_return = 1
otal_scans = 24

_90_width = 13.01[us]
_acq_time = 2.90717696[s]
_angle = 45[deg]
_atn = 4[db]
_pulse = 6.505[us]
rr_mode = Off
ri_mode = Off
ante_presat = FALSE
nitial_wait = 1[s]
ecvr_gain = 50
elaxation_delay = 5[s]
epetition_time = 7.90717696[s]
emp_get = 23.2[dc]
  
```





```

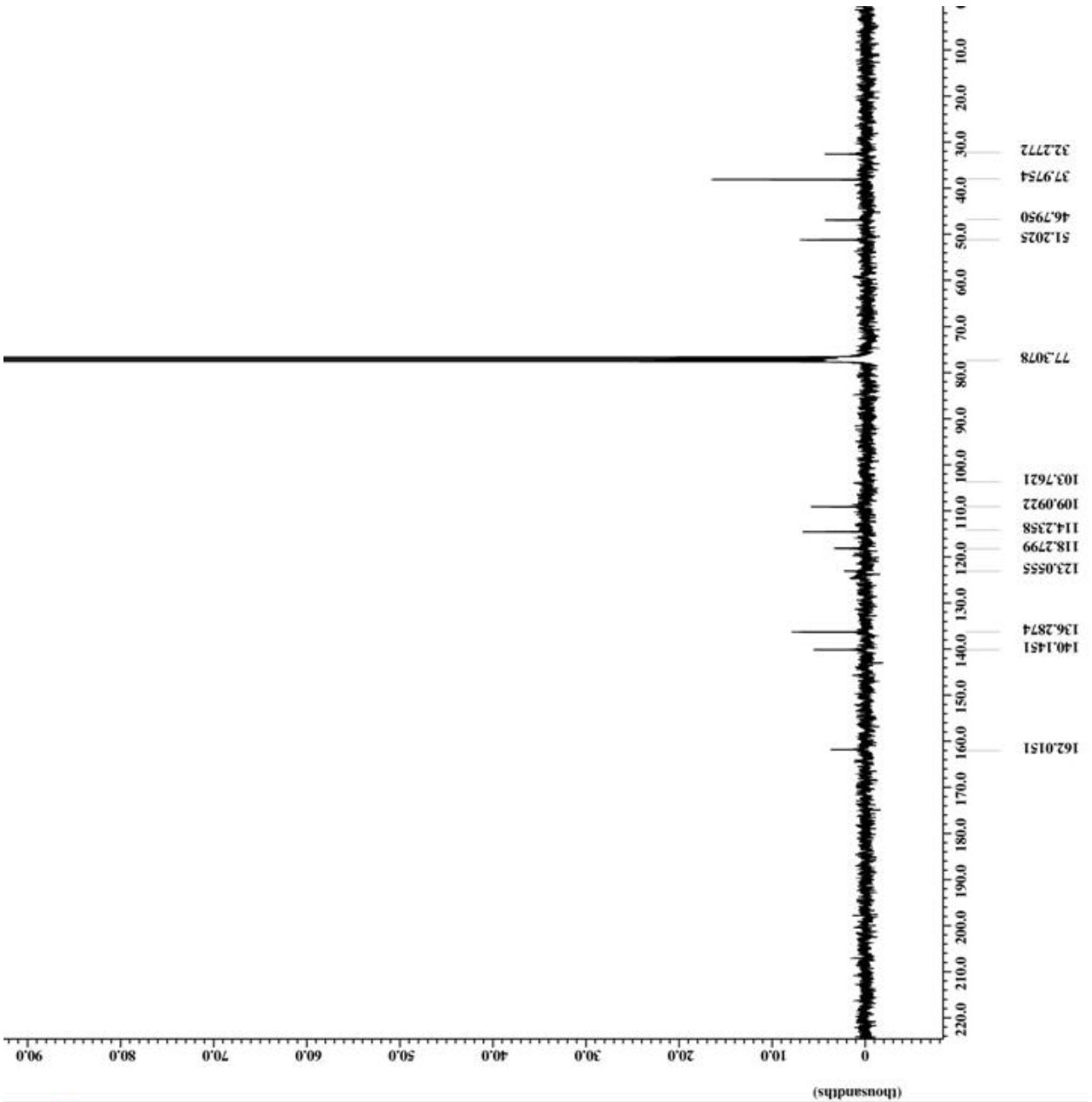
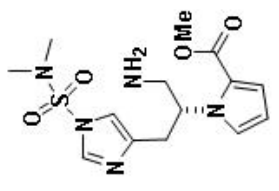
filename = sm_V_181-2.jdf
author = delta
experiment = single_pulse_dec
sample_id = S866919
solvent = CHLOROFORM-D
reaction_time = 3-AUG-2009 20:02:03
revision_time = 3-AUG-2009 22:31:12
current_time = 28-MAR-2010 11:39:24

comment = single pulse decouple
ata_format = ID COMPLEX
in_size = 52428
in_title = 13C
in_units = [ppm]
inmensions = X
ite = ECX 300
spectrometer = DELTA2_NMR

ield_strength = 7.0586013[T] (300[MHz]
acq_duration = 2.76824064[s]
domain = 13C
freq = 75.56823426[MHz]
offset = 100[ppm]
points = 65536
prescans = 4
resolution = 0.36124027[Hz]
sweep = 23.67424242[kHz]
tr_domain = 1H
tr_freq = 300.52965592[MHz]
tr_offset = 5[ppm]
lipped = FALSE
bd_return = 10
cans = 1000
atal_scans = 1000

_90_width = 9.75[us]
acq_time = 2.76824064[s]
angle = 30[deg]
atn = 8[db]
pulse = 3.25[us]
tr_atn_dec = 25[db]
tr_atn_noe = 25[db]
tr_noise = WALTZ
scoupling = TRUE
nitial_wait = 1[s]
oe_time = TRUE
oe_time = 3[s]
scr_gain = 50
relaxation_delay = 3[s]
petition_time = 5.76824064[s]
emp_get = 23.3[dc]

```



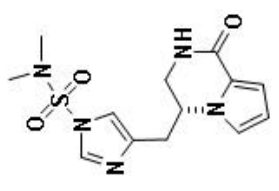
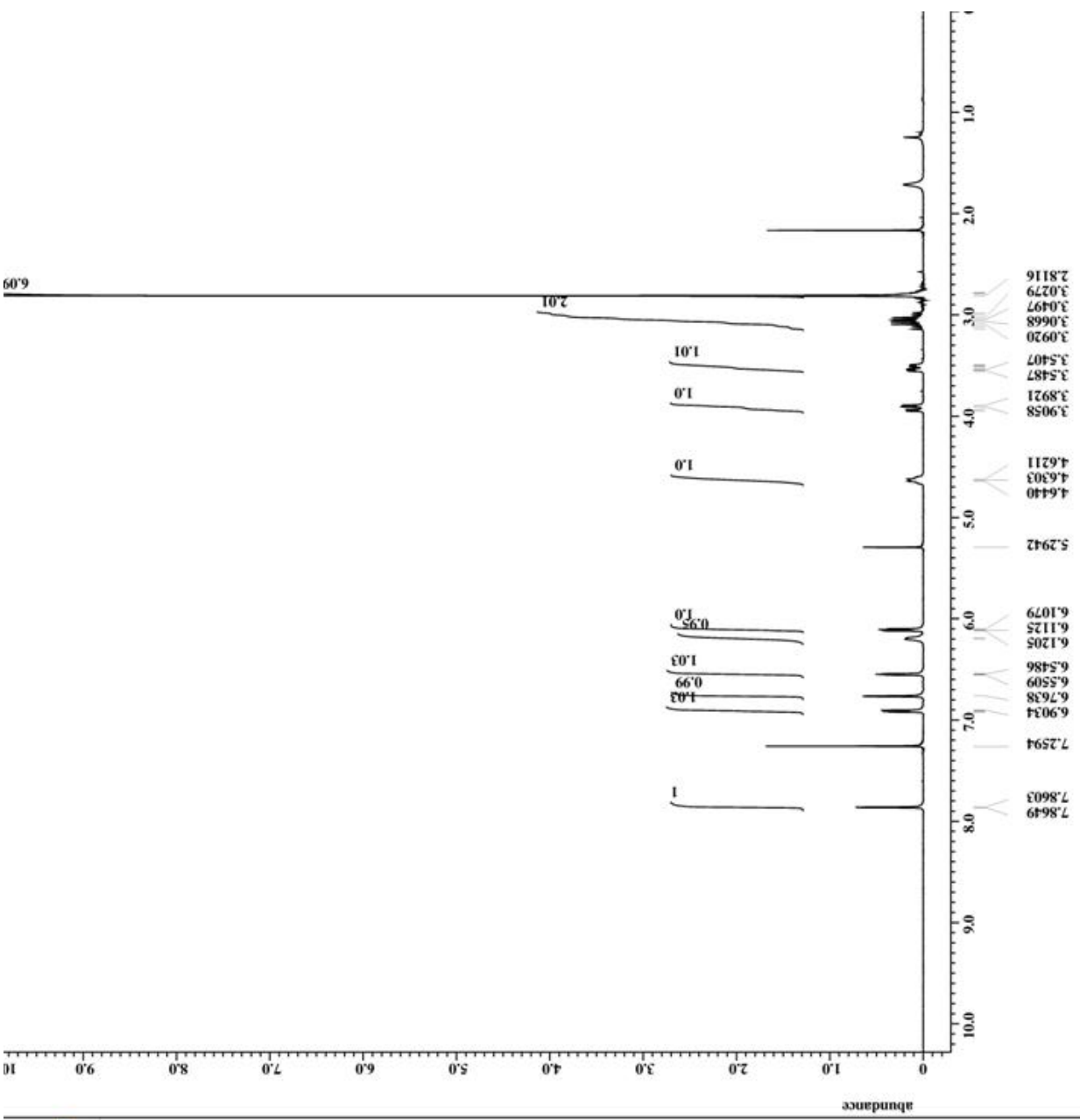
APPENDIX 25
¹H AND ¹³C NMR SPECTRUM OF
(S)-N,N-dimethyl-4-((1-oxo-1,2,3,4-tetrahydropyrrolo[1,2-a]pyrazin-4-yl)methyl)-1H-imidazole-1-sulfonamide (55)



```

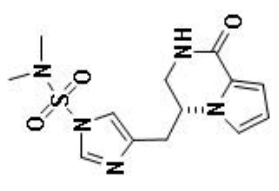
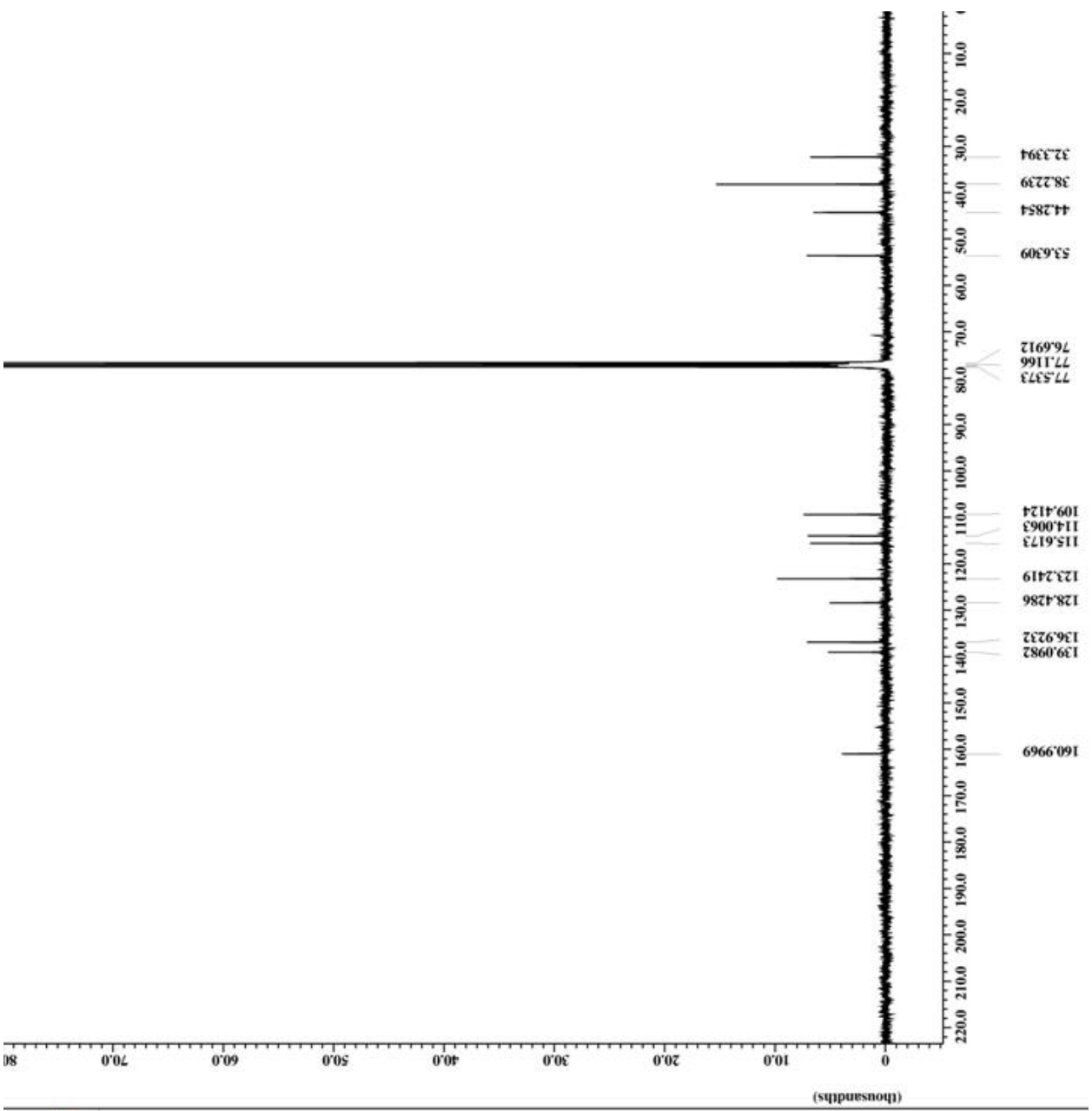
filename = sm_V_189_CRUDE-4_jdf
author = delta
experiment = single_pulse.ex2
sample_id = S9483444
solvent = CHLOROFORM-D
reaction_time = 14-AUG-2009 13:23:17
evolution_time = 28-MAR-2010 11:42:50
current_time = 28-MAR-2010 11:43:37
comment = single_pulse
ata_format = ID REAL
in_size = 13107
in_title = 1H
in_units = [ppm]
in_dimensions = X
ite = ECX 300
spectrometer = DELTA2_NMR
ield_strength = 7.0586013[T] (300[MHz]
_acq_duration = 2.90717696[s]
_domain = 1H
_freq = 300.52965592[MHz]
_offset = 5[ppm]
_points = 16384
_prescans = 0
_resolution = 0.34397621[Hz]
_sweep = 5.63570784[kHz]
tr_domain = 1H
tr_freq = 300.52965592[MHz]
tr_offset = 5[ppm]
ri_domain = 1H
ri_freq = 300.52965592[MHz]
ri_offset = 5[ppm]
lipped = FALSE
od_return = 1
atal_scans = 24
_90_width = 13.01[us]
_acq_time = 2.90717696[s]
_angle = 45[deg]
_atn = 4[dB]
_pulse = 6.505[us]
tr_mode = Off
ri_mode = Off
ante_presat = FALSE
ntial_wait = 1[s]
scvr_gain = 46
relaxation_delay = 5[s]
petition_time = 7.90717696[s]
emp_get = 22.8[dc]

```





```
filename = sm_V_189_pure-2.jdf
author = delta
experiment = single_pulse_dec
sample_id = S#806600
solvent = CHLOROFORM-D
reaction_time = 15-AUG-2009 07:55:35
evision_time = 15-AUG-2009 11:46:50
current_time = 28-MAR-2010 11:46:47
comment = single pulse decouple
ata_format = ID COMPLEX
in_size = 52428
in_title = 13C
in_units = [ppm]
inmensions = X
ite = ECX 300
pectrometer = DELTA2_NMR
ield_strength = 7.0586013[T] (300[MHz]
_acq_duration = 2.76824064[s]
_domain = 13C
_freq = 75.56823426[MHz]
_offset = 100[ppm]
_points = 65536
_prescans = 4
_resolution = 0.36124027[Hz]
_sweep = 23.67424242[kHz]
rr_domain = 1H
rr_freq = 300.52965592[MHz]
rr_offset = 5[ppm]
lipped = FALSE
od_return = 10
cans = 6000
otal_scans = 6000
_90_width = 9.75[us]
_acq_time = 2.76824064[s]
_angle = 30[deg]
_atn = 8[db]
_pulse = 3.25[us]
rr_atn_dec = 25[db]
rr_atn_noe = 25[db]
rr_noise = WALTZ
scoupling = TRUE
nitial_wait = 1[s]
oe_time = TRUE
oe_time = 3[s]
ecvr_gain = 50
elaxation_delay = 3[s]
epetition_time = 5.76824064[s]
emp_get = 23.7[dc]
```



APPENDIX 26
¹H AND ¹³C NMR SPECTRUM OF
(S)-4-((6,7-dibromo-1-oxo-1,2,3,4-tetrahydropyrrolo[1,2-a]pyrazin-4-yl)methyl)-N,N-dimethyl-1H-imidazole-1-sulfonamide (93)

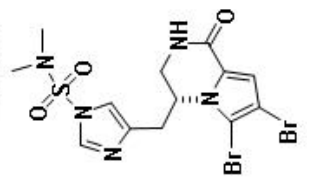
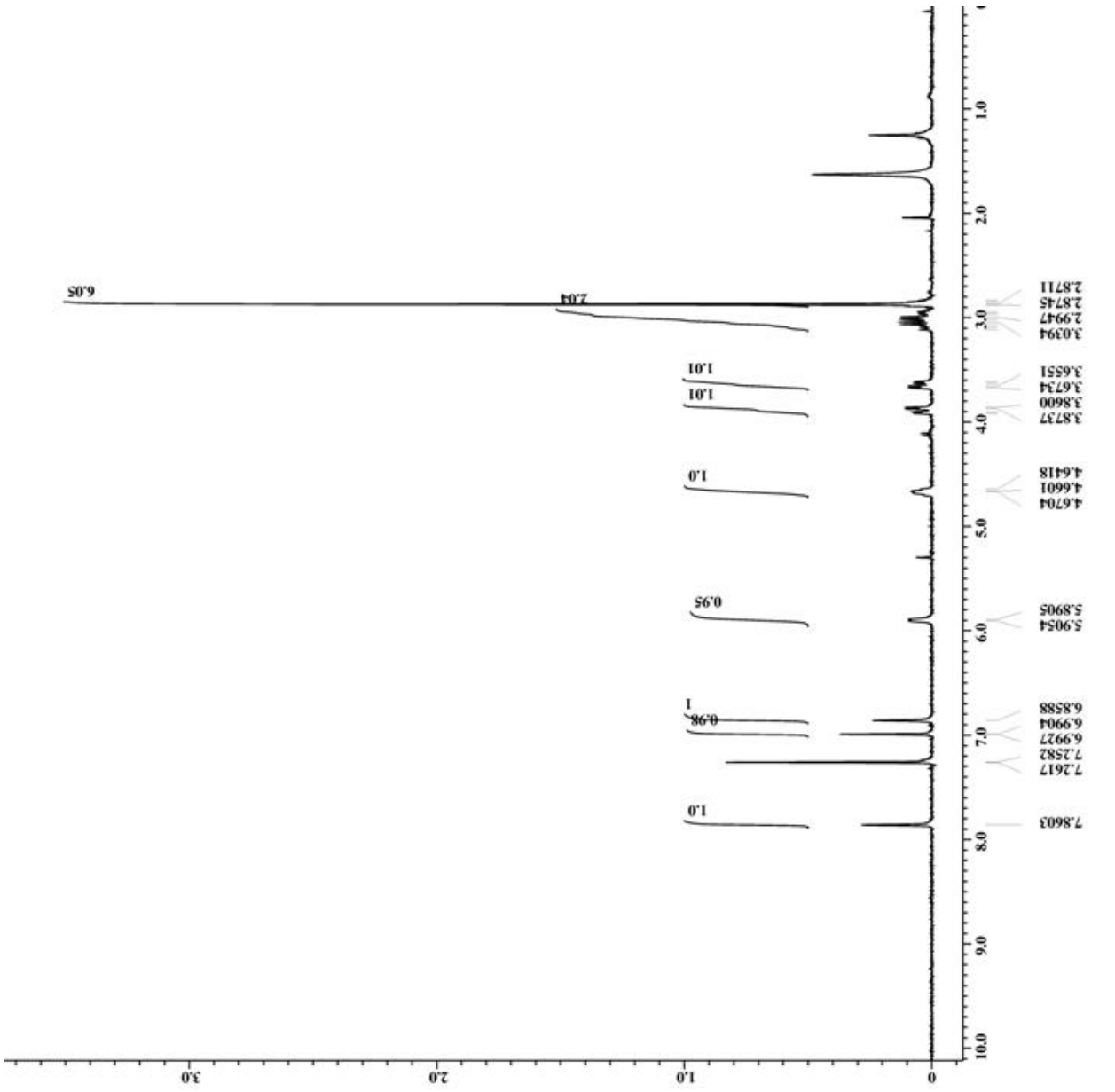


```

filename = sm_VI_7B_pure-3.jdf
author = delta
experiment = single_pulse.ex2
sample_id = S8665729
solvent = CHLOROFORM-D
reaction_time = 20-OCT-2009 18:33:57
evision_time = 20-OCT-2009 18:35:46
current_time = 28-MAR-2010 11:54:59
comment =
  = single_pulse
  = ID REAL
  = 13107
  = 1H
  = [ppm]
  = X
  = ECX 300
  = DELTA2_NMR
ield_strength = 7.0586013[T] (300[MHz])
  _acq_duration = 2.90717696[s]
  _domain = 1H
  _freq = 300.52965592[MHz]
  _offset = 5[ppm]
  _points = 16384
  _prescans = 0
  _resolution = 0.34397621[Hz]
  _sweep = 5.63570784[kHz]
  rr_domain = 1H
  rr_freq = 300.52965592[MHz]
  rr_offset = 5[ppm]
  ri_domain = 1H
  ri_freq = 300.52965592[MHz]
  ri_offset = 5[ppm]
  lipped = FALSE
  od_return = 1
  otal_scans = 16
  _90_width = 13.01[us]
  _acq_time = 2.90717696[s]
  _angle = 45[deg]
  _atn = 4[db]
  _pulse = 6.505[us]
  rr_mode = Off
  ri_mode = Off
  ante_presat = FALSE
  nitial_wait = 1[s]
  scvr_gain = 48
  relaxation_delay = 5[s]
  repetition_time = 7.90717696[s]
  temp_get = 23.5[degC]

```

abundance





```

filename = sm_VI_7B_ii_pure-2.jd
author = delta
experiment = single_pulse_dec
sample_id = S95710
solvent = CHLOROFORM-D
reaction_time = 21-OCT-2009 09:48:21
acquisition_time = 21-OCT-2009 09:50:27
current_time = 28-MAR-2010 11:57:45

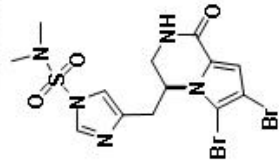
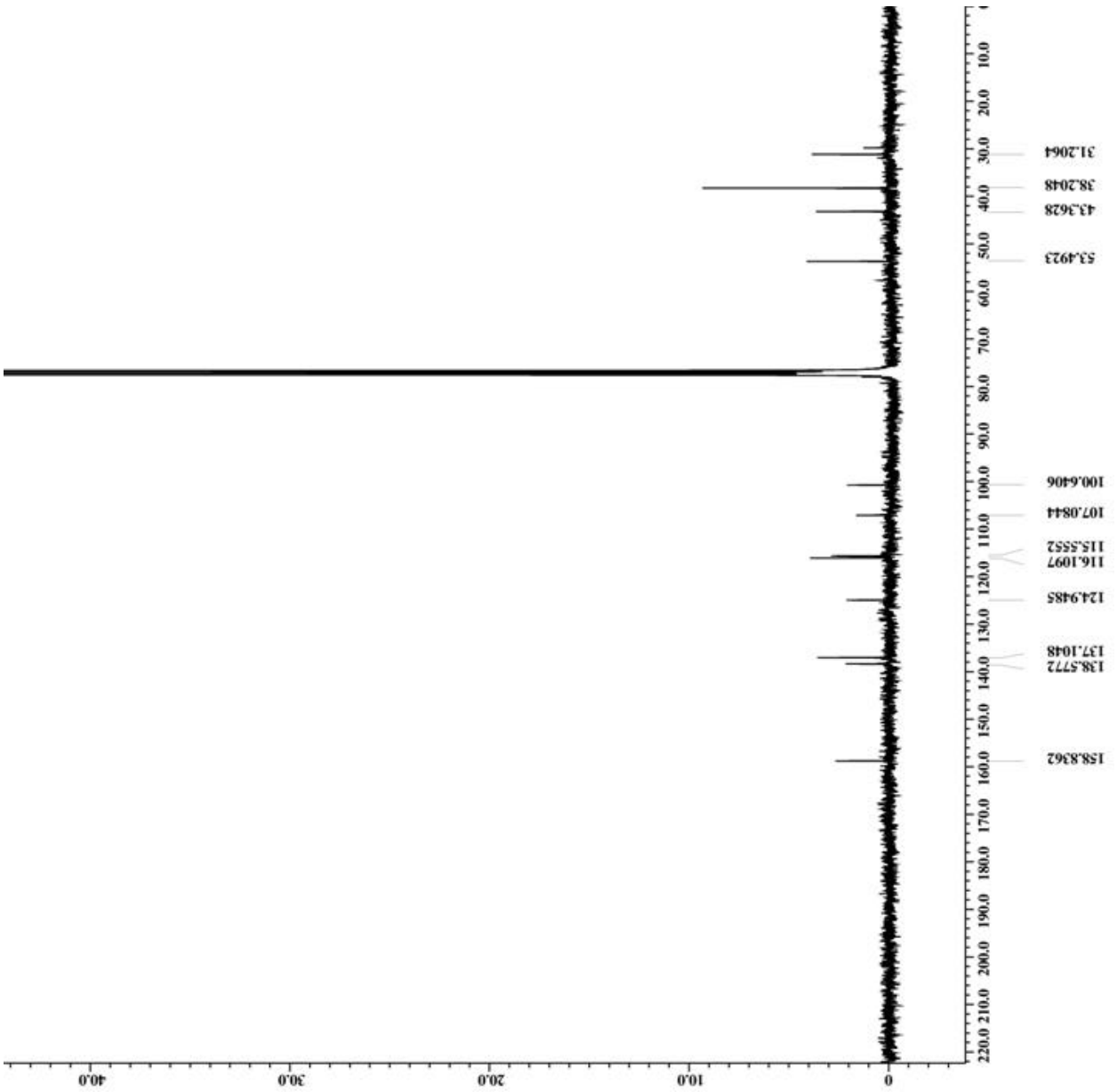
comment = single pulse decouple
ata_format = ID COMPLEX
in_size = 52428
in_title = 13C
in_units = [ppm]
in_dimensions = X
ite = ECX 300
spectrometer = DELTA2_NMR

ield_strength = 7.0586013[T] (300[MHz]
acq_duration = 2.76824064[s]
domain = 13C
freq = 75.56823426[MHz]
offset = 100[ppm]
points = 65536
prescans = 4
resolution = 0.36124027[Hz]
sweep = 23.67424242[kHz]
tr_domain = 1H
tr_freq = 300.52965592[MHz]
tr_offset = 5[ppm]
lipped = FALSE
bd_return = 10
cans = 6000
atal_scans = 6000

_90_width = 9.75[us]
acq_time = 2.76824064[s]
angle = 30[deg]
atn = 8[db]
pulse = 3.25[us]
tr_atn_dec = 25[db]
tr_atn_noe = 25[db]
tr_noise = WALTZ
scoupling = TRUE
nitial_wait = 1[s]
oe_time = TRUE
oe_time = 3[s]
ecvr_gain = 50
relaxation_delay = 3[s]
petition_time = 5.76824064[s]
emp_get = 23.7[dc]

```

(thousands)





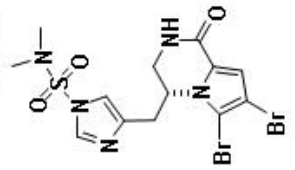
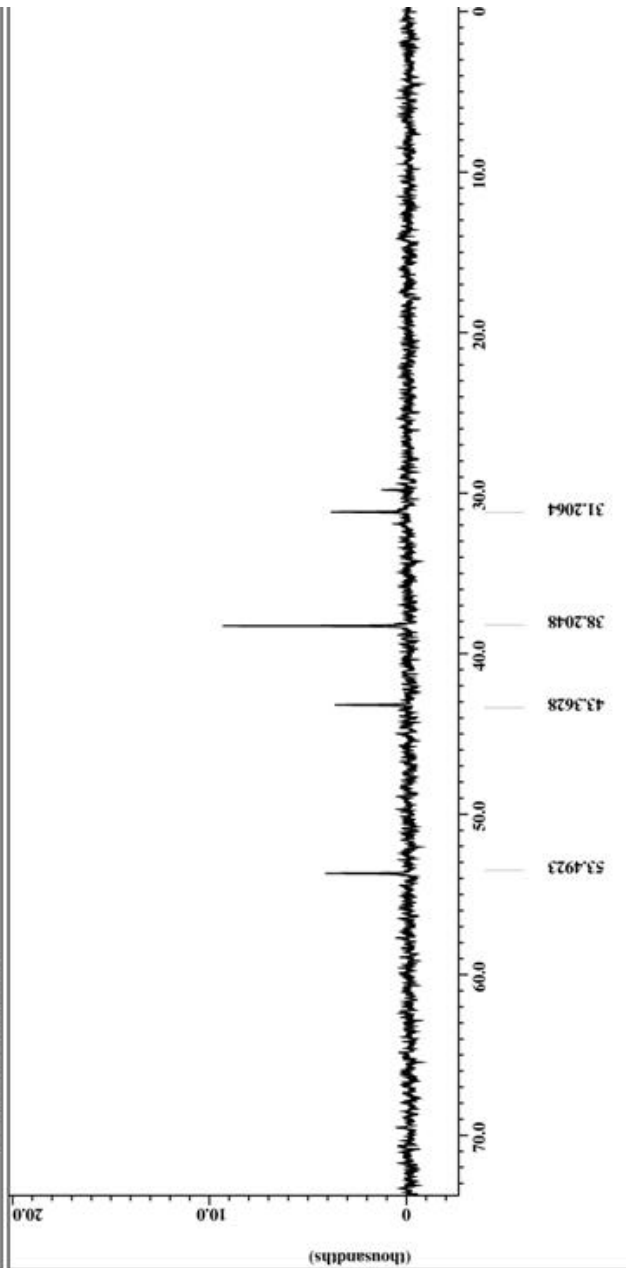
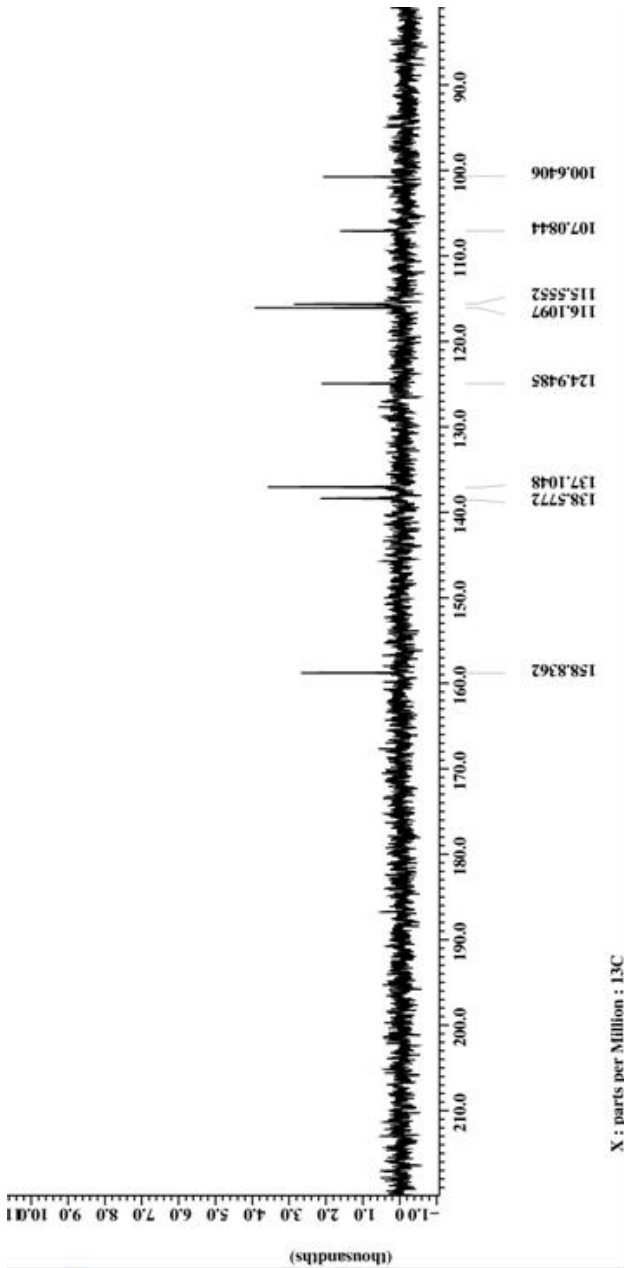
```

filename = sm_VI_7B_i1_pure-2.jd
author = delta
experiment = single_pulse_dec
sample_id = S95710
solvent = CHLOROFORM-D
reaction_time = 21-OCT-2009 09:48:21
evision_time = 21-OCT-2009 09:50:27
current_time = 28-MAR-2010 11:59:46

comment = single pulse decouple
ata_format = ID COMPLEX
in_size = 52428
in_title = 13C
in_units = [ppm]
inmensions = X
ite = ECX 300
pectrometer = DELTA2_NMR

ield_strength = 7.0586013[T] (300 [MHz]
_domain = 2.76824064[s]
_freq = 75.56823426[MHz]
_offset = 100[ppm]
_points = 65536
_prescans = 4
_resolution = 0.36124027[Hz]
_sweep = 23.67424242[kHz]
rr_domain = 1H
rr_freq = 300.52965592[MHz]
rr_offset = 5[ppm]
lipped = FALSE
od_return = 10
ctal_scans = 6000

_90_width = 9.75[us]
_acq_time = 2.76824064[s]
_angle = 30[deg]
_atn = 8[db]
_pulse = 3.25[us]
rr_atn_dec = 25[db]
rr_atn_noe = 25[db]
rr_noise = WALTZ
scoupling = TRUE
nitial_wait = 1[s]
oe_time = 3[s]
ecvr_gain = 50
elaxation_delay = 3[s]
epetition_time = 5.76824064[s]
emp_get = 23.7[dc]
  
```

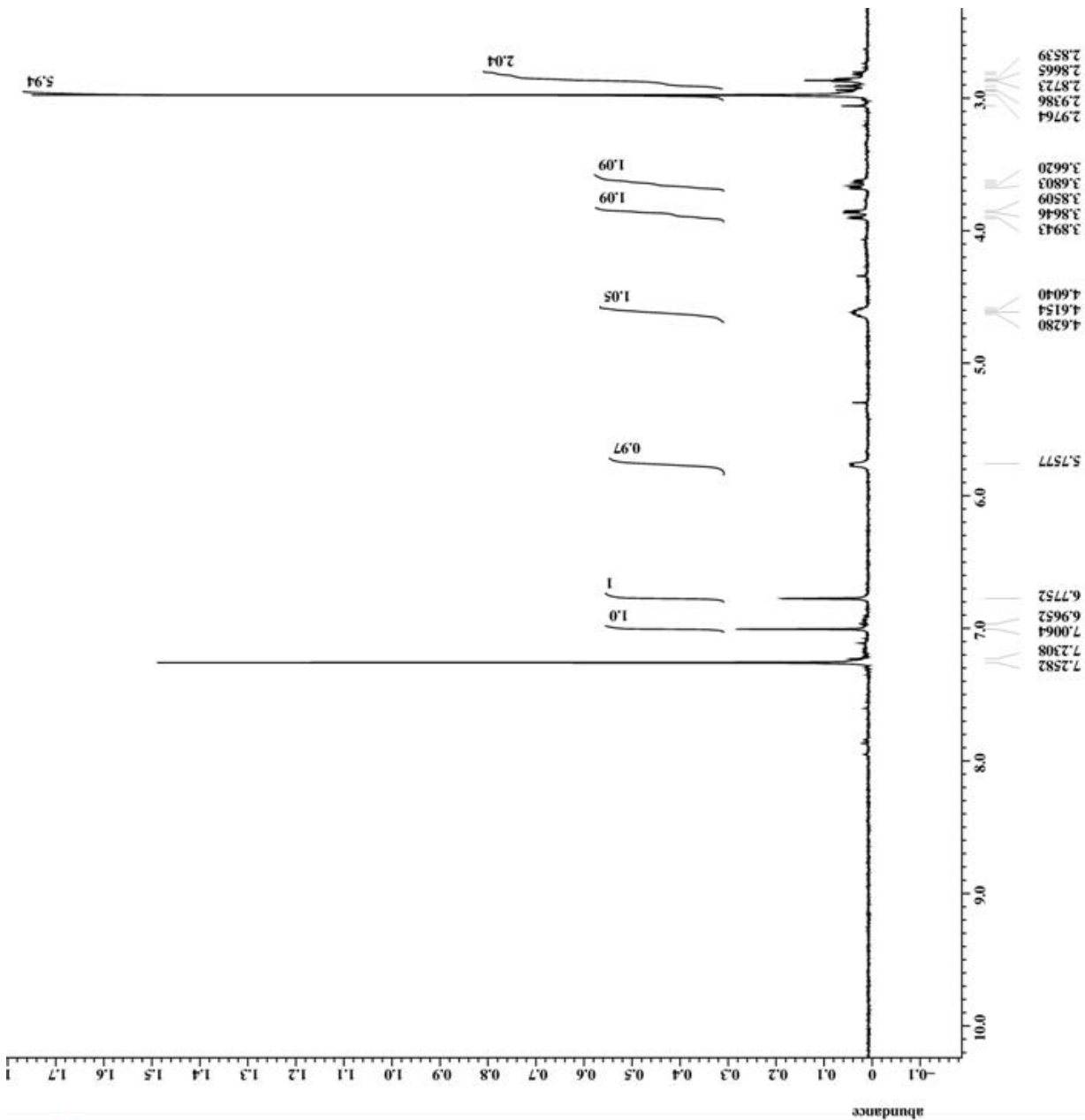


APPENDIX 27
¹H AND ¹³C NMR SPECTRUM OF
(S)-2-azido-4-((6,7-dibromo-1-oxo-1,2,3,4-tetrahydropyrrolo[1,2-a]pyrazin-4-yl)methyl)-N,N-dimethyl-1H-imidazole-1-sulfonamide (94)



```

filename = sm_VI_46_pure-3.jdf
author = delta
experiment = single_pulse.ex2
sample_id = S#783465
solvent = CHLOROFORM-D
creation_time = 30-NOV-2009 21:55:27
revision_time = 28-MAR-2010 12:15:24
current_time = 28-MAR-2010 12:16:00
comment =
  = single_pulse
  = ID COMPLEX
  = 13107
  = 1H
  = [ppm]
  = X
  = ECX 300
  = DELTA2_NMR
field_strength = 7.0586013[T] (300[MHz])
acq_duration = 2.90717696[s]
domain = 1H
freq = 300.52965592[MHz]
offset = 5[ppm]
points = 16384
prescans = 0
resolution = 0.34397621[Hz]
sweep = 5.63570784[kHz]
  = 1H
  = 300.52965592[MHz]
  = 5[ppm]
  = 1H
  = 300.52965592[MHz]
  = 5[ppm]
  = FALSE
lippped = 1
od_return = 24
otal_scans = 24
  = 13.01[us]
  = 2.90717696[s]
  = 45[deg]
  = 4[db]
  = 6.505[us]
  = Off
  = Off
  = FALSE
  = 1[s]
  = 46
  = 5[s]
  = 7.90717696[s]
  = 22.8[dc]
  
```





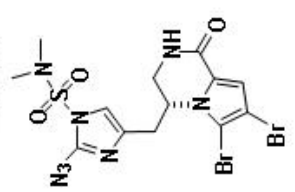
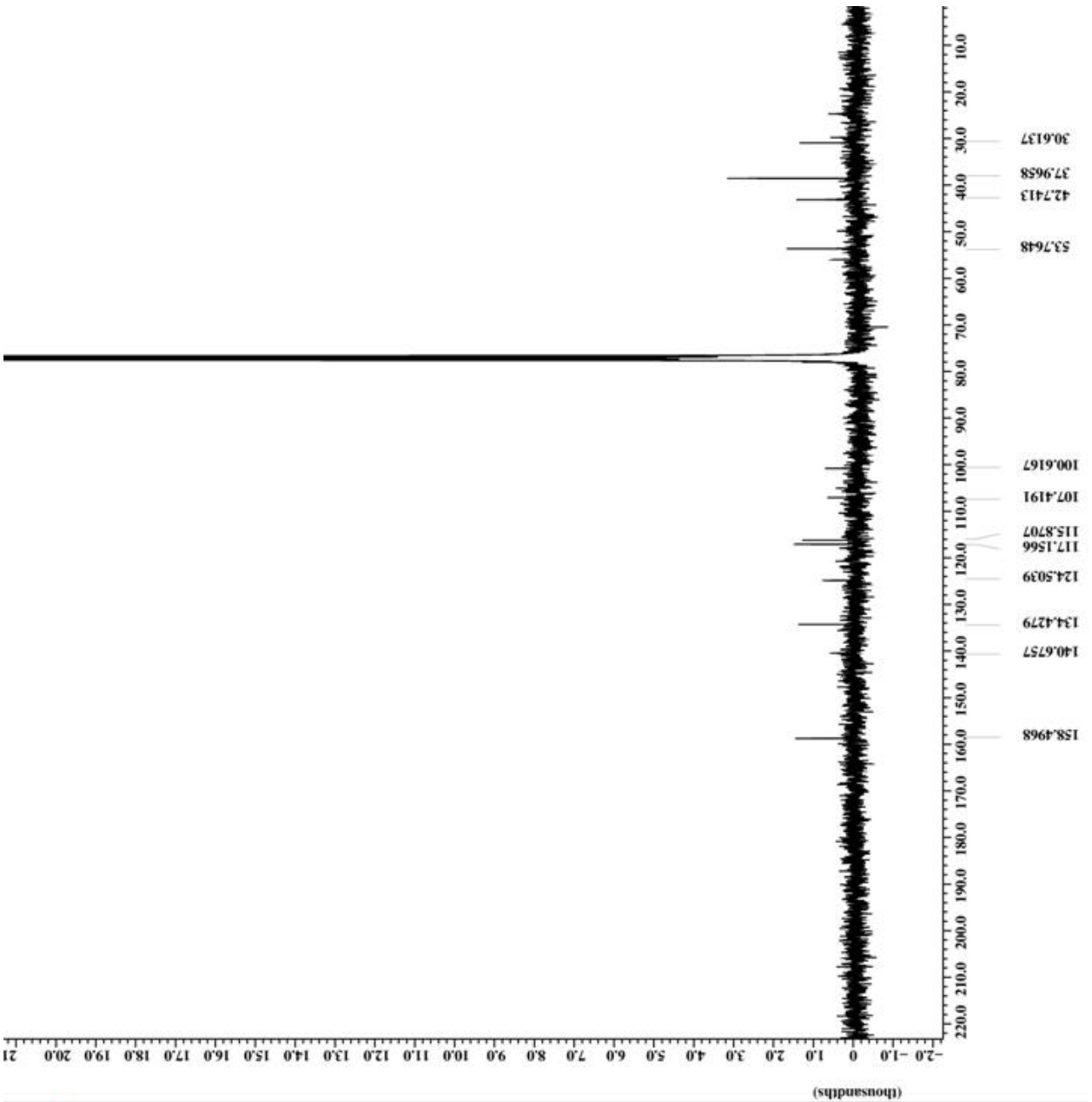
```

filename = sm_VI_46_pure-2.jdf
author = delta
experiment = single_pulse_dec
sample_id = S#788174
solvent = CHLOROFORM-D
acquisition_time = 1-DEC-2009 10:13:11
revision_time = 1-DEC-2009 10:07:54
current_time = 28-MAR-2010 12:21:26

comment = single pulse decouple
ata_format = ID COMPLEX
in_size = 52428
in_title = 13C
in_units = [ppm]
in_dimensions = X
ite = ECX 300
spectrometer = DELTA2_NMR

ield_strength = 7.0586013[T] (300[MHz]
acq_duration = 2.76824064[s]
domain = 13C
freq = 75.56823426[MHz]
offset = 100[ppm]
points = 65536
prescans = 4
resolution = 0.36124027[Hz]
sweep = 23.67424242[kHz]
rr_domain = 1H
rr_freq = 300.52965592[MHz]
rr_offset = 5[ppm]
lipped = FALSE
bd_return = 10
cans = 7625
otal_scans = 7625

_90_width = 9.75[us]
acq_time = 2.76824064[s]
angle = 30[deg]
atn = 8[db]
pulse = 3.25[us]
rr_atn_dec = 25[db]
rr_atn_noe = 25[db]
rr_noise = WALTZ
scoupling = TRUE
nitial_wait = 1[s]
oe_time = TRUE
oe_time = 3[s]
ecvr_gain = 50
elaxation_delay = 3[s]
epetition_time = 5.76824064[s]
emp_get = 23.3[dc]
  
```

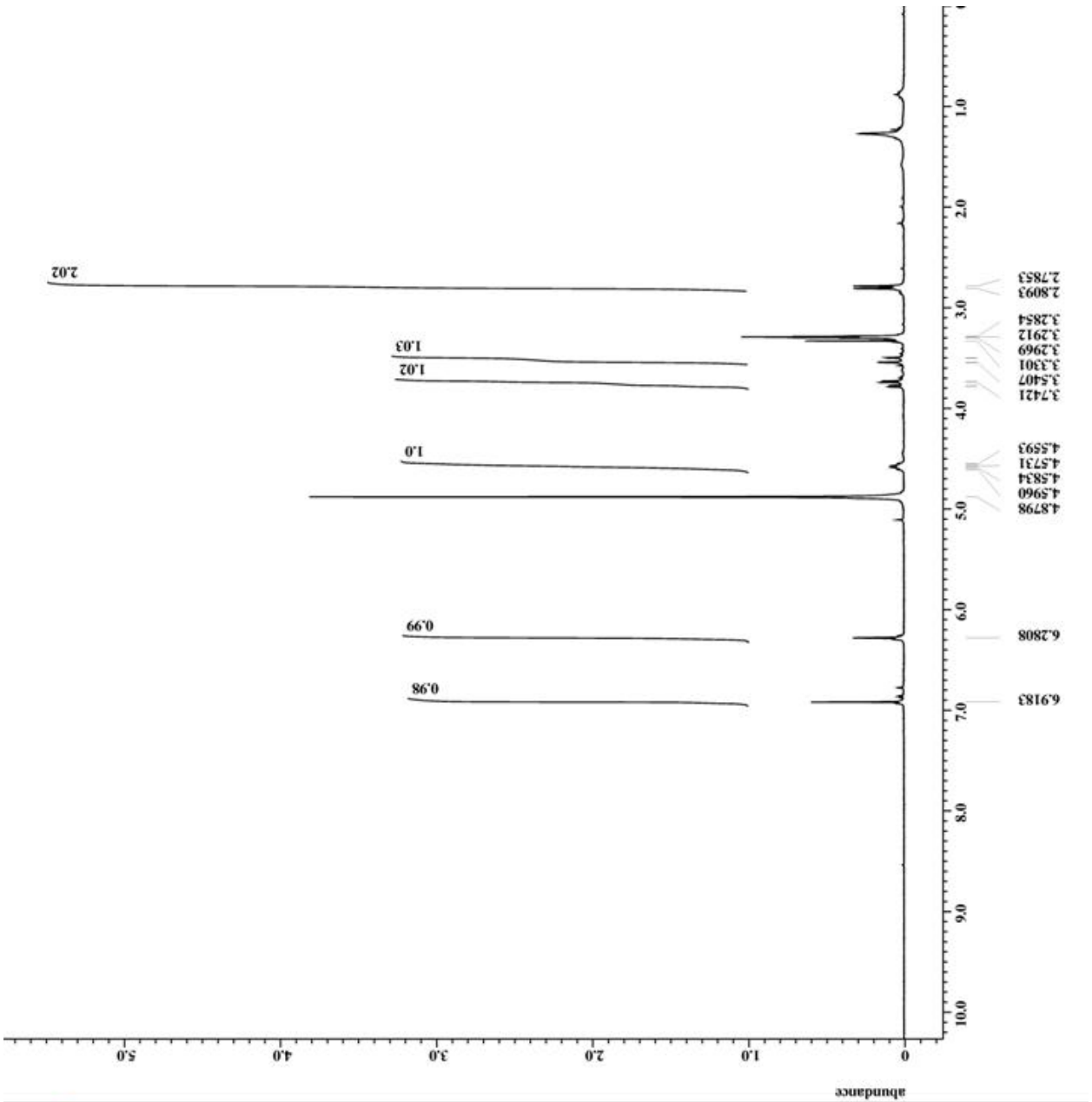
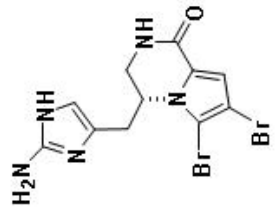


APPENDIX 28
¹H AND ¹³C NMR SPECTRUM OF
(*ent*)-Cyclooroidin (3)



```

filename = sm_VI_87_pure-2.jdf
author = delta
experiment = single_pulse.ex2
sample_id = S#15125
solvent = METHANOL-D3
creation_time = 8-JAN-2010 00:15:58
revision_time = 8-JAN-2010 00:34:38
current_time = 28-MAR-2010 12:25:51
comment = single_pulse
ata_format = ID COMPLEX
in_size = 13107
in_title = 1H
in_units = [ppm]
inmensions = X
ite = ECX 300
spectrometer = DELTA2_NMR
ield_strength = 7.0586013[T] (300[MHz]
_acq_duration = 2.90717696[s]
_domain = 1H
_freq = 300.52965592[MHz]
_offset = 5[ppm]
_points = 16384
_prescans = 0
_resolution = 0.34397621[Hz]
_sweep = 5.63570784[kHz]
rr_domain = 1H
rr_freq = 300.52965592[MHz]
rr_offset = 5[ppm]
ri_domain = 1H
ri_freq = 300.52965592[MHz]
ri_offset = 5[ppm]
lipped = FALSE
od_return = 1
otal_scans = 40
_90_width = 13.01[us]
_acq_time = 2.90717696[s]
_angle = 45[deg]
_atn = 4[db]
_pulse = 6.505[us]
rr_mode = Off
ri_mode = Off
ante_presat = FALSE
nitial_wait = 1[s]
scvr_gain = 46
relaxation_delay = 5[s]
petition_time = 7.90717696[s]
emp_get = 20.8[dc]
  
```





```

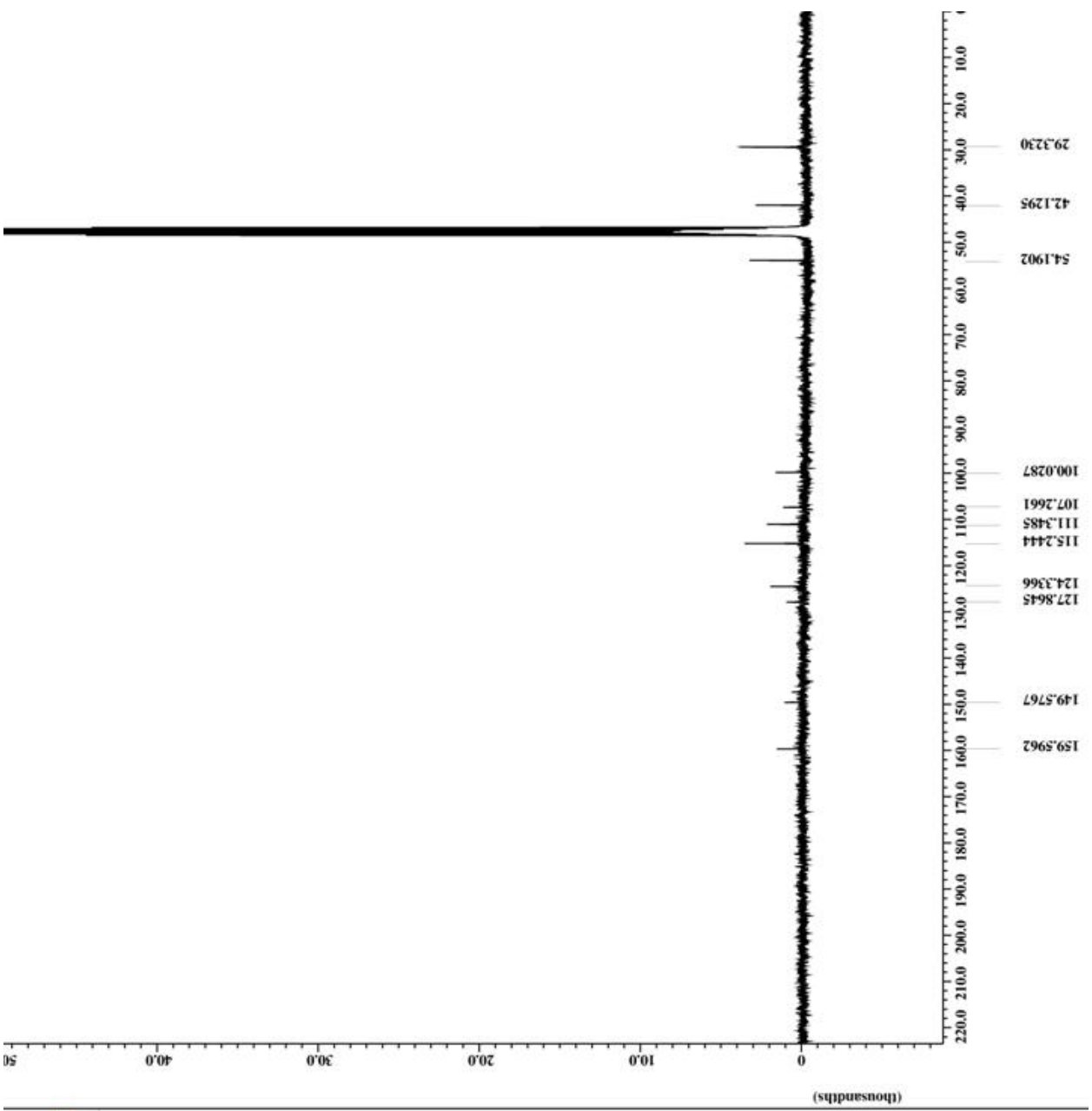
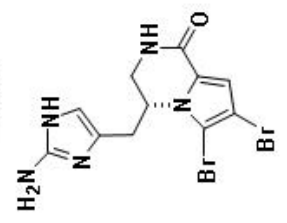
filename = sm_VI_87_pure-2.jdf
author =
experiment = single_pulse_dec
pulse_id = S#18841
solvent = METHANOL-D3
acquisition_time = 8-JAN-2010 13:31:51
revision_time = 8-JAN-2010 13:48:16
current_time = 28-MAR-2010 12:29:40

comment = single pulse decouple
ata_format = ID COMPLEX
in_size = 52428
in_title = 13C
in_units = [ppm]
in_dimensions = X
ite = ECX 300
spectrometer = DELTA2_NMR

field_strength = 7.0586013[T] (300[MHz])
acq_duration = 2.76824064[s]
domain = 13C
freq = 75.56823426[MHz]
offset = 100[ppm]
points = 65536
prescans = 4
resolution = 0.36124027[Hz]
sweep = 23.67424242[kHz]
rr_domain = 1H
rr_freq = 300.52965592[MHz]
rr_offset = 5[ppm]
lipped = TRUE
bd_return = 10
cans = 8272
total_scans = 8272

_90_width = 9.75[us]
acq_time = 2.76824064[s]
angle = 30[deg]
atn = 8[dB]
pulse = 3.25[us]
rr_atn_dec = 25[dB]
rr_atn_noe = 25[dB]
rr_noise = WALTZ
scoupling = TRUE
initial_wait = 1[s]
oe = TRUE
oe_time = 3[s]
ecvr_gain = 50
relaxation_delay = 3[s]
petition_time = 5.76824064[s]
temp_get = 21[degC]

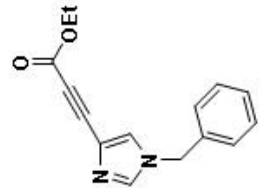
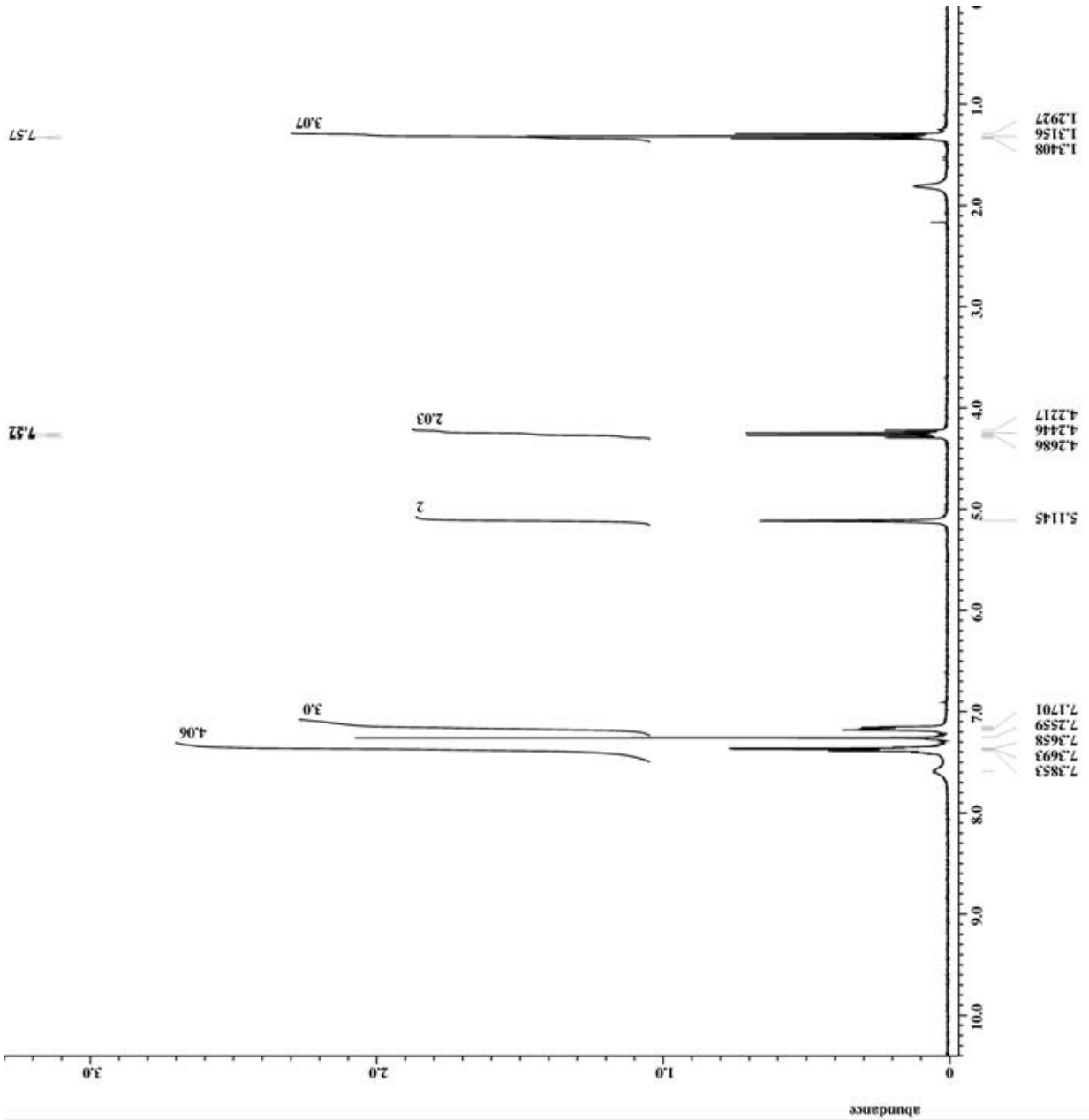
```



APPENDIX 29
¹H AND ¹³C NMR SPECTRUM OF
Ethyl 3-(1-benzyl-1H-imidazol-4-yl)propiolate (224)



```
filenname = sm_III_89_PURE_300-4.
author = delta
experiment = single_pulse.ex2
sample_id = S#576335
solvent = CHLOROFORM-D
reaction_time = 12-OCT-2007 16:24:59
acquisition_time = 28-MAR-2010 12:39:32
current_time = 28-MAR-2010 12:40:42
comment = single_pulse
ata_format = ID COMPLEX
in_size = 13107
in_title = 1H
in_units = [ppm]
inmensions = X
ite = ECX 300
pectrometer = DELTA2_NMR
ield_strength = 7.0586013[T] (300[MHz]
_acq_duration = 2.90717696[s]
_domain = 1H
_freq = 300.52965592[MHz]
_offset = 5[ppm]
_points = 16384
_prescans = 0
_resolution = 0.34397621[Hz]
_sweep = 5.63570784[kHz]
rr_domain = 1H
rr_freq = 300.52965592[MHz]
rr_offset = 5[ppm]
ri_domain = 1H
ri_freq = 300.52965592[MHz]
ri_offset = 5[ppm]
lipped = FALSE
od_return = 1
otal_scans = 24
_90_width = 13.01[us]
_acq_time = 2.90717696[s]
_angle = 45[deg]
_atn = 4[dB]
_pulse = 6.505[us]
rr_mode = Off
ri_mode = Off
ante_presat = FALSE
nitial_wait = 1[s]
scvr_gain = 46
relaxation_delay = 5[s]
epetition_time = 7.90717696[s]
emp_get = 23[dc]
```



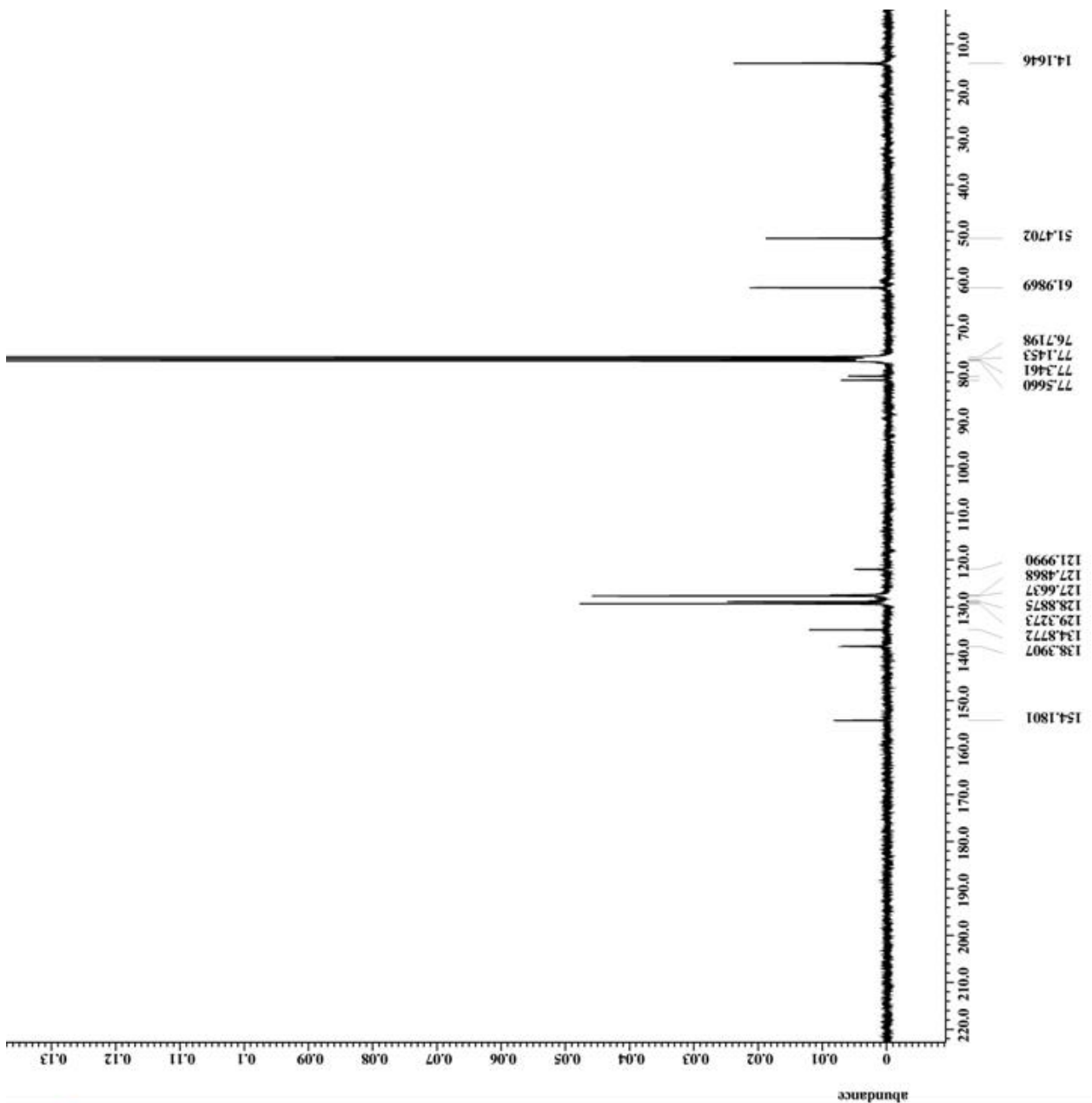
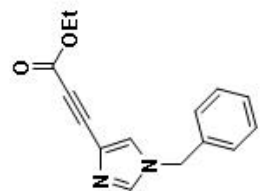


```
filname      = sm_III_88-3_3.jdf
author       =
xperiment    = single_pulse_dec
sample_id    = S8836837
solvent      = CHLOROFORM-D
reaction_time = 11-APR-2009 06:36:41
revision_time = 28-MAR-2010 12:46:18
current_time  = 28-MAR-2010 12:50:11

comment      = single pulse decouple
date_format  = ID_COMPLEX
im_size      = 52428
im_title     = 13C
im_units     = [ppm]
imensions    = X
ite          = ECX 300
spectrometer = DELTA2_NMR

field_strength = 7.0586013[T] (300[MHz]
acq_duration   = 2.76824064[s]
domain         = 13C
freq           = 75.56823426[MHz]
offset         = 100[ppm]
points         = 65536
prescans       = 4
resolution     = 0.36124027[Hz]
sweep         = 23.67424242[KHz]
rr_domain     = 1H
rr_freq       = 300.52965592[MHz]
rr_offset     = 5[ppm]
lipped        = TRUE
bd_return     = 10
cans          = 5000
stal_scans    = 5000

_90_width     = 9.75[us]
acq_time      = 2.76824064[s]
angle         = 30[deg]
atn           = 8[db]
pulse        = 3.25[us]
rr_atn_dec   = 25[db]
rr_atn_noe   = 25[db]
rr_noise     = WALTZ
scoupling    = TRUE
nit1al_wait  = 1[s]
ce           = TRUE
be_time      = 3[s]
ecvr_gain    = 50
relaxation_delay = 3[s]
apitation_time = 5.76824064[s]
emp_get      = 23.3[dc]
```



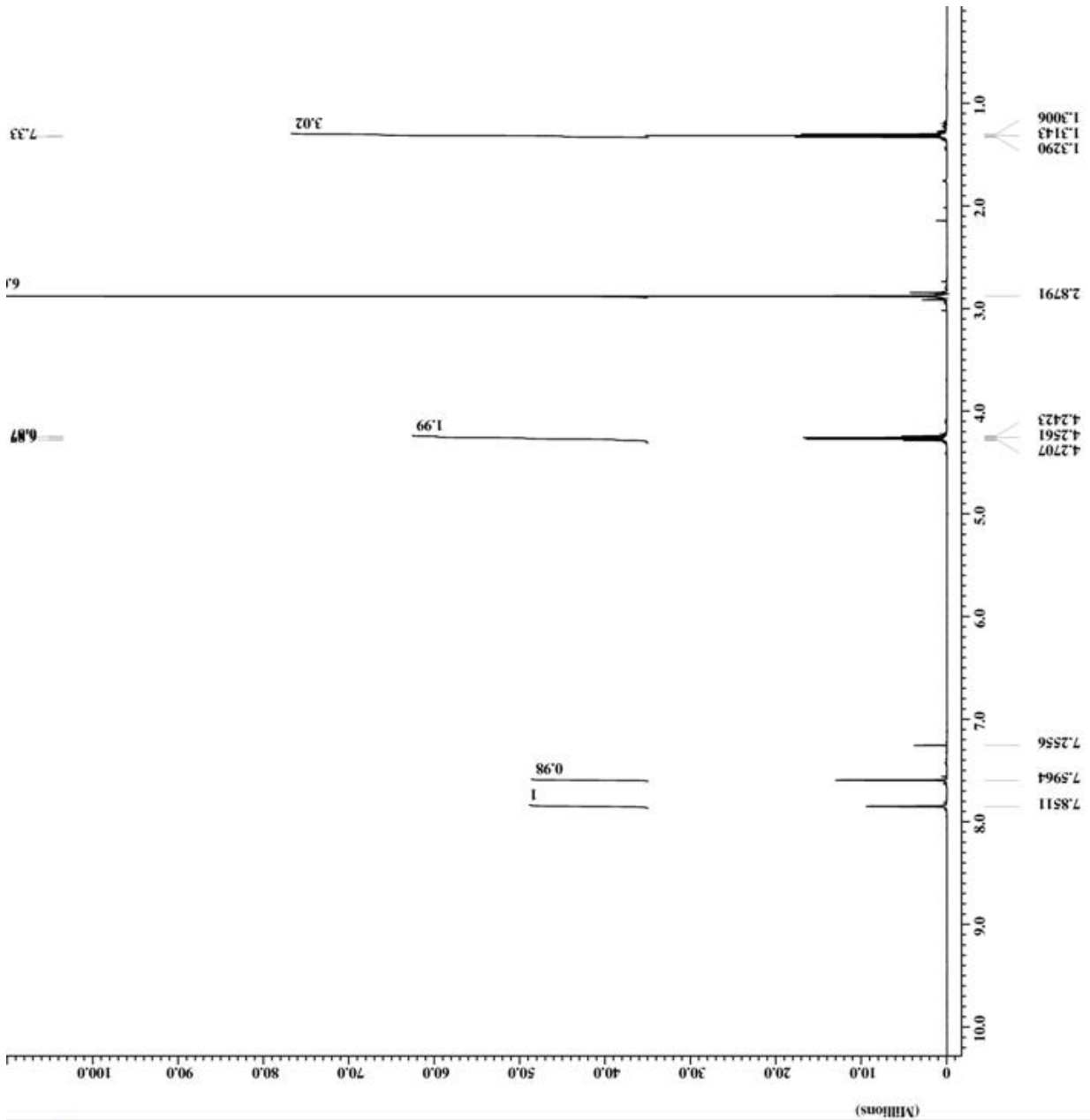
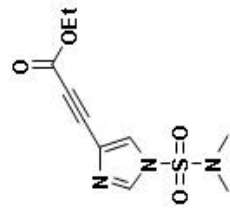
APPENDIX 30
¹H AND ¹³C NMR SPECTRUM OF
Ethyl 3-(1-(N,N-dimethylsulfamoyl)-1H-imidazol-4-yl)propiolate (223)



```

filename = sm_III_dmasproc_ester
author = delta
experiment = single_pulse_exp
sample_id = S#802675
solvent = CHLOROFORM-D
reaction_time = 10-OCT-2007 02:51:25
revision_time = 28-MAR-2010 16:41:33
current_time = 28-MAR-2010 16:42:03
comment = Single Pulse Experiment
data_format = ID COMPLEX
in_size = 16384
in_title = 1H
in_units = [ppm]
instruments = X
ite = Eclipse+ 500
nucleus1 = DELTA_NMR
p1 = 11.7473579[T] (500[MH]
p2 = 2.1823486[s]
p3 = 1H
p4 = 500.15991521[MHz]
p5 = 5[ppm]
p6 = 16384
p7 = 0
p8 = 0.45822189[Hz]
p9 = 7.50750751[kHz]
p10 = FALSE
p11 = 1
p12 = 7
p13 = 7
p14 = 18.5[us]
p15 = 2.1823486[s]
p16 = 45[deg]
p17 = 7.25[us]
p18 = 1[s]
p19 = 3[us]
p20 = 17
p21 = 4[s]
p22 = 25.2[dc]
p23 = 2[us]

```





```

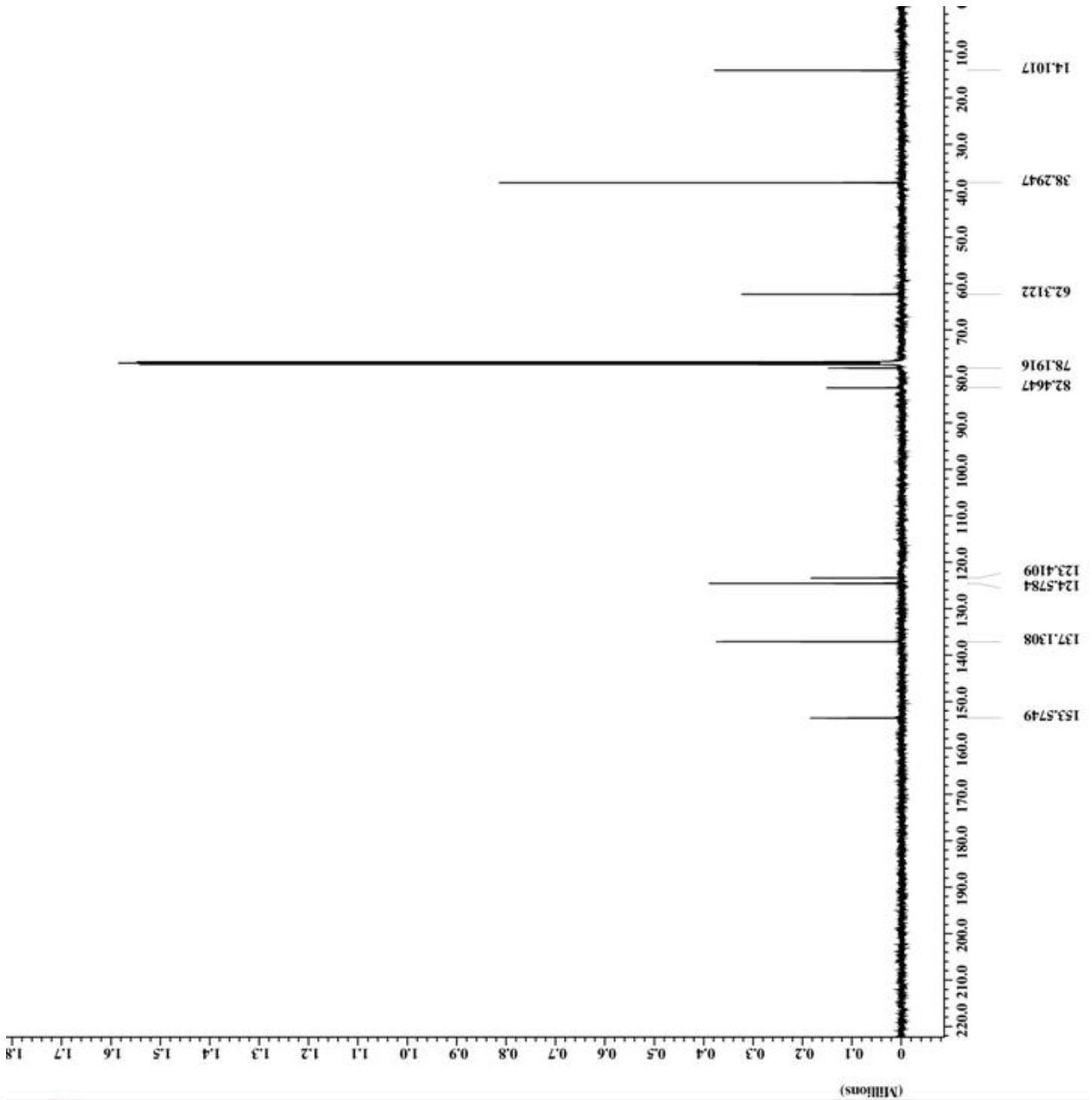
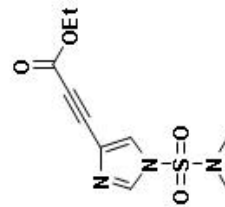
filename = sm_III_dmasproc_ester
author = delta
experiment = single_pulse_dec
sample_id = S#804229
solvent = CHLOROFORM-D
acquisition_time = 10-OCT-2007 04:18:23
revision_time = 10-OCT-2007 00:04:53
current_time = 28-MAR-2010 16:45:21

comment = single pulse decouple
ata_format = ID COMPLEX
im_size = 65536
im_title = 13C
im_units = [ppm]
imensions = X
ite = Eclipse+ 500
spectrometer = DELTA_NMR

field_strength = 11.747379[T] (500[MH
-acq_duration = 2.0840448[s]
-domain = 13C
-freq = 125.76529768[MHz]
-offset = 100[ppm]
-points = 65536
-prescans = 4
-resolution = 0.47983613[Hz]
-sweep = 31.44654088[kHz]
-tr_domain = 1H
-tr_freq = 500.15991521[MHz]
-tr_offset = 5[ppm]
-tipped = TRUE
-bd_return = 10
-cans = 1000
-etal_scans = 1000

_90_width = 14.2[us]
-acq_time = 2.0840448[s]
-angle = 30[deg]
-pulse = 4.73333333[us]
-nitral_wait = 1[s]
-pe_time = 1[s]
-base_preset = 3[us]
-ecvr_gain = 27
-relaxation_delay = 2[s]
-emp_get = 27[dc]
-nblank_time = 2[us]

```

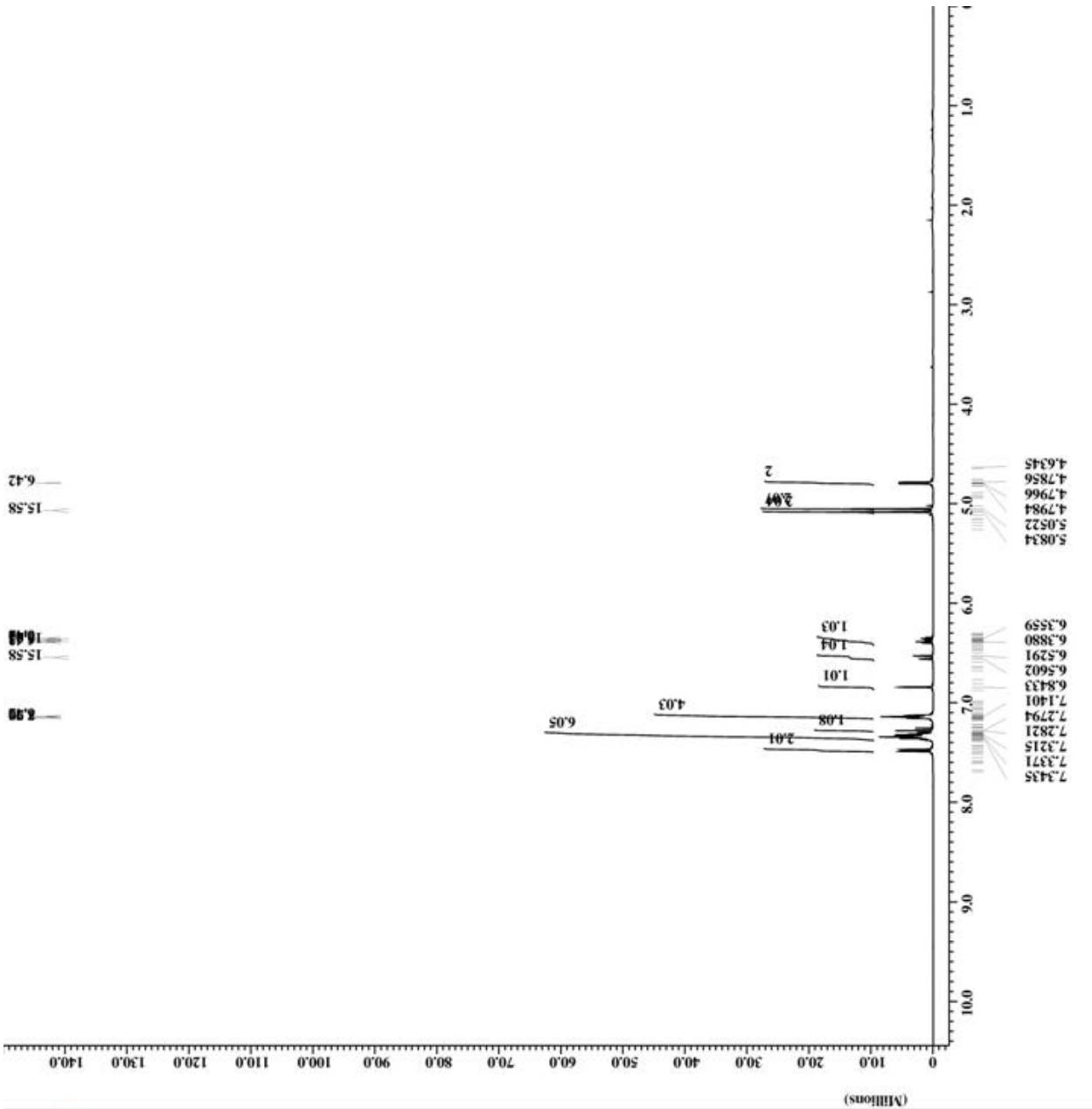
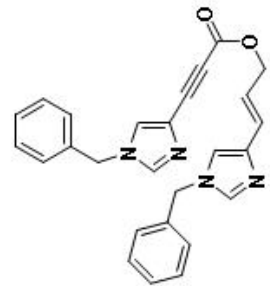


APPENDIX 31
¹H AND ¹³C NMR SPECTRUM OF
(*E*)-3-(1-benzyl-1H-imidazol-4-yl)allyl 3-(1-benzyl-1H-imidazol-4-yl)propiolate
(228)



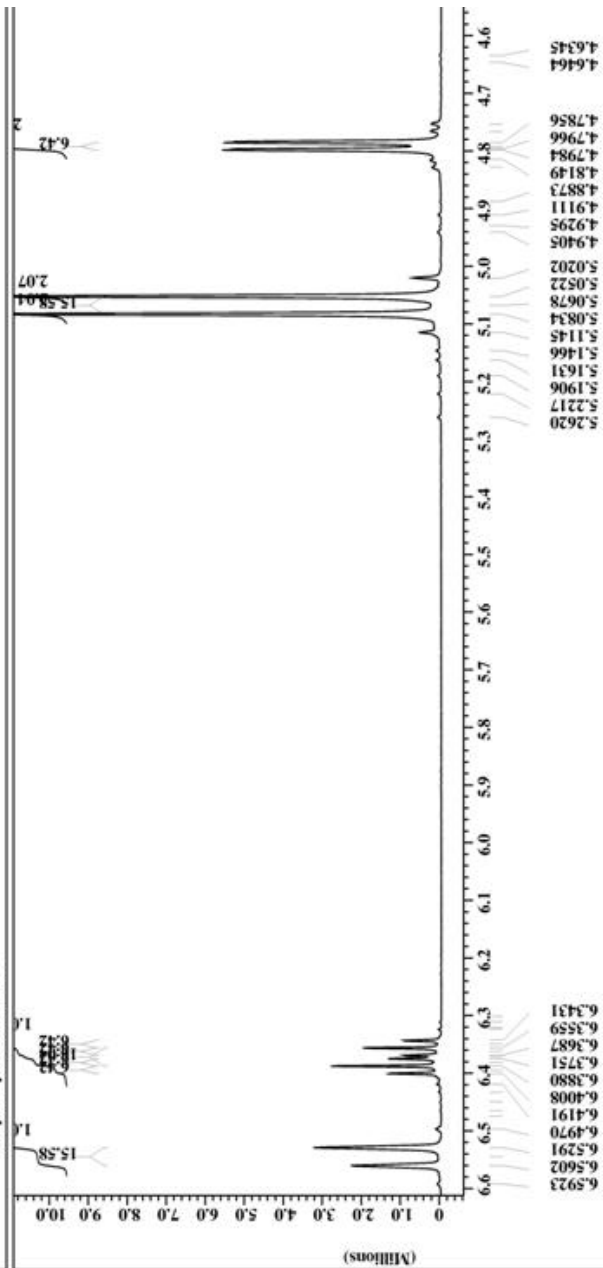
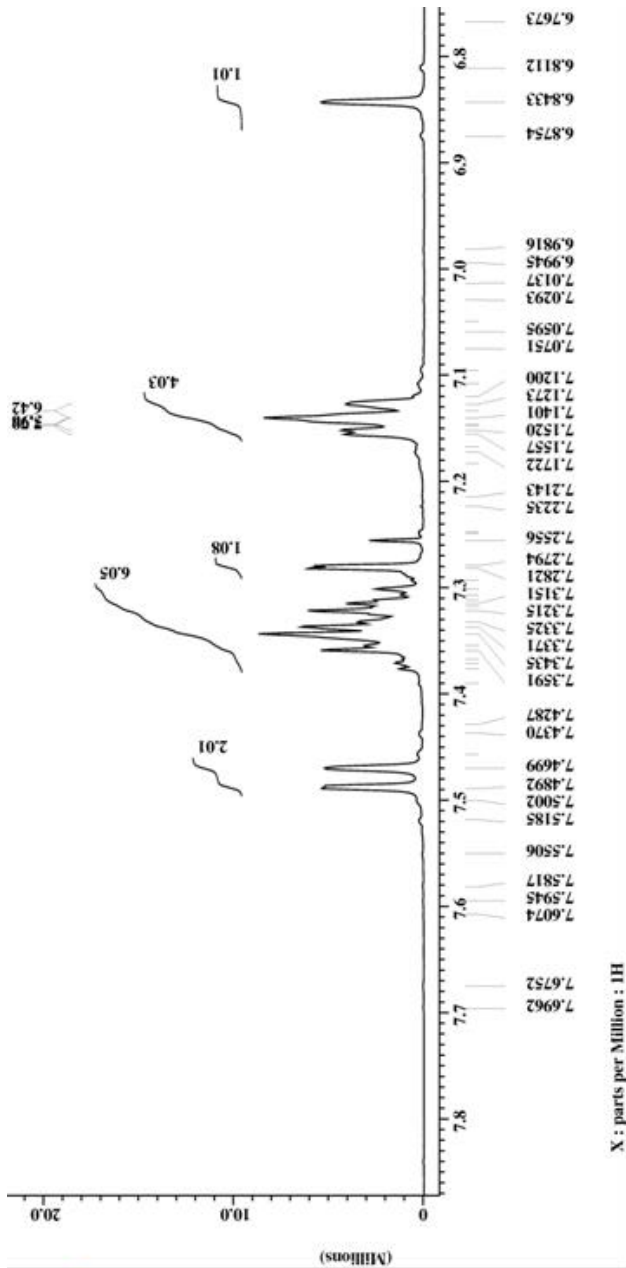
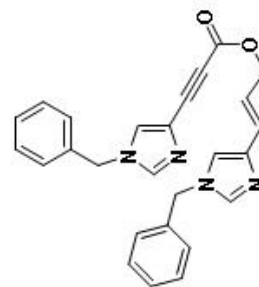
```

filename = sm_III_99_pure_500_2-
author = delta
experiment = single_pulse_exp
sample_id = S#448020
solvent = CHLOROFORM-D
reaction_time = 17-OCT-2007 17:05:04
revision_time = 28-MAR-2010 16:32:37
current_time = 28-MAR-2010 16:54:15
comment = Single Pulse Experiment
data_format = ID COMPLEX
in_size = 16384
in_title = 1H
in_units = [ppm]
instruments = X
pulse_program = Eclipse+ 500
preamplifier = DELTA_NMR
field_strength = 11.74737579[T] (500[MH]
acq_duration = 2.1823486[s]
domain = 1H
freq = 500.15991521[MHz]
offset = 5[ppm]
points = 16384
prescans = 0
resolution = 0.45822189[Hz]
sweep = 7.50750751[kHz]
tipped = FALSE
od_return = 1
cans = 8
total_scans = 8
_90_width = 18.5[us]
acq_time = 2.1823486[s]
angle = 45[deg]
pulse = 7.25[us]
nit1_wait = 1[s]
base_preset = 3[us]
ecvr_gain = 18
relaxation_delay = 4[s]
emp_get = 25.2[dc]
nblank_time = 2[us]
  
```



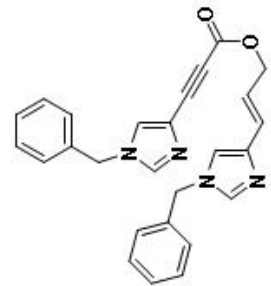
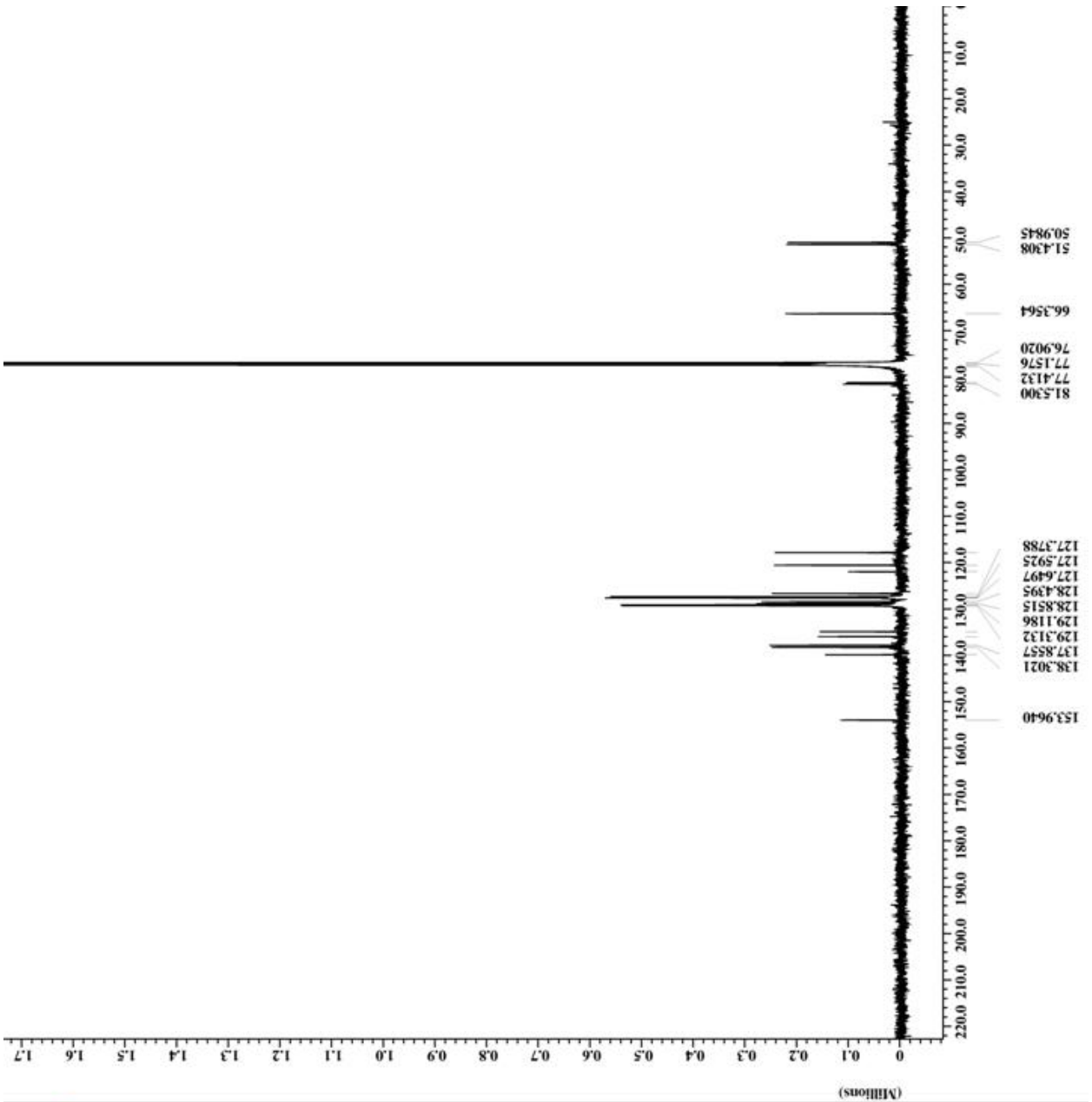


```
filename = sm_iii_99_pure_500_2-
author = delta
experiment = single_pulse_exp
sample_id = S#448020
solvent = CHLOROFORM-D
reaction_time = 17-OCT-2007 17:05:04
revision_time = 28-MAR-2010 16:32:37
current_time = 28-MAR-2010 16:58:03
comment = Single Pulse Experiment
ata_format = ID COMPLEX
im_size = 16384
im_title = 1H
im_units = [ppm]
lms = X
lms_units = Eclipse+ 500
lms_title = DELTA_NMR
spectrometer =
field_strength = 11.7473579[T] (500[MH]
acq_duration = 2.1823488[s]
domain =
freq = 500.15991521[MHz]
points = 16384
prescans = 0
resolution = 0.45822189[Hz]
sweep = 7.50750751[kHz]
lipped = FALSE
od_return = 1
cans = 8
dcal_scans = 8
_90_width = 18.5[us]
_acq_time = 2.1823488[s]
_angle = 45[deg]
_pulse = 7.25[us]
_nitai_wait = 1[s]
_base_preset = 3[us]
_scvr_gain = 18
relaxation_delay = 4[s]
emp_get = 25.2[dc]
nblank_time = 2[us]
```





```
filename = sm_III_99_PURE_500-2.
author = delta
experiment = single_pulse_dec
sample_id = S943280
solvent = CHLOROFORM-D
acquisition_time = 18-OCT-2007 22:33:52
revision_time = 19-OCT-2007 10:18:14
current_time = 28-MAR-2010 17:01:01
comment = single pulse decouple
data_format = ID COMPLEX
im_size = 65536
im_title = 13C
im_units = [ppm]
imensions = X
ite = Eclipse+ 500
spectrometer = DELTA_NMR
field_strength = 11.747379[T] (500[MH
acq_duration = 2.0840448[s]
domain = 13C
freq = 125.76529768[MHz]
offset = 100[ppm]
points = 65536
prescans = 4
resolution = 0.47983613[Hz]
sweep = 31.44654088[kHz]
tr_domain = 1H
tr_freq = 500.15991521[MHz]
tr_offset = 5[ppm]
lipped = TRUE
bd_return = 10
cans = 2000
otal_scans = 2000
_90_width = 14.2[us]
_acq_time = 2.0840448[s]
_angle = 30[deg]
_pulse = 4.73333333[us]
_nitral_wait = 1[s]
_pe_time = 1[s]
_base_preset = 3[us]
_scvr_gain = 30
_relaxation_delay = 2[s]
emp_get = 27.2[dC]
nblank_time = 2[us]
```





```

filename = sm_III_99_PURE_500-2.
author = delta
experiment = single_pulse_dec
sample_id = S9543280
solvent = CHLOROFORM-D
acquisition_time = 18-OCT-2007 22:33:52
revision_time = 19-OCT-2007 10:18:14
current_time = 28-MAR-2010 17:03:17

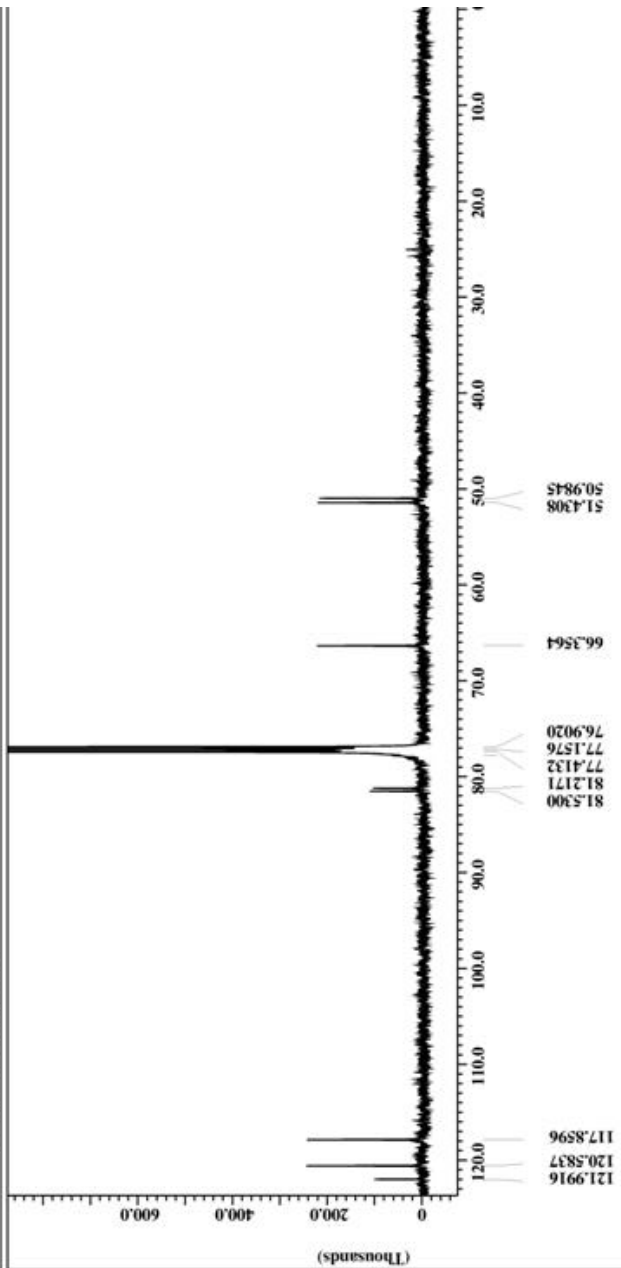
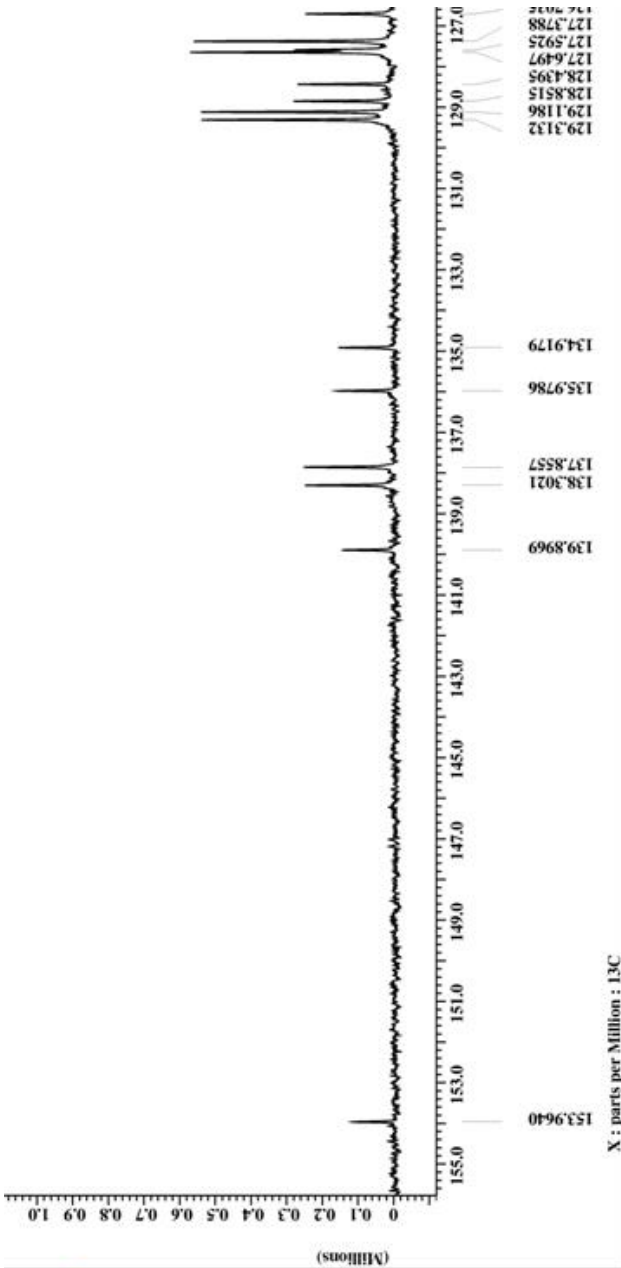
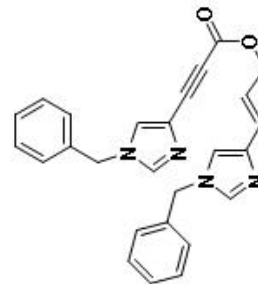
comment = single pulse decouple
          = ID COMPLEX
          = 65536
          = 13C
          = [ppm]
          = X
          = Eclipse+ 500
          = DELTA_NMR

p1 = 13C
p2 = 13C
p3 = 13C
p4 = 13C
p5 = 13C
p6 = 13C
p7 = 13C
p8 = 13C
p9 = 13C
p10 = 13C
p11 = 13C
p12 = 13C
p13 = 13C
p14 = 13C
p15 = 13C
p16 = 13C
p17 = 13C
p18 = 13C
p19 = 13C
p20 = 13C
p21 = 13C
p22 = 13C
p23 = 13C
p24 = 13C
p25 = 13C
p26 = 13C
p27 = 13C
p28 = 13C
p29 = 13C
p30 = 13C
p31 = 13C
p32 = 13C
p33 = 13C
p34 = 13C
p35 = 13C
p36 = 13C
p37 = 13C
p38 = 13C
p39 = 13C
p40 = 13C
p41 = 13C
p42 = 13C
p43 = 13C
p44 = 13C
p45 = 13C
p46 = 13C
p47 = 13C
p48 = 13C
p49 = 13C
p50 = 13C
p51 = 13C
p52 = 13C
p53 = 13C
p54 = 13C
p55 = 13C
p56 = 13C
p57 = 13C
p58 = 13C
p59 = 13C
p60 = 13C
p61 = 13C
p62 = 13C
p63 = 13C
p64 = 13C
p65 = 13C
p66 = 13C
p67 = 13C
p68 = 13C
p69 = 13C
p70 = 13C
p71 = 13C
p72 = 13C
p73 = 13C
p74 = 13C
p75 = 13C
p76 = 13C
p77 = 13C
p78 = 13C
p79 = 13C
p80 = 13C
p81 = 13C
p82 = 13C
p83 = 13C
p84 = 13C
p85 = 13C
p86 = 13C
p87 = 13C
p88 = 13C
p89 = 13C
p90 = 13C
p91 = 13C
p92 = 13C
p93 = 13C
p94 = 13C
p95 = 13C
p96 = 13C
p97 = 13C
p98 = 13C
p99 = 13C
p100 = 13C

field_strength = 11.747379[T] (500[MH
acq_duration = 2.0840448[s]
domain = 13C
freq = 125.76529768[MHz]
offset = 100[ppm]
points = 65536
prescans = 4
resolution = 0.47983613[Hz]
sweep = 31.44654088[kHz]
tr_domain = 1H
tr_freq = 500.15991521[MHz]
tr_offset = 5[ppm]
tipped = TRUE
bd_return = 10
cans = 2000
total_scans = 2000

_90_width = 14.2[us]
acq_time = 2.0840448[s]
angle = 30[deg]
pulse = 4.73333333[us]
nitai_wait = 1[s]
pe_time = 1[s]
base_preset = 3[us]
scvr_gain = 30
relaxation_delay = 2[s]
emp_get = 27.2[dc]
nblank_time = 2[us]

```



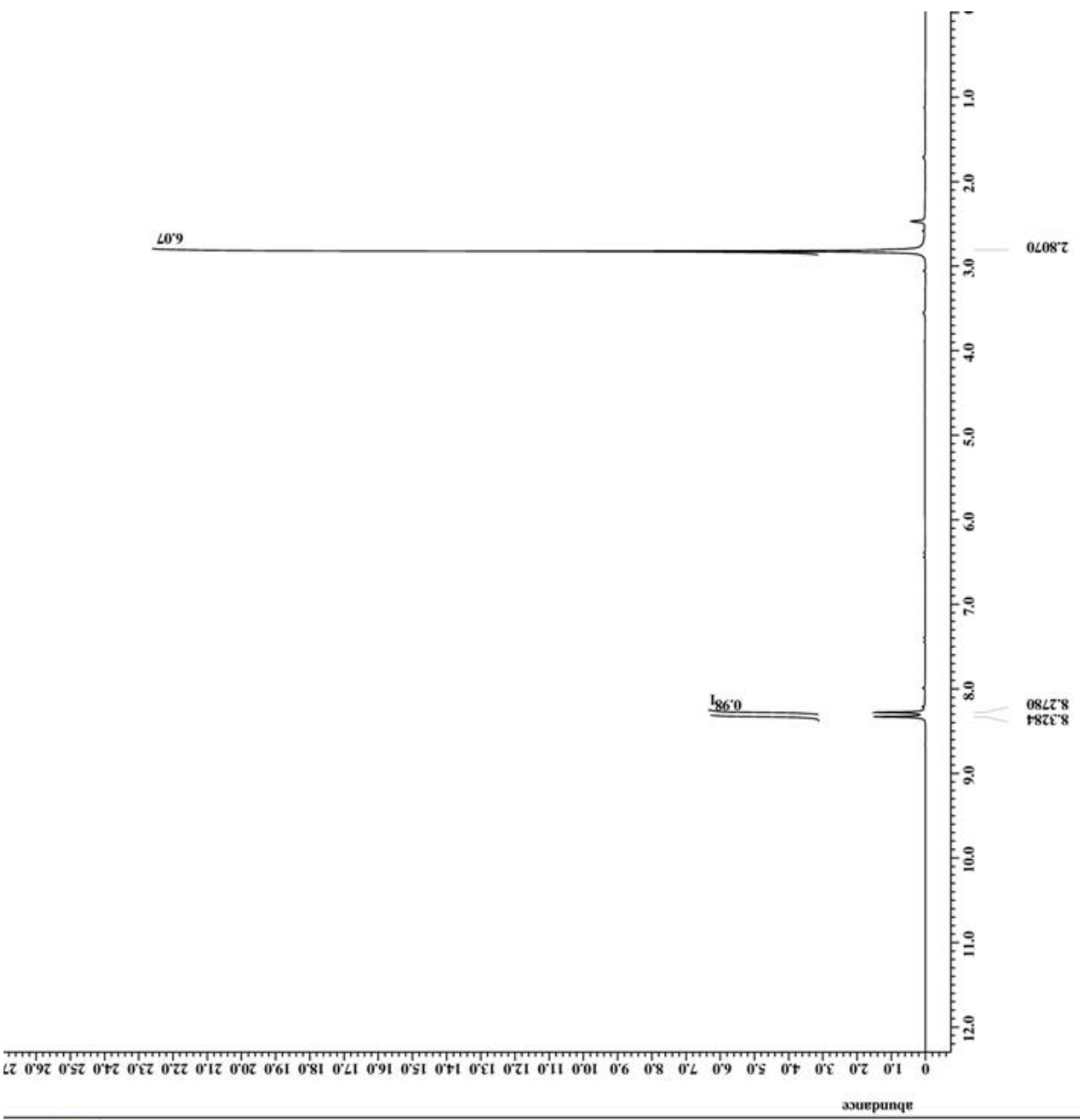
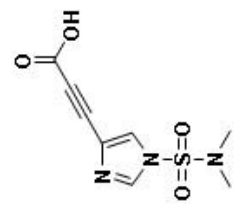
APPENDIX 32
¹H AND ¹³C NMR SPECTRUM OF
3-(1-(N,N-dimethylsulfamoyl)-1H-imidazol-4-yl)propionic acid (225)



```

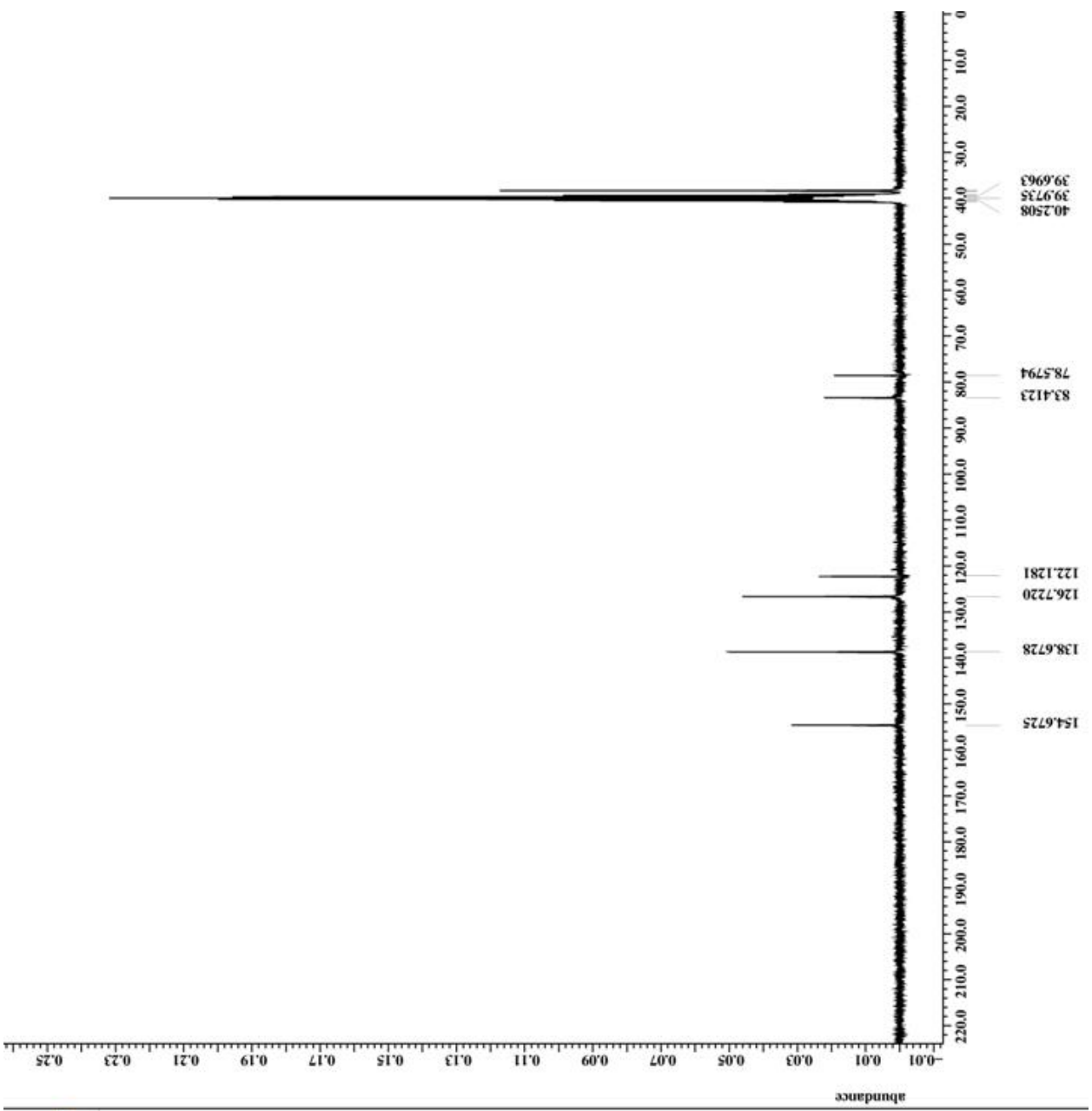
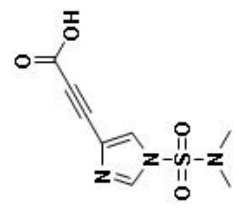
filename = sm_V_dmas_acid-4.jdf
author = delta
experiment = single_pulse.ex2
sample_id = S#679031
solvent = DMSO-d6
creation_time = 7-APR-2009 18:15:54
revision_time = 28-MAR-2010 17:14:27
current_time = 28-MAR-2010 17:22:01
comment = single_pulse
ata_format = ID COMPLEX
in_size = 13107
in_title = 1H
in_units = [ppm]
in_dimensions = X
ite = ECX 300
spectrometer = DELTA2_NMR
field_strength = 7.0586013[T] (300[MHz])
acq_duration = 2.90717696[s]
domain = 1H
freq = 300.52965592[MHz]
offset = 5[ppm]
points = 16384
prescans = 0
resolution = 0.34397621[Hz]
sweep = 5.63570784[kHz]
tr_domain = 1H
tr_freq = 300.52965592[MHz]
tr_offset = 5[ppm]
ri_domain = 1H
ri_freq = 300.52965592[MHz]
ri_offset = 5[ppm]
lipped = FALSE
od_return = 1
cans = 23
atal_scans = 23
_90_width = 13.01[us]
acq_time = 2.90717696[s]
angle = 45[deg]
atn = 4[dB]
pulse = 6.505[us]
tr_mode = Off
ri_mode = Off
ante_presat = FALSE
nitial_wait = 1[s]
ecvr_gain = 38
relaxation_delay = 5[s]
petition_time = 7.90717696[s]
emp_get = 23.2[dc]

```





```
filename = sm_v_dmas_acid-3.jdf
author = delta
experiment = single_pulse_dec
sample_id = S9681372
solvent = DMSO-D6
creation_time = 7-APR-2009 19:52:38
revision_time = 8-APR-2009 11:41:20
current_time = 28-MAR-2010 17:26:19
comment = single pulse decouple
ata_format = ID REAL
in_size = 52428
in_title = 13C
in_units = [ppm]
inmensions = X
ite = ECX 300
spectrometer = DELTA2_NMR
keld_strength = 7.0586013[T] (300[MHz]
acq_duration = 2.76824064[s]
domain = 13C
freq = 75.56823426[MHz]
offset = 100[ppm]
points = 65536
prescans = 4
resolution = 0.36124027[Hz]
sweep = 23.67424242[kHz]
tr_domain = 1H
tr_freq = 300.52965592[MHz]
tr_offset = 5[ppm]
lipped = FALSE
bd_return = 10
cans = 1000
atal_scans = 1000
_90_width = 9.75[us]
acq_time = 2.76824064[s]
angle = 30[deg]
atn = 8[dB]
pulse = 3.25[us]
tr_atn_dec = 25[dB]
tr_atn_noe = 25[dB]
tr_noise = WALTZ
scoupling = TRUE
nitral_wait = 1[s]
be_time = TRUE
be_time = 3[s]
scvr_gain = 50
relaxation_delay = 3[s]
spetition_time = 5.76824064[s]
emp_get = 23.2[dc]
```



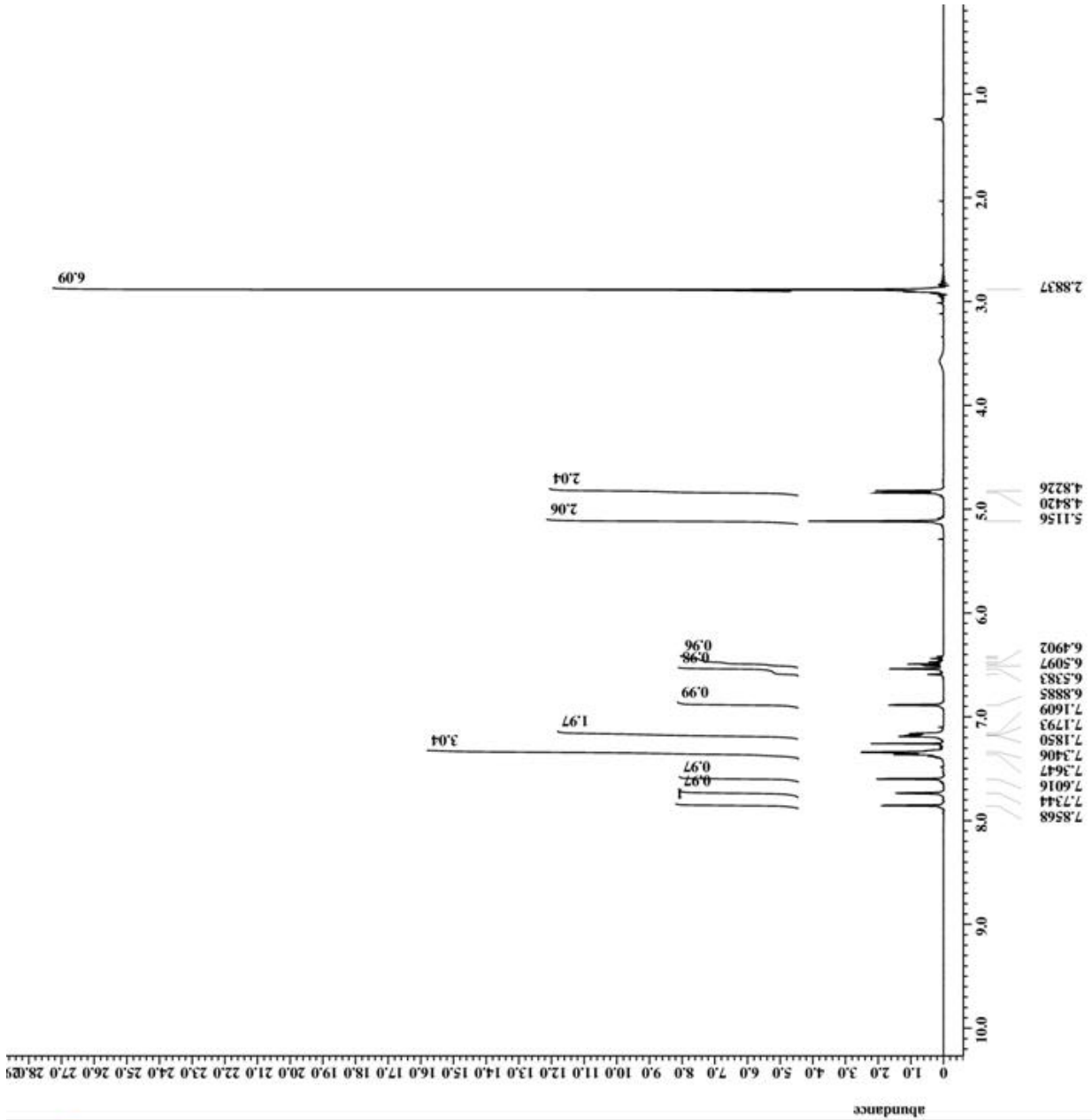
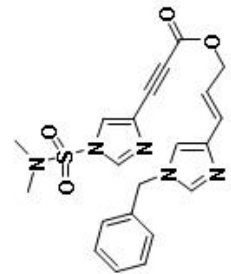
APPENDIX 33
¹H AND ¹³C NMR SPECTRUM OF
(*E*)-3-(1-benzyl-1H-imidazol-4-yl)allyl-3-(1-(N,N-dimethylsulfamoyl)-1H-
imidazol-4-yl)propiolate (227)



```

filename = sm_IV_64_pure-6.jdf
author = delta
experiment = single_pulse.ex2
sample_id = S88459
solvent = CHLOROFORM-D
creation_time = 6-MAY-2008 00:27:25
revision_time = 28-MAR-2010 17:38:25
current_time = 28-MAR-2010 17:39:10
comment =
ata_format = single_pulse
id = ID_REAL
in_size = 13107
in_title = 1H
in_units = [ppm]
inmensions = X
ite = ECX 300
spectrometer = DELTA2_NMR
ield_strength = 7.0586013[T] (3000[MHz]
_acq_duration = 2.90717696[s]
_domain = 1H
_freq = 300.52965592[MHz]
_offset = 5[ppm]
_points = 16384
_prescans = 0
_resolution = 0.34397621[Hz]
_sweep = 5.63570784[kHz]
tr_domain = 1H
tr_freq = 300.52965592[MHz]
tr_offset = 5[ppm]
ri_domain = 1H
ri_freq = 300.52965592[MHz]
ri_offset = 5[ppm]
lipped = FALSE
od_return = 1
otal_scans = 17
_90_width = 13.01[us]
_acq_time = 2.90717696[s]
_angle = 45[deg]
_atn = 4[db]
_pulse = 6.505[us]
tr_mode = Off
ri_mode = Off
ante_presat = FALSE
nitial_wait = 1[s]
scvr_gain = 50
relaxation_delay = 5[s]
petition_time = 7.90717696[s]
emp_get = 23.1[dc]

```



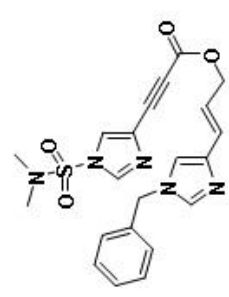
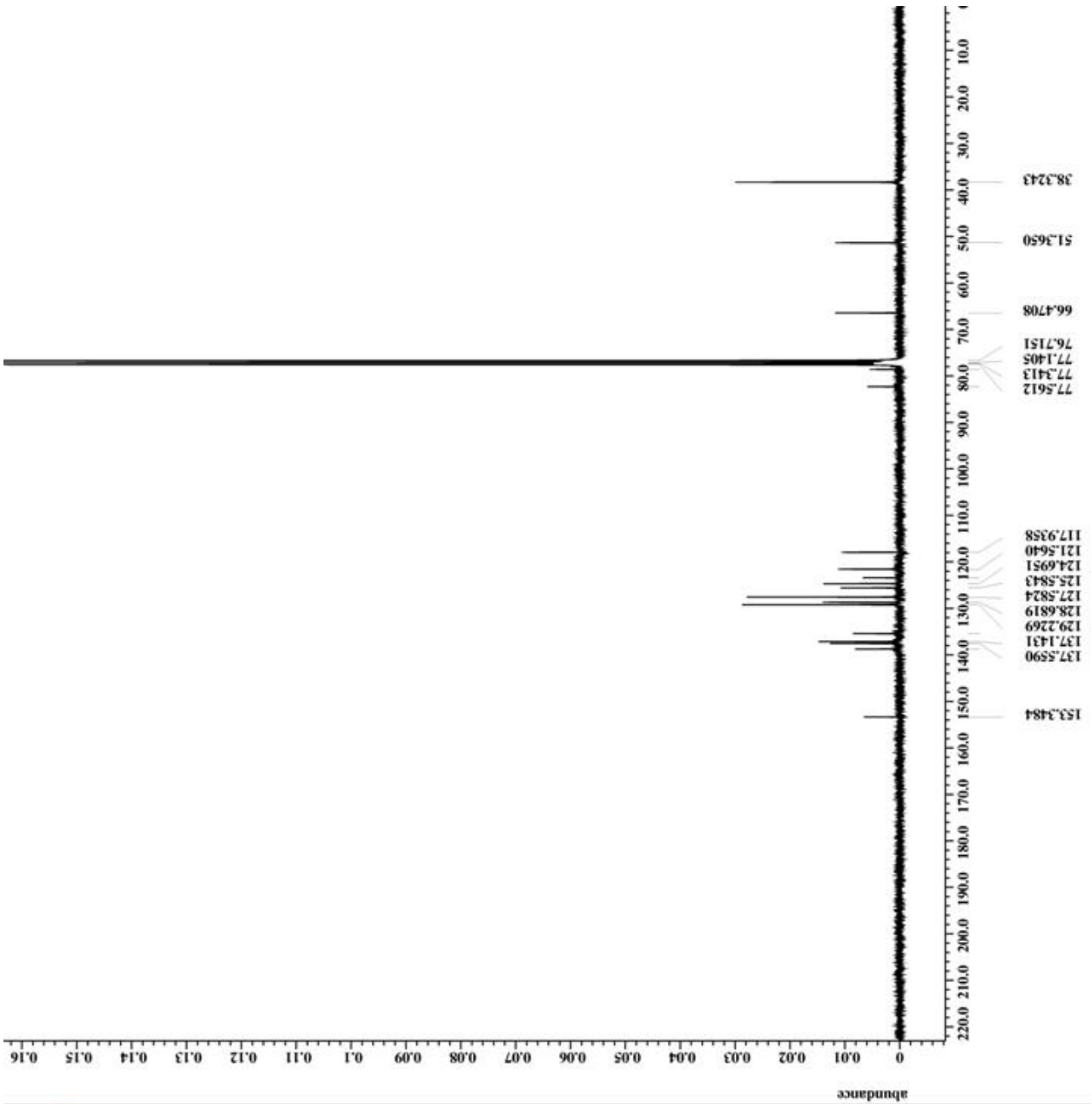


```
filename = sm_IV_64_pure-3.jdf
author = delta
experiment = single_pulse_dec
sample_id = S#10441
solvent = CHLOROFORM-D
reaction_time = 6-MAY-2008 08:29:25
revision_time = 6-MAY-2008 11:49:19
current_time = 28-MAR-2010 17:49:18

comment = single pulse decouple
ata_format = ID REAL
in_size = 52428
in_title = 13C
in_units = [ppm]
inmensions = X
ite = ECX 300
spectrometer = DELTA2_NMR

ield_strength = 7.0586013[T] (300[MHz]
acq_duration = 2.76824064[s]
domain = 13C
freq = 75.56823426[MHz]
offset = 100[ppm]
points = 65536
prescans = 4
resolution = 0.36124027[Hz]
sweep = 23.67424242[kHz]
tr_domain = 1H
tr_freq = 300.52965592[MHz]
tr_offset = 5[ppm]
lipped = TRUE
bd_return = 10
cans = 5000
otal_scans = 5000

_90_width = 9.75[us]
acq_time = 2.76824064[s]
angle = 30[deg]
atn = 8[db]
pulse = 3.25[us]
tr_atn_dec = 25[db]
tr_atn_noe = 25[db]
tr_noise = WALTZ
scoupling = TRUE
nitial_wait = 1[s]
oe_time = TRUE
oe_time = 3[s]
ecvr_gain = 50
elaxation_delay = 3[s]
epetition_time = 5.76824064[s]
emp_get = 23.3[dc]
```





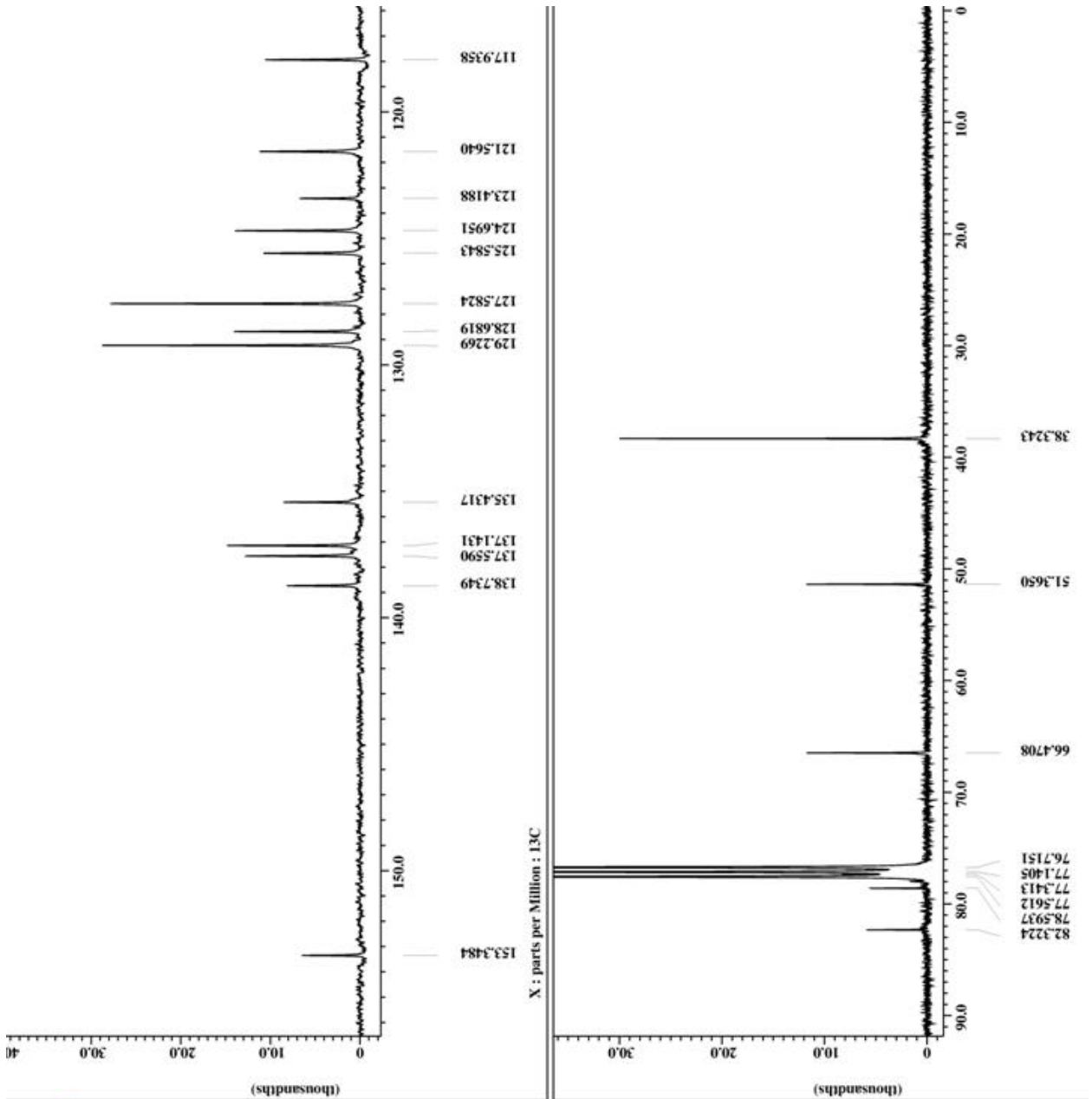
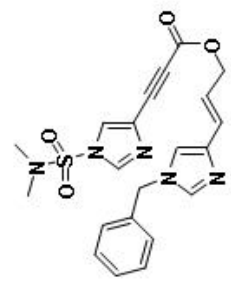
```

filename = sm_IV_64_pure-3.jdf
author = delta
experiment = single_pulse_dec
sample_id = S#10441
solvent = CHLOROFORM-D
reaction_time = 6-MAY-2008 08:29:25
revision_time = 6-MAY-2008 11:49:19
current_time = 28-MAR-2010 17:52:01

comment = single pulse decouple
ata_format = ID REAL
in_size = 52428
in_title = 13C
in_units = [ppm]
inmensions = X
ite = ECX 300
spectrometer = DELTA2_NMR

ield_strength = 7.0586013[T] (300 [MHz]
-acq_duration = 2.76824064[s]
-domain = 13C
-freq = 75.56823426[MHz]
-offset = 100[ppm]
-points = 65536
-prescans = 4
-resolution = 0.36124027[Hz]
-sweep = 23.67424242[kHz]
-tr_domain = 1H
-tr_freq = 300.52965592[MHz]
-tr_offset = 5[ppm]
-lipped = TRUE
-bd_return = 10
-cans = 5000
-otal_scans = 5000

_90_width = 9.75[us]
-acq_time = 2.76824064[s]
-angle = 30[deg]
-atn = 8[dB]
-pulse = 3.25[us]
-tr_atn_dec = 25[dB]
-tr_atn_noe = 25[dB]
-tr_noise = WALTZ
-coupling = TRUE
-nitial_wait = 1[s]
-be_time = TRUE
-be_time = 3[s]
-scvr_gain = 50
-relaxation_delay = 3[s]
-epetition_time = 5.76824064[s]
-emp_get = 23.3[dc]
  
```



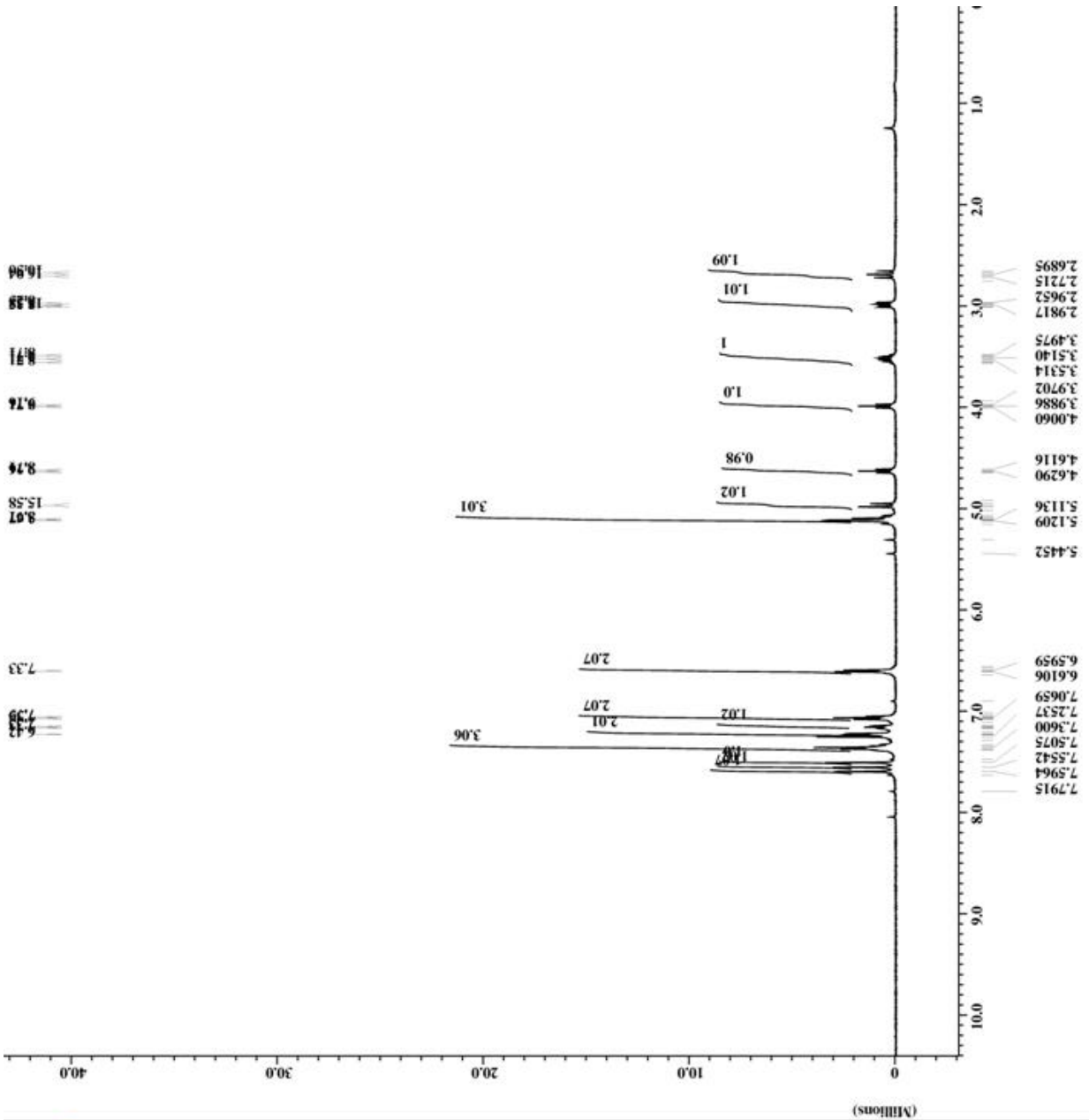
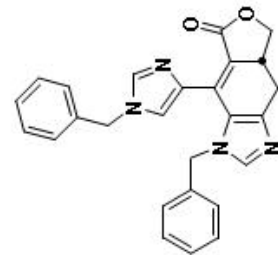
APPENDIX 34
¹H AND ¹³C NMR SPECTRUM OF
(*R*^{*})-1-benzyl-8-(1-benzyl-1H-imidazol-4-yl)-4a,5-dihydro-1H-
isobenzofuro[5,6-d]imidazol-7(4H)-one (230)



```

filename = sm_III_100_pure_2-4.j
author = delta
experiment = single_pulse_exp
sample_id = S#10791
solvent = CHLOROFORM-D
reaction_time = 20-JAN-2008 05:45:53
revision_time = 28-MAR-2010 18:01:02
current_time = 28-MAR-2010 18:01:32
comment = Single Pulse Experiment
data_format = ID COMPLEX
data_size = 16384
in_title = 1H
in_units = [ppm]
lms = X
lms_units = Eclipse+ 500
lms_preamplifier = DELTA_NMR
lms_preamplifier_gain = 500.15991521 [MHz]
lms_offset = 16384
lms_points = 0
lms_resolution = 0.45822189 [Hz]
lms_sweep = 7.50750751 [kHz]
lms_lipped = FALSE
lms_od_return = 1
lms_total_scans = 7
lms_90_width = 18.5 [us]
lms_acq_time = 2.1823486 [s]
lms_angle = 45 [deg]
lms_pulse = 4.25 [us]
lms_pulse_wait = 1 [s]
lms_base_preset = 2 [us]
lms_scvr_gain = 20
lms_relaxation_delay = 4 [s]
lms_amp_get = 25 [dc]
lms_blank_time = 2 [us]

```





```

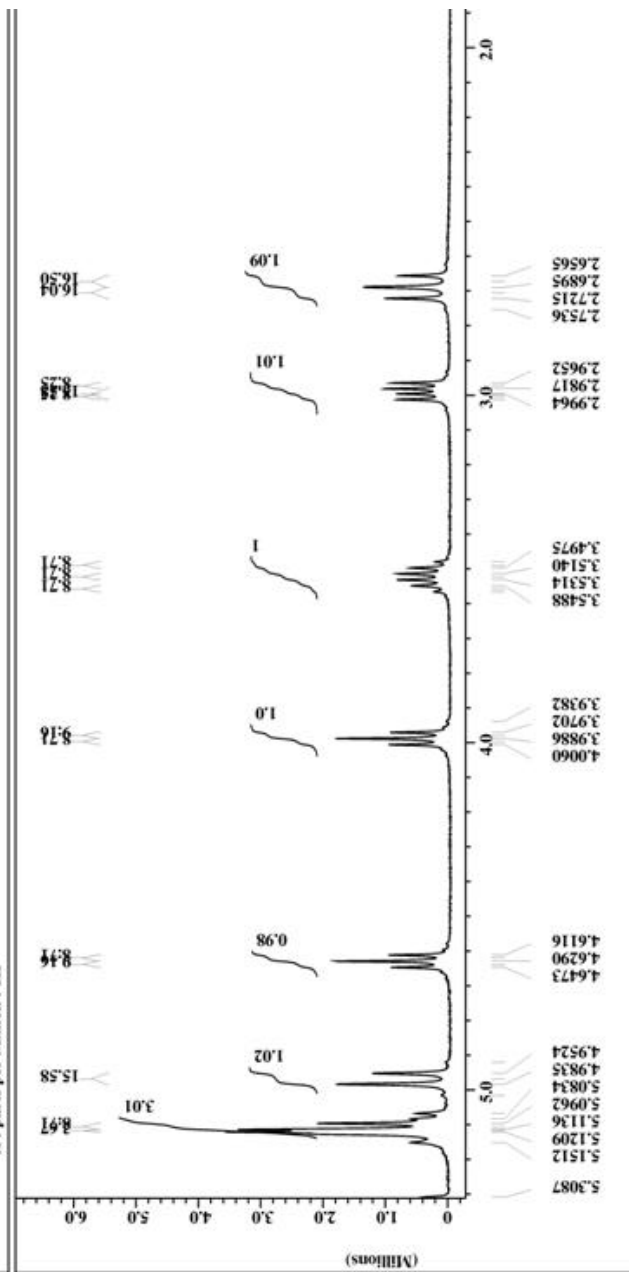
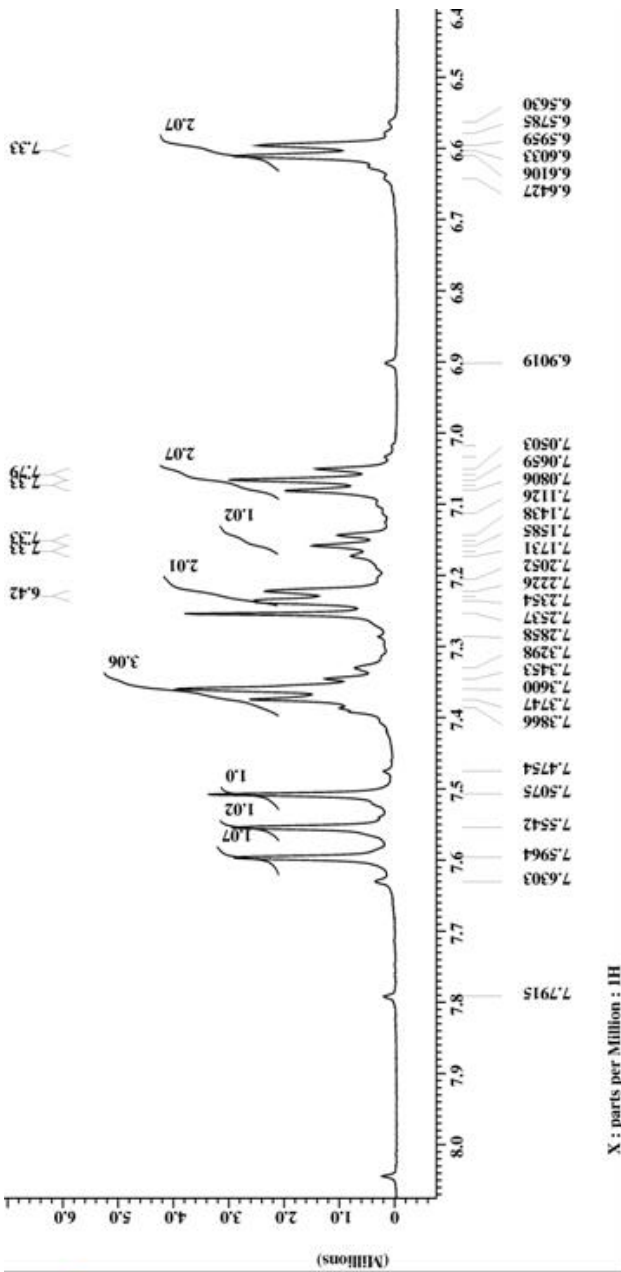
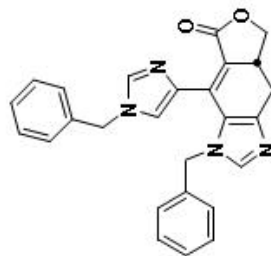
ilname      = sm_111_100_pure_2-4.j
author
experiment  = delta
sample_id   = S810791
solvent     = CHLOROFORM-D
revision    = 20-JAN-2008 05:45:53
vision_time = 28-MAR-2010 18:01:02
current_time = 28-MAR-2010 18:06:44

comment     = Single Pulse Experiment
sta_format  = ID COMPLEX
im_size     = 16384
im_title    = 1H
im_units    = [ppm]
imensions  = X
ite         = Eclipset 500
spectrometer = DELTA_RMR

field_strength = 11.7473579[T] (500[M]
acq_duration   = 2.1823488[s]
-domain        = 1H
-freq          = 500.15991521[M]
-offset        = 5[ppm]
-points        = 16384
-prescans      = 0
-resolution    = 0.45822189[Hz]
-sweep         = 7.50750751[kHz]
tipped         = FALSE
cd_return      = 1
cans           = 7
stai_scans    = 7

-60_width     = 18.5[us]
-acq_time     = 2.1823488[s]
-angle        = 45[deg]
-pulse        = 7.25[us]
-ntial_wait   = 3[s]
-base_preset  = 5[us]
-sevr_gain    = 20
-relaxation_delay = 4[s]
-emp_get      = 25[dc]
-ntblank_time = 2[us]

```





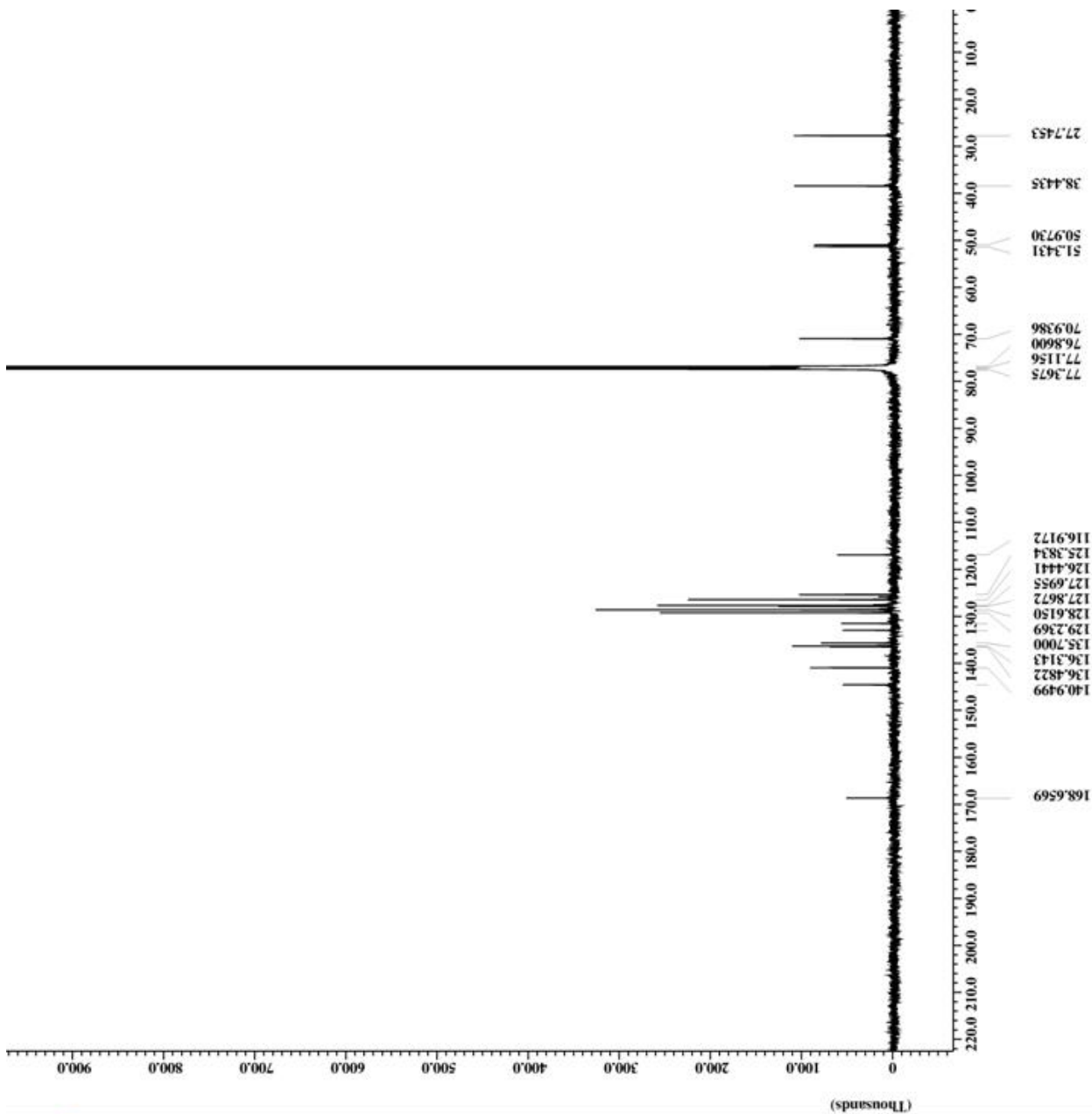
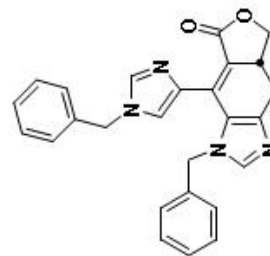
```

ilname      = sm_III_100_pure_2-2.j
author
xparment   = delta
sample_id  = S812042
solvent     = CHLOROFORM-D
reaction_time = 20-JAN-2008 14:17:55
revision_time = 20-JAN-2008 13:52:06
current_time = 28-MAR-2010 18:09:27

comment
ata_format = single pulse decouple
im_size     = 65536
im_title    = 13C
im_units    = [ppm]
imensions  = X
ite         = Eclipset 500
pctrometer = DELTA_NMR

ield_strength = 11.7473579[T] (500[MH
-acq_duration = 2.0840448[s]
-domain      = 13C
-freq        = 125.76529768[MHz]
-offset      = 100[ppm]
-points      = 65536
-prescans    = 4
-resolution  = 0.47983613[Hz]
-sweep       = 31.44654088[KHz]
-rr_domain  = 1H
-rr_freq     = 500.15991521[MHz]
-rr_offset   = 5[ppm]
-lipped      = FALSE
-bd_return   = 10
-cans        = 6000
-stal_scans  = 6000

-90_width   = 14.2[us]
-acq_time    = 2.0840448[s]
-angle       = 30[deg]
-pulse       = 4.73333333[us]
-nitral_wait = 1[s]
-be_time     = 1[s]
-base_preset = 3[us]
-scvr_gain   = 29
-relaxation_delay = 2[s]
-emp_get     = 26.9[dC]
-nblank_time = 2[us]
  
```





```

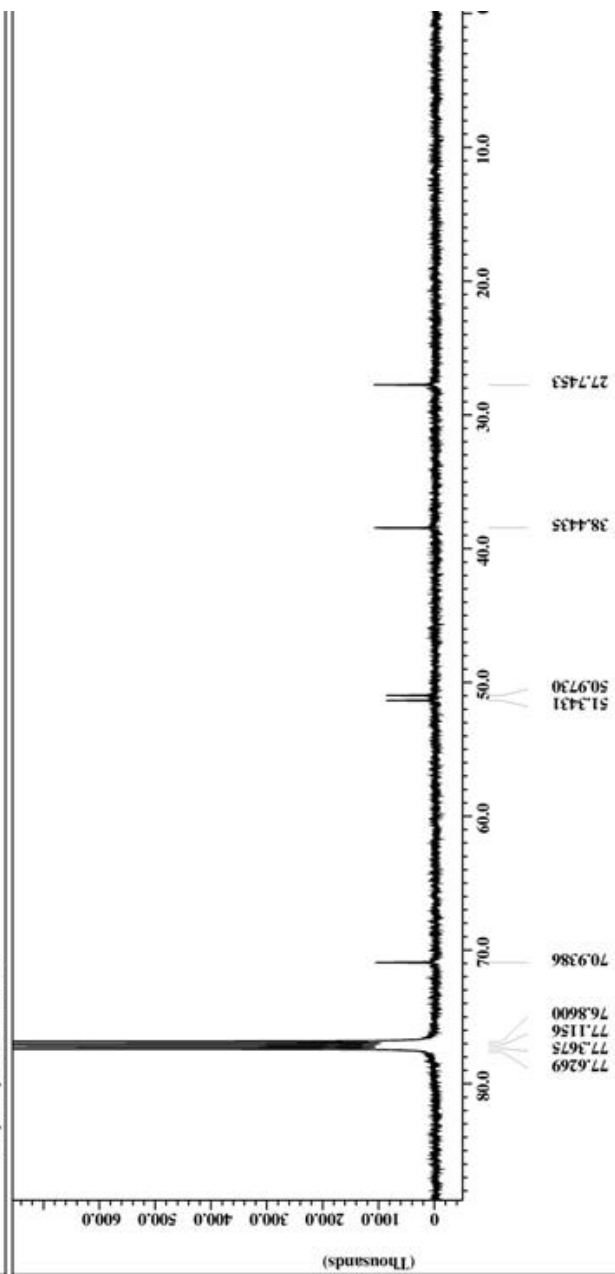
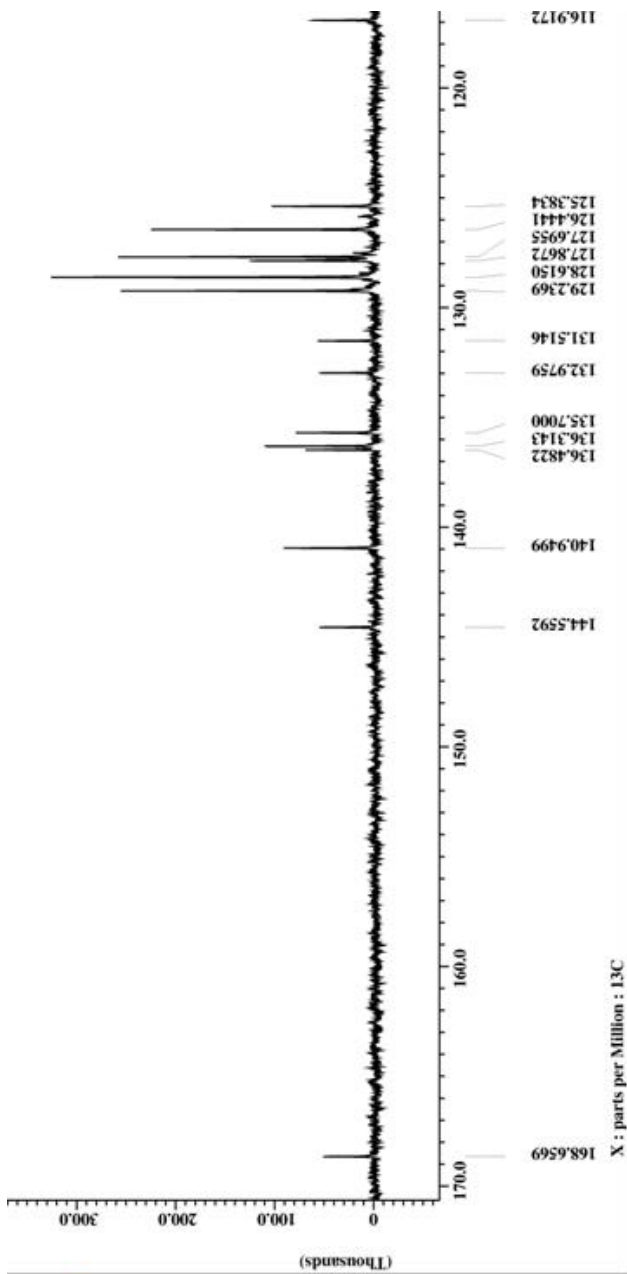
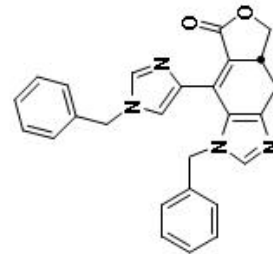
ilname      = sm_III_100_pure_2-2.j
author
xperiment   = delta
sample_id   = S812042
solvent     = CHLOROFORM-D
reaction_time = 20-JAN-2008 14:17:55
evision_time = 20-JAN-2008 13:52:06
urrent_time = 28-MAR-2010 18:12:13

omment
ata_format  = 1D COMPLEX
m_size      = 65536
m_title     = 13C
m_units     = [ppm]
mensions    = X
ite         = ECLIPSET 500
pctrometer  = DELTA_RMR

ield_strength = 11.7473579[T] (500[MH
-acq_duration = 2.0840448[s]
-domain      = 13C
-freq        = 125.76529768[MHz]
-offset      = 100[ppm]
-points      = 4
-prescans    = 4
-resolution  = 0.47983613[Hz]
-sweep       = 31.44654088[KHz]
rr_domain   = 1H
rr_freq     = 500.15991521[MHz]
rr_offset   = 5[ppm]
lipped      = FALSE
bd_return   = 10
cans        = 6000
stal_scans  = 6000

_90_width   = 14.2[us]
-acq_time   = 2.0840448[s]
-angle      = 30[deg]
-pulse      = 4.73333333[us]
nitial_wait = 1[s]
se_time     = 1[s]
base_preset = 3[us]
scvr_gain   = 29
elaxation_delay = 2[s]
emp_get     = 26.9[dc]
nblank_time = 2[us]

```

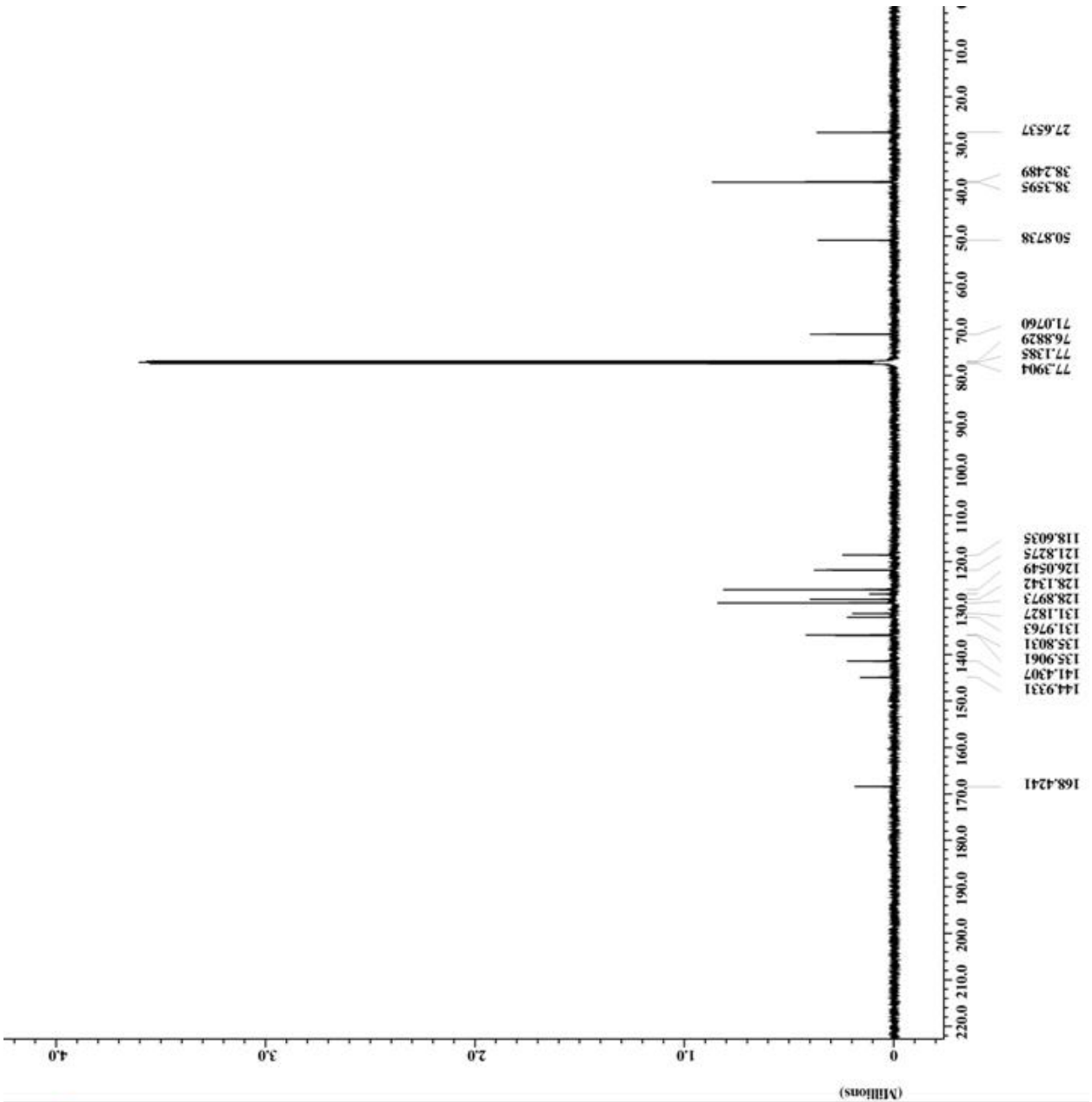
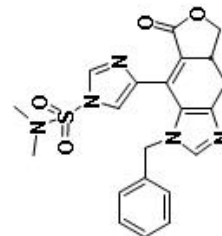


APPENDIX 35
¹H AND ¹³C NMR SPECTRUM OF
(*R*^{*})-4-(1-benzyl-7-oxo-4,4a,5,7-tetrahydro-1H-isobenzofuro[5,6-d]imidazol-8-yl)-N,N-dimethyl-1H-imidazole-1-sulfonamide (229)



```

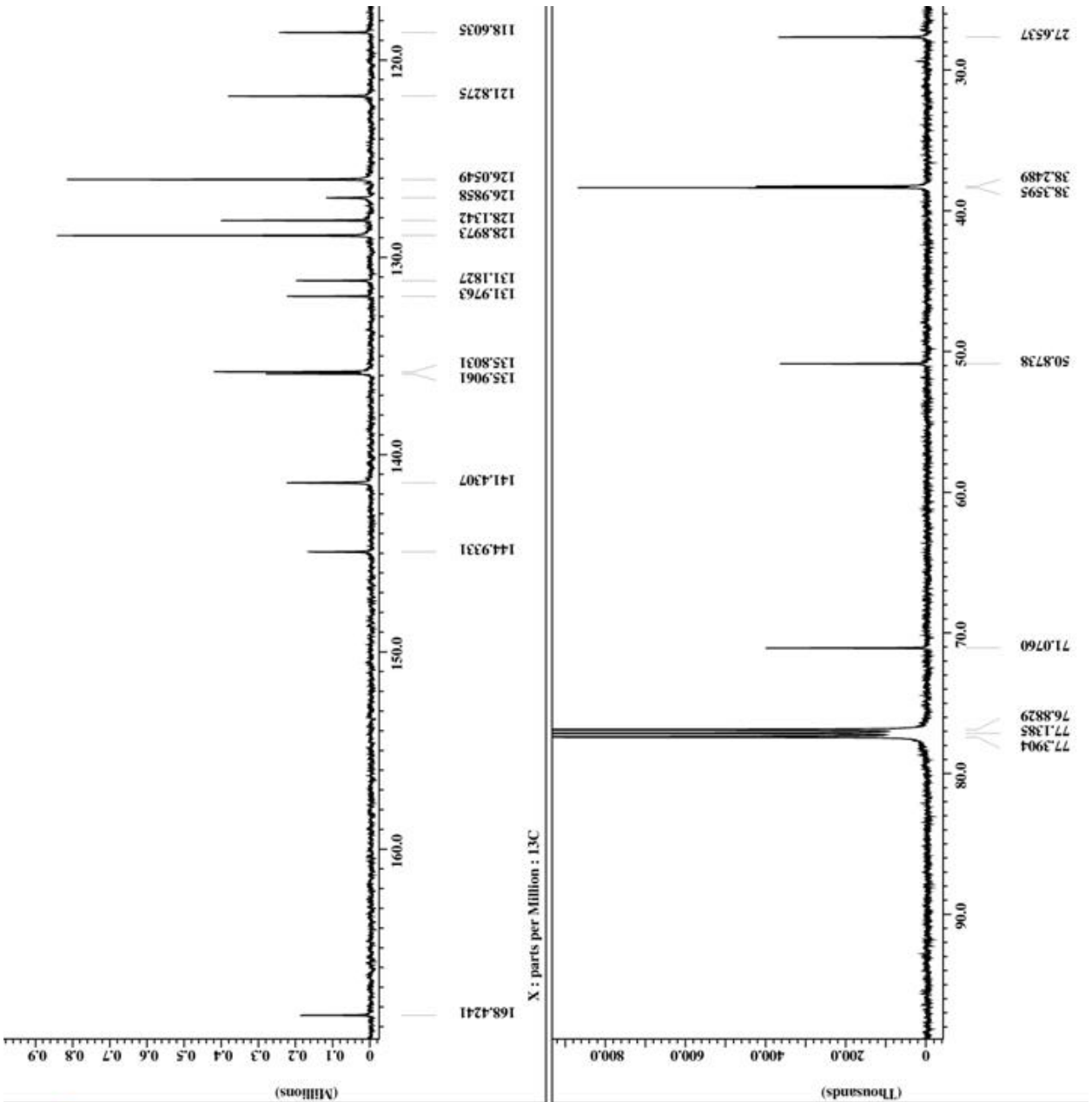
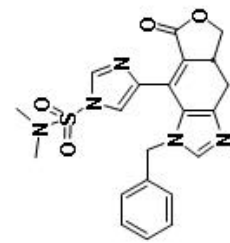
filename = sm_III_61_pure_500-2.
author = delta
experiment = single_pulse_dec
sample_id = S#827058
solvent = CHLOROFORM-D
reaction_time = 4-SEP-2007 10:23:45
revision_time = 4-SEP-2007 09:55:03
current_time = 28-MAR-2010 18:22:48
comment = single pulse decouple
ata_format = ID COMPLEX
im_size = 65536
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im_units = [ppm]
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ite = Eclipse+ 500
spectrometer = DELTA_NMR
field_strength = 11.7473579[T] (500[MH
acq_duration = 2.0840448[s]
domain = 13C
freq = 125.76529768[MHz]
offset = 100[ppm]
points = 65536
prescans = 4
resolution = 0.47983613[Hz]
sweep = 31.44654088[kHz]
tr_domain = 1H
tr_freq = 500.15991521[MHz]
tr_offset = 5[ppm]
lipped = FALSE
bd_return = 10
cans = 5000
otal_scans = 5000
_90_width = 14.2[us]
_acq_time = 2.0840448[s]
_angle = 30[deg]
_pulse = 4.73333333[us]
_nitral_wait = 1[s]
_be_time = 1[s]
_base_preset = 3[us]
_scvr_gain = 30
_relaxation_delay = 2[s]
emp_get = 25.7[dC]
nblank_time = 2[us]
  
```





```

filename = sm_III_61_pure_500-2.
author = delta
experiment = single_pulse_dec
sample_id = S827058
solvent = CHLOROFORM-D
reaction_time = 4-SEP-2007 10:23:45
revision_time = 4-SEP-2007 09:55:03
current_time = 28-MAR-2010 18:24:53
comment = single pulse decouple
          ID COMPLEX
          65536
          13C
          [ppm]
          X
          Eclipse+ 500
          DELTA_NMR
          spectrometer
          ata_format
          im_size
          im_title
          im_units
          im_dimensions
          lte
          spectrometer
          field_strength = 11.7473579[T] (500[MH
          acq_duration = 2.0840448[s]
          domain = 13C
          freq = 125.76529768[MHz]
          offset = 100[ppm]
          points = 65536
          prescans = 4
          resolution = 0.47983613[Hz]
          sweep = 31.44654088[kHz]
          tr_domain = 1H
          tr_freq = 500.15991521[MHz]
          tr_offset = 5[ppm]
          tipped = FALSE
          bd_return = 10
          cans = 5000
          otal_scans = 5000
          _90_width = 14.2[us]
          _acq_time = 2.0840448[s]
          _angle = 30[deg]
          _pulse = 4.73333333[us]
          _nitai_wait = 1[s]
          _pe_time = 1[s]
          _base_preset = 3[us]
          _scvr_gain = 30
          _relaxation_delay = 2[s]
          _emp_get = 25.7[dC]
          _nblank_time = 2[us]
  
```

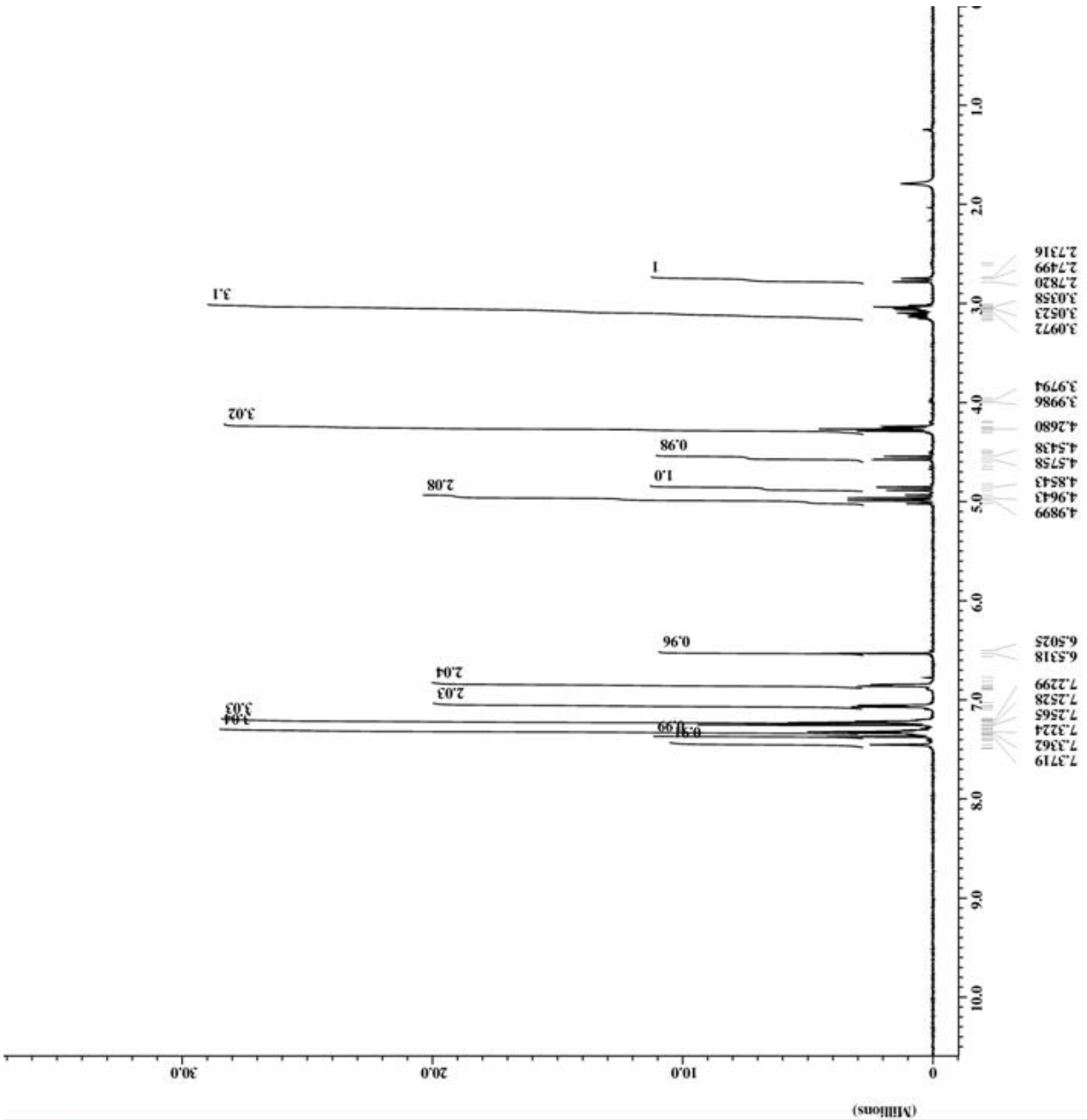
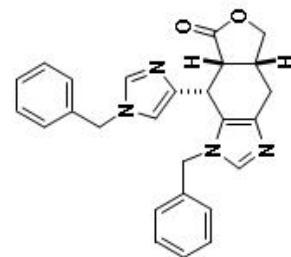


APPENDIX 36
¹H AND ¹³C NMR SPECTRUM OF
(4a*R*^{*},7a*S*^{*},8*R*^{*})-1-benzyl-8-(1-benzyl-1H-imidazol-4-yl)-4a,5,7a,8-tetrahydro-
1H-isobenzofuro[5,6-d]imidazol- 7(4H)-one (232)



```

filename = sm_V_Bn_Bn_Hy_pure-4.
author = delta
experiment = single_pulse_exp
sample_id = S818212
solvent = CHLOROFORM-D
reaction_time = 15-JAN-2009 22:46:52
acquisition_time = 28-MAR-2010 20:03:50
current_time = 28-MAR-2010 20:04:20
comment = Single Pulse Experiment
data_format = ID REAL
in_size = 16384
in_title = 1H
in_units = [ppm]
inversion = X
l1te = Eclipse+ 500
nucleus1 = DELTA_NMR
p1_frequency = 500.15991521 [MHz]
p1_offset = 5 [ppm]
p1_points = 16384
p1_prescans = 0
p1_resolution = 0.45822189 [Hz]
p1_sweep = 7.50750751 [kHz]
p1_tipped = FALSE
p1_return = 1
p1_cans = 12
p1_datal_scans = 12
p1_90_width = 18.5 [us]
p1_acq_time = 2.1823488 [s]
p1_angle = 45 [deg]
p1_pulse = 7.25 [us]
p1_pulse_wait = 1 [s]
p1_base_preset = 2 [us]
p1_scvr_gain = 22
p1_relaxation_delay = 4 [s]
p1_amp_get = 46.1 [dC]
p1_blank_time = 2 [us]
  
```

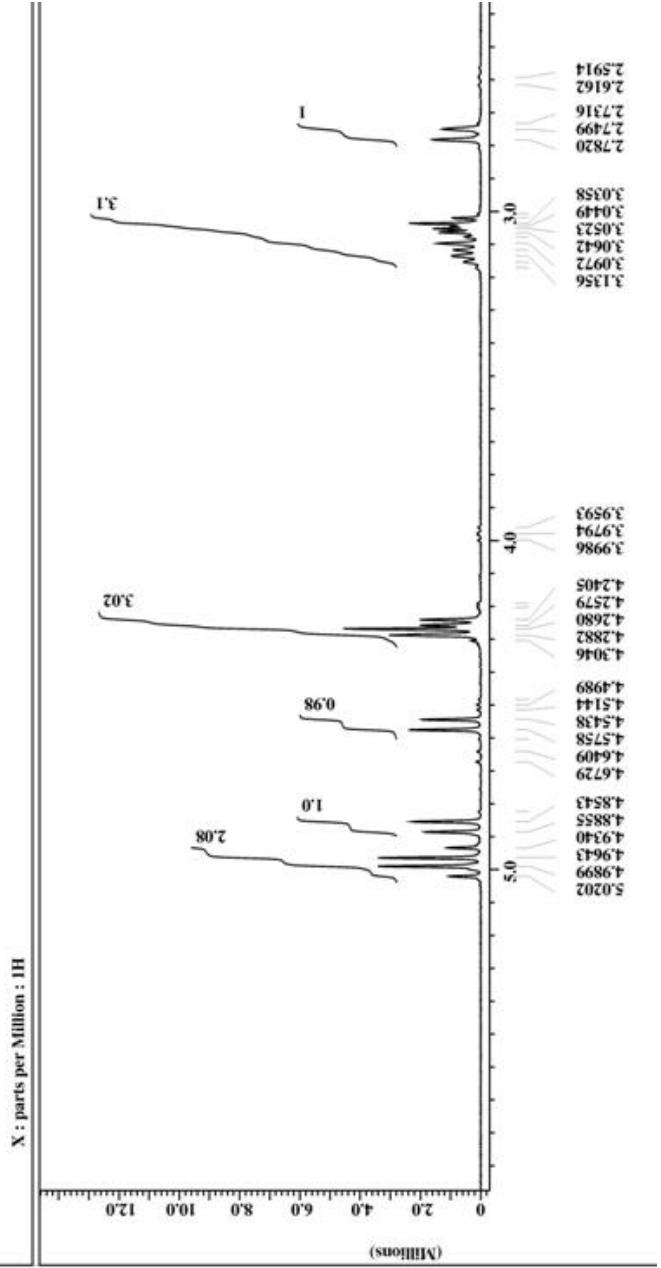
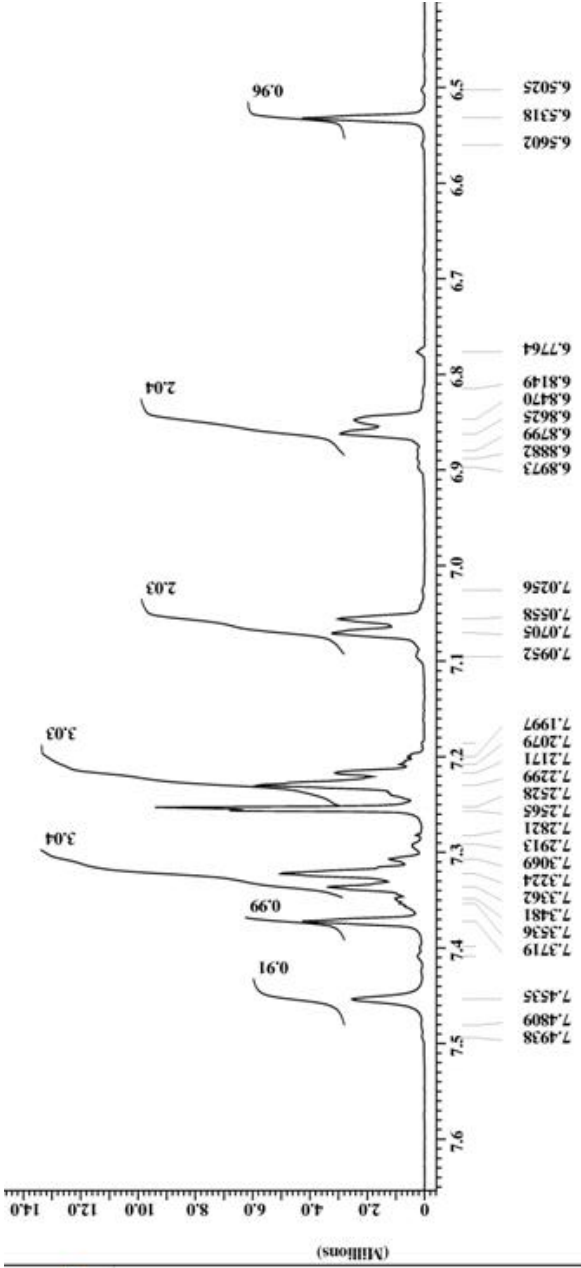
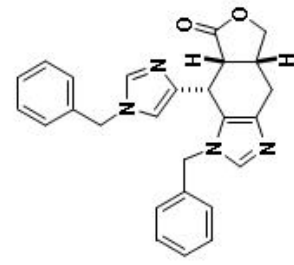




```

filename = sm_V_Bn-Bn_Hy_pure-4.
author = delta
experiment = single_pulse_exp
sample_id = S818212
solvent = CHLOROFORM-D
reaction_time = 15-JAN-2009 22:46:52
acquisition_time = 28-MAR-2010 20:03:50
current_time = 28-MAR-2010 20:06:47
comment = Single Pulse Experiment
data_format = ID REAL
in_size = 16384
in_title =
in_units =
instruments = X
ite = Eclipse+ 500
nucleus = DELTA_NMR
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```

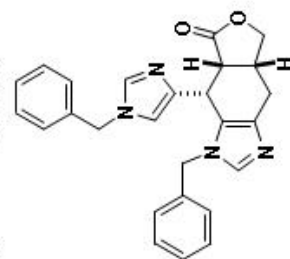
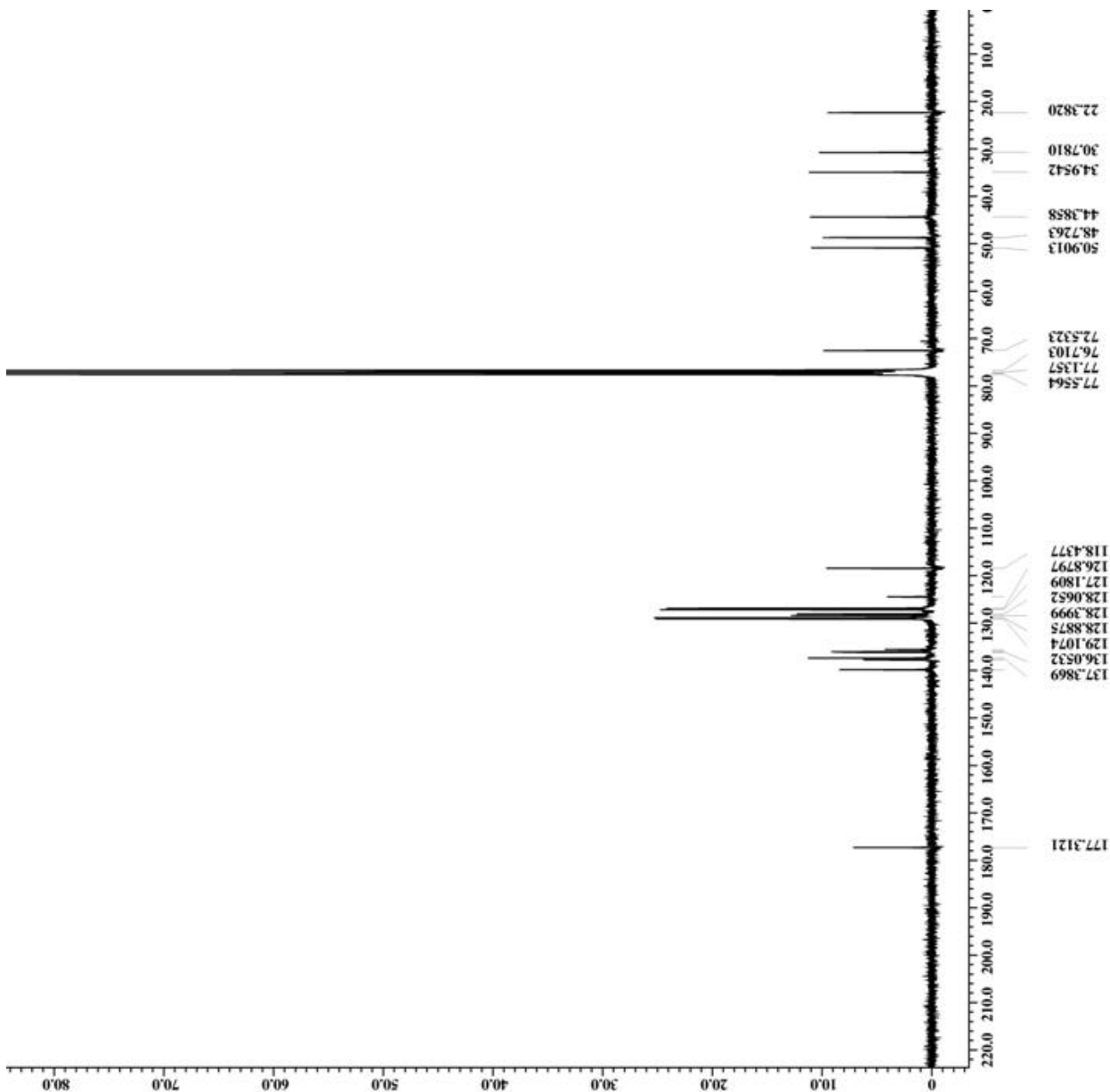




```

filename = sm_V_Bn_Bn_Hy_pure-3.
author = delta
experiment = single_pulse_dec
sample_id = S835108
solvent = CHLOROFORM-D
reaction_time = 15-JAN-2009 07:31:35
evision_time = 15-JAN-2009 11:11:15
current_time = 28-MAR-2010 20:08:52
comment = single pulse decouple
ata_format = ID REAL
in_size = 52428
in_title = 13C
in_units = [ppm]
inmensions = X
ite = ECX 300
spectrometer = DELTA2_NMR
ield_strength = 7.0586013[T] (300[MHz]
_acq_duration = 2.76824064[s]
_domain = 13C
_freq = 75.56823426[MHz]
_offset = 100[ppm]
_points = 65536
_prescans = 4
_resolution = 0.36124027[Hz]
_sweep = 23.67424242[kHz]
tr_domain = 1H
tr_freq = 300.52965592[MHz]
tr_offset = 5[ppm]
lipped = FALSE
od_return = 10
cans = 5000
otal_scans = 5000
_90_width = 9.75[us]
_acq_time = 2.76824064[s]
_angle = 30[deg]
_atn = 8[db]
_pulse = 3.25[us]
tr_atn_dec = 25[db]
tr_atn_noe = 25[db]
tr_noise = WALTZ
scoupling = TRUE
nitial_wait = 1[s]
oe_time = TRUE
oe_time = 3[s]
scr_gain = 50
elaxation_delay = 3[s]
petition_time = 5.76824064[s]
emp_get = 23.3[dc]
  
```

(thousands)

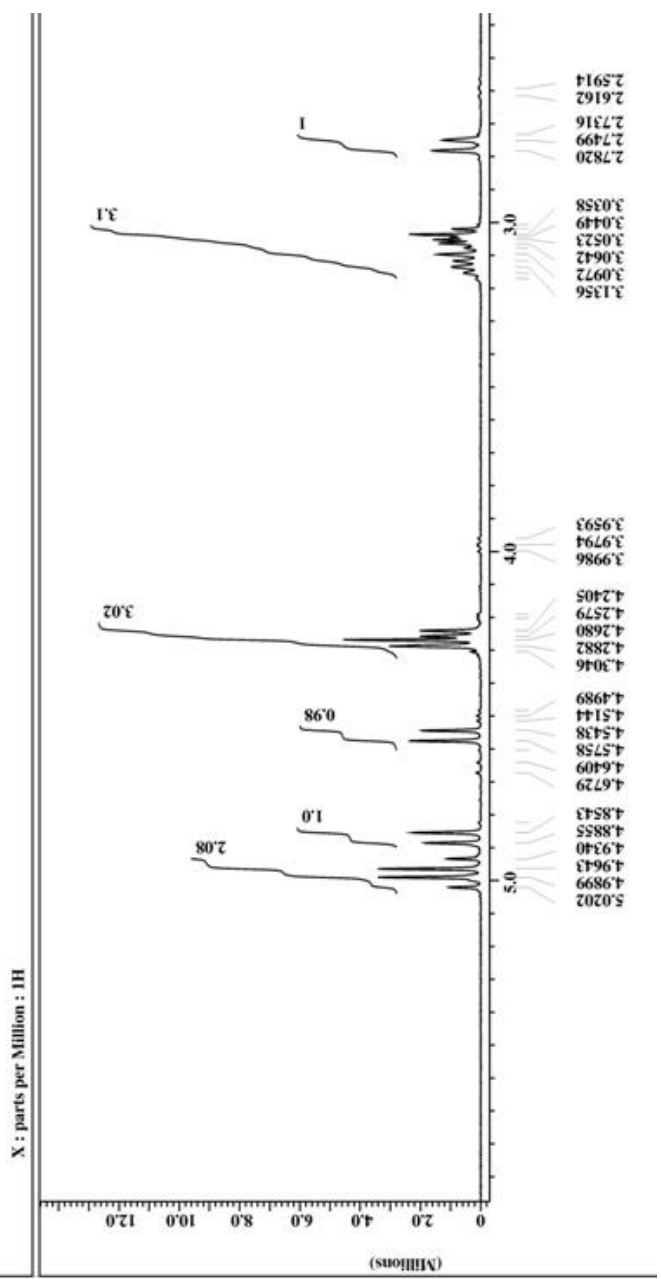
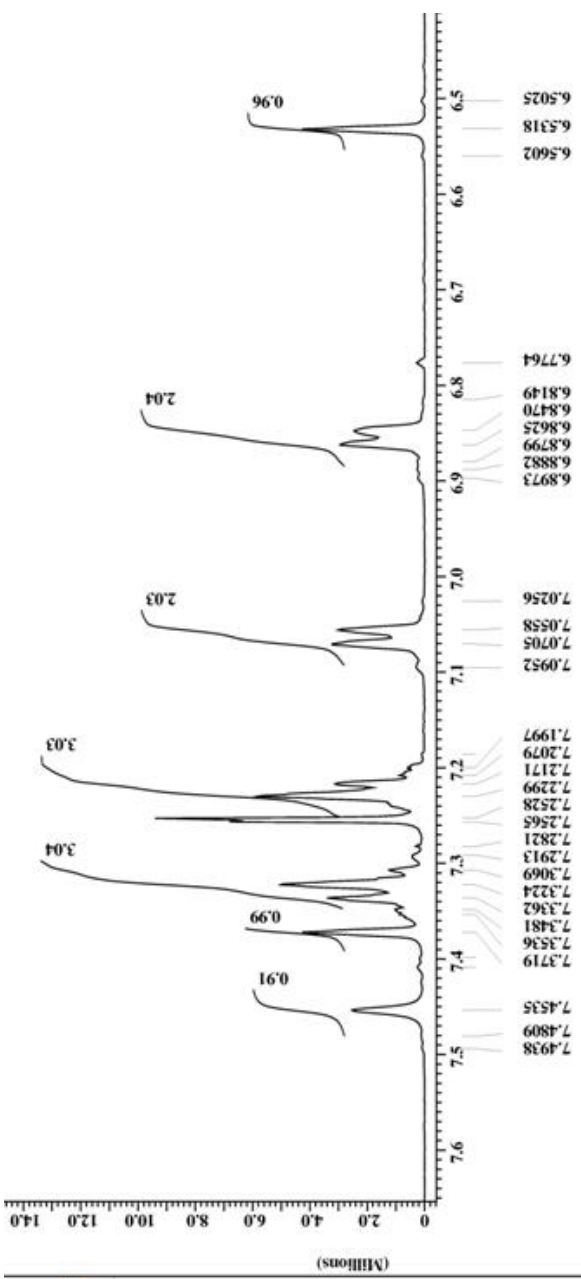
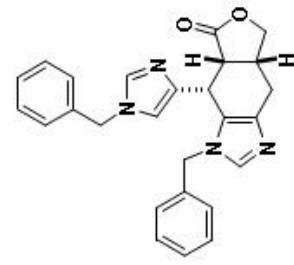




```

filename = sm_V_Bn-Bn_Hy_pure-4.
author = delta
experiment = single_pulse_exp
sample_id = S818212
solvent = CHLOROFORM-D
reaction_time = 15-JAN-2009 22:46:52
acquisition_time = 28-MAR-2010 20:03:50
current_time = 28-MAR-2010 20:06:47
comment = Single Pulse Experiment
data_format = ID REAL
in_size = 16384
in_title =
in_units =
instruments = X
ite = Eclipse+ 500
nucleus = DELTA_NMR
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```



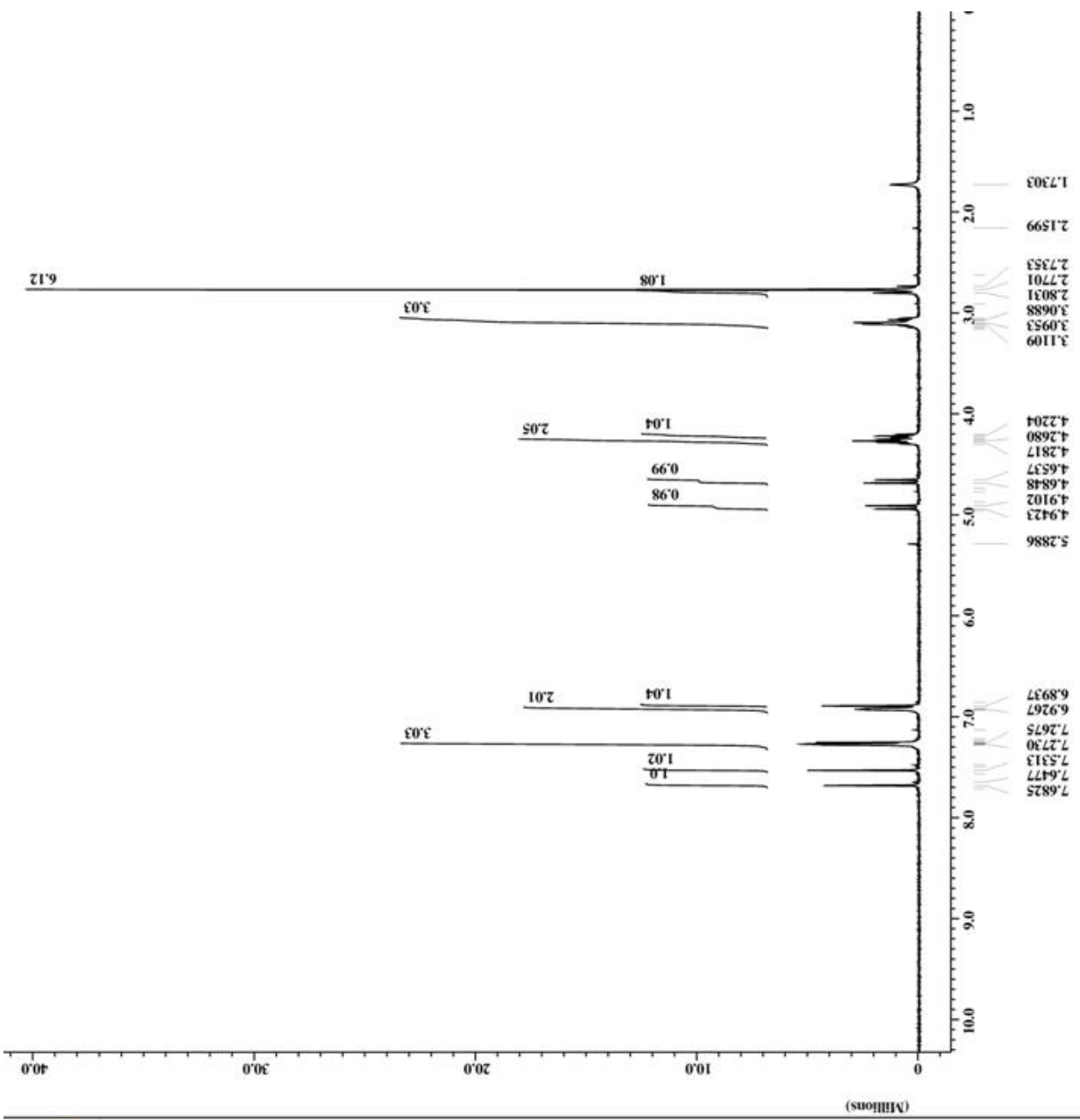
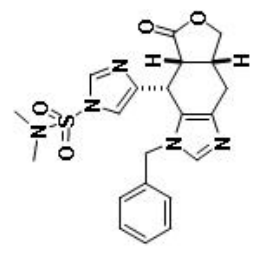
APPENDIX 37
¹H AND ¹³C NMR SPECTRUM OF
4-((4a*R*^{*},7a*S*^{*},8*R*^{*})-1-benzyl-7-oxo-4,4a,5,7,7a,8-hexahydro-1H-
isobenzofuro[5,6-d]imidazol-8-yl)-N,N-dimethyl-1H-imidazole-1-sulfonamide
(231)



```

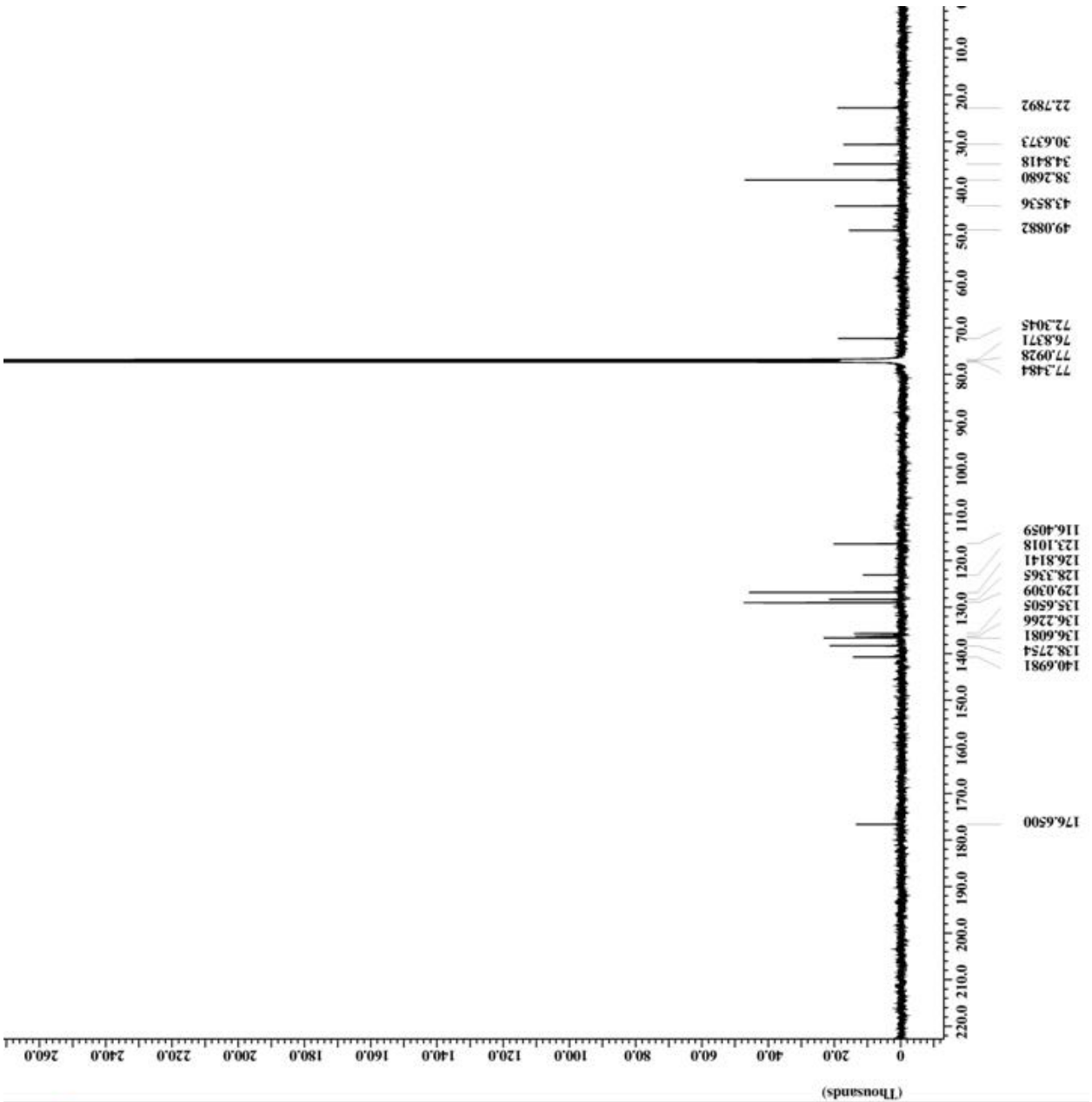
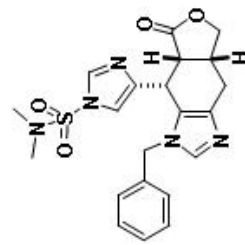
filename = sm_V_102_pure-3.jdf
author = delta
experiment = single_pulse_exp
sample_id = S#389287
solvent = CHLOROFORM-D
reaction_time = 18-MAR-2009 20:38:58
evision_time = 28-MAR-2010 20:15:37
current_time = 28-MAR-2010 20:16:08
comment = Single Pulse Experiment
ata_format = ID COMPLEX
in_size = 16384
in_title = 1H
in_units = [ppm]
inmensions = X
ite = Eclipse+ 500
spectrometer = DELTA_NMR
field_strength = 11.747379[T] (500[MH
acq_duration = 2.182348[s]
domain = 1H
freq = 500.15991521[MHz]
offset = 5[ppm]
points = 16384
prescans = 0
resolution = 0.45822189[Hz]
sweep = 7.50750751[kHz]
lipped = FALSE
od_return = 1
cans = 8
otal_scans = 8
_90_width = 18.5[us]
_acq_time = 2.182348[s]
_angle = 45[deg]
_pulse = 7.25[us]
nitel_wait = 1[s]
hase_preset = 2[us]
ecvr_gain = 23
elaxation_delay = 4[s]
emp_get = 46[dc]
nblank_time = 2[us]

```





```
filename = sm_V_67_ii_pure-2.jdf
author = delta
experiment = single_pulse_dec
sample_id = S8730303
solvent = CHLOROFORM-D
reaction_time = 19-MAR-2009 14:38:41
revision_time = 19-MAR-2009 10:47:05
current_time = 28-MAR-2010 20:18:50
comment = single pulse decouple
ata_format = ID COMPLEX
im_size = 65536
im_title = 13C
im_units = [ppm]
imensions = X
ite = Eclipse+ 500
spectrometer = DELTA_NMR
field_strength = 11.7473579[T] (500[MH
acq_duration = 2.0840448[s]
domain = 13C
freq = 125.76529768[MHz]
offset = 100[ppm]
points = 65536
prescans = 4
resolution = 0.47983613[Hz]
sweep = 31.44654088[kHz]
tr_domain = 1H
tr_freq = 500.15991521[MHz]
tr_offset = 5[ppm]
lipped = FALSE
bd_return = 10
cans = 6000
otal_scans = 6000
_90_width = 14.2[us]
_acq_time = 2.0840448[s]
_angle = 30[deg]
_pulse = 4.73333333[us]
nitai_wait = 1[s]
pe_time = 1[s]
base_preset = 3[us]
scvr_gain = 24
relaxation_delay = 2[s]
emp_get = 29.3[dC]
nblank_time = 2[us]
```





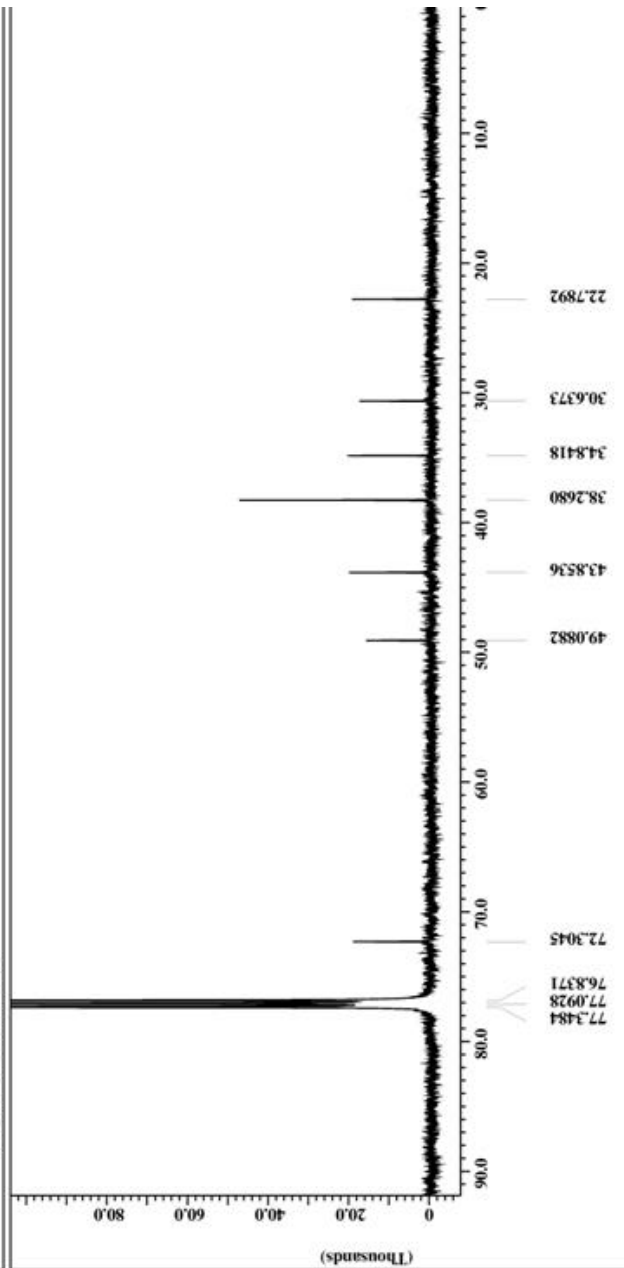
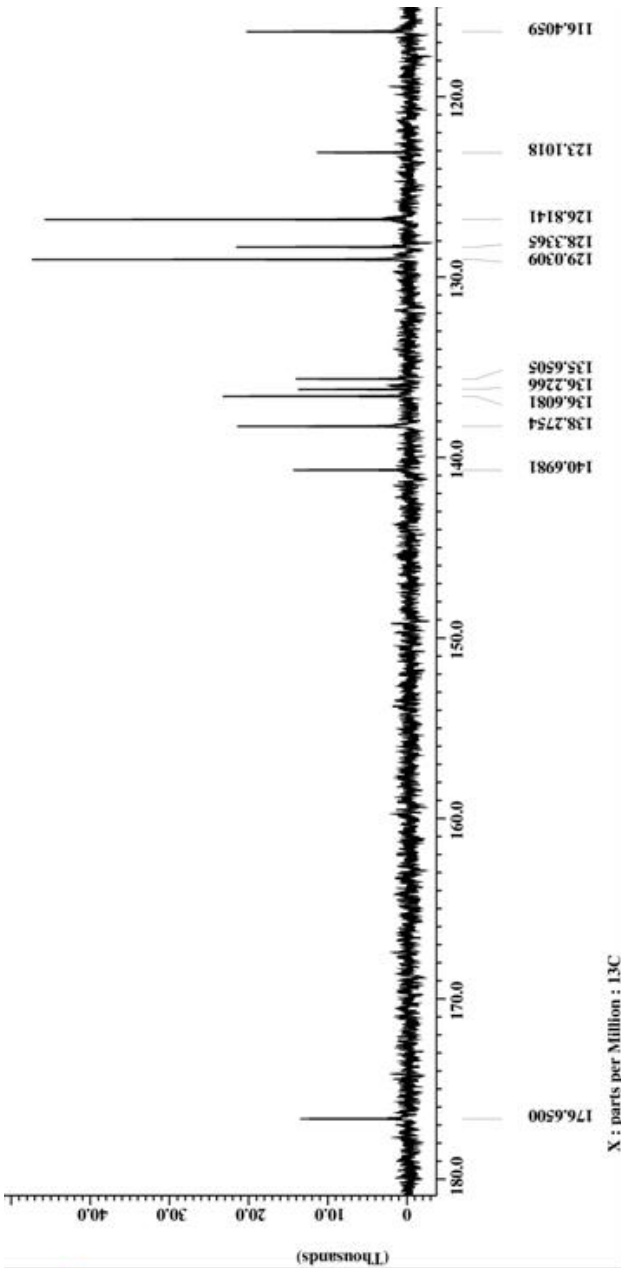
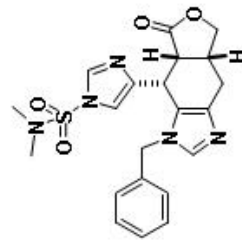
```

filename = sm_V_67_ii_pure-2.jdf
author =
experiment = single_pulse_dec
sample_id = S8730303
solvent = CHLOROFORM-D
reaction_time = 19-MAR-2009 14:38:41
evision_time = 19-MAR-2009 10:47:05
current_time = 28-MAR-2010 20:20:29

comment = single pulse decouple
ata_format = ID COMPLEX
in_size = 65536
in_title = 13C
in_units = [ppm]
inmensions = X
ite = Eclipse+ 500
spectrometer = DELTA_NMR

field_strength = 11.7473579[T] (500[MH
acq_duration = 2.0840448[s]
domain = 13C
freq = 125.76529768[MHz]
offset = 100[ppm]
points = 65536
prescans = 4
resolution = 0.47983613[Hz]
sweep = 31.44654088[kHz]
tr_domain = 1H
tr_freq = 500.15991521[MHz]
tr_offset = 5[ppm]
lipped = FALSE
bd_return = 10
cans = 6000
otal_scans = 6000

_90_width = 14.2[us]
acq_time = 2.0840448[s]
angle = 30[deg]
pulse = 4.73333333[us]
nitai_wait = 1[s]
pe_time = 1[s]
base_preset = 3[us]
scrz_gain = 24
relaxation_delay = 2[s]
emp_get = 29.3[dC]
nblank_time = 2[us]
  
```

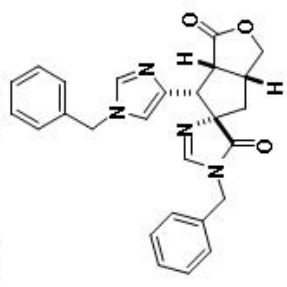
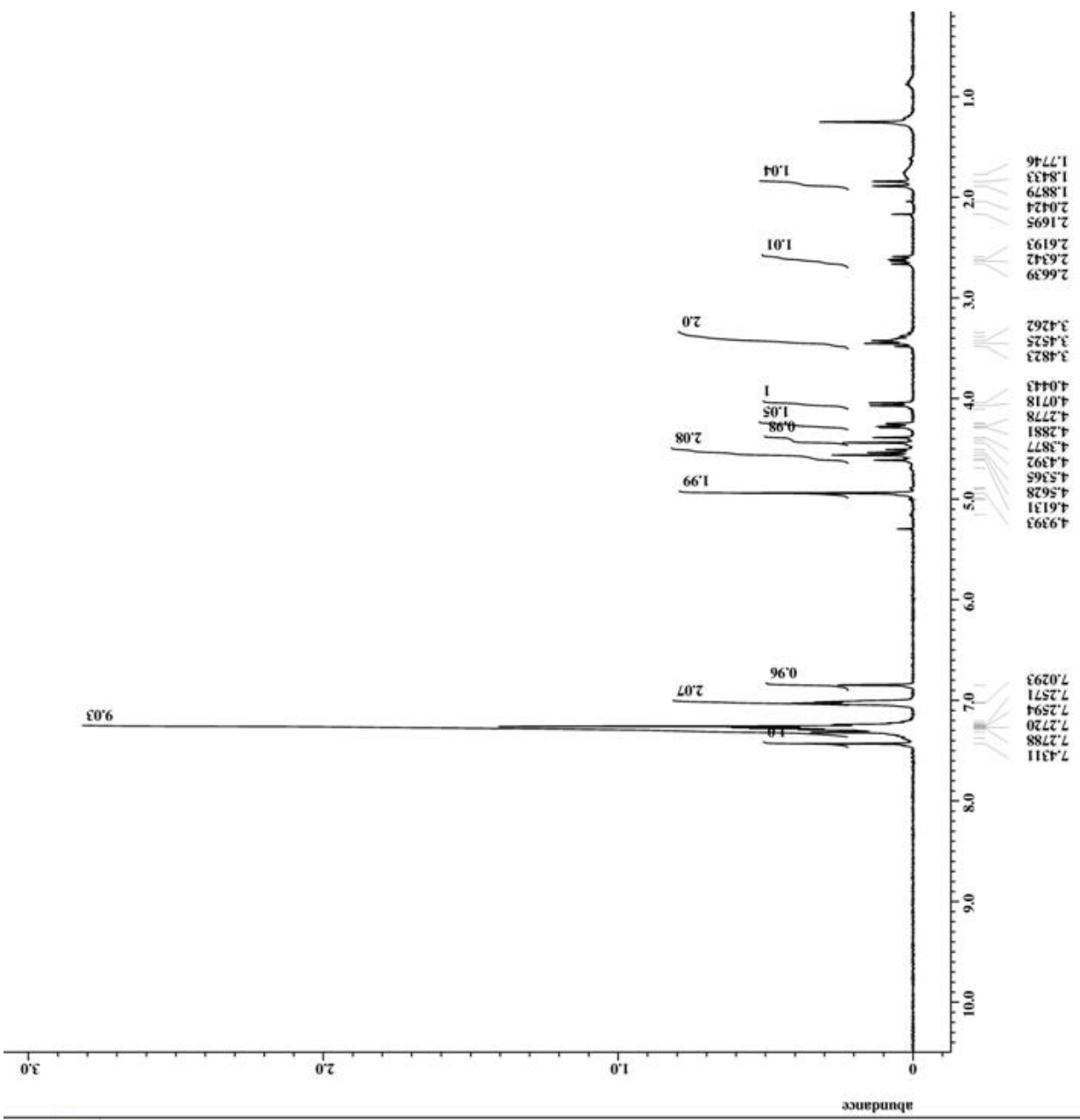


APPENDIX 38
¹H AND ¹³C NMR SPECTRUM OF
(3a*R*^{*},4*S*^{*},6*R*^{*},6a*S*^{*})-1'-benzyl-6-(1-benzyl-1H-imidazol-4-yl)-3a,4,6,6a-
tetrahydrospiro[cyclopenta[c]furan-5,4'-imidazole]-1,5'(1*H*,3*H*)-dione (234)



```

filename = sm_V_46_pure-4_.jdf
author = delta
experiment = single_pulse.ex2
sample_id = S#767039
solvent = CHLOROFORM-D
creation_time = 16-JAN-2009 21:40:51
revision_time = 28-MAR-2010 20:24:53
current_time = 28-MAR-2010 20:25:38
comment =
  = single_pulse
  = ID REAL
  = 13107
  = 1H
  = [ppm]
  = X
  = ECX 300
  = DELTA2_NMR
  = 7.0586013[T] (300[MHz]
  = 2.90717696[s]
  = 1H
  = 300.52965592[MHz]
  = 5[ppm]
  = 16384
  = 0
  = 0.34397621[Hz]
  = 5.63570784[kHz]
  = 1H
  = 300.52965592[MHz]
  = 5[ppm]
  = 1H
  = 300.52965592[MHz]
  = 5[ppm]
  = FALSE
  = 1
  = 24
  = 24
  = 13.01[us]
  = 2.90717696[s]
  = 45[deg]
  = 4[db]
  = 6.505[us]
  = Off
  = Off
  = FALSE
  = 1[s]
  = 46
  = 5[s]
  = 7.90717696[s]
  = 23[dc]
  
```





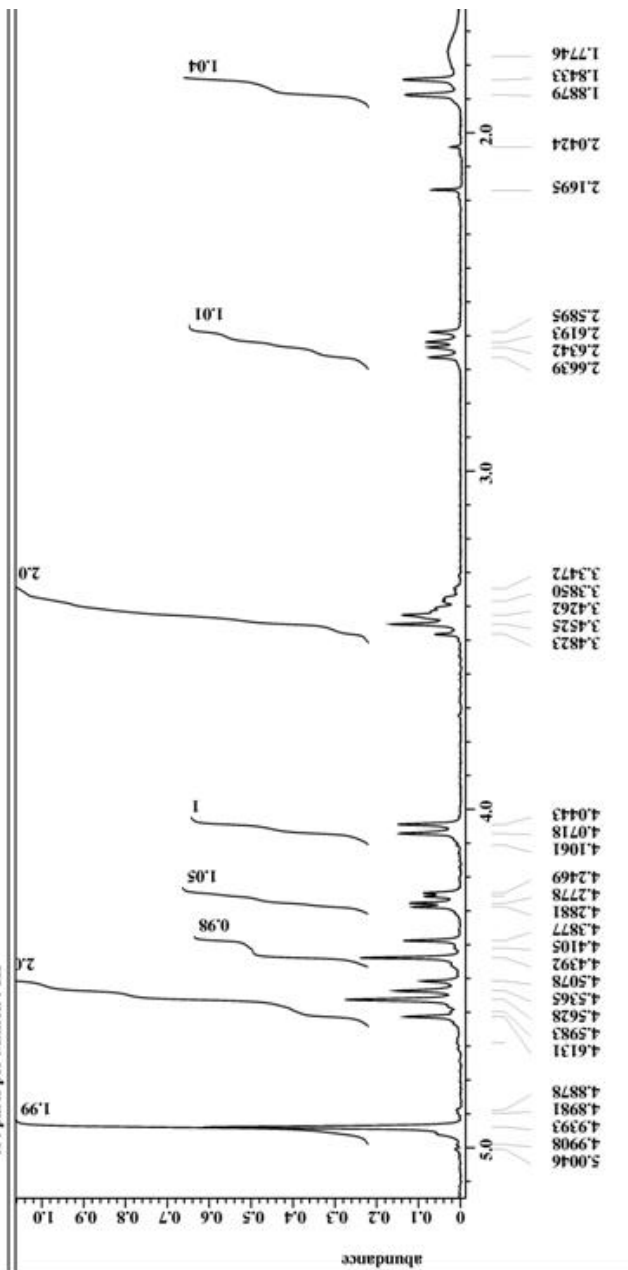
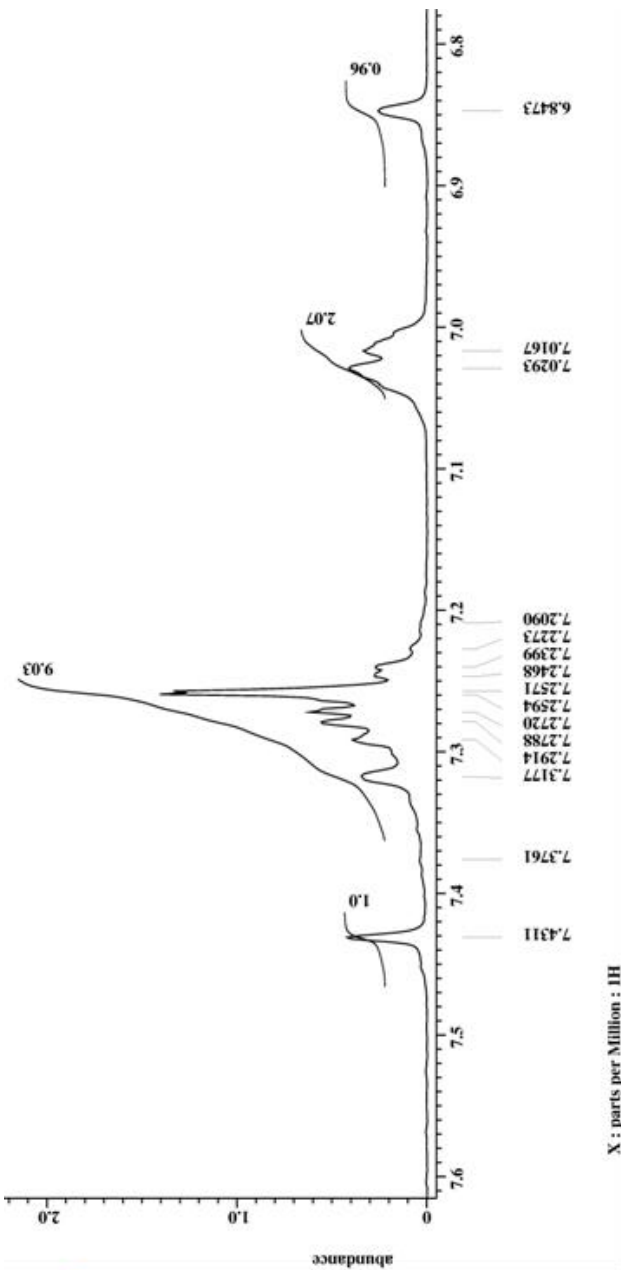
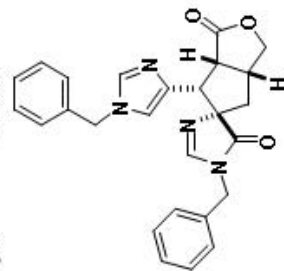
```

ilname      = sm_v_46_pure-4.jdf
author      =
experiment  = single_pulse.ex2
sample_id   = S876039
solvent     = CHLOROFORM-D
reaction_time = 16-JAN-2009 21:40:51
revision_time = 28-MAR-2010 20:24:53
current_time = 28-MAR-2010 20:27:50

comment     =
date_format = ID REAL
im_size     = 13107
im_title    =
im_units    = [ppm]
imensions   = X
ite         =
spectrometer = ECA 300
            = DELTA2_NMR

field_strength = 7.0586013[T] (300[MHz])
-acq_duration  = 2.90717696[s]
-domain        = 1H
-freq          = 300.52965592[MHz]
-offset        = 5[ppm]
-points        = 16384
-prescans      = 0
-resolution    = 0.34397631[Hz]
-sweep         = 5.63570784[kHz]
-rr_domain     = 1H
-rr_freq       = 300.52965592[MHz]
-rr_offset     = 5[ppm]
-rr_domain     = 1H
-ri_freq       = 300.52965592[MHz]
-ri_offset     = 5[ppm]
-lipped        = FALSE
-cd_return     = 1
-cans          = 24
-stal_scans    = 24

-90_width     = 13.01[us]
-acq_time     = 2.90717696[s]
-angle        = 45[deg]
-atn          = 4[db]
-pulse        = 6.505[us]
-rr_mode      = Off
-rr_mode      = Off
-ante_presat  = FALSE
-ntial_wait   = 1[s]
-scvr_gain    = 46
-relaxation_delay = 5[s]
-epitation_time = 7.90717696[s]
-emp_get      = 23[dc]
  
```





```

ilname      = sm_v_46_pure-3.jdf
author
experiment  = delta
sample_id   = S8769604
solvent     = CHLOROFORM-D
revision_time = 17-JAN-2009 07:18:45
evision_time = 17-JAN-2009 16:25:45
current_time = 28-MAR-2010 20:30:27

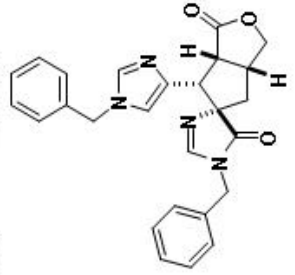
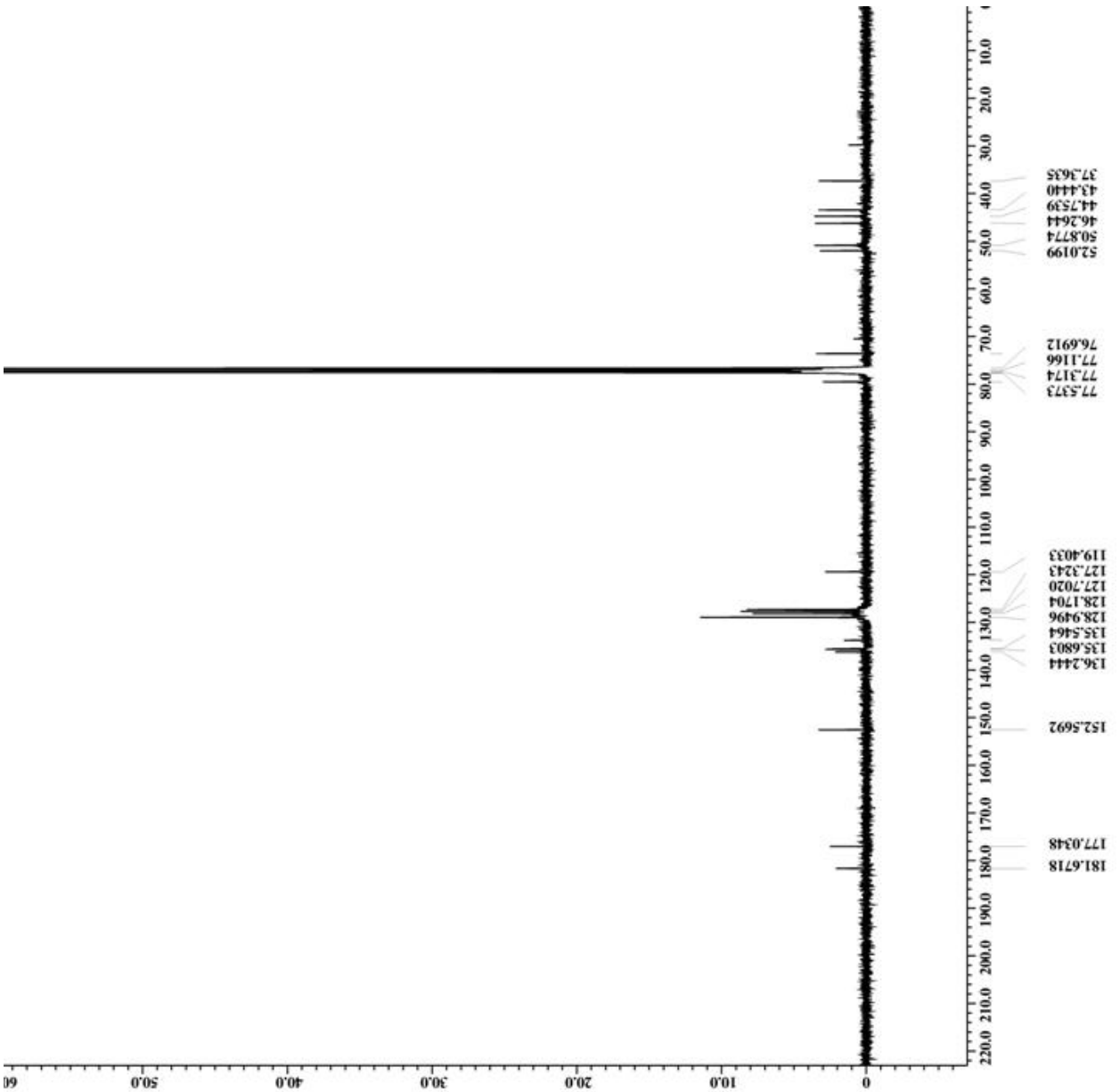
comment
ate_format  = single pulse decouple
im_size     = 52428
im_title    = 13C
im_units    = [ppm]
imensions   = X
ite         = ECX 300
pctrometer  = DELTA2_NMR

ield_strength = 7.0586013[T] (300[MHz]
-acq_duration = 2.76824064[s]
-domain      = 13C
-freq       = 75.56823426[MHz]
-offset     = 100[ppm]
-points     = 65536
-prescans   = 4
-resolution = 0.36124027[Hz]
-sweep      = 23.67424242[KHz]
-rr_domain  = 1H
-rr_freq    = 300.52965592[MHz]
-rr_offset  = 5[ppm]
-lipped     = TRUE
-bd_return  = 10
-cans       = 6000
-stal_scans = 6000

_90_width   = 9.75[us]
-acq_time   = 2.76824064[s]
-angle      = 30[deg]
-atn        = 8[db]
-pulse      = 3.25[us]
-rr_atn_dec = 25[db]
-rr_atn_noe = 25[db]
-rr_noise   = WALTZ
-acoupling  = TRUE
-nitital_wait = 1[s]
-be         = TRUE
-be_time    = 3[s]
-scvr_gain  = 50
-relaxation_delay = 3[s]
-epitation_time = 5.76824064[s]
-emp_get    = 23.3[dc]

```

(thousands)





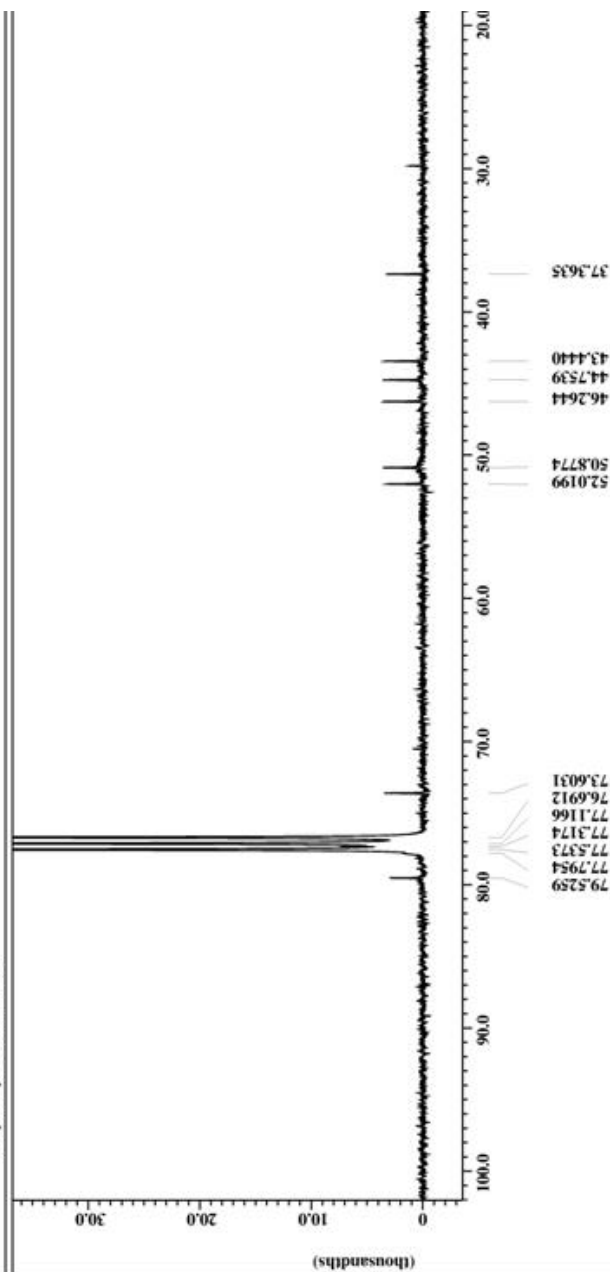
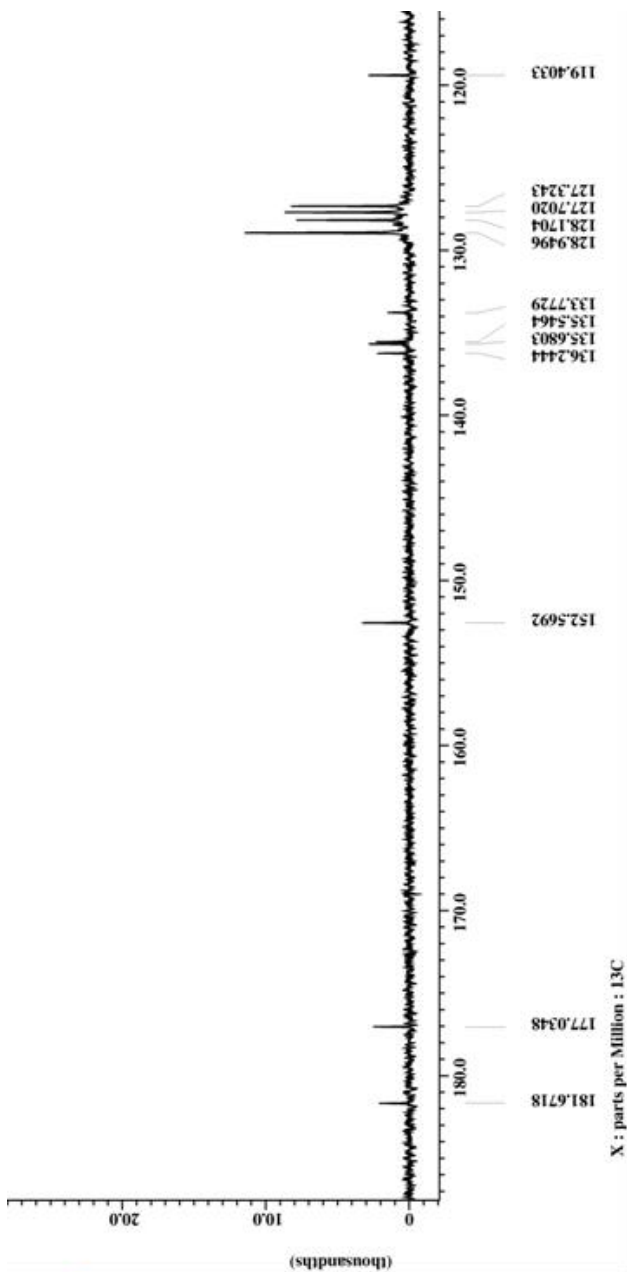
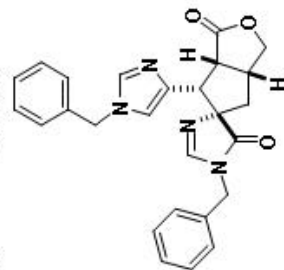
```

ilname      = sm_v_46_pure-3.jdf
author      =
experiment  = single_pulse_dec
sample_id   = S876904
solvent     = CHLOROFORM-D
reaction_time = 17-JAN-2009 07:18:45
evision_time = 17-JAN-2009 16:25:45
current_time = 28-MAR-2010 20:32:05

comment     = single pulse decouple
ate_format  = 1D REAL
im_size     = 52428
im_title    = 13C
im_units    = [ppm]
imensions  = X
ite         =
pctrometer  = ECK 300
            = DELTA2_NMR

field_strength = 7.0586013[T] (300[MHz]
-acq_duration  = 2.76824064[s]
-domain        = 13C
-freq          = 75.56823426[MHz]
-offset        = 100[ppm]
-points        = 65536
-prescans      = 4
-resolution    = 0.36124027[Hz]
-sweep         = 23.67424242[KHz]
-rr_domain    = 1H
-rr_freq       = 300.52965592[MHz]
-rr_offset     = 5[ppm]
-lipped        = TRUE
-bd_return     = 10
-cans          = 6000
-stal_scans    = 6000

-90_width     = 9.75[us]
-acq_time     = 2.76824064[s]
-angle        = 30[deg]
-atn          = 8[db]
-pulse        = 3.25[us]
-rr_atn_dec   = 25[db]
-rr_atn_noe   = 25[db]
-rr_noise     = WALTZ
-acoupling    = TRUE
-nit1al_wait  = 1[s]
-be           = TRUE
-be_time      = 3[s]
-scvr_gain    = 50
-relaxation_delay = 3[s]
-epitation_time = 5.76824064[s]
-emp_get      = 23.3[dc]
  
```



APPENDIX 39
¹H AND ¹³C NMR SPECTRUM OF
4-((3a*R,4'*S**,6*R**,6a*S*')-1'-benzyl-1,5'-dioxo-1,1',3,3a,4,5',6,6a-**
octahydrospiro[cyclopenta[c]furan-5,4'-imidazole]-6-yl)-N,N-dimethyl-1H-
imidazole-1-sulfonamide (233)



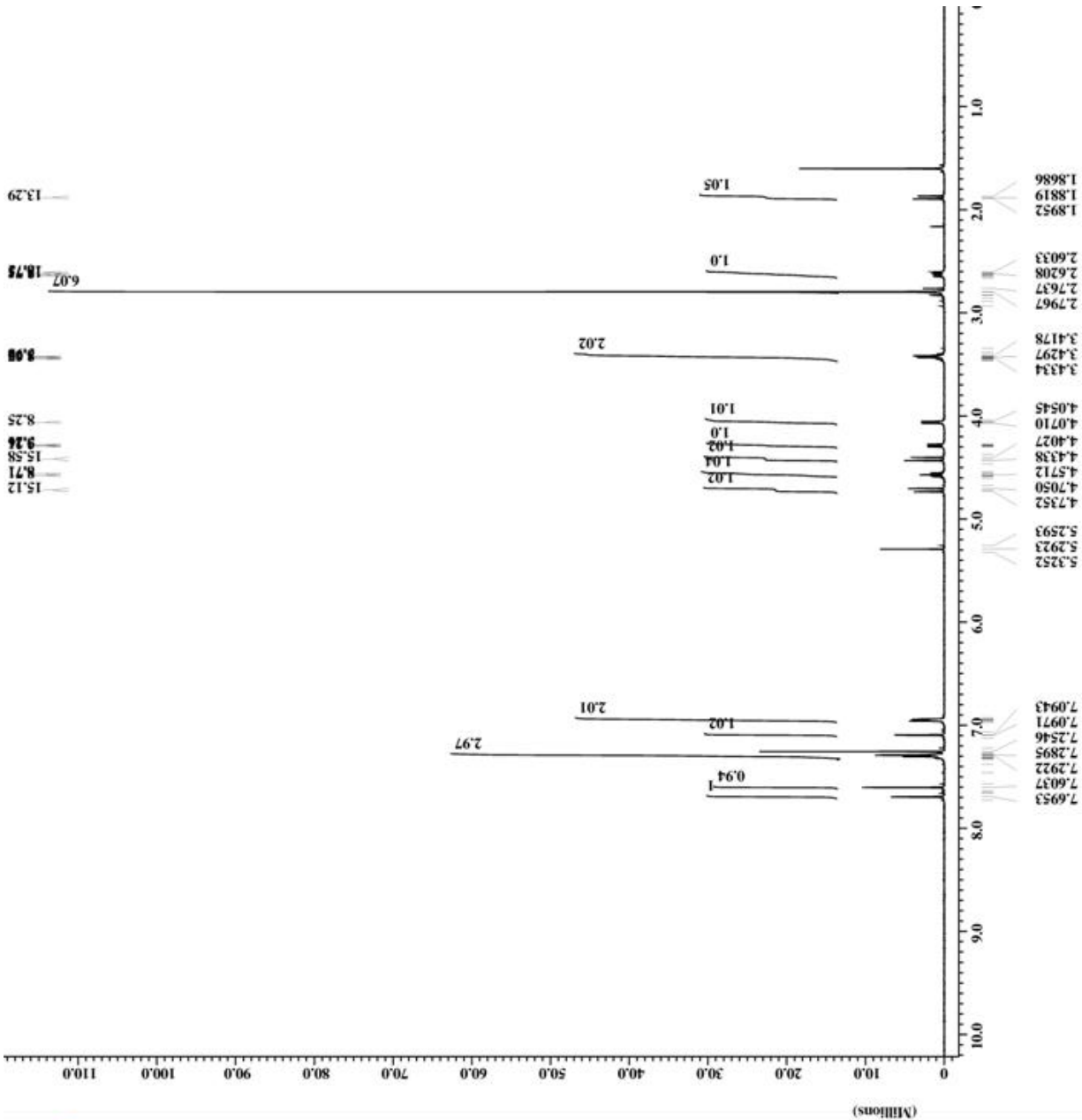
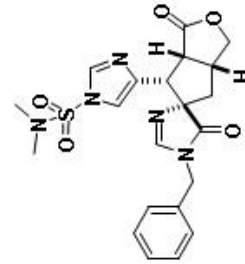
```

ilname      = sm_v_spiro_Bn_Dmas_pu
author      = delta
experiment  = single_pulse_exp
sample_id   = S8714680
solvent     = CHLOROFORM-D
revision    = 22-MAR-2009 05:49:02
evision_time = 28-MAR-2010 20:35:38
current_time = 28-MAR-2010 20:36:11

comment     = Single Pulse Experime
sta_format  = 1D REAL
im_size     = 16384
im_title    = 1H
im_units    = [ppm]
imensions  = X
ite         = Eclipse+ 500
spectrometer = DELTA_NMR

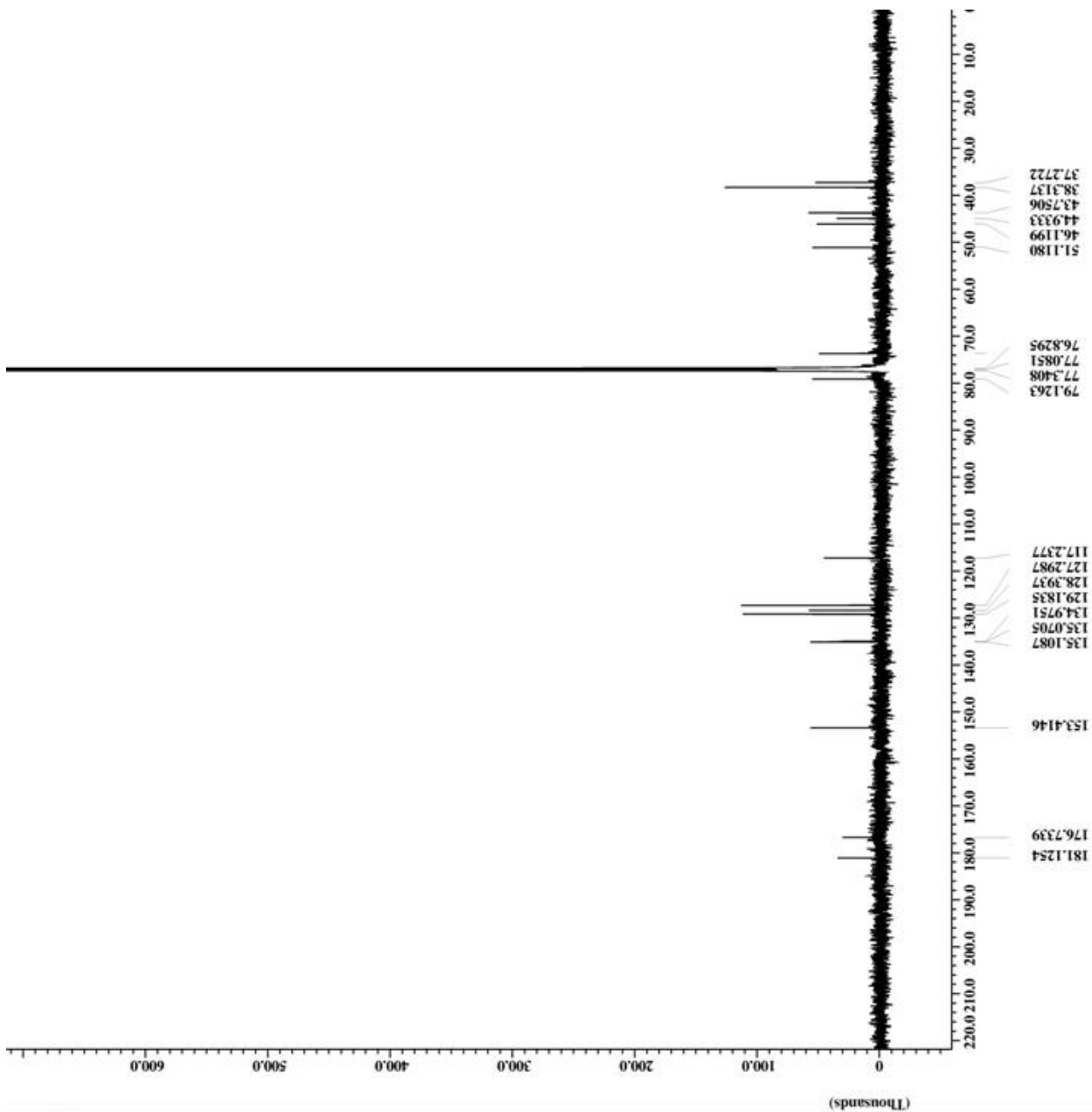
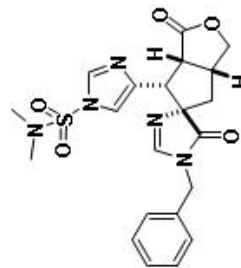
field_strength = 11.7473579[T] (500[MH
-acq_duration = 2.1823488[s]
-domain       = 1H
-freq         = 500.15991521[MHz]
-offset       = 5[ppm]
-points       = 16384
-prescans     = 0
-resolution   = 0.45822189[Hz]
-sweep        = 7.50750751[kHz]
lipped        = FALSE
bd_return     = 1
cans          = 18
stal_scans    = 18

-60_width    = 18.5[us]
-acq_time     = 2.1823488[s]
-angle        = 45[deg]
-pulse        = 9.25[us]
nitial_wait   = 3[s]
base_preset   = 3[us]
svr_gain      = 24
relaxation_delay = 4[s]
emp_get       = 26.1[dc]
nblank_time   = 2[us]
  
```





```
filename = sm_v_spiro_pure-2.jdf
author = delta
experiment = single_pulse_dec
sample_id = S871231
solvent = CHLOROFORM-D
reaction_time = 22-MAR-2009 12:55:48
revision_time = 28-MAR-2010 20:38:42
current_time = 28-MAR-2010 20:39:11
comment = single pulse decouple
ata_format = 1D COMPLEX
im_size = 65536
im_title = 13C
im_units = [ppm]
imensions = X
ite = Eclipse+ 500
spectrometer = DELTA_NMR
field_strength = 11.7473579[T] (500[MH
-acq_duration = 2.0840448[s]
-domain = 13C
-freq = 125.76529768[MHz]
-offset = 100[ppm]
-points = 65536
-prescans = 4
-resolution = 0.47983613[Hz]
-sweep = 31.44654088[KHz]
rr_domain = 1H
rr_freq = 500.15991521[MHz]
rr_offset = 5[ppm]
lipped = TRUE
bd_return = 10
cans = 5000
stal_scans = 5000
-90_width = 14.2[us]
-acq_time = 2.0840448[s]
-angle = 30[deg]
-pulse = 4.73333333[us]
nit1al_wait = 1[s]
be_time = 1[s]
base_preset = 3[us]
scvr_gain = 30
relaxation_delay = 2[s]
emp_get = 29.2[dc]
nblank_time = 2[us]
```





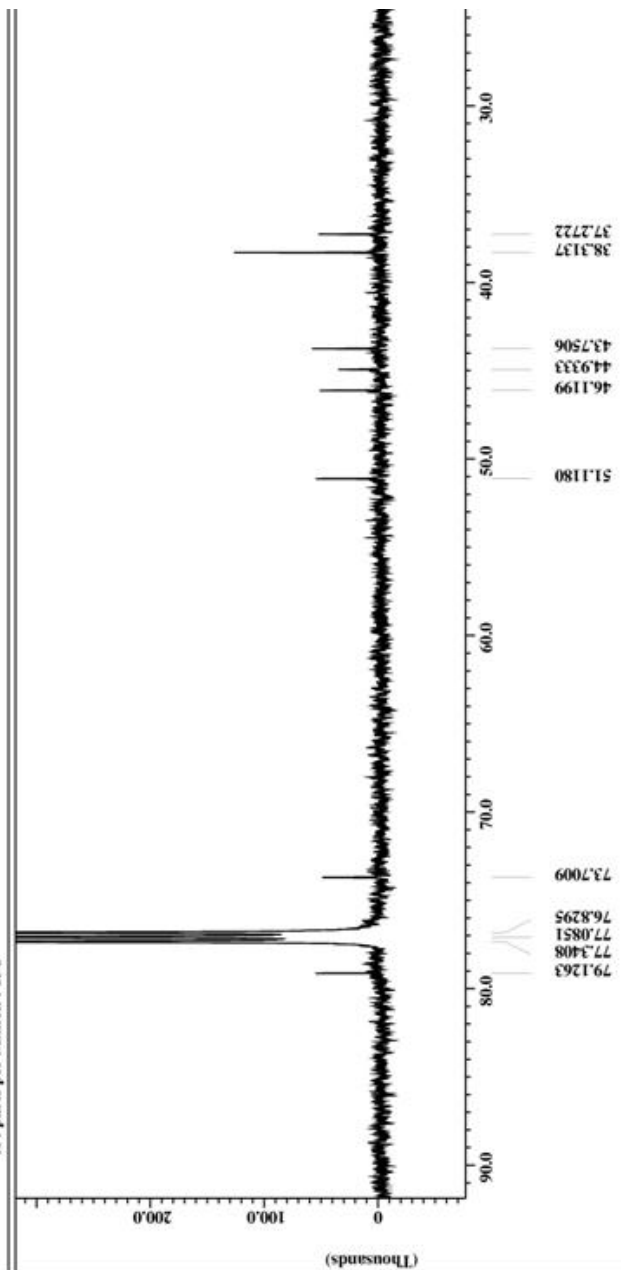
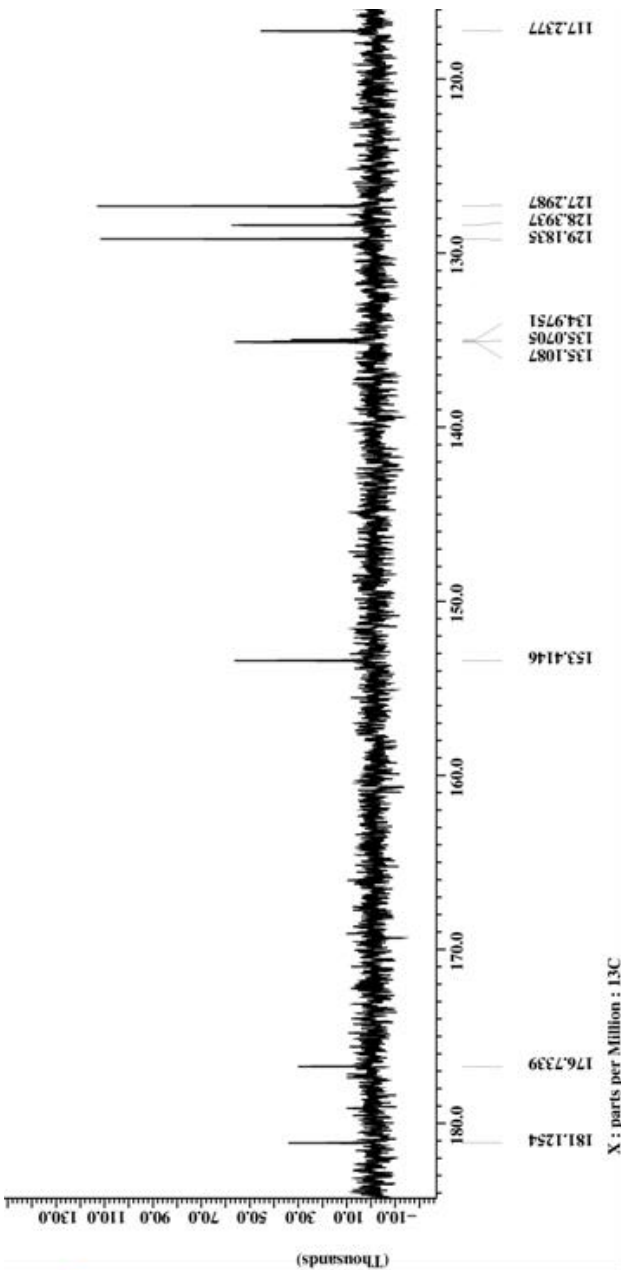
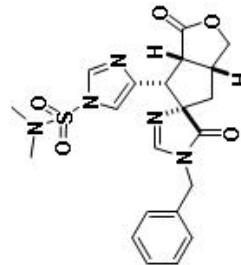
```

ilname      = sm_v_spiro_pure-2.jdf
author
experiment  = delta
sample_id   = S8717231
solvent     = CHLOROFORM-D
reaction_time = 22-MAR-2009 12:55:48
evision_time = 28-MAR-2010 20:38:42
current_time = 28-MAR-2010 20:40:42

comment     = single pulse decouple
ata_format  = ID COMPLEX
im_size     = 65536
im_title    = 13C
im_units    = [ppm]
imensions  = X
ite         = Eclipt+ 500
pctrometer = DELTA_RMR

field_strength = 11.7473579[T] (500[MH
-acq_duration = 2.0840448[s]
-domain      = 13C
-freq       = 125.76529768[MHz]
-offset     = 100[ppm]
-points     = 4
-prescans   = 4
-resolution = 0.47983613[Hz]
-sweep      = 31.44654088[KHz]
-rr_domain  = 1H
-rr_freq    = 500.15991521[MHz]
-rr_offset  = 5[ppm]
-lipped     = TRUE
-bd_return  = 10
-cans       = 5000
-stal_scans = 5000

-90_width   = 14.2[us]
-acq_time   = 2.0840448[s]
-angle      = 30[deg]
-pulse      = 4.73333333[us]
-nitral_wait = 1[s]
-be_time    = 1[s]
-base_preset = 3[us]
-scvr_gain  = 30
-relaxation_delay = 2[s]
-emp_get    = 29.2[dc]
-nblank_time = 2[us]
  
```



APPENDIX 40
¹H AND ¹³C NMR SPECTRUM OF
(5*R*^{*},6*R*^{*},7*R*^{*})-methyl-1-benzyl-7-(1-(*N,N*-dimethylsulfamoyl)-1*H*-imidazol-4-yl)-5-(hydroxymethyl)-4,5,6,7-tetrahydro-1*H*-benzo[d]imidazole-6-carboxylate (235)



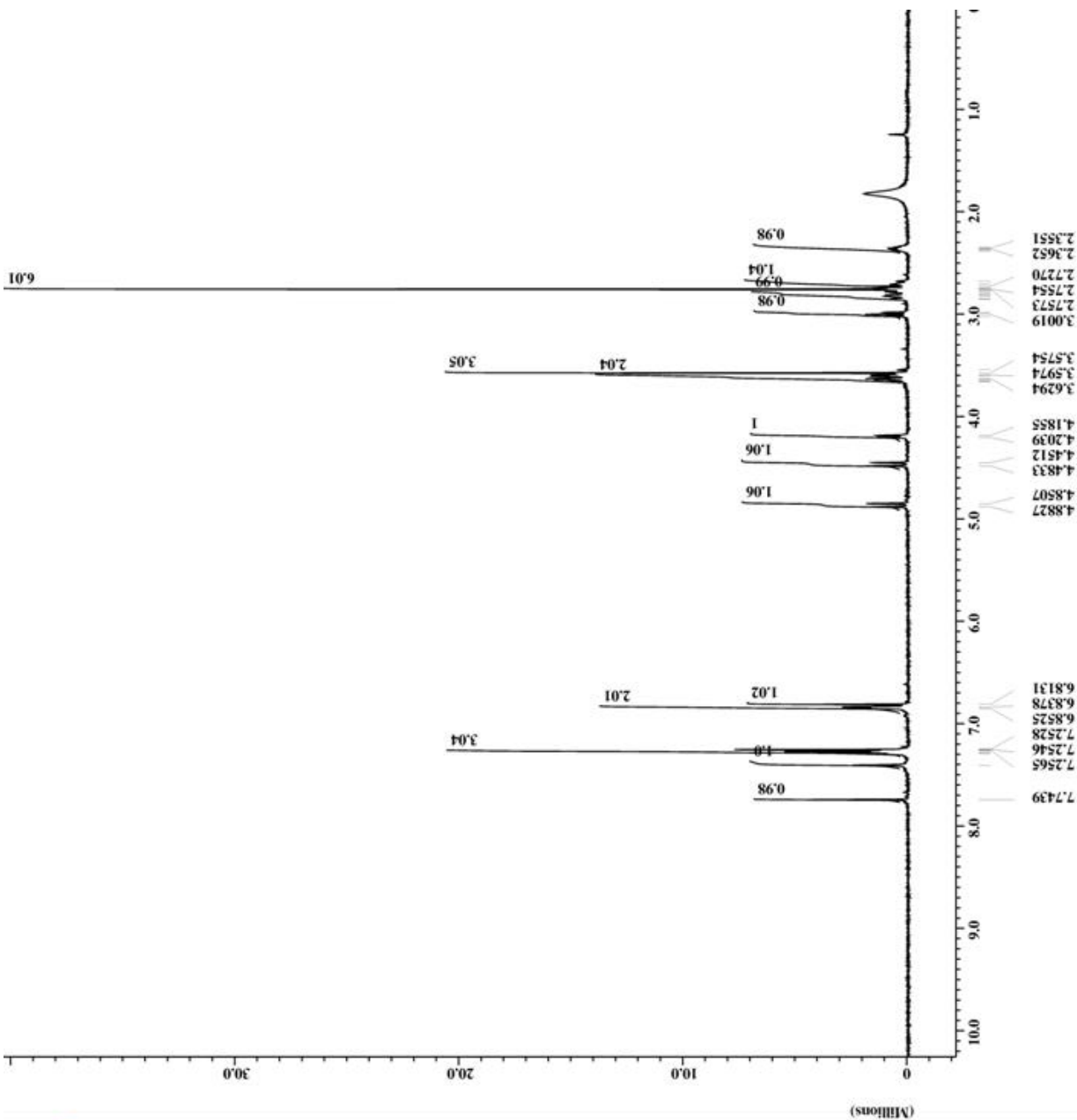
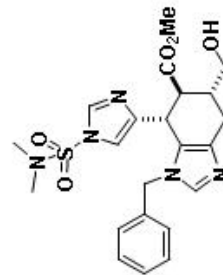
```

ilname      = sm_v_105_pure-2.jdf
author      = delta
experiment  = single_pulse_exp
sample_id   = S8763077
solvent     = CHLOROFORM-D
reaction_time = 23-MAR-2009 07:11:33
revision_time = 28-MAR-2010 20:46:28
current_time = 28-MAR-2010 20:46:53

comment     = Single Pulse Experiment
sta_format  = ID COMPLEX
im_size     = 16384
im_title    = 1H
im_units    = [ppm]
imensions  = X
ite         = Eclipse+ 500
spectrometer = DELTA_NMR

field_strength = 11.7473579[T] (500[MH
-acq_duration = 2.1823488[s]
-domain       = 1H
-freq         = 500.15991521[MHz]
-offset       = 5[ppm]
-points       = 16384
-prescans     = 0
-resolution   = 0.45822189[Hz]
-sweep        = 7.50750751[kHz]
-lipped       = FALSE
-bd_return    = 1
-cans         = 16
-stal_scans   = 16

-90_width    = 18.5[us]
-acq_time     = 2.1823488[s]
-angle        = 45[deg]
-pulse        = 7.25[us]
-pitch_wait   = 3[s]
-base_preset  = 5[us]
-svr_gain     = 24
-relaxation_delay = 4[s]
-emp_get      = 26[dc]
-nblank_time  = 2[us]
  
```





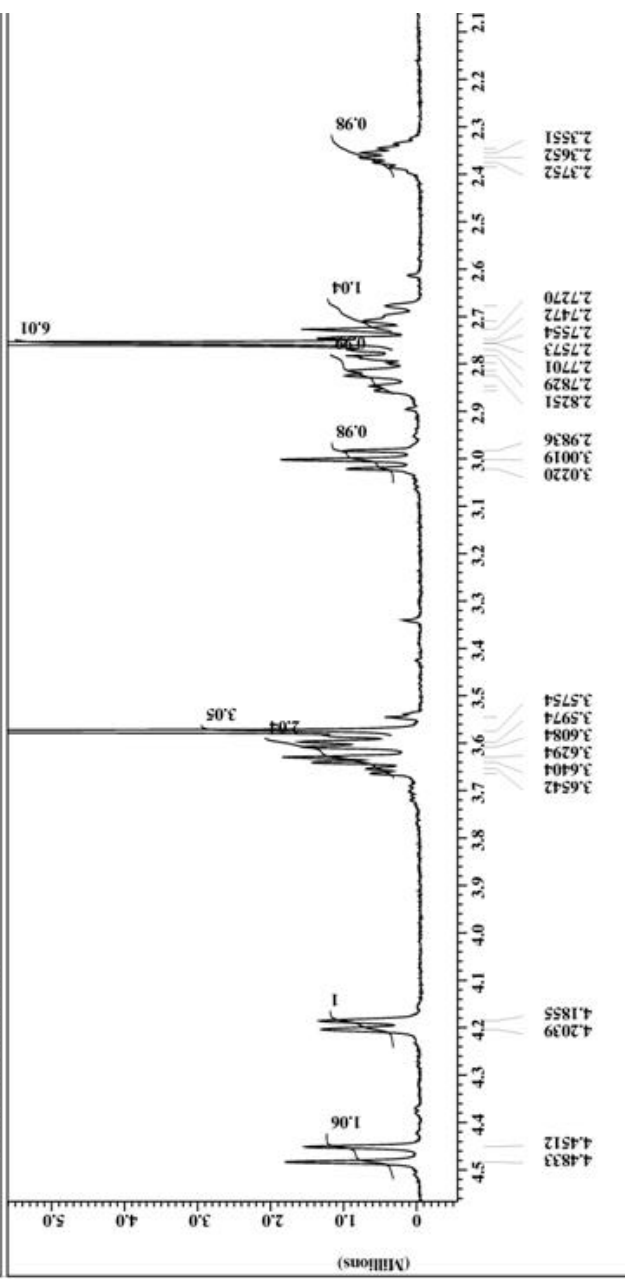
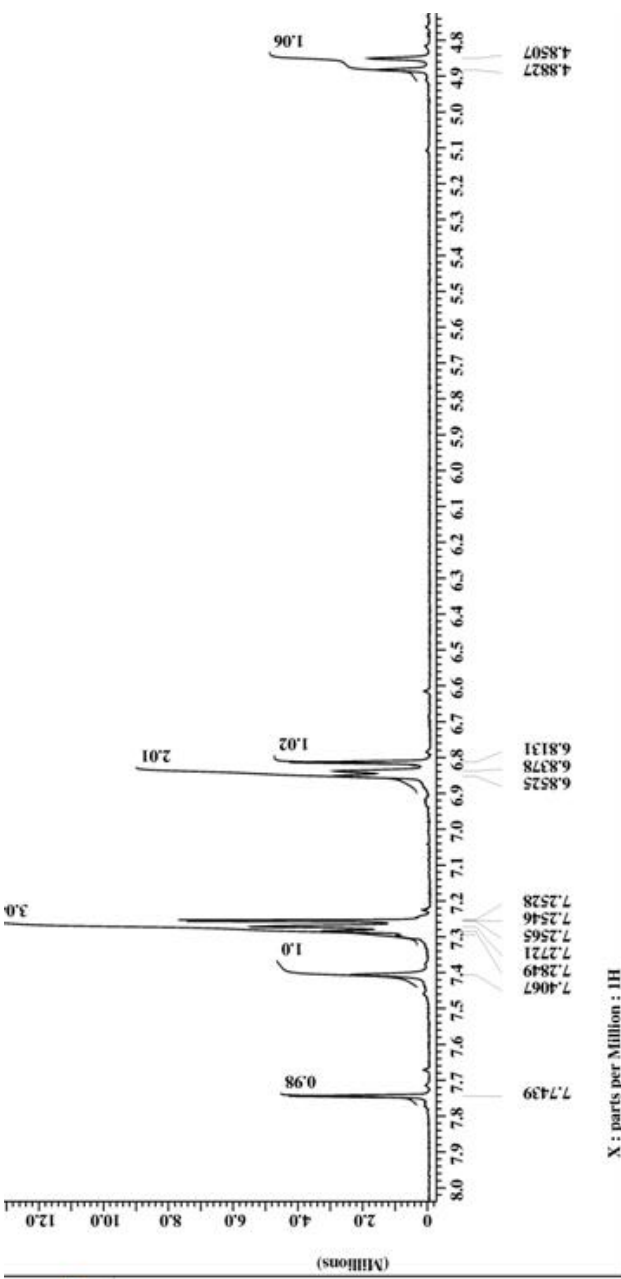
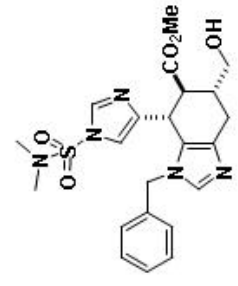
```

filename = sm_V_105_pure-2.jdf
author = delta
experiment = single_pulse_exp
sample_id = S9763077
solvent = CHLOROFORM-D
reaction_time = 23-MAR-2009 07:11:33
revision_time = 28-MAR-2010 20:46:28
current_time = 28-MAR-2010 20:48:59

comment = Single Pulse Experiment
ata_format = ID COMPLEX
in_size = 16384
in_title =
in_units =
inmensions = X
ite = Eclipse+ 500
spectrometer = DELTA_NMR

field_strength = 11.747379[T] (500[MH
acq_duration = 2.1823486[s]
domain =
freq = 500.15991521[MHz]
offset = 5[ppm]
points = 16384
prescans = 0
resolution = 0.45822189[Hz]
sweep = 7.50750751[kHz]
tipped = FALSE
od_return = 1
cans = 16
total_scans = 16

_90_width = 18.5[us]
acq_time = 2.1823486[s]
angle = 45[deg]
pulse = 7.25[us]
p1 = 1[s]
base_preset = 2[us]
ecvr_gain = 24
relaxation_delay = 4[s]
emp_get = 26[dc]
nblank_time = 2[us]
  
```





```

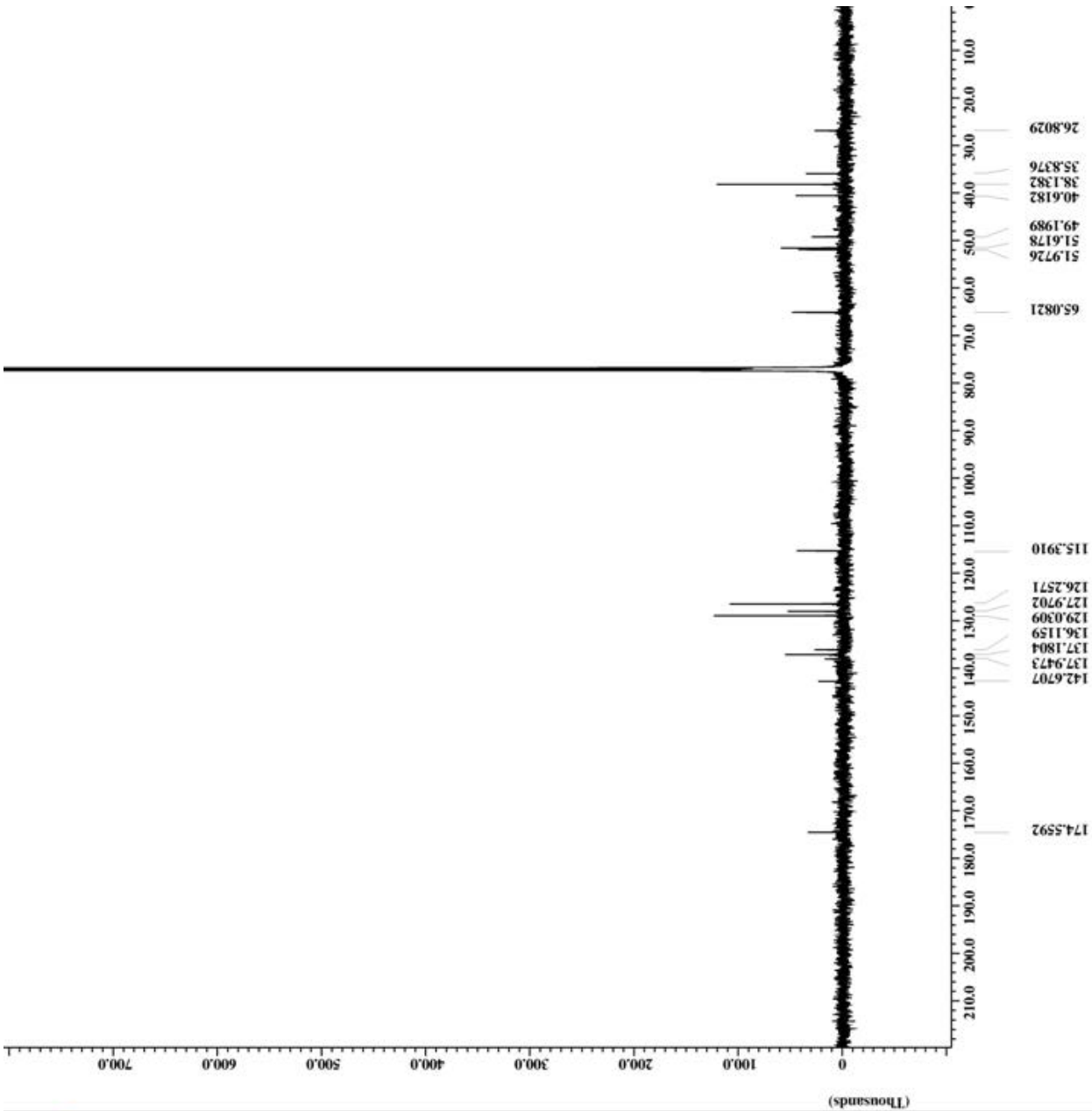
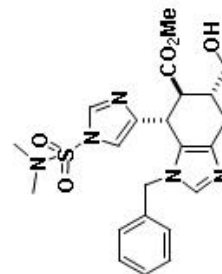
filename = sm_V_105_pure-2.jdf
author = delta
experiment = single_pulse_dec
sample_id = S8765343
solvent = CHLOROFORM-D
reaction_time = 23-MAR-2009 14:19:47
acquisition_time = 23-MAR-2009 08:30:23
current_time = 28-MAR-2010 20:51:12

comment = single pulse decouple
ata_format = ID COMPLEX
im_size = 65536
im_title = 13C
im_units = [ppm]
imensions = X
ite = Eclipse+ 500
spectrometer = DELTA_NMR

field_strength = 11.7473579[T] (500[MH
acq_duration = 2.0840448[s]
domain = 13C
freq = 125.76529768[MHz]
offset = 100[ppm]
points = 65536
prescans = 4
resolution = 0.47983613[Hz]
sweep = 31.44654088[kHz]
tr_domain = 1H
tr_freq = 500.15991521[MHz]
tr_offset = 5[ppm]
tipped = TRUE
bd_return = 10
cans = 5000
ctal_scans = 5000

_90_width = 14.2[us]
acq_time = 2.0840448[s]
angle = 30[deg]
pulse = 4.73333333[us]
nitai_wait = 1[s]
pe_time = 1[s]
base_preset = 3[us]
scvr_gain = 30
relaxation_delay = 2[s]
emp_get = 29.3[dc]
nblank_time = 2[us]

```



APPENDIX 41
¹H AND ¹³C NMR SPECTRUM OF
(5*S*^{*},6*R*^{*},7*R*^{*},8*R*^{*})-methyl-3-benzyl-6-(1-(*N,N*-dimethylsulfamoyl)-1*H*-imidazol-4-yl)-8-(hydroxymethyl)-4-oxo-1,3-diazaspiro[4.4]non-1-ene-7-carboxylate (236)



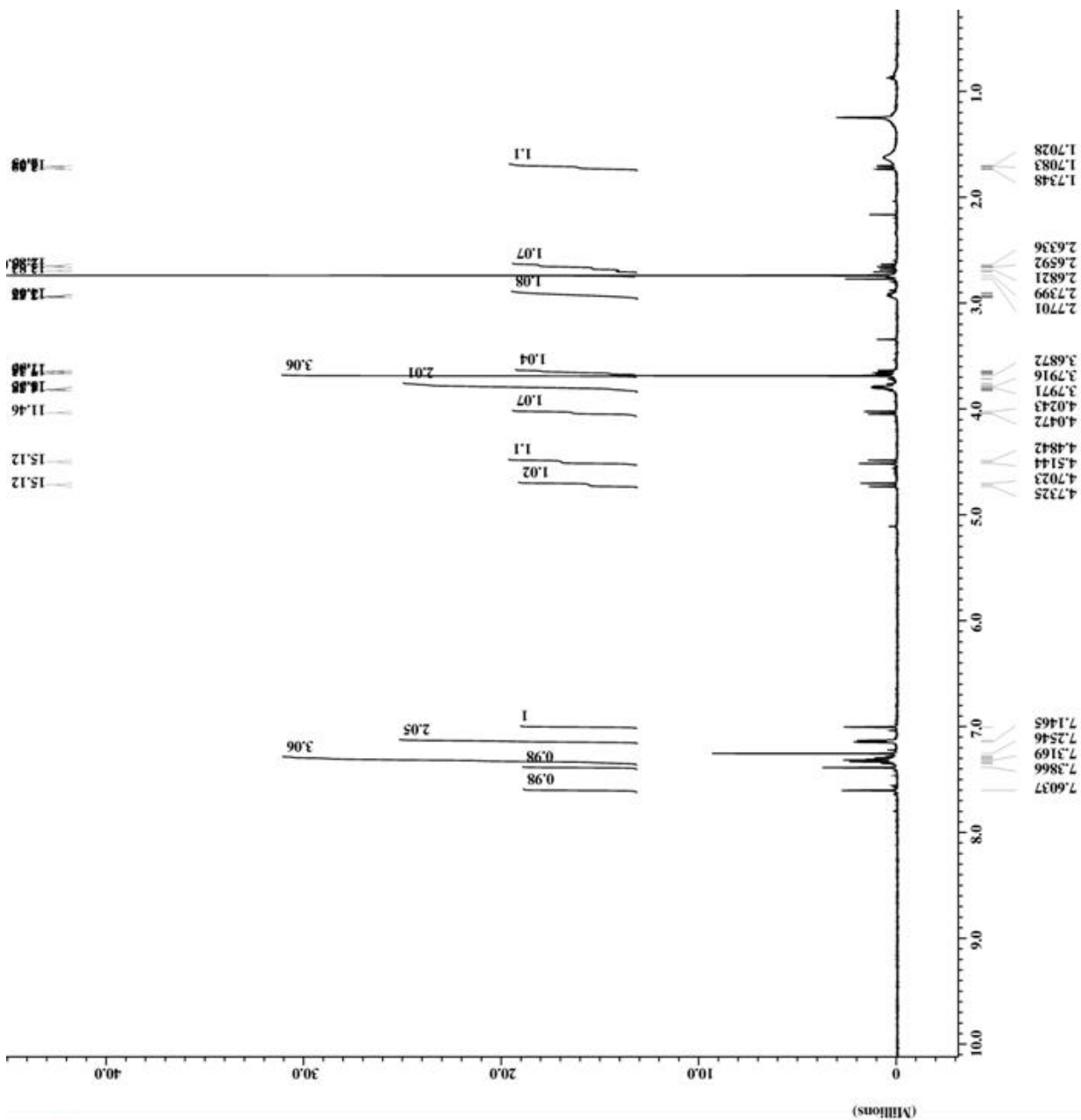
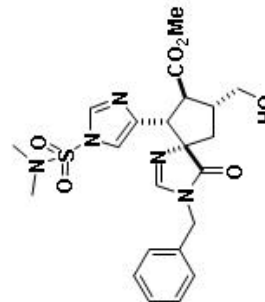
```

ilname      = sm_v_104_PURE-4.jdf
author      = delta
experiment  = single_pulse_exp
sample_id   = S#598701
solvent     = CHLOROFORM-D
reaction_time = 4-APR-2009 03:00:34
evision_time = 28-MAR-2010 20:56:54
current_time = 28-MAR-2010 20:57:24

comment     = Single Pulse Experiment
sta_format  = ID COMPLEX
im_size     = 16384
im_title    = 1H
im_units    = [ppm]
imensions  = X
ite        = Eclipse+ 500
pctrometer = DELTA_NMR

field_strength = 11.7473579[T] (500[M]
acq_duration   = 2.1823488[s]
domain        = 1H
freq          = 500.15991521[M]
offset        = 5[ppm]
points        = 16384
prescans      = 0
resolution    = 0.45822189[Hz]
sweep         = 7.50750751[kHz]
lipped        = FALSE
bd_return     = 1
cans          = 1
stal_scans    = 16

_90_width     = 18.5[us]
acq_time      = 2.1823488[s]
angle         = 45[deg]
pulse         = 9.25[us]
p1          = 3[s]
base_preset   = 5[us]
scvr_gain     = 23
relaxation_delay = 4[s]
emp_get       = 24.9[dc]
nblank_time   = 2[us]
  
```





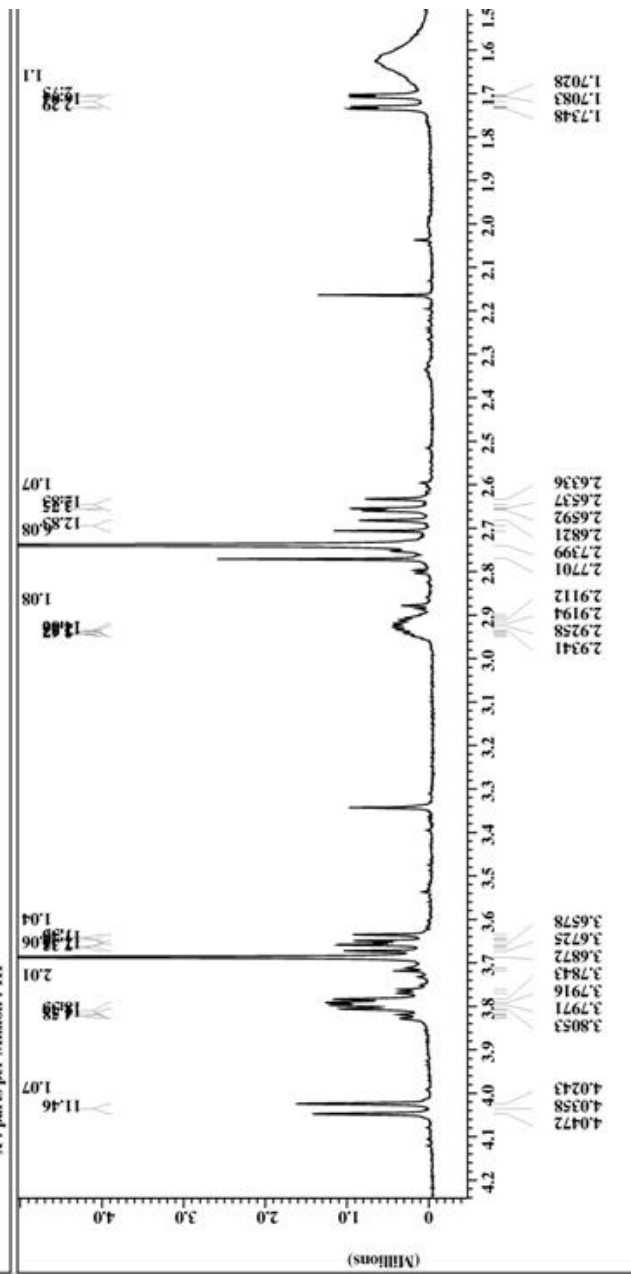
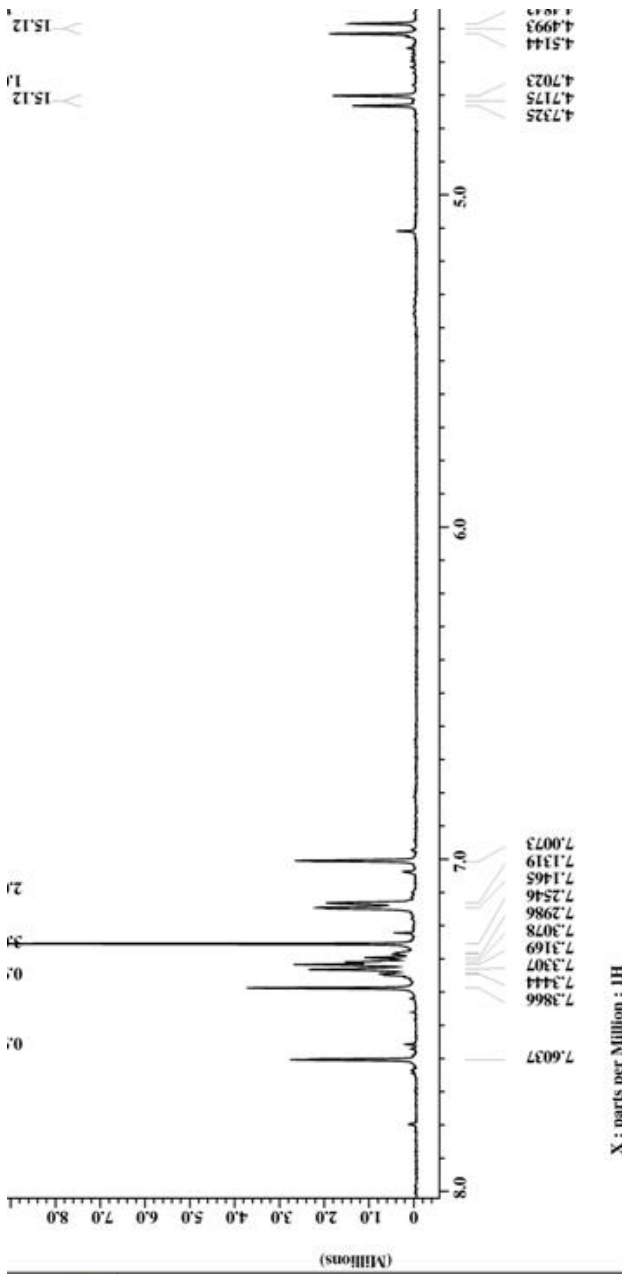
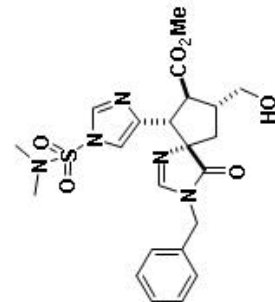
```

ilname      = sm_v_104_PURE-4_.jdf
author      = delta
experiment  = single_pulse_exp
sample_id   = S#598701
solvent     = CHLOROFORM-D
revision    = 4-APR-2009 03:00:34
evision     = 28-MAR-2010 20:56:54
current_time = 28-MAR-2010 20:59:24

comment     = Single Pulse Experime
sta_format  = ID COMPLEX
im_size     = 16384
im_title    = 1H
im_units    = [ppm]
imensions  = X
ite         = Eclipse+ 500
spectrometer = DELTA_NMR

field_strength = 11.7473579[T] (500[MH
-acq_duration  = 2.1823488[s]
-domain        = 1H
-freq          = 500.15991521[MHz]
-offset        = 16384
-points        = 0
-prescans      = 0
-resolution    = 0.45822189[Hz]
-sweep         = 7.50750751[kHz]
-tipped        = FALSE
-td_return     = 1
-cans          = 16
-stal_scans    = 1

-60_width     = 18.5[us]
-acq_time     = 2.1823488[s]
-angle        = 45[deg]
-pulse        = 9.25[us]
-pitch        = 5[s]
-base_preset  = 5[us]
-svr_gain     = 23
-relaxation_delay = 4[s]
-emp_get      = 24.9[dc]
-nblank_time  = 2[us]
  
```





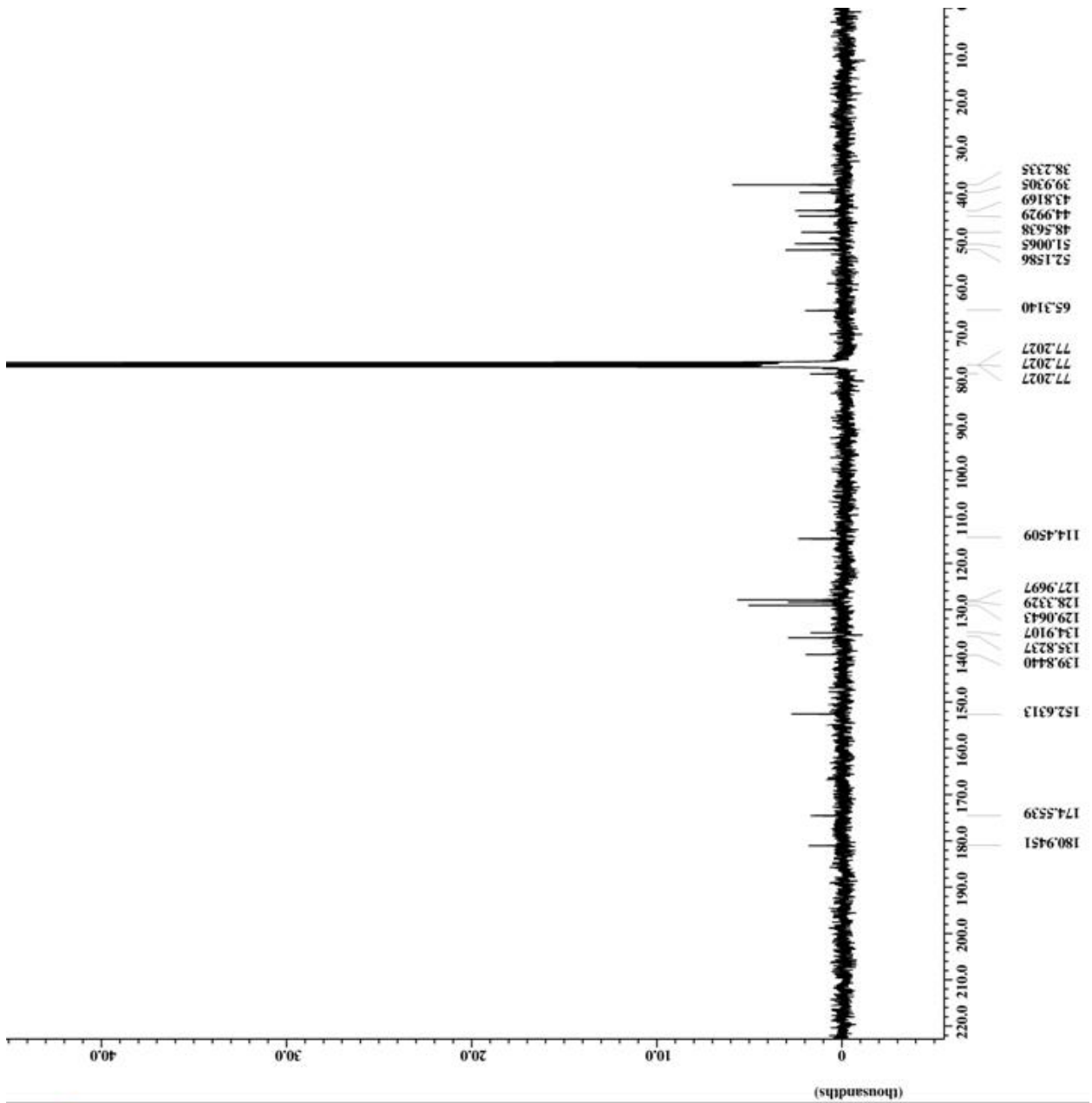
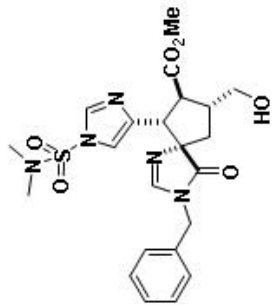
```

ilname      = sm_v_104_pure_ii-2_jd
author      = delta
experiment  = single_pulse_dec
sample_id   = S#400830
solvent     = CHLOROFORM-D
reaction_time = 21-MAR-2009 16:06:14
revision_time = 21-MAR-2009 15:45:31
current_time = 28-MAR-2010 21:01:33

comment     = single pulse decouple
ate_format  = ID COMPLEX
im_size     = 52428
im_title    = 13C
im_units    = [ppm]
imensions  = X
ite         = ECX 300
pctrometer  = DELTA2_NMR

ield_strength = 7.0586013[T] (300[MHz]
-acq_duration = 2.76824064[s]
-domain       = 13C
-freq         = 75.56823426[MHz]
-offset       = 100[ppm]
-points       = 65536
-prescans     = 4
-resolution   = 0.36124027[Hz]
-sweep        = 23.67424242[KHz]
-rr_domain    = 1H
-rr_freq      = 300.52965592[MHz]
-rr_offset    = 5[ppm]
-lipped       = FALSE
-bd_return    = 10
-cans         = 2859
-stal_scans   = 2859

-90_width    = 9.75[us]
-acq_time     = 2.76824064[s]
-angle        = 30[deg]
-atn          = 8[db]
-pulse        = 3.25[us]
-rr_atn_dec   = 25[db]
-rr_atn_noe   = 25[db]
-rr_noise     = WALTZ
-accoupling   = TRUE
-nitral_wait  = 1[s]
-be           = TRUE
-be_time      = 3[s]
-scvr_gain    = 50
-relaxation_delay = 3[s]
-epitation_time = 5.76824064[s]
-emp_get      = 23.1[dc]
  
```





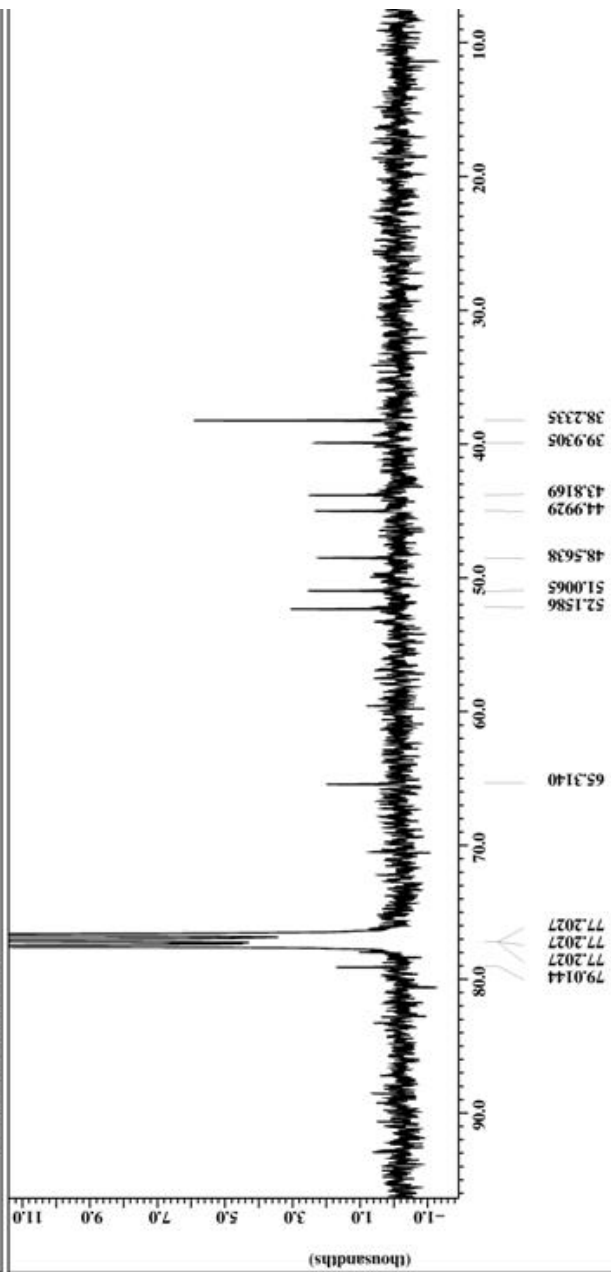
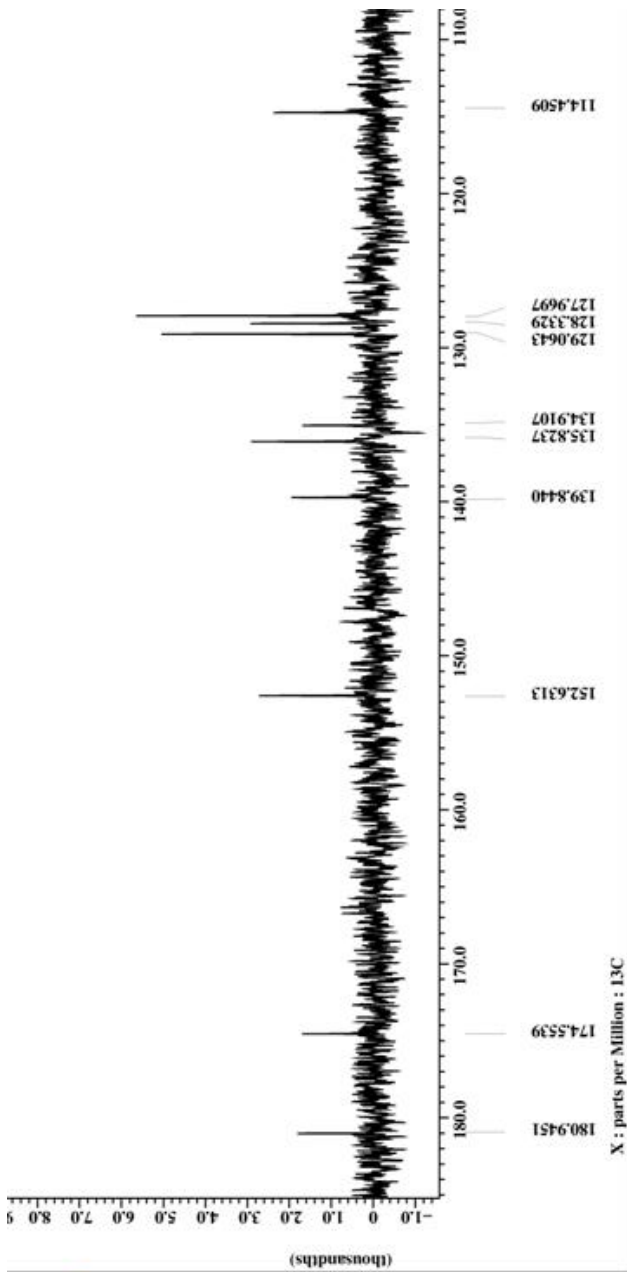
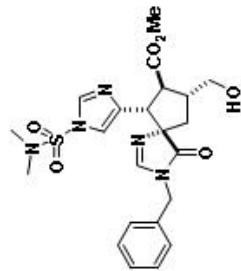
```

ilname      = sm_v_104_pure_ii-2_jd
author      = delta
experiment  = single_pulse_dec
sample_id   = S#400830
solvent     = CHLOROFORM-D
reaction_time = 21-MAR-2009 16:06:14
revision_time = 21-MAR-2009 15:45:31
current_time = 28-MAR-2010 21:03:00

comment     = single pulse decouple
ate_format  = ID COMPLEX
ate_size    = 52428
im_title    = 13C
im_units    = [ppm]
imensions   = X
ite         = ECK 300
pctrometer  = DELTA2_NMR

field_strength = 7.0586013[T] (300[MHz]
-acq_duration  = 2.76824064[s]
-domain        = 13C
-freq          = 75.56823426[MHz]
-offset        = 100[ppm]
-points        = 65536
-prescans      = 4
-resolution    = 0.36124027[Hz]
-sweep         = 23.67424242[KHz]
-rr_domain     = 1H
-rr_freq       = 300.52965592[MHz]
-rr_offset     = 5[ppm]
-lipped        = FALSE
-bd_return     = 10
-cans          = 2859
-stal_scans    = 2859

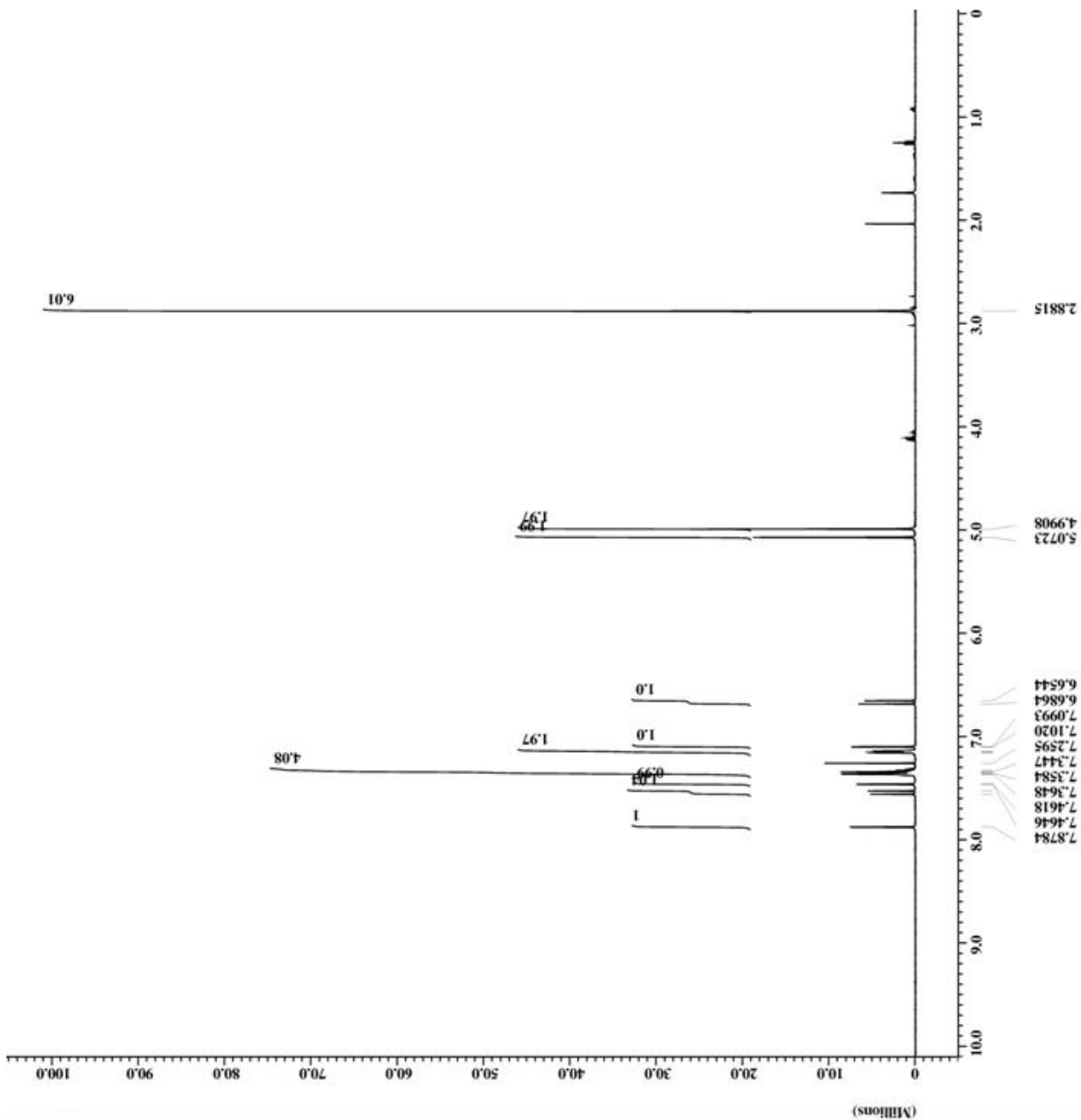
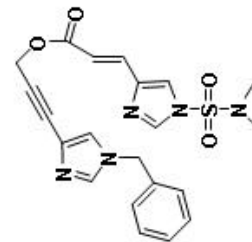
-90_width     = 9.75[us]
-acq_time     = 2.76824064[s]
-angle        = 30[deg]
-atn          = 8[db]
-pulse        = 3.25[us]
-rr_atn_dec   = 25[db]
-rr_atn_noe   = 25[db]
-rr_noise     = WALTZ
-acoupling    = TRUE
-nitail_wait  = 1[s]
-be           = TRUE
-be_time      = 3[s]
-scvr_gain    = 50
-relaxation_delay = 3[s]
-epitation_time = 5.76824064[s]
-emp_get      = 23.1[dc]
  
```



APPENDIX 42
¹H AND ¹³C NMR SPECTRUM OF
(*E*)-3-(1-benzyl-1H-imidazol-4-yl)prop-2-ynyl-3-(1-(*N,N*-dimethylsulfamoyl)-
1H-imidazol-4-yl)acrylate (239)



```
filename = sm_VI_78_pure-3.jdf
author = delta
experiment = single_pulse_exp
sample_id = S8587591
solvent = CHLOROFORM-D
revision_time = 26-DEC-2009 08:01:09
evision_time = 28-MAR-2010 21:16:38
current_time = 28-MAR-2010 21:17:02
comment = Single Pulse Experiment
ata_format = ID COMPLEX
im_size = 16384
im_title = 1H
im_units = [ppm]
ite = X
spectrometer = Eclipse+ 500
field_strength = 11.7465928[T] (500[MH
-acq_duration = 2.1839872[s]
-domain = 1H
-freq = 500.12734003[MHz]
-offset = 5[ppm]
-points = 16384
-prescans = 0
-resolution = 0.45787814[Hz]
-sweep = 7.50187547[kHz]
lipped = FALSE
bd_return = 1
cans = 8
stal_scans = 8
-90_width = 18.5[us]
-acq_time = 2.1839872[s]
-angle = 45[deg]
-pulse = 7.25[us]
nitial_wait = 3[s]
base_preset = 5[us]
svr_gain = 23
relaxation_delay = 4[s]
emp_get = 25.8[dc]
nblank_time = 2[us]
```





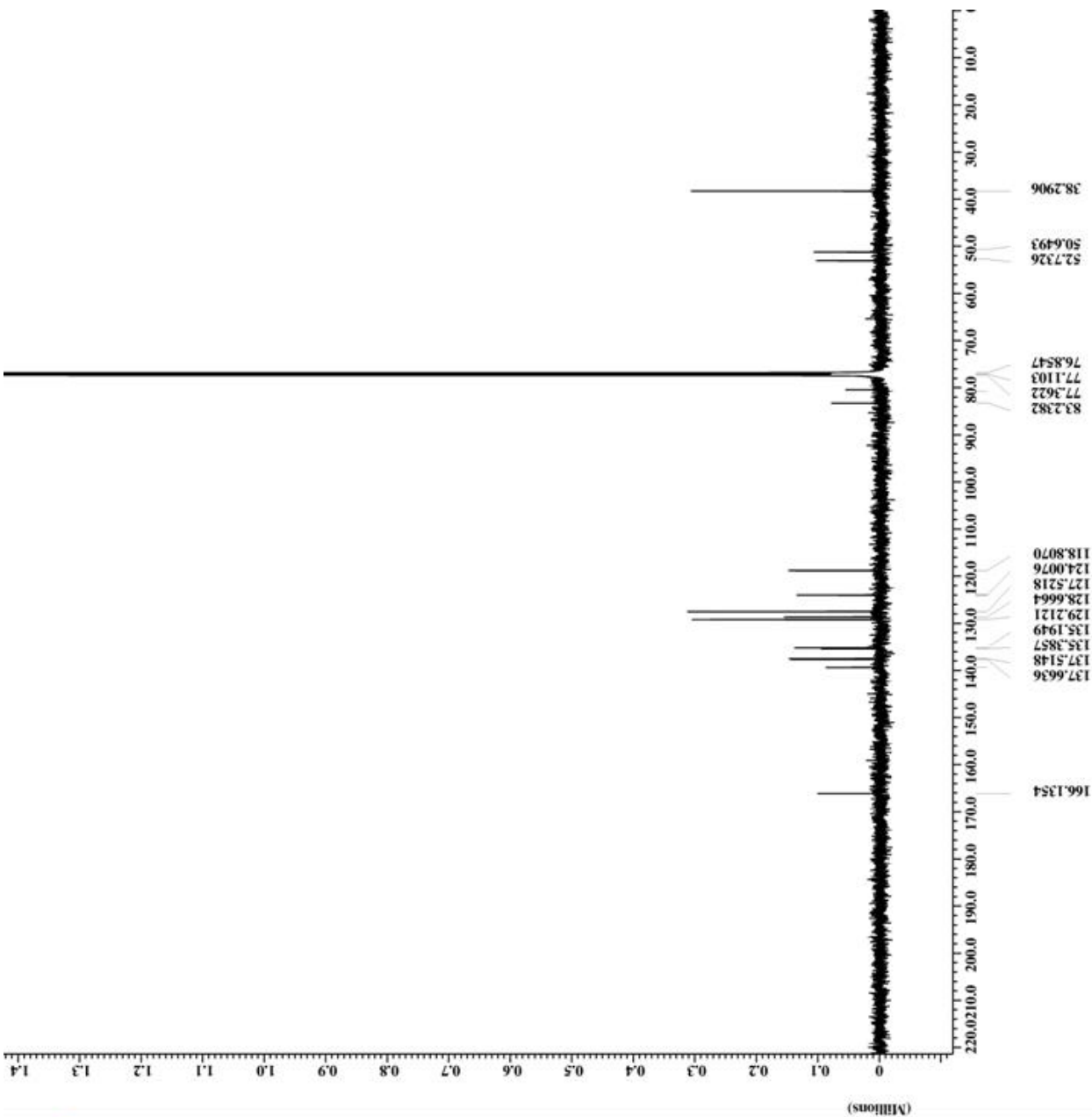
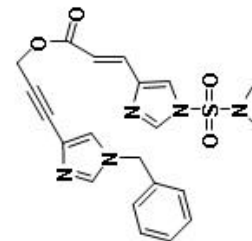
```

ilname      = sm_VI_78_pure-2.jdf
author      = delta
experiment  = single_pulse_dec
sample_id   = S858949
solvent     = CHLOROFORM-D
reaction_time = 26-DEC-2009 10:11:45
revision_time = 26-DEC-2009 16:23:16
current_time = 28-MAR-2010 21:19:48

comment     = single pulse decouple
ata_format  = ID_COMPLEX
im_size     = 65536
im_title    = 13C
im_units    = [ppm]
imensions  = X
ite        = Eclipt+ 500
pctometer  = DELTA_NMR

field_strength = 11.7465928[T] (500[MH
-acq_duration = 2.0840448[s]
-domain      = 13C
-freq       = 125.75710665[MHz]
-offset     = 100[ppm]
-points     = 65536
-prescans   = 4
-resolution = 0.47983613[Hz]
-sweep      = 31.44654088[KHz]
-rr_domain  = 1H
-rr_freq    = 500.12734003[MHz]
-rr_offset  = 5[ppm]
-lipped     = TRUE
-bd_return  = 10
-cans       = 1500
-stal_scans = 1500

-90_width   = 14.2[us]
-acq_time   = 2.0840448[s]
-angle      = 30[deg]
-pulse      = 4.73333333[us]
-nitral_wait = 1[s]
-be_time    = 1[s]
-base_preset = 3[us]
-scvr_gain  = 30
-relaxation_delay = 2[s]
-emp_get    = 28.5[dc]
-nblank_time = 2[us]
  
```





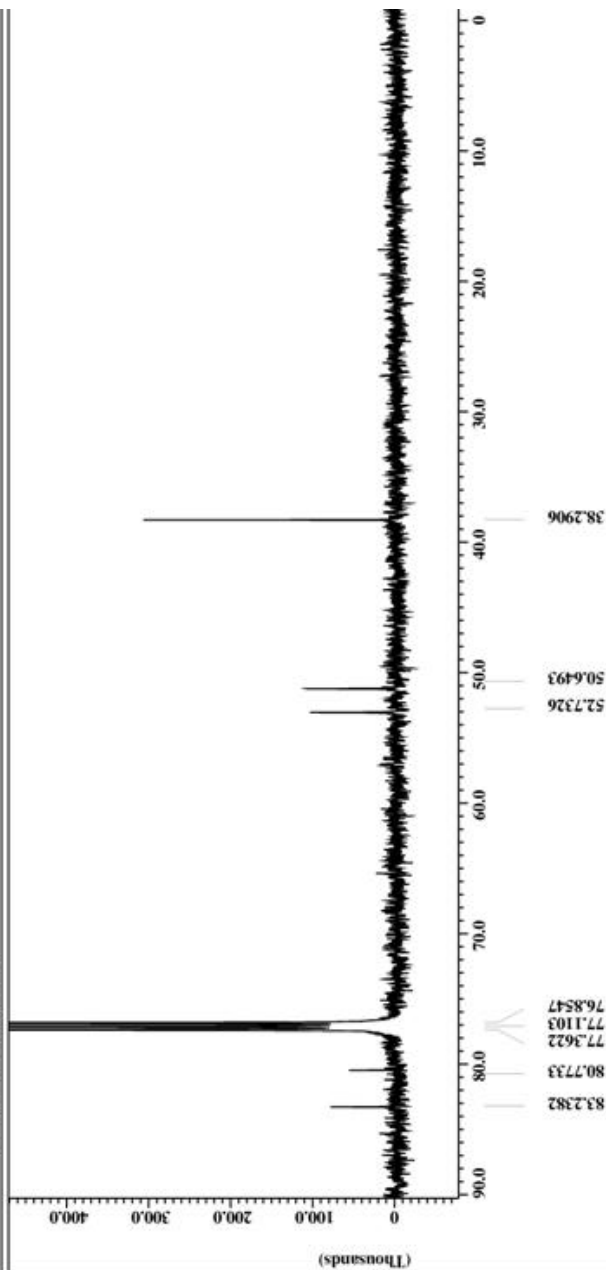
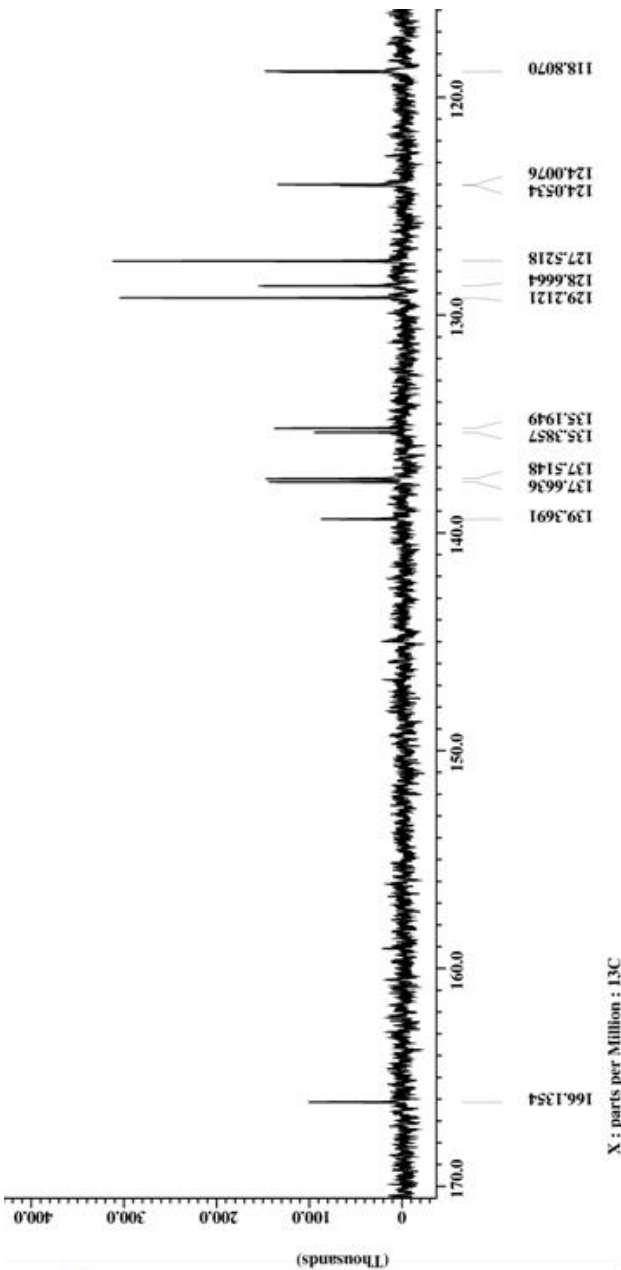
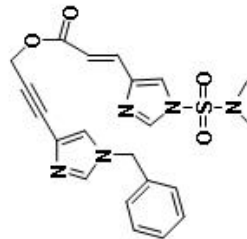
```

ilname      = sm_VI_78_pure-2.jdf
author      =
xperiment   = delta_pulse_dec
sample_id   = S858949
solvent     = CHLOROFORM-D
reaction_time = 26-DEC-2009 10:11:45
evision_time = 26-DEC-2009 16:23:16
urrent_time = 28-MAR-2010 21:21:28

cment       = single pulse decouple
ata_format  = 1D COMPLEX
im_size     = 65536
im_title    = 13C
im_units    = [ppm]
imensions   = X
ite         = Eclipset 500
pctrometer  = DELTA_RMR

ield_strength = 11.7465928[T] (500[MH
-acq_duration = 2.0840448[s]
-domain       = 13C
-freq         = 125.75710665[MHz]
-offset       = 100[ppm]
-points       = 65536
-prescans     = 4
-resolution   = 0.47983613[Hz]
-sweep        = 31.44654088[KHz]
rr_domain    = 1H
rr_freq      = 500.12734003[MHz]
rr_offset    = 5[ppm]
lipped       = TRUE
bd_return    = 10
cans         = 1500
stal_scans   = 1500

_90_width    = 14.2[us]
-acq_time     = 2.0840448[s]
-angle        = 30[deg]
-pulse        = 4.73333333[us]
nitial_wait  = 1[s]
be_time      = 1[s]
base_preset  = 3[us]
scvr_gain    = 30
elaxation_delay = 2[s]
emp_get      = 28.5[dc]
nblank_time  = 2[us]
  
```



APPENDIX 43
¹H AND ¹³C NMR SPECTRUM OF
(*E*)-3-(1-(*N,N*-dimethylsulfamoyl)-1*H*-imidazol-4-yl)prop-2-ynyl-3-(1-benzyl-1*H*-imidazol-4-yl)acrylate (242)



```

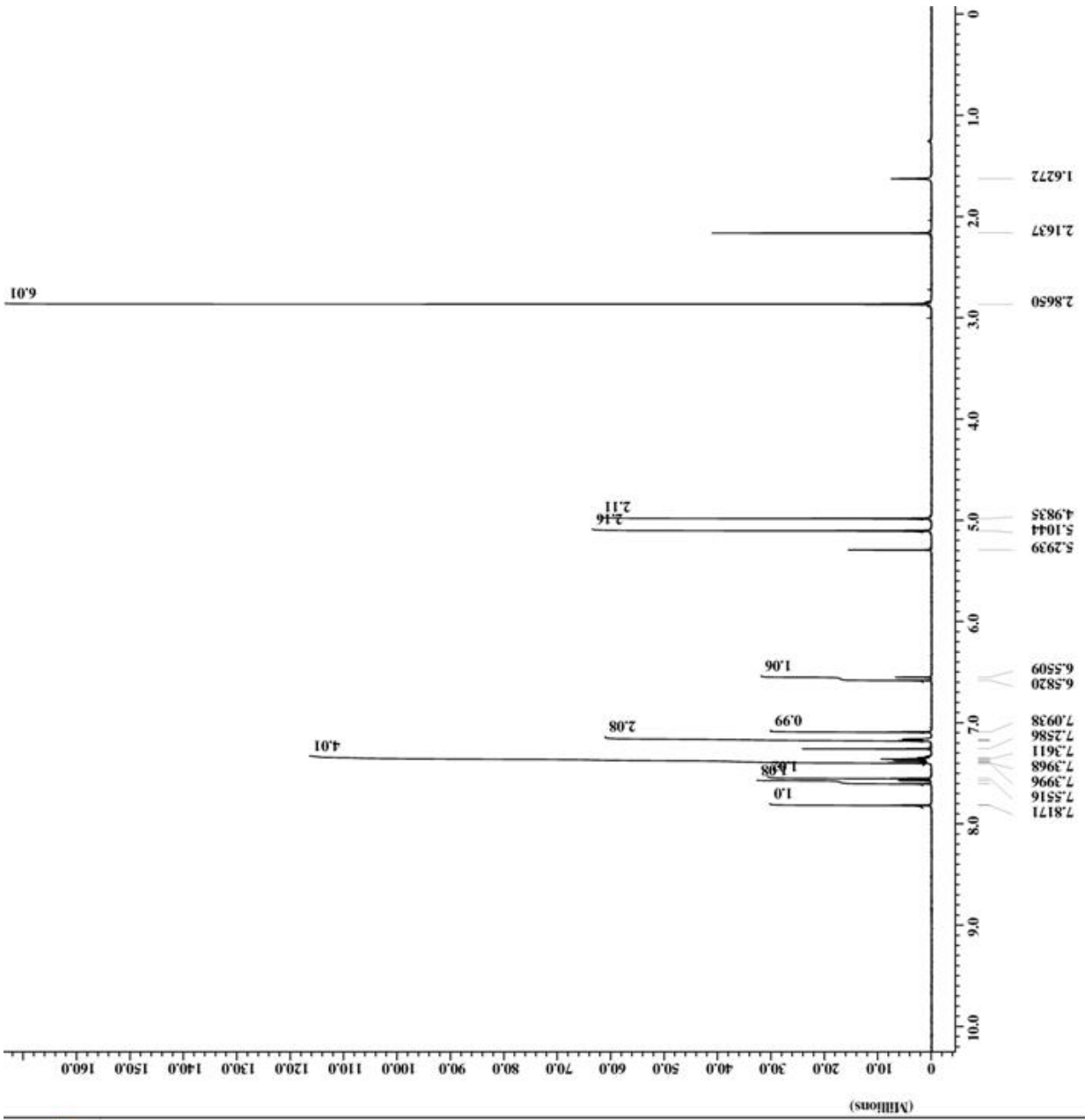
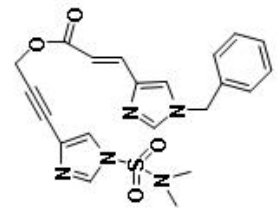
filename = sm_VI_80_pure-4.jdf
author = delta
experiment = single_pulse.exp
sample_id = S#505736
solvent = CHLOROFORM-D
reaction_time = 30-DEC-2009 05:46:49
revision_time = 28-MAR-2010 21:26:19
current_time = 28-MAR-2010 21:26:52

comment = Single Pulse Experiment
ata_format = ID COMPLEX
in_size = 16384
in_title = 1H
in_units = [ppm]
imsions = X
ite = Eclipse+ 500
spectrometer = DELTA_NMR

field_strength = 11.7465928[T] (500[MH
-acq_duration = 2.1839872[s]
-domain = 1H
-freq = 500.12734003[MHz]
-gamma = 5[ppm]
-points = 16384
-prescans = 0
-resolution = 0.45787814[Hz]
-sweep = 7.50187547[kHz]
-lipped = FALSE
od_return = 1
cans = 8
otal_scans = 8

_90_width = 18.5[us]
-acq_time = 2.1839872[s]
-angle = 45[deg]
-pulse = 7.25[us]
ntial_wait = 1[s]
base_preset = 2[us]
ecvr_gain = 25
axiation_delay = 4[s]
emp_get = 29.8[dc]
nblank_time = 2[us]

```



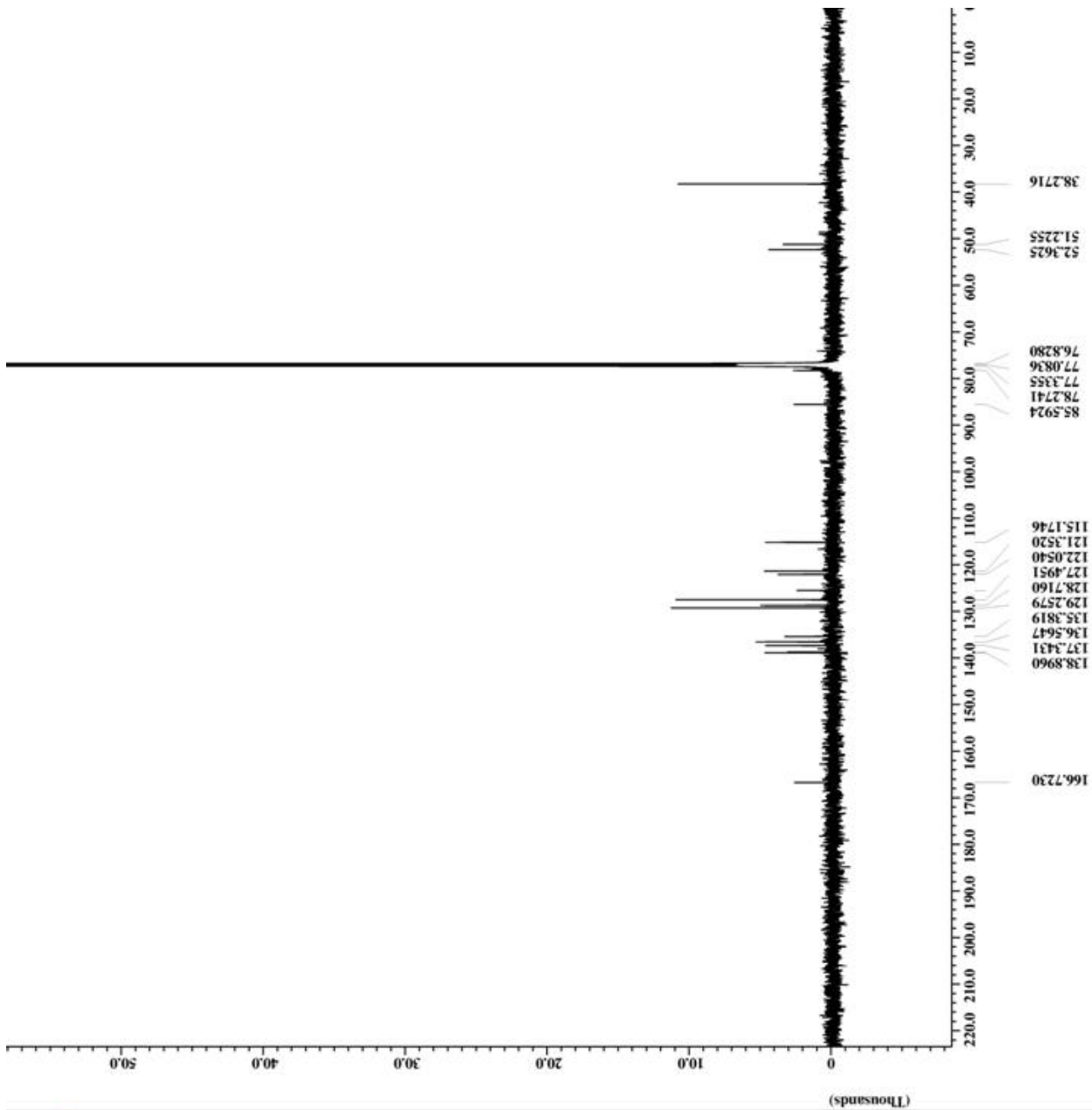
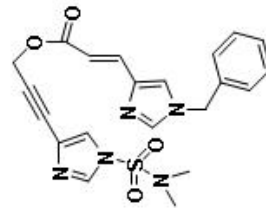


```
filename = sm_VI_80_pure-2.jdf
author = delta
experiment = single_pulse_dec
sample_id = S87522
solvent = CHLOROFORM-D
reaction_time = 28-DEC-2009 22:58:07
revision_time = 28-DEC-2009 14:09:09
current_time = 28-MAR-2010 21:29:38

comment = single pulse decouple
ata_format = ID COMPLEX
im_size = 65536
im_title = 13C
im_units = [ppm]
imensions = X
ite = Eclipse+ 500
spectrometer = DELTA_NMR

field_strength = 11.7465928[T] (500[MH
acq_duration = 2.0840448[s]
domain = 13C
freq = 125.75710665[MHz]
offset = 100[ppm]
points = 65536
prescans = 4
resolution = 0.47983613[Hz]
sweep = 31.44654088[kHz]
tr_domain = 1H
tr_freq = 500.12734003[MHz]
tr_offset = 5[ppm]
lipped = FALSE
bd_return = 10
cans = 5000
ctal_scans = 5000

_90_width = 14.2[us]
acq_time = 2.0840448[s]
angle = 30[deg]
pulse = 4.73333333[us]
nitia1_wait = 1[s]
pe_time = 1[s]
base_preset = 3[us]
scvr_gain = 22
relaxation_delay = 2[s]
emp_get = 32.1[dC]
nblank_time = 2[us]
```





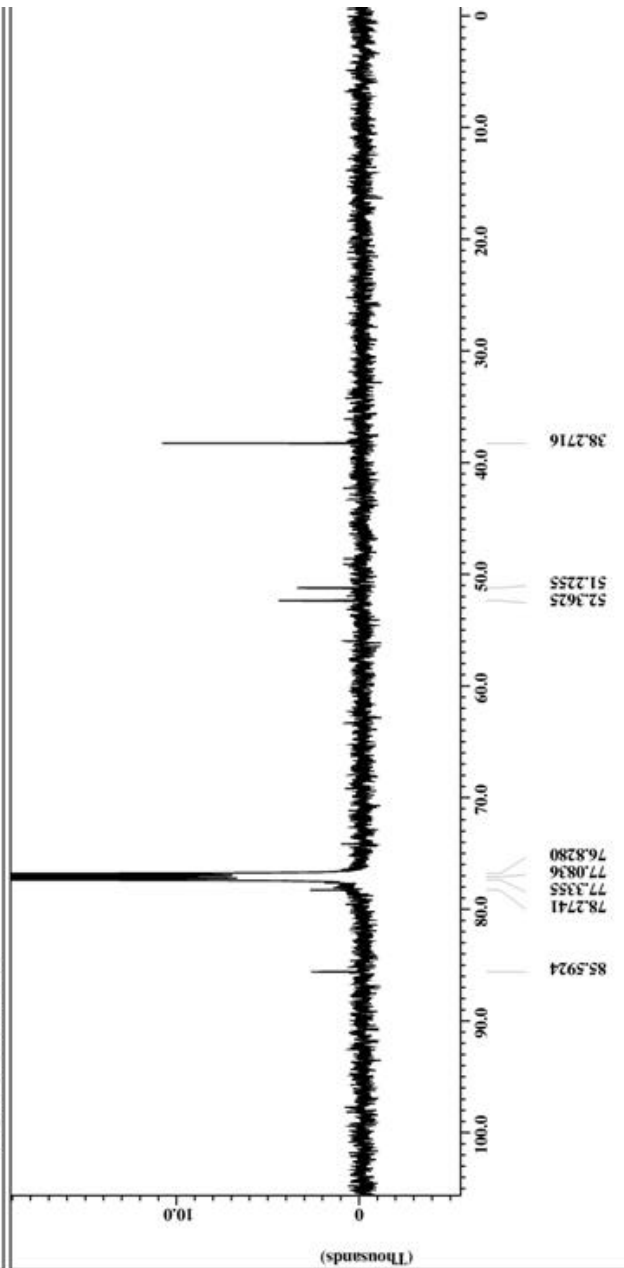
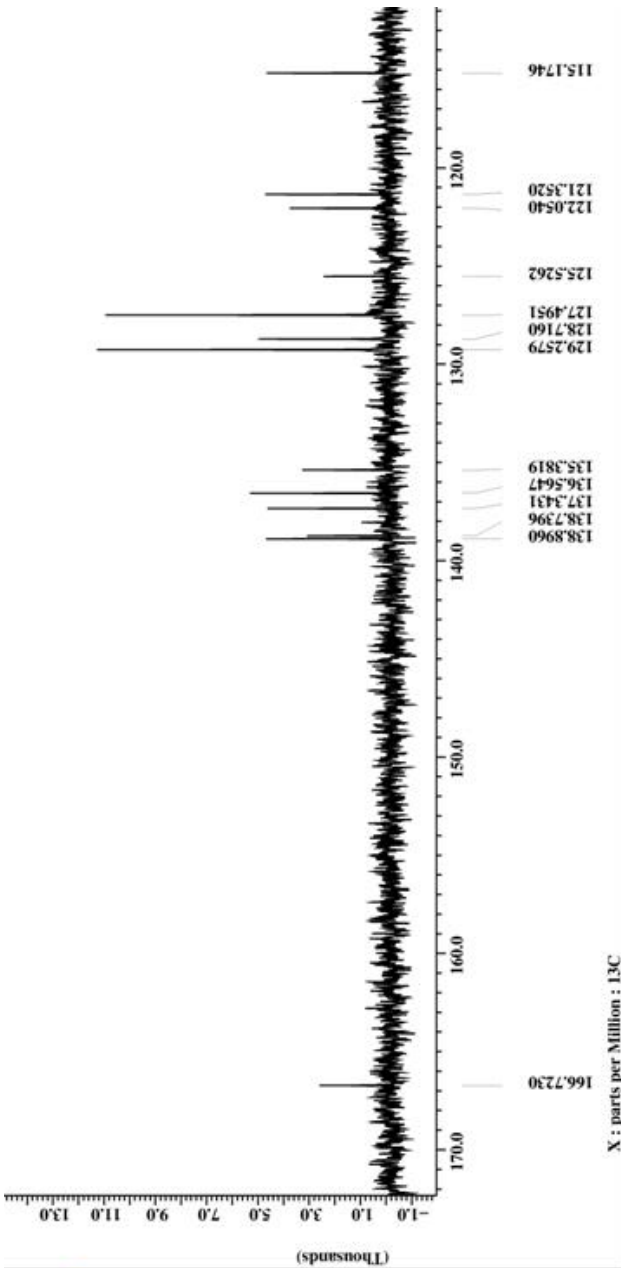
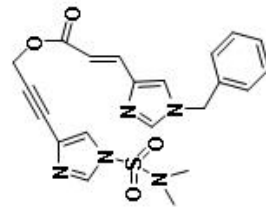
```

filename = sm_VI_80_pure-2.jdf
author = delta
experiment = single_pulse_dec
sample_id = S87522
solvent = CHLOROFORM-D
reaction_time = 28-DEC-2009 22:58:07
evision_time = 28-DEC-2009 14:09:09
current_time = 28-MAR-2010 21:31:55

comment = single pulse decouple
          = ID COMPLEX
          = 65536
          = 13C
          = [ppm]
          = X
          = Eclipse+ 500
          = DELTA_NMR

p1 = 11.7465928[T] (500 [MH
p2 = 2.0840448[s]
p3 = 125.75710665 [MHZ]
p4 = 100 [ppm]
p5 = 65536
p6 = 4
p7 = 0.47983613 [Hz]
p8 = 31.44654088 [kHz]
p9 = IR
p10 = 500.12734003 [MHZ]
p11 = 5 [ppm]
p12 = FALSE
p13 = 10
p14 = 5000
p15 = 5000

p16 = 14.2 [us]
p17 = 2.0840448 [s]
p18 = 30 [deg]
p19 = 4.73333333 [us]
p20 = 1 [s]
p21 = 1 [s]
p22 = 3 [us]
p23 = 22
p24 = 2 [s]
p25 = 32.1 [dC]
p26 = 2 [us]
  
```



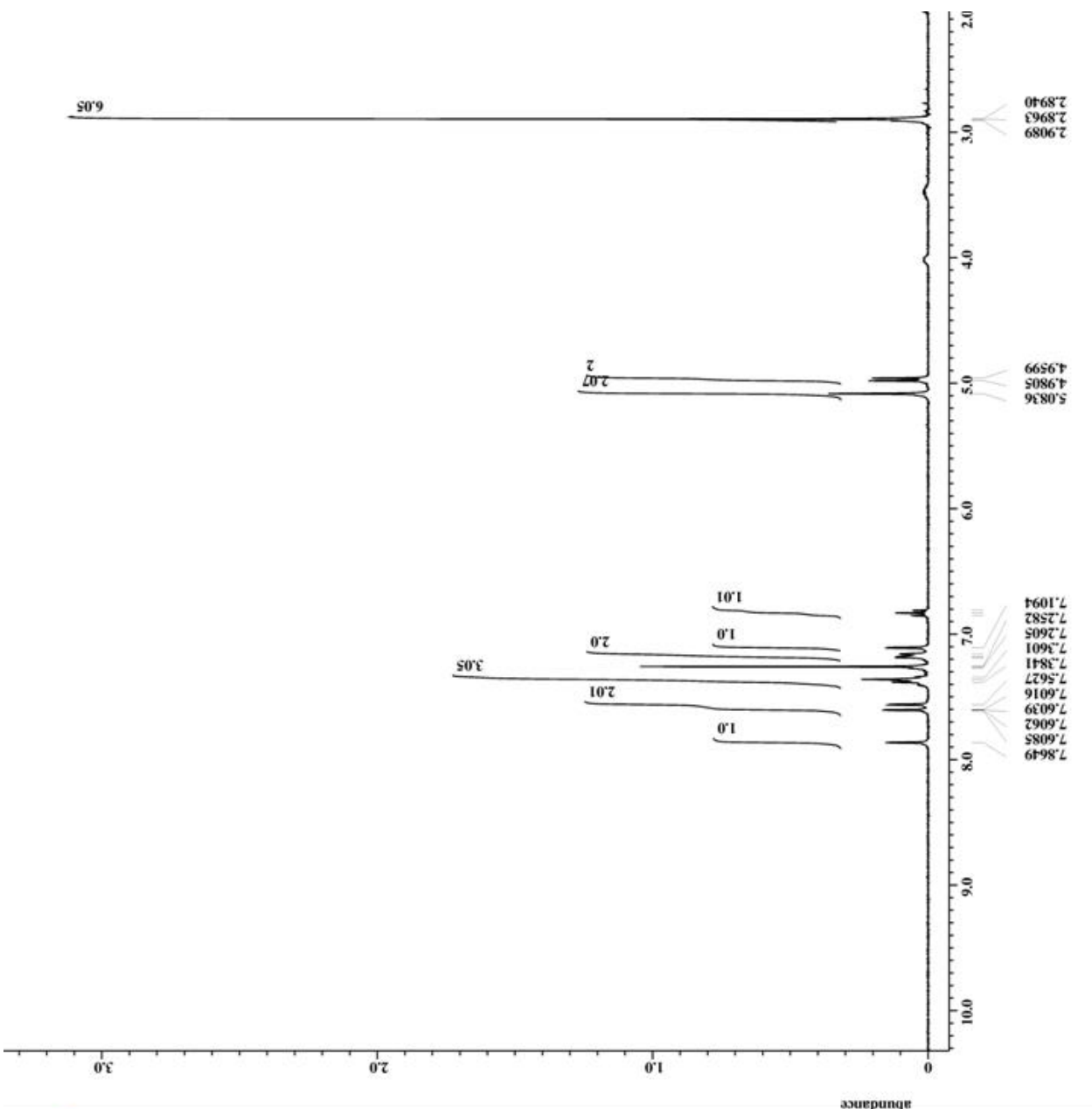
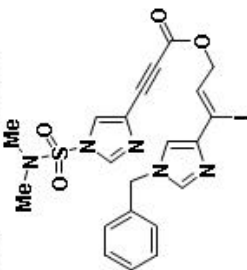
APPENDIX 44
¹H AND ¹³C NMR SPECTRUM OF
(Z)-3-(1-benzyl-1H-imidazol-4-yl)-3-iodoallyl-3-(1-(N,N-dimethylsulfamoyl)-
1H-imidazol-4-yl)propiolate (250)



```

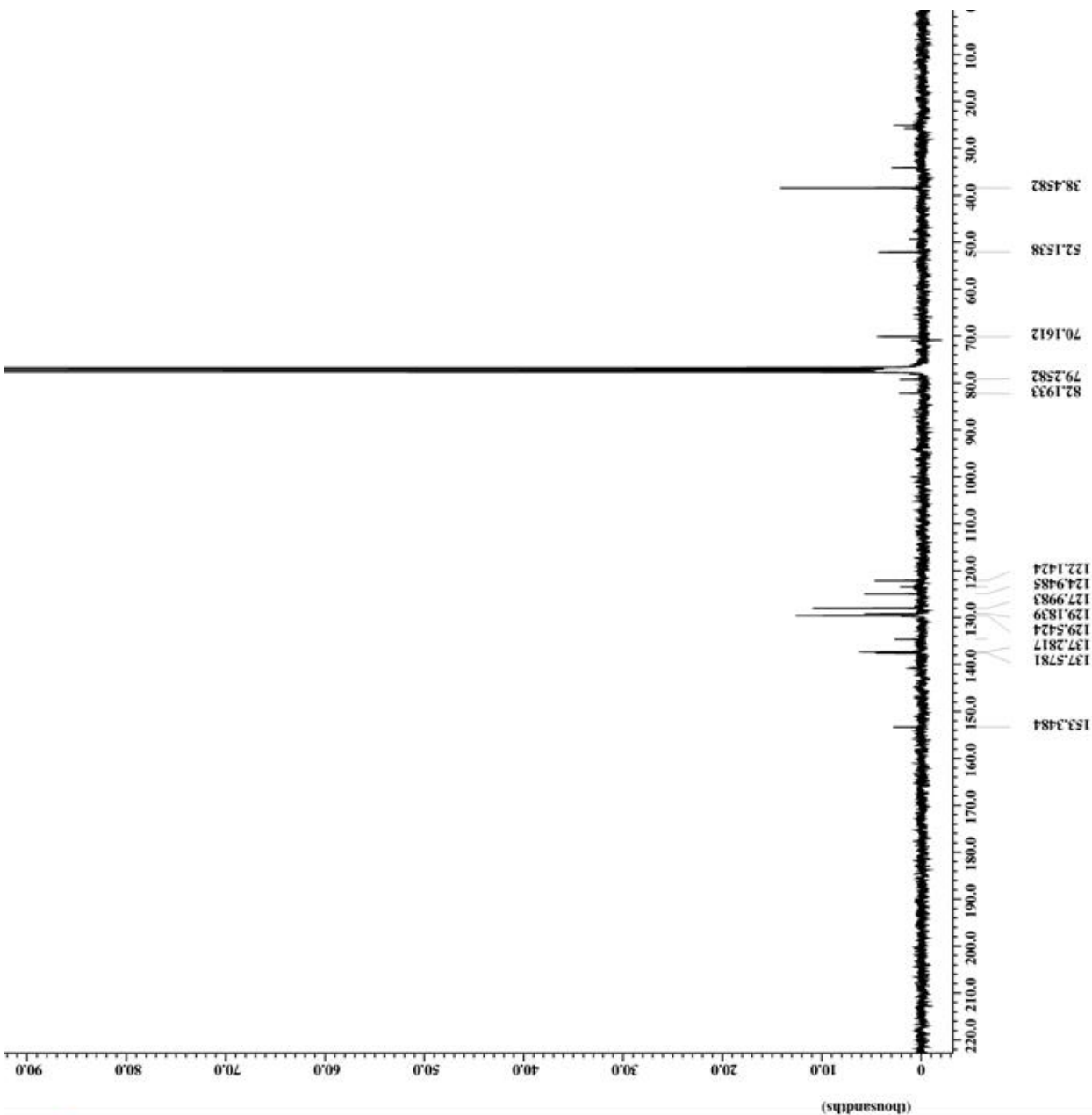
filename = sm_V_193_crude-4.jdf
author = delta
experiment = single_pulse.ex2
sample_id = S#732653
solvent = CHLOROFORM-D
reaction_time = 1-SEP-2009 20:20:29
acquisition_time = 28-MAR-2010 21:36:43
current_time = 28-MAR-2010 21:37:24
comment =
  = single_pulse
  = ID REAL
  = 13107
  = 1H
  = [ppm]
  = X
  = ECX 300
  = DELTA2_NMR
p1
p2
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```





```
filename = sm_VI_iodo_DCC_coupli
author = delta
experiment = single_pulse_dec
sample_id = S#793596
solvent = CHLOROFORM-D
reaction_time = 20-JAN-2010 01:50:29
acquisition_time = 19-APR-2010 19:25:03
current_time = 19-APR-2010 19:26:04
comment = single pulse decouple
ata_format = ID COMPLEX
in_size = 52428
in_title = I3C
in_units = [ppm]
inmensions = X
ite = ECX 300
spectrometer = DELTA2_NMR
ield_strength = 7.0586013[T] (300[MHz]
acq_duration = 2.76824064[s]
domain = I3C
freq = 75.56823426[MHz]
offset = 100[ppm]
points = 65536
prescans = 4
resolution = 0.36124027[Hz]
sweep = 23.67424242[kHz]
tr_domain = 1H
tr_freq = 300.52965592[MHz]
tr_offset = 5[ppm]
lipped = FALSE
bd_return = 10
cans = 2500
otal_scans = 2500
_90_width = 9.75[us]
acq_time = 2.76824064[s]
angle = 30[deg]
atn = 8[dB]
pulse = 3.25[us]
tr_atn_dec = 25[dB]
tr_atn_noe = 25[dB]
tr_noise = WALTZ
scoupling = TRUE
nitial_wait = 1[s]
oe_time = TRUE
oe_time = 3[s]
ecvr_gain = 50
elaxation_delay = 3[s]
epetition_time = 5.76824064[s]
emp_get = 21.3[dc]
```



APPENDIX 45
¹H AND ¹³C NMR SPECTRUM OF
3-(1-benzyl-1H-imidazol-4-yl)prop-2-ynyl-3-(1-(N,N-dimethylsulfamoyl)-1H-
imidazol-4-yl)propiolate (251)



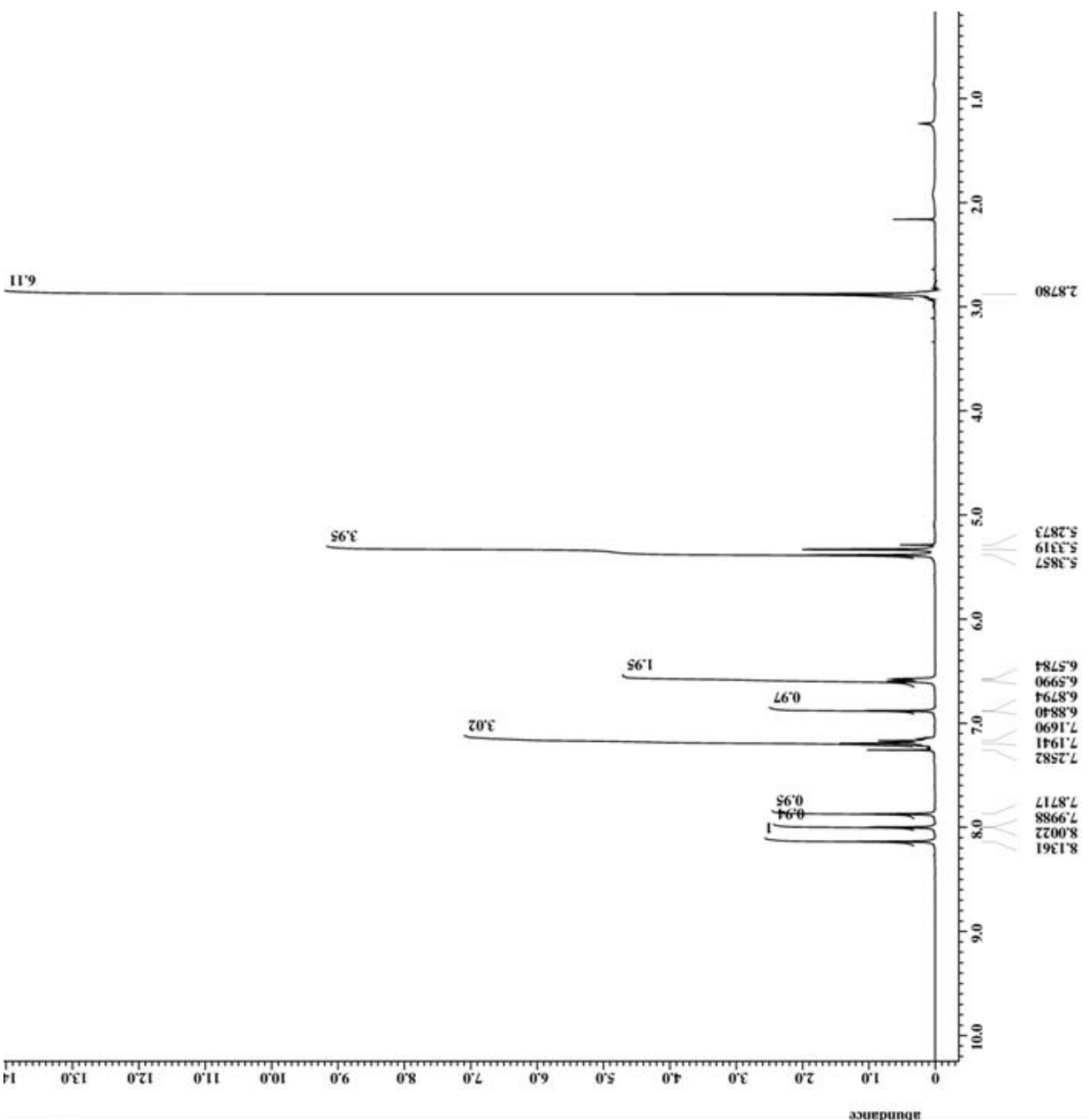
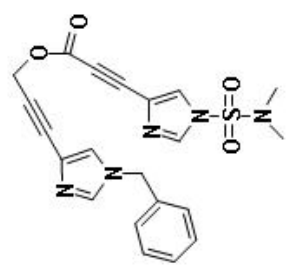
```

filename = sm_VI_ynp_pure-2.
author = delta
experiment = single_pulse.ex2
sample_id = S8658718
solvent = CHLOROFORM-D
reaction_time = 2-FEB-2010 18:09:37
acquisition_time = 2-FEB-2010 18:23:03
current_time = 19-APR-2010 19:31:36

comment = single_pulse
ata_format = ID COMPLEX
in_size = 13107
in_title = 1H
in_units = [ppm]
in_dimensions = X
ite = ECX 300
spectrometer = DELTA2_NMR

field_strength = 7.0586013[T] (300[MHz])
acq_duration = 2.90717696[s]
domain = 1H
freq = 300.52965592[MHz]
offset = 5[ppm]
points = 16384
prescans = 0
resolution = 0.34397621[Hz]
sweep = 5.63570784[kHz]
tr_domain = 1H
tr_freq = 300.52965592[MHz]
tr_offset = 5[ppm]
ri_domain = 1H
ri_freq = 300.52965592[MHz]
ri_offset = 5[ppm]
lipped = FALSE
od_return = 1
cans = 19
atal_scans = 19

_90_width = 13.01[us]
acq_time = 2.90717696[s]
angle = 45[deg]
atn = 4[dB]
pulse = 6.505[us]
tr_mode = Off
ri_mode = Off
ante_presat = FALSE
nit1_wait = 1[s]
scvr_gain = 42
relaxation_delay = 5[s]
petition_time = 7.90717696[s]
emp_get = 19.8[dc]
  
```

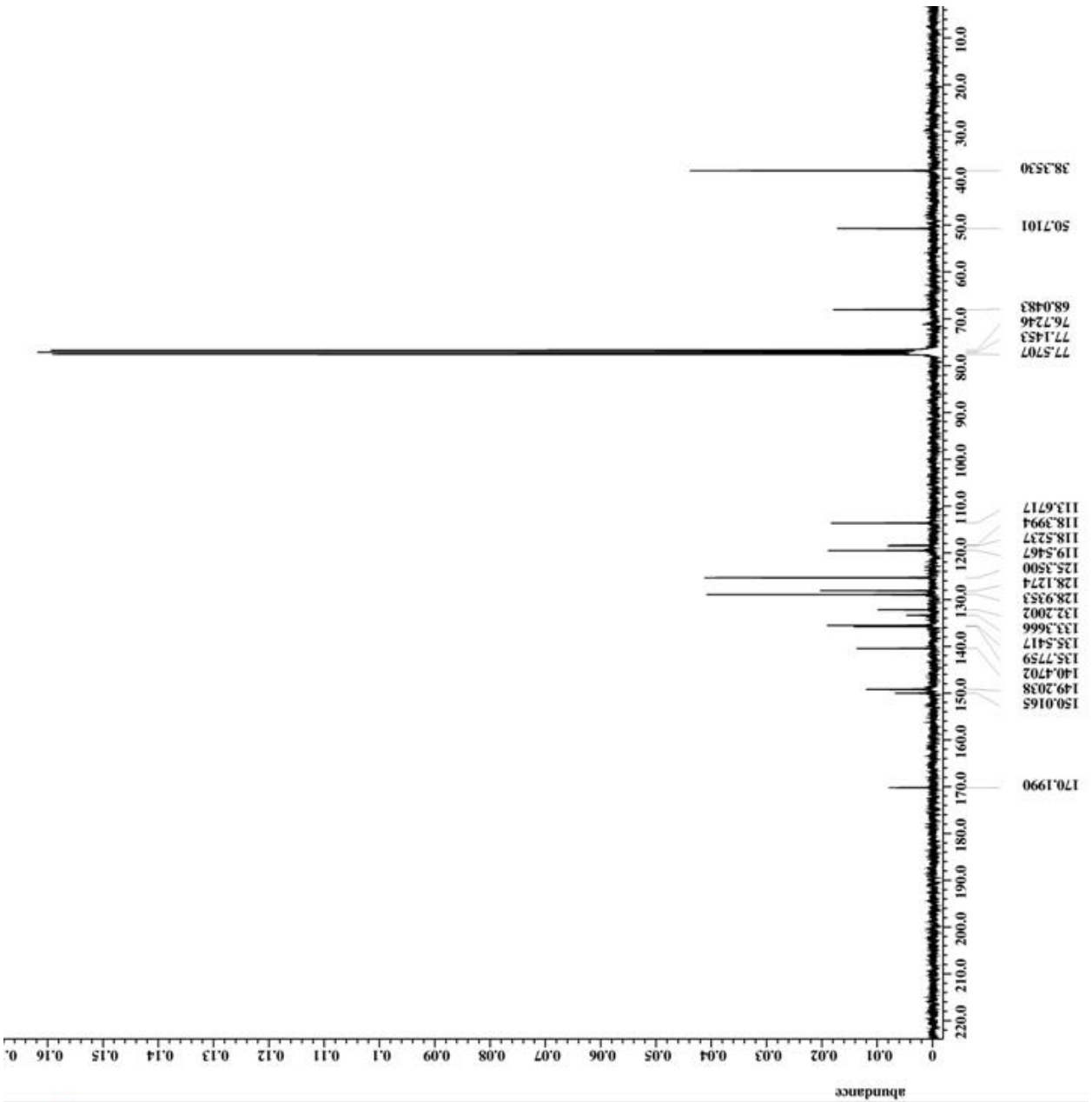
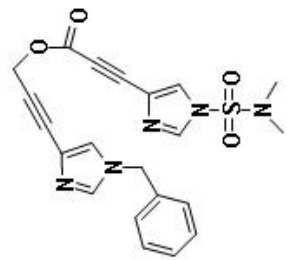




```

filename = sm_VI_ynp_pure-2.
author = delta
experiment = single_pulse_dec
pulse_id = S8660727
solvent = CHLOROFORM-D
reaction_time = 2-FEB-2010 20:35:31
evolution_time = 2-FEB-2010 23:41:18
current_time = 19-APR-2010 19:34:35
comment = single pulse decouple
ata_format = ID COMPLEX
in_size = 52428
in_title = 13C
in_units = [ppm]
in_dimensions = X
ite = ECX 300
ite_preamplifier = DELTA2_NMR
ield_strength = 7.0586013[T] (300[MHz])
_acq_duration = 2.76824064[s]
_domain = 13C
_freq = 75.56823426[MHz]
_offset = 100[ppm]
_points = 65536
_prescans = 4
_resolution = 0.36124027[Hz]
_sweep = 23.67424242[kHz]
tr_domain = 1H
tr_freq = 300.52965592[MHz]
tr_offset = 5[ppm]
lipped = FALSE
bd_return = 10
cans = 1500
otal_scans = 1500
_90_width = 9.75[us]
_acq_time = 2.76824064[s]
_angle = 30[deg]
_atn = 8[dB]
_pulse = 3.25[us]
tr_atn_dec = 25[dB]
tr_atn_noe = 25[dB]
tr_noise = WALTZ
scoupling = TRUE
nitial_wait = 1[s]
oe_time = TRUE
oe_time = 3[s]
scr_gain = 50
slaxation_delay = 3[s]
petition_time = 5.76824064[s]
emp_get = 19.8[GC]

```



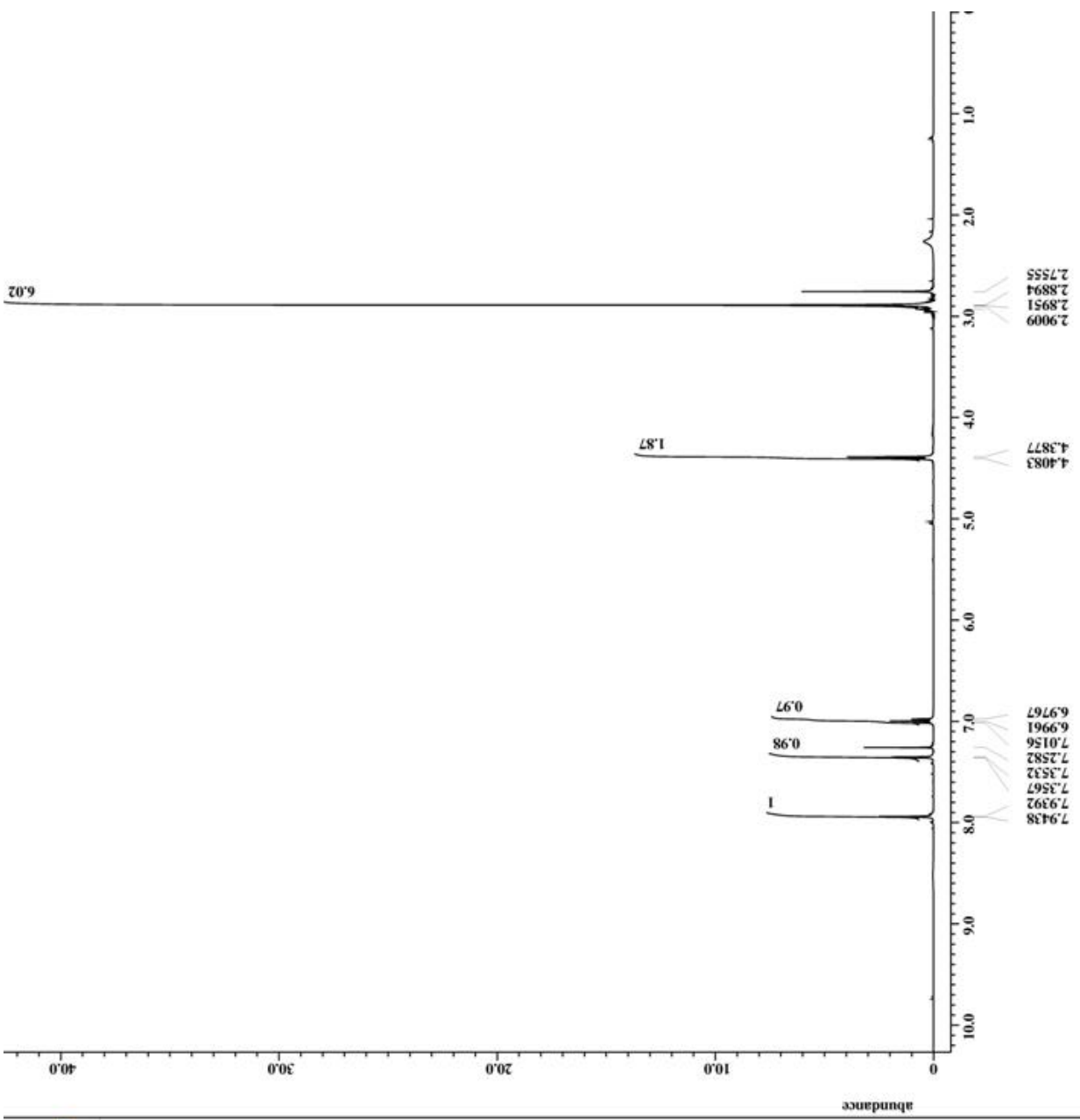
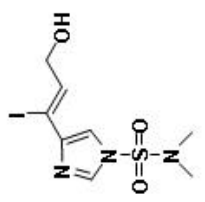
APPENDIX 46
¹H AND ¹³C NMR SPECTRUM OF
(Z)-4-(3-hydroxy-1-iodoprop-1-enyl)-N,N-dimethyl-1H-imidazole-1-
sulfonamide (244)



```

filename = sm_VI_88_pure-4_jdf
author = delta
experiment = single_pulse.ex2
sample_id = S#543823
solvent = CHLOROFORM-D
creation_time = 11-JAN-2010 14:55:29
revision_time = 28-MAR-2010 21:54:44
current_time = 28-MAR-2010 21:55:17
comment =
  = single_pulse
  = ID COMPLEX
  = 13107
  = 1H
  = [ppm]
  = X
  = ECX 300
  = DELTA2_NMR
field_strength = 7.0586013[T] (300[MHz])
  _acq_duration = 2.90717696[s]
  _domain = 1H
  _freq = 300.52965592[MHz]
  _offset = 5[ppm]
  _points = 16384
  _prescans = 0
  _resolution = 0.34397621[Hz]
  _sweep = 5.63570784[kHz]
  rr_domain = 1H
  rr_freq = 300.52965592[MHz]
  rr_offset = 5[ppm]
  ri_domain = 1H
  ri_freq = 300.52965592[MHz]
  ri_offset = 5[ppm]
  lipped = FALSE
  od_return = 1
  otal_scans = 24
  _90_width = 13.01[us]
  _acq_time = 2.90717696[s]
  _angle = 45[deg]
  _atn = 4[db]
  _pulse = 6.505[us]
  rr_mode = Off
  ri_mode = Off
  ante_presat = FALSE
  nitia_wait = 1[s]
  scvr_gain = 50
  relaxation_delay = 5[s]
  repetition_time = 7.90717696[s]
  temp_get = 21[degC]

```





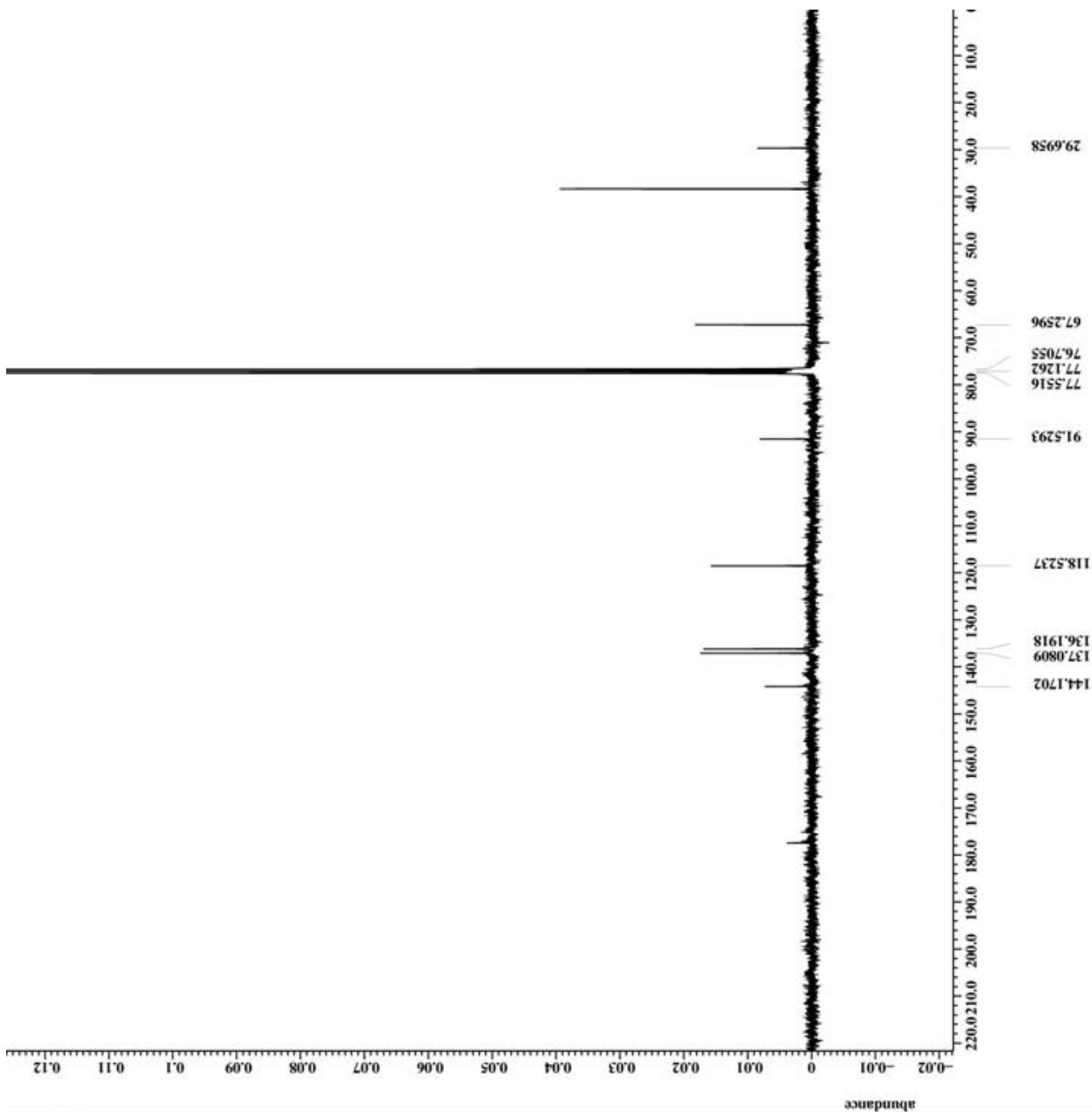
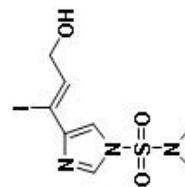
```

ilename      = sm_VI_88_PURE-3.jdf
author       = delta
experiment   = single_pulse_dec
sample_id    = S#549621
solvent      = CHLOROFORM-D
reaction_time = 11-JAN-2010 16:38:22
revision_time = 28-MAR-2010 21:58:07
current_time = 28-MAR-2010 21:58:25

comment      = single pulse decouple
ate_format   = 1D COMPLEX
im_size      = 52428
im_title     = 13C
im_units     = [ppm]
imensions    = X
ite          = ECX 300
pctrometer   = DELTA2_NMR

ield_strength = 7.0586013[T] (300[MHz]
-acq_duration = 2.76824064[s]
-domain       = 13C
-freq         = 75.56823426[MHz]
-offset       = 100[ppm]
-points       = 65336
-prescans     = 4
-resolution   = 0.36124027[Hz]
-sweep        = 23.67424242[KHz]
rr_domain    = 1H
rr_freq       = 300.52965592[MHz]
rr_offset    = 5[ppm]
lipped       = FALSE
bd_return    = 10
cans         = 1000
stal_scans   = 1000

_90_width    = 9.75[us]
-acq_time     = 2.76824064[s]
-angle        = 30[deg]
-atn          = 8[db]
-pulse        = 3.25[us]
rr_atn_dec   = 25[db]
rr_atn_noe   = 25[db]
rr_noise     = WALTZ
scoupling    = TRUE
nit1al_wait  = 1[s]
ce           = TRUE
be_time      = 3[s]
scvr_gain    = 50
relaxation_delay = 3[s]
apitation_time = 5.76824064[s]
emp_get      = 21.2[dc]
  
```

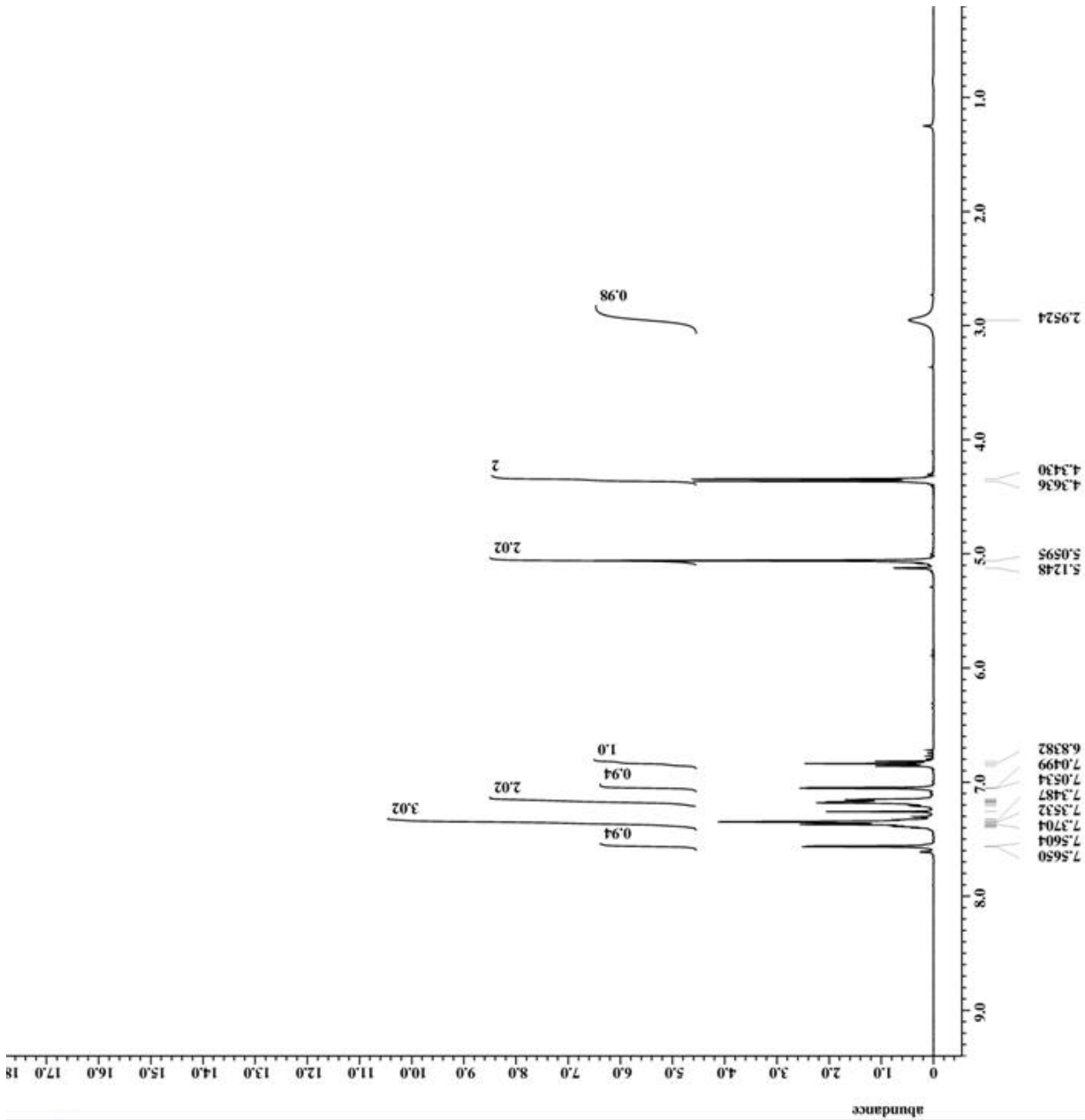
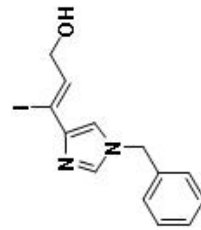


APPENDIX 47
¹H AND ¹³C NMR SPECTRUM OF
(Z)-3-(1-benzyl-1H-imidazol-4-yl)-3-iodoprop-2-en-1-ol (245)



```

filename = sm_V_180_pure-7_jdf
author = delta
experiment = single_pulse.ex2
sample_id = S811089
solvent = CHLOROFORM-D
acquisition_time = 3-AUG-2009 22:27:59
revision_time = 28-MAR-2010 22:03:00
current_time = 28-MAR-2010 22:03:34
comment = single_pulse
ata_format = ID REAL
in_size = 13107
in_title = 1H
in_units = [ppm]
in_dimensions = X
ite = ECX 300
spectrometer = DELTA2_NMR
field_strength = 7.0586013[T] (300[MHz])
acq_duration = 2.90717696[s]
domain = 1H
freq = 300.52965592[MHz]
offset = 5[ppm]
points = 16384
prescans = 0
resolution = 0.34397621[Hz]
sweep = 5.63570784[kHz]
tr_domain = 1H
tr_freq = 300.52965592[MHz]
tr_offset = 5[ppm]
ri_domain = 1H
ri_freq = 300.52965592[MHz]
ri_offset = 5[ppm]
lipped = FALSE
od_return = 1
cans = 24
atal_scans = 24
_90_width = 13.01[us]
acq_time = 2.90717696[s]
angle = 45[deg]
atn = 4[dB]
pulse = 6.505[us]
ri_mode = Off
ri_mode = Off
ante_presat = FALSE
initial_wait = 1[s]
scvr_gain = 50
relaxation_delay = 5[s]
petition_time = 7.90717696[s]
emp_get = 23.2[dc]
  
```



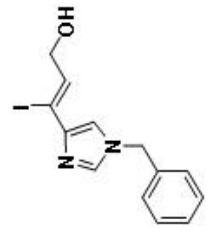
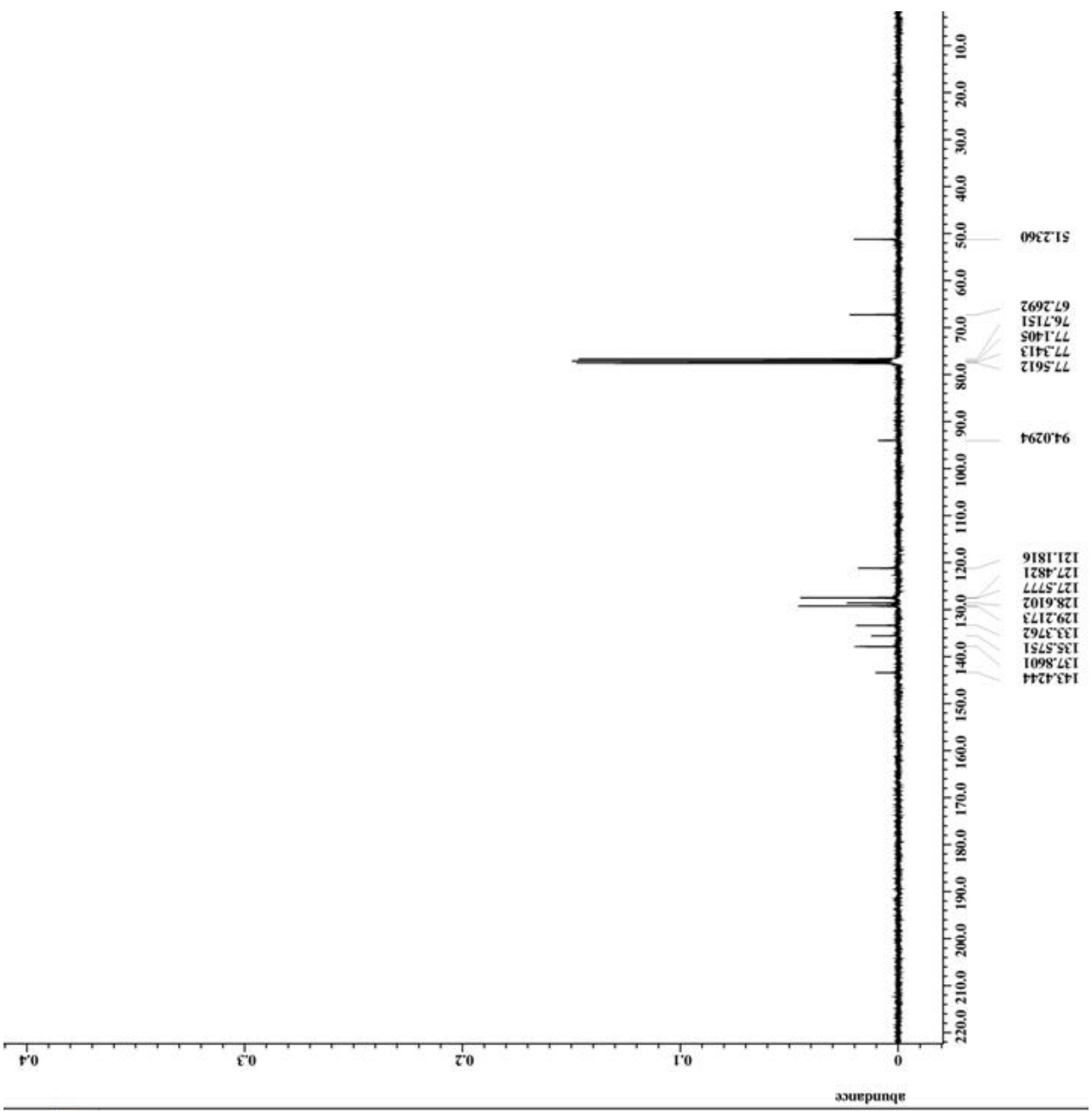


```
filename = sm_V_180_pure-2.jdf
author = delta
experiment = single_pulse_dec
sample_id = S813546
solvent = CHLOROFORM-D
acquisition_time = 4-AUG-2009 03:17:32
revision_time = 4-AUG-2009 12:18:54
current_time = 28-MAR-2010 22:05:58

comment = single pulse decouple
ata_format = ID COMPLEX
in_size = 52428
in_title = 13C
in_units = [ppm]
in_dimensions = X
ite = ECX 300
spectrometer = DELTA2_NMR

field_strength = 7.0586013[T] (300[MHz])
acq_duration = 2.76824064[s]
domain = 13C
freq = 75.56823426[MHz]
offset = 100[ppm]
points = 65536
prescans = 4
resolution = 0.36124027[Hz]
sweep = 23.67424242[kHz]
tr_domain = 1H
tr_freq = 300.52965592[MHz]
tr_offset = 5[ppm]
lipped = FALSE
bd_return = 10
cans = 3000
total_scans = 3000

_90_width = 9.75[us]
acq_time = 2.76824064[s]
angle = 30[deg]
atn = 8[dB]
pulse = 3.25[us]
tr_atn_dec = 25[dB]
tr_atn_noe = 25[dB]
tr_noise = WALTZ
scoupling = TRUE
initial_wait = 1[s]
be_time = TRUE
be = 3[s]
ecvr_gain = 50
relaxation_delay = 3[s]
petition_time = 5.76824064[s]
emp_get = 23.2[dc]
```



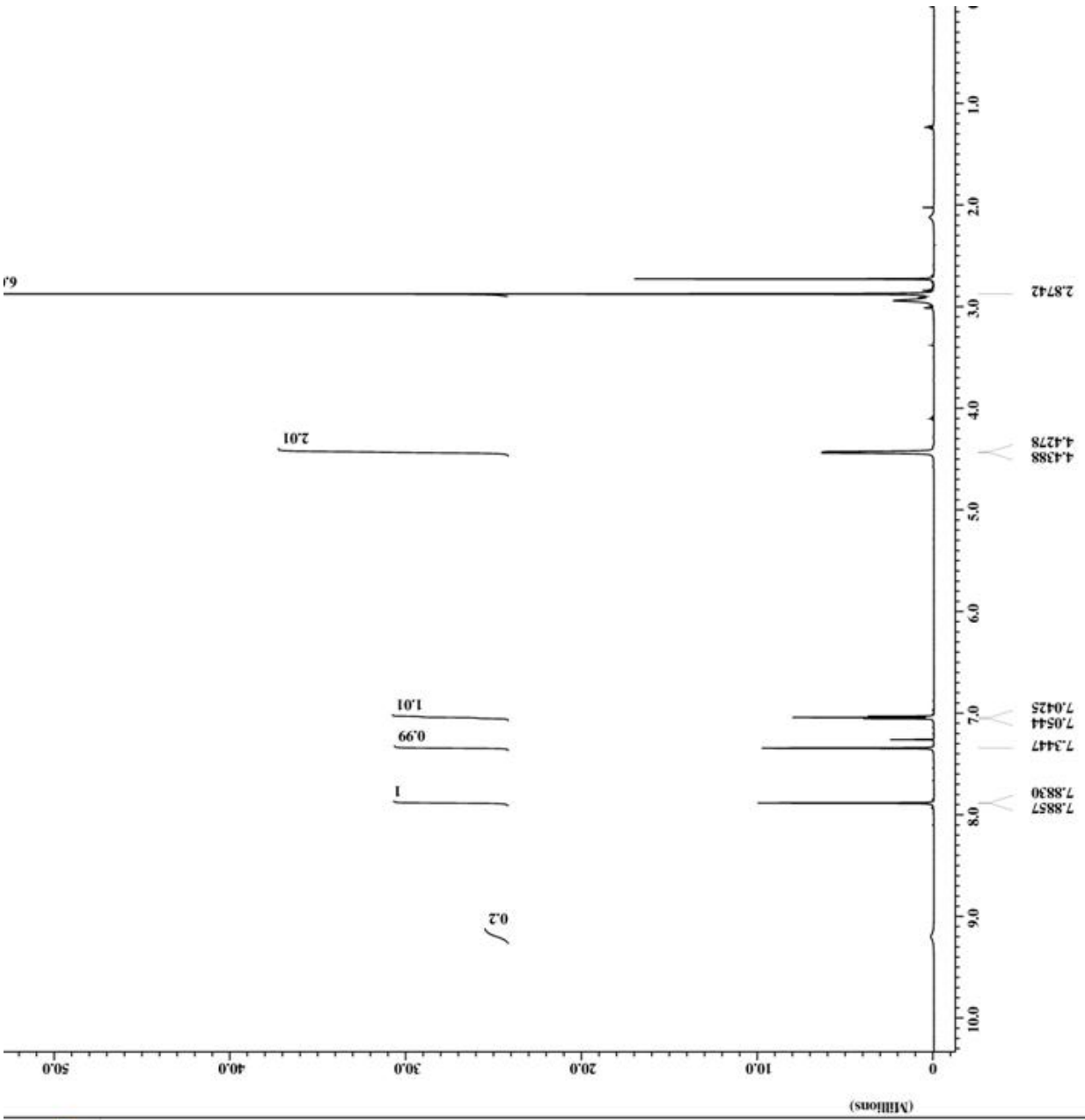
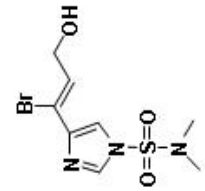
APPENDIX 48
¹H AND ¹³C NMR SPECTRUM OF
(Z)-4-(1-bromo-3-hydroxyprop-1-enyl)-N,N-dimethyl-1H-imidazole-1-
sulfonamide (259)



```

filename = sm_VI_74_pure-3.jdf
author = delta
experiment = single_pulse_exp
sample_id = S#862202
solvent = CHLOROFORM-D
reaction_time = 22-DEC-2009 15:36:16
revision_time = 28-MAR-2010 22:10:17
current_time = 28-MAR-2010 22:10:54
comment = Single Pulse Experiment
data_format = ID COMPLEX
in_size = 16384
in_title = 1H
in_units = [ppm]
in_dimensions = X
ite = Eclipse+ 500
spectrometer = DELTA_NMR
field_strength = 11.7465928[T] (500[MH
acq_duration = 2.1839872[s]
domain = 1H
freq = 500.12734003[MHz]
offset = 5[ppm]
points = 16384
prescans = 0
resolution = 0.45787814[Hz]
sweep = 7.50187547[kHz]
tipped = FALSE
od_return = 1
total_scans = 8
_90_width = 18.5[us]
acq_time = 2.1839872[s]
angle = 45[deg]
pulse = 7.25[us]
p1 = 1[s]
base_preset = 3[us]
ecvr_gain = 17
relaxation_delay = 4[s]
emp_get = 25.5[dc]
nblank_time = 2[us]

```





```

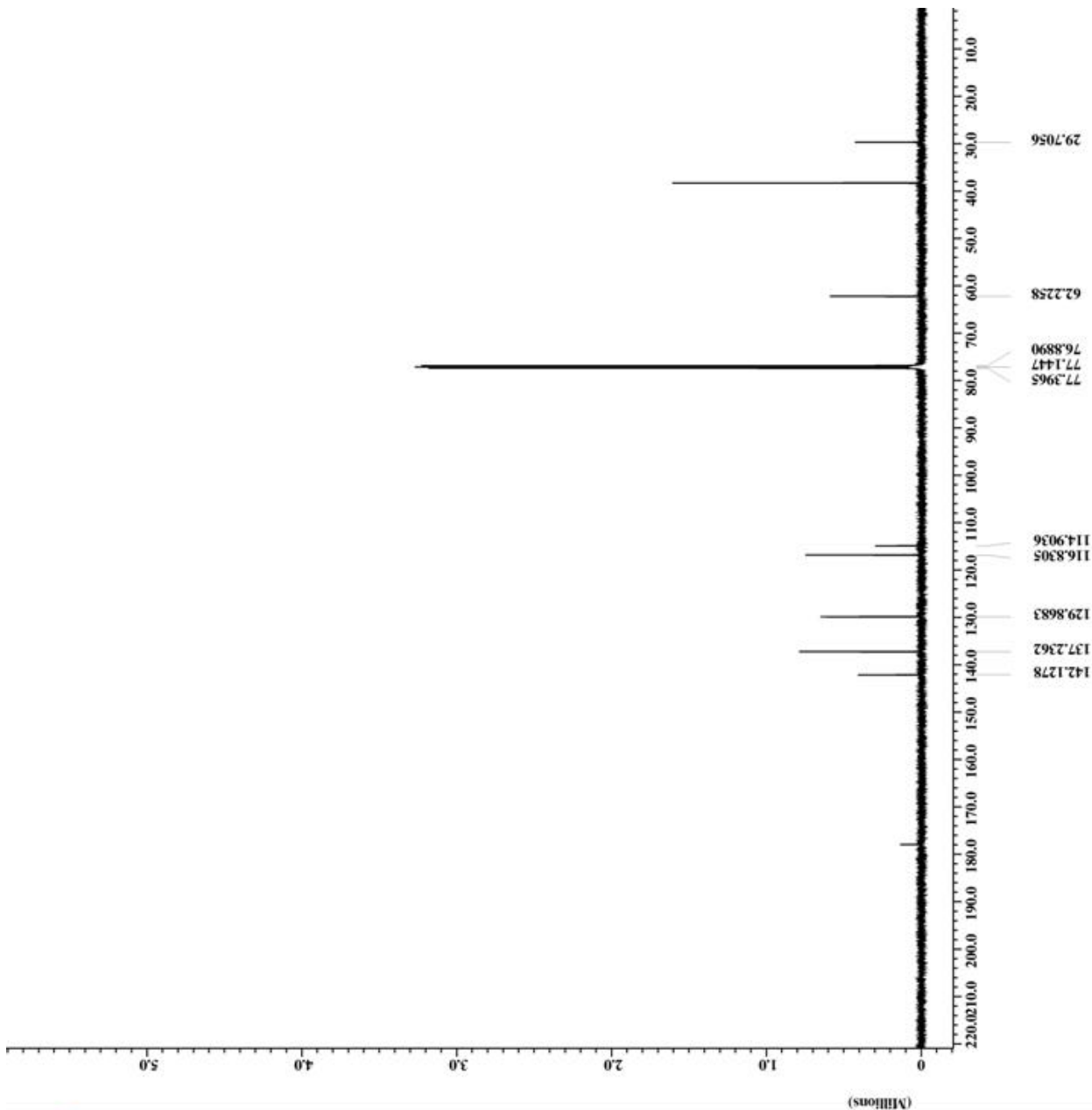
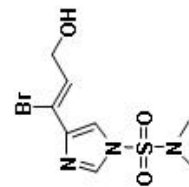
filename = sm_VI_74_pure-3.jdf
author =
experiment = single_pulse_dec
sample_id = S863468
solvent = CHLOROFORM-D
reaction_time = 22-DEC-2009 22:42:01
evision_time = 28-MAR-2010 22:13:15
current_time = 28-MAR-2010 22:13:28

comment = single pulse decouple
ata_format = ID COMPLEX
im_size = 65536
im_title = 13C
im_units = [ppm]
imensions = X
ite = Eclipse+ 500
spectrometer = DELTA_NMR

field_strength = 11.7465928[T] (500[MH
-acq_duration = 2.0840448[s]
-domain = 13C
-freq = 125.75710665[MHz]
-offset = 100[ppm]
-points = 65536
-prescans = 4
-resolution = 0.47983613[Hz]
-sweep = 31.44654088[kHz]
tr_domain = 1H
tr_freq = 500.12734003[MHz]
tr_offset = 5[ppm]
lipped = TRUE
bd_return = 10
cans = 5000
otal_scans = 5000

_90_width = 14.2[us]
-acq_time = 2.0840448[s]
-angle = 30[deg]
-pulse = 4.73333333[us]
nitai_wait = 1[s]
pe_time = 1[s]
base_preset = 3[us]
scvr_gain = 30
relaxation_delay = 2[s]
emp_get = 28.5[dc]
nblank_time = 2[us]

```



APPENDIX 49
¹H AND ¹³C NMR SPECTRUM OF
(Z)-3-(1-benzyl-1H-imidazol-4-yl)-3-bromoprop-2-en-1-ol (260)



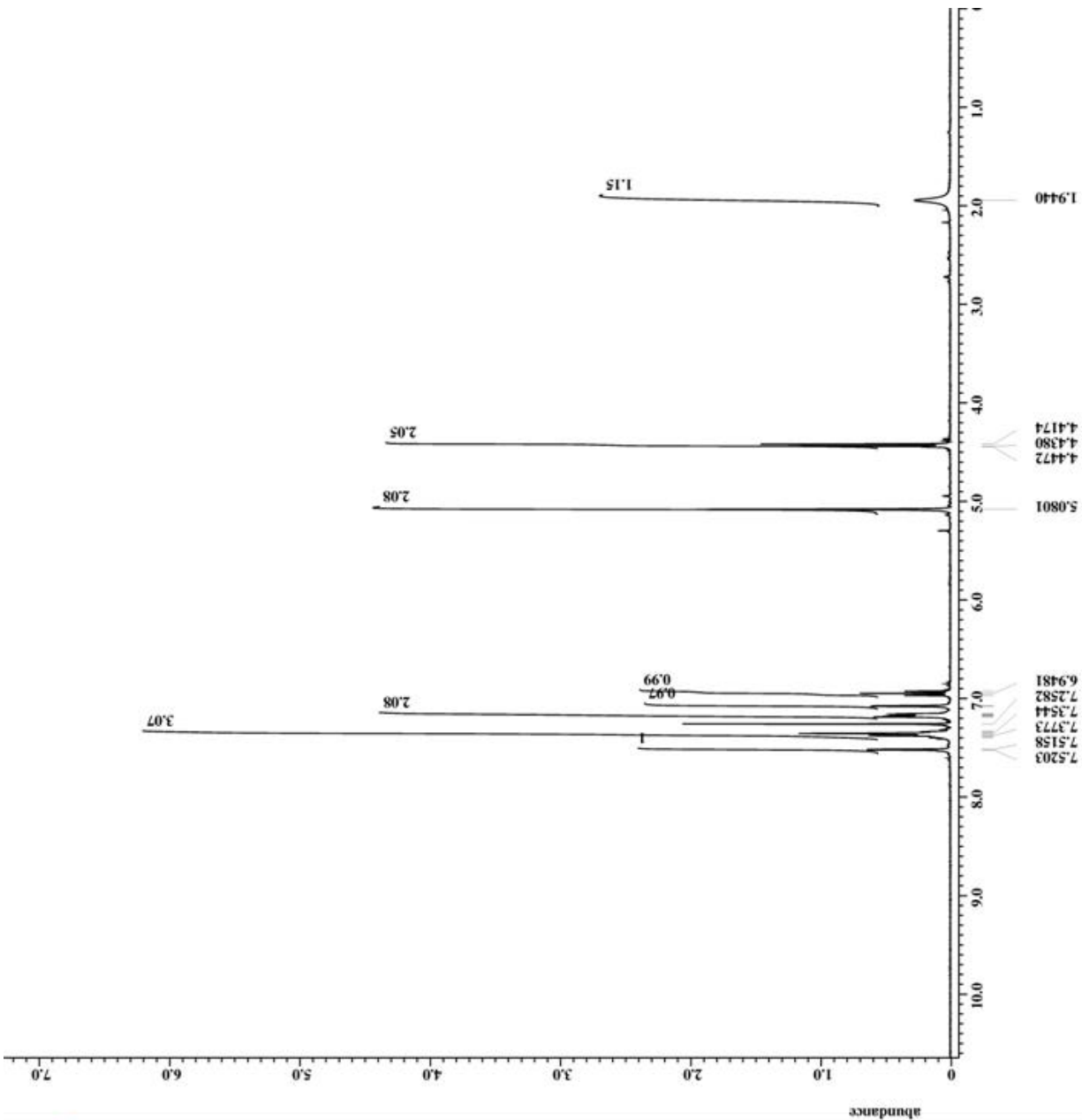
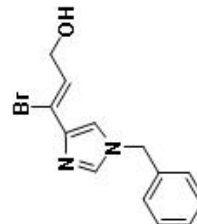
```

filename = sm_VI_67_pure-3.jdf
author = delta
experiment = single_pulse.ex2
sample_id = S#610342
solvent = CHLOROFORM-D
reaction_time = 15-DEC-2009 17:08:45
acquisition_time = 28-MAR-2010 22:16:54
current_time = 28-MAR-2010 22:17:17

comment = single_pulse
ata_format = ID COMPLEX
in_size = 13107
in_title = 1H
in_units = [ppm]
inmensions = X
ite = ECX 300
spectrometer = DELTA2_NMR

field_strength = 7.0586013[T] (300[MHz])
acq_duration = 2.90717696[s]
domain = 1H
freq = 300.52965592[MHz]
offset = 5[ppm]
points = 16384
prescans = 0
resolution = 0.34397621[Hz]
sweep = 5.63570784[kHz]
tr_domain = 1H
tr_freq = 300.52965592[MHz]
tr_offset = 5[ppm]
ri_domain = 1H
ri_freq = 300.52965592[MHz]
ri_offset = 5[ppm]
lipped = FALSE
od_return = 1
cans = 24
atal_scans = 24

_90_width = 13.01[us]
acq_time = 2.90717696[s]
angle = 45[deg]
atn = 4[dB]
pulse = 6.505[us]
tr_mode = Off
ri_mode = Off
ante_presat = FALSE
nitial_wait = 1[s]
scvr_gain = 48
relaxation_delay = 5[s]
petition_time = 7.90717696[s]
emp_get = 23.1[dc]
  
```





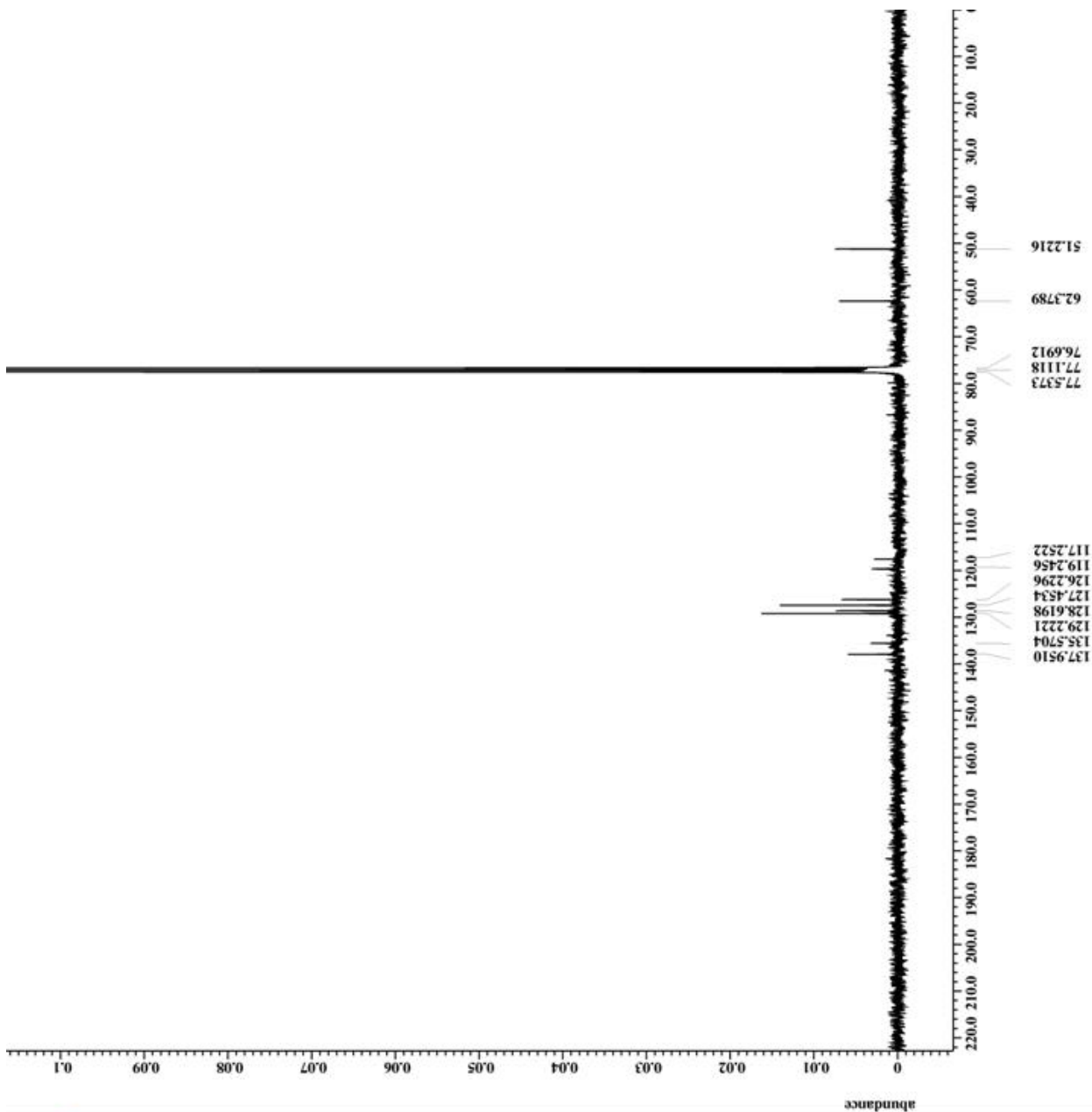
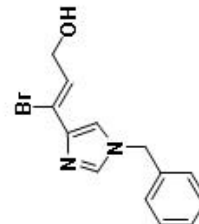
```

ilname      = sm_VI_67_pure-2.jdf
uthor       = delta
xperiment   = single_pulse_dec
ampl_id     = S8612850
solvent     = CHLOROFORM-D
reacion_time = 15-DEC-2009 19:06:01
evision_time = 15-DEC-2009 21:41:28
urrent_time = 28-MAR-2010 22:19:39

omment      = single pulse decouple
ate_format  = ID COMPLEX
im_size     = 52428
im_title    = 13C
im_units    = [ppm]
imensions   = X
ite         = ECX 300
ictrometer  = DELTA2_NMR

ield_strength = 7.0586013[T] (300[MHz]
.acq_duration = 2.76824064[s]
-domain       = 13C
-freq         = 75.56823426[MHz]
-offset       = 100[ppm]
-points       = 65336
-prescans     = 4
-resolution   = 0.36124027[Hz]
-sweep        = 23.67424242[KHz]
rr_domain    = 1H
rr_freq       = 300.52965592[MHz]
rr_offset     = 5[ppm]
lipped       = FALSE
bd_return    = 10
cans         = 1200
stal_scans   = 1200

_90_width    = 9.75[us]
.acq_time     = 2.76824064[s]
-angle       = 30[deg]
.atn         = 8[db]
_pulse       = 3.25[us]
rr_atn_dec   = 25[db]
rr_atn_noe   = 25[db]
rr_noise     = WALTZ
scoupling    = TRUE
nitital_wait = 1[s]
ce           = TRUE
ce_time      = 3[s]
ecvr_gain    = 50
elaxation_delay = 3[s]
epitation_time = 5.76824064[s]
emp_get      = 23.4[dC]
  
```



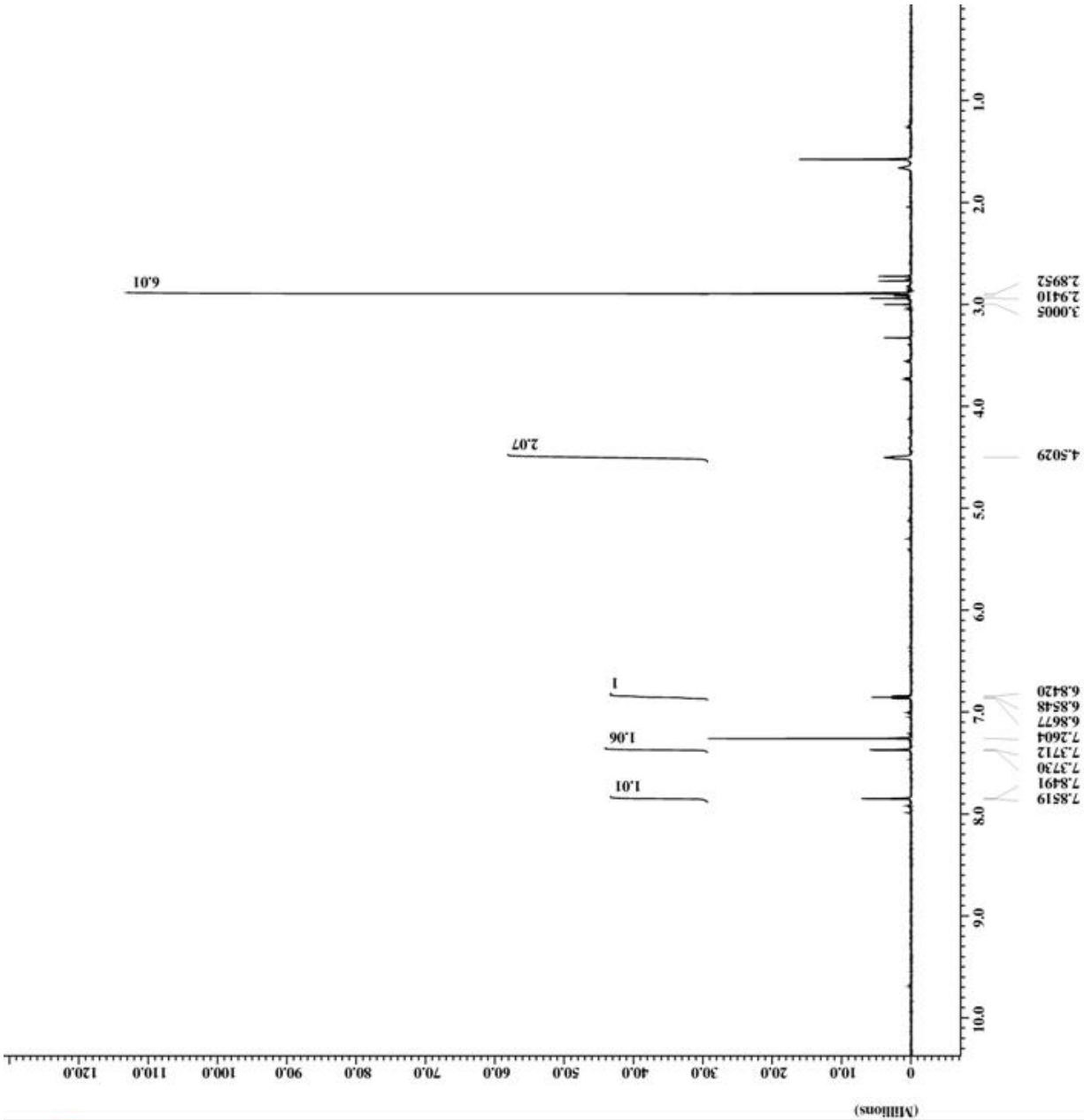
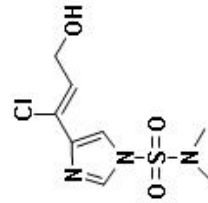
APPENDIX 50
¹H AND ¹³C NMR SPECTRUM OF
(Z)-4-(1-chloro-3-hydroxyprop-1-enyl)-N,N-dimethyl-1H-imidazole-1-
sulfonamide (263)



```

filename = sm_VI_RED_Al_Chloro_c
author = delta
experiment = single_pulse.exp
sample_id = S#545276
solvent = CHLOROFORM-D
reaction_time = 15-DEC-2009 06:43:31
acquisition_time = 28-MAR-2010 22:22:59
current_time = 28-MAR-2010 22:23:30
comment = Single Pulse Experiment
data_format = ID COMPLEX
in_size = 16384
in_title = 1H
in_units = [ppm]
in_dimensions = X
ite = Eclipse+ 500
spectrometer = DELTA_NMR
field_strength = 11.7465928[T] (500[MH
acq_duration = 2.1839872[s]
domain = 1H
freq = 500.12734003[MHz]
offset = 5[ppm]
points = 16384
prescans = 0
resolution = 0.45787814[Hz]
sweep = 7.50187547[kHz]
tipped = FALSE
od_return = 1
total_scans = 8
_90_width = 18.5[us]
_acq_time = 2.1839872[s]
_angle = 45[deg]
_pulse = 7.25[us]
_nitral_wait = 1[s]
_base_preset = 2[us]
_scvr_gain = 25
relaxation_delay = 4[s]
emp_get = 25.7[dc]
nblank_time = 2[us]

```





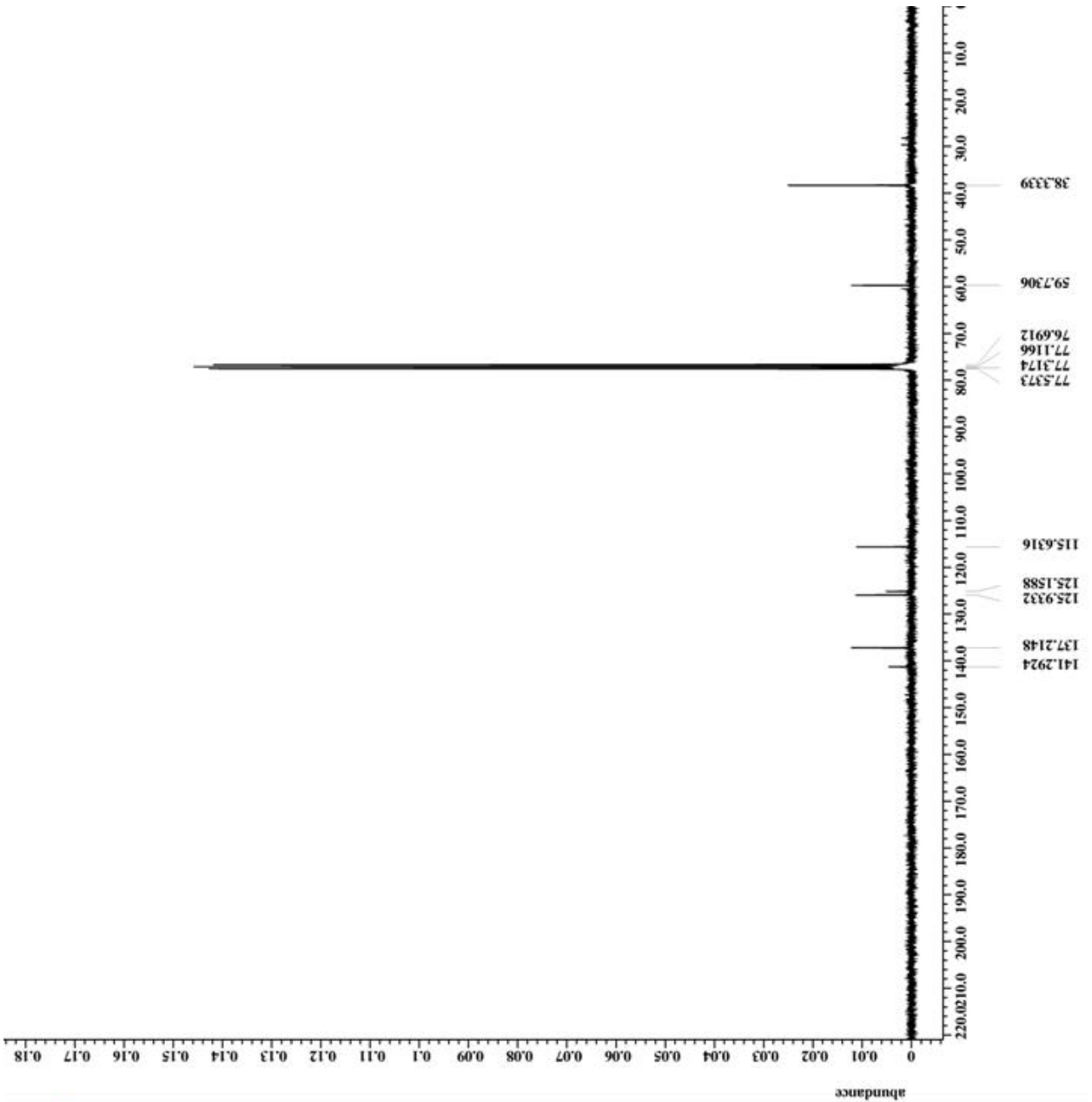
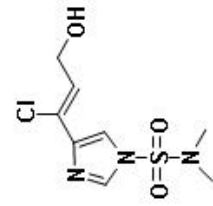
```

filename = sm_VI_64_pure-2.jdf
author = delta
experiment = single_pulse_dec
sample_id = S808732
solvent = CHLOROFORM-D
reaction_time = 15-DEC-2009 06:36:33
revision_time = 28-MAR-2010 22:26:29
current_time = 28-MAR-2010 22:26:51

comment = single pulse decouple
date_format = ID_COMPLEX
im_size = 52428
im_title = 13C
im_units = [ppm]
imensions = X
ite = ECX 300
spectrometer = DELTA2_NMR

field_strength = 7.0586013[T] (300[MHz]
-acq_duration = 2.76824064[s]
-domain = 13C
-freq = 75.56823426[MHz]
-offset = 100[ppm]
-points = 65536
-prescans = 4
-resolution = 0.36124027[Hz]
-sweep = 23.67424242[KHz]
rr_domain = 1H
rr_freq = 300.52965592[MHz]
rr_offset = 5[ppm]
lipped = FALSE
bd_return = 10
cans = 5000
stal_scans = 5000

_90_width = 9.75[us]
-acq_time = 2.76824064[s]
-angle = 30[deg]
-atn = 8[db]
-pulse = 3.25[us]
rr_atn_dec = 25[db]
rr_atn_noe = 25[db]
rr_noise = WALTZ
scoupling = TRUE
nit1al_wait = 1[s]
ce = TRUE
se_time = 3[s]
svr_gain = 50
relaxation_delay = 3[s]
acquisition_time = 5.76824064[s]
emp_get = 23.1[dc]
  
```



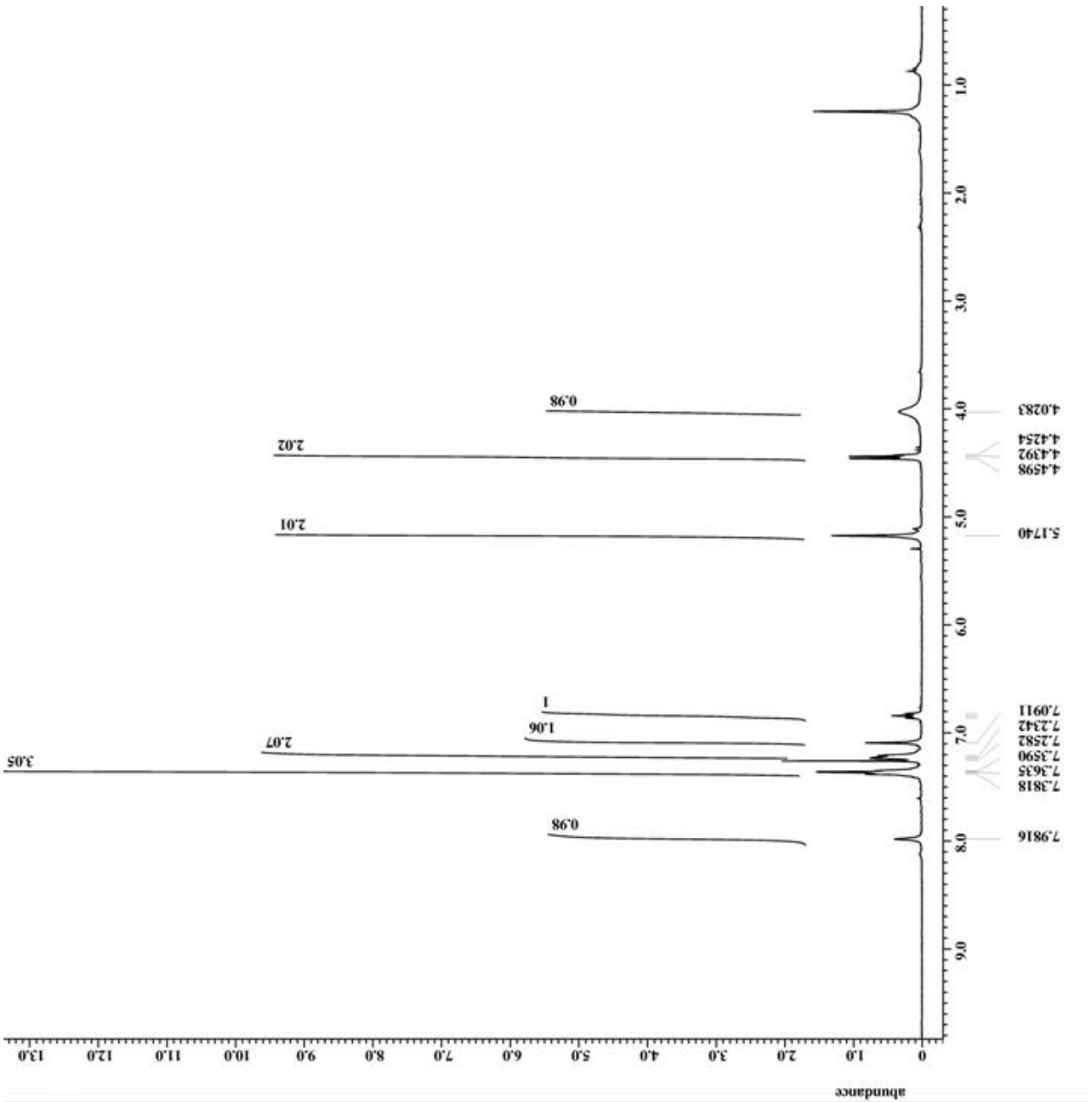
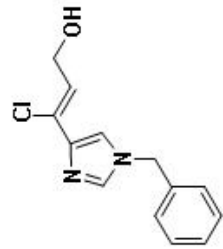
APPENDIX 51
¹H AND ¹³C NMR SPECTRUM OF
(Z)-3-(1-benzyl-1H-imidazol-4-yl)-3-chloroprop-2-en-1-ol (264)



```

filename = sm_VI_Bn_Chloro_alcoh
author = delta
experiment = single_pulse.ex2
sample_id = S8652470
solvent = CHLOROFORM-D
reaction_time = 2-FEB-2010 17:59:21
acquisition_time = 19-APR-2010 19:08:12
current_time = 19-APR-2010 19:11:47
comment = single_pulse
ata_format = ID COMPLEX
in_size = 13107
in_title = 1H
in_units = [ppm]
inmensions = X
ite = ECX 300
spectrometer = DELTA2_NMR
ield_strength = 7.0586013[T] (300[MHz]
_acq_duration = 2.90717696[s]
_domain = 1H
_freq = 300.52965592[MHz]
_offset = 5[ppm]
_points = 16384
_prescans = 0
_resolution = 0.34397621[Hz]
_sweep = 5.63570784[kHz]
tr_domain = 1H
tr_freq = 300.52965592[MHz]
tr_offset = 5[ppm]
ri_domain = 1H
ri_freq = 300.52965592[MHz]
ri_offset = 5[ppm]
lipped = FALSE
od_return = 1
otal_scans = 24
_90_width = 13.01[us]
_acq_time = 2.90717696[s]
_angle = 45[deg]
_atn = 4[dB]
_pulse = 6.505[us]
tr_mode = Off
ri_mode = Off
ante_presat = FALSE
nitial_wait = 1[s]
ecvr_gain = 50
relaxation_delay = 5[s]
petition_time = 7.90717696[s]
emp_get = 19.8[dc]

```





```

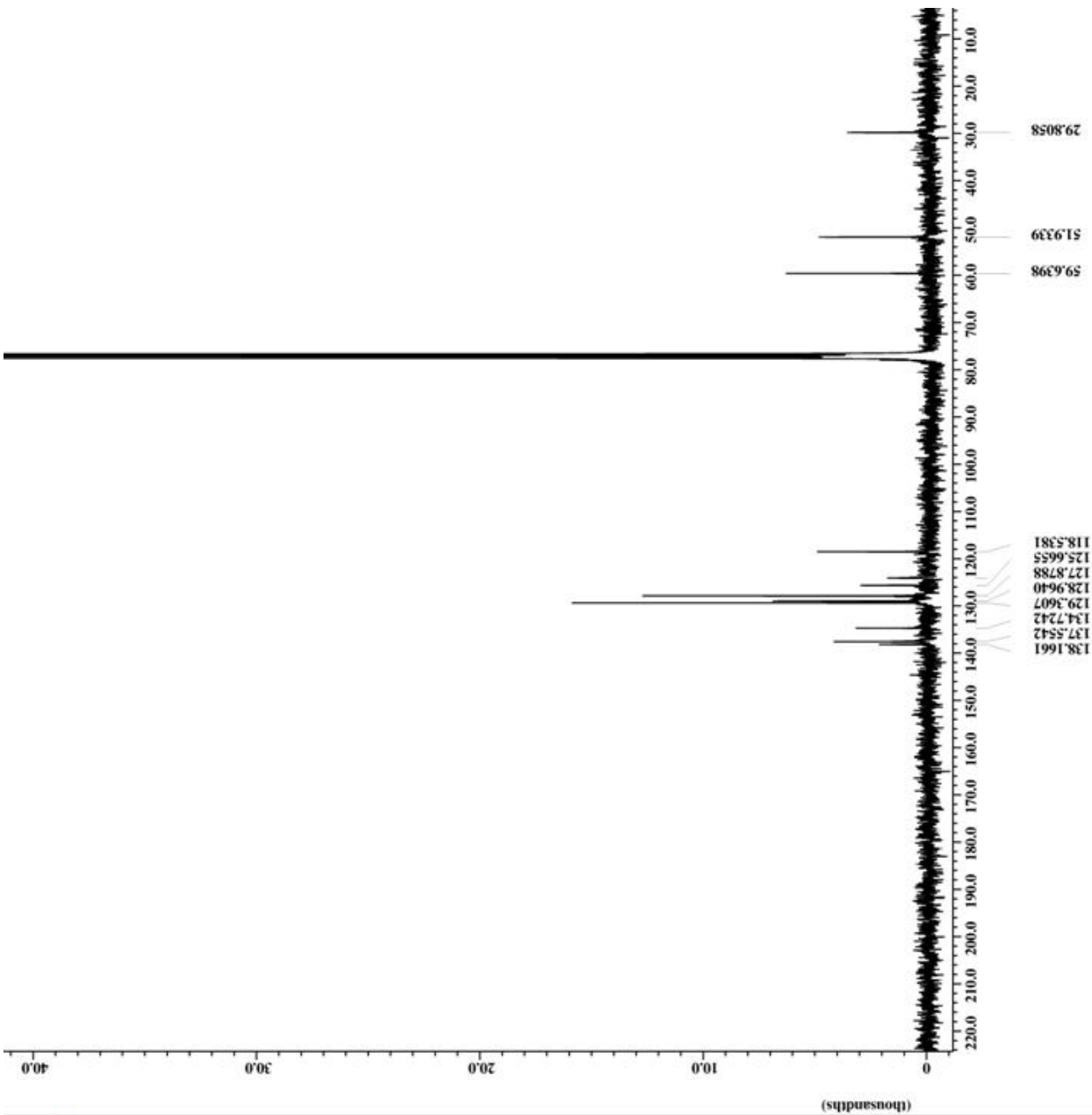
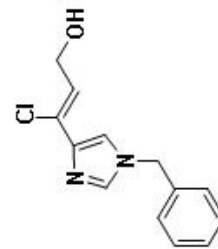
filename = sm_VI_Bn_Chloro_alcoh
author = delta
experiment = single_pulse_dec
sample_id = S854224
solvent = CHLOROFORM-D
acquisition_time = 3-FEB-2010 05:09:11
revision_time = 19-APR-2010 19:19:07
current_time = 19-APR-2010 19:20:08

comment = single pulse decouple
ata_format = ID COMPLEX
in_size = 52428
in_title = 13C
in_units = [ppm]
in_dimensions = X
ite = ECX 300
spectrometer = DELTA2_NMR

field_strength = 7.0586013[T] (300[MHz]
acq_duration = 2.76824064[s]
domain = 13C
freq = 75.56823426[MHz]
offset = 100[ppm]
points = 65536
prescans = 4
resolution = 0.36124027[Hz]
sweep = 23.67424242[kHz]
tr_domain = 1H
tr_freq = 300.52965592[MHz]
tr_offset = 5[ppm]
lipped = FALSE
bd_return = 10
cans = 3500
atal_scans = 3500

_90_width = 9.75[us]
acq_time = 2.76824064[s]
angle = 30[deg]
atn = 8[db]
pulse = 3.25[us]
tr_atn_dec = 25[db]
tr_atn_noe = 25[db]
tr_noise = WALTZ
scoupling = TRUE
nitral_wait = 1[s]
oe_time = TRUE
oe_time = 3[s]
ecvr_gain = 50
relaxation_delay = 3[s]
repetition_time = 5.76824064[s]
temp_get = 19.5[deg]

```

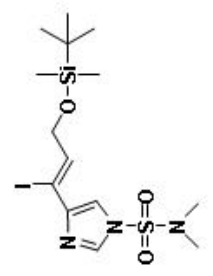
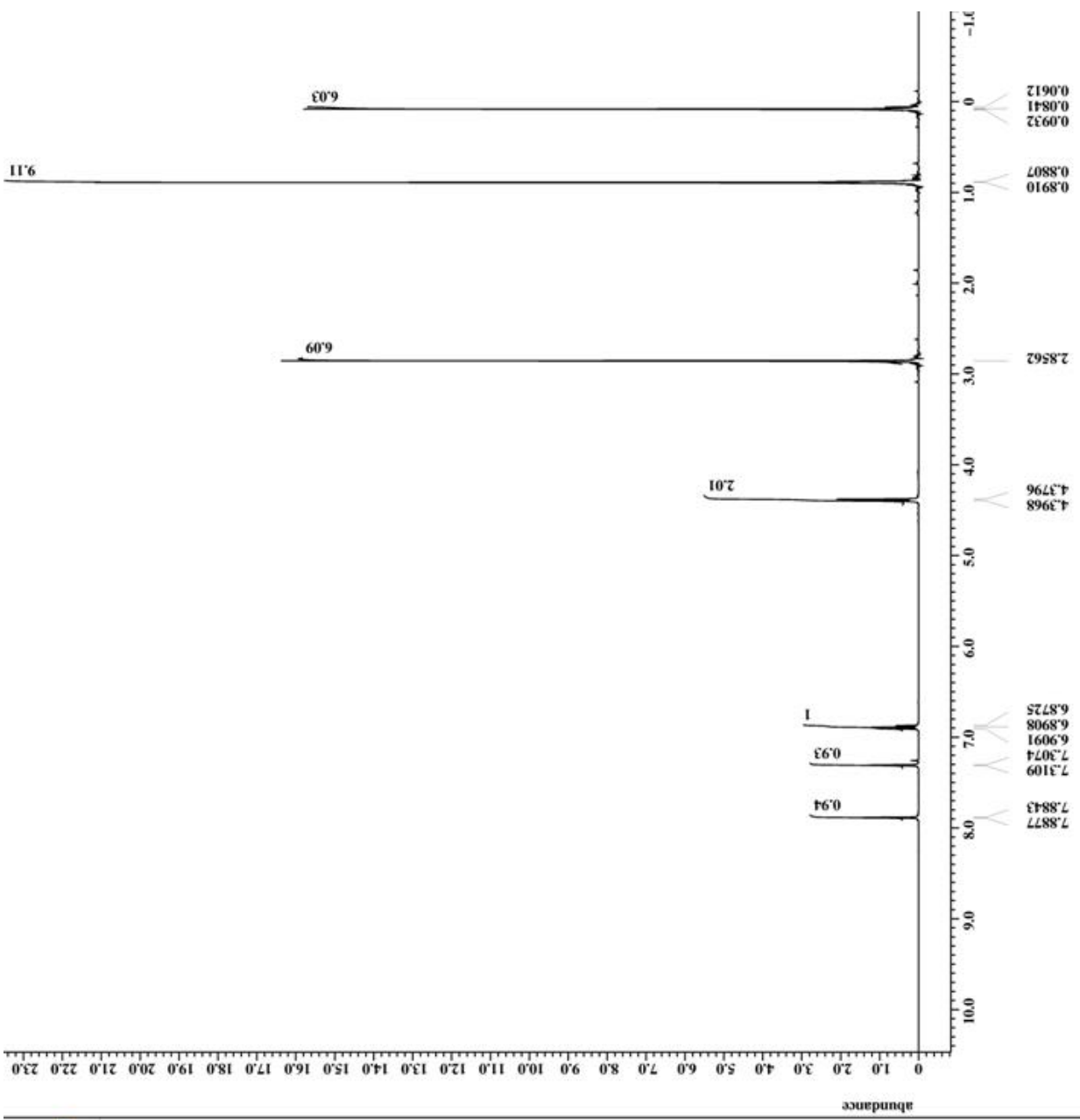


APPENDIX 52
¹H AND ¹³C NMR SPECTRUM OF
(Z)-4-(3-(tert-butyldimethylsilyloxy)-1-iodoprop-1-enyl)-N,N-dimethyl-1H-
imidazole-1-sulfonamide (255)



```

filename = sm_VI_65_pure-2.jdf
author = delta
experiment = single_pulse.ex2
sample_id = S834849
solvent = CHLOROFORM-D
creation_time = 15-DEC-2009 23:22:18
revision_time = 28-MAR-2010 22:33:10
current_time = 28-MAR-2010 22:33:37
comment =
  = single_pulse
  = ID COMPLEX
  = 13107
  = 1H
  = [ppm]
  = X
  = ECX 300
  = DELTA2_NMR
  = 7.0586013[T] (300[MHz]
  = 2.90717696[s]
  = 1H
  = 300.52965592[MHz]
  = 5[ppm]
  = 16384
  = 0
  = 0.34397621[Hz]
  = 5.63570784[kHz]
  = 1H
  = 300.52965592[MHz]
  = 5[ppm]
  = 1H
  = 300.52965592[MHz]
  = 5[ppm]
  = FALSE
  = 1
  = 20
  = 20
  = 13.01[us]
  = 2.90717696[s]
  = 45[deg]
  = 4[db]
  = 6.505[us]
  = Off
  = Off
  = FALSE
  = 1[s]
  = 30
  = 5[s]
  = 7.90717696[s]
  = 23.1[dc]
  
```





```

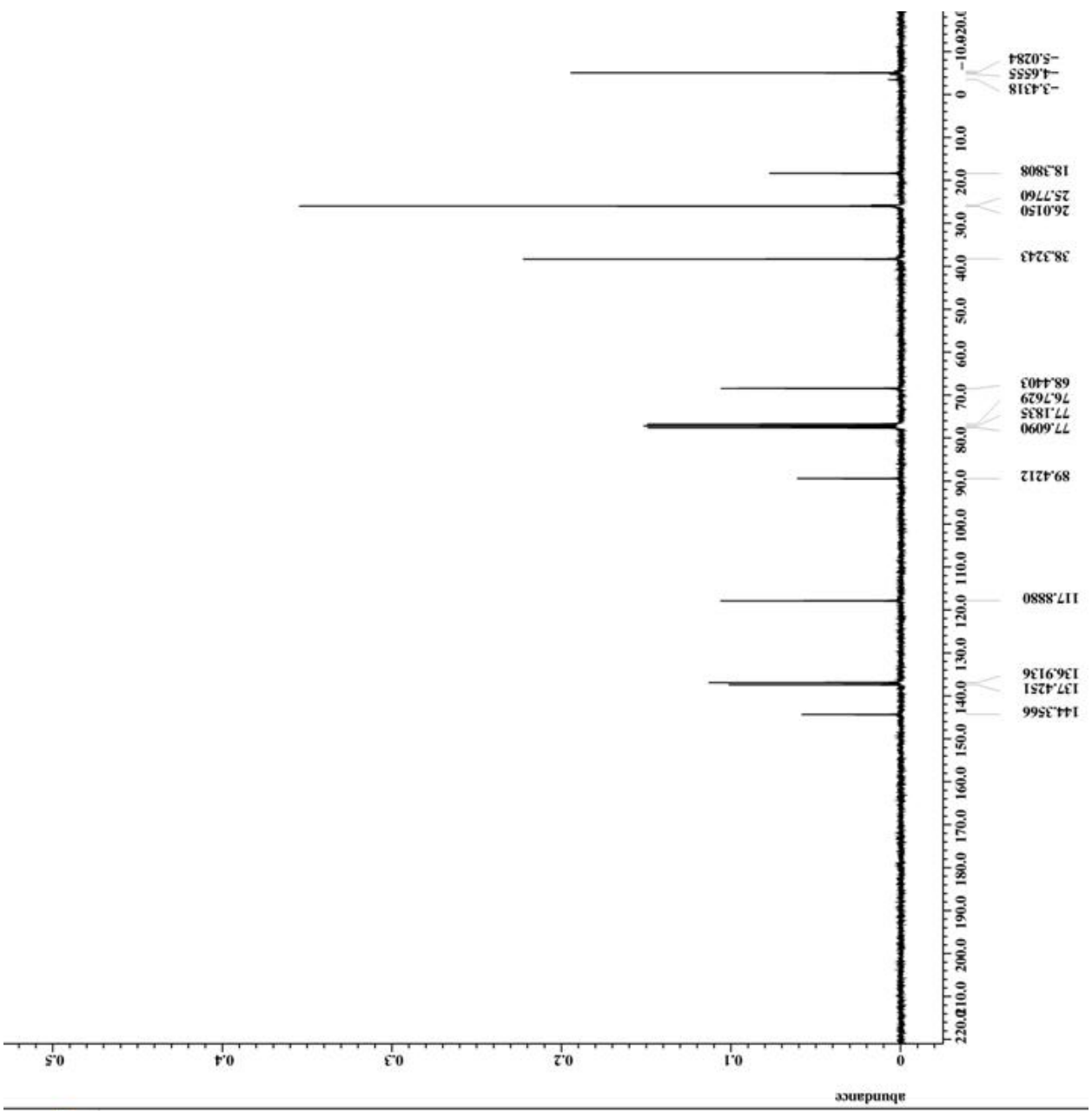
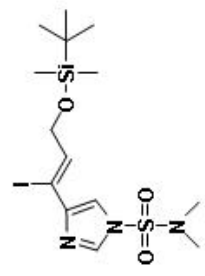
filename = sm_VI_65_pure-2.jdf
author = delta
experiment = single_pulse_dec
pulse_id = S836865
solvent = CHLOROFORM-D
acquisition_time = 16-DEC-2009 02:38:36
revision_time = 16-DEC-2009 13:10:09
current_time = 28-MAR-2010 22:36:13

comment = single pulse decouple
ata_format = ID COMPLEX
in_size = 52428
in_title = 13C
in_units = [ppm]
in_dimensions = X
ite = ECX 300
spectrometer = DELTA2_NMR

ield_strength = 7.0586013[T] (300[MHz])
acq_duration = 2.76824064[s]
domain = 13C
freq = 75.56823426[MHz]
offset = 100[ppm]
points = 65536
prescans = 40
resolution = 0.36124027[Hz]
sweep = 23.67424242[kHz]
tr_domain = 1H
tr_freq = 300.52965592[MHz]
tr_offset = 5[ppm]
lipped = TRUE
bd_return = 10
cans = 2000
total_scans = 2000

_90_width = 9.75[us]
acq_time = 2.76824064[s]
angle = 30[deg]
atn = 8[db]
pulse = 3.25[us]
tr_atn_dec = 25[db]
tr_atn_noe = 25[db]
tr_noise = WALTZ
scoupling = TRUE
nitral_wait = 1[s]
be_time = 3[s]
be_time = 3[s]
scvr_gain = 50
relaxation_delay = 3[s]
petition_time = 5.76824064[s]
emp_get = 24.4[dc]

```



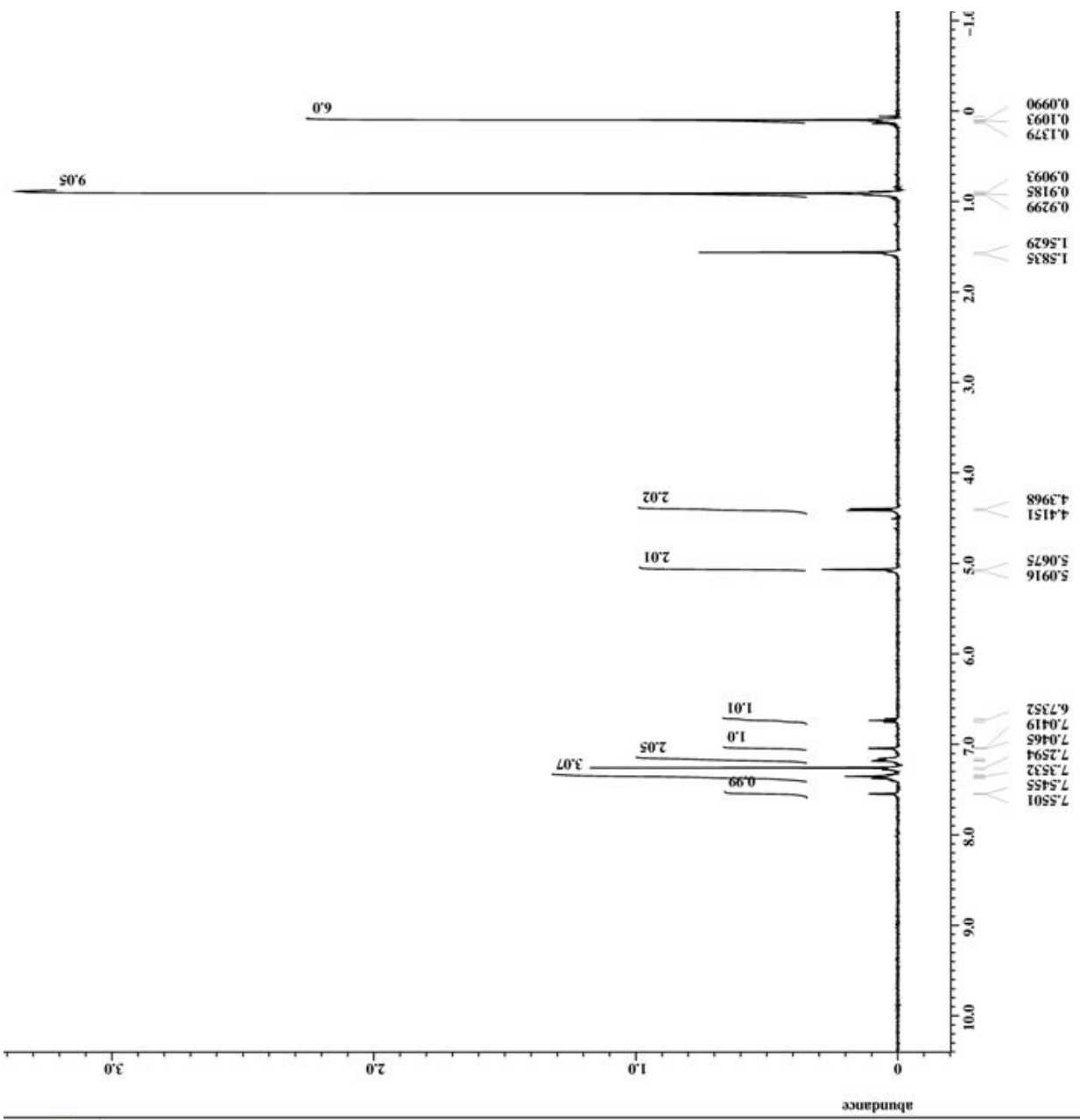
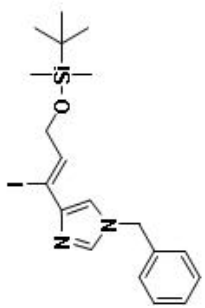
APPENDIX 53
¹H AND ¹³C NMR SPECTRUM OF
(Z)-1-benzyl-4-(3-(tert-butyldimethylsilyloxy)-1-iodoprop-1-enyl)-1H-
imidazole (253)



```

filename = sm_VI_15_pure-4.jdf
author = delta
experiment = single_pulse.ex2
sample_id = S#799735
solvent = CHLOROFORM-D
reaction_time = 14-OCT-2009 22:16:13
acquisition_time = 28-MAR-2010 22:39:23
current_time = 28-MAR-2010 22:39:45
comment = single_pulse
ata_format = ID REAL
in_size = 13107
in_title =
in_units = [ppm]
inmensions = X
ite = ECX 300
spectrometer = DELTA2_NMR
ield_strength = 7.0586013[T] (300[MHz]
_acq_duration = 2.90717696[s]
_domain = 1H
_freq = 300.52965592[MHz]
_offset = 5[ppm]
_points = 16384
_prescans = 0
_resolution = 0.34397621[Hz]
_sweep = 5.63570784[kHz]
tr_domain = 1H
tr_freq = 300.52965592[MHz]
tr_offset = 5[ppm]
tr_domain = 1H
ri_domain = 1H
ri_freq = 300.52965592[MHz]
ri_offset = 5[ppm]
lipped = FALSE
od_return = 1
cans = 16
otal_scans = 16
_90_width = 13.01[us]
_acq_time = 2.90717696[s]
_angle = 45[deg]
_atn = 4[db]
_pulse = 6.505[us]
tr_mode = Off
ri_mode = Off
ante_presat = FALSE
nitial_wait = 1[s]
scvr_gain = 46
relaxation_delay = 5[s]
petition_time = 7.90717696[s]
emp_get = 23.1[dc]

```





```

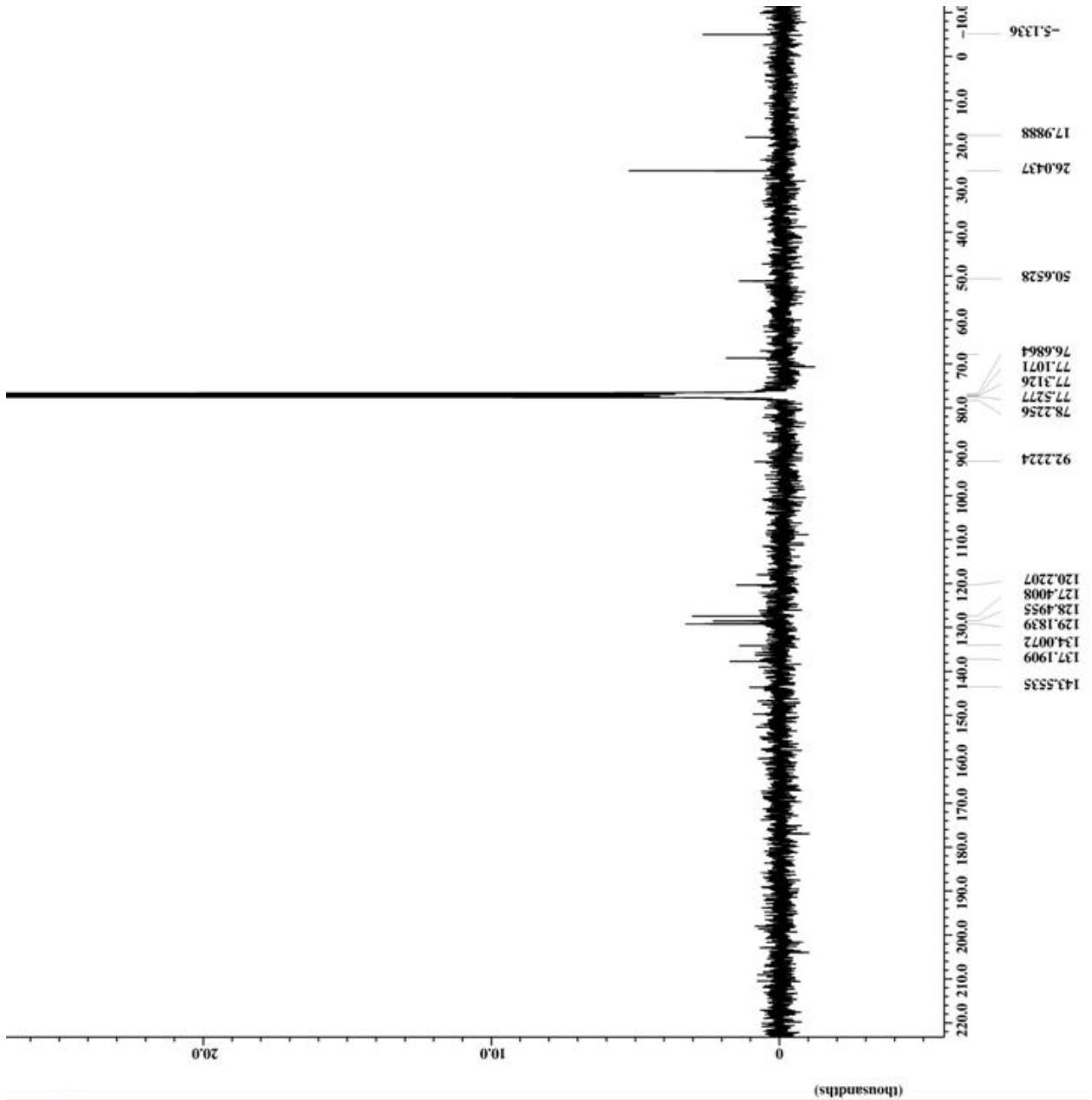
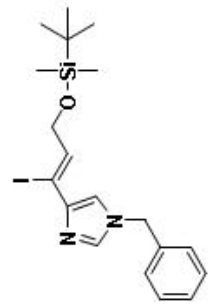
filename = sm_VI_15_pure-3.jdf
author = delta
experiment = single_pulse_dec
pulse_id = S#801383
solvent = CHLOROFORM-D
reaction_time = 15-OCT-2009 03:05:14
revision_time = 28-MAR-2010 22:43:04
current_time = 28-MAR-2010 22:43:39

comment = single pulse decouple
ata_format = ID COMPLEX
in_size = 52428
in_title = 13C
in_units = [ppm]
inmensions = X
ite = ECX 300
spectrometer = DELTA2_NMR

ield_strength = 7.0586013[T] (300[MHz]
acq_duration = 2.76824064[s]
domain = 13C
freq = 75.56823426[MHz]
offset = 100[ppm]
points = 65536
prescans = 4
resolution = 0.36124027[Hz]
sweep = 23.67424242[kHz]
tr_domain = 1H
tr_freq = 300.52965592[MHz]
tr_offset = 5[ppm]
lipped = FALSE
bd_return = 10
cans = 3000
atal_scans = 3000

_90_width = 9.75[us]
acq_time = 2.76824064[s]
angle = 30[deg]
atn = 8[db]
pulse = 3.25[us]
tr_atn_dec = 25[db]
tr_atn_noe = 25[db]
tr_noise = WALTZ
scoupling = TRUE
nitiaal_wait = 1[s]
be_time = TRUE
be_time = 3[s]
ecvr_gain = 50
relaxation_delay = 3[s]
spetition_time = 5.76824064[s]
emp_get = 23.2[dc]

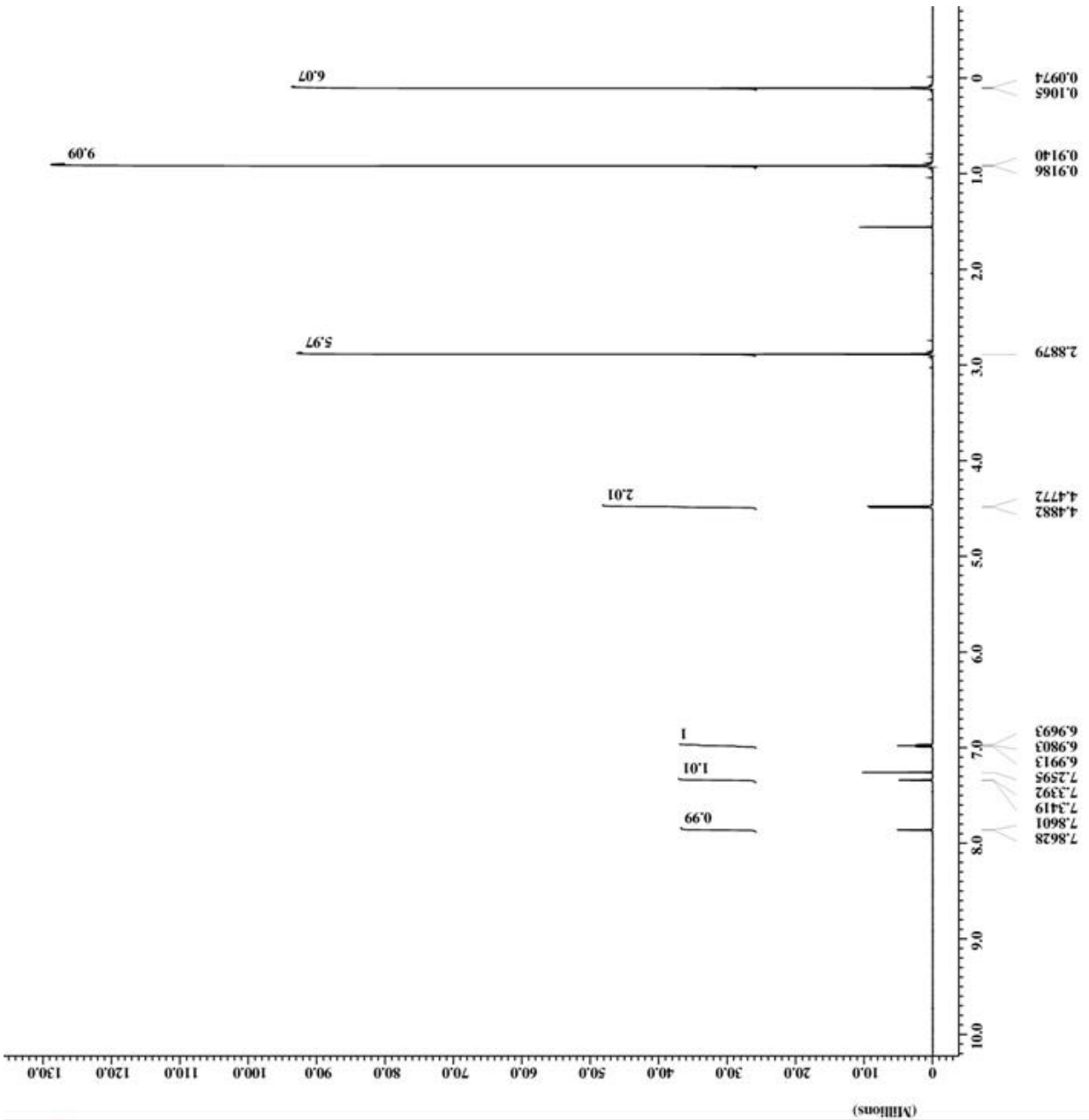
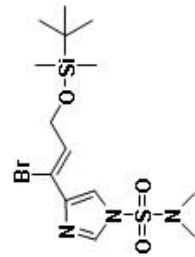
```



APPENDIX 54
¹H AND ¹³C NMR SPECTRUM OF
(Z)-4-(1-bromo-3-(tert-butyldimethylsilyloxy)prop-1-enyl)-N,N-dimethyl-1H-
imidazole-1-sulfonamide (261)



```
filename = sm_VI_76_pure-3.jdf
author = delta
experiment = single_pulse.exp
sample_id = S#7037
solvent = CHLOROFORM-D
reaction_time = 29-DEC-2009 15:55:28
revision_time = 28-MAR-2010 22:46:51
current_time = 28-MAR-2010 22:47:13
comment = Single Pulse Experiment
data_format = ID COMPLEX
in_size = 16384
in_title = 1H
in_units = [ppm]
inmensions = X
ite = Eclipse+ 500
spectrometer = DELTA_NMR
field_strength = 11.7465928[T] (500[MH
-acq_duration = 2.1839872[s]
-domain = 1H
-freq = 500.12734003[MHz]
-gamma = 5[ppm]
-points = 16384
-prescans = 0
-resolution = 0.45787814[Hz]
-sweep = 7.50187547[kHz]
-tipped = FALSE
-od_return = 1
-cans = 8
-otat_scans = 8
-90_width = 18.5[us]
-acq_time = 2.1839872[s]
-angle = 45[deg]
-pulse = 7.25[us]
-nitral_wait = 1[s]
-base_preset = 2[us]
-ecvr_gain = 22
-relaxation_delay = 4[s]
-emp_get = 29.6[dc]
-nblank_time = 2[us]
```





```

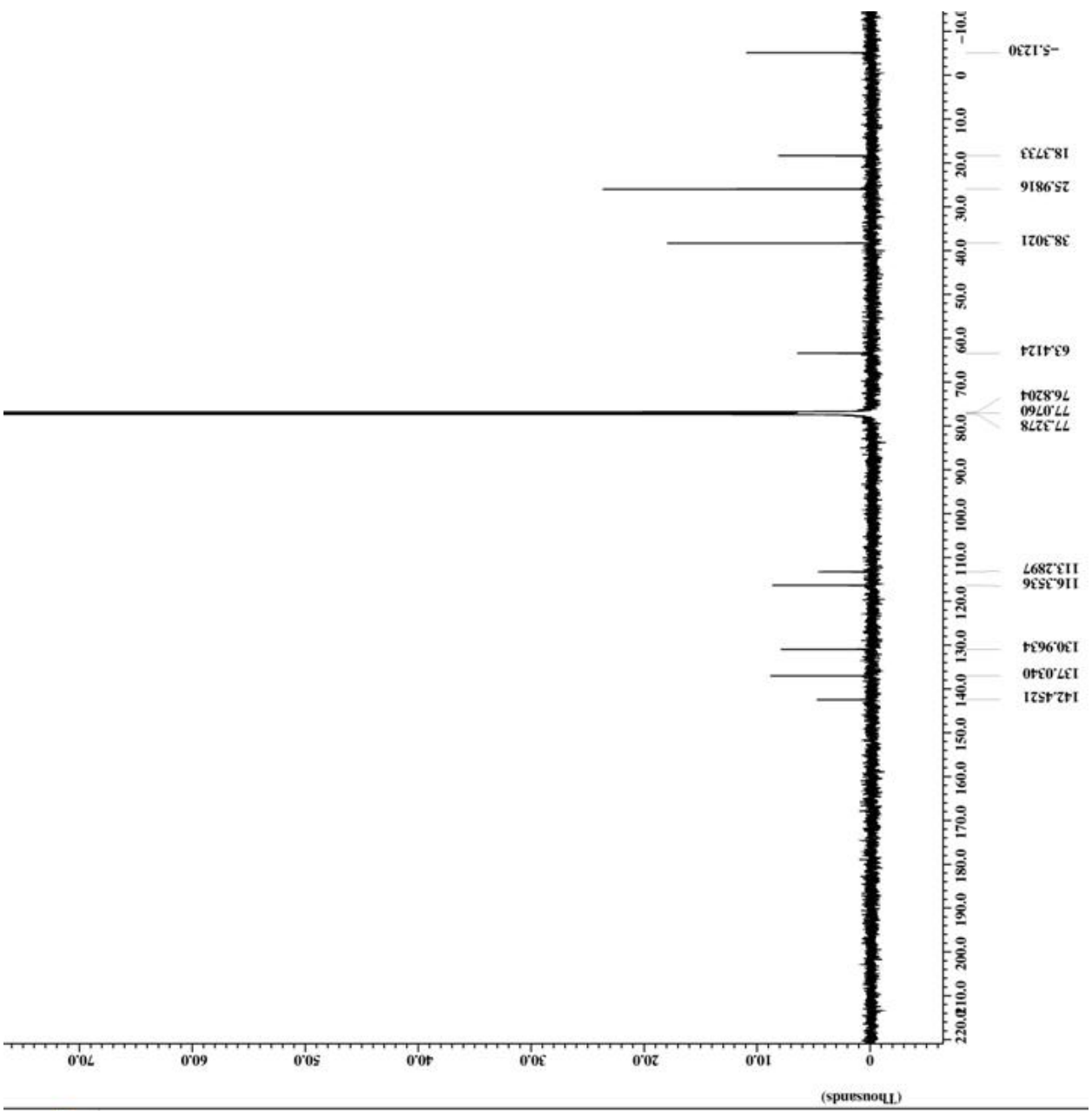
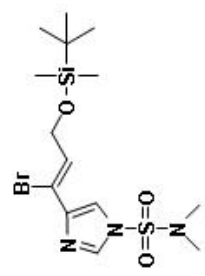
filename = sm_VI_76_pure-2.jdf
author = delta
experiment = single_pulse_dec
sample_id = S8266
solvent = CHLOROFORM-D
reaction_time = 29-DEC-2009 23:00:01
revision_time = 29-DEC-2009 13:52:02
current_time = 28-MAR-2010 22:49:53

comment = single pulse decouple
ata_format = ID COMPLEX
im_size = 65536
im_title = 13C
im_units = [ppm]
imensions = X
ite = Eclipse+ 500
spectrometer = DELTA_NMR

field_strength = 11.7465928[T] (500[MH
-acq_duration = 2.0840448[s]
-domain = 13C
-freq = 125.75710665[MHz]
-offset = 100[ppm]
-points = 65536
-prescans = 4
-resolution = 0.47983613[Hz]
-sweep = 31.44654088[kHz]
-tr_domain = 1H
-tr_freq = 500.12734003[MHz]
-tr_offset = 5[ppm]
-tipped = FALSE
-bd_return = 10
-cans = 5000
-etal_scans = 5000

_90_width = 14.2[us]
-acq_time = 2.0840448[s]
-angle = 30[deg]
-pulse = 4.73333333[us]
-nitral_wait = 1[s]
-pe_time = 1[s]
-base_preset = 3[us]
-ecvr_gain = 22
-relaxation_delay = 2[s]
-emp_get = 32.5[dC]
-nblank_time = 2[us]

```



APPENDIX 55
¹H AND ¹³C NMR SPECTRUM OF
(Z)-1-benzyl-4-(1-bromo-3-(tert-butyldimethylsilyloxy)prop-1-enyl)-1H-
imidazole (262)



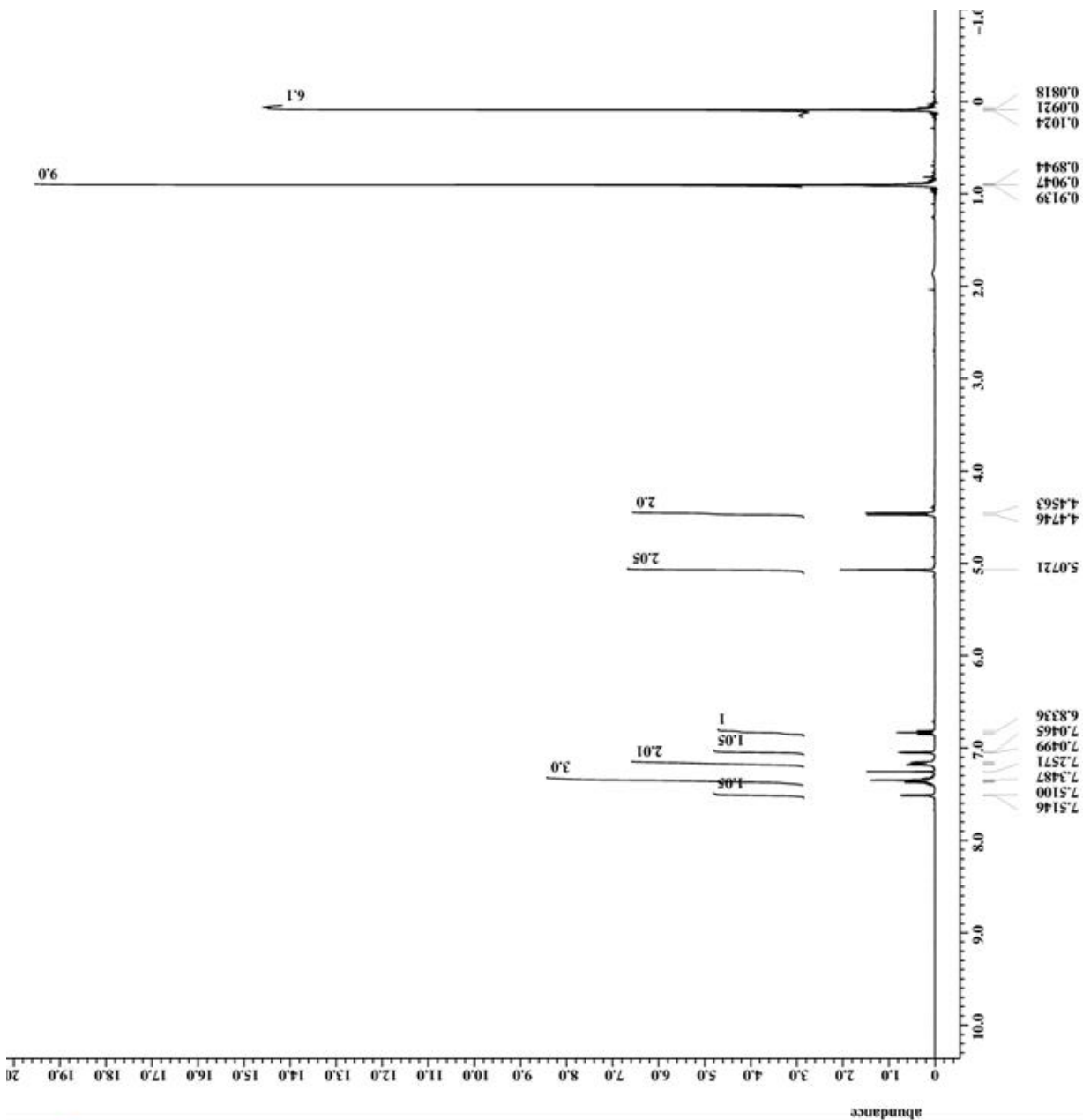
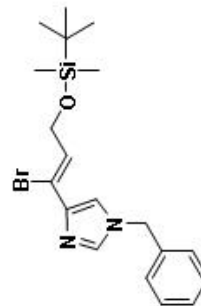
```

ilname      = sm_VI_70_pure-3.jdf
author      = delta
experiment  = single_pulse.ex2
sample_id   = S807529
solvent     = CHLOROFORM-D
reaction_time = 17-DEC-2009 22:37:15
revision_time = 28-MAR-2010 22:53:12
current_time = 28-MAR-2010 22:53:32

comment     = single_pulse
ate_format  = ID_COMPLEX
im_size     = 13107
im_title    = 1H
im_units    = [ppm]
imensions   = X
ite         = ECX 300
pctrometer  = DELTA2_NMR

field_strength = 7.0586013[T] (300[MHz])
-acq_duration = 2.90717696[s]
-domain       = 1H
-freq        = 300.52965592[MHz]
-offset      = 5[ppm]
-points      = 16384
-prescans    = 0
-resolution  = 0.34397631[Hz]
-sweep       = 5.63570784[kHz]
-rr_domain   = 1H
-rr_freq     = 300.52965592[MHz]
-rr_offset   = 5[ppm]
-ri_domain   = 1H
-ri_freq     = 300.52965592[MHz]
-ri_offset   = 5[ppm]
-lipped      = FALSE
-bd_return   = 1
-cans        = 21
-stal_scans  = 21

-90_width    = 13.01[us]
-acq_time    = 2.90717696[s]
-angle       = 45[deg]
-atn         = 4[dB]
-pulse       = 6.505[us]
-rr_mode     = Off
-ri_mode     = Off
-ante_presat = FALSE
-nit1al_wait = 1[s]
-scvr_gain   = 44
-relaxation_delay = 5[s]
-epitation_time = 7.90717696[s]
-emp_get     = 23.2[dC]
  
```

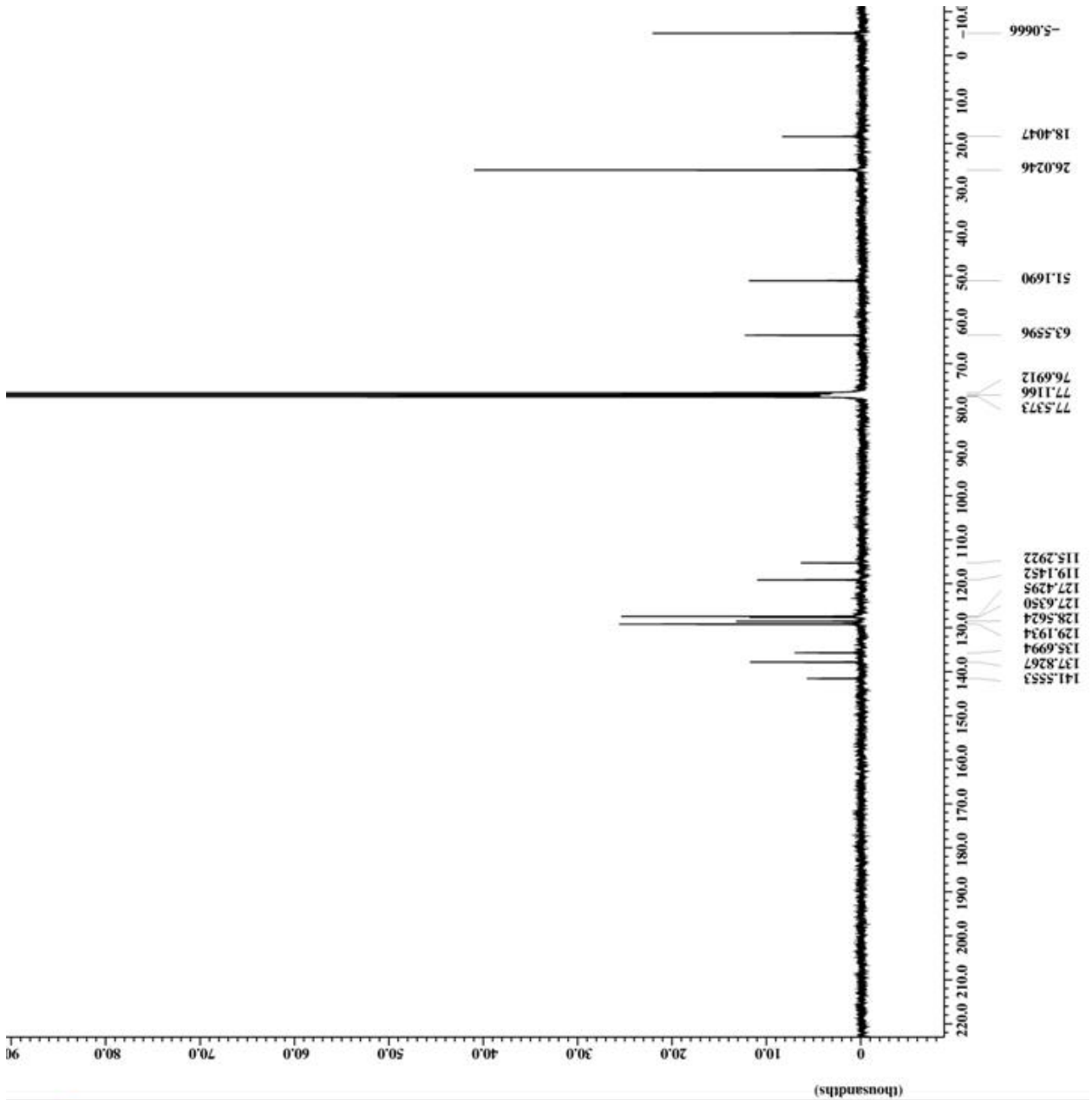
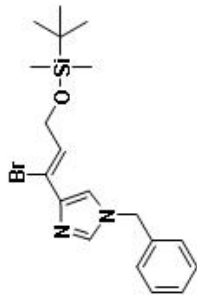




```

filename = sm_VI_70_pure-2.jdf
author = delta
experiment = single_pulse_dec
pulse_id = S#809631
solvent = CHLOROFORM-D
reaction_time = 18-DEC-2009 06:38:27
revision_time = 18-DEC-2009 13:11:57
current_time = 28-MAR-2010 22:56:31
comment = single pulse decouple
ata_format = ID COMPLEX
in_size = 52428
in_title = 13C
in_units = [ppm]
inmensions = X
ite = ECX 300
spectrometer = DELTA2_NMR
ield_strength = 7.0586013[T] (300[MHz]
_acq_duration = 2.76824064[s]
_domain = 13C
_freq = 75.56823426[MHz]
_offset = 100[ppm]
_points = 65536
_prescans = 4
_resolution = 0.36124027[Hz]
_sweep = 23.67424242[kHz]
tr_domain = 1H
tr_freq = 300.52965592[MHz]
tr_offset = 5[ppm]
lipped = FALSE
bd_return = 10
cans = 5000
atal_scans = 5000
_90_width = 9.75[us]
_acq_time = 2.76824064[s]
_angle = 30[deg]
_atn = 8[db]
_pulse = 3.25[us]
tr_atn_dec = 25[db]
tr_atn_noe = 25[db]
tr_noise = WALTZ
scoupling = TRUE
nitial_wait = 1[s]
oe_time = TRUE
oe_time = 3[s]
scvr_gain = 50
relaxation_delay = 3[s]
petition_time = 5.76824064[s]
emp_get = 23.2[dc]

```



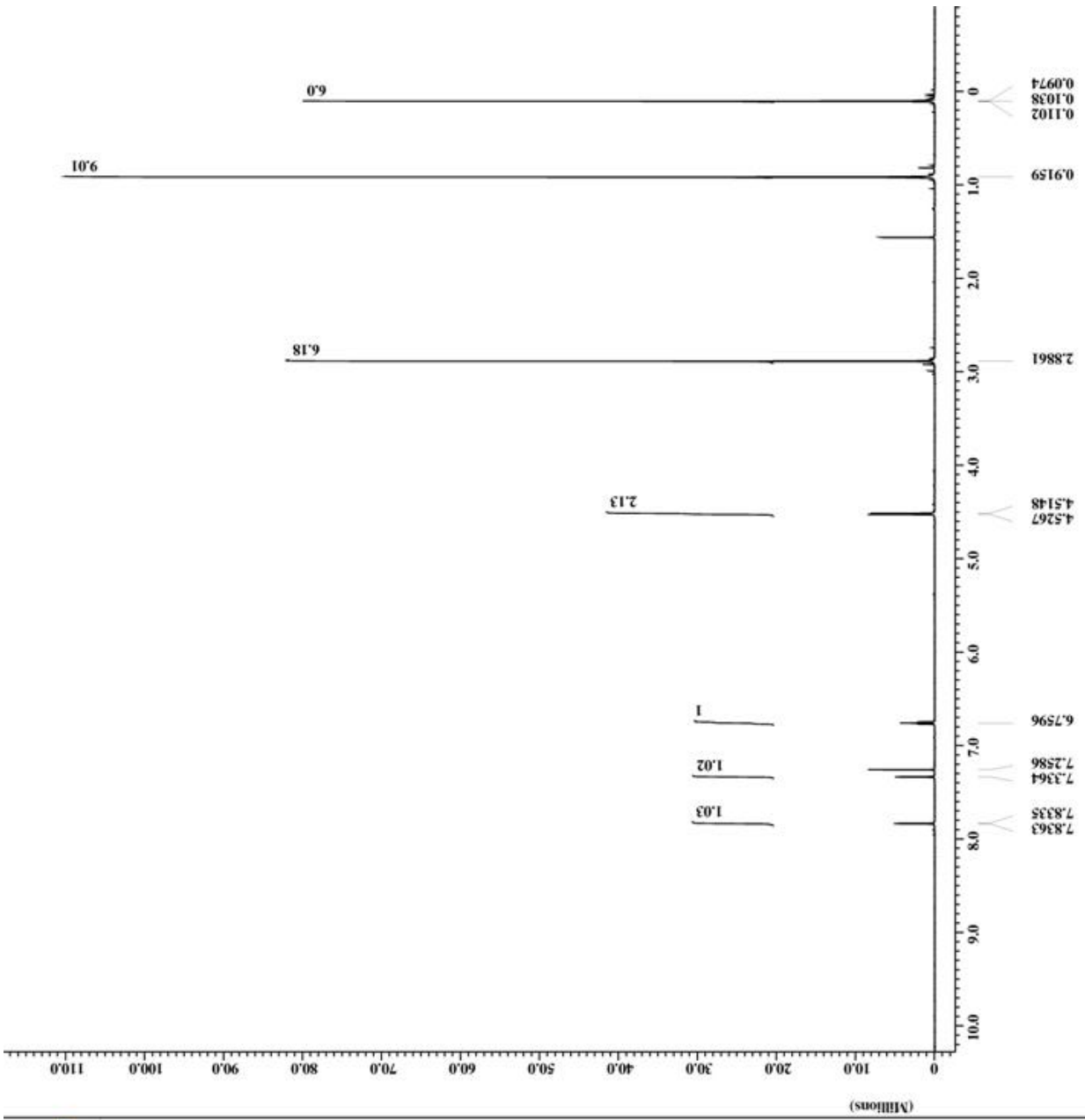
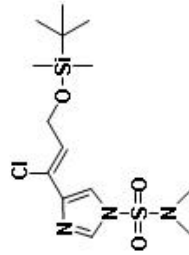
APPENDIX 56
¹H AND ¹³C NMR SPECTRUM OF
(Z)-4-(3-(tert-butyl)dimethylsilyloxy)-1-chloroprop-1-enyl)-N,N-dimethyl-1H-
imidazole-1-sulfonamide (265)



```

filename = sm_VI_77_pure-3.jdf
author = delta
experiment = single_pulse_exp
sample_id = S#529045
solvent = CHLOROFORM-D
reaction_time = 28-DEC-2009 06:24:34
revision_time = 28-MAR-2010 22:58:44
current_time = 28-MAR-2010 22:59:05
comment = Single Pulse Experiment
data_format = ID COMPLEX
in_size = 16384
in_title = 1H
in_units = [ppm]
instruments = X
ite = Eclipse+ 500
nucleus1 = DELTA_NMR
p1 = 11.7465928[T] (500[MH
-acq_duration = 2.1839872[s]
-domain = 1H
-freq = 500.12734003[MHz]
-gamma = 5[ppm]
-points = 16384
-prescans = 0
-resolution = 0.45787814[Hz]
-sweep = 7.50187547[kHz]
-tipped = FALSE
-od_return = 1
-cans = 8
-otol_scans = 8
-90_width = 18.5[us]
-acq_time = 2.1839872[s]
-angle = 45[deg]
-pulse = 7.25[us]
-nitral_wait = 1[s]
-base_preset = 2[us]
-ecvr_gain = 22
-relaxation_delay = 4[s]
-emp_get = 29[dc]
-nblank_time = 2[us]

```





```

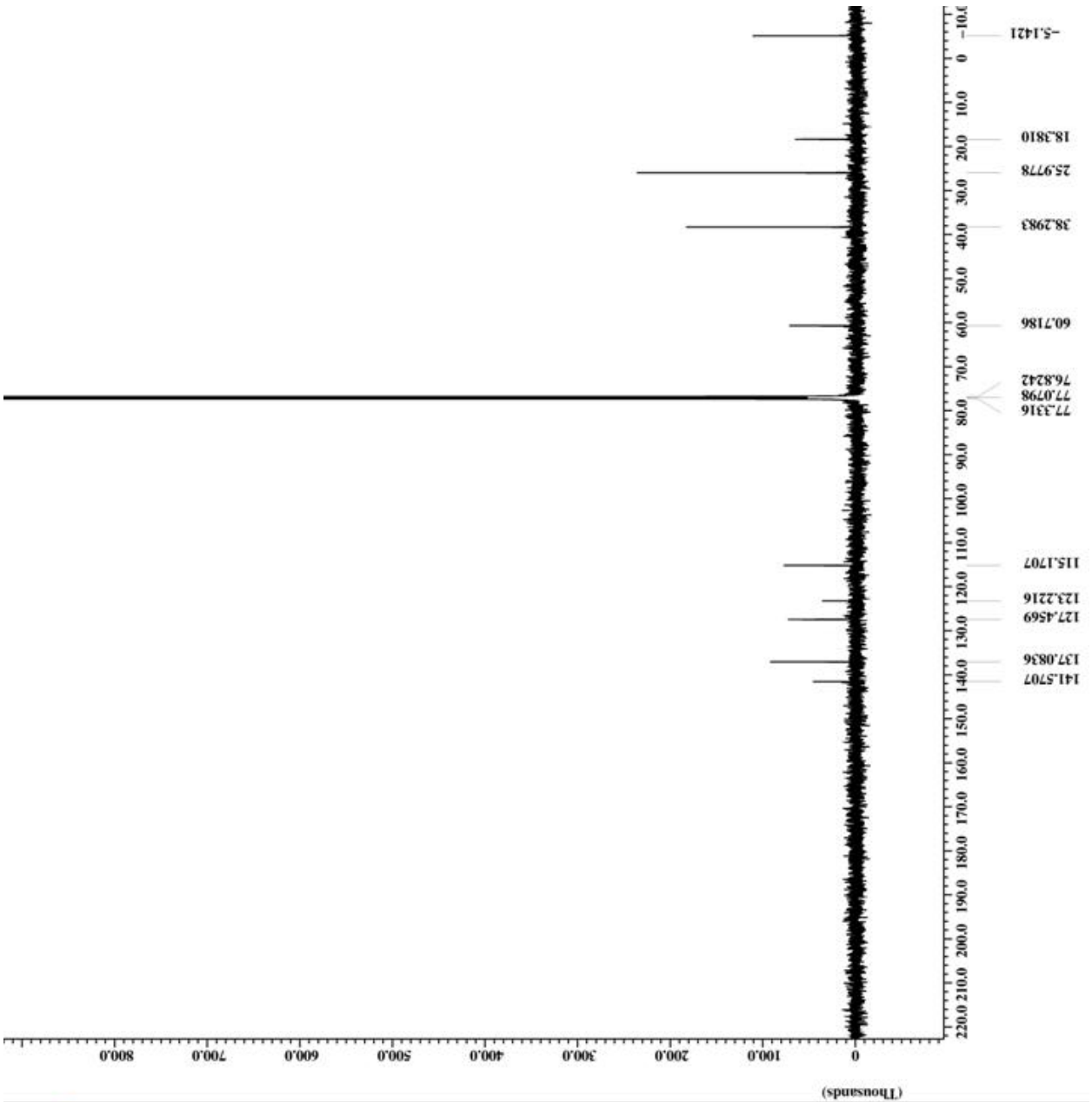
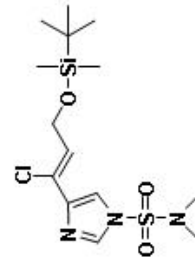
filename = sm_VI_77_pure-2.jdf
author = delta
experiment = single_pulse_dec
sample_id = S9530285
solvent = CHLOROFORM-D
acquisition_time = 28-DEC-2009 08:33:31
revision_time = 28-DEC-2009 00:06:42
current_time = 28-MAR-2010 23:01:53

comment = single pulse decouple
ata_format = ID COMPLEX
im_size = 65536
im_title = 13C
im_units = [ppm]
imensions = X
ite = Eclipse+ 500
spectrometer = DELTA_NMR

field_strength = 11.7465928[T] (500[MH
-acq_duration = 2.0840448[s]
-domain = 13C
-freq = 125.75710665[MHz]
-offset = 100[ppm]
-points = 65536
-prescans = 4
-resolution = 0.47983613[Hz]
-sweep = 31.44654088[kHz]
-tr_domain = 1H
-tr_freq = 500.12734003[MHz]
-tr_offset = 5[ppm]
-tipped = TRUE
-bd_return = 10
-cans = 1500
-etal_scans = 1500

_90_width = 14.2[us]
-acq_time = 2.0840448[s]
-angle = 30[deg]
-pulse = 4.73333333[us]
-nitral_wait = 1[s]
-be_time = 1[s]
-base_preset = 3[us]
-ecvr_gain = 29
-relaxation_delay = 2[s]
-emp_get = 31.9[dC]
-nblank_time = 2[us]

```

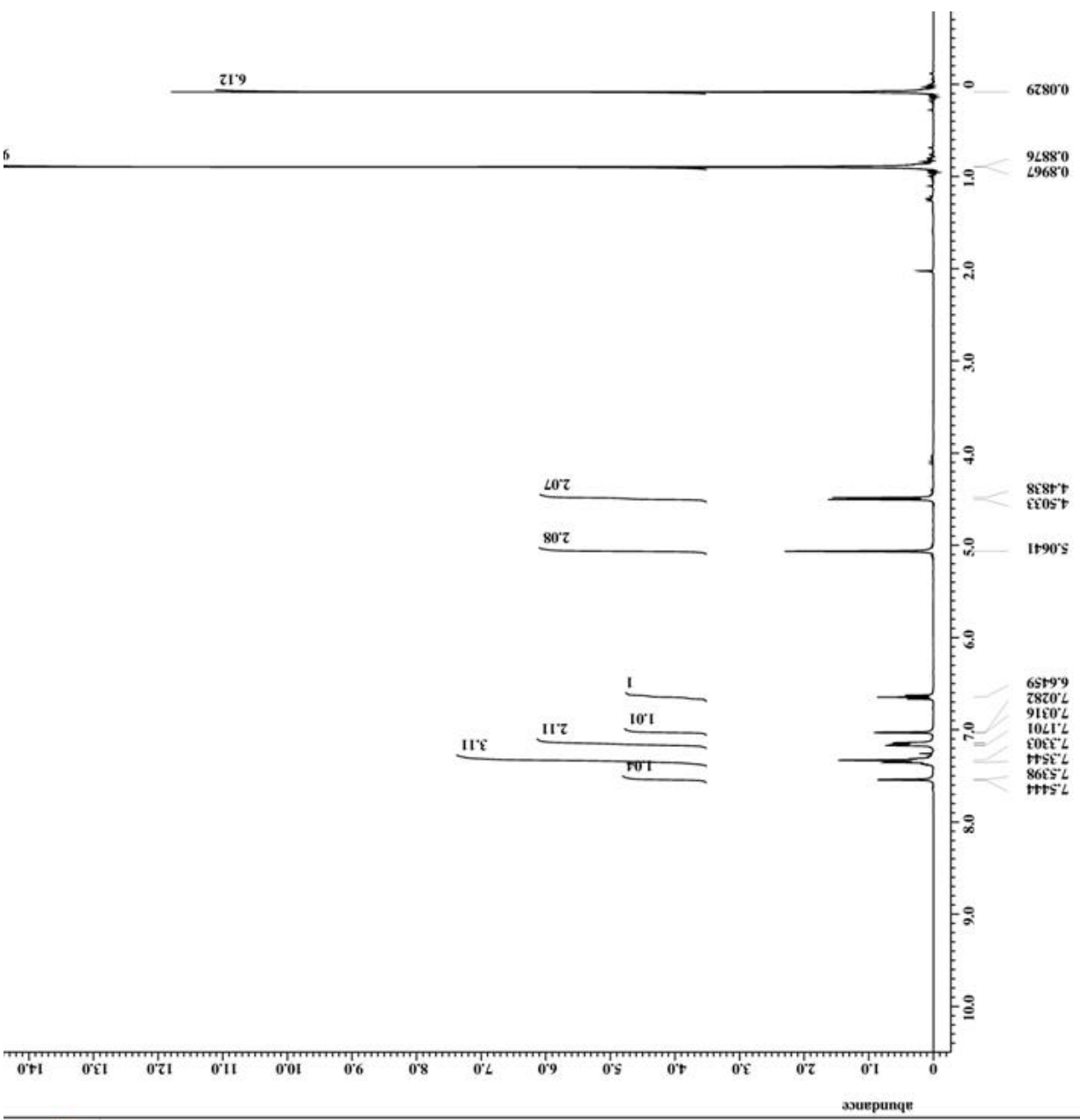
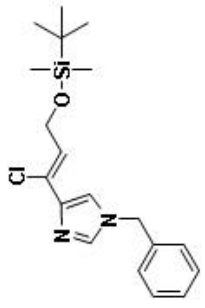


APPENDIX 57
¹H AND ¹³C NMR SPECTRUM OF
(Z)-1-benzyl-4-(3-(tert-butyldimethylsilyloxy)-1-chloroprop-1-enyl)-1H-
imidazole (266)



```

filename = sm_VI_86_pure-3_jdf
author = delta
experiment = single_pulse.ex2
sample_id = S#860471
solvent = CHLOROFORM-D
creation_time = 10-JAN-2010 23:42:17
revision_time = 28-MAR-2010 23:04:15
current_time = 28-MAR-2010 23:04:40
comment = single_pulse
ata_format = ID COMPLEX
in_size = 13107
in_title = 1H
in_units = [ppm]
inmensions = X
ite = ECX 300
spectrometer = DELTA2_NMR
ield_strength = 7.0586013[T] (300[MHz]
_acq_duration = 2.90717696[s]
_domain = 1H
_freq = 300.52965592[MHz]
_offset = 5[ppm]
_points = 16384
_prescans = 0
_resolution = 0.34397621[Hz]
_sweep = 5.63570784[kHz]
rr_domain = 1H
rr_freq = 300.52965592[MHz]
rr_offset = 5[ppm]
ri_domain = 1H
ri_freq = 300.52965592[MHz]
ri_offset = 5[ppm]
lipped = FALSE
od_return = 1
otal_scans = 16
_90_width = 13.01[us]
_acq_time = 2.90717696[s]
_angle = 45[deg]
_atn = 4[db]
_pulse = 6.505[us]
rr_mode = Off
ri_mode = Off
ante_presat = FALSE
nitial_wait = 1[s]
scrvr_gain = 30
relaxation_delay = 5[s]
petition_time = 7.90717696[s]
emp_get = 21.2[dc]
  
```



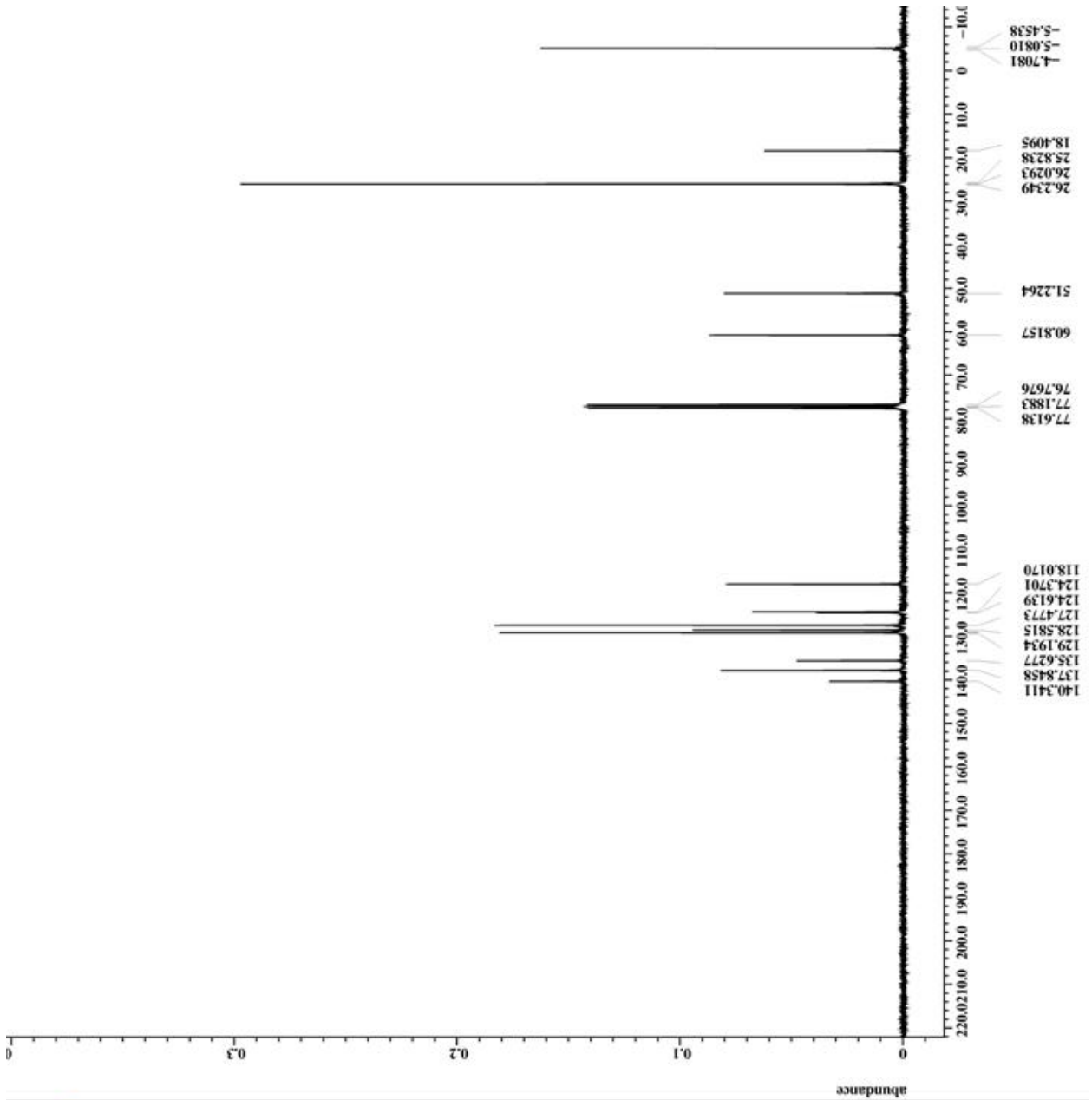
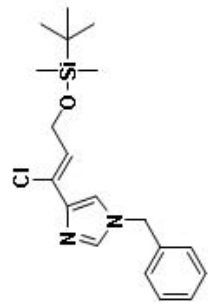


```
filename = sm_VI_86-pure-2.jdf
author = delta
experiment = single_pulse_dec
pulse_id = S#862122
solvent = CHLOROFORM-D
acquisition_time = 11-JAN-2010 05:20:31
revision_time = 28-MAR-2010 23:06:56
current_time = 28-MAR-2010 23:07:05

comment = single pulse decouple
ata_format = ID COMPLEX
in_size = 52428
in_title = 13C
in_units = [ppm]
in_dimensions = X
ite = ECX 300
spectrometer = DELTA2_NMR

ield_strength = 7.0586013[T] (300[MHz])
acq_duration = 2.76824064[s]
domain = 13C
freq = 75.56823426[MHz]
offset = 100[ppm]
points = 65536
prescans = 4
resolution = 0.36124027[Hz]
sweep = 23.67424242[kHz]
tr_domain = 1H
tr_freq = 300.52965592[MHz]
tr_offset = 5[ppm]
lipped = FALSE
bd_return = 10
cans = 3500
atal_scans = 3500

_90_width = 9.75[us]
acq_time = 2.76824064[s]
angle = 30[deg]
atn = 8[db]
pulse = 3.25[us]
tr_atn_dec = 25[db]
tr_atn_noe = 25[db]
tr_noise = WALTZ
scoupling = TRUE
nitral_wait = 1[s]
be_time = TRUE
be_time = 3[s]
ecvr_gain = 50
relaxation_delay = 3[s]
petition_time = 5.76824064[s]
emp_get = 21[dc]
```



APPENDIX 58
¹H AND ¹³C NMR SPECTRUM OF
(5a*R*^{*},8a*S*^{*})-1-benzyl-5-((tert-butyldimethylsilyloxy)methyl)-7-phenyl-7,8a-
dihydroimidazo[4,5-*e*]isoindole-6,8(1*H*,5a*H*)-dione (257)



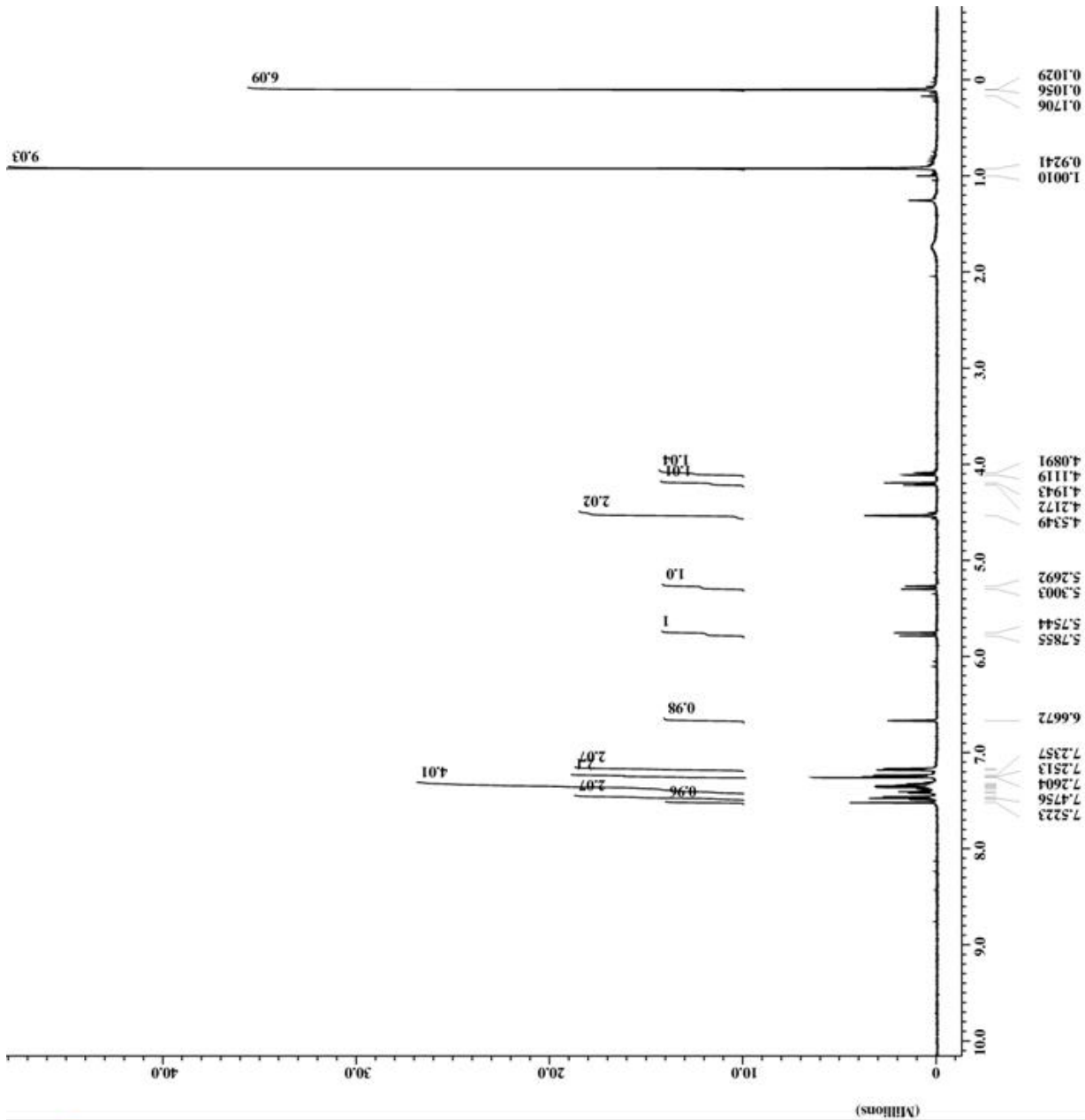
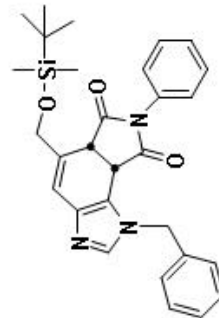
```

ilname      = sm_VI_44_pure_i1-3.jd
author      = delta
experiment  = single_pulse_exp
sample_id   = S#696981
solvent     = CHLOROFORM-D
reaction_time = 29-NOV-2009 10:46:29
revision_time = 28-MAR-2010 23:11:33
current_time = 28-MAR-2010 23:11:51

comment     = Single Pulse Experiment
ata_format  = ID COMPLEX
in_size     = 16384
in_title    = 1H
in_units    = [ppm]
inmensions  = X
ite         = Eclipse+ 500
spectrometer = DELTA_NMR

field_strength = 11.7465928[T] (500[MH]
acq_duration   = 2.1839872[s]
domain        = 1H
freq          = 500.12734003[MHz]
offset        = 5[ppm]
points        = 16384
prescans      = 0
resolution    = 0.45787814[Hz]
sweep         = 7.50187547[kHz]
lipped        = FALSE
od_return     = 1
cans          = 8
otai_scans    = 8

_90_width     = 18.5[us]
acq_time      = 2.1839872[s]
angle         = 45[deg]
pulse         = 4.25[us]
nitai_wait   = 1[s]
hase_preset  = 2[us]
ecvr_gain    = 22
relaxation_delay = 4[s]
emp_get      = 25.9[dc]
nblank_time  = 2[us]
  
```





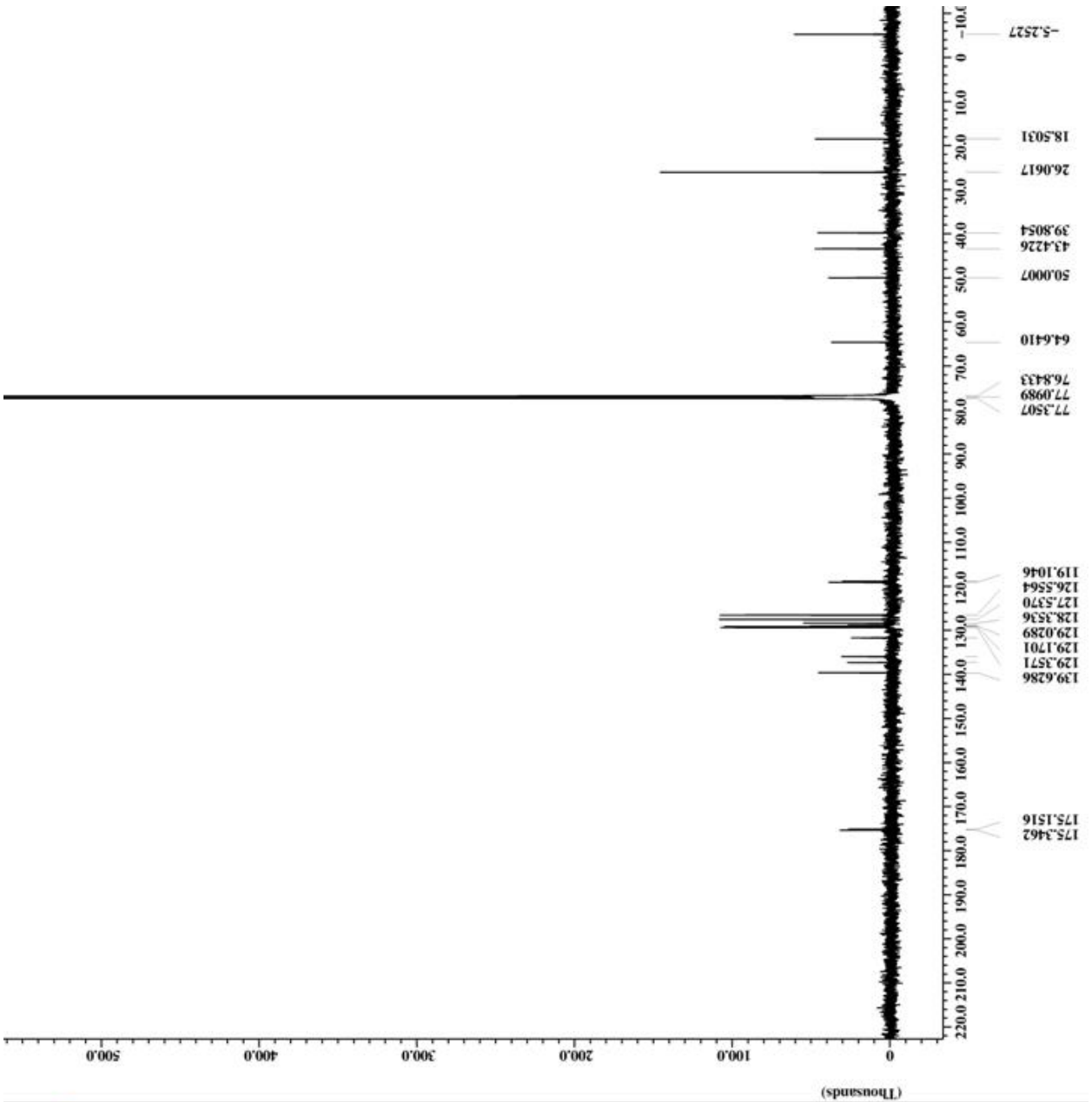
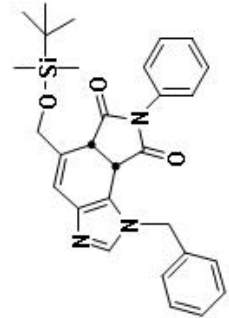
```

filename = sm_VI_44_pure_i1-2.jd
author = delta
experiment = single_pulse_dec
sample_id = S8698993
solvent = CHLOROFORM-D
reaction_time = 29-NOV-2009 16:28:27
revision_time = 30-NOV-2009 00:33:02
current_time = 28-MAR-2010 23:15:26

comment = single pulse decouple
ata_format = ID COMPLEX
in_size = 65536
in_title = 13C
in_units = [ppm]
in_dimensions = X
ite = Eclipse+ 500
spectrometer = DELTA_NMR

field_strength = 11.7465928[T] (500[MH
acq_duration = 2.0840448[s]
domain = 13C
freq = 125.75710665[MHz]
offset = 100[ppm]
points = 65536
prescans = 4
resolution = 0.47983613[Hz]
sweep = 31.44654088[kHz]
tr_domain = 1H
tr_freq = 500.12734003[MHz]
tr_offset = 5[ppm]
lipped = TRUE
bd_return = 10
cans = 4000
otal_scans = 4000

_90_width = 14.2[us]
acq_time = 2.0840448[s]
angle = 30[deg]
pulse = 4.73333333[us]
nitai_wait = 1[s]
pe_time = 1[s]
base_preset = 3[us]
scvr_gain = 28
relaxation_delay = 2[s]
emp_get = 28.6[dc]
nblank_time = 2[us]
  
```





```

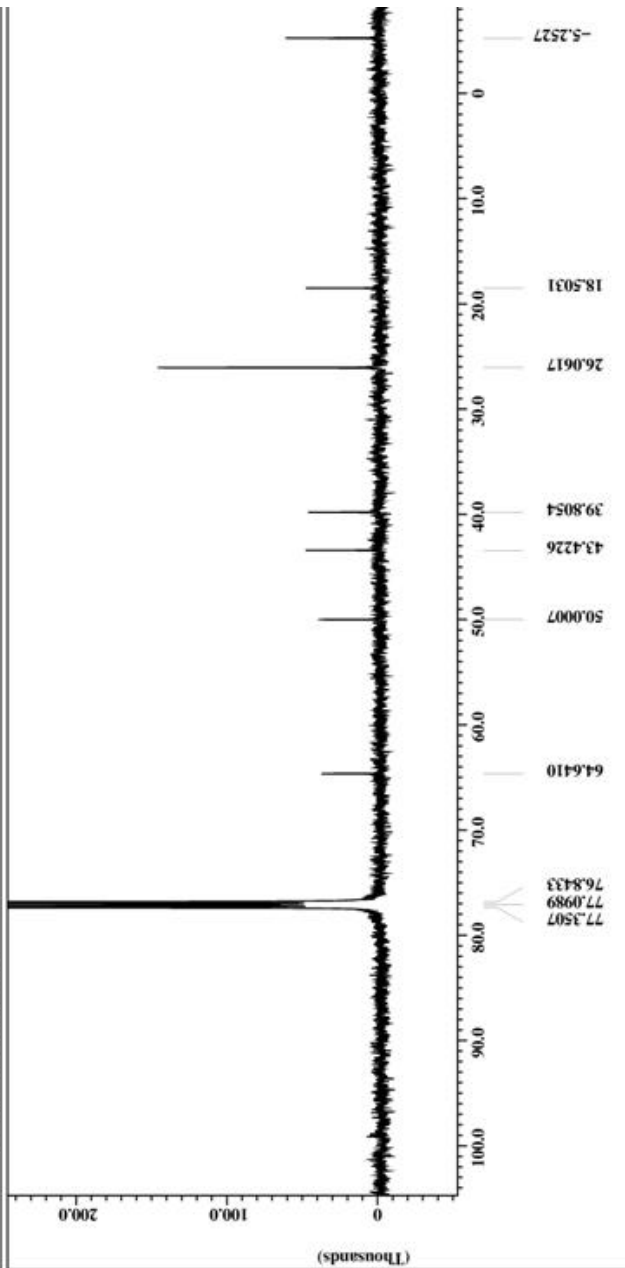
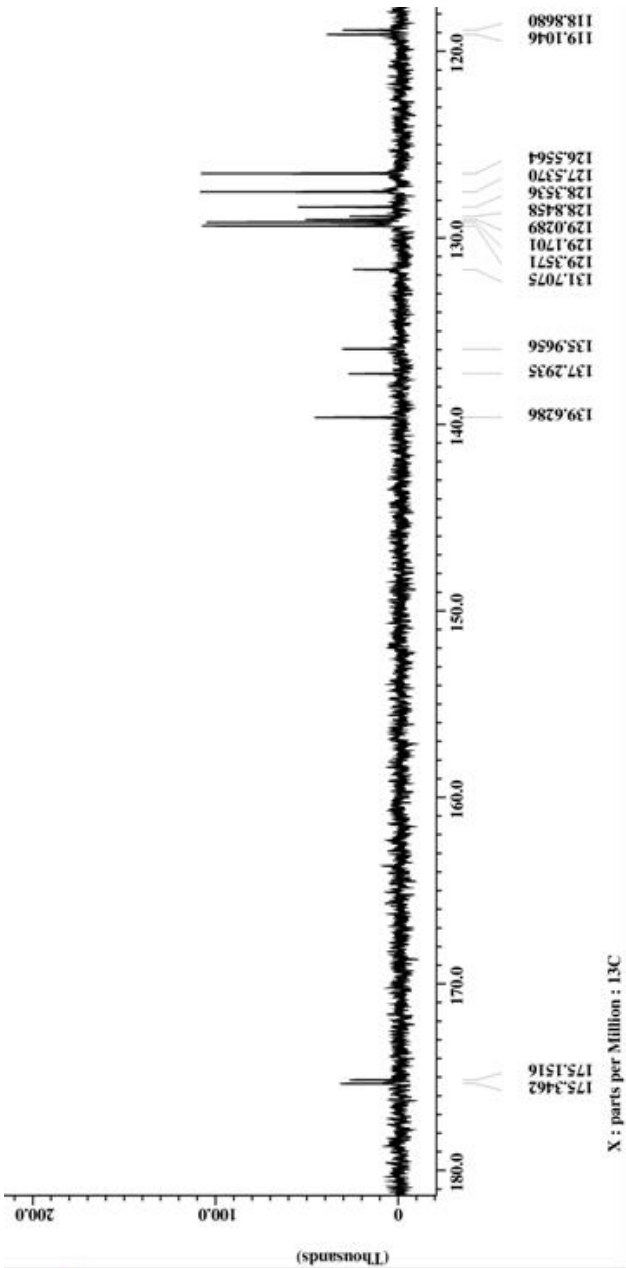
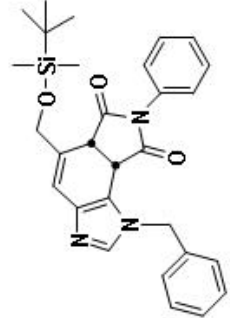
filename = sm_VI_44_pure_i1-2.jd
author = delta
experiment = single_pulse_dec
sample_id = S#698993
solvent = CHLOROFORM-D
reaction_time = 29-NOV-2009 16:28:27
revision_time = 30-NOV-2009 00:33:02
current_time = 28-MAR-2010 23:18:45

comment = single pulse decouple
          = ID COMPLEX
          = 65536
          = 13C
          = [ppm]
          = X
          = Eclipse+ 500
          = DELTA_NMR

field_strength = 11.7465928[T] (500[MH
acq_duration = 2.0840448[s]
domain = 13C
freq = 125.75710665[MHz]
offset = 100[ppm]
points = 65536
prescans = 4
resolution = 0.47983613[Hz]
sweep = 31.44654088[kHz]
tr_domain = 1H
tr_freq = 500.12734003[MHz]
tr_offset = 5[ppm]
tipped = TRUE
bd_return = 10
cans = 4000
total_scans = 4000

_90_width = 14.2[us]
acq_time = 2.0840448[s]
angle = 30[deg]
pulse = 4.73333333[us]
p1_width = 1[s]
p2_time = 1[s]
p3_time = 3[us]
base_preset = 28
scvr_gain = 2[s]
relaxation_delay = 2[s]
emp_get = 28.6[dC]
nblank_time = 2[us]

```



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