

Biotransformation of Chalcones by the Endophytic Fungus *Aspergillus flavus* Isolated from *Paspalum maritimum* Trin.

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O fungo *Aspergillus flavus* isolado como endofítico da planta *Paspalum maritimum* Trin. foi avaliado quanto ao seu potencial de aplicação em reações de biotransformações. Os compostos chalcona (**1**), 3,4,5-trimetoxichalcona (**2**) e 2,3,4,4'-tetrametoxichalcona (**3**) foram biotransformados, respectivamente, na diidrochalcona (**4**), 3,4,5-trimetoxidiidrochalcona (**5**) e 2,3,4,4'-tetrametoxidiidrochalcona (**6**). As estruturas destes compostos foram determinadas por análises de RMN uni e bidimensionais e por espectrometria de massas. As diidrochalconas **5** e **6** são substâncias inéditas.

The fungus *Aspergillus flavus* isolated as endophytic of the plant *Paspalum maritimum* Trin. was evaluated for its potential application in biotransformation reactions. The compounds chalcone (**1**), 3,4,5-trimethoxychalcone (**2**) and 2,3,4,4'-tetramethoxychalcone (**3**) were biotransformed, respectively, in dihydrochalcone (**4**), 3,4,5-trimethoxydihydrochalcone (**5**) and 2,3,4,4'-tetramethoxydihydrochalcone (**6**). The structures were elucidated by spectroscopic methods including 1D and 2D NMR techniques, and MS analysis. The dihydrochalcones **5** and **6** are new compounds.

Keywords: biotransformations, *Aspergillus flavus*, *Paspalum maritimum*, chalcones, dihydrochalcones

Introduction

Chemical reactions performed by microorganisms have been used as an alternative to obtain products of chemical and biological interest. Taking into account the modern concepts and requirements for a safer and environmentally friendly chemistry, the use of reactions promoted by microorganisms tends to increase in the near future.¹⁻³ Such reactions, usually known as biotransformation, have certain advantages over the conventional reactions. Biotransformation is usually performed in aqueous systems and at neutral pH, preventing the common use of dangerous solvents in conventional synthesis.⁴⁻⁹ In addition, microorganisms are capable of performing a wide variety of reactions, including some almost inaccessible by chemical means.¹⁰⁻¹³

Chalcones are precursors of the flavonoids and are defined as α,β -unsaturated ketones in which both the carbonyl and the olefinic fragment are linked to an aromatic ring. They possess a large spectrum of action, such as antiviral activity,¹⁴ inhibition of NS3 protease of dengue virus,¹⁵ activity against herpes simplex virus,¹⁶ HIV-1 replication inhibition in lymphocytes-H9 in mice,¹⁷ allelopathic activity,¹⁸ anti-inflammatory and antibacterial properties.¹⁹ These activities have been raising the interest in synthesizing and characterizing different chalcones, in order to obtain the corresponding dihydrochalcones to correlate structure *versus* biological activity.

Biotransformation of bioactive natural and synthetic compounds by many microorganisms including species of *Aspergillus* have been examined and investigated.^{20,21} Regioselective *O*-demethylation of flavones into their 4'-*O*-demethylated metabolites was performed using an *A. niger* strain.²² Biotransformation of flavanones by one wild

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strain of *A. niger* KB and three UV mutants of *A. niger* resulted in the reduction of carbonyl group and dehydrogenation at C-2 and C-3 positions; reduction of carbonyl group and hydroxylation at C-7; and dehydrogenation at C-2 and C-3 and hydroxylation at C-3 producing flavonol.²³ In the high molecular weight fraction of the culture extract of *Aspergillus oryzae*, genistein was transformed into shoyuflavone B in the presence of (\pm)-transepoxy succinic acid.²⁴ Epigallocatechin gallate (EGCG) has been reported to be converted into epigallocatechin (EGC) by a hydrolase from *Aspergillus oryzae* and the yield of EGC could reach at least 70%.²⁵ Biotransformation of the flavanone by means of *Aspergillus niger* MB culture led to formation of three products: 6-hydroxyflavanone, 2',5'-dihydroxydihydrochalcone and 2'-hydroxydihydrochalcone. Their yields were 18%, 6% and 7.5% after 9 days, respectively.²⁶

Due to the potential ability of *Aspergillus* to promote biotransformation reactions, we decided to study biotransformation reactions of chalcone (**1**), 3,4,5-trimethoxychalcone (**2**) and 2,3,4,4'-tetramethoxychalcone (**3**), using the endophytic fungus *Aspergillus flavus* isolated from the plant *Paspalum maritimum*.

Results and Discussion

The chalcones **1** (68.5% yield),²⁷ **2** (75.0% yield)²⁸ and **3** (60.0% yield)²⁹ were synthesized using methodology described in the literature¹⁸ and their structures were determined by the analysis of 1D- and 2D-NMR, mass spectral information and comparison with reported physical and spectroscopic data. These chalcones were bioreduced by *Aspergillus flavus* producing the dihydrochalcones **4** (74.5% yield), **5** (36.3% yield) and **6** (15.6% yield), respectively, after 10 days of reaction (Figure 1). The known dihydrochalcone (**4**) was identified by comparison of the spectroscopic data with those

published in literature.³⁰ The dihydrochalcones **5** and **6** are new compounds.

The bioreduction reaction of the chalcones was regioselective, showing only alteration in carbon-carbon double bond of the olefin, with no reaction at the carbonyl double bond. Several reactions of chalcones are reported, however, none of them showed regioselectivity for the double bond C-2 and C-3. Among the reactions of biotransformation reported directly on chalcones, *Aspergillus alliaceus* cyclized the chalcone 2'-hydroxy-2,3-dimethoxychalcone to three flavanones and to *O*-demethylated (6.5% yield) and hydroxylated (7.2% yield) chalcones.³¹ The unsubstituted chalcone was converted to 2''-hydroxychalcone and 2'',3''-dihydroxychalcone in 25% and 59% yields, respectively, by *Escherichia coli*.³² Microbial transformation of chalcone xanthohumol, using the culture broth of *Pichia membranifaciens* afforded three metabolites, one isomeric prenylflavanone in 3.3% yield, and two modified prenylchalcones in 0.55 and 0.58% yield.³³ The yields obtained in the biotransformation reactions of chalcones to dihydrochalcones in this work are higher (15.6-75.5% yield) than those reported in the literature, although no reaction can be directly compared, because none of them performed this type of bioreduction (direct formation of dihydrochalcones from chalcone).

The biotransformation products (dihydrochalcones **4**, **5** and **6**) were characterized mainly by the presence of two triplets in the ¹H NMR spectra that indicate the reduction of the double bond of the conjugated olefinic system. In the ¹³C NMR spectrum, the chemical shift of the carbonyl carbon increased about 10 ppm due to the loss of conjugation.

Compound **5** was identified as 3,4,5-trimethoxydihydrochalcone. The analysis of the LR-MS spectrum showed that compound **5** has a molecular ion peak at *m/z* 300 (64%) and fragments at *m/z* 195 (100%), 181 (36%), 165 (42%) and 77 (36%), suggesting a molecular formula

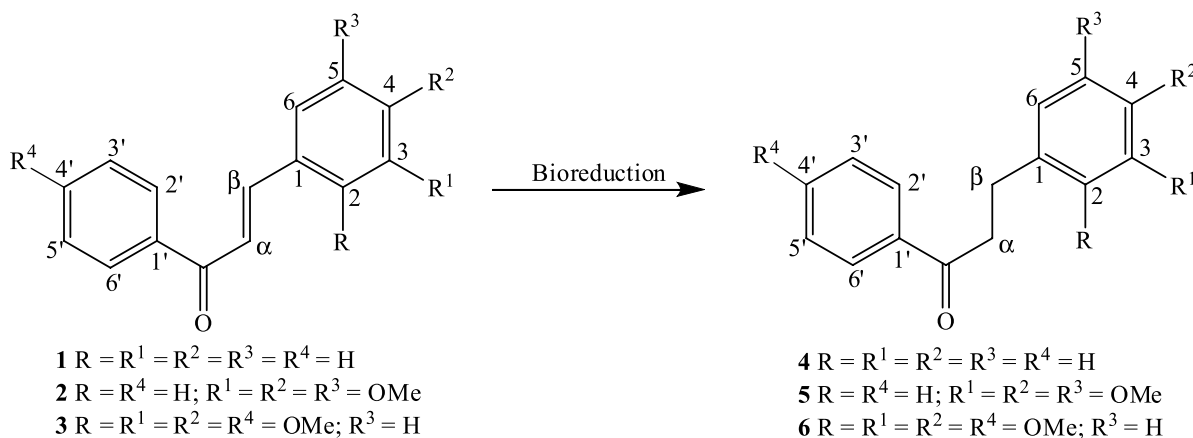


Figure 1. Bioreduction of the chalcones **1**, **2** and **3** by *Aspergillus flavus*.

$C_{18}H_{20}O_4$ (9 degrees of unsaturation). This molecular formula was confirmed on the basis of HRESIMS [m/z 323.1270 (M+Na)⁺; Δ -1.1 mmu] analysis and NMR data (Table 1). The 1H NMR spectrum of **5** (Table 1) showed signals due to the benzylic methylene protons H β at δ_H 3.01 (2H, *t*, *J* 7.5 Hz) and to the α -carbonyl protons H α at δ_H 3.29 (2H, *t*, *J* 7.5 Hz). The coupling of the protons H α and H β was confirmed by analysis of $^1H \times ^1H$ -COSY spectrum. HETCOR spectrum ($^1J_{CH}$) showed correlations of the signals of the protons at δ_H 3.29 (H α) and 3.01 (H β) with signals at δ_C 40.6 (C α) and 30.6 (C β), respectively. The triplet at δ_H 3.01, assigned to the signal of hydrogen H β , showed a correlation in the HMBC spectrum with the signals at δ_C 137.1 (C1) and at δ_C 105.4 (C2/C6). The singlet at δ_H 6.46 (2H, *s*, ArH) assigned to the equivalent protons H2/H6 of ring B showed correlation with the signal at δ_C 105.4 (C2/C6) in the HETCOR spectrum. The HMBC spectrum showed the correlation of the signal at δ_H 6.46 (H2/H6) with the signals at δ_C 153.2 (C3/C5), 136.9 (C4), 105.4 (C2/C6) and 30.6 (C β). The assignment of the protons and carbons signals of ring A was proposed from the analysis of $^1H \times ^1H$ -COSY and HETCOR spectra.

The HETCOR spectrum showed the correlations of signals at δ_C 128.6 (C2'/C6') 128.0 (C3'/C5') and 133.1 (C4') with the signals at δ_H 7.96 (H2'/H6'), 7.45 (H3'/H5') and 7.56 (H4'), respectively. The HMBC spectrum showed correlation of the signal at δ_H 7.45 (H3'/H5') with signals at δ_C 128.6 (C2'/C6') and δ_C 136.9 (C1') and of the signal at δ_H 7.96 (H2'/H6') with the signals at δ_C 128.0 (C3'/C5'), 133.1 (C4') and 199.2 (C=O).

Compound **6** was identified as 2,3,4,4'-tetramethoxy-dihydrochalcone. The analysis of LR-MS spectrum of **6** showed a molecular ion peak at m/z 330 (82%) and the peaks at m/z 195 (24%), 181 (98%), 166 (48%), 135 (100%) and 107 (16%), which are consistent with the molecular formula $C_{19}H_{22}O_5$ (9 degrees of unsaturation). This molecular formula was confirmed on the basis of HRESIMS [m/z 353.1358 (M+Na)⁺; Δ +0.7 mmu] analysis and NMR data (Table 1). The 1H NMR spectrum of **6** showed signals attributed to the benzylic methylene protons H β and H α similar to those of compound **5**. Two *ortho*-related doublets in the aromatic region (1H each, *J* 8.4 Hz) at δ_H 6.87 and 6.59, were assigned to H6 and H5, respectively. The signal at δ_H 6.59 correlated with the signals of carbons at δ_C 127.2 and 142.3 in the

Table 1. 1H and ^{13}C NMR chemical shifts (δ_H and δ_C in ppm) of compounds **5** and **6** in $CDCl_3$ ^{a,b}

Position	5			6		
	δ_H	δ_C	HMBC ^c	δ_H	δ_C	HMBC
1		137.1 ^d			127.2	5, 6, α
2	6.46 (s)	105.4	6, β		151.9	OMe-2
3		153.2	2, OMe-3		142.3	5
4		136.9 ^d	2, 6, OMe-4		152.3	6, OMe-4
5		153.2	6, OMe-5	6.59 (d, 1H, <i>J</i> 8.4)	107.2	
6	6.46	105.4	2, β	6.87 (d, 1H, <i>J</i> 8.4)	123.9	3
α	3.29 (t, 2H, <i>J</i> 7.5)	40.6	β	3.18 (t, 2H, <i>J</i> 8.4)	39.5	β
β	3.01 (t, 2H, <i>J</i> 7.5)	30.6	2, 6, α	2.96 (t, 2H, <i>J</i> 8.4)	25.3	6, α
C=O		199.2	2', 6'		198.4	α , β , 2', 6'
1'		136.9	3', 5'		129.9	
2'	7.96 (dd, 2H, <i>J</i> 7.5 and 1.5)	128.6	6'	7.95 (d, 2H, <i>J</i> 9.3)	130.6	3', 6'
3'	7.45 (t, 2H, <i>J</i> 7.5)	128.0		6.92 (d, 2H, <i>J</i> 9.3)	113.6	
4'	7.56 (tt, 1H, <i>J</i> 13.2, 7.5 and 1.5)	133.1	2', 6'		163.3	2', 6', OMe-4'
5'	7.45 (t, 2H, <i>J</i> 7.5)	128.0		6.92 (d, 2H, <i>J</i> 9.3)	113.6	
6'	7.96 (dd, 2H, <i>J</i> 1.5 and 7.5)	128.6	2'	7.95 (d, 2H, <i>J</i> 9.3)	130.6	2', 5'
OMe-2				3.90 (s)	60.8	
OMe-3	3.84 (s)	56.1		3.88 (s)	60.6	
OMe-4	3.82 (s)	60.8		3.83 (s)	55.3	
OMe-5	3.84 (s)	56.1				
OMe-4'				3.85 (s)	55.9	

^a 1H NMR data were recorded at 300 MHz. ^b ^{13}C NMR data were recorded at 75 MHz. ^c 1H - ^{13}C HMBC correlations are from the carbon(s) specified to the protons indicated. ^d Values can be reversed.

HMBC spectra, what allowed the assignment of these signals to C1 and C3, respectively. In the same NMR spectrum, it was observed that the signal at δ_{H} 6.87 (H6) showed correlation with the signals at δ_{C} 151.9 and 152.3, which indicated that these signals can be attributed to C2 and C4 carbons, respectively. The analysis of the HETCOR ($^1J_{\text{CH}}$) spectrum of **6** confirms the assignments of the signals of the protons H5 and H6, since they showed correlations with the signals of carbons at δ_{C} 107.2 (C5) and 123.9 (C6). The ^1H NMR spectrum showed two *ortho*-related doublets at δ_{H} 7.95 and 6.92 (2H each, J 8.4 Hz), due to the protons of a system AA'-XX', assigned to H2'/H6' and H3'/H5', respectively. This coupling was confirmed by analysis of spectrum $^1\text{H} \times ^1\text{H}$ -COSY. The analysis of HETCOR spectrum of **6**, relating to the ring A, was important to assign accurately the signals at δ_{C} 130.3 and 113.6 to carbons C2'/C6' and C3'/C5', respectively, that showed $^1J_{\text{CH}}$ correlations with signals at δ_{H} 7.95 and 6.92. The HMBC spectrum showed correlations of the signals of carbons at δ_{C} 129.9 (C1') and 163.3 (C4') with the signals at δ_{H} 6.92 (H3'/H5') and δ_{H} 7.95 (H2'/H6'). The assignment of carbonyl carbon signal (δ_{C} 198.4) was based on the analysis of the correlations with the signals due to H β and H α .

Experimental

General experimental procedures

Low-resolution mass spectra data were obtained on a Thermo DSQII GC-MS instrument equipped with fused silica capillary column (30 m \times 0.25 mm id) with DB-5MS stationary phase (0.25 μm thickness of the film); carrier gas helium gas, temperature of the source and other parts 200 °C; type of injection: splitless; programmed temperature to 100-300 °C, with a gradient of 5 °C min^{-1} . The quadrupole filter swept range 39-400 daltons every second. Ionization was obtained by the technique of electron impact with energy of 70 eV, and high resolution mass spectra were recorded on an LC-IT-TOF mass spectrometer model 225-07100-34-SHIMADZU, by positive ionization mode of the ESI. NMR spectra, including HMBC experiment, were recorded on a Varian Mercury-300 spectrometer, operating at 300 MHz for ^1H and 75 MHz for ^{13}C NMR, using solvent signal (d-chloroform) as reference. The chemical shifts are given in delta (δ) values and the coupling constants (J) in Hertz (Hz).

Plant material

The leaves of *Paspalum maritimum* were collected in January 2009 in the forest reserve of EMBRAPA-Amazônia

Oriental in Belém, Pará State, Brazil. A voucher specimen (IAN 183337) was deposited at the herbarium of this institution.

Microorganisms and culture

The general procedures adopted for isolation of the microorganism followed the methodology described by Araújo *et al.*³⁴ Epiphytic hyphae and spores were eliminated by immersion of the plant material in 70% ethanol (1 min), followed by 5% aqueous sodium hypochlorite (4 min) and finally with 70% aqueous ethanol (30 s), before being rinsed in sterile distilled water. The material was then deposited on a Petri dish containing Sabouraud (SDA), composition meat peptone 10 g L^{-1} (HIMEDIA), D-glucose anhydrous 20.0 g L^{-1} (NUCLEAR), Agar 20.0 g L^{-1} (VETEC) and chloramphenicol 1.0 g L^{-1} and incubated in the dark at 30 °C. Samples of the fast growing fungi were subcultured onto other plates with the culture medium used. The microorganisms were conserved as suspension in distilled water. The aflpm3D1 strain was identified as *Aspergillus flavus* by Dra. Maria Inês de Moura Sarquis, Fiocruz-RJ. A voucher specimen has been deposited at "Laboratório de Química de Microorganismos e Bioensaios-UFPA".

Synthesis of substrate:¹⁸ chalcones 1-3

In a flat-bottomed flask (125 mL), placed in an ice bath, 15 mL of EtOH, 11 mmol of acetophenone or derivatives, 15 mL of 10% NaOH solution and 12 mmol of benzaldehyde or derivatives were added. The reaction mixture was kept on magnetic stirring at 40 °C for 40 min. After this period, the mixture was cooled and left in freezer for 48 h and then filtered under vacuum. The product was crystallized in methanol. Chalcone **1** was obtained in 68.5% yield (1.47 g), chalcone **2** was obtained in 75.0% yield (2.42 g) and chalcone **3** was obtained in 60.0% yield (2.62 g).

General procedure for biotransformation reactions

One Erlenmeyer flask (250 mL), containing 100 mL SDA modified medium (meat peptone 10 g L^{-1} (HIMEDIA), D-glucose anhydrous 10 g L^{-1} (NUCLEAR) and chloramphenicol 1.0 g L^{-1}) was previously autoclaved at 121 °C for 15 min. Three small disc cut from Petri dishes containing mycelium of *A. flavus* in SDA medium and the appropriate chalcones (**1-3**), 0.05 g dissolved in 0.5 mL DMSO, were added under steril conditions to the Erlenmeyer flask and shaken (160 rpm) at 30 °C. After 10 days the reaction mixture was extracted with EtOAc (3 \times 100 mL), and organic phases were grouped, dried (Na_2SO_4), filtered

and evaporated at reduced pressure. Experiments were performed in triplicate and were reproducible according to TLC analysis. Control flasks consisted of cultures containing substrate in DMSO without fungus and cultures containing fungus and DMSO without substrate. The purifications were carried out from one experiment. The crude residue (102.0 mg) obtained from the biotransformation reaction of chalcone **1** was chromatographed on silica gel by elution with EtOAc (300 mL) and MeOH (100 mL). Evaporation of solvent furnished EtOAc extract (62.0 mg) and MeOH extract (30.6 mg). TLC analysis indicated that the substances of interest were in the extract EtOAc. This extract was chromatographed sequentially on silica gel eluting with hexanes and increasing polarity with EtOAc, to give eleven fractions (A1-A11). Fractions A3-A5 (hexanes:EtOAc 9:1; 8:2 and 8:2, respectively) were grouped and recrystallized with MeOH giving dihydrochalcone **4** (37.2 mg). The crude residues obtained from the biotransformation reactions of chalcones **2** and **3** after the same treatment, provided EtOAc extract (113.0 mg) and MeOH extract (90.6 mg) to the chalcone **2** reaction, and EtOAc extract (119.0 mg) and MeOH extract (97.3 mg) to the chalcone **3** reaction. The EtOAc extract obtained from the biotransformation reactions of chalcone **2** was chromatographed sequentially on silica gel eluting with hexanes and increasing polarity with EtOAc, to give fourteen fractions (B1-B14). Fractions B6-B10 (hexanes:EtOAc 8:2; 8:2; 7:3; 7:3 and 6:4, respectively) were grouped and recrystallized with MeOH giving dihydrochalcone **5** (18.3 mg). The EtOAc extract obtained from the biotransformation reactions of chalcone **3** was chromatographed sequentially on silica gel eluting with hexanes and increasing polarity with EtOAc, to give seventeen fractions (C1-C17). Fractions C12 (hexane:EtOAc 6:4) and C13 (hexane:EtOAc 6:4) were grouped and recrystallized with MeOH giving dihydrochalcone **6** (7.8 mg).

3,4,5-Trimethoxydihydrochalcone (5)

Crystalline solid; mp 59-61 °C (MeOH); ¹H NMR (300 MHz, CDCl₃): see Table 1; ¹³C NMR (75 MHz, CDCl₃): see Table 1. LR-MS: *m/z* (rel. int.): 300 (64), 195 (100), 181 (36), 165 (42), 77 (36). HRESIMS [M+Na]⁺; Found: 323.1270 [M +Na]⁺; Calc. for C₁₈H₂₀O₄Na: 323.1259.

2,3,4,4'-Tetramethoxydihydrochalcone (6)

Crystalline solid; mp 75-77 °C (MeOH); ¹H NMR (300 MHz, CDCl₃): see Table 1; ¹³C NMR (75 MHz, CDCl₃): see Table 1. LR-MS: *m/z* (rel. int.): 330 (82), 195 (24), 181 (98), 166 (48), 135 (100), 107 (16). HRESIMS [M+Na]⁺ Found: 353.1358 [M+Na]⁺; Calc. for C₁₉H₂₂O₅Na: 353.1365.

Conclusions

Microbial transformation is an effective tool for the structural modification of bioactive natural and synthetic compounds, including chalcones. Microbial factories pose advantages such as rapid growth, ease of cultivation, convenient genetic manipulations, and high level production on natural products biotransformation. Moreover, microbial production increases product selectivity and reduces the usage of toxic chemicals while conserving energy usage. The use of microorganisms in bioreduction reactions of chalcones has led to the formation of several products in mixture, formed by cyclization, reduction, *O*-demethylation, hydroxylation reactions, among others. Generally, these reactions have shown low yield and stereoselectivity. In this work we performed the biotransformation reaction of chalcones to dihydrochalcones in excellent yields, compared to other biotransformation reactions of flavonoids. The bioreduction reaction was regioselective, showing only alteration in carbon-carbon double bond of the olefin, not occurring any alteration in the carbonyl double bond of the α,β -unsaturated ketone system. In addition, to the best of our knowledge this is the first report of the obtention of dihydrochalcones directly from biotransformation reaction of chalcones.

Supplementary Information

Supplementary Information (Figure S1-S19) is available free of charge at <http://jbcs.org.br> as PDF file.

Acknowledgments

The authors would like to thank CAPES, CNPq and FAPESPA for financial support; Dra. Maria Inês de Moura Sarquis (Fiocruz-RJ) by identification of the fungus; Laboratory of Mass Spectrometry of Northeast (LEMANOR) by acquisition of mass spectra (LCMS-IT-TOF: quadrupole ion trap-time of flight mass spectrometer).

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Submitted: July 8, 2010

Published online: March 22, 2011

Supplementary Information

Biotransformation of Chalcones by the Endophytic Fungus *Aspergillus flavus* Isolated from *Paspalum maritimum* Trin.

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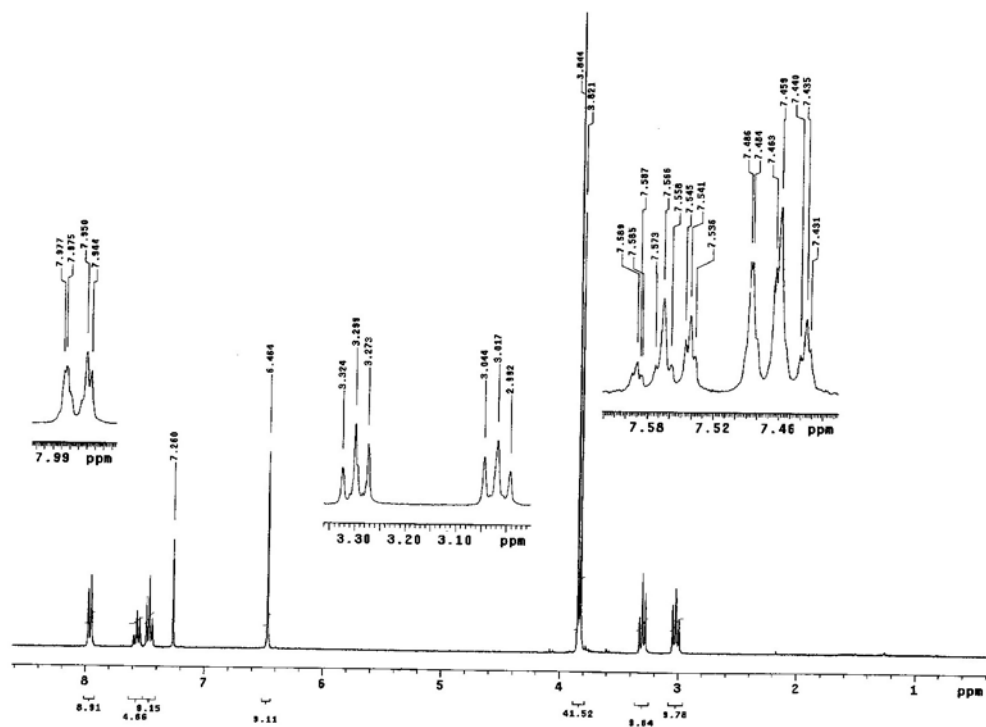


Figure S1. ¹H NMR spectrum (300 MHz, CDCl₃) of compound 5.

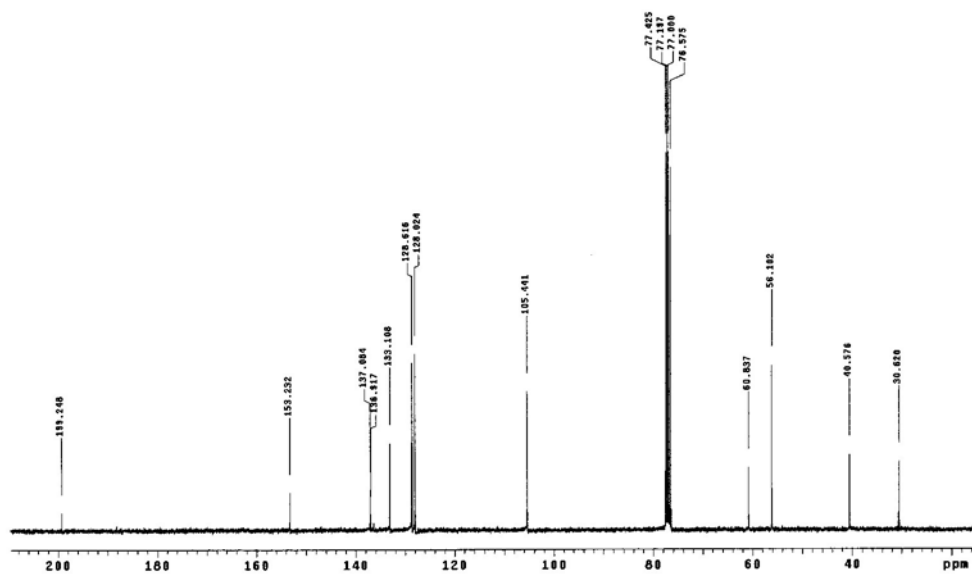


Figure S2. ^{13}C NMR spectrum (75 MHz, CDCl_3) of compound 5.

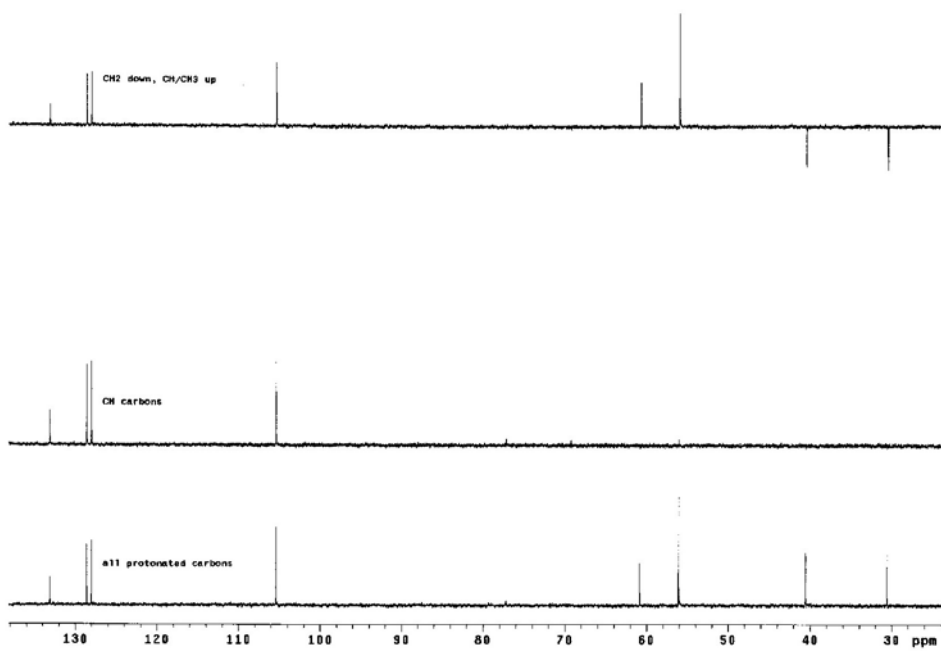


Figure S3. DEPT spectrum (75 MHz, CDCl_3) of compound 5.

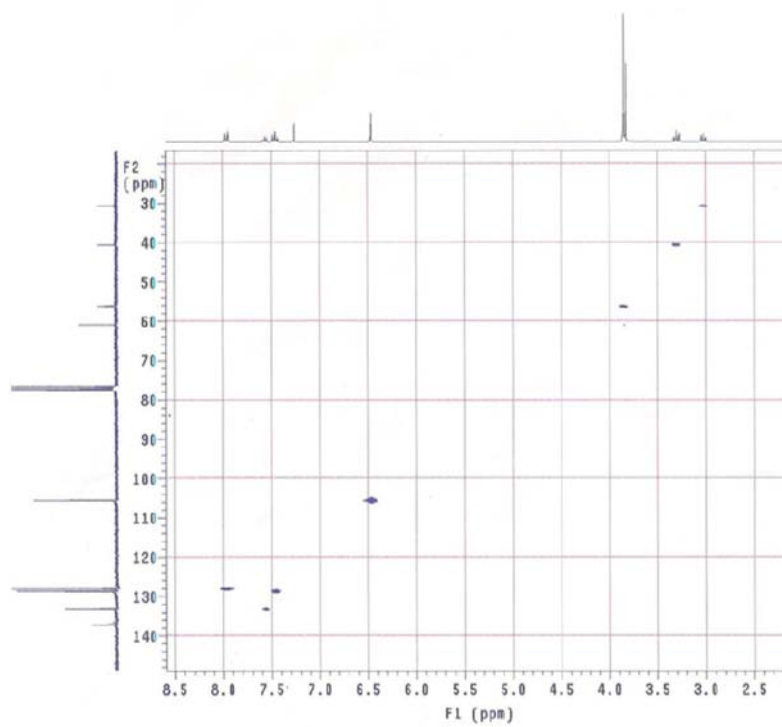


Figure S4. HETCOR spectrum (CDCl₃) of compound **5**.

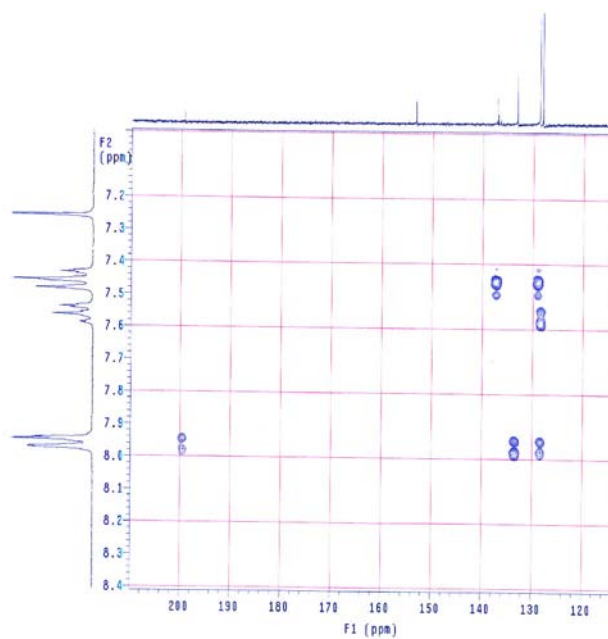


Figure S5. HMBC spectrum of compound **5** (expansion).

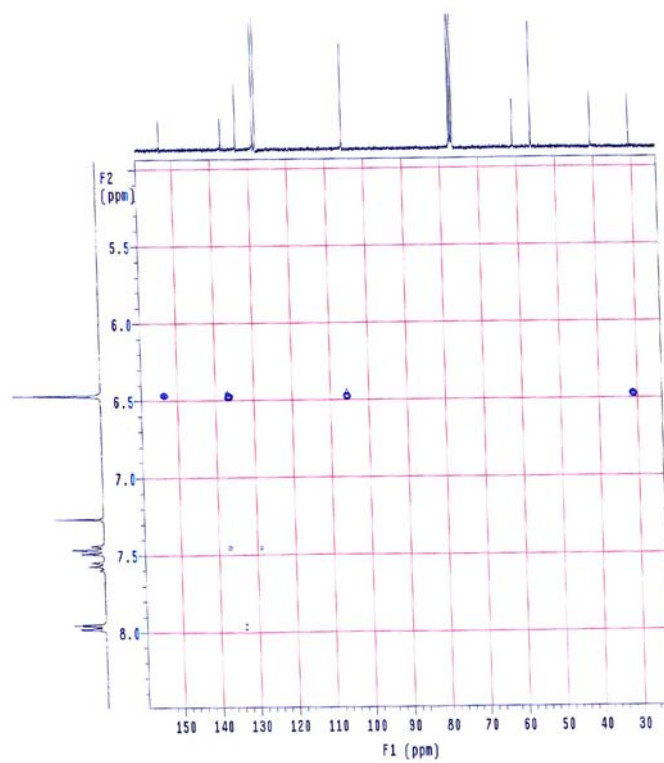


Figure S6. HMBC spectrum of compound **5** (expansion).

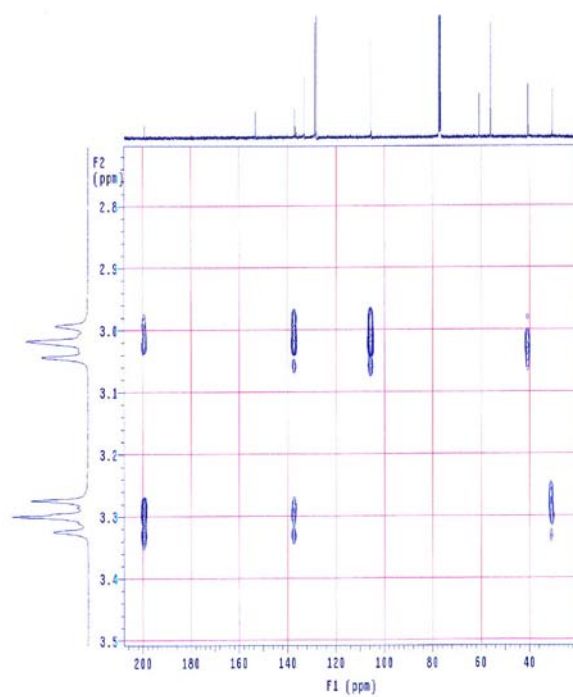


Figure S7. HMBC spectrum of compound **5** (expansion).

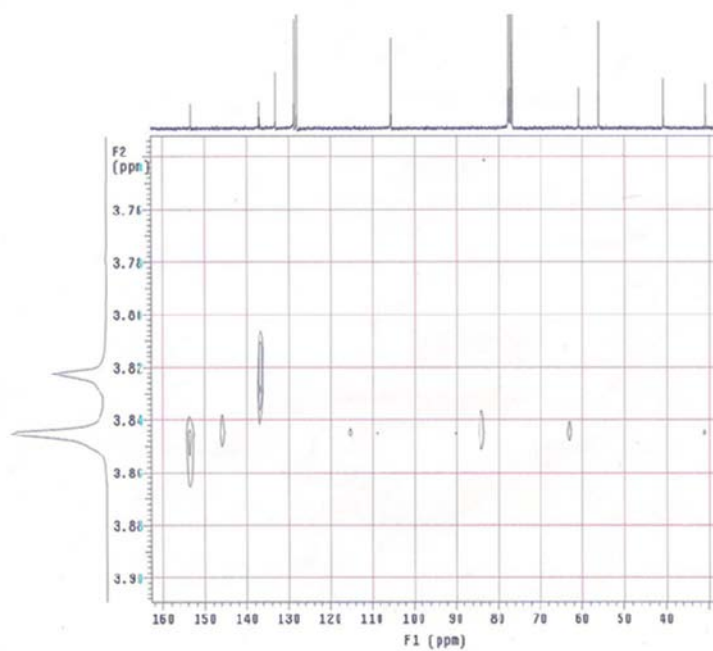


Figure S8. HMBC spectrum of compound **5** (expansion).

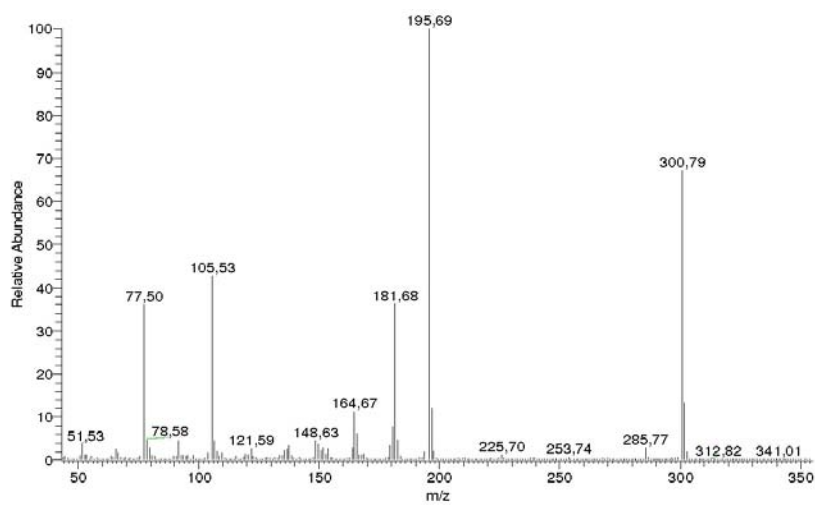


Figure S9. Low resolution mass spectrum of compound **5**.

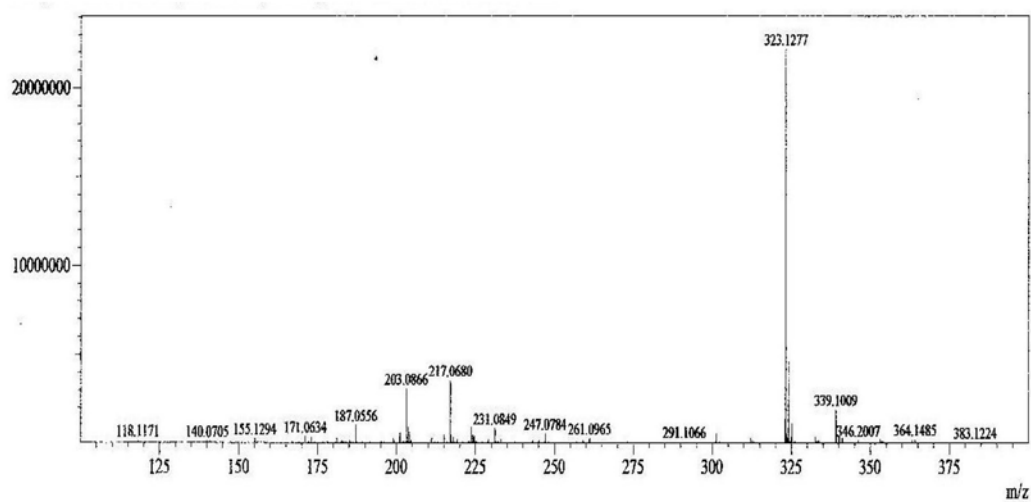


Figure S10. High resolution mass spectrum of compound 5.

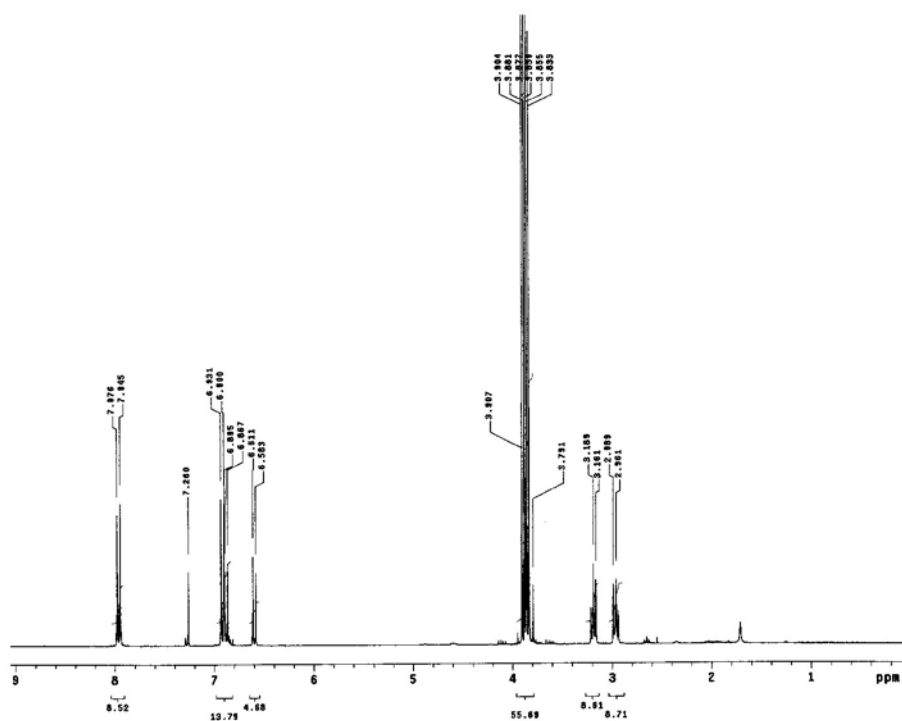


Figure S11. ^1H NMR spectrum (300 MHz, CDCl_3) of compound 6.

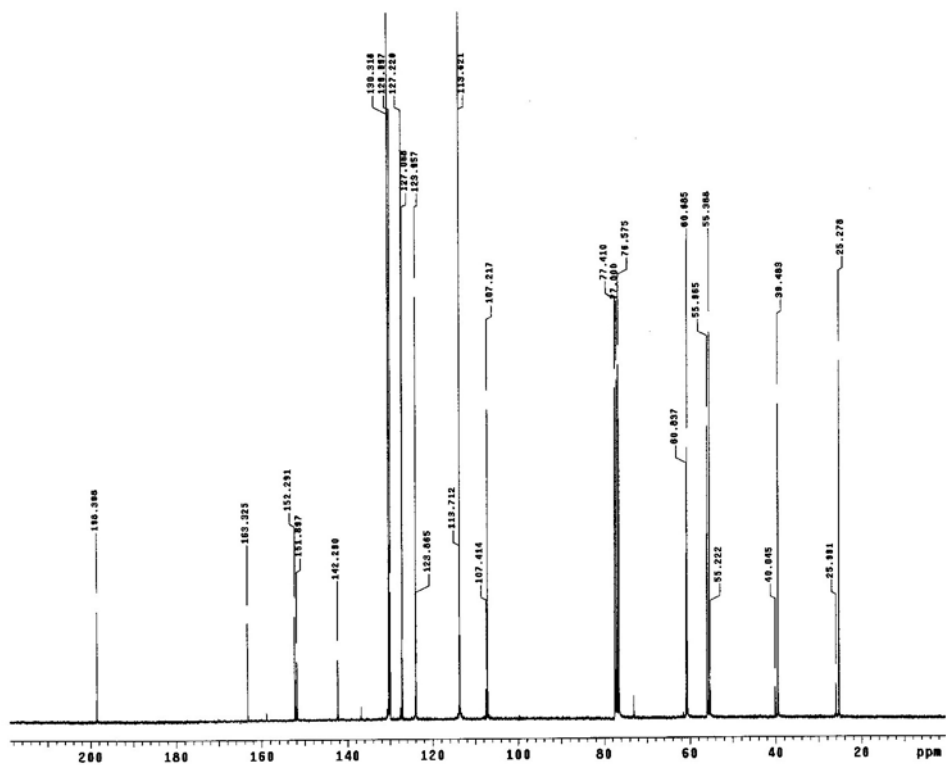


Figure S12. ^{13}C NMR spectrum (75 MHz, CDCl_3) of compound 6.

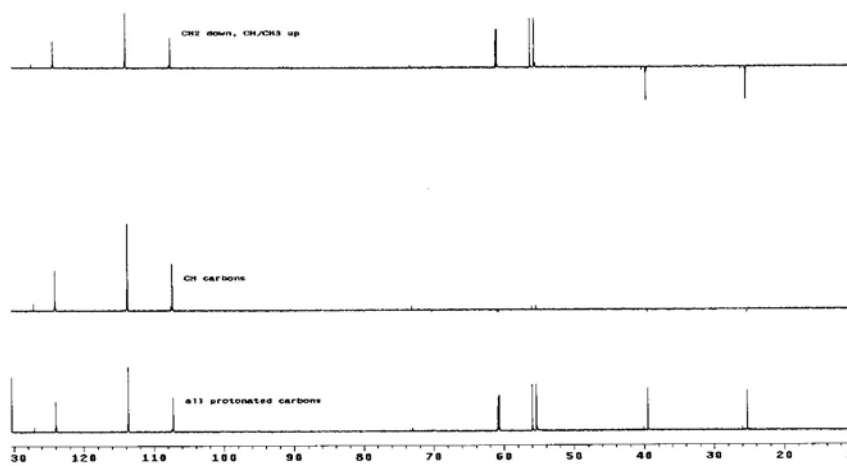


Figure S13. DEPT spectrum (75 MHz, CDCl_3) of compound 6.

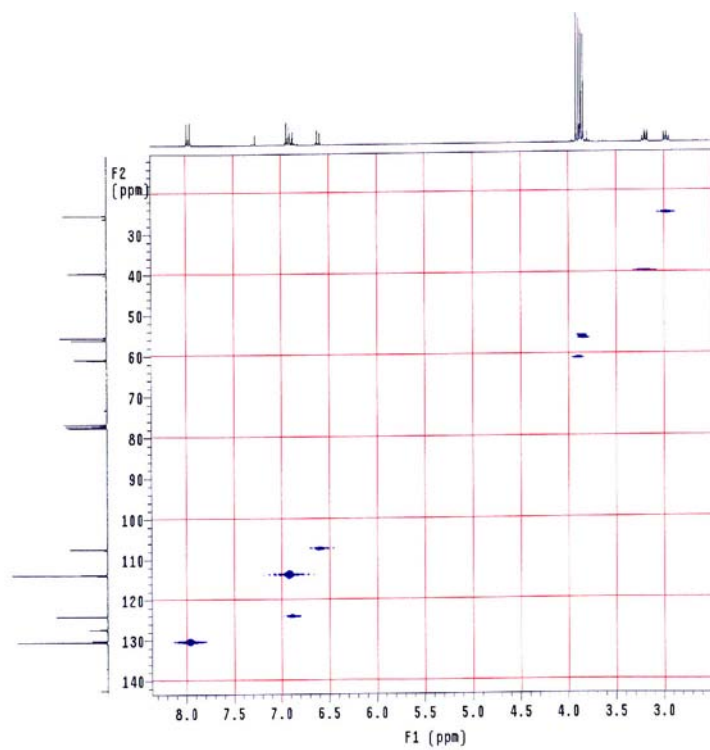


Figure S14. HETCOR spectrum (75 MHz, CDCl₃) of compound **6**.

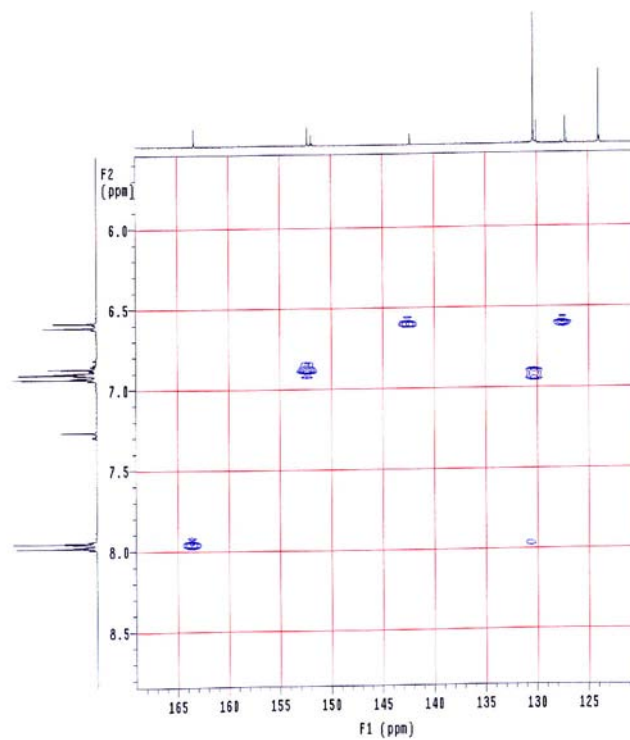


Figure S15. HMBC spectrum of compound **6** (expansion).

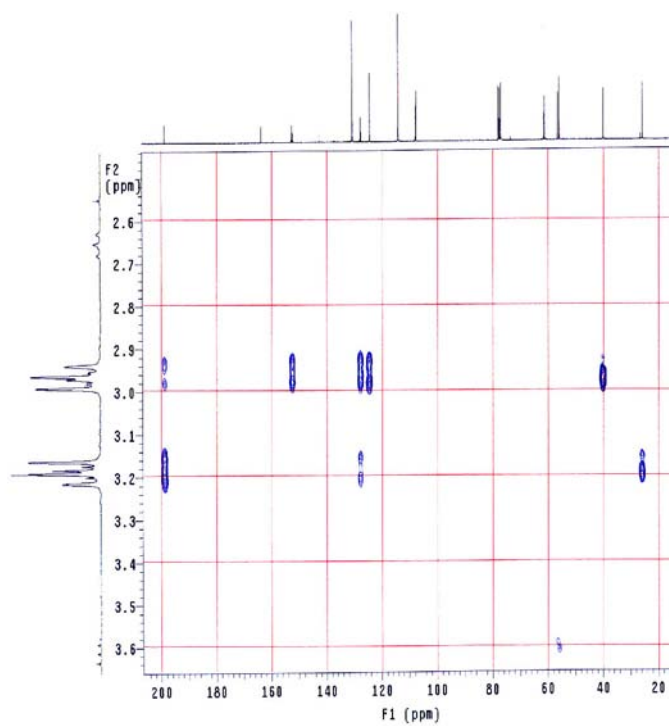


Figure S16. HMBC spectrum of compound 6 (expansion).

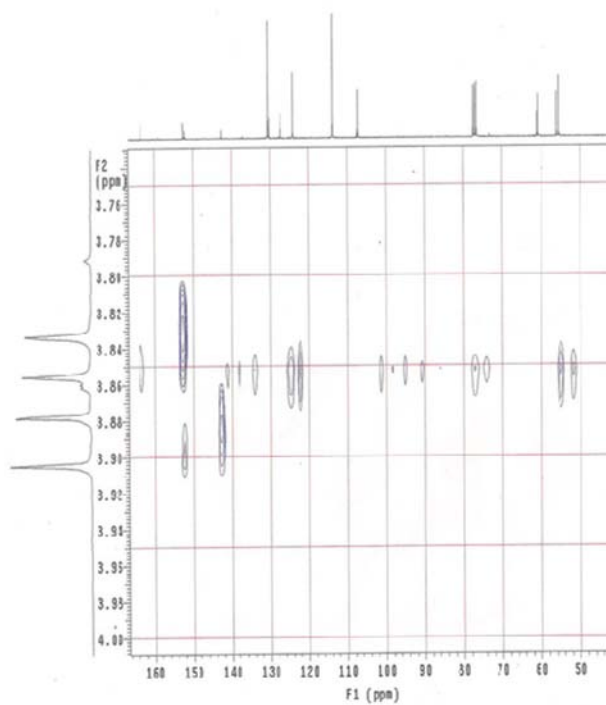


Figure S17. HMBC spectrum of compound 6 (expansion).

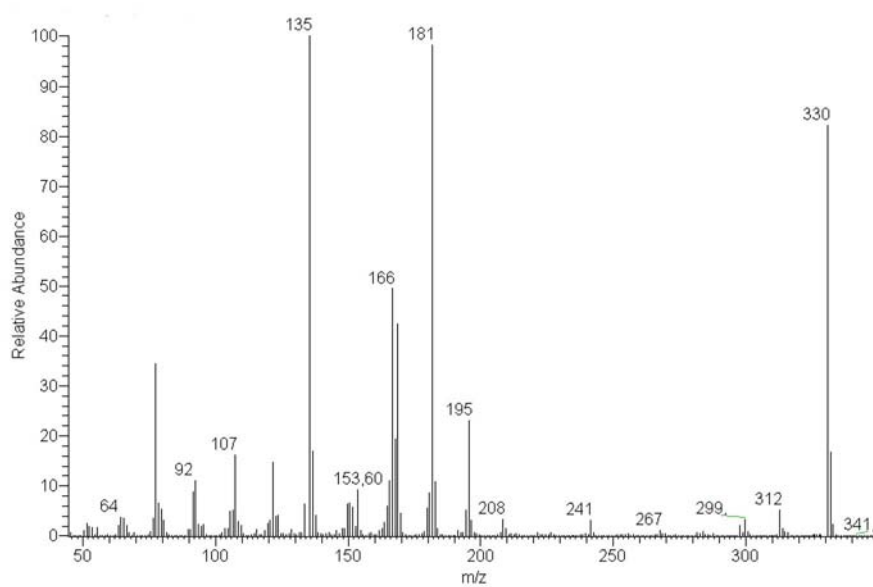


Figure S18. Mass spectrum of compound 6 (CG-MS).

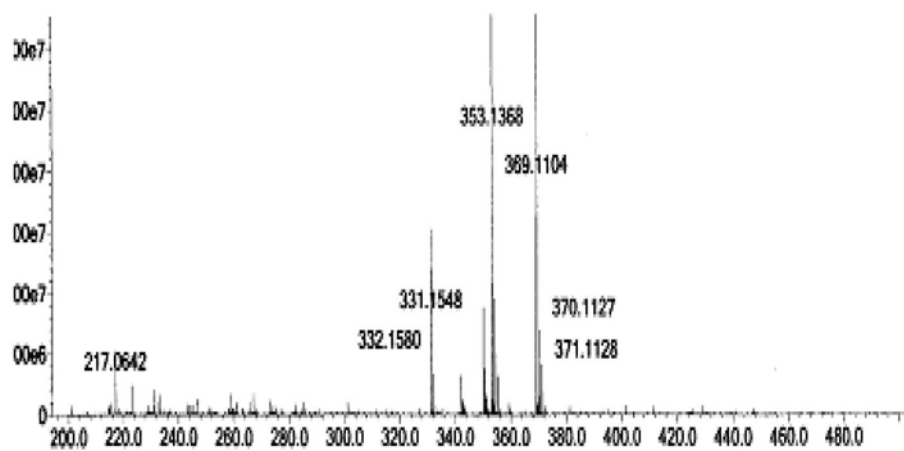


Figure S19. High resolution mass spectrum of compound 6.