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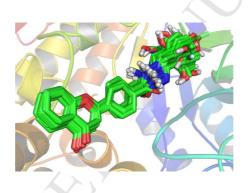


Graphical Abstract

Synthesis of Novel Flavone Hydrazones: *In-Vitro* Evaluation of α-Glucosidase Inhibition, QSAR Analysis and Docking Studies

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A new series of potent novel flavone derivatives 5-34 as α -glucosidase inhibitors was identified.

Synthesis of Novel Flavone Hydrazones: *In-Vitro* Evaluation of α -Glucosidase Inhibition, QSAR Analysis and Docking Studies

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ABSTRACT

Thirty derivatives of flavone hydrazone (5-34) had been synthesized through a five-step reaction and screened for their α -glucosidase inhibition activity. Chalcone 1 was synthesized through aldol condensation then subjected through oxidative cyclization, esterification, and condensation reaction to afford the final products. The result for baker's yeast α -glucosidase (EC 3.2.1.20) inhibition assay showed that all compounds are active with reference to the IC_{50} value of the acarbose (standard drug) except for compound 3. Increase in activity observed for compounds 2 to 34 clearly highlights the importance of flavone, hydrazide and hydrazone linkage in suppressing the activity of α -glucosidase. Additional functional group on Nbenzylidene moiety further enhances the activity significantly. Compound 5 (15.4 \pm 0.22 μ M), a 2,4,6-trihydroxy substituted compound, is the most active compound in the series. Other compounds which were found to be active are those having chlorine, fluorine, and nitro substituents. Compounds with methoxy, pyridine, and methyl substituents are weakly active. Further studies showed that they are not active in inhibiting histone deacetylase activity and do not possess any cytotoxic properties. QSAR model was being developed to further identify the structural requirements contributing to the activity. Using Discovery Studio (DS) 2.5, various 2D descriptors were being used to develop the model. The QSAR model is able to predict the pIC₅₀ and could be used as a prediction tool for compounds having the same skeletal framework. Molecular docking was done for all compounds using homology model of α glucosidase to identify important binding modes responsible for inhibition activity.

Keywords: Flavone, hydrazone, α-glucosidase, 2D-QSAR, docking.

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1. Introduction

Diabetes mellitus is a disease involving metabolic disorder resulting from defects in insulin secretion and action. This result in increase of blood glucose level (hyperglycemia) and causes dysfunction to important organs like blood vessels and nerves [1]. Study shows that the number of diabetes mellitus patients increases each year. The trend displayed that patient with diabetes increases from 153 million to 347 million between the year 1980 to 2008 [2]. Based on World Health Organization (WHO) projection, diabetes is predicted to become the seventh leading cause of death worldwide by 2030. One of the approaches, which can be taken to reduce diabetes is to inhibit enzymes like α -glucosidase by controlling the postprandial glucose levels and suppressing postprandial hyperglycemia [3].

Inhibitors for α -glucosidase function by reversibly inhibit digestive α -glucosidase that retards glucose liberation from complex starch and carbohydrate. This results in delay for glucose to be absorbed into bloodstream and, therefore, decreases plasma glucose level. Inhibition of a-glucosidase has attracted plenty of interest by pharmaceutical industry as a treatment method of diseases like diabetics, viral infections, hepatitis, and cancer [4-6]. Inhibitors for α -Glucosidase are found to show antitumor, antidiabetic, antiviral, and immunoregulatory activities [7-9]. Research on α -glucosidase inhibitors such as castanospermine, *N*-butyl-deoxynojirimycin and deoxynojirimycin showed that they are potent against HIV replication [10].

Currently, glycosidic based α -glucosidase inhibitors such as miglitol [11], voglibose [12], acarbose [13] and nojirimycin [14] are used to control glucose level in blood of diabetic patients. Despite being potent, these inhibitors have some limitations such as diarrhea, abdominal distension, meteorism and flatulence [15]. Acarbose, deoxynojirimycin, miglitol, and voglibose are among the key known candidates which are used extensively to inhibit α -glucosidase activity. Due to side effects and absorptivity problems associated with these inhibitors [16, 17], new potent α -glucosidase inhibitors are highly desired. Hence, a lot of efforts have been put in to design and develop non-glycosidic based inhibitors for safer [18] and effective α -glucosidase inhibitors [19].

 α -Glucosidase (EC3.2.1.20) is an enzyme commonly found in small intestine. α -Glucosidase hydrolyzes carbohydrates and produces α -d-glucose during food digestion. Carbohydrates and α -d-glucose that has been produced are being absorbed into blood stream, thus increases postprandial blood glucose level and leads to diabetes. Therefore, α -glucosidase inhibitors are of significant importance to control diabetes since they could reduce carbohydrate digestion

and monosaccharide absorption [20, 21]. Glucose concentration in blood is extremely critical for diabetes mellitus management and it must be between an acceptable ranges (70-100 mg/dl) [22]. In addition to preventing diabetes, blood glucose level which is controlled may avoid hyperlipoproteinemia, obesity, and hyperlipidemia [23].

 α -Glucosidase also enables monosaccharide removal from viral glycoproteins, thus, its inhibitors could alter cell-to-cell signaling, virus recognition by the cell and could be used in the treatment of viral diseases, cancer, and immune-regulations [24-28].

Flavone is one of the class of flavonoids that is present in fruits and vegetables. Flavones are being consumed in daily diet and they improve health without giving any major side effect [29]. In order to explore diverse roles of flavones, investigating various methods for their synthesis and structural modification of flavones ring has now become important goals of several research groups. Thus, naturally obtained flavone moiety having a variety of biological activities can be taken as lead compound for the synthesis of semi- and purely synthetic flavone derivatives with different functional groups at different positions of flavone skeleton.

Flavone scaffold is an important structure found in many pharmaceutically active compounds. They are structurally diverse and possess a variety of biological activity. This reason has increased the interest of medicinal chemists to further study flavones as lead molecules to treat various diseases. Recently, researchers had focused a lot on flavonoids bioactivities like free radicals scavenging ability and protection against peroxidation of lipid [30, 31]. Researchers also showed interest in some flavonoids due to their ability to modulate NADPH oxidase activity and endothelial nitric oxide metabolism [32-38]. Various research indicate that flavonoids could reduce hyperglycemia, improve sensitivity and enhance secretion of insulin [39]. It has been established that flavonoids like kaempferol, luteolin, apigenin, chrysin and baicalein have the capability of inhibiting α -glucosidase activity [40, 41], taking into consideration of the importance of ring A, B, and C towards the inhibition of α -glucosidase activity.

Benzohydrazides had been reported to possess various biological activities, which includes antileishmanial [42], antioxidant [43], antiglycation [44], antibacterial [45], and α -glucosidase inhibition [46] activities. In this paper, flavone derivatives possessing hydrazone linkage synthesized and evaluated for their α -glucosidase inhibition activity. We attempted to rationalize the effect of various substituents at different position on the inhibitory effect in terms of how the molecules bind with α -glucosidase protein.

2. Result and discussion

2.1. Chemistry

The title compounds were synthesized through scheme 1 and 2. Flavone hydrazones 5-34 were synthesized through a single step reaction of 4-(4-oxo-4H-chromen-2-yl)benzohydrazide 4 with appropriate aryl aldehydes, in the presence of catalytic amount of glacial acetic acid. Compound 4 which is the key intermediate for the synthesis of target compounds 5-34 was prepared through a four-step sequence. Initially 2'-hydroxyacetophenone undergoes Aldol condensation with 4-formylbenzoic acid using aqueous KOH in ethanol to form (*E*)-4-(3-(2-hydroxyphenyl)-3-oxoprop-1-en-1-yl)benzoic acid 1. Chalcone 1 was cyclized into flavone through oxidative cyclization by using well-known I₂-DMSO mixture as the oxidizing agent. Carboxylic acid of the resulting product 2 was converted into ester 3 before finally reacted it with hydrazine to form compound 4.

Insert Scheme 1 here

Insert Scheme 2 here

The structure of newly synthesized compounds were confirmed by the various analytical techniques, ¹H-NMR, ¹3C-NMR, IR, MS and elemental analysis. All spectral data were in accordance with assumed structures. In the ¹H-NMR spectra of chalcone **1**, olefinic protons H_{α} and H_{β} (Figure-1) appeared as doublets at 7.86 ppm and 8.12 ppm respectively. The formation of chalcone **1** having *trans* conformation was verified using the coupling constant values (15.6 Hz) of both vinyl hydrogens (H_{α} and H_{β}). Oxidation of chalcone **1** leads to the formation of flavone **2** which involves removal of hydrogen from hydroxyl group and H_{β} . This was being confirmed through several changes to the ¹H-NMR spectra. Splitting produced by vinylic protons (H_{α} and H_{β}) were no longer observed. This deduction was further confirmed by the appearance of singlet proton at 7.13 ppm representing H_{α} on the chromone ring.

Insert Figure 1 here

Insert Table 1 here

2.2. *In-vitro* α -glucosidase inhibition

In continuation of our work on enzyme inhibition [47], we had synthesized compounds 1-34 and evaluate them for their α -glucosidase inhibition activity. The assay was carried out using baker's yeast α -glucosidase (EC 3.2.1.20). The result in table 2 showed, with the exception of compound 3, the compounds synthesized in this study are more active than the standard drug. Compounds 1 to 4, which are the intermediate compounds for synthesis of the final compounds 5-34 were also evaluated for their inhibition activity. Comparison between results for intermediates 1 and 2 showed that flavone 2 has better activity as compared to chalcone 1. This result indicates that the chromone moiety is playing a significant role in inhibiting the enzyme activity. Conversion of carboxylic acid to hydrazide 3 enhances the inhibition activity. Converting hydrazide 4 into hydrazones 5-34 further improves the inhibition activity. Comparison between results for intermediate compounds 1-4 with flavone-hydrazone derivatives 5-34 showed a significant improvement in the inhibitory activity, which indicates the importance of hydrazone linkage in inhibiting α -glucosidase activity. Compounds having hydroxyl, chloro and fluoro substituent showed potent activity with IC₅₀ ranging between 15.4 to 86.3 μ M while the other compounds having methyl, methoxy, nitro and bromo substituent are weakly active. Compounds 28, 29 and 30 that are having pyridine ring also showed weak inhibitory activity against α -glucosidase. In order to understand further the effect of structural features and mechanism of action, the compounds were subjected through 2D QSAR and docking studies, which studies enable us to identify key structural features responsible for the inhibition activity as well as rationalizing the best binding modes in the active site of α glucosidase. The compounds were not showing any activity against histone deacetylase inhibition and cytotoxicity activity.

Insert Table 2 here

2.3. 2D QSAR study2.3.1. QSAR model development and validation

Quantitative structural activity relationship (QSAR) for α -glucosidase inhibition activity was performed to determine the factors/descriptors which correlates the bioassay results obtained with flavone derivatives (5-34), and to determine structural features contributing towards the inhibition activity. In this study, the compounds for training set and test set were selected using "cluster analysis" protocol in Discovery Studio 2.5. Cluster selection was done by allowing

cluster to have fixed number of compounds for each cluster. Compounds selection was done using predefined set consisting of items such as AlogP, molecular weight, number of hydrogen donors and acceptors. The training set consisting of 21 compounds from the newly synthesized flavone hydrazones together with their measured pIC_{50} (-logIC₅₀) was used to develop the QSAR model. The molecular properties for the compounds in the training set were calculated using "Calculate Molecular Properties" protocol. The protocol includes calculation for 2D molecular properties, energies of highest occupied and lowest unoccupied molecular orbitals (HOMO and LUMO) for compounds of the training set. In this study, 2D descriptors like AlogP, molecular weight, number of hydrogen donor, and number of rotatable bonds were utilized in developing the model.

Notably, AlogP is a measure of the hydrophobicity of the molecule and calculated in Discovery Studio as the Log of the octanol-water partition coefficient, while Molecular Fractional Polar Surface Area is the ratio of the polar surface area divided by the total surface area of the molecule.

In this study, Multiple Linear Regression (MLR) analysis method was being used to obtain the model. Descriptors were selected based on the results of the intercorrelation matrix between the descriptors, which should intercorrelate more than 0.6. Descriptors that were selected for this study have intercorrelation values lower than 0.5 (table 3a) to prevent model overfitting. These value also showed that these descriptors are independent. Robustness of the established QSAR models were verified using Leave-one-out (LOO) internal validation or cross-validation (q^2), where r^2 (squared correlation coefficient value) equals 0.848, while the r^2 (pred), which is equivalent to q^2 from a leave-one-out cross validation, is 0.705 (table 3b).

Insert Table 3 here

In addition, validation was done by measuring the residuals between the experimental and the predicted activity of the training set (Table 4). A correlation plot representing the correlation between predicted pIC_{50} against experimental pIC_{50} of test set (Fig. 2b) showed that the plot has a regression correlation coefficient (r²) of 0.783, which is considered quite good. Despite showing good correlation, the QSAR model can be further improved to be used applied for prediction of more effective hits having the same skeletal framework.

Insert Figure 2 here

Insert Table 4 here

2.3.2. 2D QSAR model analysis

Equation 1 represents the model obtained using Multiple Linear Regression and it is graphically visualized using a scatter plot of predicted pIC₅₀ against experimental pIC₅₀ values for the training set and test set compounds as shown in Figure 2. Equation 1 describes that α glucosidase inhibition activity of flavone derivatives 5-34 are affected by four descriptors, which are AlogP, molecular weight, number of hydrogen bond donor, and number of rotatable bonds. Comparison between descriptors in the equation showed that AlogP, number of hydrogen bond donor, and number of rotatable bonds are among those which contributed significantly towards the high correlation as compared to molecular weight. Despite showing some effect on the pIC_{50} , small molecular variation that only involves substitution on the extended benzylidene ring has affected the molecular weight descriptor. Having the lowest constant value, the equation suggest that small changes in molecular weight could slightly affect the IC₅₀ value as compared to the other descriptors. Based on the equation, it was observed that increasing the number of rotatable bonds descriptor, which has the highest constant value, in the molecule could significantly increase the pIC_{50} value and, therefore, gives a lower IC₅₀. However, other descriptors in the equation, which are AlogP and molecular weight, suggest that increasing the lipophilicity and molecular weight of the compounds will only reduce the pIC_{50} value and gives a higher IC_{50} value. This observation supports the actual activity which showed that adding rotatable and non-polar substituents methyl and methoxy substituent to compounds could increase the lipophilicity and molecular weight of the compounds and will significantly reduce the pIC₅₀ value to give a higher IC₅₀ value. On the other hand, the equation also suggest that increasing the number of hydrogen bond donor could increases the activity of the compounds by lowering the IC₅₀ value, which also supports the general observation results from *in-vitro* activity, which show that increasing hydroxyl group in the structure could enhance the activity by giving lower IC₅₀ value. Docking results further validate this QSAR model by taking into accounts the effect of all descriptors used in this model.

Equation 1. Equation representing the QSAR model;

 $pIC_{50} = -16.827 - 0.509 (AlogP) - 5.819 \times 10^{-2} (Molecular_Weight) + 0.213 (Num_H_Donors) + 1.334 (Num_RotatableBonds)$

2.4. Docking study

Molecular docking was performed on all flavones derivatives (5-34) to identify possible binding mode which explains the reason for their potency. Since crystal structure for α glucosidase of *S. cerevisiae* is still not available, docking study was conducted using a homology model for α -glucosidase. Preliminary results on sequence analysis of α -glucosidase from *S. cerevisiae* showed that the most suitable template for homology modelling is isomaltase (EC 3.2.1.10, oligo-1,6-glucosidase, MALX3)(PDB ID: 3A4A) from baker's yeast which shares 71% identity and 84% similarity with the target enzyme, α -glucosidase of *S. cerevisiae*. Sequence analysis on the homology model's sequence showed high sequence homology with the target enzyme. The active site contains important and highly conserved amino acids amino acids (Fig. 3) [48]. The final structure of α -glucosidase generated from homology modeling was evaluated using PROCHECK. The Ramachandran plot obtained from PROCHECK showed that 87.5% of residues of the final 3D structure lied in most favored regions (Fig. 4).

Insert Figure 3 here Insert Figure 4 here

Prior to docking and analysis of the binding mode of the most active compound **5**, the docking method was validated through control docking of native inhibitor. Acarbose was docked into α -glucosidase from Sugar beet (PDB code: 3W37) and compared by superimposing with the native ligand in the protein (Figure 5a). The rmsd value between docked and actual pose of acarbose was found to be 0.65Å. Even though α -glucosidase from Sugar beet share relatively low homology with baker's yeast α -glucosidase (16% identity), the active site is highly conserved and main interactions of the ligand remain the same [48]. Another control docking was done on the target protein, isomaltase from Baker's yeast, using α -D-glucopyranose which is the native ligand located in the active site (Fig. 5b). The rmsd value for docked pose and native ligand was found to be 0.93Å.

Insert Figure 5 here

All flavones were docked and aligned in the binding pocket. Based on docking results of the aligned molecules (Fig. 6), the benzylidene moiety of the compounds are oriented towards the core of the binding pocket. This enables the substituents on the benzylidene moiety to interact

closely with important residues in the active site. Introduction of hydrophilic substituents with H-bond donating properties like hydroxyl groups on the benzylidene moiety showed the most interaction with active residues which led to extra interactions and increase of the activity. Benzohydrazone linkage provides torsional degree of freedom (rotatability) that substantially decreases the entropic penalty for the formation of enzyme-inhibitor complex. This feature validates one of the descriptors used in the QSAR model, which suggest that increase in number of rotatable bonds could enhance the inhibition activity. Four hydrogen bonds are established between the phenolic and carbonyl oxygen of compound **5** and the side chains of His239, His245, His279, and Thr275. Benzohydrazone moiety resides in close proximity to the catalytic residues including Asp214, Glu276, His348, and Asp349, indicating that it can serve as surrogate for the terminal glucose in the substrate.

Insert Figure 6 here

The main flavone scaffold of all derivatives (**5-34**) are better aligned when compared to the benzylidene moiety. The flavone scaffold which is positioned in a hydrophobic pocket allows formation of highly stable complexes. Several observations were made based on the interaction of flavone moiety with the enzyme. The main interaction which played the most significant role in stabilizing the enzyme-inhibitor complex is the interaction of benzopyrone ring of flavone moiety with Phe311 and Lys155 through a Pi-Pi stacking and Pi-cation interaction, respectively. Ring B of flavone forms additional Pi-Pi stacking with Phe311.

Two hydrogen bonds were observed for the most active compound, **5**, in the series (Fig. 7a). Besides forming pi-cation interaction, NH fragment of Lys155 also forms hydrogen bonding with oxygen of the ether linkage on ring C which further stabilizes the enzyme inhibitor complex. On the other hand, hydroxyl forms hydrogen bonding with catalytic residue His279, which is believed to have some reducing effect in the enzyme's activity. In figure 6b, nitrogen of benzohydrazone backbone forms a Pi-cation interaction with another catalytic residue His239.

Insert Figure 7 here

Attempt to validate the QSAR model using docking results showed a high degree of correlation in terms of substitutent's lipophilicity, ability to form hydrogen bonding, and rotatability.

Docking studies showed that increase in number of hydrogen bond allows more interaction to form between between active residues and the ligand. This observation satisfy the number of hydrogen bond donor descriptor, which was observed to reduce the IC₅₀ value when number of hydroxyl substituent was increased. In a different observation, it is also clearly visualized that increasing lipophilicity properties of the substituents by replacing hydroxyl with methoxy substituents causes shifting of benzylidene ring and prevents optimal interaction from taking place. Inhibition activity also decreases due to steric hindrance imposed by methoxy and inability to function as a hydrogen bond donor. The effect from steric hindrance by methoxy is visualized in figure 8c and 8d. Compounds 14 and 15, which contains both hydroxyl and methoxy substituents, displayed better activity as compared to compounds 30 and 31 which are single substituted with methoxy substituent. For compounds 14 and 15, the hydroxyl groups for both compounds are oriented towards the hydrophilic pocket (Fig. 8a and 8b). It was observed that substituents for both compounds 14 and 15 are not capable of interacting with any catalytic residues due to steric hindrance imposed by methoxy substituent. For compound 14, methoxy group forms hydrogen bonding with Thr217, while for compound 15, hydroxyl group interacts with Asn241.

Insert Figure 8 here

Eventhough the activity for compound 14 and 15 are considered quite good, hydrogen bond between methoxy of compound 14 and hydroxyl of compound 15 were not enough to increase the activity in a significant way. Rather than adding more hydroxyl substituents, it seems more important to place hydroxyl substituents on *N*-benzylidene ring at the correct position for the compounds to be able to inhibit α -glucosidase activity.

3. Conclusion

All flavone-hydrazones synthesized in this study are active in inhibiting α -glucosidase. The result strongly suggest that chromone moiety and hydrazone linkage played important role in suppressing the activity of α -glucosidase activity as the activity of all the compounds improve significantly as compared to the starting material **4**. 2D QSAR model established showed correlation and contribution of various structural features in the activity. Descriptors such as AlogP, number of H bond acceptor/donor, molecular weight, and number of rotatable bonds are statistically important in this study. AlogP and number of rotatable bonds play an important

role as compared to other descriptors like H-bond donors and molecular weight. QSAR equation suggest that by increasing number of hydrogen bond donor, molecular weight, and number of rotatable bonds, the inhibition activity could be improved. The validated QSAR model is suitable to predict α -glucosidase inhibition activity for compounds having similar skeletal structure in the future. Docking studies showed that benzopyrone ring of flavone moiety that is located in hydrophobic pocket stabilizes the enzyme-inhibitor complex by forming Pi-Pi stacking with Phe311 and Pi-cation interaction with Lys155. N-benzylidene moiety resides in close proximity to catalytic residues like Asp214, Glu276, His348, and Asp349 indicates that it can serve as a surrogate for terminal glucose in α -glucosidase. On the other hand, benzohydrazone linkage provides torsional degree of freedom in terms of its rotatablility, which had substantially decreased the entropic penalty for the formation of enzyme-inhibitor complex. Phenolic and carbonyl oxygens of compound 5, which is the most active derivative, forms 4 hydrogen bonds with side chains of His239, His245, His279, and Thr275. Therefore, the compounds are most likely to be capable of inhibiting the catalytic action of α -glucosidase by a tight binding in the active site through the multiple hydrogen bond and hydrophobic interactions in a cooperative fashion. In addition to a-glucosidase inhibition activity, the compounds were also tested for their cytotoxicity and ability to inhibit histone deacetylase. It was found that they do not show any activity for histone deacetylase inhibition and cytotoxicity activity.

4. Experimental

4.1. Chemistry

Melting points were determined using Sinosource SGW X-4 melting point apparatus with microscope (Guangzhou, China). IR spectra obtained using PerkinElmer Spectrum 100 FTIR Spectrometer (Waltham, MA, USA) equipped with a diamond crystal Attenuated Total Reflectance (ATR) accessory by PIKE Technologies (Madison, WI, USA). NMR spectroscopy was obtained using Bruker Ultra Shield FT NMR 500 MHz and Avance III 600 Ascend spectrometer (Wissembourg, France). EI-MS spectroscopic analysis had been obtained using Finnigan-MAT-311-A instrument (Bremen, Germany). Thin layer chromatography (TLC) was performed using precoated silica gel plates (Merck, Kieselgel 60 F-254, 0.20 mm).

4.2. Synthesis of (*E*)-4-(3-(2-hydroxyphenyl)-3-oxoprop-1-en-1-yl)benzoic acid (1)

In a 250 ml flask, 2'-hydroxyacetophenone (30 mmole) was being mixed with 4-formylbenzoic acid (30 mmole). The mixture was dissolved in 100 ml of 15% (w/v) sodium hydroxide in

ethanol and stirred at room temperature. Reaction progress was monitored using TLC and upon completion, diluted sulphuric acid was added to allow precipitation. The precipitate was filtered and crystallized in ethanol to afford pure product. Yellow solid; yield 78 %; m.p. 250-252 °C; IR (ATR) cm⁻¹: 2990, 2831, 1678, 1643, 1568, 1291; ¹H-NMR (600 MHz, DMSO-*d6*): δ 7.02-7.05 (m, 2H), 7.58 (t, *J* = 7.2 Hz, 1H), 7.86 (d, *J* = 15.6 Hz, 1H), 8.00-8.05 (m, 4H), 8.13 (d, *J* = 15.6 Hz, 1H), 8.24 (d, *J* = 7.5 Hz, 1H), 11.15 (s, 1H), 12.36 (s, 1H); ¹³C-NMR (125 MHz, DMSO-*d6*): δ 117.7, 119.5, 120.9, 122.3, 126.3, 128.6, 130.6, 130.6, 133.5, 135.5, 137.5, 143.3, 161.4, 168.5, 193.5; Anal. Calcd for C₁₆H₁₂O₄: C = 71.64, H = 4.51, Found: C = 71.63, H = 4.52; EI MS *m*/*z* (% rel. abund.): 268.07 (M⁺, 68.2 %).

4.3. Synthesis of 4-(4-oxo-4H-chromen-2-yl)benzoic acid (2)

Compound **1** (22.4 mmole) was mixed with iodine (0.23 mmole). The mixture was dissolved in 50 mL of DMSO and refluxed at 170 °C. After 3 hours, sodium thiosulfate was added to the reaction mixture followed by excess amount of water to allow precipitation. The product was rinsed and allowed to dry at room temperature to afford pure product. Light yellow solid; yield 90%; m.p. 298-300 °C; IR (ATR) cm⁻¹: 3420, 3081, 2885, 1707, 1623, 1588, 1567, 1415, 1242; ¹H-NMR (500 MHz, DMSO-*d*6): δ 7.13 (s, 1H), 7.51(t, *J* = 7.5 Hz, 1H), 7.80 (d, *J* = 8.5 Hz, 1H), 7.84 (t, *J* = 7.0 Hz, 1H), 8.05 (d, *J* = 8.0 Hz, 1H), 8.09 (d, *J* = 8.5 Hz, 2H), 8.22 (d, *J* = 8.5 Hz, 2H), 12.21 (s, 1H); ¹³C-NMR (125 MHz, DMSO-*d*6): δ 105.9, 118.4, 124.6, 124.6, 124.9, 125.7, 126.0, 128.9, 130.3, 130.3, 132.6, 134.4, 156.8, 163.7, 168.8, 178.0; Anal. Calcd for C₁₆H₁₀O₄: C = 72.18, H = 3.79, Found: C = 72.19, H = 3.81; EI MS *m/z* (% rel. abund.): 266.06 (M⁺, 73.6%)

4.4. Synthesis of methyl 4-(4-oxo-4H-chromen-2-yl)benzoate (3)

Compound **2** (18.8 mmole) was mixed with 1 ml of concentrated sulphuric acid and dissolved in 100 ml methanol. The reaction was refluxed for more than 16 hours and monitored using TLC. Upon completion, methanol was being removed using rotavap. Residue was collected and rinsed using excessive amount of water and crystallized in methanol to afford pure product. White solid; yield 95%; m.p. 268-270 °C; IR (ATR) cm⁻¹: 3325, 3073, 2873, 1742, 1707, 1648, 1556, 1412, 1262; ¹H-NMR (500 MHz, DMSO-*d*6): δ 3.91 (s, 3H), 7.16 (s, 1H), 7.52(t, *J* = 7.0 Hz, 1H), 7.82 (d, *J* = 8.0 Hz, 1H), 7.86 (t, *J* = 7.0 Hz, 1H), 8.07(d, *J* = 8.0 Hz, 1H), 8.13(d, *J* = 8.5 Hz, 2H), 8.27(d, *J* = 8.5 Hz, 2H); ¹³C-NMR (125 MHz, DMSO-*d*6): δ 55.6, 105.91, 118.2, 124.7, 125.1, 125.1, 125.7, 126.5, 128.2, 128.2, 131.0, 131.5, 133.2, 156.8, 163.7, 167.4, 178.0;

Anal. Calcd for $C_{17}H_{12}O_4$: C = 72.85, H = 4.32, Found: C = 72.87, H = 4.35; EI MS *m*/*z* (% rel. abund.): 280.07 (M⁺, 76.3 %),

4.5. Synthesis of 4-(4-oxo-4H-chromen-2-yl)benzohydrazide (4)

Compound **3** (17.7 mmole) was dissolved in 100 ml methanol containing 25 ml hydrazine hydrate. The reaction mixture was refluxed for at least 6 hours. Reaction progress was monitored using TLC. Methanol was removed using rotavap and product was rinsed with plenty of water. The product was rinsed and allowed to dry at room temperature to afford pure product. White solid; yield 88%; m.p. 197-199 °C; IR (ATR) cm⁻¹: 3327, 3139, 2997, 1630, 1558, 1469, 1341, 1251; ¹H-NMR (500 MHz, DMSO-*d6*): δ 4.51 (s, 2H), 6.91 (t, J = 7.5 Hz, 1H), 6.96 (d, J = 7.5 Hz, 1H), 7.19 (t, J = 7.0 Hz, 1H), 7.33 (s, 1H), 7.74 (d, J = 7.5 Hz, 1H), 7.918 (s, 4H), 9.82 (s, 1H); ¹³C-NMR (125 MHz, DMSO-*d6*): δ 105.8, 118.3, 124.3, 124.8, 125.2, 125.6, 125.6, 126.3, 126.3, 133.8, 137.3, 137., 157.2, 163.3, 167.4, 178.2; Anal. Calcd for C₁₆H₁₂N₂O₃: C = 68.56, H = 4.32, N = 9.99, Found: C = 68.57, H = 4.34, N = 9.97; EI MS *m/z* (% rel. abund.): 280.17 (M⁺, 58.2%)

4.6. General procedure for synthesis of flavone hydrazones (5-34)

Compound 4 (0.5 mmole) and substituted benzaldehydes (0.5 mmole) were being mixed with 1 ml of glacial acetic acid in a 50 mL round bottom flask. The mixture was being dissolved in 20 mL of methanol and refluxed. Progress monitored using TLC and upon completion, solvent was being removed using rotary evaporator. Product was collected and rinsed with diethyl ether to afford pure product.

4.6.1. (*E*)-**4**-(**4**-oxo-**4**H-chromen-2-yl)-*N*'-(**2**,**4**,**6**-trihydroxybenzylidene)benzohydrazide (**5**) Light yellow solid; Yield 76 %; m.p. 283-285 °C; IR (ATR) cm⁻¹: 3353, 3139, 2994, 1657, 1611, 1556, 1519, 1470, 1372, 1269, 1186, 760; ¹H-NMR (500 MHz, DMSO-*d*6): δ 5.86 (s, 2H), 6.93 (t, *J* = 7.5 Hz, 1H), 6.93 (d, *J* = 8.0 Hz, 1H), 6.93 (t, *J* = 7.0 Hz, 1H), 7.38 (s, 1H), 7.75 (s, 1H), 7.99-8.04 (m, 4H), 8.83 (s, 1H), 9.85 (s, 1H), 11.13 (s, 2H), 11.94 (s, 1H); ¹³C-NMR (125 MHz, DMSO-*d*6): δ 95.8, 95.8, 105.5, 106.8, 118.2, 125.1, 125.4, 125.4, 125.6, 126.2, 126.8, 126.8, 133.7, 137.2, 138.8, 144.6, 156.2, 161.7, 161.7, 163.2, 163.5, 164.3, 178.1; Anal. Calcd for C₂₃H₁₆N₂O₆: C = 66.34, H = 3.87, N = 6.73, Found: C = 66.33, H = 3.88, N = 6.71; EI MS *m*/*z* (% rel. abund.): 416.20 (M⁺, 72.5%)

4.6.2. (*E*)-4-(4-oxo-4H-chromen-2-yl)-*N*'-(2,3-dihydroxybenzylidene)benzohydrazide (6)

White solid; Yield 81 %; m.p. 288-290 °C; IR (ATR) cm⁻¹: 3373, 3159, 2835, 1609, 1539, 1469, 1357, 1257, 1186, 754; ¹H-NMR (500 MHz, DMSO-*d6*): δ 6.75 (t, *J* = 7.5 Hz, 1H), 6.87 (d, *J* = 7.0 Hz, 1H), 6.92 (t, *J* = 7.5 Hz, 1H), 6.98 (d, *J* = 7.5 Hz, 2H), 7.20 (t, *J* = 7.0 Hz, 1H), 7.76 (s, 1H), 8.01-8.04 (m, 5H), 8.63 (s, 1H), 9.25 (s, 1H), 11.14 (s, 1H), 12.19 (s, 1H); ¹³C-NMR (125 MHz, DMSO-*d6*): δ 105.9, 118.2, 119.7, 119.9, 121.3, 122.0, 125.0, 125.2, 125.4, 125.4, 126.2, 126.9, 126.9, 133.9, 137.1, 138.7, 144.4, 146.4, 149.3, 155.9, 163.4, 164.5, 178.3; Anal. Calcd for C₂₃H₁₆N₂O₅: C = 69.00, H = 4.03, N = 7.00, Found: C = 69.01, H = 4.05, N = 6.98; EI MS *m*/*z* (% rel. abund.): 400.12 (M⁺, 64.7%)

4.6.3. (E)-4-(4-oxo-4H-chromen-2-yl)-N'-(2,4-dihydroxybenzylidene)benzohydrazide (7)

Light yellow solid; Yield 88 %; m.p. 294-296 °C; IR (ATR) cm⁻¹: IR (ATR) cm⁻¹: 3354, 3212, 3021, 1626, 1606, 1495, 1466, 1259, 1166, 749; ¹H-NMR (500 MHz, DMSO-*d6*): δ 6.34 (s, 1H), 6.37 (d, *J* = 8.0 Hz, 1H), 6.91 (t, *J* = 7.5 Hz, 1H), 6.97 (d, *J* = 8.0 Hz, 1H), 7.20 (t, *J* = 7.5 Hz, 1H), 7.32 (d, *J* = 8.5 Hz, 1H), 7.36 (s, 1H), 7.75 (d, *J* = 7.0 Hz, 1H), 8.00-8.03 (m, 4H), 8.53 (s, 1H), 9.23 (s, 1H), 10.14 (s, 1H), 11.53 (s, 1H); ¹³C-NMR (125 MHz, DMSO-*d6*): δ 103.0, 105.9, 109.2, 113.1, 118.2, 125.1, 125.4, 125.4, 125.4, 126.2, 126.9, 126.9, 130.8, 133.9, 137.1, 138.7, 151.1, 156.2, 160.5, 160.5, 163.4, 164.3, 178.1; Anal. Calcd for C₂₃H₁₆N₂O₅: C = 69.00; H = 4.03; N = 7.00; Found: C = 69.02; H = 4.04; N = 7.01; EI MS *m/z* (% rel. abund.): 400.25 (M⁺, 69.2%)

4.6.4. (E)-4-(4-oxo-4H-chromen-2-yl)-N'-(2,5-dihydroxybenzylidene)benzohydrazide (8)

Light orange solid; Yield 74%; m.p. > 320 °C; IR (ATR) cm⁻¹: IR (ATR) cm⁻¹: 3244, 3164, 3029, 1647, 1580, 1545, 1487, 1272, 1256, 1141, 747; ¹H-NMR (500 MHz, DMSO-*d6*): δ 6.73-6.78 (m, 2H), 6.91 (t, *J* = 7.0 Hz, 1H), 6.97-6.99 (m, 2H), 7.20 (t, *J*= 8.5 Hz, 1H), 7.37 (s, 1H), 7.75 (d, *J*= 7.5 Hz, 1H), 8.00-8.05 (m, 4H), 8.60 (s, 1H), 9.04 (s, 1H), 10.48 (s, 1H), 12.04 (s, 1H); ¹³C-NMR (125 MHz, DMSO-*d6*): δ 105.9, 116.4, 118.1, 118.2, 121.6, 122.0, 125.0, 125.2, 125.4, 125.4, 126.2, 126.9, 126.9, 133.9, 137.1, 138.7, 149.6, 151.2, 152.7, 156.2, 163.4, 164.3, 178.1; Anal. Calcd for C₂₃H₁₆N₂O₅: C = 69.00, H = 4.03, N = 7.00; Found: C = 68.69, H = 4.04, N = 6.99; EI MS *m/z* (% rel. abund.): 400.15 (M⁺, 71.5%)

4.6.5. (*E*)-4-(4-oxo-4H-chromen-2-yl)-*N*'-(3,4-dihydroxybenzylidene)benzohydrazide (9)

Yellowish white solid; Yield 92 %; m.p. 293-295 °C; IR (ATR) cm⁻¹: IR (ATR) cm⁻¹: 3381, 3062, 2898, 1613, 1590, 1558, 1470, 1445, 1277, 1180, 752; ¹H-NMR (500 MHz, DMSO-*d6*): δ 6.79 (d, J = 8.0 Hz, 1H), 6.91-6.98 (m, 4H), 7.19 (t, J = 7.5 Hz, 1H), 7.35 (s, 1H), 7.75 (d, J = 7.0 Hz, 1H), 7.76 (s, 1H), 7.99 (m, 4H), 8.29 (s, 1H), 10.23 (s, 1H), 11.59 (s, 1H); ¹³C-NMR (125 MHz, DMSO-*d6*): δ 105.96, 115.90, 116.62, 118.24, 121.06, 125.0, 125.2, 125.44, 125.4, 126.9, 127.9, 127.9, 133.9, 137.1, 138.7, 145.5, 148.1, 148.6, 149.6, 156.2, 163.4, 164.3, 178.9; Anal. Calcd for C₂₃H₁₆N₂O₅: C = 69.00, H = 4.03, N = 7.00, Found: C = 69.01, H = 4.02, N = 6.99; EI MS *m/z* (% rel. abund.): 400.27 (M⁺, 76.8%)

4.6.6. (E)-4-(4-oxo-4H-chromen-2-yl)-N'-(2-hydroxybenzylidene)benzohydrazide (10)

White solid; Yield 76 %; m.p. 305-306 °C; IR (ATR) cm⁻¹: 3329, 3210, 3021, 1636, 1606, 1537, 1488, 1348, 1259, 1183, 1115, 746; ¹H-NMR (500 MHz, DMSO-*d*6): δ 6.91-6.99 (m, 4H), 7.20 (t, *J* = 7.5 Hz, 1H), 7.31 (t, *J*= 8.0 Hz, 1H), 7.37 (s, 1H), 7.56 (d, *J*= 7.5 Hz, 1H), 7.75 (d, *J*= 7.0 Hz, 1H), 8.01-8.06 (m, 4H), 8.67 (s, 1H), 10.28 (s, 1H), 11.32 (s, 1H); ¹³C-NMR (125 MHz, DMSO-*d*6): δ 105.9, 117.3, 118.2, 120.3, 121.1, 125.0, 125.2, 125.4, 125.4, 126.2, 126.9, 126.9, 128.5, 129.7, 133.9, 137.1, 138.7, 151.0, 156.2, 158.3, 163.6, 164.6, 178.1; Anal. Calcd for C₂₃H₁₆N₂O₄: C = 71.87, H = 4.20, N = 7.29, Found: C = 71.88, H = 4.22, N = 7.30; EI MS *m*/*z* (% rel. abund.): 384.32 (M⁺, 67.4%)

4.6.7. (E)-4-(4-oxo-4H-chromen-2-yl)-N'-(3-hydroxybenzylidene)benzohydrazide (11)

Light red solid; Yield 83 %; m.p. 299-301 °C; IR (ATR) cm-1: 3385, 3212, 3021, 1626, 1606, 1495, 1466, 1259, 1166, 756; ¹H-NMR (500 MHz, DMSO-*d6*): δ 6.84 (d, J = 8.0 Hz, 1H), 6.92 (t, J = 7.5 Hz, 1H), 6.97 (d, J = 8.0 Hz, 1H), 7.12 (d, J = 7.5 Hz, 1H), 7.20-7.23 (m, 2H), 7.26 (t, J = 7.5 Hz, 1H), 7.75 (s, 1H), 8.01 (m, 5H), 8.39 (s, 1H), 9.67 (s, 1H), 11.85 (s, 1H); ¹³C-NMR (125 MHz, DMSO-*d6*): δ 105.95, 117.32, 118.20, 119.35, 120.08, 125.03, 125.21, 125.4, 125.4, 126.2, 126.9, 126.9, 130.6, 133.9, 136.8, 137.2, 138.7, 148.7, 156.2, 156.7, 163.4, 164.3, 178.1; Anal. Calcd for C₂₃H₁₆N₂O₄: C = 71.87, H = 4.20, N = 7.29; Found: C = 71.85, H = 4.22, N = 7.28; EI MS *m*/*z* (% rel. abund.): 384.13 (M⁺, 69.2%);

4.6.8. (*E*)-4-(4-oxo-4H-chromen-2-yl)-*N*'-(4-hydroxybenzylidene)benzohydrazide (12)

Brown solid; Yield 85 %; m.p. 292-294 °C; IR (ATR) cm⁻¹: 3429, 3254, 3062, 1607, 1587, 1507, 1467, 1275, 1236, 1180, 755; ¹H-NMR (500 MHz, DMSO-*d*6): δ 6.85 (d, *J* = 8.0 Hz,

1H), 6.91 (t, J = 7.5 Hz, 1H), 6.97 (d, J = 8.0 Hz, 1H), 7.20 (t, J = 7.5 Hz, 1H), 7.40 (s, 1H), 7.58 (d, J = 8.5 Hz, 2H), 7.75 (s, 1H), 8.00 (m, 5H), 8.38 (s, 1H), 9.98 (s, 1H), 11.70 (s, 1H); ¹³C-NMR (125 MHz, DMSO-*d6*): δ 105.9, 115.6, 115.6, 118.2, 124.3, 124.8, 125.4, 125.4, 125.9, 126.3, 126.8, 126.8, 129.7, 133.8, 137.1, 137.1, 138.6, 149.3, 156.3, 158.4, 163.4, 164.2, 178.0; Anal. Calcd for C₂₃H₁₆N₂O₄: C = 71.87, H = 4.20, N = 7.29, Found: C = 71.88, H = 4.21, N = 7.27; EI MS *m*/*z* (% rel. abund.): 384.18 (M⁺, 61.5%)

4.6.9. (E)-N'-(2-hydroxy-4-methoxybenzylidene)-4-(4-oxo-4H-chromen-2-

yl)benzohydrazide (13)

Dark yellow solid; Yield 94 %; m.p. 311-312 °C; IR (ATR) cm⁻¹: 3347, 3182, 3075, 2838, 1633, 1603, 1489, 1465, 1260, 1184, 750; ¹H-NMR (500 MHz, DMSO-*d6*): δ 3.79 (s, 3H), 6.52 (s, 1H), 6.54 (d, *J* = 8.5 Hz, 1H), 6.92 (t, *J* = 7.5 Hz, 1H), 6.97 (d, *J*= 8.0 Hz, 1H), 7.20 (t, *J* = 8.0 Hz, 1H), 7.44 (d, *J* = 8.5, 1H), 7.75 (s, 1H), 8.02 (m, 5H), 8.58 (s, 1H), 11.65 (s, 1H), 12.04 (s, 1H); ¹³C-NMR (125 MHz, DMSO-*d6*): δ 56.3, 105.9, 107.2, 113.9, 118.2, 124.9, 125.2, 125.4, 125.4, 126.2, 126.9, 126.9, 130.8, 133.9, 137.1, 138.7, 151.0, 156.0, 156.2, 161.2, 162.0, 163.4, 164.3, 178.1; Anal. Calcd for C₂₄H₁₈N₂O₅: C = 69.56, H = 4.38, N = 6.76, Found: 69.57, H = 4.39, N = 6.74; EI MS *m/z* (% rel. abund.): 414.03 (M⁺, 86.9%)

4.6.10. (*E*)-*N*'-(3-hydroxy-4-methoxybenzylidene)-4-(4-oxo-4H-chromen-2-yl) benzohydrazide (14)

Dark orange solid; Yield 87 %; m.p. 293-295 °C; IR (ATR) cm⁻¹: 3389, 3249, 3064, 2843, 1603, 1584, 1561, 1520, 1470, 1440, 1276, 1246, 1171, 752; ¹H-NMR (500 MHz, DMSO-*d6*): δ 3.82 (s, 3H), 6.91 (t, *J* = 7.5 Hz, 1H), 6.97 (t, *J* = 8.0 Hz, 2H), 7.08 (d, *J* = 8.0 Hz, 1H), 7.20 (t, *J* = 8.0 Hz, 1H), 7.29 (s, 1H), 7.39 (s, 1H), 7.76 (s, 1H), 8.00 (s, 4H), 8.33 (s, 1H), 9.35 (s, 1H), 11.72 (s, 1H); ¹³C-NMR (125 MHz, DMSO-*d6*): δ 56.4, 105.8, 115.0, 115.2, 118.2, 120.3, 125.0, 125.3, 125.5, 126.2, 126.8, 126.8, 129.5, 133.8, 137.2, 138.6, 146.6, 148.5, 149.7, 156.8, 163.4, 164.4, 178.8; Anal. Calcd for C₂₄H₁₈N₂O₅: C = 69.56, H = 4.38, N = 6.76, Found: C = 69.55, H = 4.39, N = 6.74; EI MS *m/z* (% rel. abund.): 414.24 (M⁺, 65.0%)

4.6.11. (*E*)-*N*'-(2-hydroxy-5-methoxybenzylidene)-4-(4-oxo-4H-chromen-2-yl) benzohydrazide (15)

Dark orange solid; Yield 68 %; m.p. 305-307 °C; IR (ATR) cm⁻¹: 3143, 2962, 2835, 1642, 1614, 1583, 1490, 1462, 1266, 1164, 756; ¹H-NMR (500 MHz, DMSO-*d*6): δ 3.75 (s, 3H),

6.87-6.99 (m, 4H), 7.14 (s, 1H), 7.20 (t, J = 7.5 Hz, 1H), 7.37 (s, 1H), 7.76 (s, 1H), 8.02 (m, 4H), 8.66 (s, 1H), 10.71 (s, 1H), 12.02 (s, 1H); ¹³C-NMR (125 MHz, DMSO-*d*6): δ 56.3, 105.9, 112.8, 116.3, 116.5, 118.2, 121.7, 124.9, 125.1, 125.3, 125.3, 126.2, 126.9, 126.9, 133.9, 137.1, 138.7, 149.6, 154.2, 154.4, 156.2, 163.4, 164.3, 178.2; Anal. Calcd for C₂₄H₁₈N₂O₅: C = 69.56, H = 4.38, N = 6.76, Found: C = 69.57, H = 4.36, N = 6.78; EI MS *m*/*z* (% rel. abund.): 414.19 (M⁺, 72.5%)

4.6.12. (E)-N'-(3,5-dimethoxybenzylidene)-4-(4-oxo-4H-chromen-2-yl)benzohydrazide (16)

Light orange solid; Yield 71 %; m.p. 232-234 °C; IR (ATR) cm⁻¹: 3394, 3296, 3002, 2837, 1684, 1652, 1585, 1462, 1266, 1205, 1157, 736; ¹H-NMR (500 MHz, DMSO-*d6*): δ 3.86 (s, 6H), 6.57 (s, 1H), 6.95 (m, 2H), 7.09 (s, 2H), 7.21 (m, 2H), 7.74 (d, *J* = 7.5 Hz, 1H), 8.00 (d, *J* = 8.0 Hz, 2H), 8.06 (d, *J* = 8.0 Hz, 2H), 8.30 (s, 1H), 11.97 (s, 1H); ¹³C-NMR (125 MHz, DMSO-*d6*): δ 56.2, 56.2, 98.5, 105.4, 105.4, 105.9, 118.4, 124.9, 125.2, 125.5, 125.5, 126.2, 126.9, 126.9, 133.9, 137.1, 138.2, 138.7, 147.9, 156.3, 161.2, 161.2, 163.4, 164.3, 178.3; Anal. Calcd for C₂₅H₂₀N₂O₅: C = 70.09, H = 4.71, N = 6.54, Found: C = 70.07, H = 4.70, N = 6.55; EI MS *m/z* (% rel. abund.): 428.35 (M⁺, 54.1%).

4.6.13. (E)-N'-(3-bromo-4-hydroxybenzylidene)-4-(4-oxo-4H-chromen-2-

yl)benzohydrazide (17)

Brown solid; Yield 79 %; m.p. 254-255 °C; IR (ATR) cm⁻¹: 3352, 3140, 3054, 2897, 1650, 1591, 1496, 1282, 1205, 750; ¹H-NMR (500 MHz, DMSO-*d6*): δ 6.91 (t, *J* = 7.5 Hz, 1H), 6.99 (s, 1H), 7.03 (d, *J* = 8.0 Hz, 1H), 7.20 (t, *J*= 8.5 Hz, 1H), 7.58 (d, *J* = 8.5 Hz, 1H), 7.73 (s, 1H), 7.89 (s, 1H), 8.00 (m, 5H), 8.35 (s, 1H), 10.78 (s, 1H), 11.83 (s, 1H); ¹³C-NMR (125 MHz, DMSO-*d6*): δ 105.9, 109.9, 116.3, 118.2, 125.1, 125.4, 125.4, 125.4, 125.7, 126.2, 126.9, 126.9, 128.2, 130.4, 133.9, 137.1, 138.7, 148., 153.7, 156.2, 163.4, 164.3, 178.1; Anal. Calcd for C₂₃H₁₅BrN₂O₄: C = 59.63, H = 3.26, N = 6.05, Found: C = 59.64, H = 3.24, N = 6.06; EI MS *m/z* (% rel. abund.): 462.02 (M+, 54.7%)

4.6.14. (E)-N'-(2-methylbenzylidene)-4-(4-oxo-4H-chromen-2-yl)benzohydrazide (18)

White solid; Yield 85 %; m.p. 225-226 °C; IR (ATR) cm⁻¹: 3433, 3253, 3021, 2361, 1609, 1546, 1466, 1277, 755; ¹H-NMR (500 MHz, DMSO-*d6*): δ 2.47 (s, 3H), 6.91 (t, *J* = 7.5 Hz, 1H), 6.98 (s, 1H), 7.20 (t, *J* = 7.5 Hz, 1H), 7.27-7.34 (m, 4H), 7.87 (d, *J* = 7.5 Hz, 1H), 8.02 (m, 5H), 8.78 (s, 1H), 11.88 (s, 1H); ¹³C-NMR (125 MHz, DMSO-*d6*): δ 21.1, 105.9, 118.2,

125.1, 125.3, 125.5, 125.5, 126.0, 126.3, 126.6, 126.9, 126.9, 128.2, 130.5, 133.5, 133.8, 135.4, 137.1, 138.6, 145.4, 156.3, 163.5, 164.3, 178.1; Anal. Calcd for $C_{24}H_{18}N_2O_3$: C = 75.38, H = 4.74, N = 7.33, Found: C = 75.39, H = 4.75, N = 7.31; EI MS *m*/*z* (% rel. abund.): 382.29 (M⁺, 70.3%).

4.6.15. (E)-N'-(3-methylbenzylidene)-4-(4-oxo-4H-chromen-2-yl)benzohydrazide (19)

White solid; Yield 93 %; m.p. 210-211 °C; IR (ATR) cm⁻¹: 3406, 3193, 3023, 1625, 1557, 1309, 1256, 747; ¹H-NMR (500 MHz, DMSO-*d6*): δ 2.37 (s, 3H), 6.91 (t, *J* = 7.0 Hz, 1H), 6.98 (d, *J* = 7.5 Hz, 1H), 7.20 (t, *J* = 8.0 Hz, 1H), 7.27(d, *J* = 7.5 Hz, 1H), 7.35 (t, *J* = 7.5 Hz, 1H), 7.45 (s, 1H), 7.53 (d, *J* = 7.5 Hz, 1H), 7.59 (s, 1H), 7.75 (s, 1H), 8.01 (m, 4H), 8.45 (s, 1H), 11.89 (s, 1H); ¹³C-NMR (125 MHz, DMSO-*d6*): δ 21.4, 105.9, 118.2, 123.5, 124.8, 125.2, 125.5, 125.5, 126.0, 126.8, 126.8, 128.2, 128.6, 129.7, 133.8, 136.6, 137.2, 137.9, 138.8, 148.7, 156.2, 163.4, 164.3, 178.3; Anal. Calcd for C₂₄H₁₈N₂O₃: C = 75.38, H = 4.74, N = 7.33, Found: C = 75.39, H = 4.73, N = 7.31; EI MS *m/z* (% rel. abund.): 382.12 (M⁺, 61.7%).

4.6.16. (E)-N'-(4-methylbenzylidene)-4-(4-oxo-4H-chromen-2-yl)benzohydrazide (20)

Light yellow solid; Yield 74 %; m.p. 232-234 °C; IR (ATR) cm⁻¹: 3422, 3382, 3242, 3164, 1609, 1464, 1270, 749; ¹H-NMR (MeOD- d_4 , 500 MHz): δ 2.41 (s, 3H), 6.94-6.98 (m, 2H), 7.21-7.24 (m, 2H), 7.29 (d, J = 8.0 Hz, 2H), 7.75-7.77 (m, 3H), 7.99 (d, J = 7.5 Hz, 2H), 8.06 (d, J = 7.5 Hz, 2H), 8.36 (s, 1H), 12.27 (s, 1H); ¹³C-HMR (125 MHz, DMSO- d_6): δ 21.15, 105.9, 118.2, 125.0, 118.2, 125.0, 125.3, 125.3, 125.8, 126.1, 126.8, 126.8, 127.3, 129.2, 131.8, 133.9, 137.2, 138.2, 138.7, 149.4, 156.3, 163.5, 164.5, 178.0; Anal. Calcd for C₂₄H₁₈N₂O₃: C = 75.38, H = 4.74, N = 7.33, Found: C = 75.39, H = 4.76, N = 7.31; EI MS *m/z* (% rel. abund.): 382.07 (M⁺, 62.5%).

4.6.17. (E)-N'-(2-chlorobenzylidene)-4-(4-oxo-4H-chromen-2-yl)benzohydrazide (21)

White solid; Yield 78 %; m.p. 294-296 °C; IR (ATR) cm⁻¹: 3432, 3183, 3065, 1613, 1589, 1543, 1467, 1280, 757; ¹H-NMR (500 MHz, DMSO-*d6*): δ 1H-NMR (MeOD-d4, 500 MHz): δ 6.94-6.98 (m, 2H), 7.21-7.25 (m, 2H), 7.42-7.46 (m, 2H), 7.48 (d, J = 8.0 Hz, 1H), 7.75 (dd, J = 7.5 Hz, 1.5 Hz, 1H), 8.00 (d, J = 7.0 Hz, 2H), 8.08 (d, J = 7.5 Hz, 2H), 8.31 (dd, J = 7.5 Hz, 1.5 Hz, 1H), 8.91 (s, 1H), 12.23 (s, 1H); ¹³C-NMR (125 MHz, DMSO-*d6*): δ 105.9, 118.2, 125.0, 125.2, 125.4, 125.4, 126.5, 126.9, 126.9, 127.3, 127.5, 128.6, 130.9, 131.3, 132.7, 133.9, 137.1, 138.7, 149.8, 156.2, 163.2, 164.3, 178.1; Anal. Calcd for C₂₃H₁₅ClN₂O₃: C = 68.58, H =

3.75, N = 6.95, Found: C = 68.59, H = 3.76, N = 6.93; EI MS *m*/*z* (% rel. abund.): 402.08 (M+, 74.2%), 404.08 (M+2, 13.8%).

4.6.18. (E)-N'-(3-chlorobenzylidene)-4-(4-oxo-4H-chromen-2-yl)benzohydrazide (22)

Light yellow solid; Yield 69 %; m.p. 304-306 °C; IR (ATR) cm⁻¹: 3273, 3120, 2915, 1660, 1542, 1469, 1253, 743; ¹H-NMR (500 MHz, DMSO-*d6*): δ 6.89 (t, J = 7.5 Hz, 1H), 6.96 (d, J = 8.0 Hz, 1H), 7.18 (t, J = 8.0 Hz, 1H), 7.34 (s, 1H), 7.51 (d, J = 4.0 Hz, 2H), 7.72-7.75 (m, 2H), 7.81 (s, 1H), 8.02 (s, 4H), 8.47 (s, 1H), 12.06 (s, 1H); ¹³C-NMR (125 MHz, DMSO-*d6*): δ 105.9, 118.2, 124.4, 124.8, 125.2, 125.4, 125.4, 126.2, 126.9, 126.9, 127.6, 128.0, 130.2, 133.7, 134.0, 135.3, 137.1, 138.7, 148.6, 156.2, 163.4, 164.3, 178.2; Anal. Calcd for C₂₃H₁₅ClN₂O₃: C = 68.58, H = 3.75, N = 6.95, Found: C = 68.57, H = 3.77, N, 6.97; EI MS *m/z* (% rel. abund.): 402.08 (M+, 35.4%), 404.36 (M+2, 13.8%).

4.6.19. (*E*)-*N*'-(4-chlorobenzylidene)-4-(4-oxo-4H-chromen-2-yl)benzohydrazide (23)

Yellow solid; Yield 91 %; m.p. 295-297 °C; IR (ATR) cm⁻¹: 3427, 3126, 2947, 1608, 1589, 1466, 1273, 752; ¹H-NMR (MeOD- d_4 , 500 MHz): δ 6.94-6.98 (m, 2H), 7.21 (td, J = 8.5 Hz, 1.0 Hz, 1H), 7.24 (s, 1H), 7.48 (d, J = 8.5 Hz, 2H), 7.75 (dd, J = 7.5 Hz, 1.5 Hz, 1H), 7.87 (d, J = 8.5 Hz, 2H), 8.00 (d, J = 7.5 Hz, 2H), 8.02 (d, J = 8.0 Hz, 2H), 8.38 (s, 1H), 12.03 (s, 1H); ¹³C-NMR (125 MHz, DMSO-d6): δ 105.9, 118.4, 125.1, 125.4, 125.4, 125.8, 126.3, 126.8, 126.8, 129.0, 129.0, 129.4, 129.4, 132.3, 134.1, 135.1, 137.1, 138.7, 149.4, 156.3, 163.3, 164.3, 178.3. Anal. Calcd for C₂₃H₁₅ClN₂O₃: C = 68.58, H = 3.75, N = 6.95, Found: C = 68.59, H = 3.74, N = 6.94; EI MS *m*/*z* (% rel. abund.): 402.08 (M+, 64.7%), 404.21 (M+2, 13.8%).

4.6.20. (E)-N'-(2-nitrobenzylidene)-4-(4-oxo-4H-chromen-2-yl)benzohydrazide (24)

Yellow solid; Yield 65 %; m.p. 305-306 °C; IR (ATR) cm⁻¹: 3431, 3255, 3153, 3024, 1623, 1543, 1519, 1468, 1274, 755; ¹H-NMR (MeOD- d_4 , 500 MHz): δ 6.91 (t, J = 7.5 Hz, 1H), 6.98 (d, J = 7.5 Hz, 1H), 7.20 (t, J = 7.5 Hz, 1H), 7.37 (s, 1H), 7.69-7.93 (m, 3H), 8.03-8.18 (m, 6H), 8.91 (s, 1H), 12.26 (s, 1H); ¹³C-NMR (125 MHz, DMSO- d_6): δ 105.91, 118.28, 125.00, 125.1, 125.2, 125.4, 125.4, 126.2, 126.9, 126.9, 128.0, 129.3, 130.4, 132.6, 133.7, 137.1, 138.7, 142.9, 147.9, 156.9, 163.4, 164.3, 178.1; Anal. Calcd for C₂₃H₁₅N₃O₅: C = 66.83, H = 3.66, N

= 10.17, Found: C = 66.84, H = 3.65, N = 10.19; EI MS m/z (% rel. abund.): 413.07 (M⁺, 42.7%).

4.6.21. (E)-N'-(3-nitrobenzylidene)-4-(4-oxo-4H-chromen-2-yl)benzohydrazide (25)

Light orange solid; Yield 62 %; m.p. 312-314 °C; IR (ATR) cm⁻¹: 3312, 3179, 3143, 2886, 1657, 1519, 1469, 1342, 1274, 1145, 732; ¹H-NMR (500 MHz, DMSO-*d*6): δ 6.91 (t, *J* = 7.5 Hz, 1H), 6.98 (d, *J* = 8.0 Hz, 1H), 7.20 (t, *J* = 7.5 Hz, 1H), 7.37 (s, 1H), 7.75-7.77 (m, 2H), 8.03 (m, 4H), 8.17 (d, *J* = 6.5 Hz, 1H), 8.27(d, *J* = 7.5 Hz, 1H), 8.57-8.60 (m, 3H), 12.16 (s, 1H); ¹³C-NMR (125 MHz, DMSO-*d*6): δ 105.94, 118.22, 122.37, 124.18, 125.05, 125.24, 125.5, 125.5, 126.2, 126.9, 126.9, 129.9, 132.6, 133.9, 137.1, 137.7, 138.7, 147.41, 148.6, 156.2, 163.4, 164.3, 178.1; Anal. Calcd for C₂₃H₁₅N₃O₅: C = 66.83, H = 3.66, N = 10.17, Found: C = 66.84, H = 3.65, N = 10.18; EI MS *m*/*z* (% rel. abund.): 413.15 (M⁺, 52.6%).

4.6.22. (E -(4-nitrobenzylidene)-4-(4-oxo-4H-chromen-2-yl)benzohydrazide (26)

Dark brown solid; Yield 88 %; m.p. 297-299 °C; IR (ATR) cm⁻¹: 3413, 3075, 2833, 1633, 1514, 1466, 1341, 1269, 1113, 756; ¹H-NMR (500 MHz, DMSO-*d6*): δ 6.92 (t, J = 7.5 Hz, 1H), 6.98 (d, J = 8.0 Hz, 1H), 7.20 (t, J = 7.5 Hz, 1H), 7.37 (s, 1H), 7.75 (d, J = 7.0 Hz, 1H), 8.03 (m, 6H), 8.32-8.33 (m, 2H), 8.59 (s, 1H), 12.22 (s, 1H); ¹³C-NMR (125 MHz, DMSO-*d6*): δ 106.5, 118.2, 124.4, 124.4, 125.0, 125.2, 125.4, 125.4, 125.6, 126.2, 126.9, 126.9, 127.9, 127.9, 133.9, 137.1, 138.7, 148.0, 149.3, 156.2, 163.4, 164.3, 178.1; Anal. Calcd for C₂₃H₁₅N₃O₅: C = 66.83, H = 3.66, N = 10.17, Found: C = 66.82, H = 3.67, N = 10.18; EI MS *m/z* (% rel. abund.): 413.19 (M⁺, 64.8%).

4.6.23. (E)-N'-(2-fluorobenzylidene)-4-(4-oxo-4H-chromen-2-yl)benzohydrazide (27)

White solid; Yield 84 %; m.p. 257-259 °C; IR (ATR) cm⁻¹: 3434, 3232, 3019, 2861, 1660, 1608, 1519, 1466, 1360, 1272, 1187, 752; ¹H-NMR (MeOD- d_4 , 500 MHz): δ 6.96-6.98 (m, 2H), 7.19-7.24 (m, 3H), 7.28 (t, J = 7.5 Hz, 1H), 7.48 (d, J = 8.0 Hz, 1H), 7.74 (dd, J = 7.5 Hz, 1.5 Hz, 1H), 8.00 (d, J = 8.5 Hz, 2H), 8.07 (d, J = 8.5 Hz, 2H), 8.24 (t, J = 7.5 Hz, 1H), 8.70 (s, 1H), 12.11 (s, 1H); ¹³C-NMR (125 MHz, DMSO- d_6): δ 105.9, 117.4, 118.3, 124.6, 125.1, 125.4, 125.6, 125.8, 126.3, 126.9, 126.9, 129.4, 130.9, 133.9, 137.1, 138.7, 149.3, 156.1, 159.3 (d, J = 282.5 Hz), 163.4, 164.3, 178.3; Anal. Calcd for C₂₃H₁₅FN₂O₃: C = 71.50, H = 3.91, F = 4.92, N = 7.25, Found: C = 71.51, H = 3.89, F = 4.91, N = 7.26; EI MS *m*/*z* (% rel. abund.): 386.06 (M⁺, 72.6%).

4.6.24. (E)-N'-(3-fluorobenzylidene)-4-(4-oxo-4H-chromen-2-yl)benzohydrazide (28)

White solid; Yield 78 %; m.p. 254-256 °C; IR (ATR) cm-1: 3119, 3026, 2828, 1658, 1539, 1449, 1269, 745; ¹H-NMR (MeOD- d_4 , 500 MHz): δ 6.94-6.98 (m, 2H), 7.18-7.24 (m, 3H), 7.46 (d, J = 8.0 Hz, 1H), 7.62 (d, J = 7.5 Hz, 1H), 7.72-7.76 (m, 2H), 8.00 (d, J = 7.5 Hz, 2H), 8.06 (d, J = 8.0 Hz, 2H), 8.38 (s, 1H), 12.20 (s, 1H); ¹³C-NMR (125 MHz, DMSO-d6): δ 105.8, 114.4, 117.1, 118.4, 121.8, 124.7, 125.2, 125.7, 125.7, 126.2, 126.9, 126.9, 129.9, 133.9, 137.1, 138.2, 138.7, 148.6, 156.2, 161.6 (d, J = 193.75 Hz), 163.7, 164.4, 178.1; Anal. Calcd for C₂₃H₁₅FN₂O₃: C = 71.50, H = 3.91, N = 7.25. Found: C = 71.51, H = 3.93, F = 4.91, N = 7.24; EI MS *m*/*z* (% rel. abund.): 386.11 (M⁺, 62.7%).

4.6.25. (E)-N'-(4-fluorobenzylidene)-4-(4-oxo-4H-chromen-2-yl)benzohydrazide (29)

Light yellow solid; Yield 95 %; m.p. 252-254 °C; IR (ATR) cm-1: 3421, 3282, 3061, 2947, 1633, 1609, 1505, 1467, 1233, 752; ¹H-NMR (500 MHz, DMSO-*d*6): δ 6.92 (t, *J* = 7.0 Hz, 1H), 6.98 (d, *J* = 8.0 Hz, 1H), 7.20 (t, *J* = 7.5 Hz, 1H), 7.30-7.36 (m, 3H), 7.75-7.82(m, 3H), 8.02(m, 4H), 8.49(s, 1H), 11.92(s, 1H); ¹³C-NMR (125 MHz, DMSO-*d*6): δ 105.9, 115.3, 115.3, 115.6, 118.2, 124.6, 125.3, 125.7, 125.7, 126.2, 126.9, 126.9, 130.3, 130.8, 130.8, 133.8, 137.1, 138.7, 149.3, 156.2, 160.3 (d, *J* = 245.00 Hz), 163.3, 164.3, 178.2; Anal. Calcd for C₂₃H₁₅FN₂O₃: C = 71.50, H = 3.91, N = 7.25, Found: C = 71.49, H = 3.93, N = 7.26; EI MS *m/z* (% rel. abund.): 386.41 (M⁺, 47.1%).

4.6.26. (E)-N'-(3-methoxybenzylidene)-4-(4-oxo-4H-chromen-2-yl)benzohydrazide (30)

Light brown solid; Yield 91 %; m.p. 231-233 °C; IR (ATR) cm-1: 3539, 3252, 3051, 2835, 1640, 1564, 1469, 1253, 1141, 747; ¹H-NMR (MeOD- d_4 , 500 MHz): δ 3.90 (s, 3H), 6.94-6.98 (m, 2H), 7.02 (d, J = 8.0 Hz, 1H), 7.21-7.25 (m, 2H), 7.33-7.39 (m, 2H), 7.62 (s, 1H), 7.75 (d, J = 6.5 Hz, 1H), 8.00 (d, J = 7.0 Hz, 2H), 8.07 (d, J = 7.5 Hz, 2H), 8.36 (s, 1H), 12.08 (s, 1H); ¹³C-NMR (125 MHz, DMSO-d6): δ 56.2, 105.9, 112.2, 114.8, 118.2, 121.2, 124.9, 125.2, 125.4, 125.4, 126.2, 126.9, 126.9, 129.2, 133.9, 136.9, 137.1, 138.7, 148.6, 156.3, 160.2, 163.2, 164.2, 178.3; Anal. Calcd for C₂₄H₁₈N₂O₄: C = 72.35; H = 4.55; N = 7.03, Found: C = 72.33; H = 4.56; N = 7.01; EI MS *m*/*z* (% rel. abund.): 398.26 (M⁺, 51.7%).

4.6.27. (*E*)-*N*'-(4-methoxybenzylidene)-4-(4-oxo-4H-chromen-2-yl)benzohydrazide (31)

Yellow solid; Yield 89 %; m.p. 237-239 °C; IR (ATR) cm-1: 3424, 3269, 3065, 2843, 1610, 1507, 1466, 1356, 1244, 755; ¹H-NMR (500 MHz, DMSO-*d6*): δ 3.83 (s, 3H), 6.92 (t, *J* = 7.0 Hz, 1H), 6.97 (d, *J* = 8.0 Hz, 1H), 7.04 (d, *J* = 8.5 Hz, 2H), 7.20 (t, *J* = 7.0 Hz, 1H), 7.36 (s,

1H), 7.70-7.77 (m, 3H), 8.00 (s, 4H), 8.44 (s, 1H), 11.77 (s, 1H); ¹³C-NMR (125 MHz, DMSOd6): δ 56.0, 105.9, 114.3, 114.3, 118.2, 125.0, 125.2, 125.4, 125.4, 126.2, 126.6, 127.0, 127.0, 129.1, 129.1, 133.9, 137.1, 138.7, 149.3, 156.2, 160.1, 163.2, 164.6, 178.3. Anal. Calcd for $C_{24}H_{18}N_2O_4$: C = 72.35, H = 4.55, N = 7.03, Found: C = 72.36, H = 4.53, N = 7.02; EI MS *m*/*z* (% rel. abund.): 398.17 (M⁺, 69.4%).

4.6.28. (E)-4-(4-oxo-4H-chromen-2-yl)-N'-(pyridin-2-ylmethylene)benzohydrazide (32)

Orange solid; Yield 81 %; m.p. 170-172 °C; IR (ATR) cm-1: 3395, 3170, 3058, 2849, 1643, 1564, 1465, 1255, 734; ¹H-NMR (MeOD- d_4 , 500 MHz): δ 6.95-6.98 (m, 2H), 7.21-7.26 (m, 2H), 7.46 (d, J = 5.0 Hz, 1H), 7.75 (d, J = 8.0 Hz, 1H), 7.93 (t, J = 7.5 Hz, 1H), 8.01-8.02 (m, 2H), 8.07-8.13 (m, 3H), 8.34 (d, J = 7.5 Hz, 1H), 8.45 (s, 1H), 11.78 (s, 1H); ¹³C-NMR (125 MHz, DMSO-d6): δ 105.9, 118.2, 119.0, 123.0, 125.0, 125.2, 125.4, 125.4, 126.3, 126.9, 126.9, 133.9, 137.1, 137.6, 138.7, 146.1, 147.9, 154.6, 156.2, 163.2, 164.5, 178.0. Anal. Calcd for C₂₂H₁₅N₃O₃: C = 71.54, H = 4.09, N = 11.38, Found: C = 71.55, H = 4.10, N = 11.39; EI MS *m/z* (% rel. abund.): 369.21 (M⁺, 47.0%).

4.6.29. (*E*)-4-(4-oxo-4H-chromen-2-yl)-*N*'-(pyridin-3-ylmethylene)benzohydrazide (33)

Orange solid; Yield 79 %; m.p. 210-212 °C; IR (ATR) cm-1: 3417, 3037, 2837, 1641, 1556, 1467, 1280, 755; ¹H-NMR (MeOD- d_4 , 500 MHz): δ 6.96 (m, 2H), 7.21-7.24 (m, 2H), 7.55 (d, J = 5.0 Hz, 1H), 7.75 (d, J = 5.5 Hz, 1H), 8.00-8.07 (m, 4H), 8.43-8.45 (m, 2H), 8.61 (d, J = 3.5 Hz, 1H), 8.95 (s, 1H), 12.21 (s, 1H); ¹³C-NMR (125 MHz, DMSO-d6): δ 105.9, 118.2, 124.41, 125.0, 125.2, 125.4, 125.4, 126.2, 126.9, 126.9, 133.8, 134.6, 135.6, 137.1, 138.7, 145.5, 146.3, 149.8, 156.2, 163.4, 164.2, 178.1; Anal. Calcd for C₂₂H₁₅N₃O₃: C = 71.54, H = 4.09, N = 11.38, Found: C = 71.53, H = 4.08, N = 11.37; EI MS *m*/*z* (% rel. abund.): 369.15 (M⁺, 67.0%).

4.6.30. (E)-4-(4-oxo-4H-chromen-2-yl)-N'-(pyridin-4-ylmethylene)benzohydrazide (34)

Dark red solid; Yield 76 %; m.p. 198-200 °C; IR (ATR) cm-1: 3289, 3172, 2850, 1657, 1544, 1466, 1262, 1113, 754; ¹H-NMR (MeOD- d_4 , 500 MHz): δ 6.94-6.98 (m, 2H), 7.21 (t, J = 7.5 Hz, 1H), 7.25 (s, 1H), 7.74 (dd, J = 8.0 Hz, 1.5 Hz, 1H), 7.88 (d, J = 5.0 Hz, 2H), 8.01 (d, J = 8.5 Hz, 2H), 8.08 (d, J = 8.0 Hz, 2H), 8.39 (s, 1H), 8.64 (d, J = 5.5 Hz, 2H), 12.76 (s, 1H); ¹³C-NMR (125 MHz, DMSO-d6): δ 105.9, 118.2, 122.3, 122.3, 125.0, 125.28, 125.4, 125.4, 126.2, 126.9, 126.9, 133.9, 137.1, 138.7, 140.7, 149.3, 150.0, 150.0, 156.2, 163.4, 164.3, 178.1.

Anal. Calcd for $C_{22}H_{15}N_3O_3$: C = 71.54, H = 4.09, N = 11.38, Found: C = 71.56, H = 4.11, N = 11.36; EI MS *m*/*z* (% rel. abund.): 369.07 (M+, 64.5%)

4.4 α-Glucosidase Inhibition Assay

The α -glucosidase inhibition assay had been carried out using baker's yeast α -glucosidase (EC 3.2.1.20) and *p*-nitrophenyl- α -D-glucopyranoside [49]. The samples (5 μ g/mL) were prepared by dissolving the compounds **1-35** in DMSO. Test samples (10 μ L) which had been prepared were reconstituted in 100 μ L of phosphate buffer (100 mM) at pH 6.8 in 96-well micro-plate and incubated with 50 μ L of baker's yeast α -glucosidase for 5 minutes before 50 μ L of *p*-nitrophenyl- α -D-glucopyranoside (5 mM) was added. After incubating for 5 minutes, the absorbance was measured at 405 nm using SpectraMax plus384 (Molecular Devices Corporation, Sunnyvale, CA, USA). Blank in which the substrate was changed with 50 μ L of buffer were analysed to accurately determine the background absorbance. Control sample was prepared to contain 10 μ L DMSO instead of test samples. Percentage of enzyme inhibition was measured using the following formula.

% inhibition = $[(A - B)/A] \times 100$

[A] represents absorbance of control samples, and [B] corresponding to absorbance in presence of test samples.

4.5 In Vitro HDAC Inhibition Assay

This assay had been performed in accordance with method in Li *et al.* (2014) [50]. In this assay, 10 μ L of enzyme solution was mixed with 50 μ L of tested compound at various concentrations. The mixture was incubated at 37 °C for 5 minutes prior to addition of 40 μ L fluorogenic substrate Boc-Lys(acetyl)-AMC. After further incubation at 37 °C for 30 min, the reaction was stopped by adding 100 μ L of trypsin and TSA. Fluorescence intensity was measured after 20 minutes using a microplate reader at excitation and emission wavelengths of 390 and 460 nm, respectively.

4.6 Cytotoxicity

The Neutral Red cytotoxicity assay is based on the initial protocol described by Borenfreund *et al.* (1998) [51] with some modifications. Briefly, the cells (1x104/well) were seeded in 96-well

microtiter plates (Nunc) and allowed to grow for 24 hours before treatment. After 24 hours of incubation, the cells were treated with six different concentrations (0.1-100 μ g/mL) of test compounds, in three replicates. The plates were incubated for 72 h at 37°C in a 5% CO₂ incubator. A stock solution was obtained by dissolving the test compounds in DMSO. Further dilution to different tested concentrations were then carried out ensuring that the final concentration of DMSO in the test and control wells was not in excess of 1% (v/v). No effect due to the DMSO was observed. Doxorubicin was used as the positive control. The well containing untreated cells was the negative control. At the end of the incubation period, the media were replaced with medium containing 50 μ g/mL of Neutral Red. The plates were incubated for another 3 hours to allow for uptake of the vital dye into the lysosomes of viable and injured cells. After the incubation period, the media were removed and cells were washed with the neutral red washing solution. The dye was eluted from the cells by adding 200 μ L of Neutral Red resorb solution and incubated for 30 minutes at room temperature with rapid agitation on a microtiter plate shaker. Dye absorbance was measured at 540 nm using a spectrophotometer ELISA plate reader.

4.7 2D QSAR

Training set and test set compounds were being selected using "Cluster Ligands" protocol in Discovery Studio DS 2.5 software (Discovery Studio 2.5, Accelrys, Co. Ltd). Another module, "Calculate Molecular Properties", was used to calculate the 2D molecular properties as well as energies of highest occupied and lowest unoccupied molecular orbitals (HOMO and LUMO) of the training set compounds. All semi-empirical calculation and density functional theory was carried out using software package VAMP and DMol3 in Discovery Studio 2.5. Different 2D descriptors such as AlogP, molecular properties (molecular weight), molecular property counts (Num aromatic Rings, Num_-H_Acceptors, Num H Donors, Num Rings and Num_Rotatable Bonds) and surface area and volume (Molecular_Fractional Polar Surface Area), were utilized in our model. The model was validated using test set correlation and Leave-one-out (LOO) cross validation.

4.8 Docking studies

The structure of all compounds were prepared using Chem3D by CambridgeSoft. The geometry and energy of the structures were being optimized using Steepest-Descent and Polak-Ribiere algorithm in HyperChem. AutoDock 4.2 [52] was used to identify the binding modes of flavone derivatives (5-34) responsible for the activity. Genetic Algorithm (GA) with default

settings was employed for the studies. Protein sequence for Baker's yeast α -glucosidase (MAL12) was obtained from uniprot (<u>http://www.uniprot.org</u>). Homology model for *Saccharomyces cerevisiae* α -glucosidase was built using crystal structure of α -D-glucose bound isomaltase from *Saccharomyces cerevisiae* (PDB ID: 3A4A) which shares 72% identical and 85% similar sequence with α -glucosidase. Sequence alignment and homology modelling were performed using Swiss-Model, which is a fully automated homology modelling pipeline SWISS-MODEL, managed by Swiss Institute of Bioinformatics [53-55]. The docking results had been visualized using Discovery Studio visualizer 3.5 [56] and PyMol [57]. The homology model was being evaluated by using PROCHECK [58, 59].

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Schemes, Figure, and Table Caption

Scheme 1. Reaction scheme for synthesis of compound 4. Reagents and condition: (a) Ethanolic-KOH, room temperature; (b) I₂-DMSO, reflux; (c) MeOH, H₂SO₄, reflux; (d) Hydrazine hydrate, MeOH, reflux.

Scheme 2. Reaction scheme for synthesis of flavone hydrazones 5-34

Figure 1. Representation of H_{α} and H_{β} of chalcone 1 and flavone 2.

Figure 2. Predicted versus experimental pIC_{50} of the (a) training set and (b) test set against α -glucosidase.

Figure 3. Sequence analysis between 3A4A and MAL12 showing high sequence homology of important catalytic residues monitored throughout docking studies.

Figure 4. Ramachandran plot for homolog model of 3A4A.

Figure 5. Superimpose of (a) acarbose in 3W37 and (b) α -D-glucopyranose in 3A4A.

Figure 6. Flavone derivatives (5-34) aligned in the binding pocket

Figure 7. (a) Docking of the most active compound 5 and (b) its 2D interaction diagram.

Figure 8. Hydrophobic and steric surfaces of (a) compound 14, (b) compound 15, (c) compound 30 and (d) compound 31.

Table 1. Synthesis of flavone hydrazone derivatives 5-34.

Table 2. In-vitro α-glucosidase Inhibition Activity of Compounds 1-34

Table 3. a) Intercorrelation Data of Descriptors Used to Develop QSAR Model Equation 1 andb) Regression Statistics Table

Table 4. Experimental activity of the synthesized hybrids against the predicted activity according to Equation 1.

Compound	R		
5	2,4,6-OH		
6	2,3-OH		
7	2,4-OH		
8	2,5-OH		
9	3,4-OH		
10	2-OH		
11	3-OH		
12	4-OH		
13	2-OH, 4-OCH ₃		
14	3-OH, 4-OCH ₃		
15	2-OH, 5-OCH ₃		
16	3,5-OCH ₃		
17	2-Br, 4-OH		
18	2-CH ₃		
19	3-CH ₃		
20	4-CH ₃		
21	2-C1		
22	3-C1		
23	4-C1		
24	2-NO ₂		
25	3-NO ₂		
26	4-NO ₂		
27	2-Fl		
28	3-Fl		
29	4-Fl		
30	3-OCH ₃		
31	4-OCH ₃		
32	2-Pyr		
33	3-Pyr		
34	4-Pyr		

Table 1. Synthesis of flavone hydrazone derivatives5-34.

Compound	$IC_{50}(\mu M \pm SEM^{a})$	Compound	$IC_{50}(\mu M \pm SEM^a)$	
1	840 ± 2.50	19	497.4 ± 0.70	
2	802.6 ± 1.34	20	367.4 ± 0.54	
3	N.A. ^b	21	29.6 ± 0.26	
4	730 ± 0.50	22	64.5 ± 2.08	
5	15.4 ± 0.22	23	38.3 ± 0.34	
6	16.9 ± 0.65	24	123.4 ± 1.69	
7	27.3 ± 0.36	25	98.0 ± 1.10	
8	37.8 ± 0.44	26	88.4 ± 0.97	
9	17.2 ± 1.21	27	17.1 ± 0.24	
10	37.4 ± 0.50	28	22.8 ± 1.23	
11	86.3 ± 0.98	29	19.4 ± 0.20	
12	27.4 ± 1.46	30	680.5 ± 1.27	
13	29.4 ± 0.32	31	690.3 ± 2.04	
14	34.4 ± 1.18	32	487.4 ± 1.36	
15	37.4 ± 0.33	33	520.7 ± 0.80	
16	623 .1 ± 0.61	34	430.2 ± 0.28	
17	587.4 ± 1.42	Acarbose	860.23 ± 6.10	
18	487.4 ± 1.08			

 Table 2. In-vitro α-glucosidase Inhibition Activity of Compounds 1-34

^aSEM is the standard error of the mean, ^bN.A. No activity

C,

Table 3. a) Intercorrelation Data of Descriptors Used to Develop QSAR Model Equation 1 andb) Regression Statistics Table

a) Intercorrelation Data of Descriptors Us	sed to Develop QSAR Model Equation 1
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	pIC ₅₀	ALogP	Molecular_Weight	Num_H_Donors	Num_RotatableBonds
pIC50	1	-0.034	0.265	0.271	-0.221
ALogP		1	0.343	-0.422	0.032
Molecular_Weight			1	0.502	0.237
Num_H_Donors				1	-0.192
Num_RotatableBonds					1
b) Regression Statistic	cs Table	e			
Statistic		V	alue		
N			21		
r			.921		
R2			.848		
r^2_2 (Adjusted)			0.840		
r ² (predicted)			.705		
Least-squared Erro	or	0	.198		
		_			

Compound	Experimental Activity (-log IC ₅₀)	Predicted Activity (-log IC ₅₀)	Residual
11	-1.936	-1.445	-0.491
13	-1.468	-1.539	0.071
15	-1.572	-1.539	-0.034
18	-2.688	-2.632	-0.056
19	-2.696	-2.632	-0.065
24	-2.091	-1.957	-0.134
27	-1.240	-1.313	0.073
28	-1.350	-1.313	-0.037
33	-2.717	-2.684	-0.032

Table 4. Experimental activity of the synthesized hybrids against the predicted activityaccording to Equation 1.

COOH

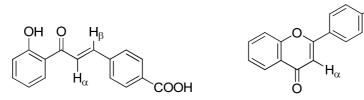


Figure 1. Representation of H_{α} and H_{β} of chalcone 1 and flavone 2.

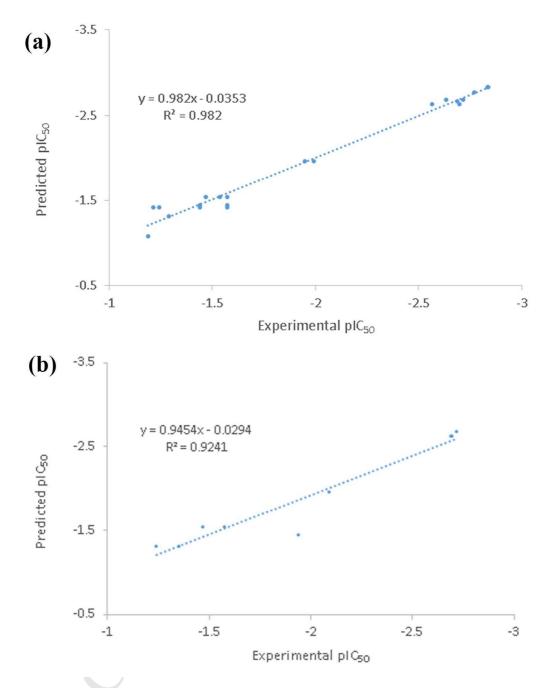


Figure 2. Predicted versus experimental pIC₅₀ of the (a) training set and (b) test set against α -glucosidase.

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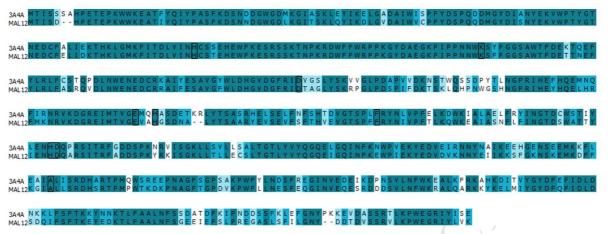


Figure 3. Sequence analysis between 3A4A and MAL12 showing high sequence homology of important catalytic residues monitored throughout docking studies.

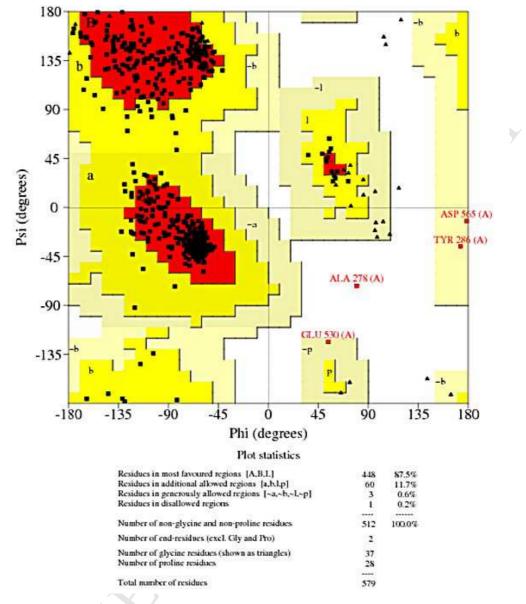


Figure 4. Ramachandran plot for homolog model of 3A4A.

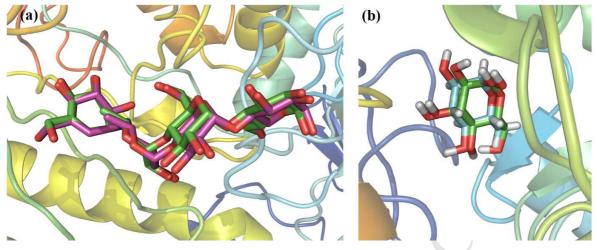


Figure 5. Superimpose of (a) acarbose in 3W37 and (b) α -*D*-glucopyranose in 3A4A.

CER HA

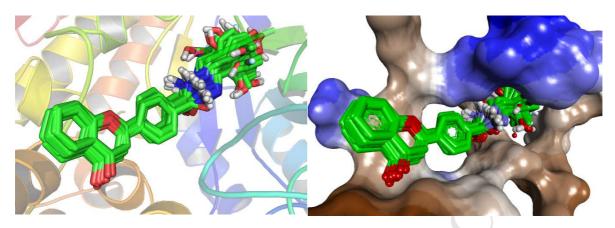


Figure 6. Flavone derivatives (5-34) aligned in the binding pocket

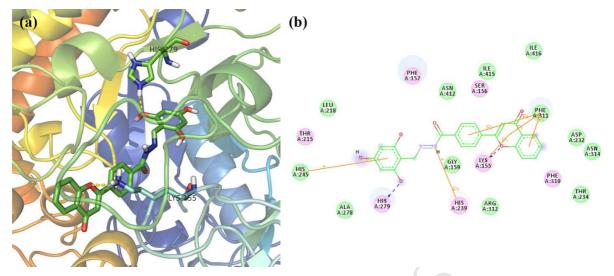


Figure 7. (a) Docking of the most active compound 5 and (b) its 2D interaction diagram.

CER CER

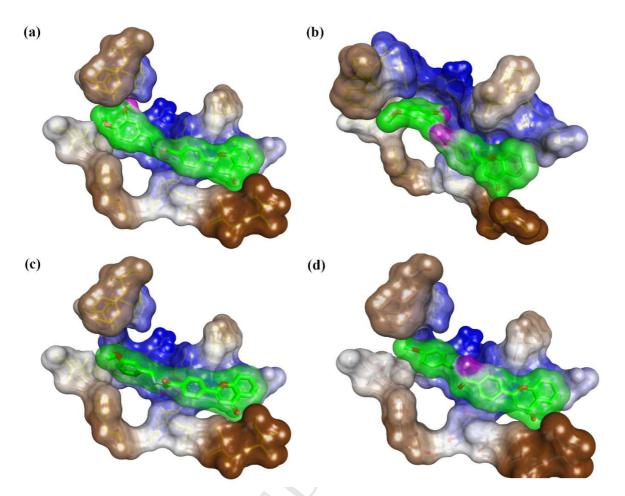
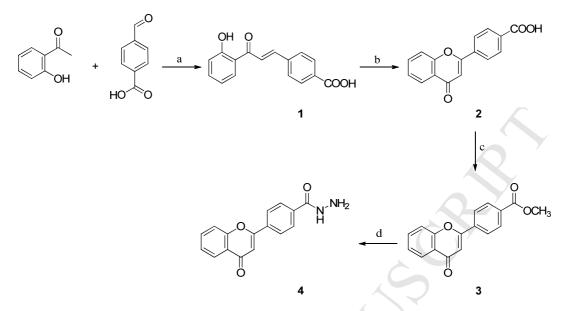
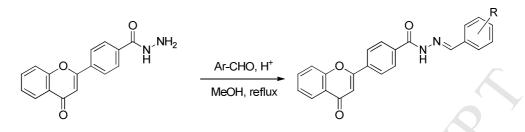


Figure 8. Hydrophobic and steric surfaces of (a) compound 14, (b) compound 15, (c) compound 30 and (d) compound 31.



Scheme 1. Reaction scheme for synthesis of compound 4. Reagents and condition: (a) Ethanolic-KOH, room temperature; (b) I₂-DMSO, reflux; (c) MeOH, H₂SO₄, reflux; (d) Hydrazine hydrate, MeOH, reflux.



4 5-34 Scheme 2. Reaction scheme for synthesis of flavone hydrazones **5-34**.

CHER MAN

Research Highlights

- Synthesis of 30 novel flavone derivatives
- > In vitro α -glucosidase inhibitory activity.
- > All hydrazone derivatives are more active than standard drug
- > Quantitative Structure-activity relationship model has been established
- Docking studies were carried out to confirm binding of active compounds with enzyme

Chillip Mark

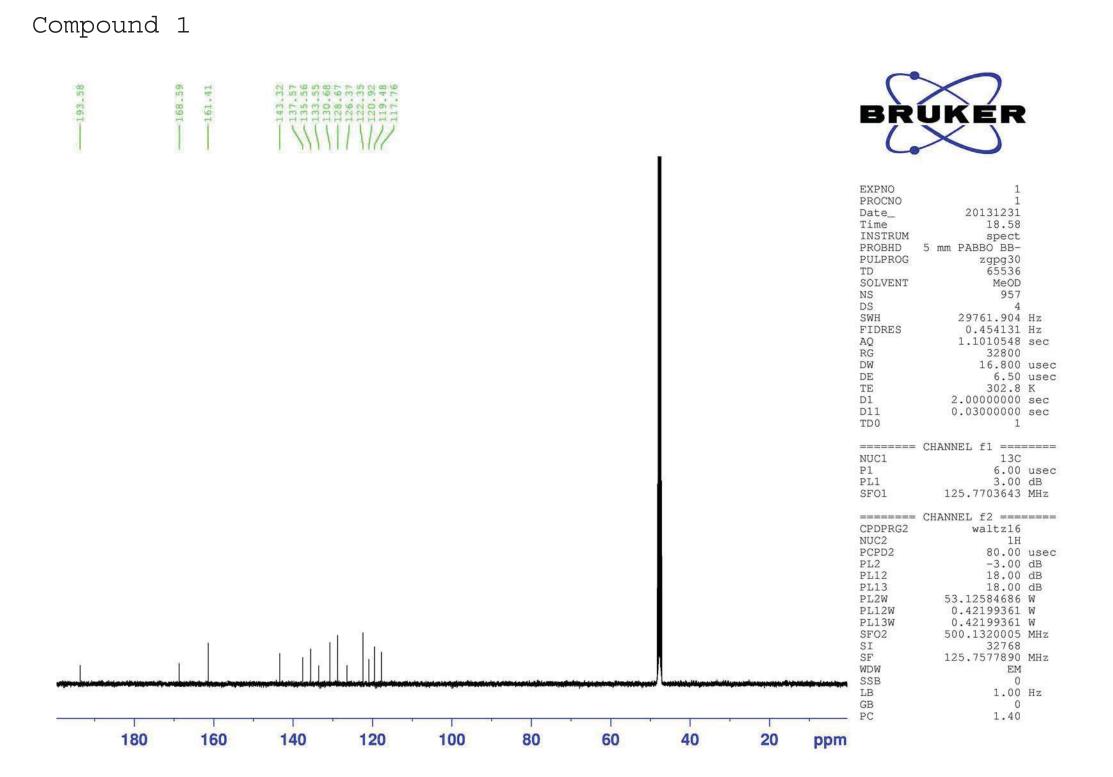
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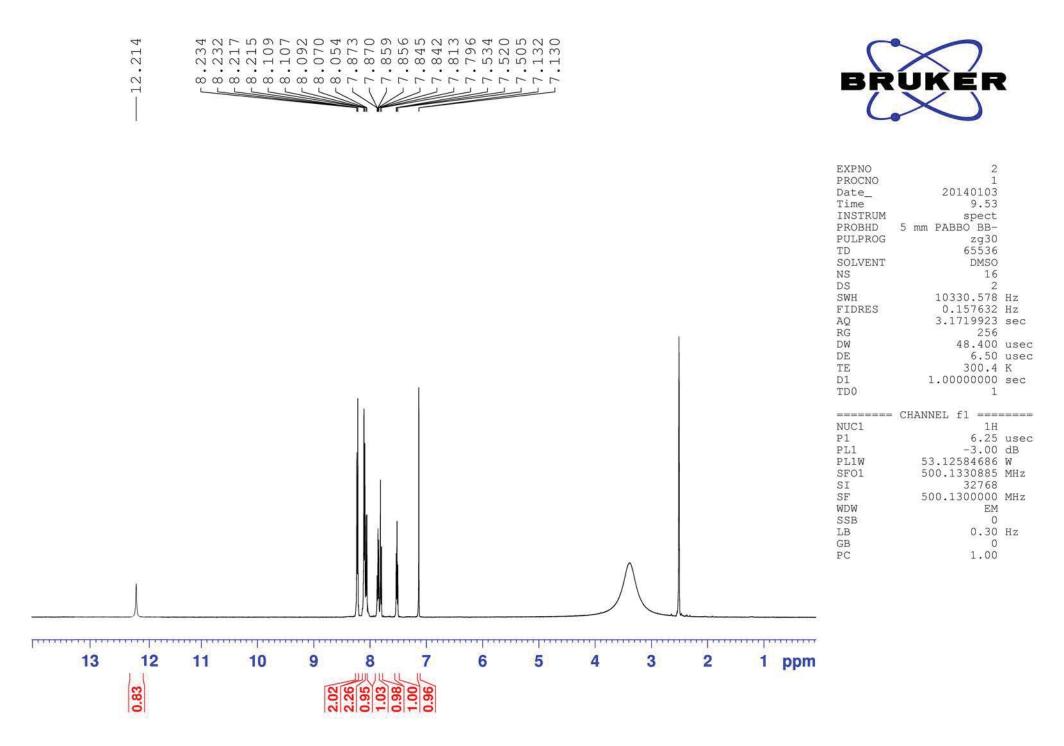
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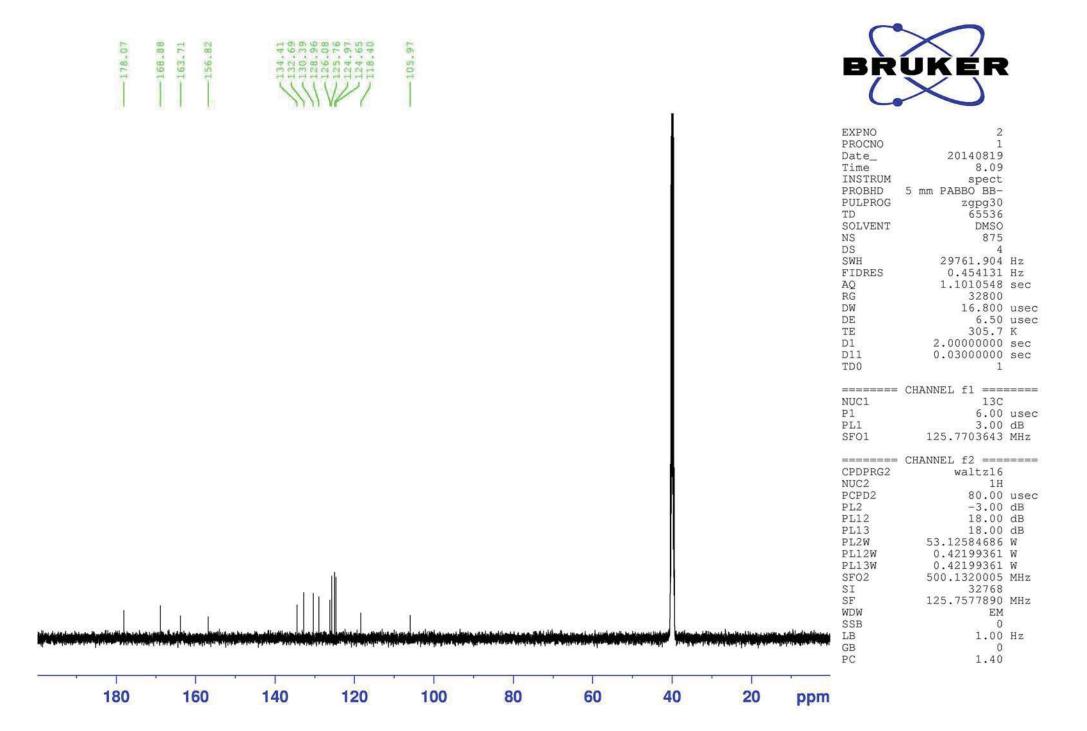
1.00 1.06 1.01 1.01

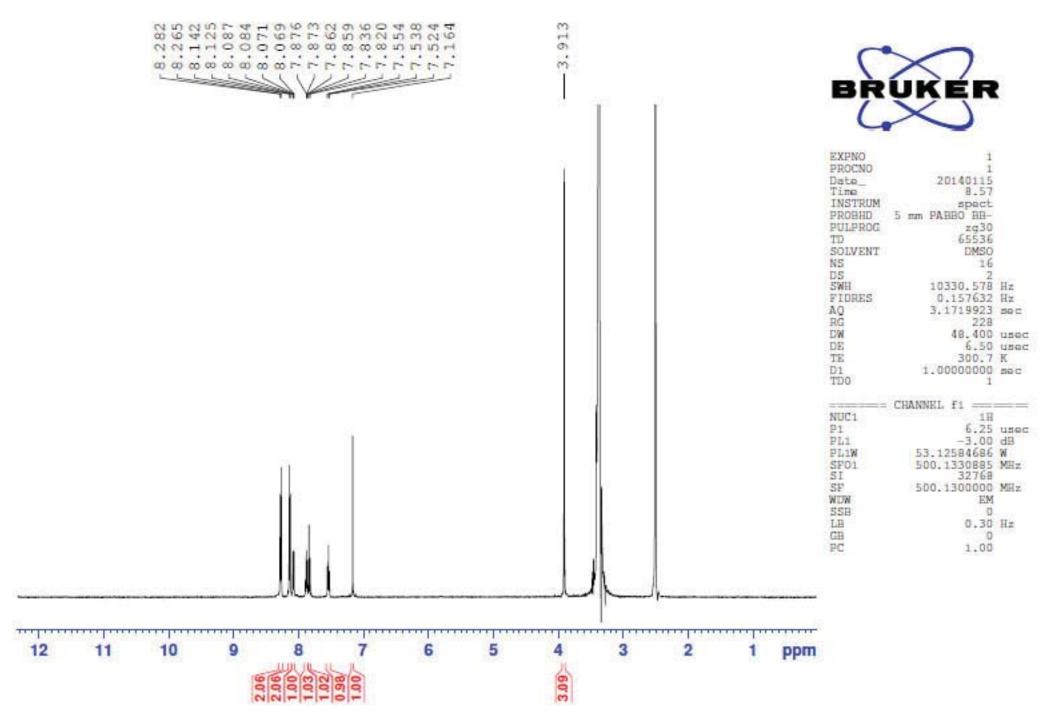
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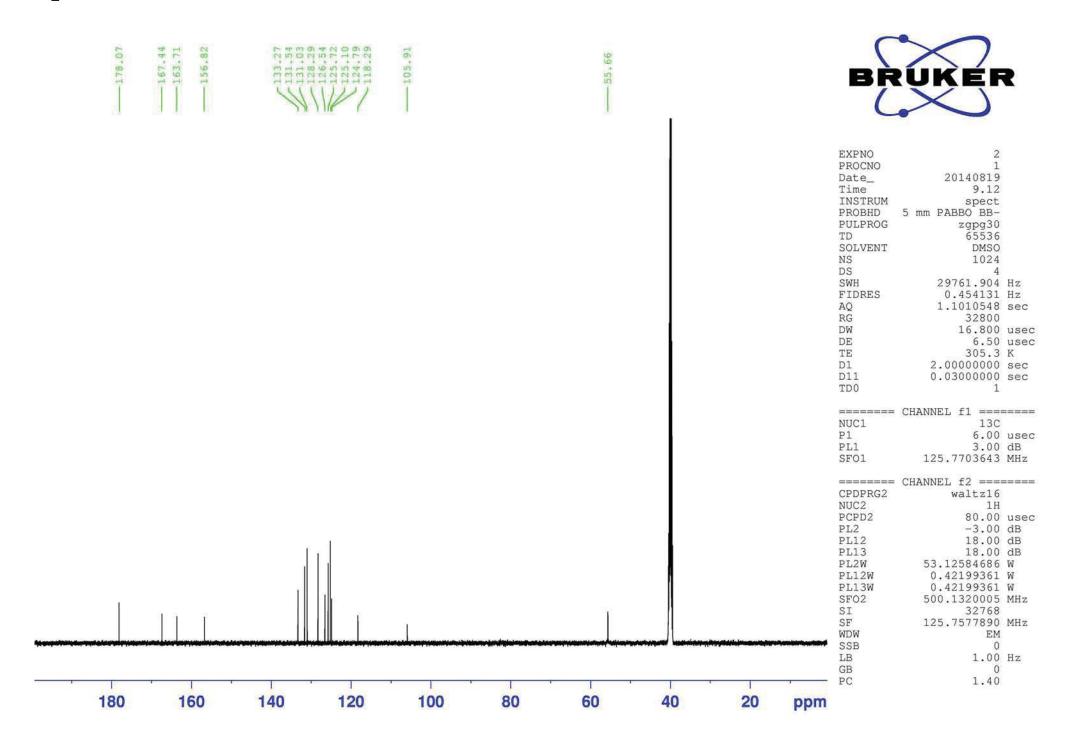
	$\begin{array}{c} 11.152 \\ 8.267 \\ 8.265 \\ 8.265 \\ 8.265 \\ 8.265 \\ 8.265 \\ 8.265 \\ 8.265 \\ 8.265 \\ 8.265 \\ 8.265 \\ 8.020 \\ 8.020 \\ 8.020 \\ 8.020 \\ 8.033 \\ 7.859 \\ 7.7 \\ 7.859 \\ 7.7 \\ 7.593 \\ 7.033 \\ 7.033 \\ 7.019 \\ 7.019 \\ 7.019 \\ 7.019 \\ 8.265 \\ 7.021 \\ 7.019 \\ 17.019 \\ 17.019 \\ 17.019 \\ 17.019 \\ 10.101 \\ 10.$	BRUKER
		EXPNO 3 PROCNO 1 F2 - Acquisition Parameters Date_ 20131227 Time 8.56 INSTRUM spect PROBHD 5 mm PABBO BB- PULPROG zg30 TD 65536 SOLVENT DMSO NS 16 DS 2 SWH 12335.526 Hz FIDRES 0.188225 Hz AQ 2.6564426 sec RG 179.65 DW 40.533 usec DE 6.50 usec TE 294.4 K D1 1.00000000 sec TD0 1 THE 11.67 usec PLW1 25.94199944 W SFO1 600.3037071 MHz F2 - Processing parameters SI 65536 SF 600.3000000 MHz WDW EM SSB 0 UB 0.30.47
13 12	11 10 9 8 7 6 5 4 3 2 1 pp	



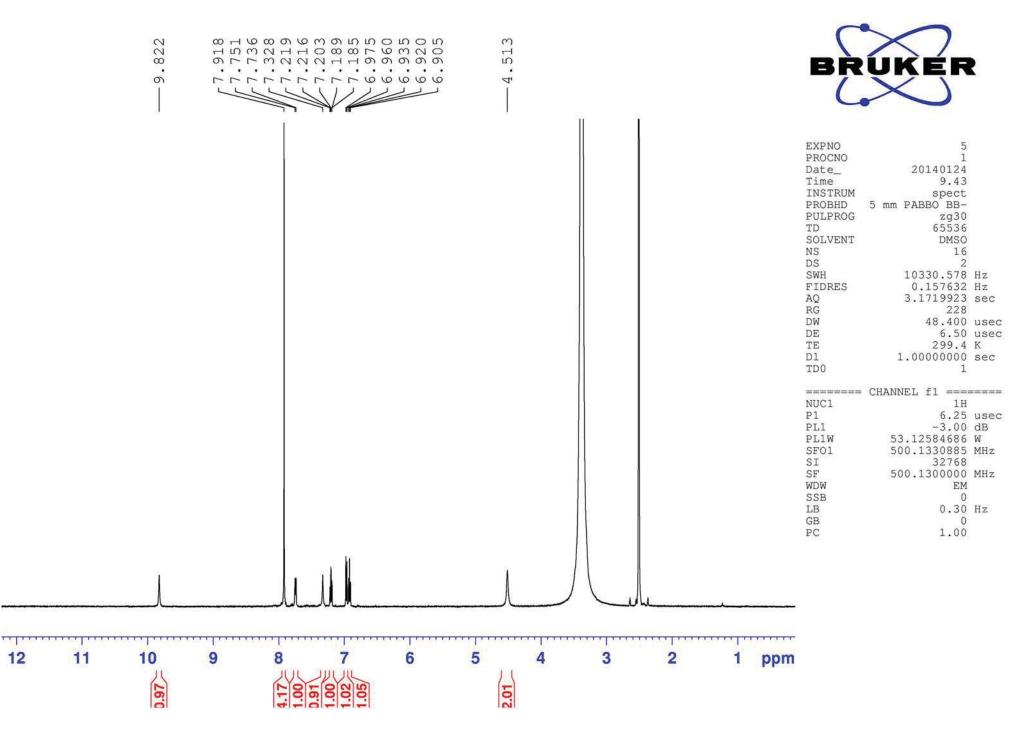


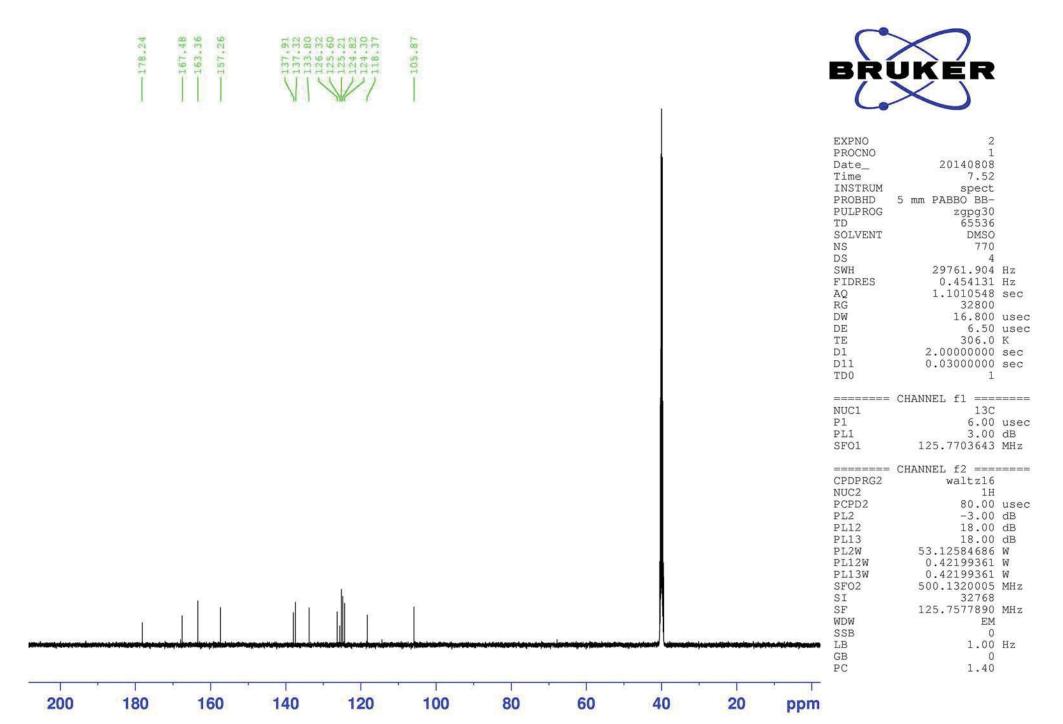


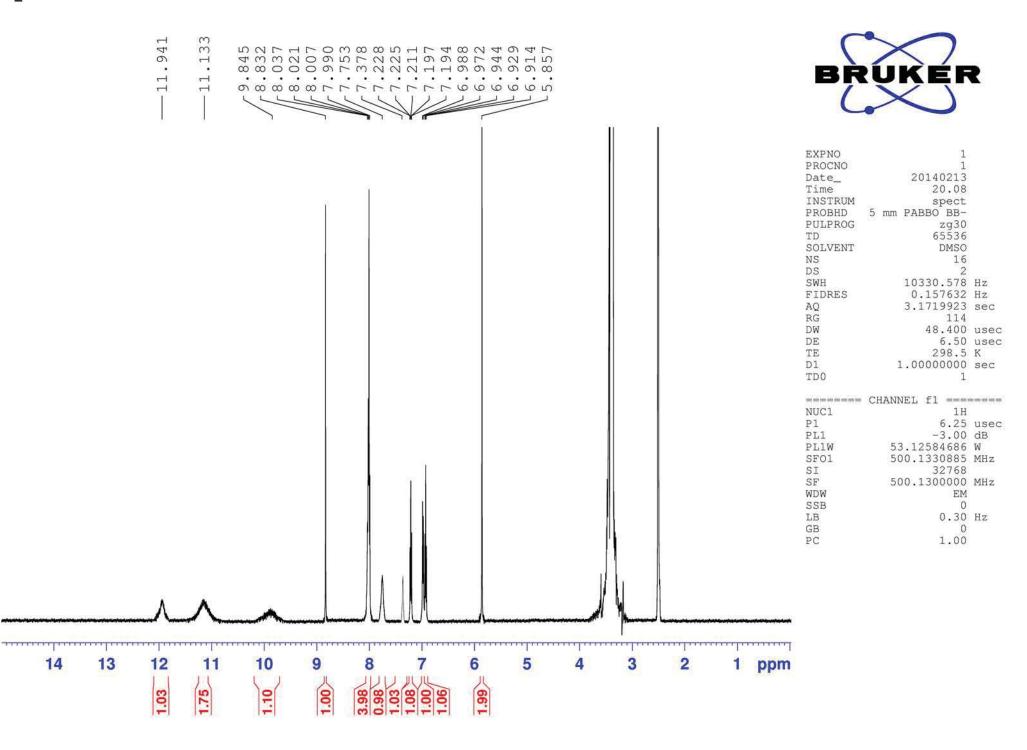


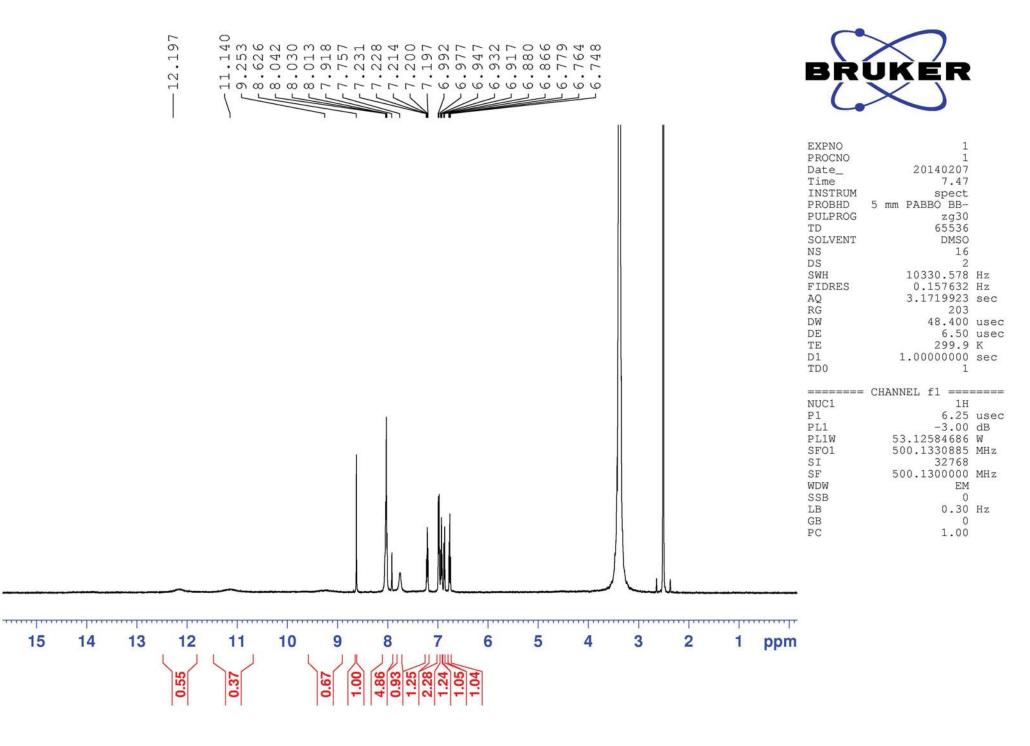


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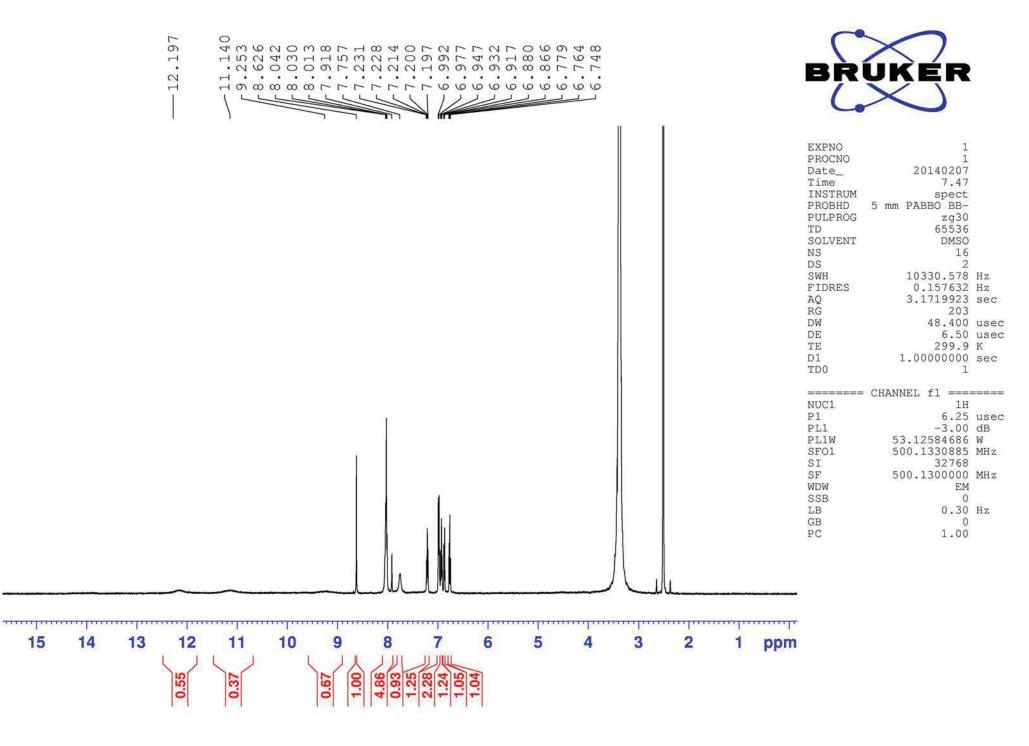








15





-178.36

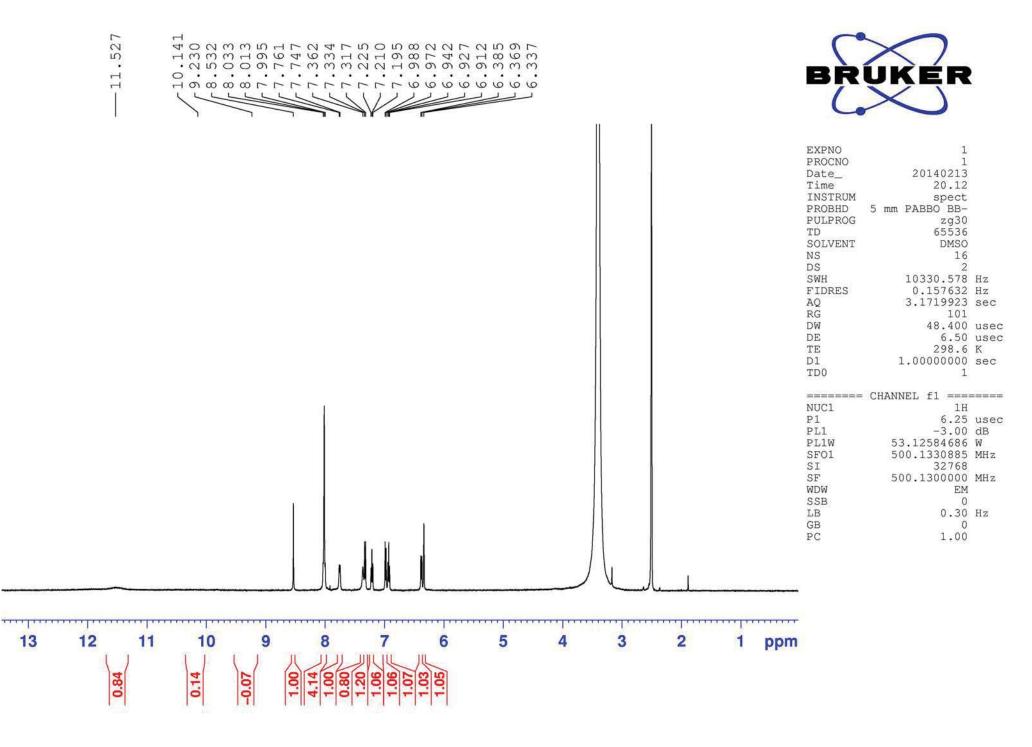
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-155.95

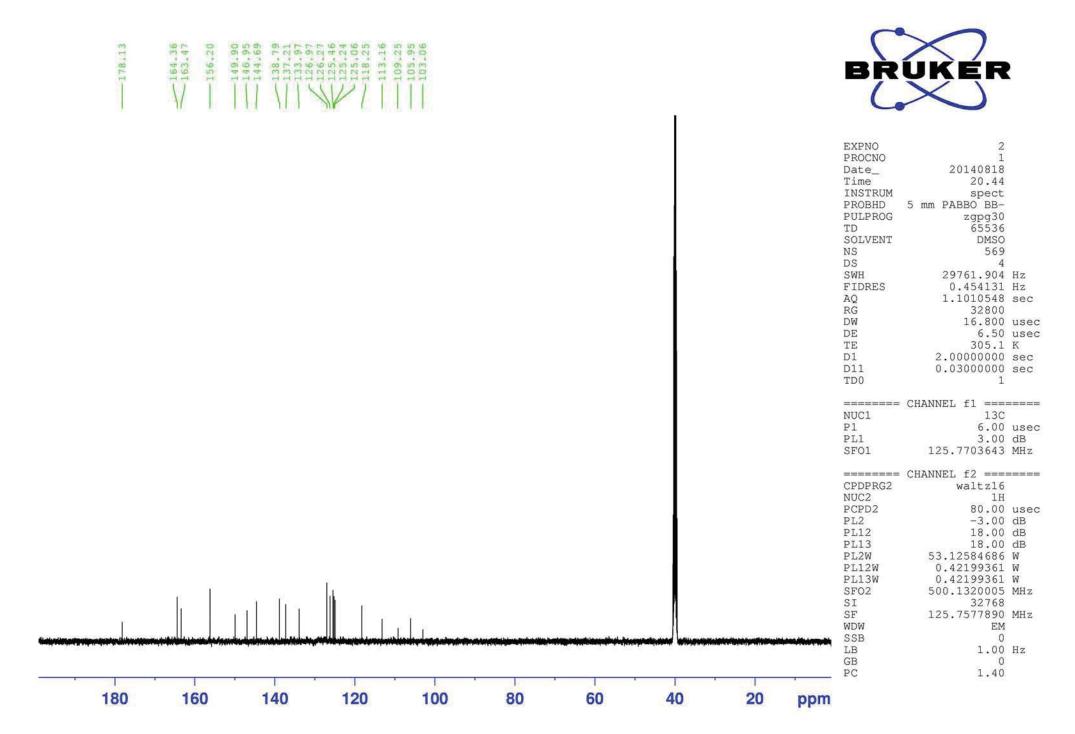
149.35 146.48 144.47 138.72 138.72 137.13 137.13 126.97 125.97 125.03 119.93 119.93 119.95 119.73 119.73

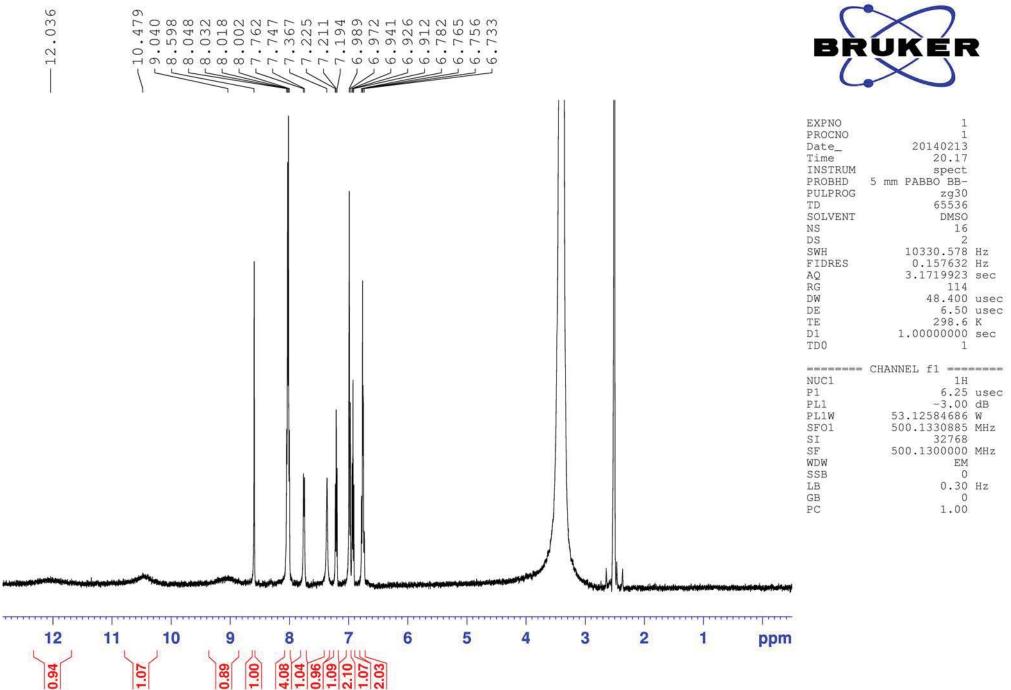


DE 6.50 usec TE 305.5 K D1 2.0000000 sec D11 0.0300000 sec TD0 1 			
AQ 1.1010548 sec RG 32800 DW 16.800 usec DE 6.50 usec TE 305.5 K D1 2.0000000 sec D1 0.03000000 sec D1 0.0300000 sec TD0 1 		PROCNO Date_ Time INSTRUM PROBHD PULPROG TD SOLVENT NS DS SWH	1 20140819 19.42 spect 5 mm PABBO BB- zgpg30 65536 DMSO 859 4 29761.904 Hz
NUC1 13C P1 6.00 used PL1 3.00 dB SF01 125.7703643 MHz ==================================		AQ RG DW DE TE D1 D11 TD0	1.1010548 sec 32800 16.800 usec 6.50 usec 305.5 K 2.00000000 sec 0.03000000 sec 1
PCPD2 80.00 used PL2 -3.00 dB PL12 18.00 dB PL13 18.00 dB PL2W 53.12584666 W PL12W 0.42199361 W PL13W 0.42199361 W PL13W 0.42199361 W SF02 500.1320005 MHz		NUC1 P1 PL1 SF01 CPDPRG2	13C 6.00 used 3.00 dB 125.7703643 MHz CHANNEL f2 ===================================
		PCPD2 PL2 PL12 PL13 PL2W PL12W PL13W	1H 80.00 used -3.00 dB 18.00 dB 18.00 dB 53.12584686 W 0.42199361 W



Compound	7
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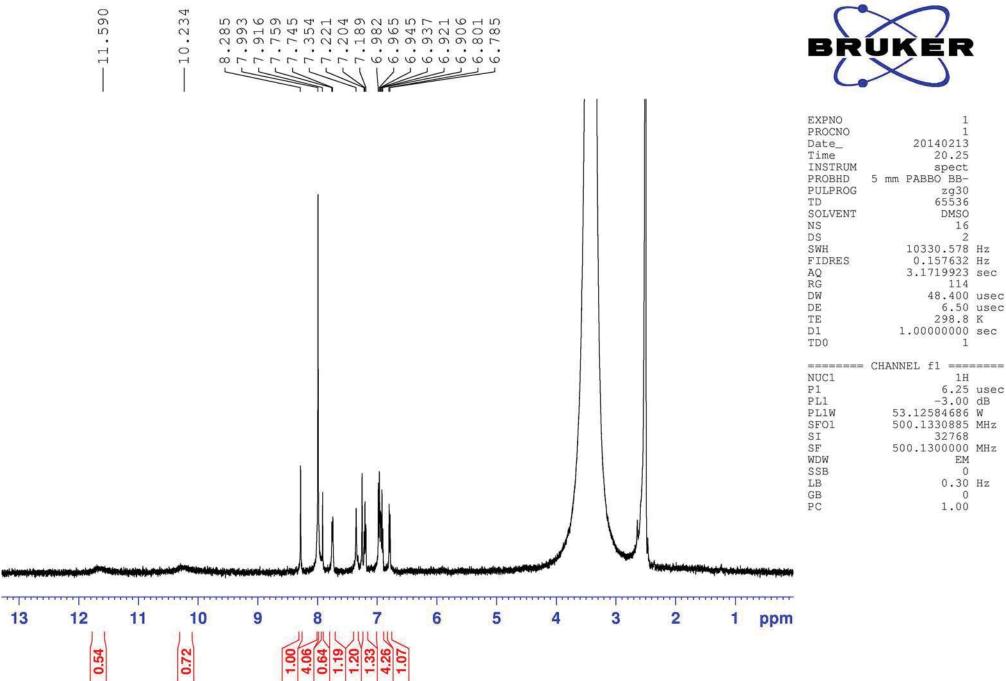


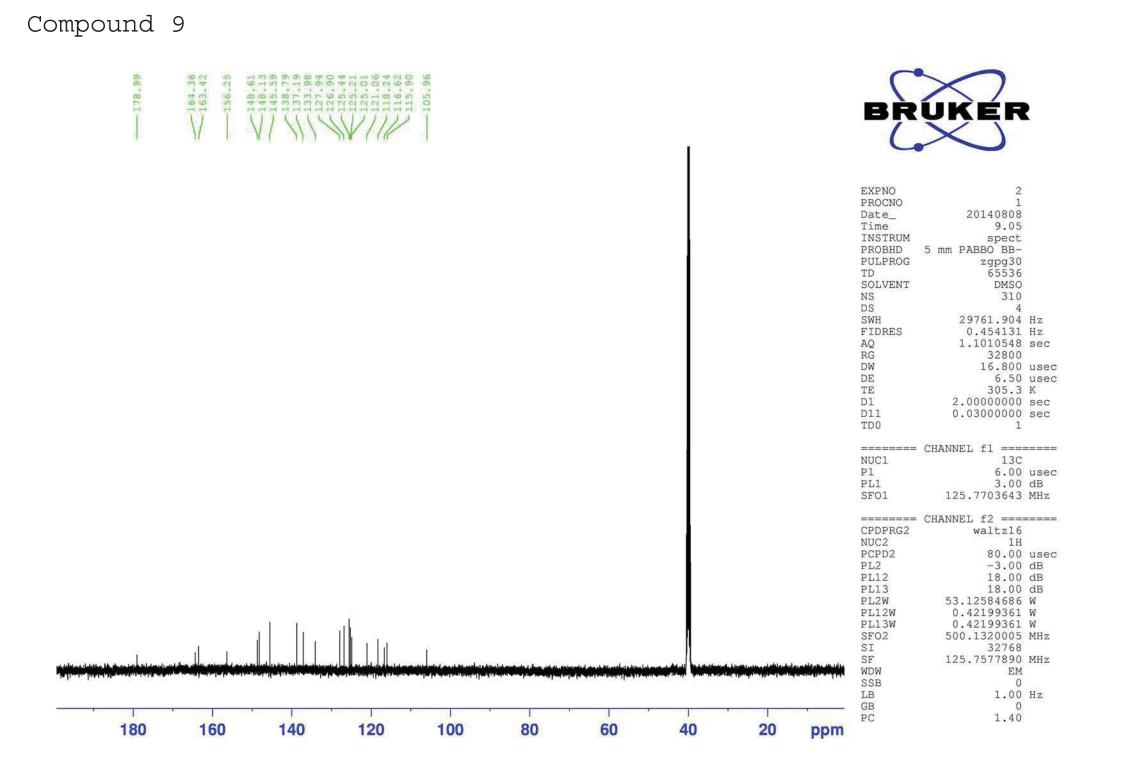
-178.14

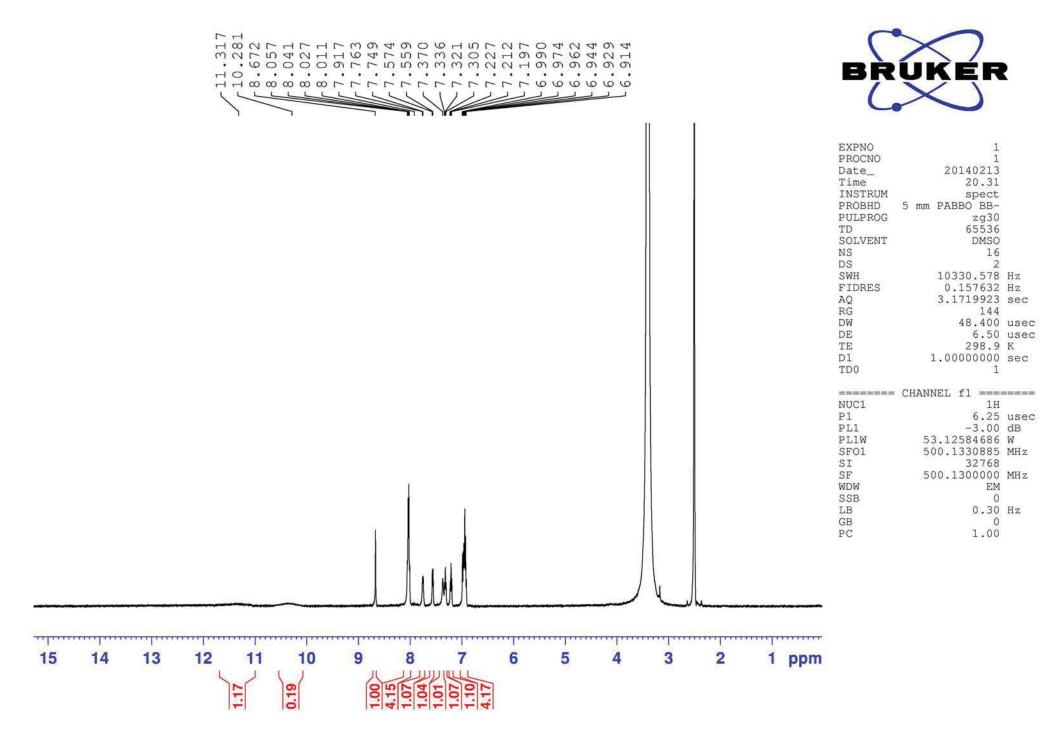
164.36 156.25 156.25 156.25 156.25 138.76 138.70 138.70 138.70 138.70 125.25 12



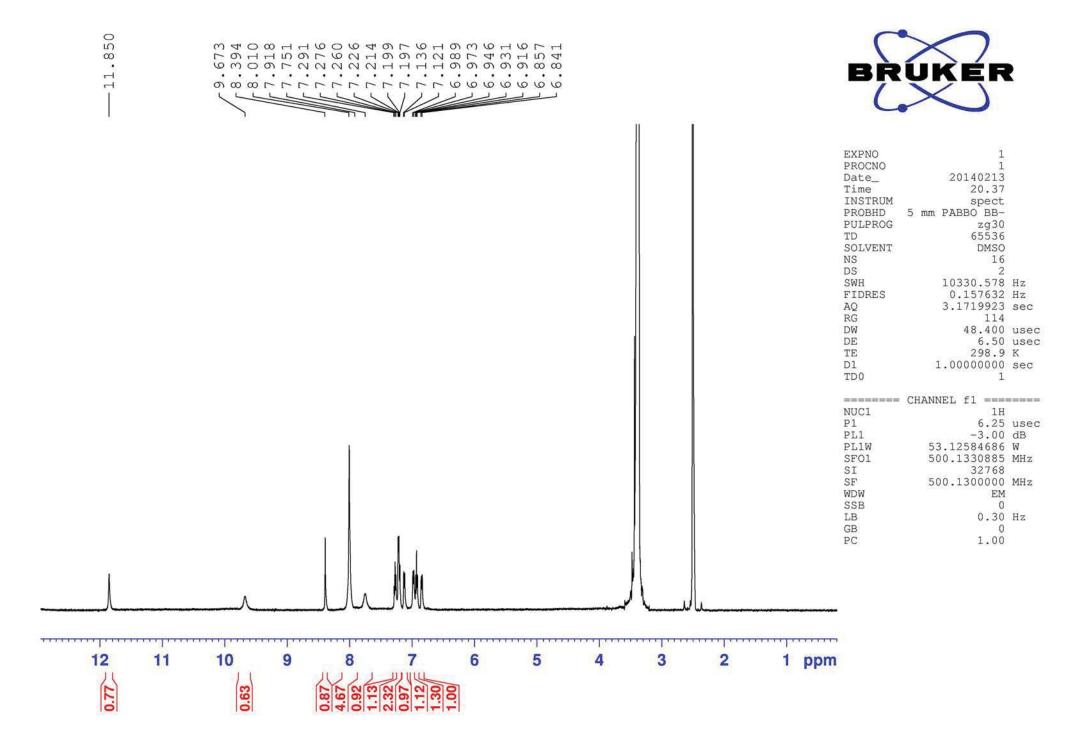
		EXPNO 2 PROCNO 1 Date_ 20140819 Time 10.02 INSTRUM spect PROBHD 5 mm PABBO BB- PULPROG zgpg30 TD 65536 SOLVENT DMSO NS 852 DS 4 SWH 29761.904 FIDRES 0.454131 AQ 1.1010548 RG 32800 DW 16.800 usec DE 6.50 usec TE 305.4 K D1 2.00000000 sec
		D11 0.03000000 sec TD0 1
200 180 160 140 120 100 80 60	40 20 0 ppm	SI 32768 SF 125.7577890 MHz WDW EM SSB 0 LB 1.00 Hz GB 0 PC 1.40



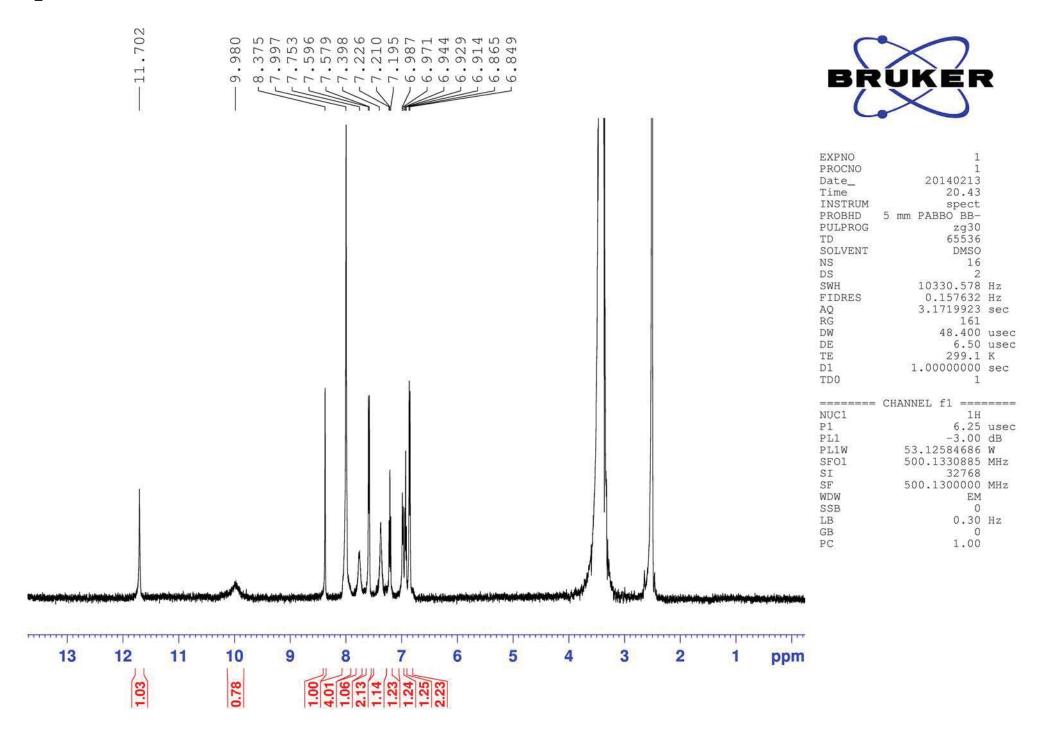




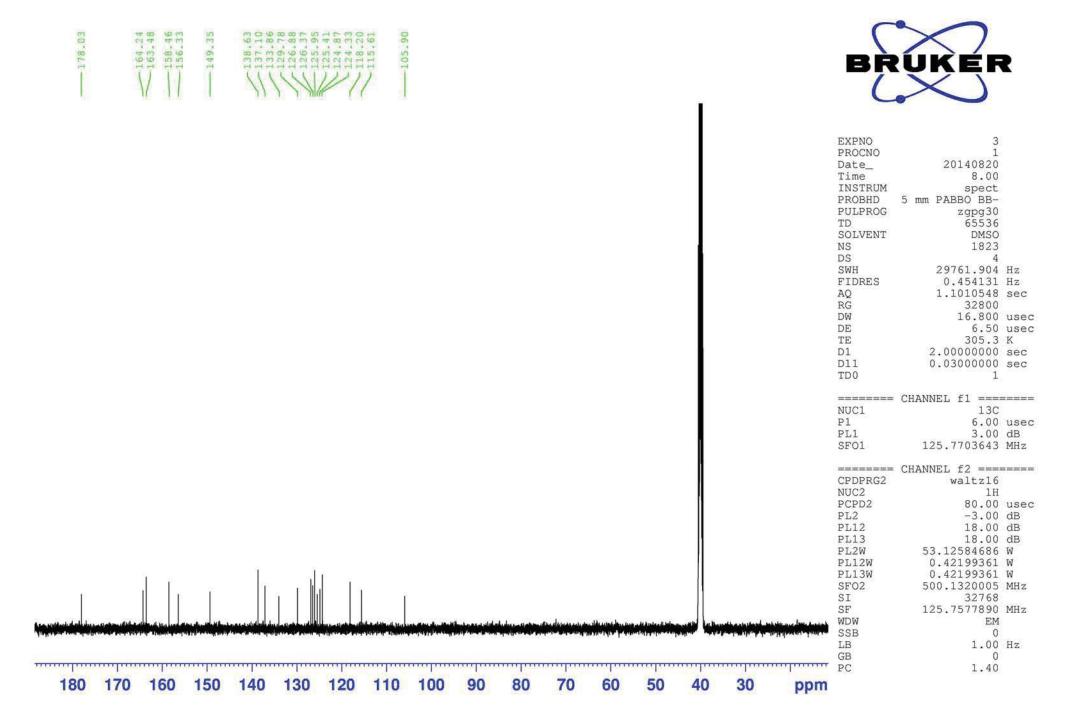
178.18		138.72 133.71 133.97 129.93 126.93 126.93 126.93 125.25 125.05 125.05 125.05 125.05 125.05 118.21 118.21	105.92			T		BR	UKER
								EXPNO PROCNO Date_ Time INSTRUM PROBHD PULPROG TD SOLVENT NS DS SWH FIDRES AQ RG DW DE TE D1 D1 D11 TD0	2 1 20140818 18.38 spect 5 mm PABBO BB- zgpg30 65536 DMSO 975 4 29761.904 Hz 0.454131 Hz 1.1010548 sec 32800 16.800 usec 6.50 usec 304.6 K 2.0000000 sec 0.0300000 sec 1
								NUC1 P1 PL1 SF01	CHANNEL f1 ======= 13C 6.00 usec 3.00 dB 125.7703643 MHz
			l Han na cyclosod a than a star a st	ya ponan wakata ka	n ble tenne tyr standyriten benef	utility with an angle of a	ula ji bala da daga kati kati kati kati kati kati kati kat	CPDPRG2 NUC2 PCPD2 PL2 PL13 PL2W PL12W PL12W PL12W SFO2 SI SF WDW SSB LB	CHANNEL f2 ======= waltz16 1H 80.00 usec -3.00 dB 18.00 dB 53.12584686 W 0.42199361 W 0.42199361 W 500.1320005 MHz 32768 125.7577890 MHz EM 0 1.00 Hz
	70 160 150	140 130 120 110		80 70	60 50	40 30		GB PC	1.40

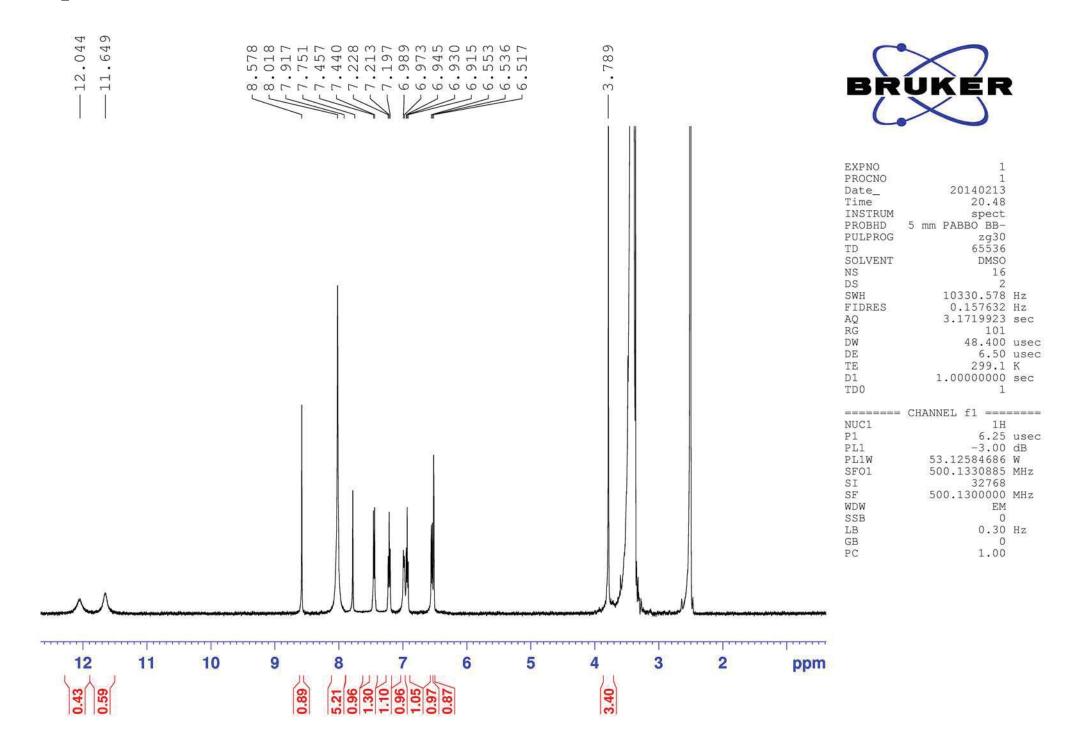


00 74 00	148,71 138,72 137,20 135,88 126,95 126,95 119,35 119,35 119,35 119,35 119,35 119,35 119,35 119,35	BRUKER
		EXPNO 2 PROCNO 1 Date_ 20140818 Time 20.10 INSTRUM spect PROBHD 5 mm PROBHD 5 mm PULPROG zgpg30 TD 65536 SOLVENT DMSO NS 594 DS 4 SWH 29761.904 FIDRES 0.454131 AQ 1.1010548 RG 32800 DW 16.800 DE 6.50 DE 6.50 DE 6.50 DI 2.00000000 SEC TE JUL 2.00000000

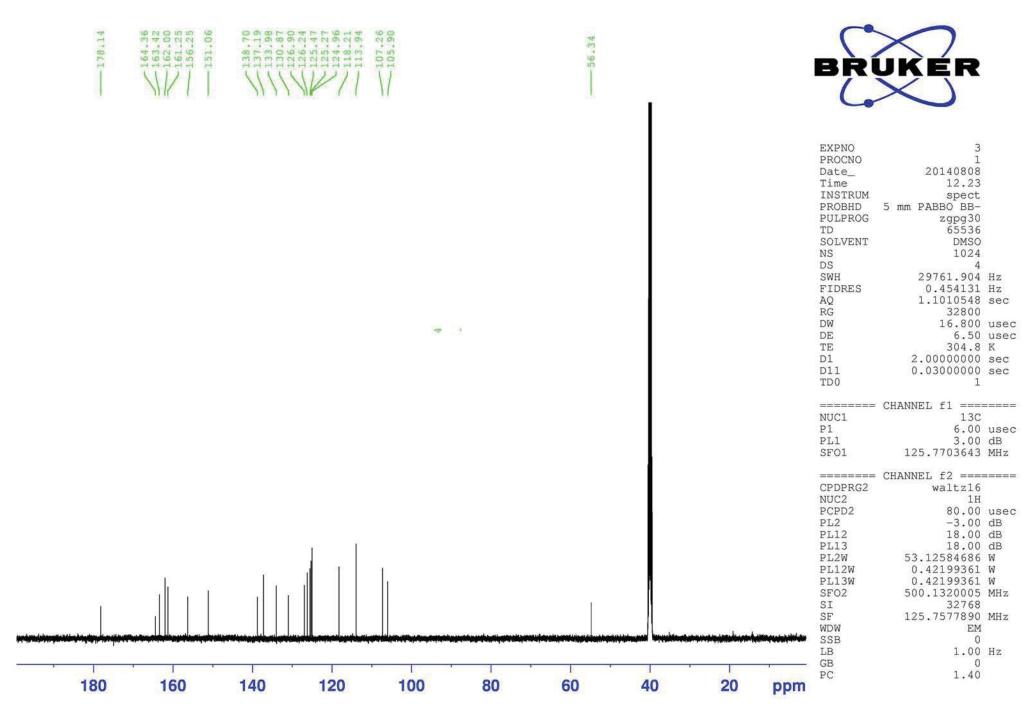


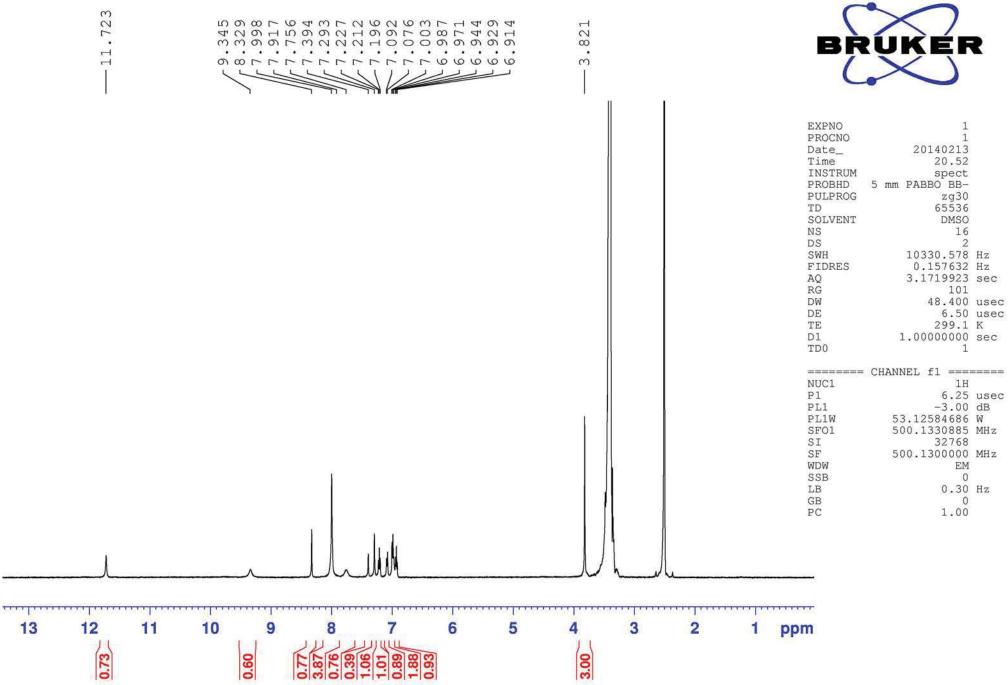
Compound	12
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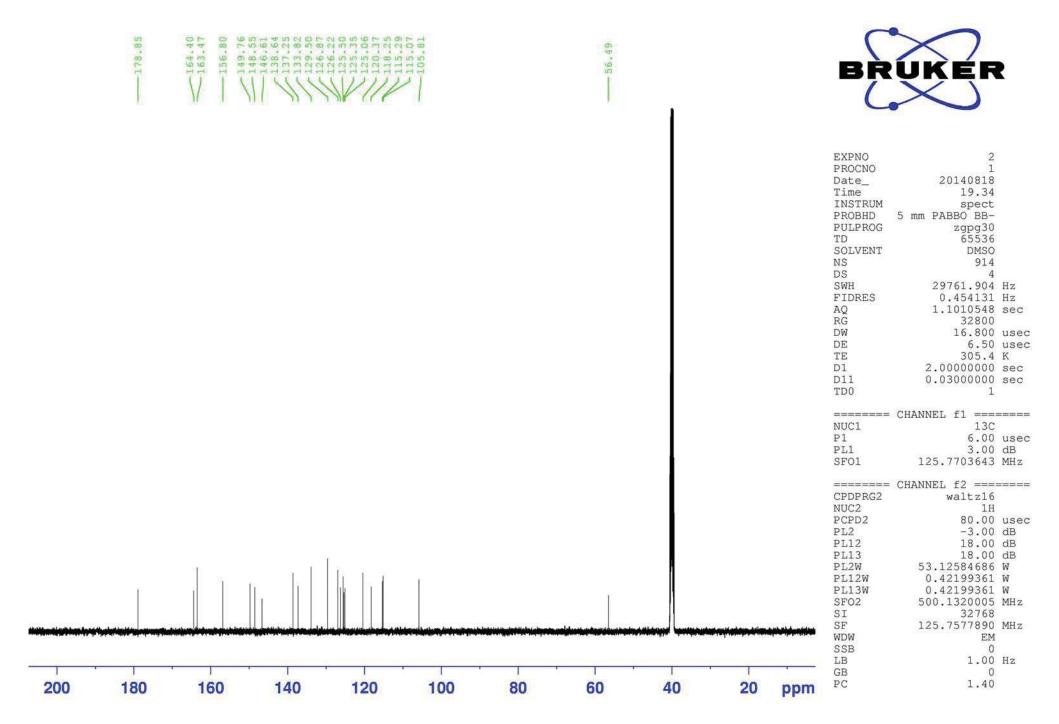


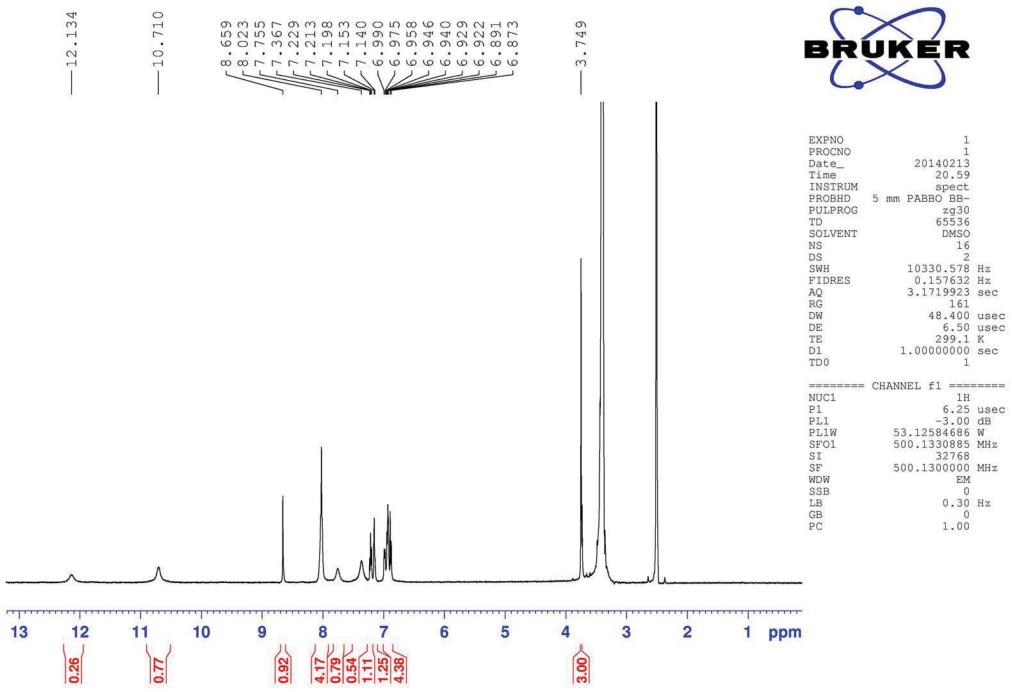
Compound 13



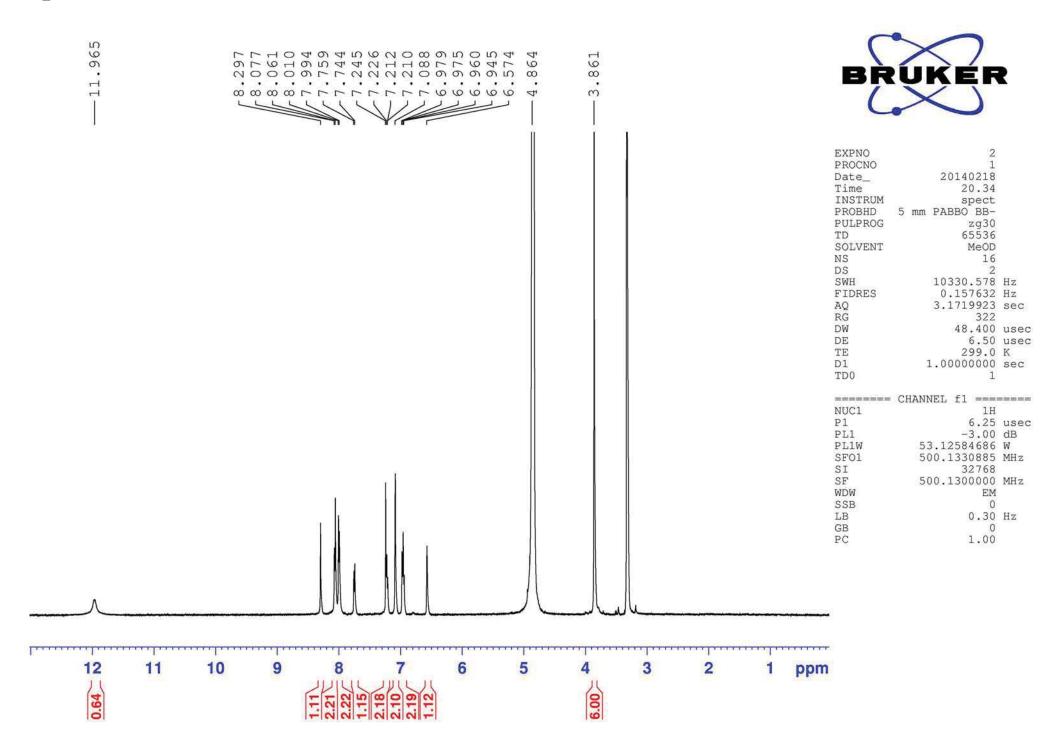


Compound	14
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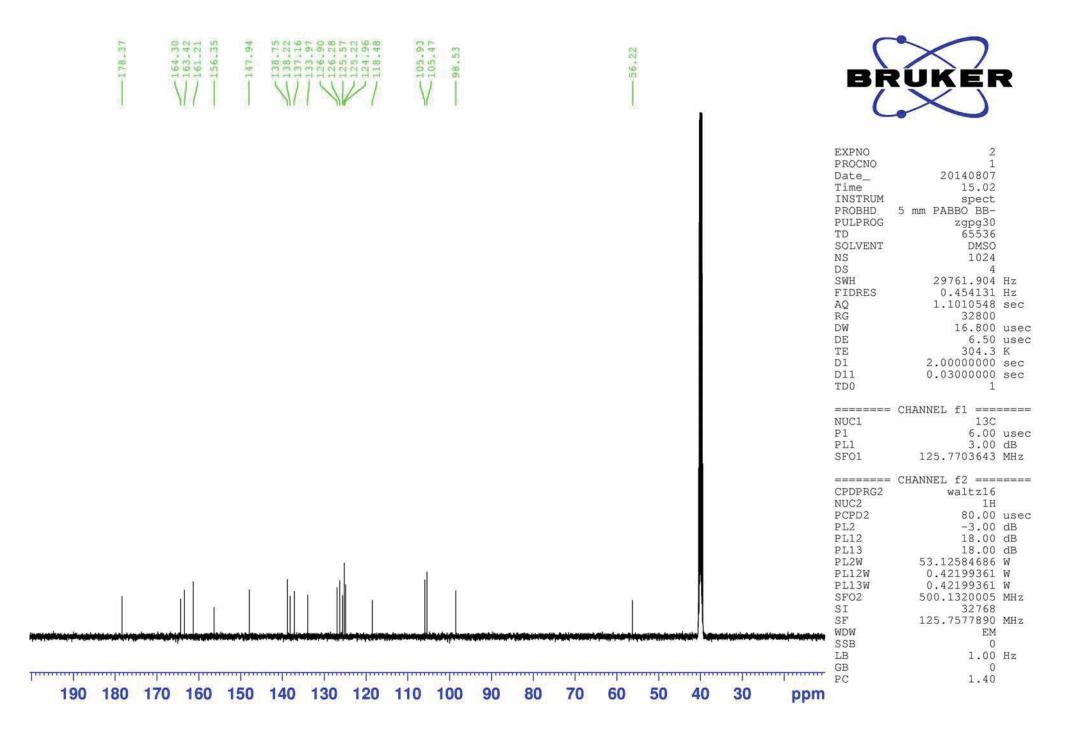


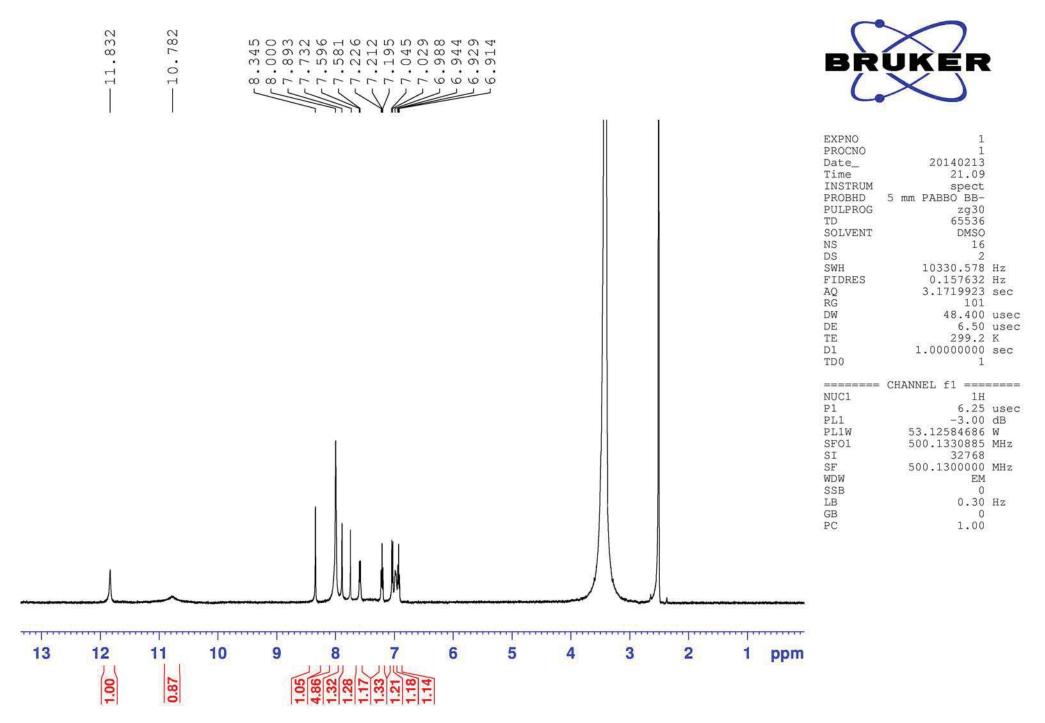


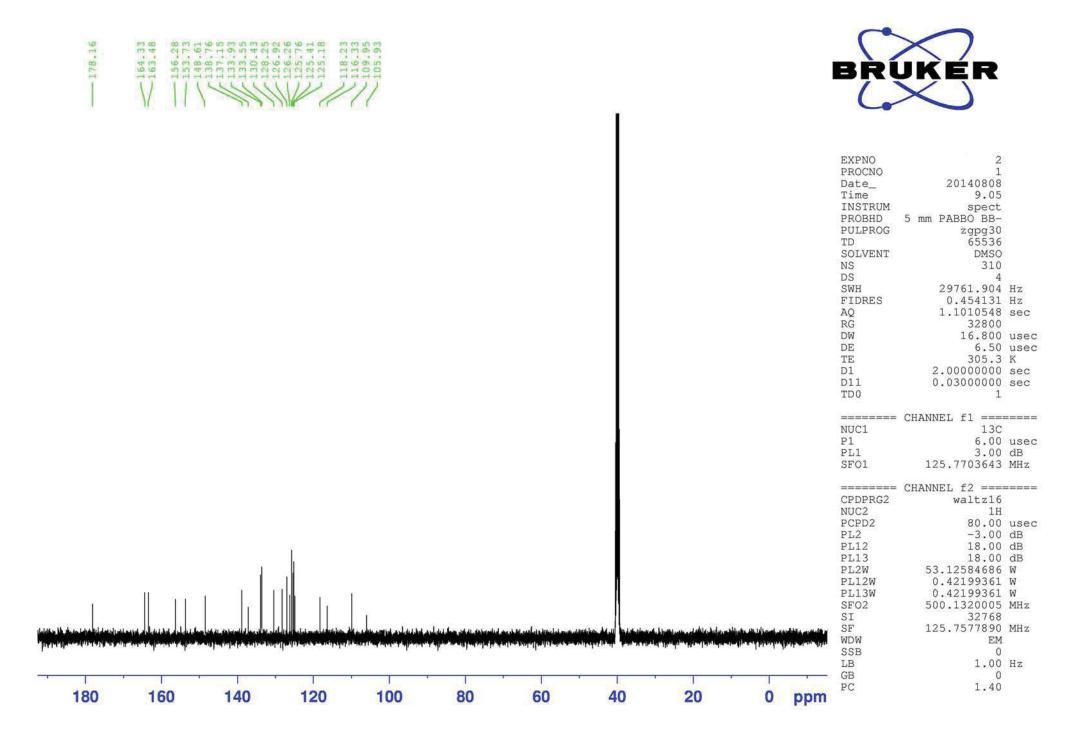
178.26	164.33 163.48 156.28 154.48 154.20 154.20	138.74 137.16 133.95 126.98 126.98 124.90 121.74 118.23 118.23 118.23 116.54 116.54 116.53 112.82	200	BRUKER
				EXPNO 1 PROCNO 1 Date_ 20140808 Time 12.23 INSTRUM spect PROBHD 5 mm PABBO BB- PULPROG zgpg30 TD 65536 SOLVENT DMSO NS 1024 DS 4 SWH 29761.904 Hz FIDRES 0.454131 Hz AQ 1.1010548 sec RG 32800 DW 16.800 usec DE 6.50 usec TE 304.8 K D1 2.00000000 sec D11 0.03000000 sec D11 0.03000000 sec D11 13C P1 6.00 usec P1 6.00 usec
				SF01 125.7703643 MHz

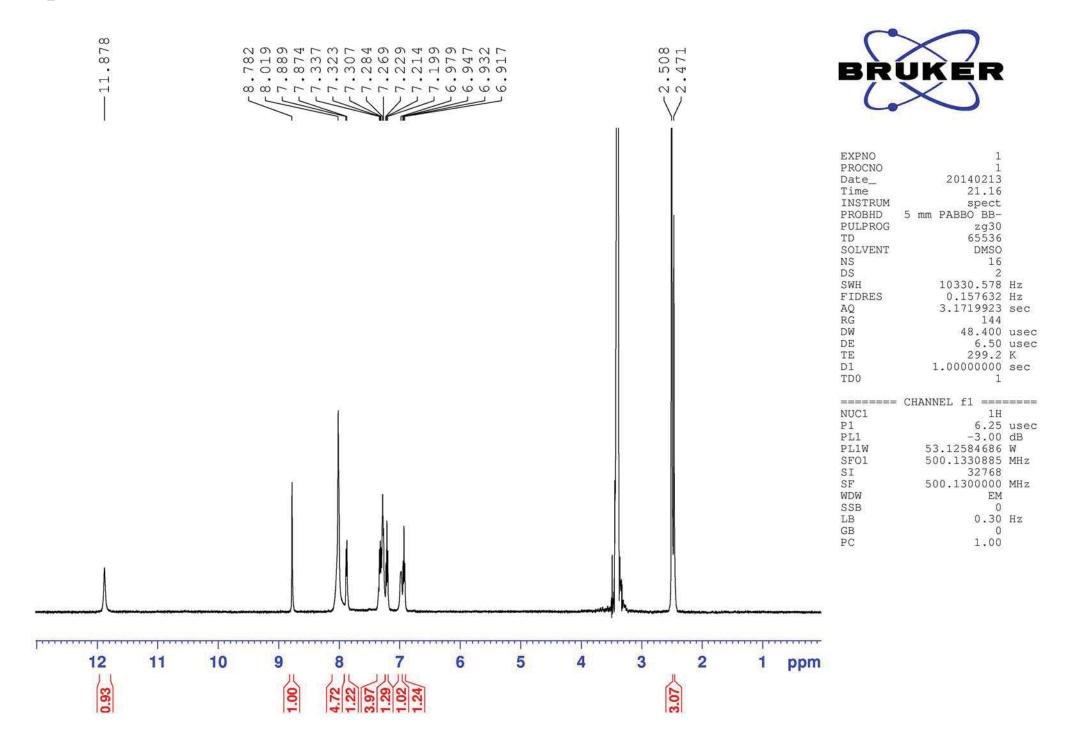


Compound	16
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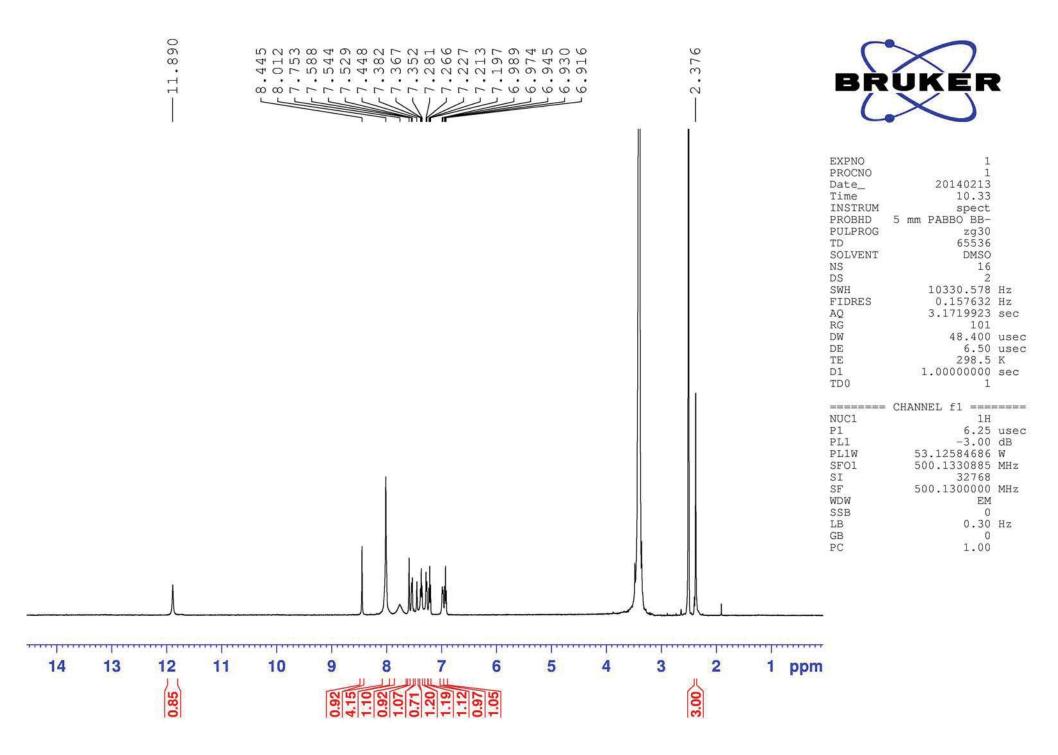




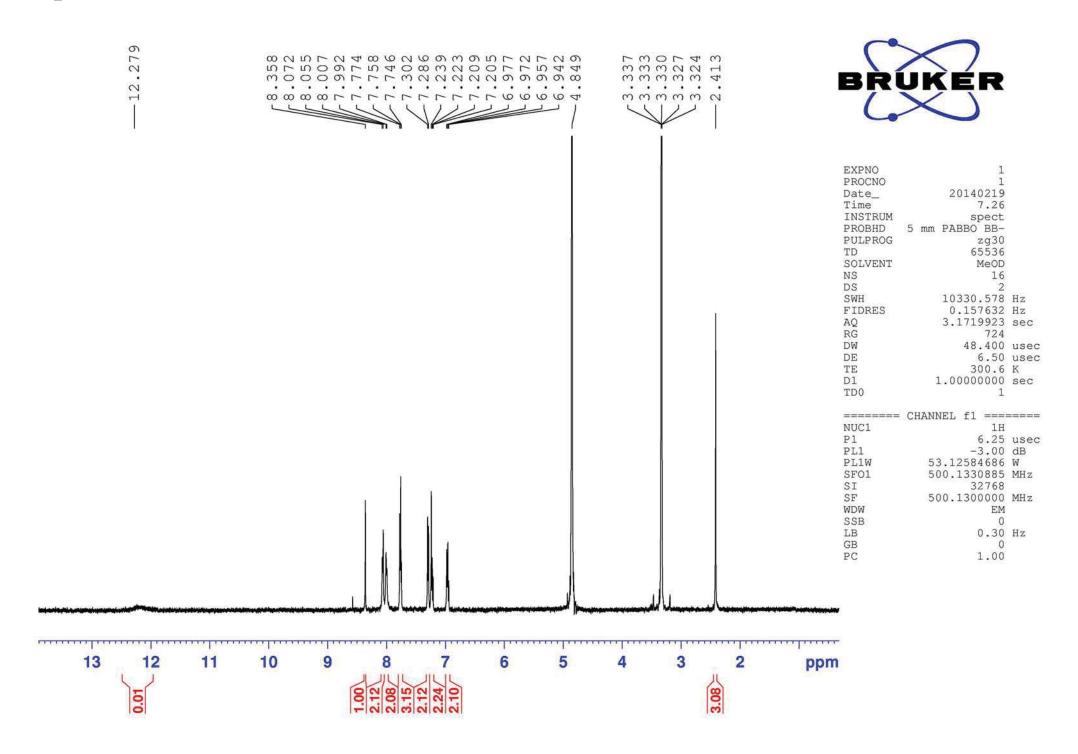


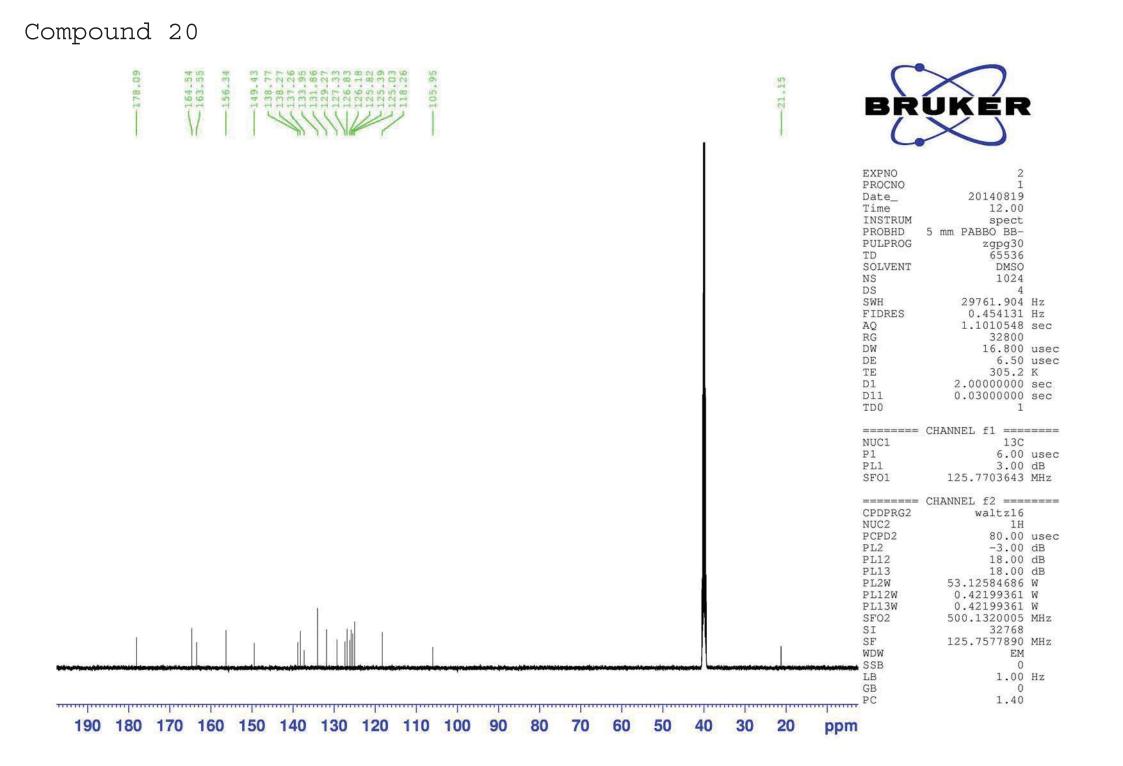


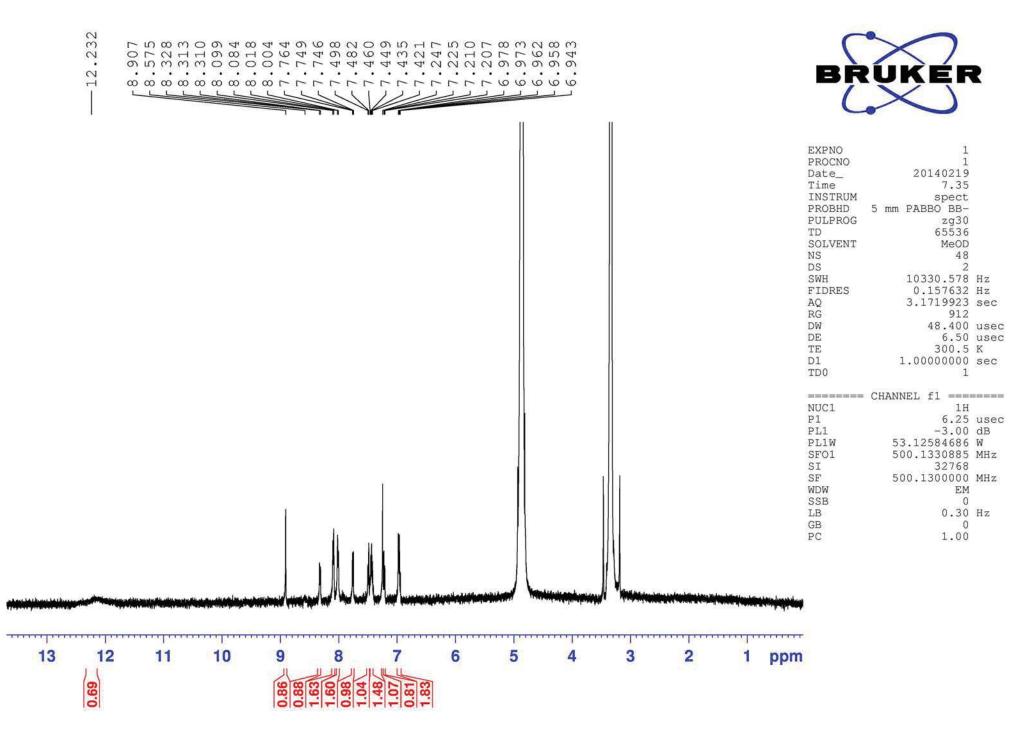
178.18 164.33 164.33 164.33 165.33 135.42 137.16 133.53 133.53 133.53 126.91 126.91 126.63 126.63 126.63 126.63 125.15 125.15 105.98	BRUKER
	EXPNO 2 PROCNO 1 Date_ 20140819 Time 17.58 INSTRUM spect PROBHD 5 mm PABBO BB- PULPROG zgpg30 TD 65536 SOLVENT DMSO NS 1837 DS 4 SWH 29761.904 Hz FIDRES 0.454131 Hz AQ 1.1010548 sec RG 32800 DW 16.800 usec DE 6.50 usec TE 304.8 K D1 2.0000000 sec D11 0.03000000 sec TD0 1
190 180 170 160 150 140 130 120 110 100 90 80 70	истрати water and an and an and an and an



Compound 1	9	
178.32	164.38 163.45 163.45 156.29 137.97 137.97 137.97 126.07 128.68 128.68 128.68 128.68 128.69 125.55 126.07 126.07 126.07 126.07 126.07 126.07 126.07 126.07 126.07 127.25 127.25 126.07 127.25 126.07 127.25 127.55	BRUKER
		EXPNO 2 PROCNO 1 Date_ 20140819 Time 20.23 INSTRUM spect PROBHD 5 mm SUP 10MSO NS 724 DS 4 SWH 29761.904 AQ 1.1010548 Sec RG G 32800 DW 16.800 DE 6.50 DI 2.00000000 Sec 0.11
		Example CHANNEL f1 f1 <thf1< th=""> <thf1< th=""> <thf1< th=""></thf1<></thf1<></thf1<>
		CHANNEL f2 f2 CPDPRG2 waltz16 NUC2 1H PCPD2 80.00 usec PL2 -3.00 dB PL12 18.00 dB PL13 18.00 dB PL2W 53.12584686 W PL12W 0.42199361 W PL13W 0.42199361 W SFO2 500.1320005 MHz SI 32768 SF 125.7577890 MHz WDW EM SSB 0 LB 1.00 Hz GB 0
		PC 1.40

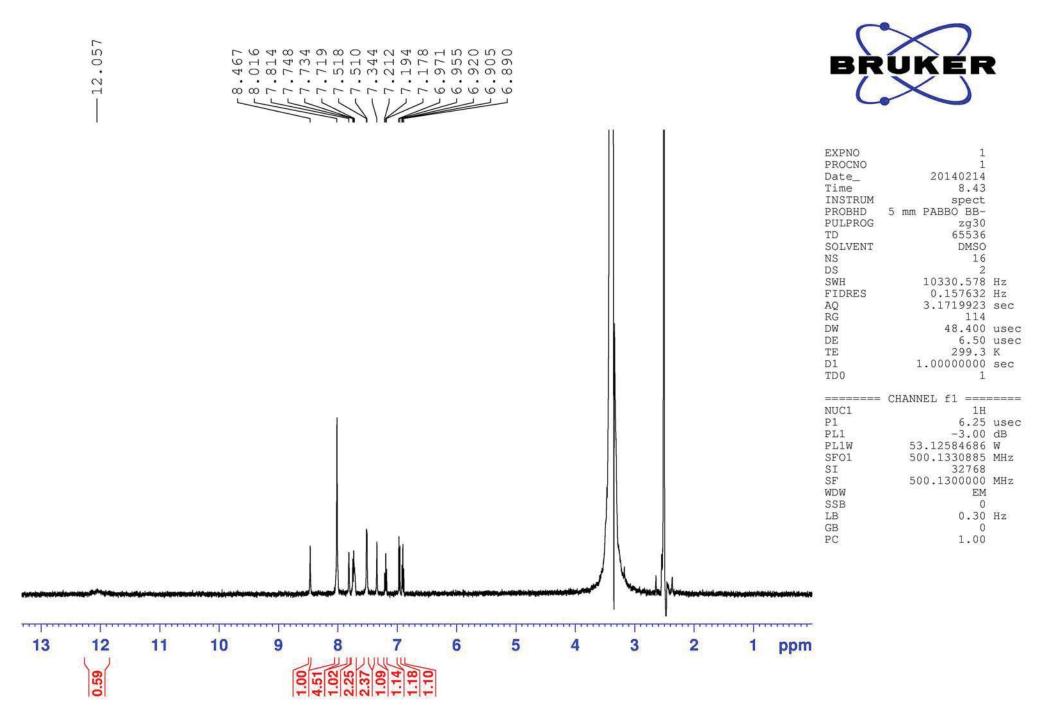




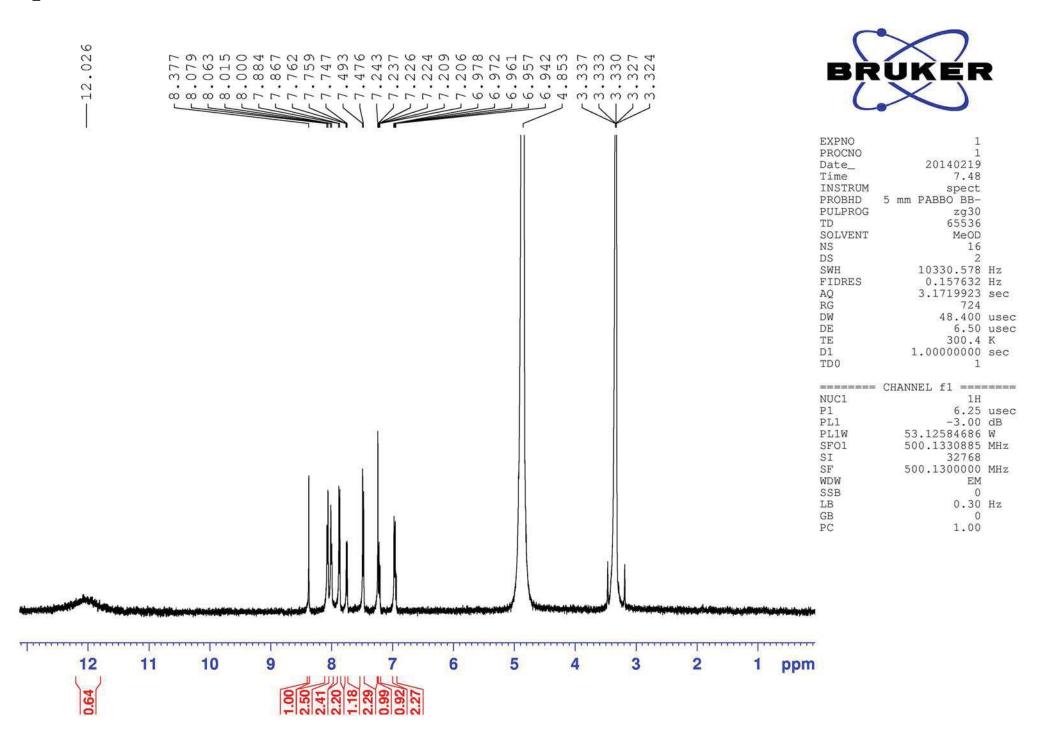


81,871	164.32 163.49 163.49 156.29 138.74 131.35 131.35 131.35 131.35 131.35 132.91 126.90 125.48 125.48 125.48 125.48 125.48 125.48 125.48 125.48 125.48 125.48 125.48 125.48 125.48	BRUKER
		EXPNO 3 PROCNO 1 Date_ 20140820 Time 8.00 INSTRUM spect PROBHD 5 mm PABBO BB- PULPROG zgpg30 TD 65536 SOLVENT DMSO NS 1823 DS 4 SWH 29761.904 FIDRES 0.454131 AQ 1.1010548 RG 32800 DW 16.800 DE 6.50 TE 305.3 D1 2.00000000 D1 0.03000000 TD0 1
		====== CHANNEL f1 NUC1 13C P1 6.00 usec PL1 3.00 dB SFO1 125.7703643 MHz ====== CHANNEL f2 CPDPRG2 waltz16 NUC2 1H PCPD2 80.00 usec PL12 18.00 dB PL13 18.00 dB PL12W 53.12584686 W PL13W 0.42199361 W SFO2 500.1320005 MHz SI 32768 SF 125.7577890 MHz
	70 160 150 140 130 120 110 100 90 80 70 60 50 40 30	PC 1.40





178.24	164.32 163.46 156.24 138.79 137.15 137.15 133.79 133.79 133.79 133.79 133.79 133.79 133.79 133.79 133.79 126.25 127.63 127.45 126.25 127.45 126.25 127.91 126.29 105.91	BR	UKER
		EXPNO PROCNO Date_ Time INSTRUM PROBHD PULPROG TD SOLVENT NS DS SWH FIDRES AQ RG DW DE TE D1 D1 D11	2 1 20140818 18.38 spect 5 mm PABBO BB- zgpg30 65536 DMSO 975 4 29761.904 Hz 0.454131 Hz 1.1010548 sec 32800 16.800 usec 6.50 usec 304.6 K 2.0000000 sec 0.03000000 sec
		NUC1 P1 PL1 SF01	1 = CHANNEL f1 ======= 13C 6.00 usec 3.00 dB 125.7703643 MHz = CHANNEL f2 =======
n transference de sete stat d'audas		CPDPRG2 NUC2 PCPD2 PL12 PL13 PL2W PL13W PL12W PL13W SF02 SI SF WDW	waltz16 1H 80.00 usec -3.00 dB 18.00 dB 18.00 dB 53.12584686 W 0.42199361 W 0.42199361 W 500.1320005 MHz 32768 125.7577890 MHz EM
	70 160 150 140 130 120 110 100 90 80 70 60 50 40 30 20 pp	FC	0 1.00 Hz 0 1.40



200

180

138.78 137.19 137.19 137.19 132.19 122.49 122.88 122.88 125.39 125.40 125.42 125.43

-105.97

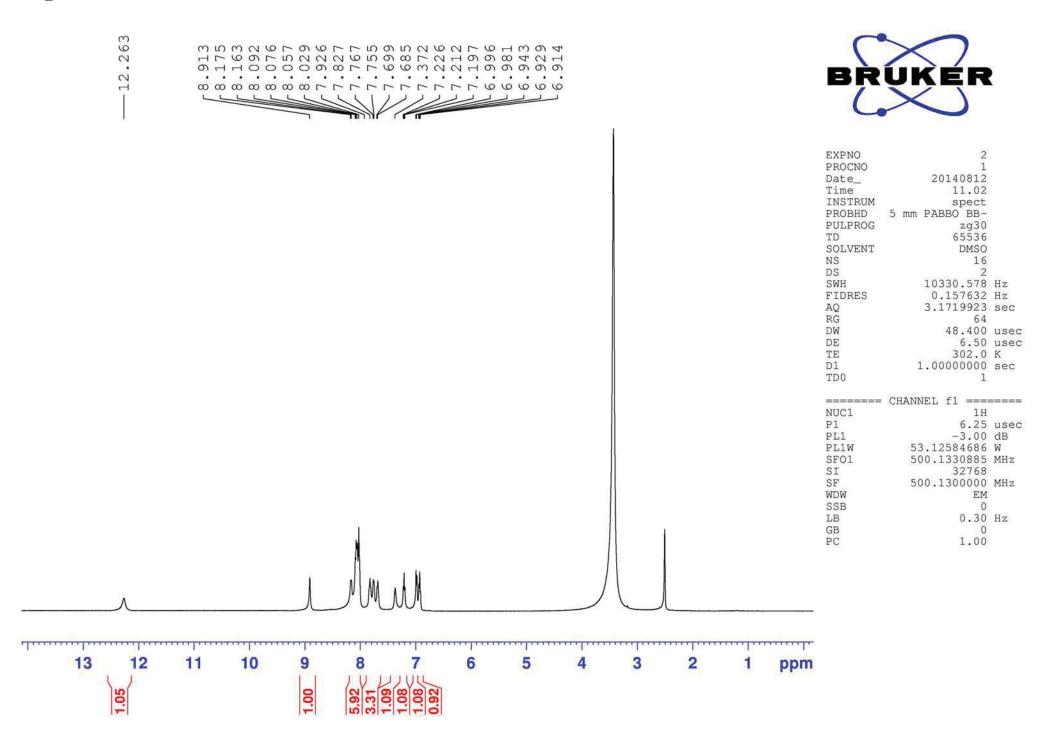
164.39

-156.31

-178.38



	TE 305.0 D1 2.00000000 D11 0.03000000 TD0 1 ======= CHANNEL f1 === NUC1 130	Hz Hz sec usec usec K sec sec
40 ppm	PL1 3.00 SF01 125.7703643 ====== CHANNEL f2 === CPDPRG2 waltz16 NUC2 1F	MHz Usec dB dB dB W W MHz MHz Hz

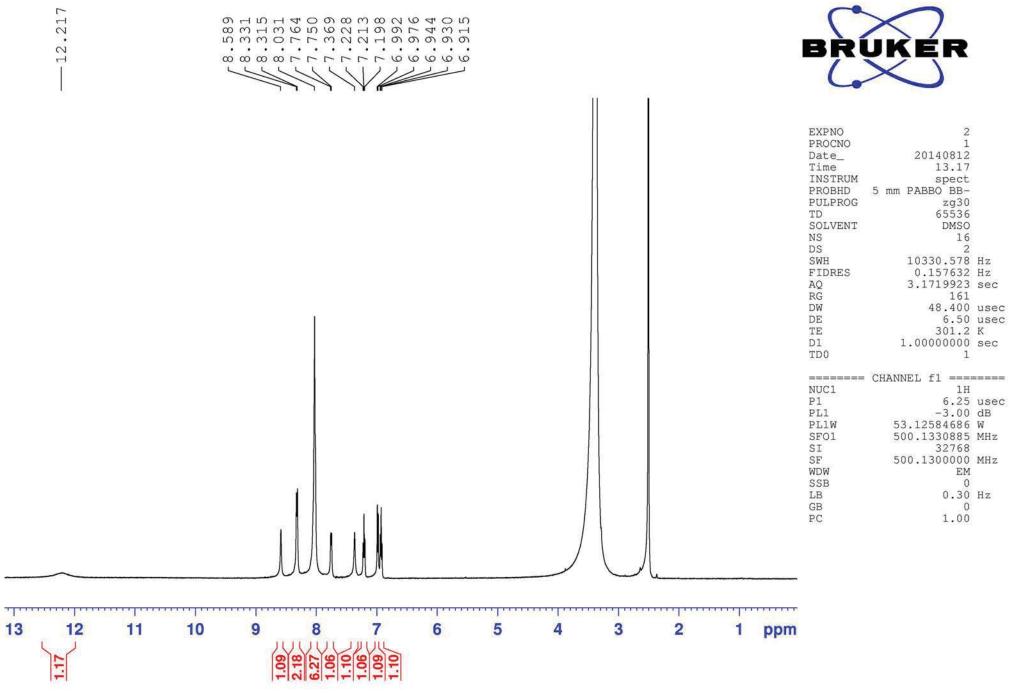


105.95 142.90 138.76 138.76 133.79 133.79 133.79 133.79 133.79 133.79 133.79 133.79 133.79 133.79 133.79 135.69 125.09 125.00 118.28 125.00 118.28	BRUKER
	EXPNO 2 PROCNO 1 Date_ 20140818 Time 21.14 INSTRUM spect PROBHD 5 mm PABBO BB- PULPROG zgpg30 TD 65536 SOLVENT DMSO NS 10386 DS 4 SWH 29761.904 Hz FIDRES 0.454131 Hz AQ 1.1010548 sec RG 32800 DW 16.800 usec DE 6.50 usec TE 304.9 K D1 2.00000000 sec D11 0.03000000 sec TD0 1
	NUC1 13C P1 6.00 used PL1 3.00 dB SF01 125.7703643 MHz ====== CHANNEL f2 ======= CPDPRG2 waltz16 NUC2 1H PCPD2 80.00 used PL2 -3.00 dB PL12 18.00 dB PL13 18.00 dB PL2W 53.12584686 W PL12W 0.42199361 W PL12W 0.42199361 W SF02 500.1320005 MHz SI 32768 SF 125.7577890 MHz WDW EM SSB 0 LB 1.000 Hz GB 0 PC 1.40

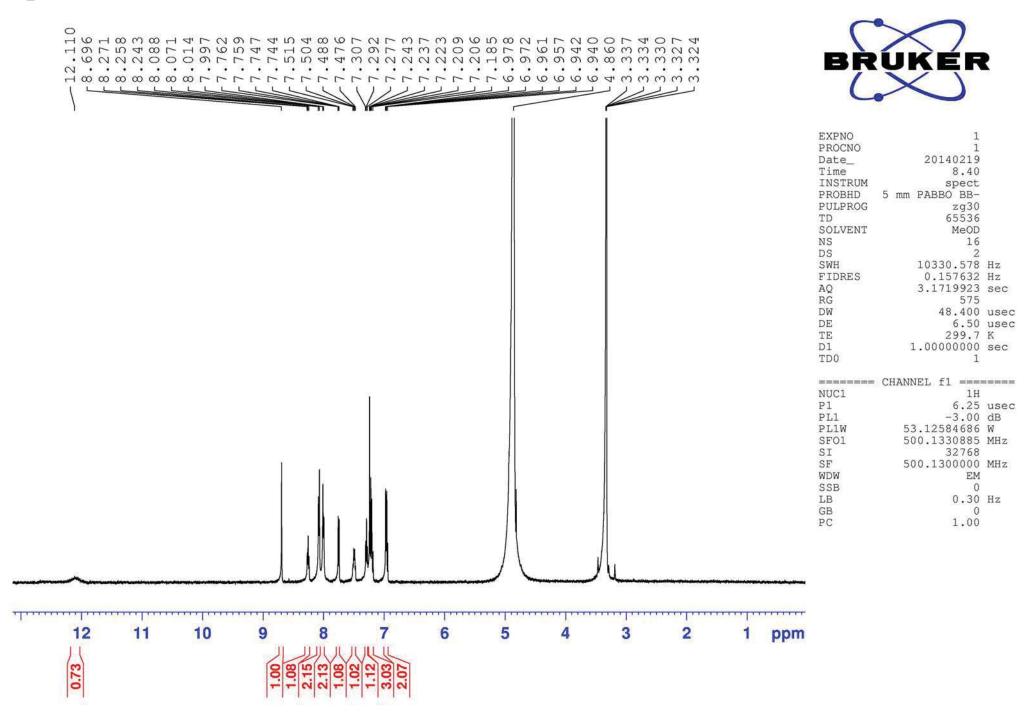
	8.600 8.574 8.574 8.280 8.280 8.285 8.185 7.77 7.753 7.753 7.753 7.753 6.979 6.979 6.979 6.979	. 6 9 1 9 1 9	BRUKER
			EXPNO 2 PROCNO 1 Date_ 20140812 Time 13.11 INSTRUM spect PROBHD 5 mm PABBO BB- PULPROG zg30 TD 65536 SOLVENT DMSO NS 16 DS 2 SWH 10330.578 Hz FIDRES 0.157632 Hz AQ 3.1719923 sec RG 90.5 DW 48.400 usec DE 6.50 usec TE 301.3 K D1 1.00000000 sec TD0 1 PL1 -3.00 dB PL1W 53.12584686 W SFO1 500.1300000 MHz WDW EM SSB 0 LB 0.30 Hz GB 0 PC 1.00
14 13 12 11	9 2 2 8 6 01 0.99 2 8 6 01	5 4 3 2	1 ppm

-178,15 	144.61 147.41 137.75 137.75 137.75 133.98 133.98 133.99 122.99 125.55 12	1	BR	UKER
			EXPNO PROCNO Date_ Time INSTRUM PROBHD PULPROG TD SOLVENT NS DS SWH FIDRES AQ RG DW DE TE D1 D11	2 1 20140818 20.10 spect 5 mm PABBO BB- zgpg30 65536 DMSO 594 4 29761.904 Hz 0.454131 Hz 1.1010548 sec 32800 16.800 usec 6.50 usec 305.3 K 2.00000000 sec 0.03000000 sec
			NUC1 P1 PL1 SF01 CPDPRG2 NUC2 PCPD2 PL2	1 CHANNEL f1 ======= 13C 6.00 usec 3.00 dB 125.7703643 MHz CHANNEL f2 ======= waltz16 1H 80.00 usec -3.00 dB
			PL12 PL13 PL2W PL12W PL13W SF02 SI SF WDW SSB LB GB GB F PC	18.00 dB 18.00 dB 53.12584686 W 0.42199361 W 0.42199361 W 500.1320005 MHz 32768 125.7577890 MHz EM 0 1.00 Hz 0 1.40

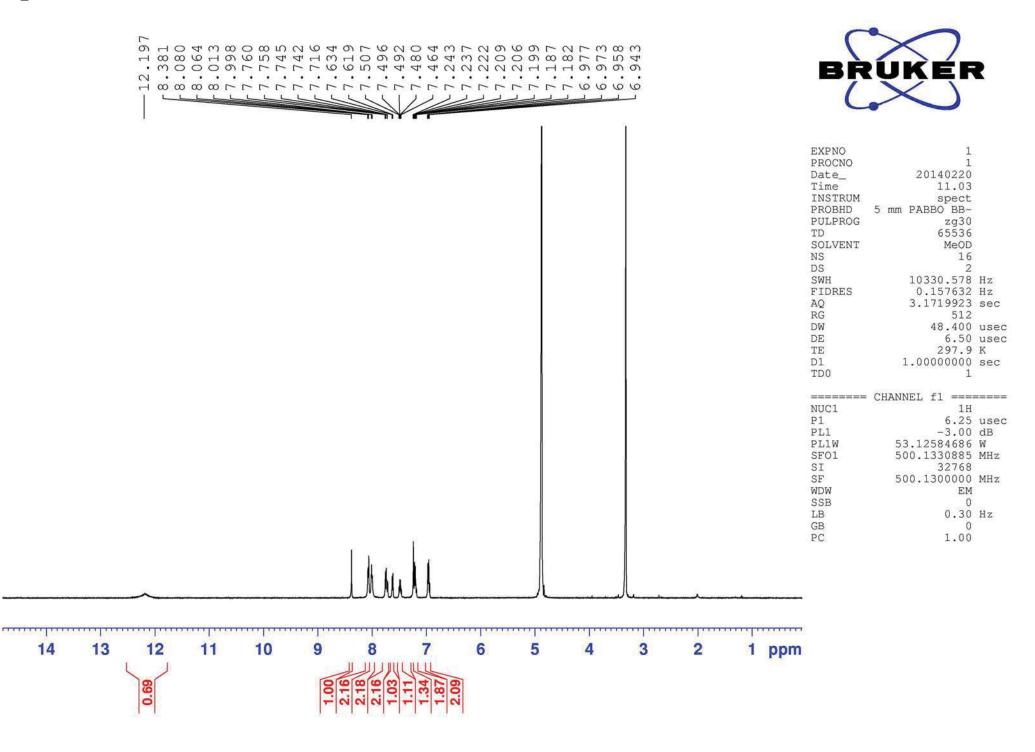




178.18	164.34 156.29 156.29 148.08 148.08 138.76 137.16 125.49 125.50	BRUKER
		EXPNO 2 PROCNO 1 Date_ 20140818 Time 20.44 INSTRUM spect PROBHD 5 mm PROBHD 65536 SOLVENT DMSO NS 569 DS 4 SWH 29761.904 FIDRES 0.454131 AQ 1.1010548 DE 6.50 DE 6.50 DI 2.00000000 Sec 0.11 D1 2.00000000 D1 <

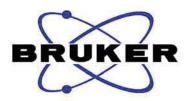


178.32 164.31 163.44 163.44 163.44 161.61 159.35 137.19 137.19 137.19 137.19 137.19 137.19 137.19 137.19 137.19 137.19 137.19 137.19 137.19 137.19 137.19 137.19 125.45	BRUKER
	EXPNO 2 PROCNO 1 Date_ 20140807 Time 10.01 INSTRUM spect PROBHD 5 mm PABBO BB- PULPROG zgpg30 TD 65536 SOLVENT DMSO NS 523 DS 4 SWH 29761.904 Hz FIDRES 0.454131 Hz AQ 1.1010548 sec RG 32800 DW 16.800 use DE 6.50 use TE 305.6 K D1 2.0000000 sec D11 0.03000000 sec
	TD0 1 ====== CHANNEL f1 NUC1 13C P1 6.00 use PL1 3.00 dB SF01 125.7703643 MHz ====== CHANNEL f2 CPDPRG2 waltz16 NUC2 1H PCPD2 80.00 use PL2 -3.00 dB PL12 18.00 dB PL13 18.00 dB PL12W 0.42199361 W PL12W 0.42199361 W SF02 500.1320005 MHz SF 125.7577890 MHz WDW EM SSB 0
190 180 170 160 150 140 130 120 110 100 90 80 70 60	the second s



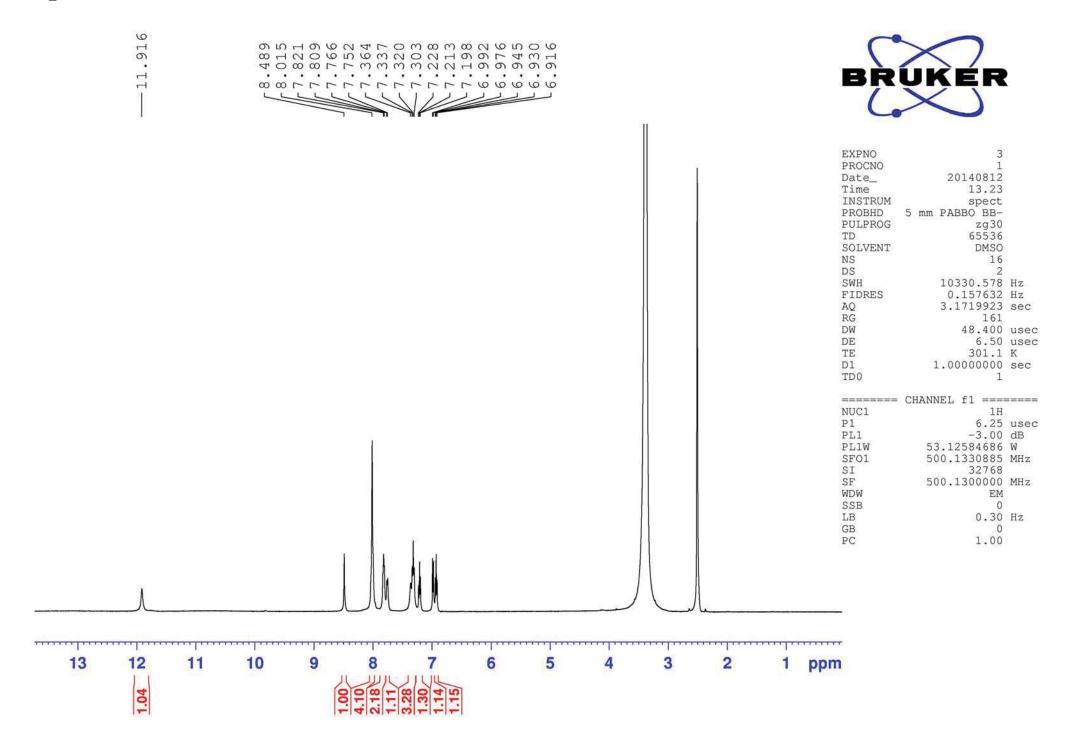
13C/MT-B-9/07-08-2014/DMSO



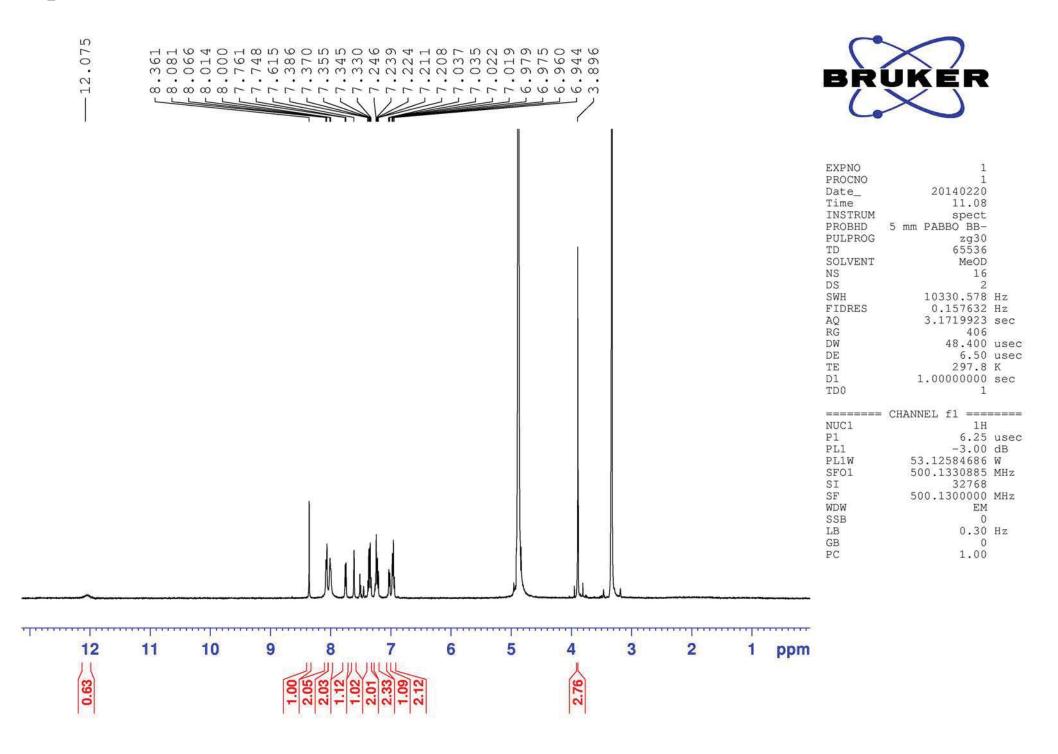


EXPNO	2	
PROCNO	1	
Date_	20140807	
Time	10.08	
INSTRUM	spect	
	5 mm PABBO BB-	
PULPROG	zgpg30	
TD	65536	
SOLVENT	DMSO	
NS	987	
DS	4	
SWH	29761.904	Hz
FIDRES	0.454131	Hz
AQ	1.1010548	sec
RG	32800	
DW	16.800	usec
DE	6.50	usec
TE	304.7	K
D1	2.00000000	sec
D11	0.03000000	sec
TD0	1	
	CHANNEL f1 ====	
NUC1	13C	
TIOOT	LUC	
P1		usec
		usec dB
Pl	6.00	dB
P1 PL1 SFO1	6.00 3.00 125.7703643 CHANNEL f2 ====	dB MHz
P1 PL1 SFO1 CPDPRG2	6.00 3.00 125.7703643 CHANNEL f2 ===== waltz16	dB MHz
P1 PL1 SFO1	6.00 3.00 125.7703643 CHANNEL f2 ===== waltz16 1H	dB MHz
P1 PL1 SFO1 CPDPRG2	6.00 3.00 125.7703643 CHANNEL f2 ==== waltz16 1H 80.00	dB MHz
P1 PL1 SF01 CPDPRG2 NUC2	6.00 3.00 125.7703643 CHANNEL f2 ===== waltz16 1H	dB MHz
P1 PL1 SF01 CPDPRG2 NUC2 PCPD2	6.00 3.00 125.7703643 CHANNEL f2 ==== waltz16 1H 80.00	dB MHz usec dB
P1 PL1 SF01 CPDPRG2 NUC2 PCPD2 PL2	6.00 3.00 125.7703643 CHANNEL f2 ==== waltz16 1H 80.00 -3.00	dB MHz usec dB
P1 PL1 SF01 CPDPRG2 NUC2 PCPD2 PL2 PL12	6.00 3.00 125.7703643 CHANNEL f2 ==== waltz16 1H 80.00 -3.00 18.00	dB MHz usec dB dB
P1 PL1 SF01 CPDPRG2 NUC2 PCPD2 PL2 PL12 PL13	6.00 3.00 125.7703643 CHANNEL f2 ===== waltz16 1H 80.00 -3.00 18.00 18.00	dB MHz usec dB dB dB
P1 PL1 SF01 CPDPRG2 NUC2 PCPD2 PL2 PL12 PL13 PL2W	6.00 3.00 125.7703643 CHANNEL f2 ==== waltz16 1H 80.00 -3.00 18.00 18.00 53.12584686	dB MHz usec dB dB dB W
P1 PL1 SF01 CPDPRG2 NUC2 PCPD2 PL2 PL12 PL12 PL13 PL2W PL12W	6.00 3.00 125.7703643 CHANNEL f2 ==== waltz16 1H 80.00 -3.00 18.00 18.00 53.12584686 0.42199361	dB MHz usec dB dB dB dB W W
P1 PL1 SF01 ————— CPDPRG2 NUC2 PCPD2 PL2 PL12 PL13 PL2W PL13W PL13W	6.00 3.00 125.7703643 CHANNEL f2 ==== waltz16 1H 80.00 -3.00 18.00 53.12584686 0.42199361 0.42199361 500.1320005	dB MHz usec dB dB dB dB W W W W
P1 PL1 SF01 CPDPRG2 NUC2 PCPD2 PL2 PL12 PL13 PL13 PL12W PL12W PL13W SF02	6.00 3.00 125.7703643 CHANNEL f2 ==== waltz16 1H 80.00 -3.00 18.00 53.12584686 0.42199361 0.42199361 500.1320005 32768	dB MHz usec dB dB dB dB W W W W
P1 PL1 SF01 CPDPRG2 NUC2 PCPD2 PL2 PL12 PL13 PL13W PL12W PL12W PL13W SF02 SI	6.00 3.00 125.7703643 CHANNEL f2 ==== waltz16 1H 80.00 -3.00 18.00 53.12584686 0.42199361 0.42199361 500.1320005 32768	dB MHz usec dB dB dB dB W W W W W MHz
P1 PL1 SF01 CPDPRG2 NUC2 PCPD2 PL2 PL12 PL13 PL2W PL12W PL12W PL12W SF02 SI SF WDW	6.00 3.00 125.7703643 CHANNEL f2 ==== waltz16 1H 80.00 -3.00 18.00 53.12584686 0.42199361 0.42199361 500.1320005 32768 125.7577890	dB MHz usec dB dB dB dB W W W W W MHz
P1 PL1 SF01 ======= CPDPRG2 NUC2 PCPD2 PL2 PL12 PL13 PL2W PL13W SF02 SI SF	6.00 3.00 125.7703643 CHANNEL f2 ==== waltz16 1H 80.00 -3.00 18.00 53.12584686 0.42199361 0.42199361 0.42199361 500.1320005 32768 125.7577890 EM	dB MHz usec dB dB dB dB W W W W W MHz
P1 PL1 SF01 CPDPRG2 NUC2 PCPD2 PL12 PL13 PL2W PL12W PL12W PL13W SF02 SI SF WDW SSB	6.00 3.00 125.7703643 CHANNEL f2 ==== waltz16 1H 80.00 -3.00 18.00 53.12584686 0.42199361 0.42199361 0.42199361 500.1320005 32768 125.7577890 EM 0	dB MHz usec dB dB dB dB W W W W MHz MHz

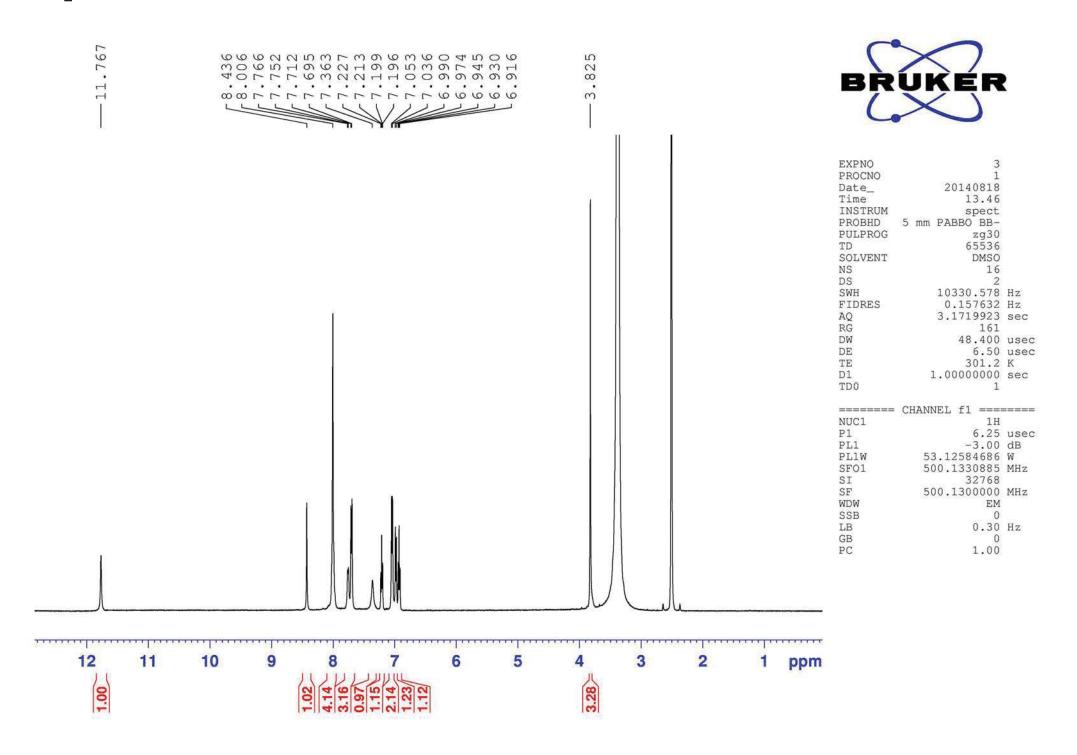
ppm

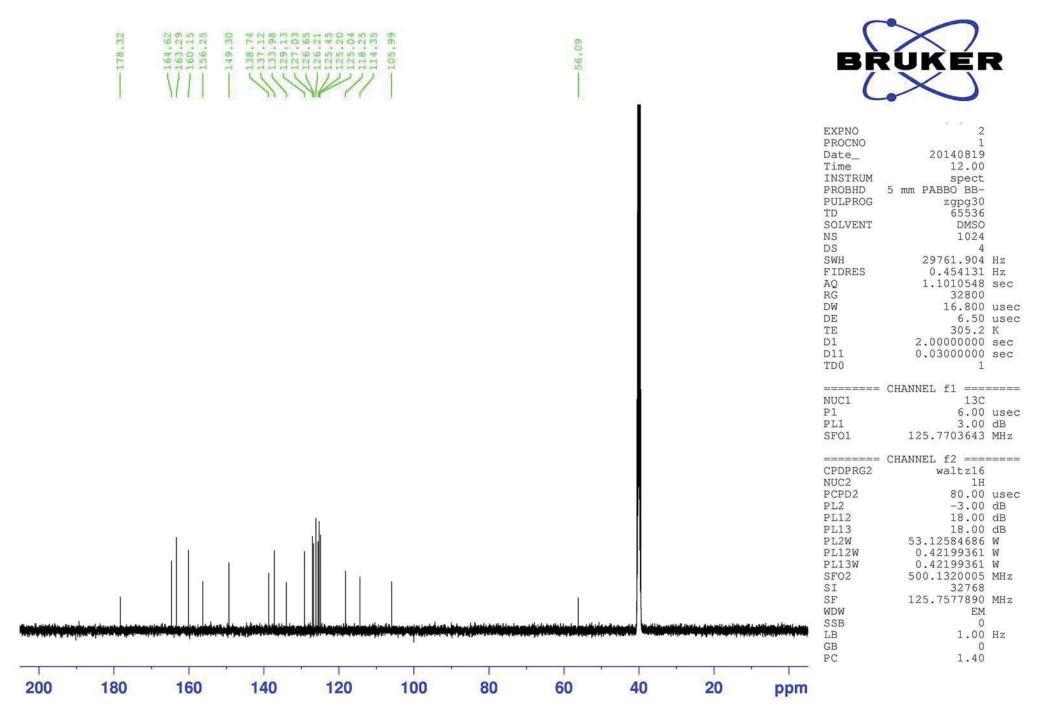


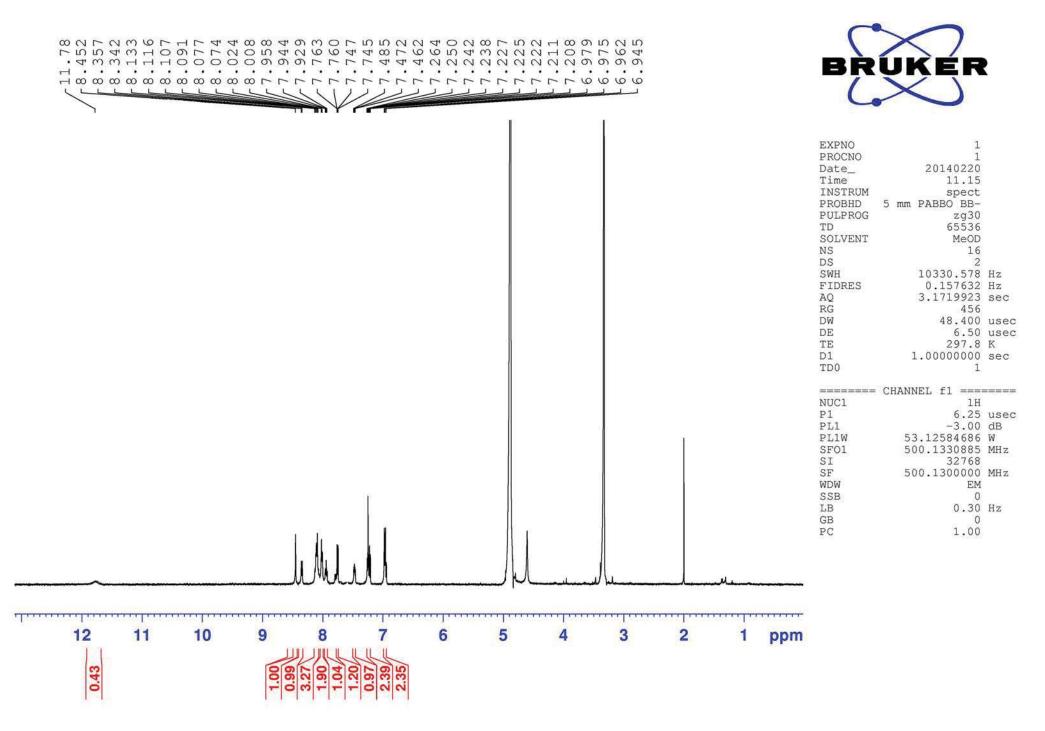
- 178.23 164.30 164.30 162.35 162.55 162.	BRUKER
	EXPNO 3 PROCNO 1 Date_ 20140829 Time 9.46 INSTRUM spect
	PROBHD 5 mm PABBO BB- PULPROG zgpg30 TD 65536 SOLVENT DMSO NS 2167 DS 4 SWH 29761.904 Hz
	FIDRES 0.454131 Hz AQ 1.1010548 sec RG 32800 DW 16.800 usec DE 6.50 usec TE 305.0 K D1 2.0000000 sec
	D11 0.03000000 sec TD0 1 ======= CHANNEL f1 ====== NUC1 13C P1 6.00 usec PL1 3.00 dB
	SF01 125.7703643 MHz ====== CHANNEL f2 CPDPRG2 waltz16 NUC2 1H PCPD2 80.00 usec PL2 -3.00 dB
	PL12 18.00 dB PL13 18.00 dB PL2W 53.12584686 W PL12W 0.42199361 W PL13W 0.42199361 W SFO2 500.1320005 MHz SI 32768 SF 125.7577890 MHz
	WDW EM SSB 0 LB 1.00 Hz GB 0 PC 1.40



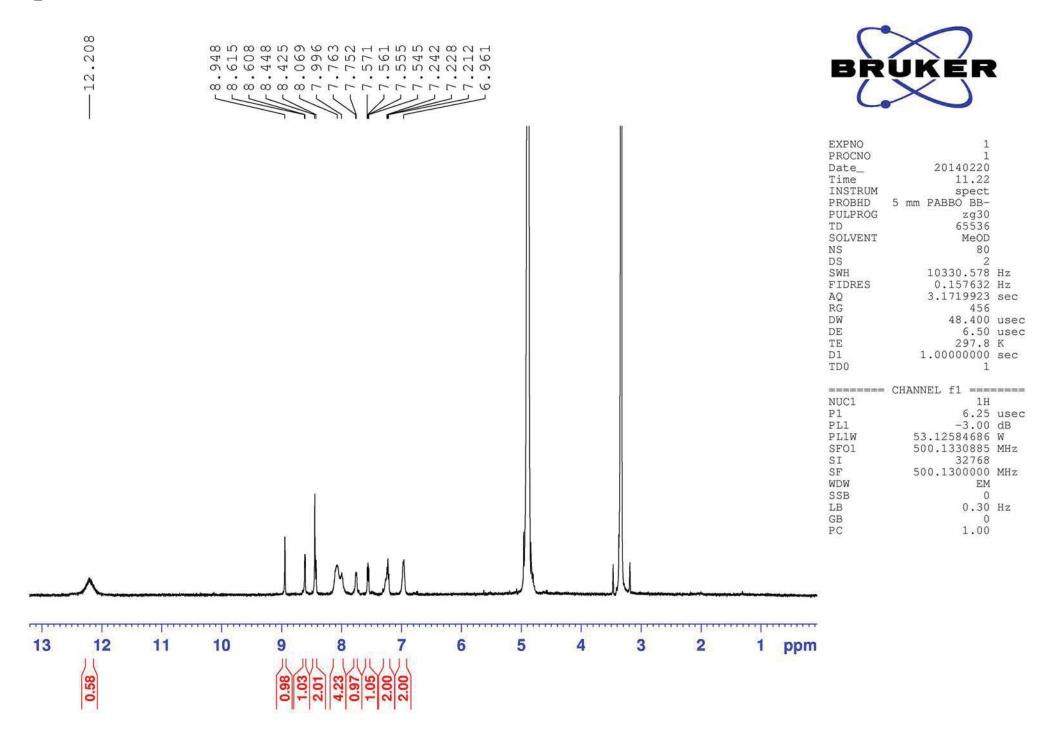
Compound 30			
	148.69 138.72 136.94 133.94 136.94 129.94 126.21 126.21 126.21 126.21 126.21 114.85 112.20 112.20	-16.29	BRUKER
			EXPNO 3 PROCNO 1 Date_ 20140829 Time 9.46 INSTRUM spect PROBHD 5 mm PABBO BB- PULPROG zgpg30 TD 65536 SOLVENT DMSO NS 2167 DS 4 SWH 29761.904 FIDRES 0.454131 AQ 1.1010548 RG 32800 DW 16.800 DE 6.50 TE 305.0 K D1 2.00000000 sec TD 13C P1 6.00 VUC1 13C P1 6.00 SFO1 125.7703643
mantanantanantanantana			PC 1.40







178.09	164.50 163.24 156.26 154.69	147.97 146.14 137.60 137.60 133.98 126.95	14000N				ĩ		BR	UKER
									EXPNO PROCNO Date_ Time INSTRUM PROBHD PULPROG	2 1 20140819 17.58 spect 5 mm PABBO BB- zgpg30
									TD SOLVENT NS DS SWH FIDRES AQ RG	65536 DMSO 1837 4 29761.904 Hz 0.454131 Hz 1.1010548 sec 32800
									DW DE TE D1 D11 TD0	16.800 usec 6.50 usec 304.8 K 2.00000000 sec 0.03000000 sec 1 CHANNEL f1 =======
									NUC1 P1 PL1 SF01	13C 6.00 usec 3.00 dB 125.7703643 MHz CHANNEL f2 ===================================
Ť	a h	र्गनी		Ť					NUC2 PCPD2 PL2 PL12 PL13 PL2W PL12W PL12W PL13W SF02	1H 80.00 usec -3.00 dB 18.00 dB 18.00 dB 53.12584686 W 0.42199361 W 0.42199361 W 500.1320005 MHz
land in picture and shares a share with a star of the second star in t			a a da a da a a da Anna a da	a en en else aner is rue a else en else else	li alian ya taka bila a da ka wa	(Ann bar of an bird and barrange a bird af na Tar payna an an an aif an barrange a s	and and a second a	na dan selan na kanan galikan Manganan garapatan selan kana	SF02 SF WDW SSB LB GB PC	32768 125.7577890 MHz EM 0 1.00 Hz 0 1.40



Compound	33										
178.15	164.29	149.84 145.21 145.30 138.58 137.11 137.11 133.66 133.69	126.21 125.42 125.42 125.04 124.41	102.91						BR	UKER
										EXPNO PROCNO Date_ Time INSTRUM PROBHD PULPROG TD SOLVENT NS DS SWH FIDRES AQ RG DW DE TE D1 D11 TD0	2 1 20140819 19.42 spect 5 mm PABBO BB- zgpg30 65536 DMSO 859 4 29761.904 Hz 0.454131 Hz 1.1010548 sec 32800 16.800 usec 6.50 usec 305.5 K 2.0000000 sec 0.03000000 sec 1
										NUC1 P1 PL1 SFO1	CHANNEL f1 ====== 13C 6.00 usec 3.00 dB 125.7703643 MHz
					nlande die jaken is der ste die ste ste die were die gebourde waar ste ste die ste ste	a di katalarki ka filo bikut a kuta	al lange dige			CPDPRG2 NUC2 PCPD2 PL12 PL13 PL2W PL13W PL12W PL13W SF02 SF SF WDW SSB	CHANNEL f2 ====== waltz16 1H 80.00 usec -3.00 dB 18.00 dB 18.00 dB 53.12584686 W 0.42199361 W 0.42199361 W 500.1320005 MHz 32768 125.7577890 MHz EM 0 1000 Hz
180	160		120	100	80	60	40	20	ppm	LB GB PC	1.00 Hz 0 1.40

	8.647 8.647 8.636 8.095 8.025 8.079 8.079 8.079 8.079 8.079 8.079 7.745 7.745 7.745 7.745 7.745 7.745 7.241 7.241 7.238	1000000	BRUKER
			EXPNO 1 PROCNO 1 Date_ 20140220 Time 8.32 INSTRUM spect PROBHD 5 mm PABBO BB- PULPROG 2g30 TD 65536 SOLVENT MeOD NS 16 DS 2 SWH 10330.578 Hz FIDRES 0.157632 Hz AQ 3.1719923 sec RG 575 DW 48.400 usec DE 6.50 usec TE 299.9 K D1 1.0000000 sec TD0 1 ====== CHANNEL f1 ===== NUC1 1H P1 6.25 usec PL1 -3.00 dB PL1W 53.12584686 W SFO1 500.1330885 MHz SI 32768 SF 500.1300000 MHz WDW EM SSB 0 LB 0.30 Hz GB 0
			PC 1.00
13 12 83	2.21 2.21 2.21 2.21 2.21 2.21 2.21 2.21	6 5 4 3 2	1 ppm

compound												
178,14	164.31		140.77 138.72 138.72 133.95 126.95	adoud	-105.93			I			BR	UKER
											EXPNO PROCNO Date_ Time INSTRUM PROBHD PULPROG TD SOLVENT NS DS SWH FIDRES AQ RG DW DE TE D1 D11 TD0	2 1 20140819 20.23 spect 5 mm PABBO BB- zgpg30 65536 DMSO 724 4 29761.904 Hz 0.454131 Hz 1.1010548 sec 32800 16.800 usec 6.50 usec 305.1 K 2.00000000 sec 0.03000000 sec 1
											NUC1 P1 PL1 SF01 CPDPRG2 NUC2 PCPD2	CHANNEL f1 ====== 13C 6.00 usec 3.00 dB 125.7703643 MHz CHANNEL f2 ====== waltz16 1H 80.00 usec
testerete honorale por por que encoder	ter a Wingship of Mari					de kalle van de blangelege oper klanster	tele and and post in the second second	fut of march owner and	hogo kunder og stade	Maiphely	PL2 PL12 PL13 PL2W PL12W PL13W SF02 SF SF WDW SSB LB	-3.00 dB 18.00 dB 18.00 dB 53.12584686 W 0.42199361 W 0.42199361 W 500.1320005 MHz 32768 125.7577890 MHz EM 0 1.00 Hz
180	1(6 <mark>0</mark>	140	120	100	80	60	40	20	ppm	GB PC	0 1.40