

Supporting Information

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Inhibition of iNOS Induction and NF- κ B Activation by Taste Compounds from the Edible Mushroom *Tricholoma caligatum* (Viv.) Ricken

Ebru Erol^{1,2*}, Zulfiqar Ali², Mehmet Oztürk¹, Shabana Khan^{2,3} and
Ikhlas A. Khan^{2,4}

¹Mugla Sitki Kocman University, Faculty of Science, Department of Chemistry, Mugla 48121,
Türkiye

²National Center for Natural Products Research, School of Pharmacy, The University of
Mississippi, University, MS 38677, USA

³Department of Biomolecular Sciences, School of Pharmacy, The University of Mississippi,
University, MS 38677, USA

⁴Department of Pharmacognosy, School of Pharmacy, The University of Mississippi, University,
MS 38677, USA

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* Corresponding author: E-Mail: e.ebrusimya@gmail.com

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Table S1. Gradient method of the semi-preparative HPLC system.

Time	Flow	%MeOH	%H ₂ O
0.01	3.00	10.0	90.0
10.00	3.00	20.0	80.0
20.00	3.00	30.0	70.0
25.00	3.00	50.0	50.0
35.00	3.00	100.0	0.0
60.00	3.00	100.0	0.0

Compounds **1-3** were purified on a reversed-phase Phenomenex C18 column (Gemini® 5 µm C18 110 Å, LC Column 250 x 21.2 mm, AXIA™ Packed) using a Waters 2795 HPLC system equipped with Photodiode Array detector. The column was eluted using a linear gradient mode with two different solvent: (A) MeOH, (B) H₂O, at a flow rate of 3.0 mL/min. The initial percent of solvent A was 20%. The percent of solvent A was constant for the initial 2 min, increased to 20% at 0–10 min, increased to 30% at 10–20 min, increased to 50% at 20–25 min, increased to 100% at 25–35 min, and remained constant at 100% at 35–60 min (Table S1). The elution was monitored using a wavelength of 354, 254 and 210 nm.

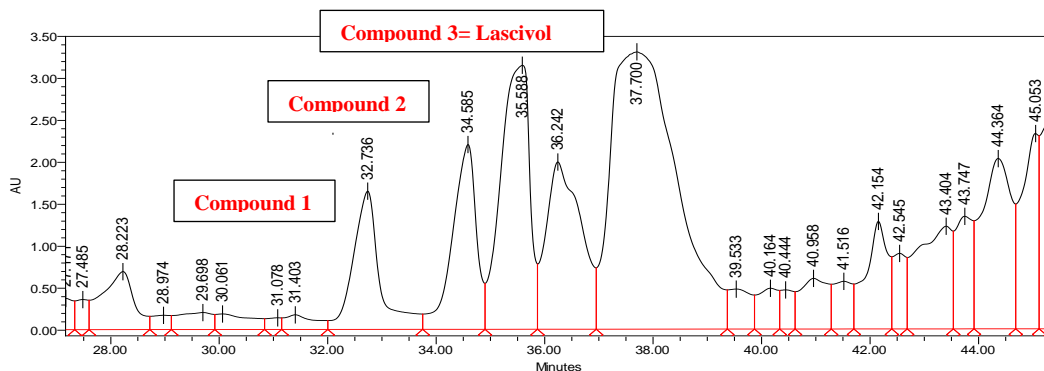


Figure S1: HPLC chromatogram of compounds **1-3**.

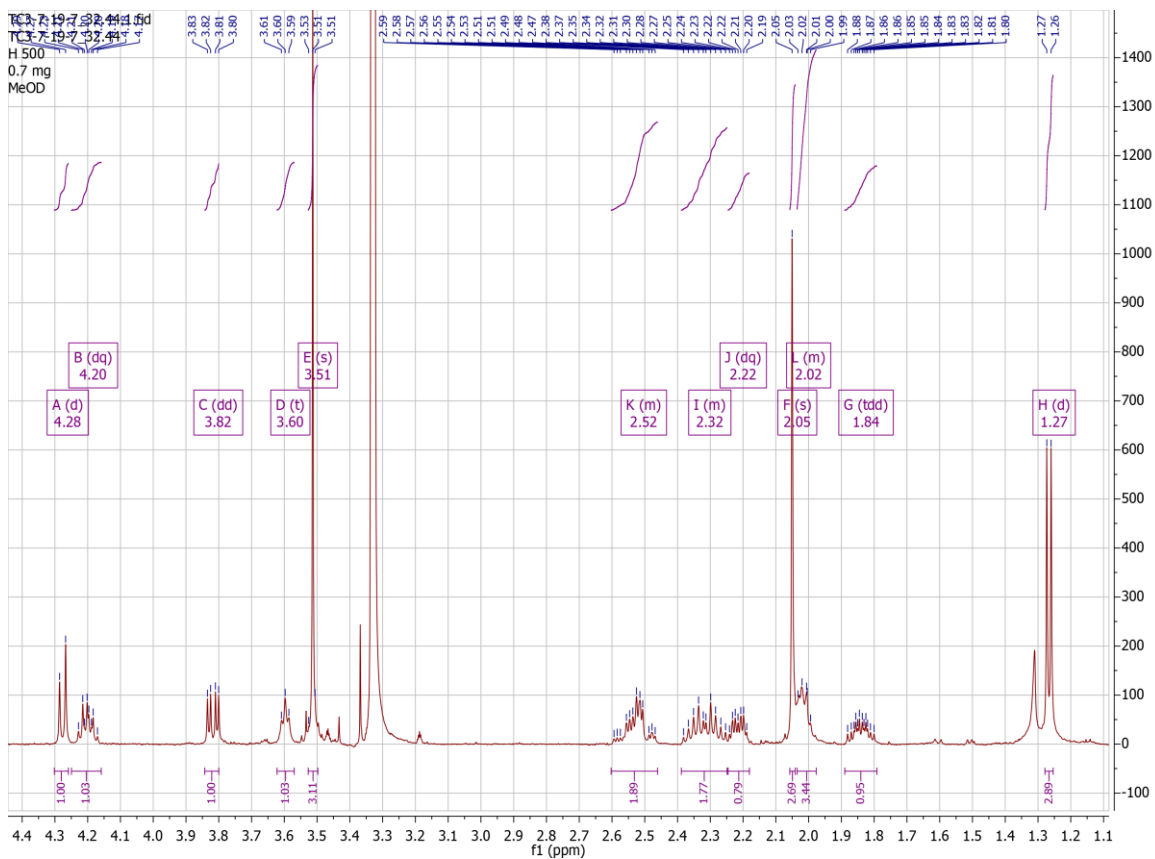


Figure S3: ^1H NMR (500 MHz, CD_3OD) spectrum of **1** expanded from 1.1 to 4.4 ppm

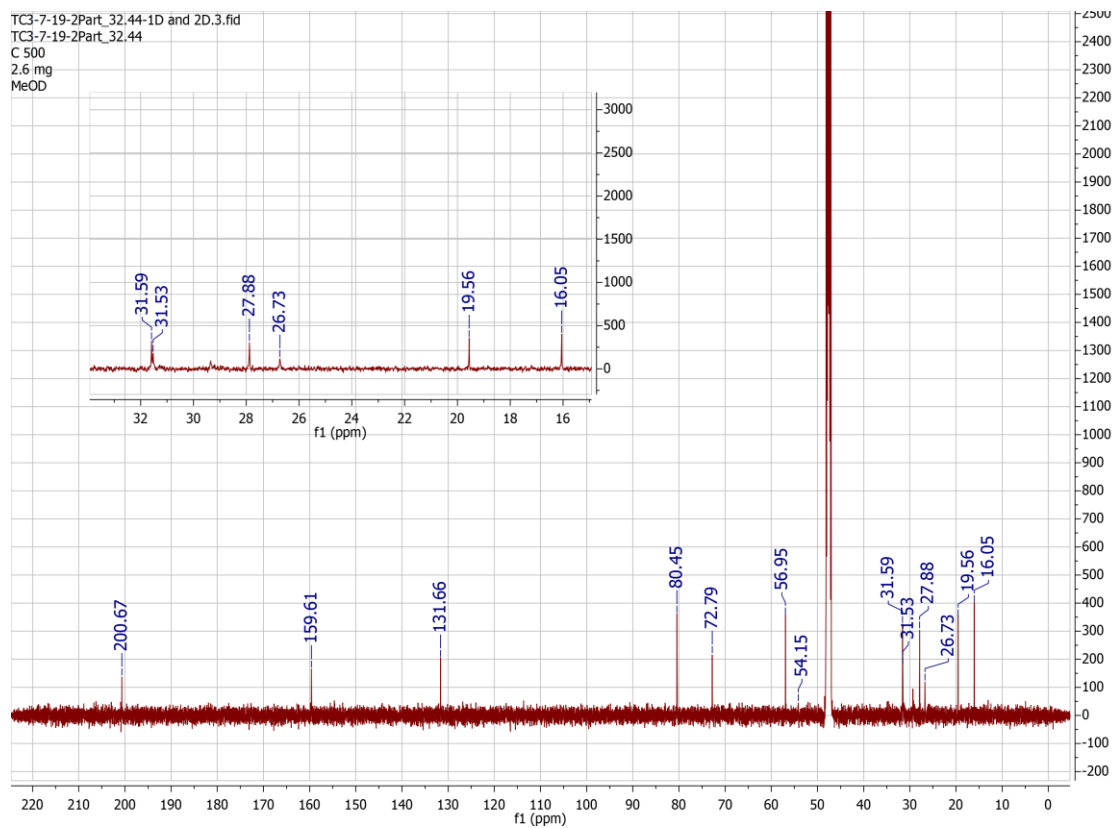


Figure S4: ^{13}C NMR (500 MHz, CD_3OD) spectrum of **1**.

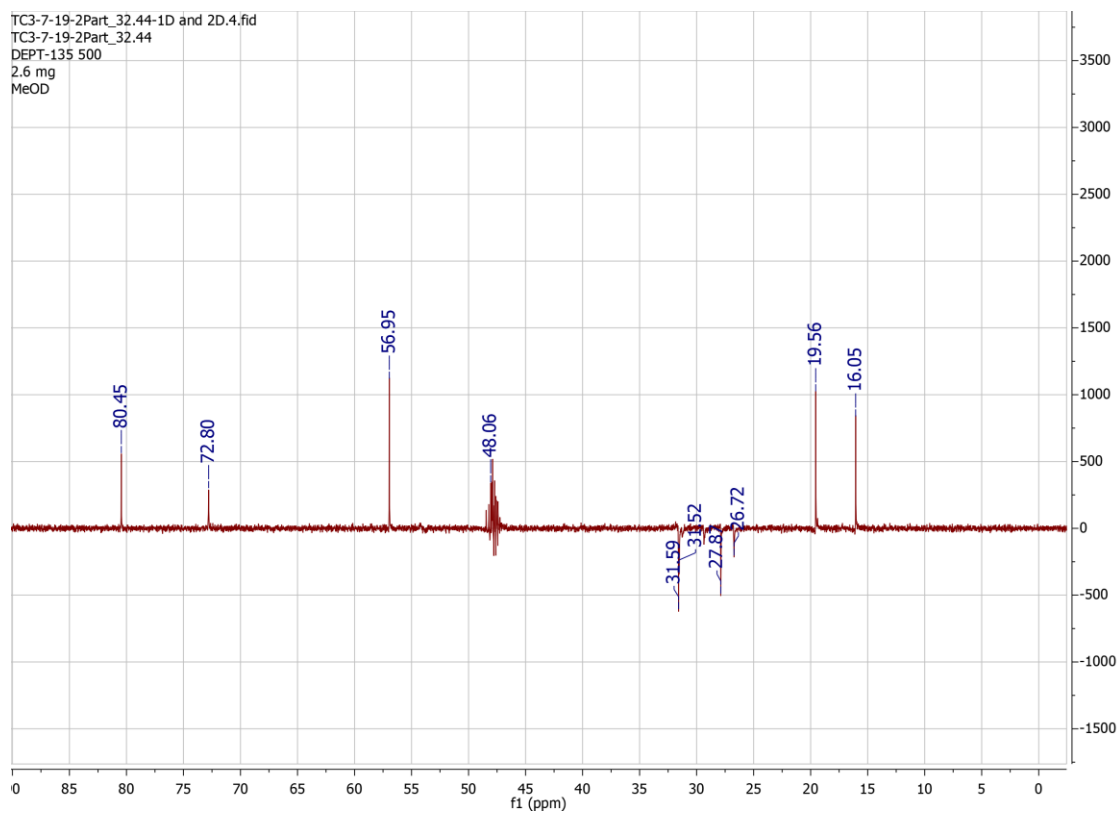


Figure S5: DEPT 135 spectrum of **1**.

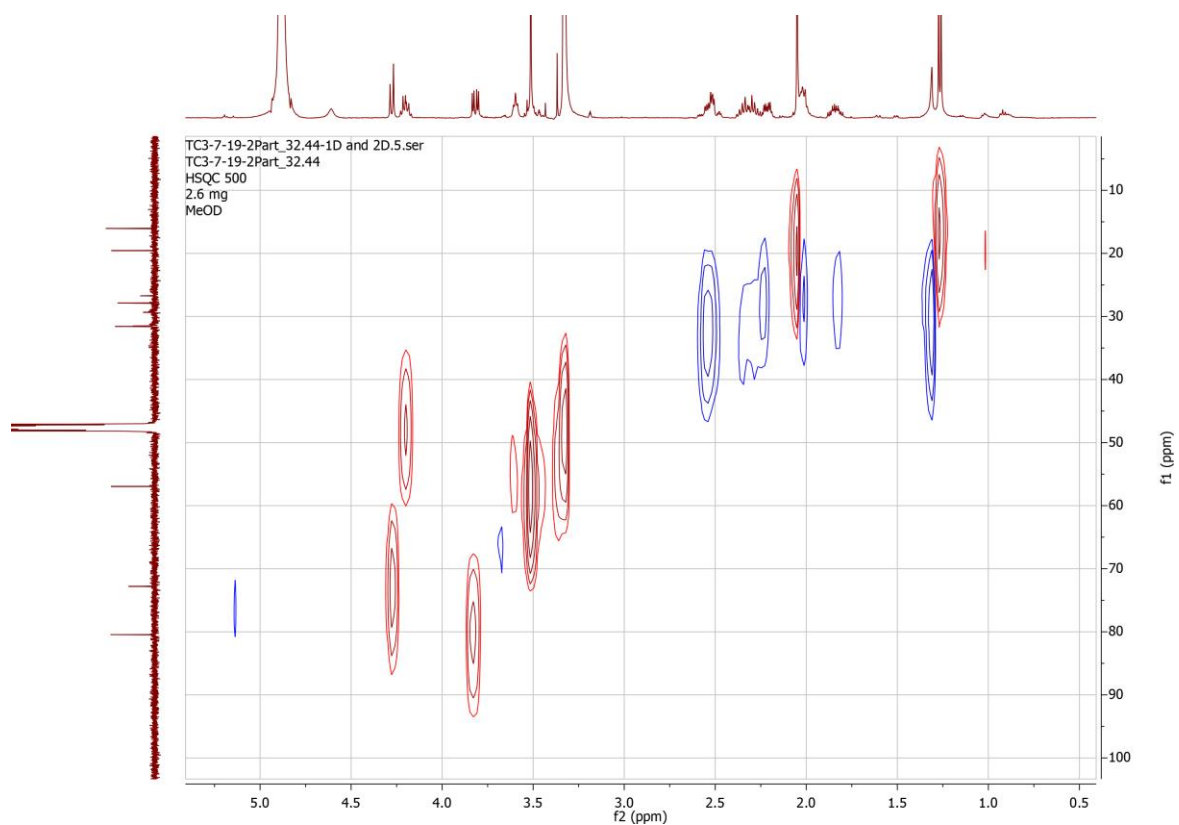


Figure S6: HSQC spectrum of 1.

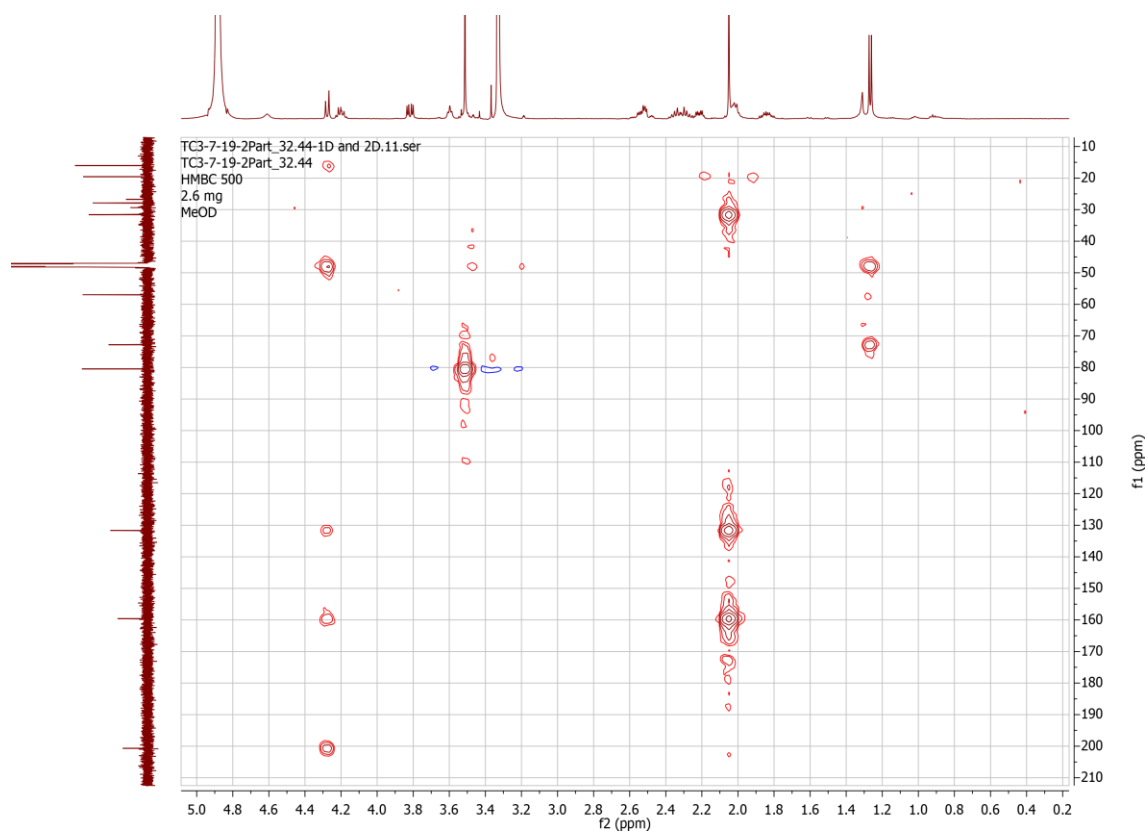


Figure S7: HMBC spectrum of 1.

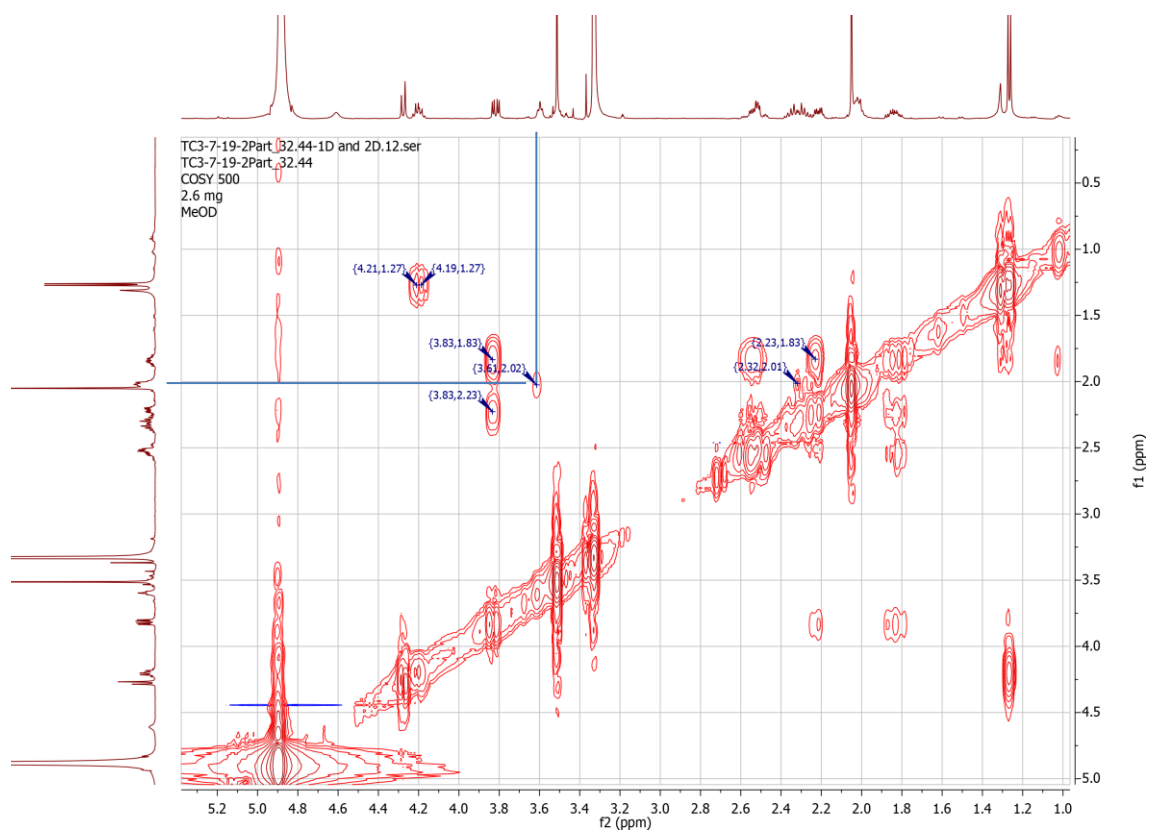


Figure S8: COSY spectrum of **1**.

Sample Name (+)Ebru-TC3-7-19-2Pa Position Vial 4 Instrument Name Instrument 1 User Name
Inj Vol 10 InjPosition Sample IRM Calibration Status Success
Data Filename (+)Ebru-TC3-7-19-2Pa ACQ Method No Column Positive.m Comment 3/5/2018 8:51:02 AM

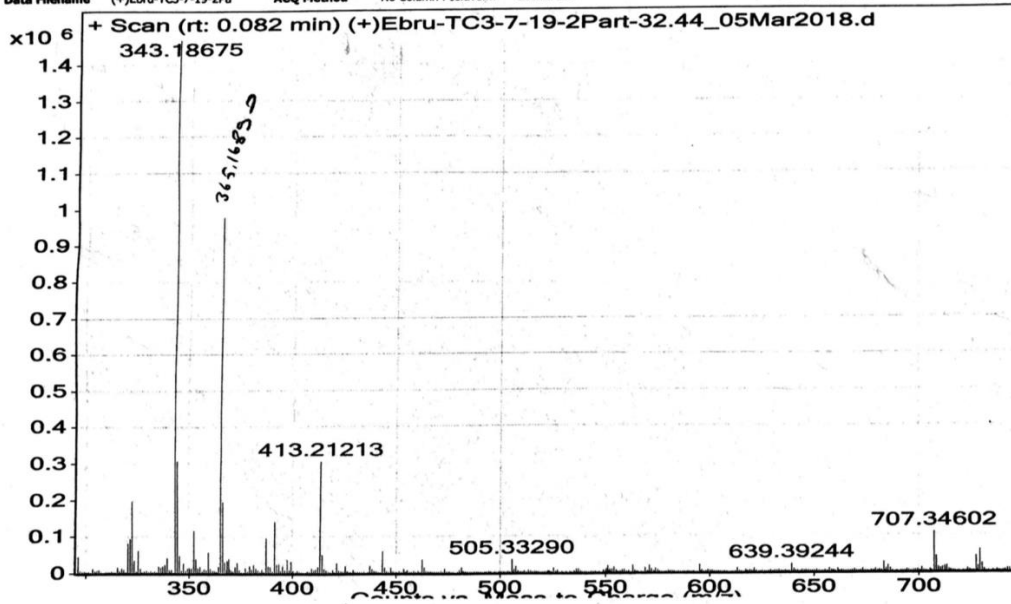


Figure S9: Mass spectrum of 1.

Sample ID:TC3-7-19-2Part_32.44 Method Name:galal 1
Sample Scans:8 User:galal
Background Scans:8 Date/Time:03/07/2018 3:23:39 PM
Resolution:4 Range:4000 - 650
System Status:Good Apodization:Happ-Genzel
File Location:C:\Program Files\Agilent\MicroLab PC\Results\galal 1\TC3-7-19-2Part_32.44_2018-03-07T15-23-39.a2r

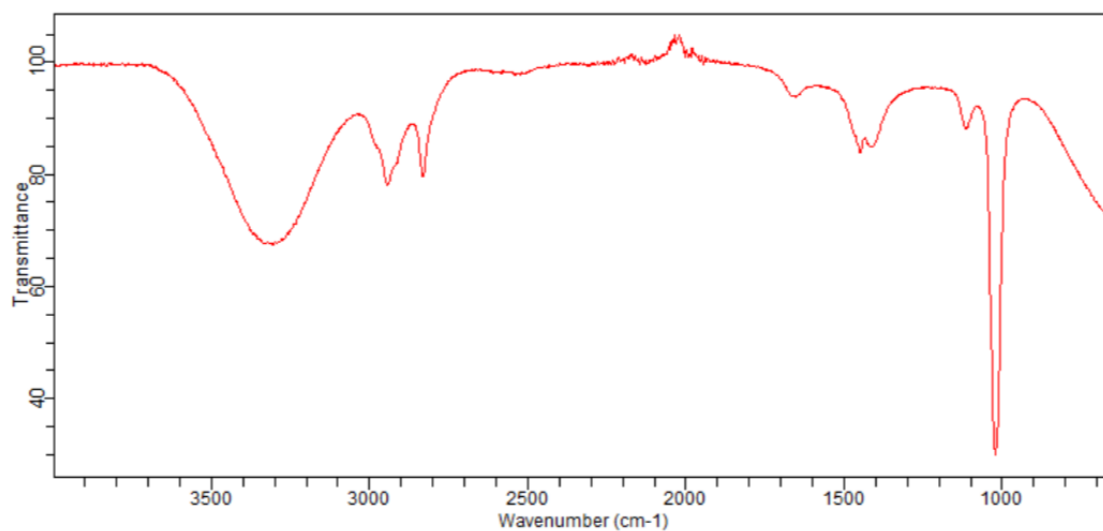


Figure S10: IR spectrum of **1**.

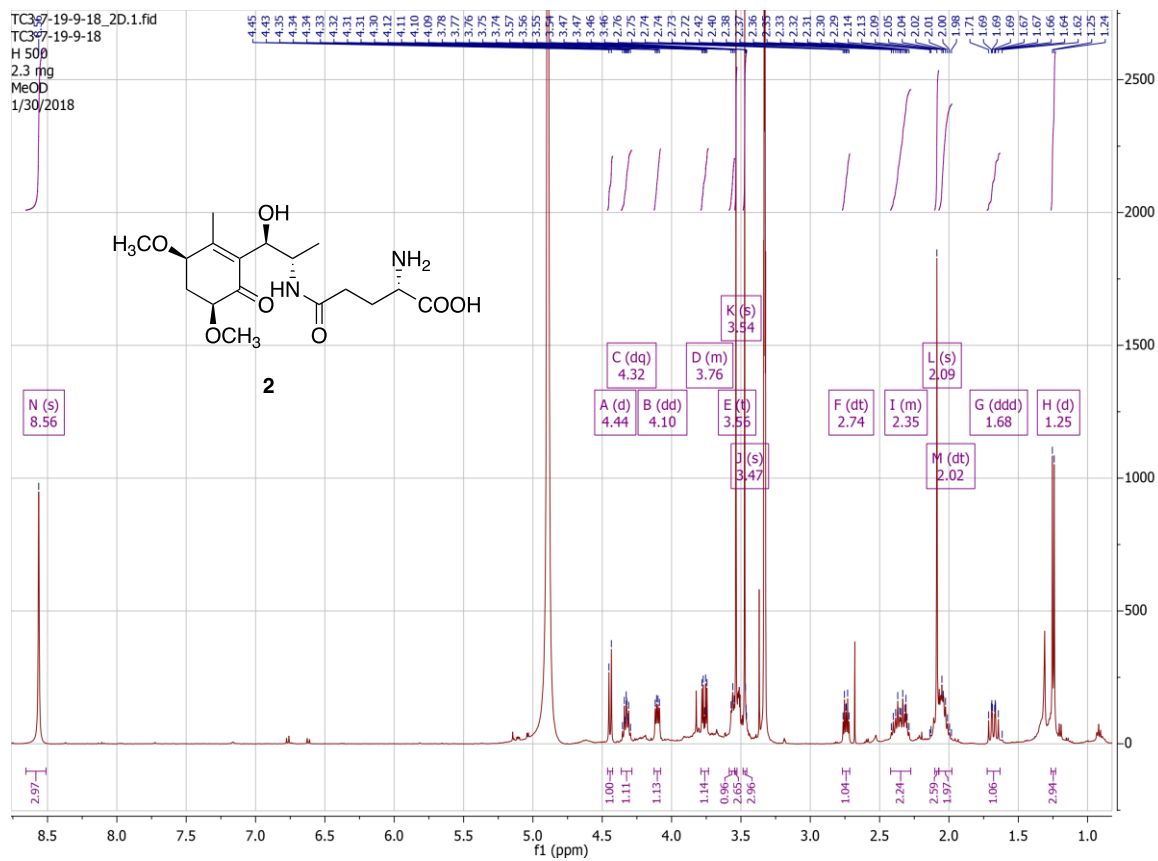


Figure S11: ^1H NMR (500 MHz, CD_3OD) spectrum of **2**.

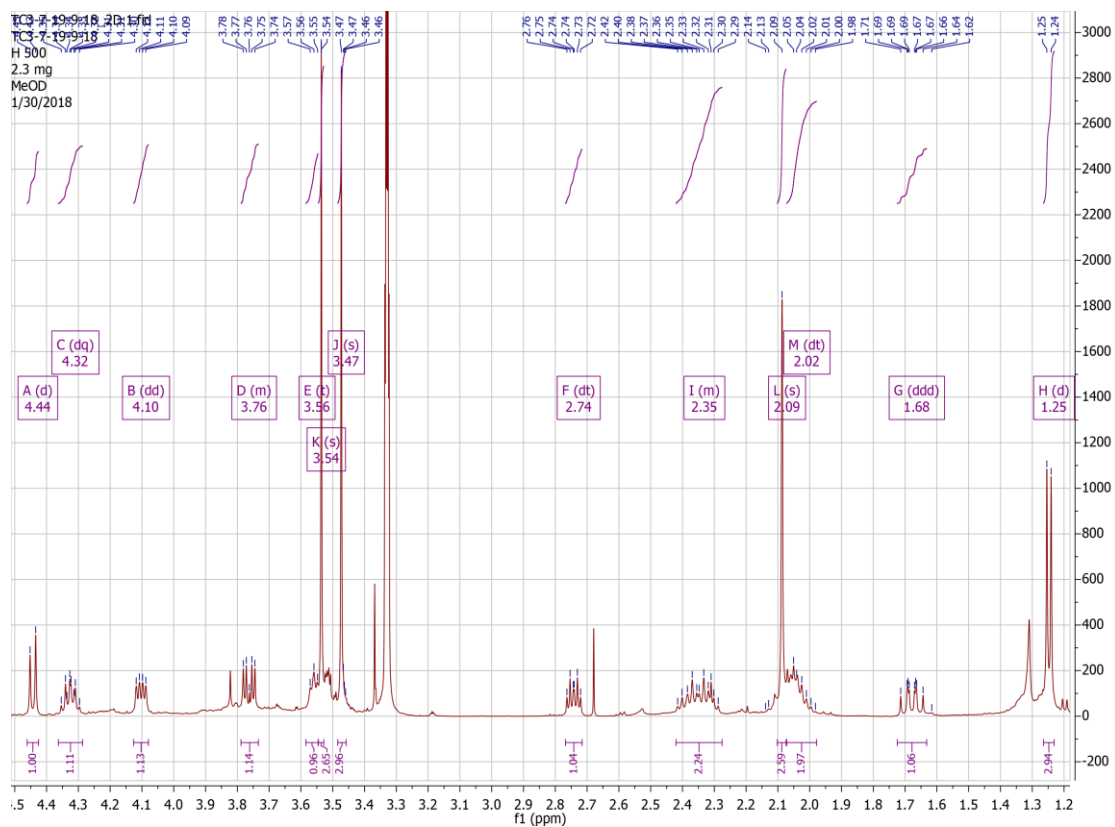


Figure S12: ^1H NMR (500 MHz, CD_3OD) spectrum of **2** expanded from 1.2 to 4.5 ppm.

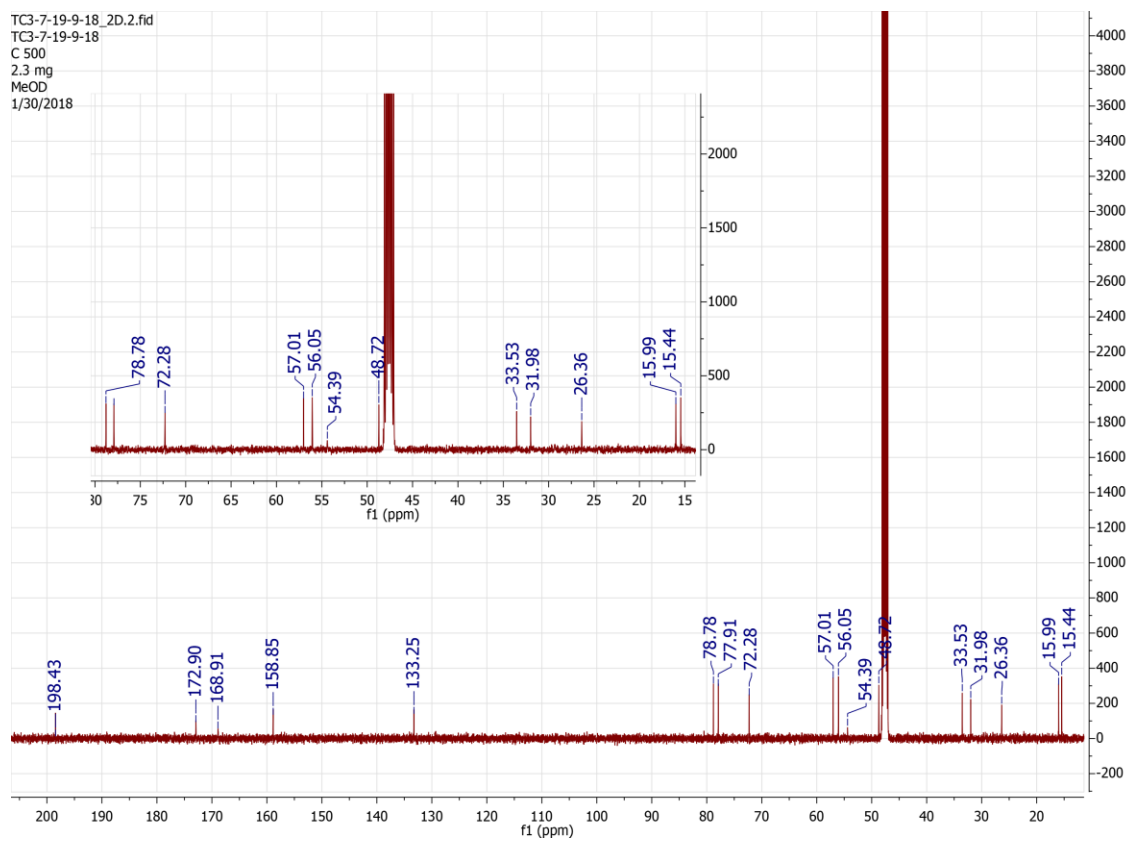


Figure S13: ^{13}C NMR (500 MHz, CD_3OD) spectrum of **2**.

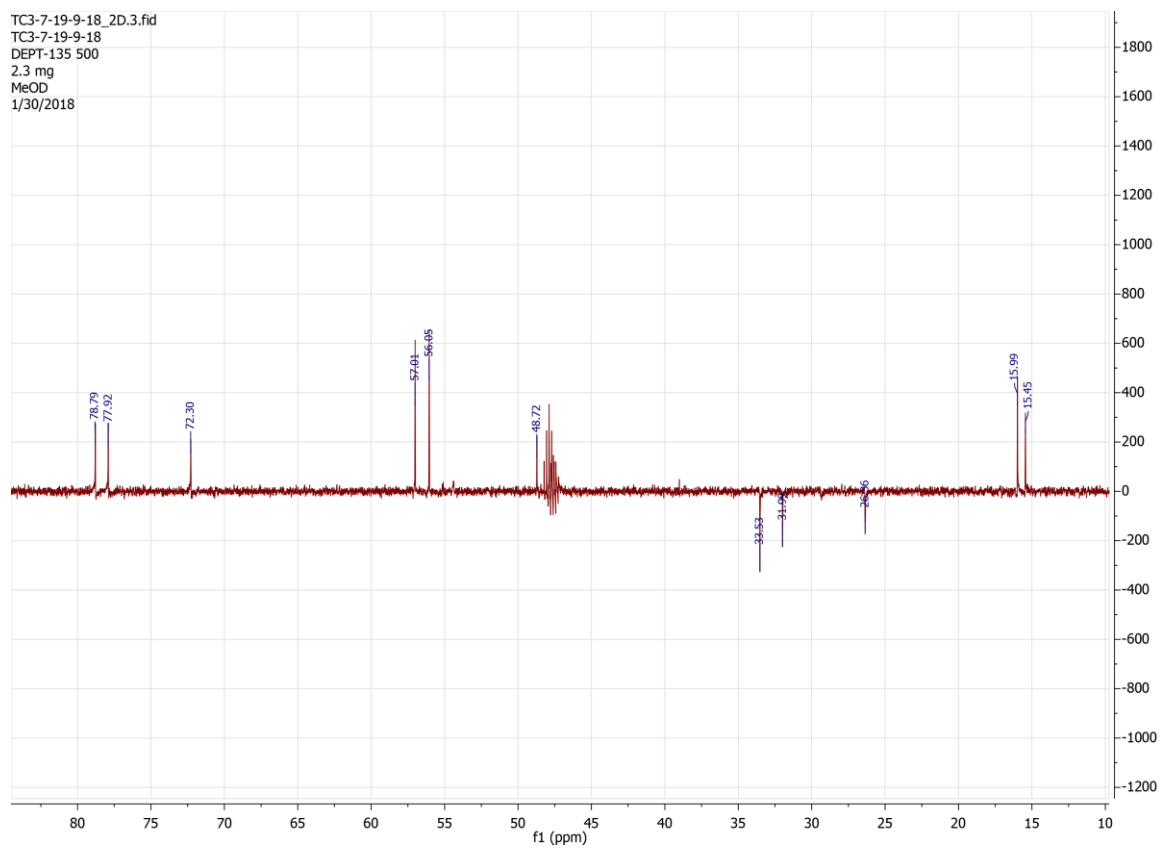


Figure S14: DEPT 135 spectrum of **2**.

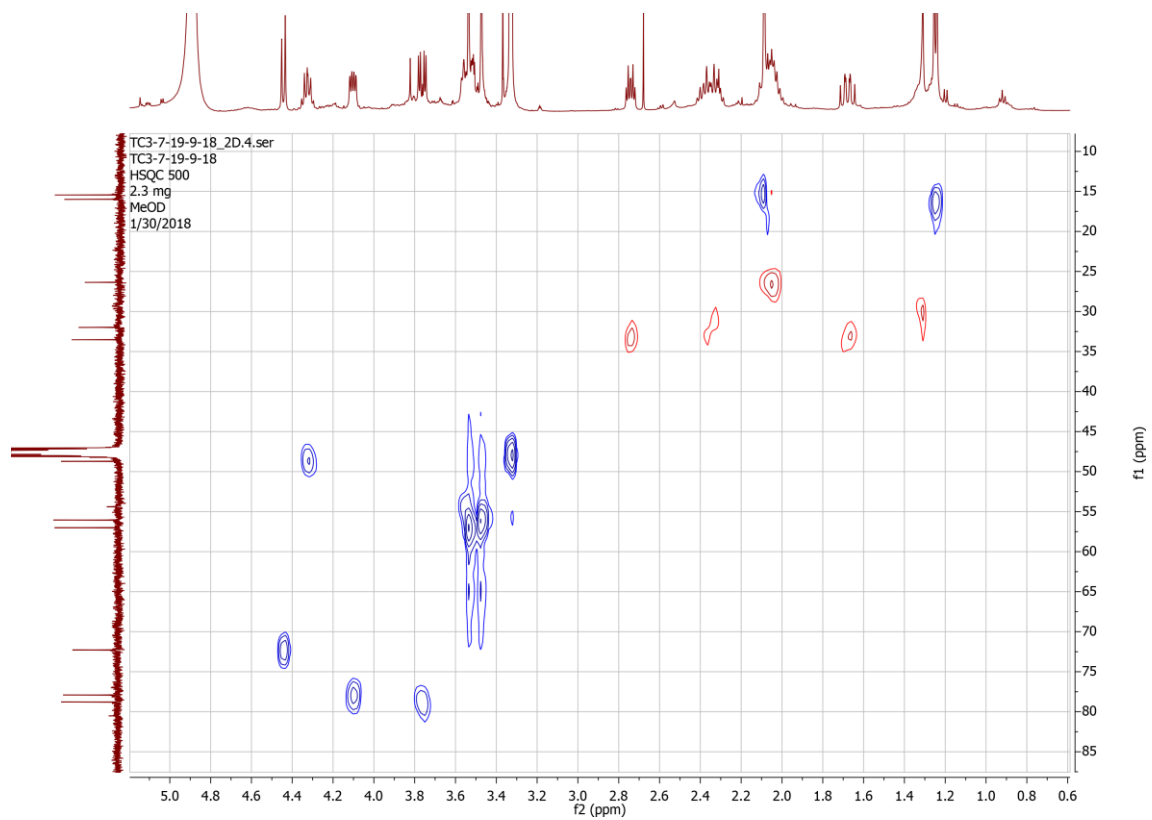


Figure S15: HSQC spectrum of 2.

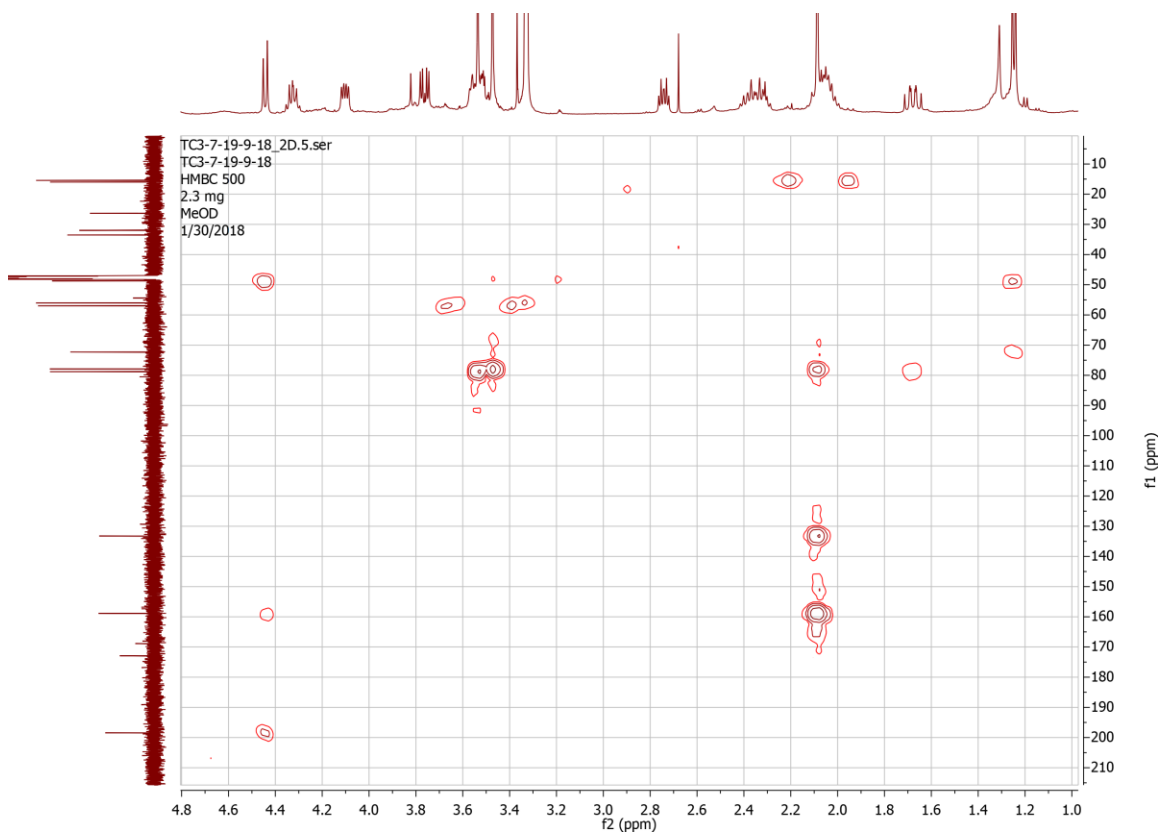


Figure S16: HMBC spectrum of **2**.

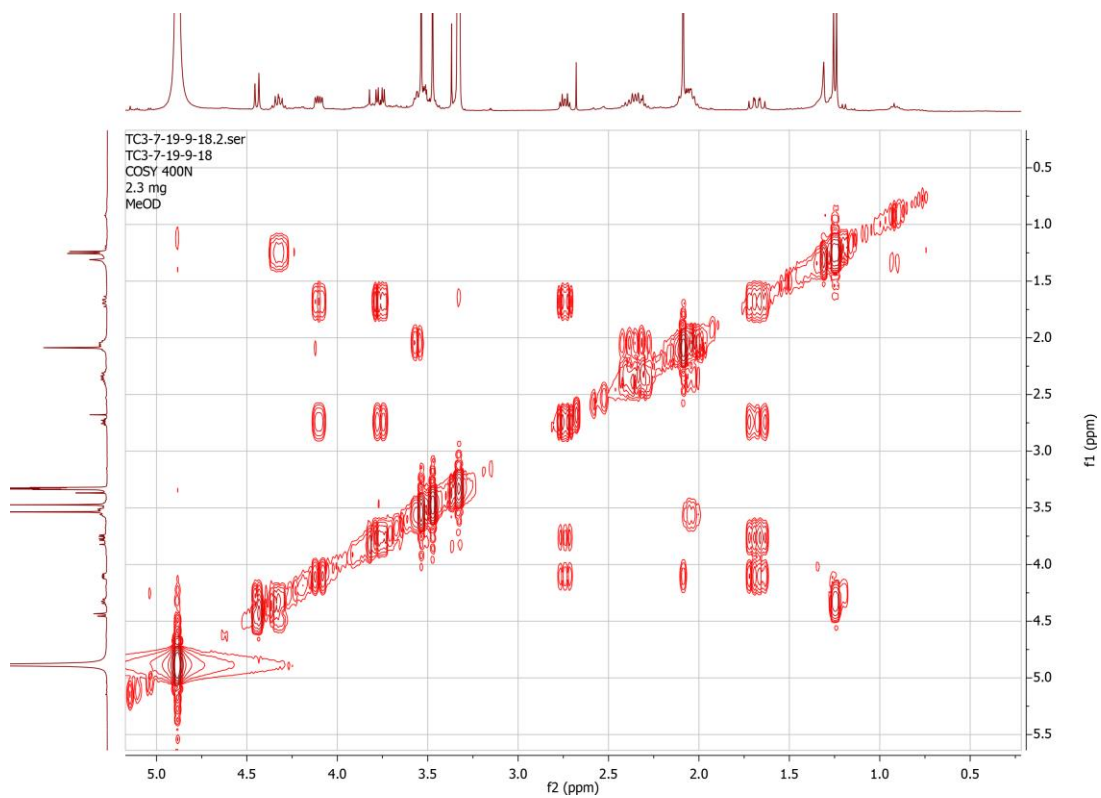


Figure S17: COSY spectrum of **2**.

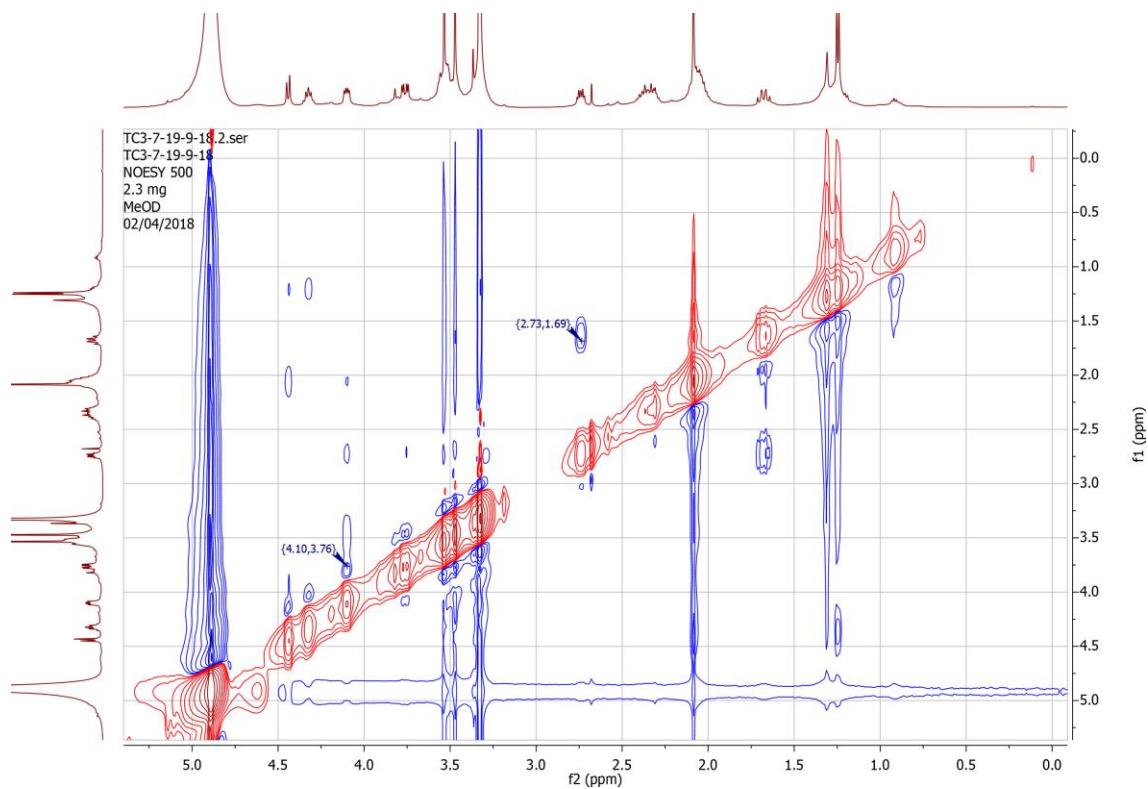


Figure S18: NOESY spectrum of **2**.

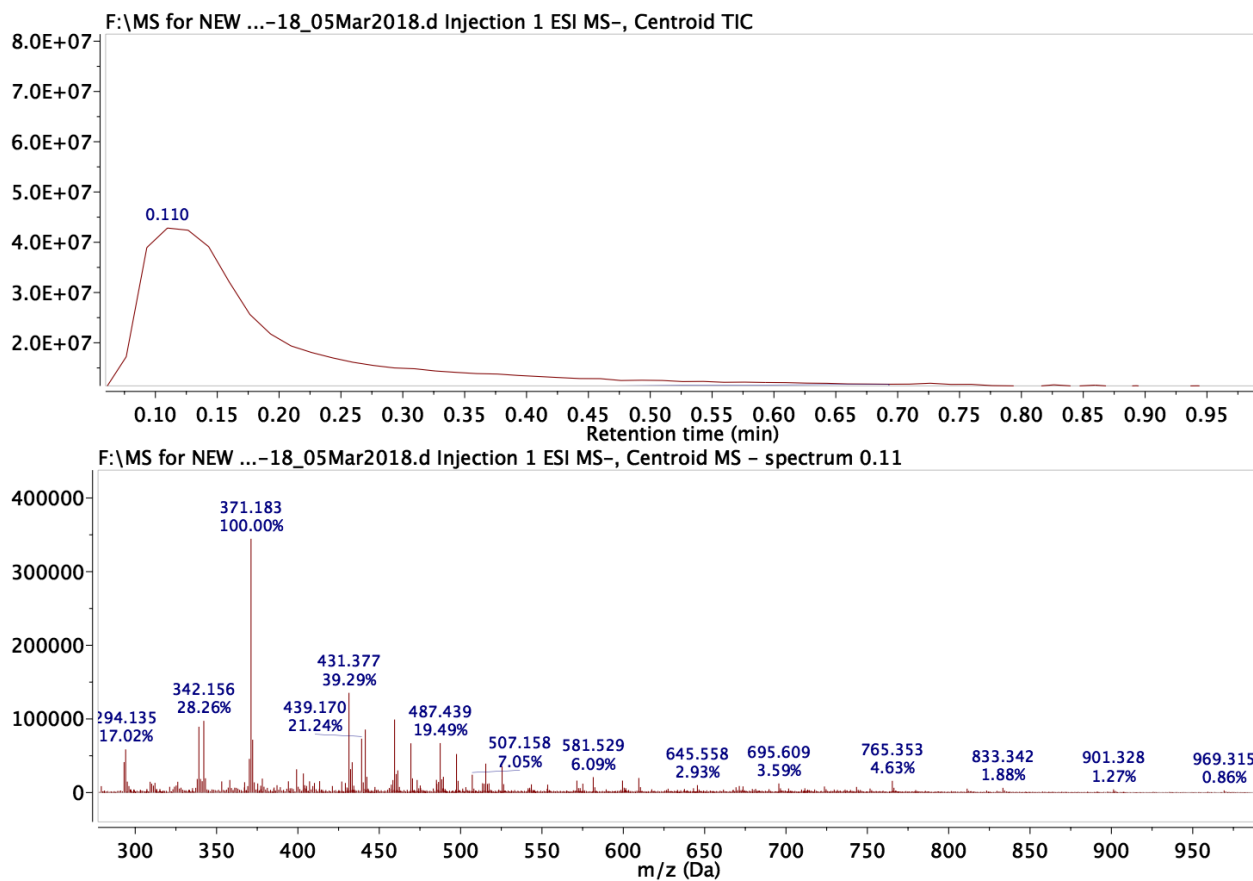


Figure S19: Mass spectrum of 2.



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Sample Scans:8
Background Scans:8
Resolution:4
System Status:Good
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Method Name:galal 1
User:galal
Date/Time:02/20/2018 5:54:50 PM
Range:4000 - 650
Apodization:Happ-Genzel

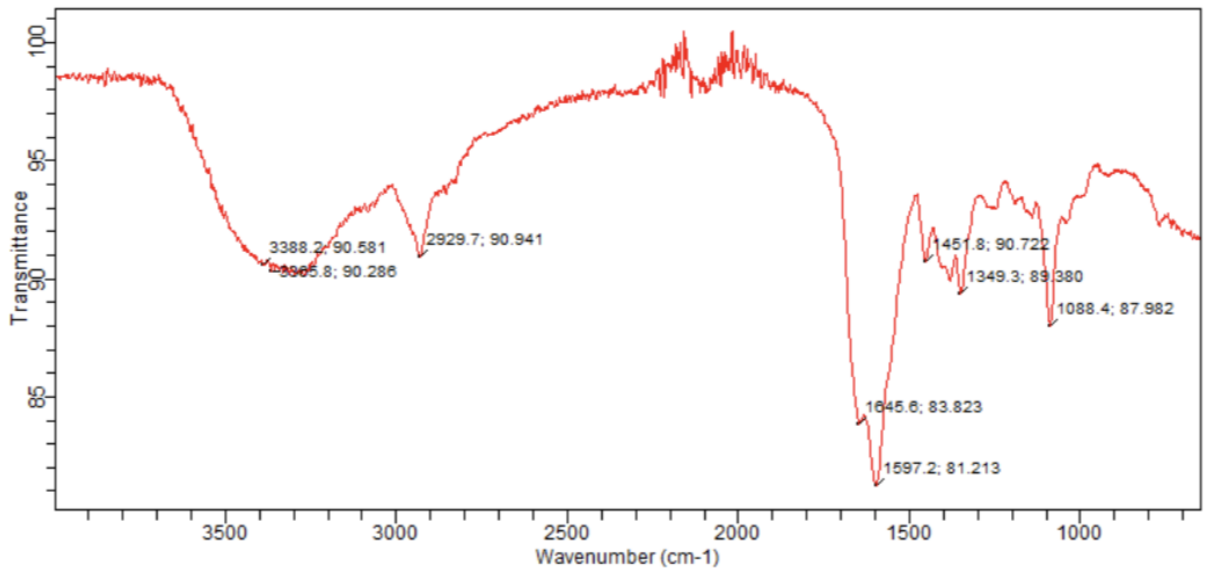


Figure S20: IR spectrum of **2**.

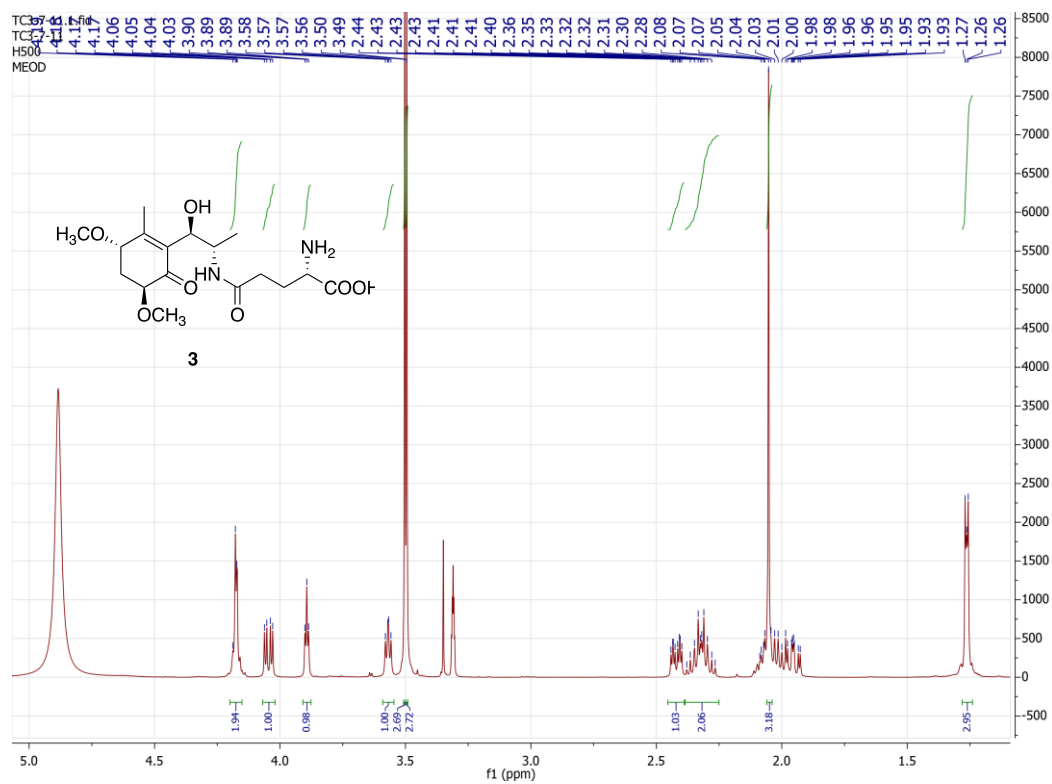


Figure S21: ¹H NMR (500 MHz, CD₃OD) spectrum of **3**.

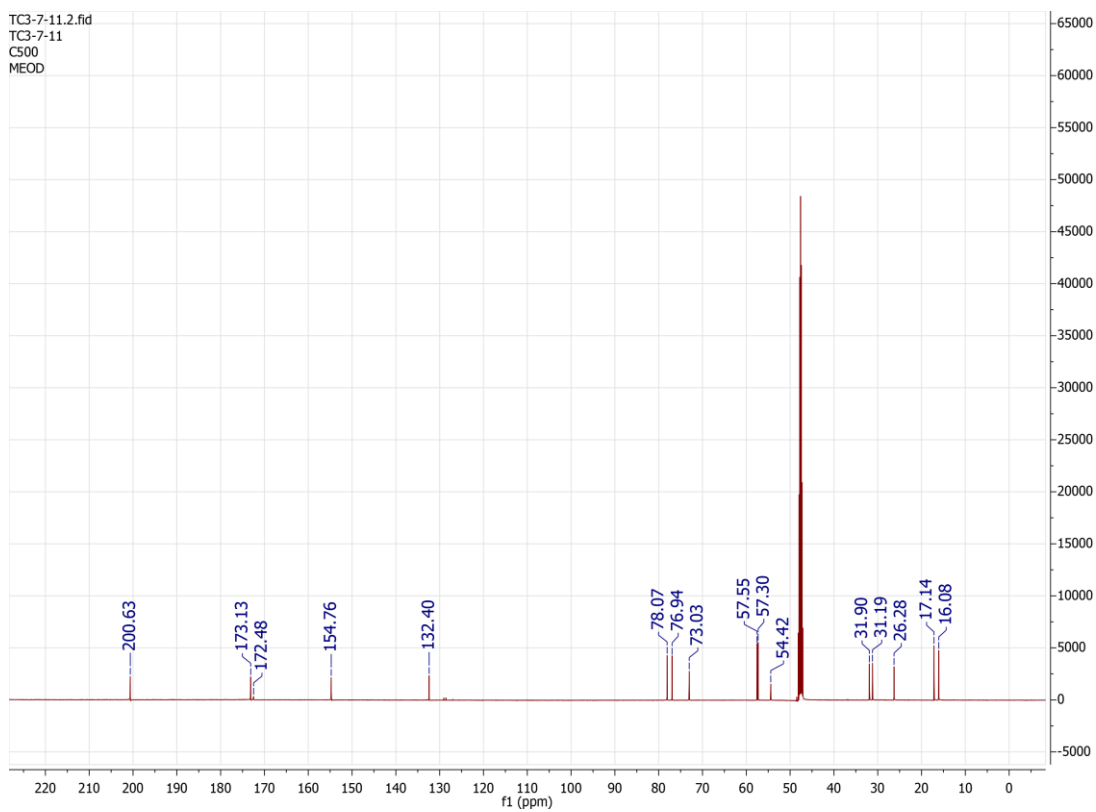


Figure S22: ^{13}C NMR (500 MHz, CD_3OD) spectrum of **3**.

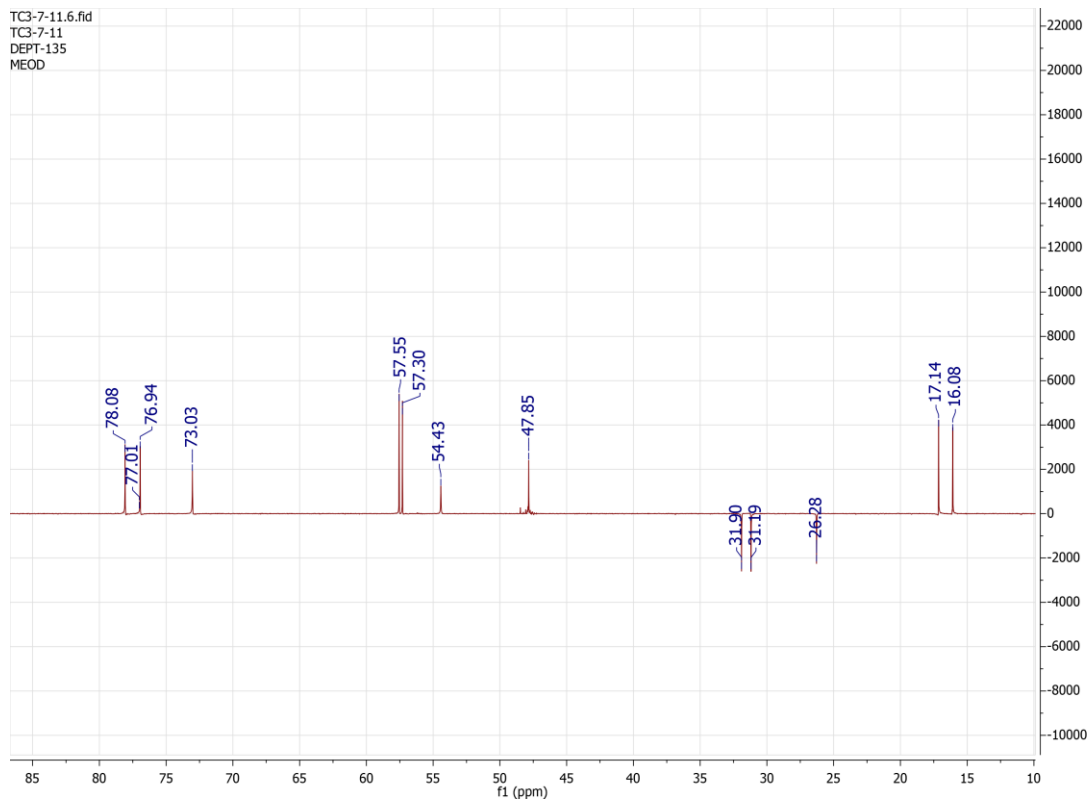


Figure S23: DEPT 135 spectrum of **3**.

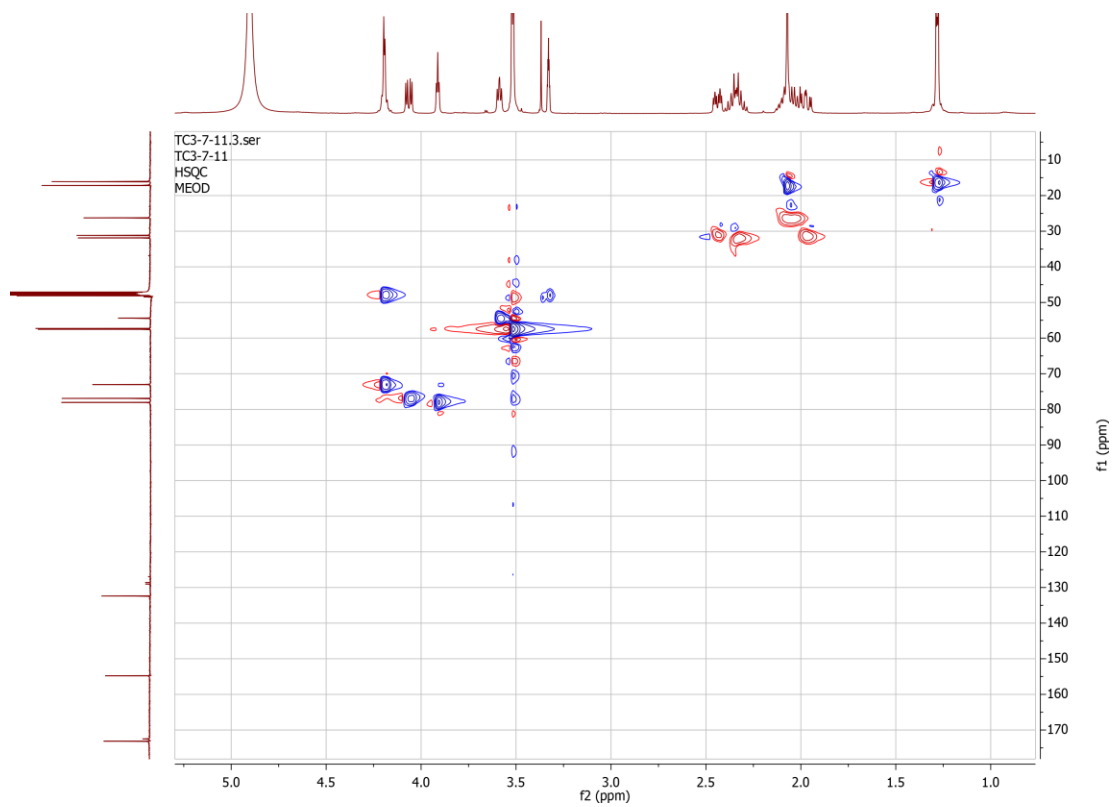


Figure 24: HSQC spectrum of **3**.

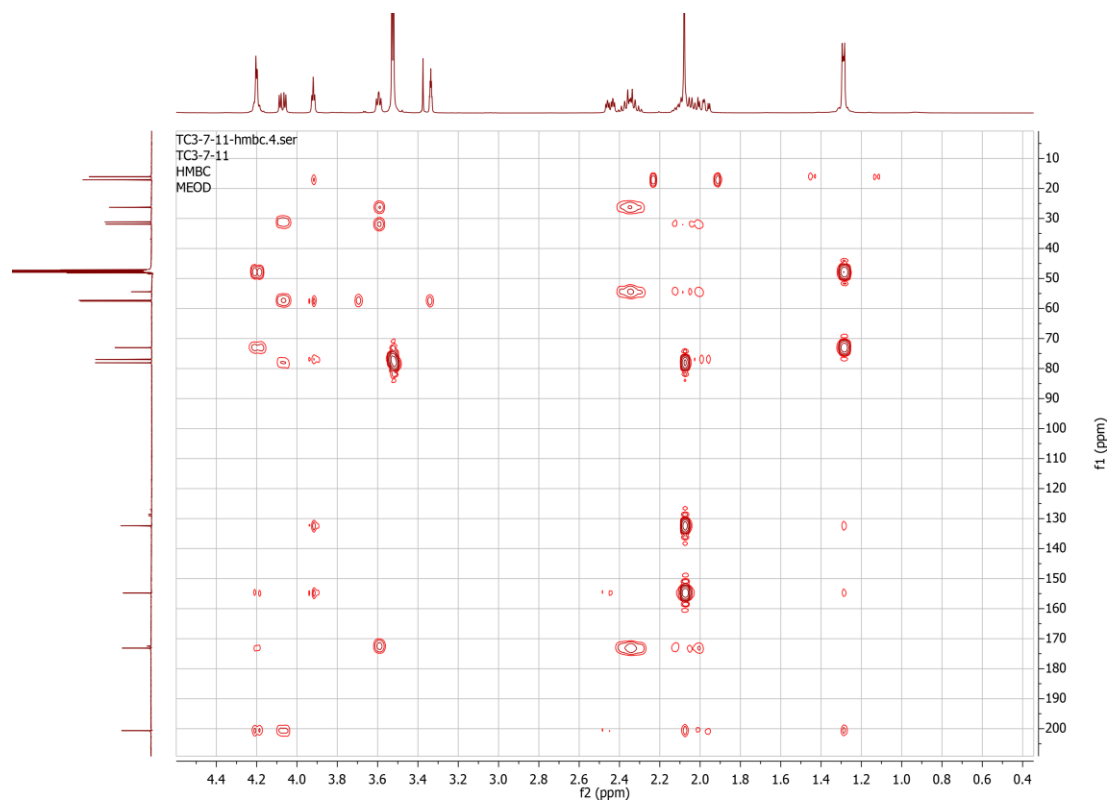


Figure S25: HMBC spectrum of **3**.

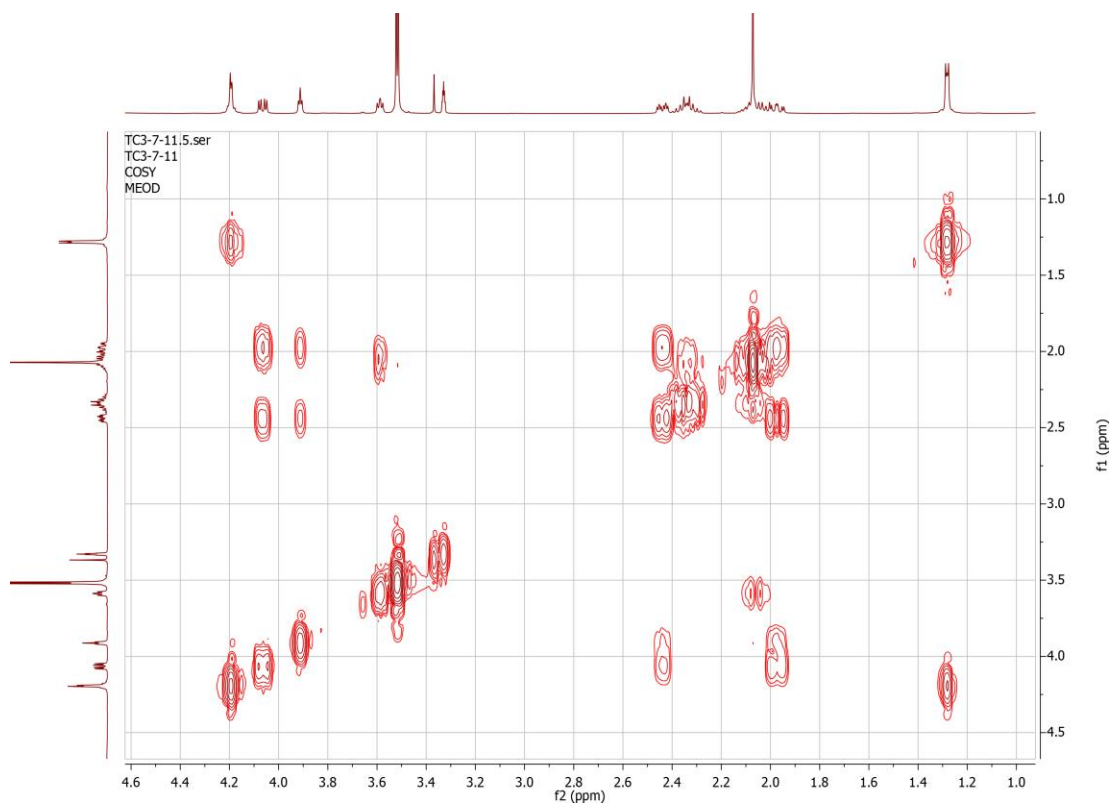


Figure S26: COSY spectrum of **3**.

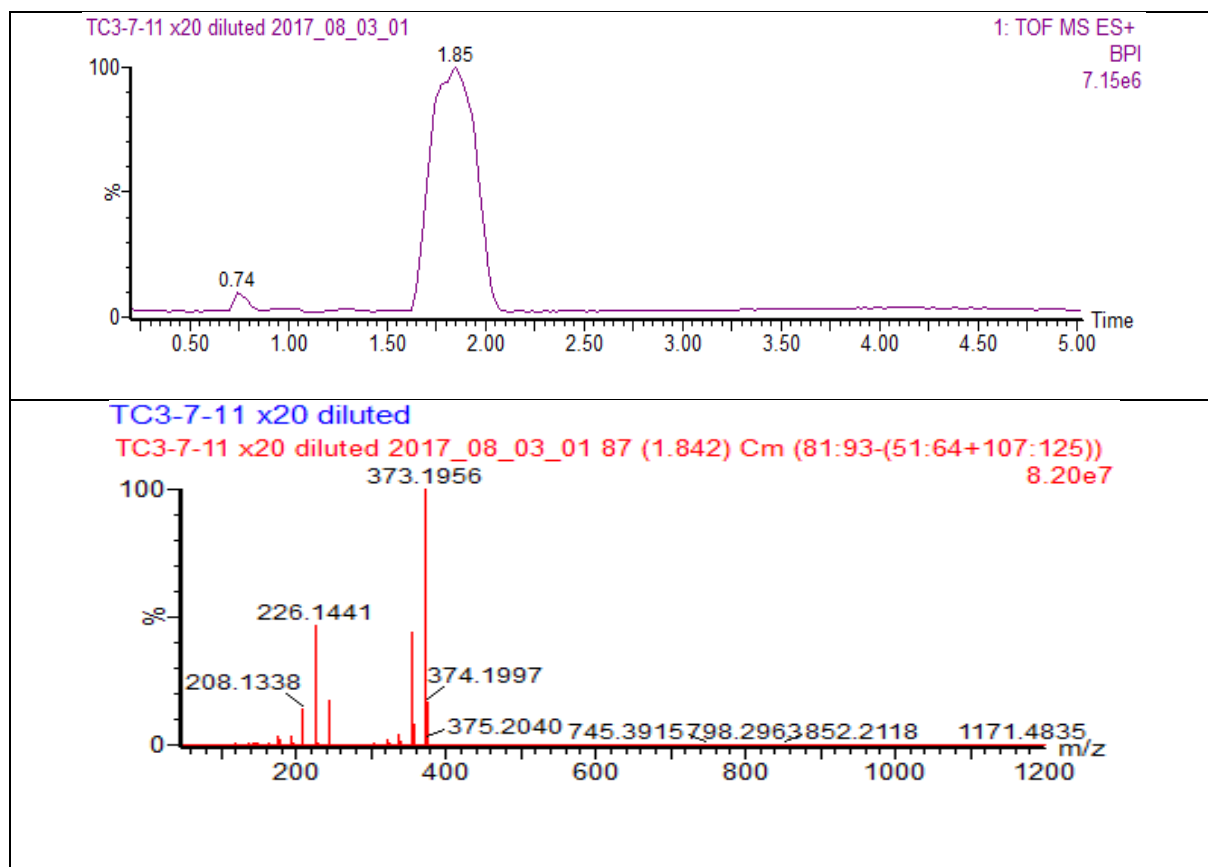


Figure S27: Mass spectrum of **3**.



Sample ID:TC3-7-11

Sample Scans:8

Background Scans:8

Resolution:4

System Status:Good

File Location:C:\Program Files\Agilent\MicroLab PC\Results\galal 1\TC3-7-11_2018-02-20T17-42-57.a2r

Method Name:galal 1

User:galal

Date/Time:02/20/2018 5:42:57 PM

Range:4000 - 650

Apodization:Happ-Genzel

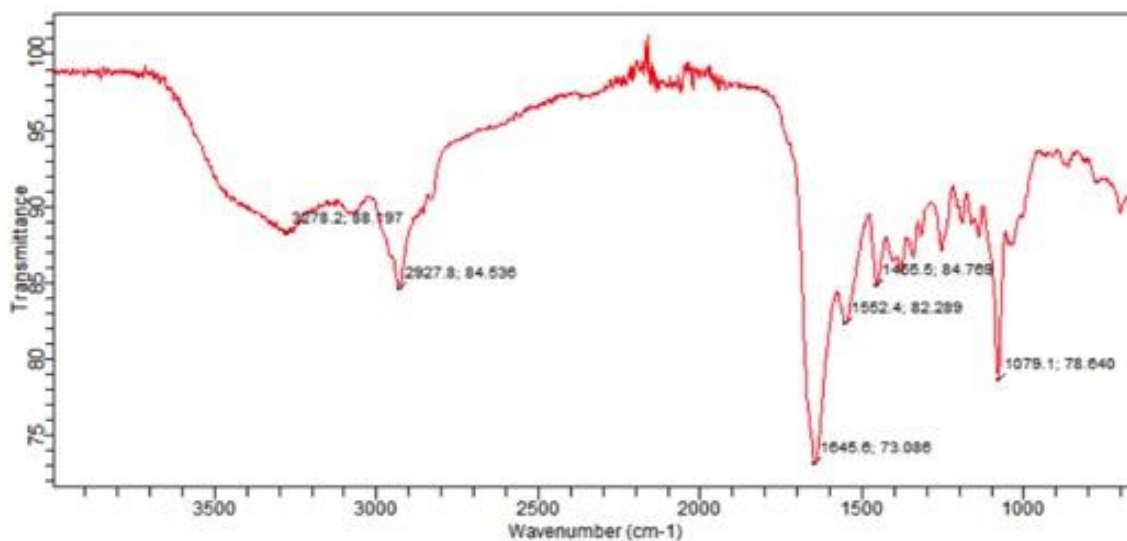


Figure S28: IR spectrum of 3.