

Supplementary data

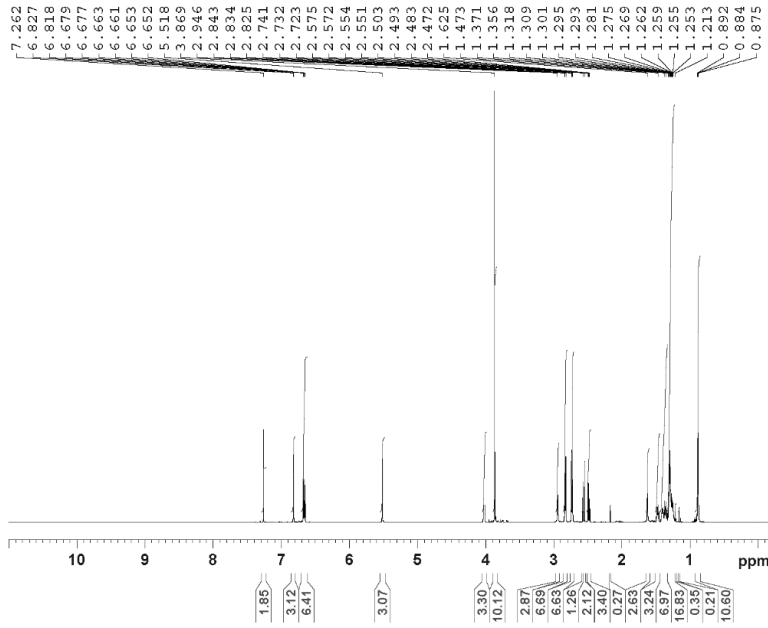
Phenolics isolated from *Aframomum meleguta* enhance proliferation and ossification markers in bone cells

Ashraf B. Abdel-Naim ^{1,2}, Abdullah A. Alghamdi ³, Mardi M. Algandaby ^{1,4},

Fahad A. Al-Abbasi ^{1,3}, Ahmed M. Al-Abd ^{2,5}, Hossam M. Abdallah ^{6,7},

Ali M. El-Halawany ⁷, Masao Hattori ⁸

Dr. Hossam
Sample : A-5 CDCL3



Current Data Parameters
NAME HOGSAM A-5 18-05-2015
EXPNO 10
PROCNO 1

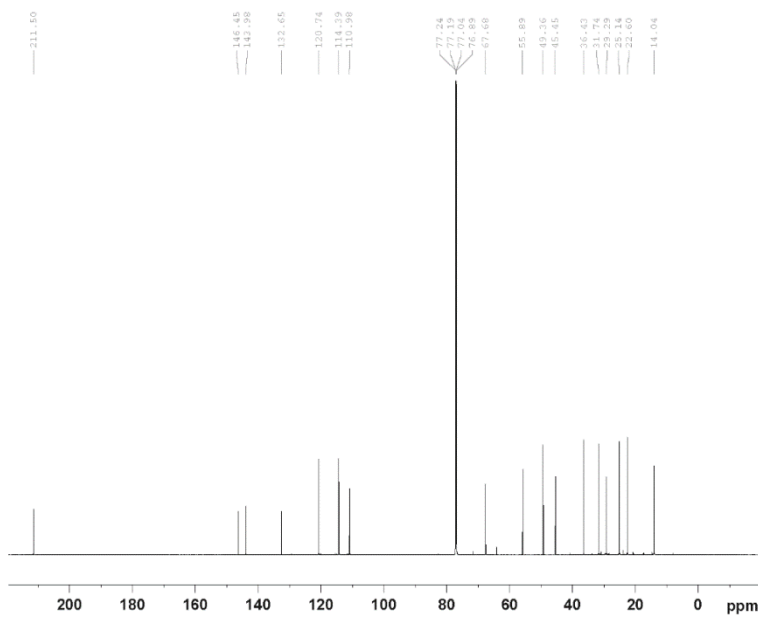
F2 - Acquisition Parameters
Date_ 20160528
Time 20.25
INSTRUM spect
PROBHD 5 mm CPQCI 1H-
PULPROG zg30
TD 65536
SOLVENT CDCl3
NS 20
DS 2
SWH 17006.803 Hz
FIDRES 0.255003 Hz
AQ 1.9267584 sec
RG 9.04
DM 29.400 usec
DE 10.00 usec
TE 298.2 K
D1 1.00000000 sec
TDO

***** CHANNEL f1 *****
SFO1 850.1552500 MHz
NUC1 1H
P1 6.00 usec
PLM1 15.30000019 W

F2 - Processing parameters
SI 65536
SF 850.1500200 MHz
WDW EM
SSB 0
LB 0.30 Hz
GB 0
PC 2.00



Dr. Hossam
Sample : A-5 CDCL3



Current Data Parameters
NAME HOGSAM A-5 04-06-2015
EXPNO 10
PROCNO 1

F2 - Acquisition Parameters
Date_ 20160605
Time 18.40
INSTRUM spect
PROBHD 5 mm CPQCI 1H-
PULPROG zgpg30
TD 65536
SOLVENT CDCl3
NS 2500
DS 4
SWH 51020.406 Hz
FIDRES 0.770810 Hz
AQ 0.6422528 sec
RG 186.93
DM 9.800 usec
DE 18.00 usec
TE 298.2 K
D1 2.00000000 sec
L11 0.03000000 sec
TDO 1

***** CHANNEL f1 *****
SFO1 213.7217436 MHz
NUC1 13C
P1 12.00 usec
PLM1 130.00000000 W

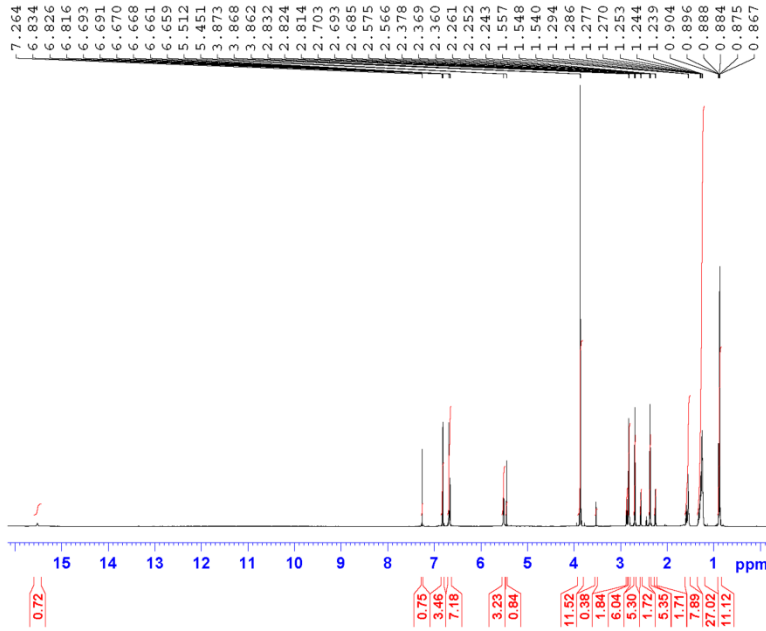
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NUC2 1H
CDPRG2 waltz16
SFO2 80.00 usec
PLM2 13.00000019 W
PLM12 0.13800000 W
PLM13 0.09832000 W

F2 - Processing parameters
SI 32768
SF 213.7703875 MHz
WDW EM
SSB 0
LB 1.50 Hz
GB 0
PC 2.00



Figure S1. ¹H and ¹³C NMR charts of 6-gingerol
(Samples were recorded on DRX-850 spectrometer (Bruker BioSpin, Billerica, MA, USA)).

Dr.Hossam
Sample : PARADOL CDCL3



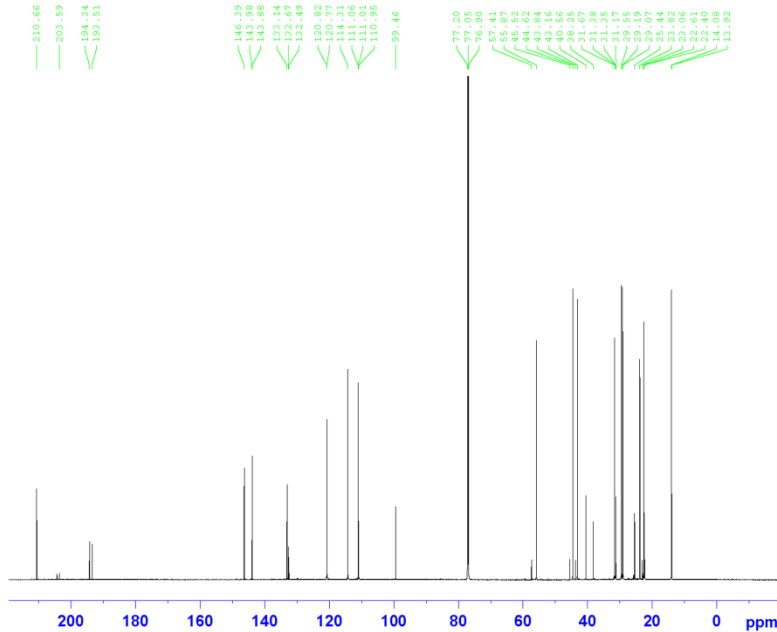
Current Data Parameters
NAME HOSSAM PARADOL 21-01-2015
EXPNO 10
PROCNO 1

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PROBHD 5 mm CPQCI 1H-
PULPROG zgpg30
TD 65536
SOLVENT CDCL3
NS 32
DS 2
SFO1 17006.803 Hz
FIDRES 0.159653 Hz
AQ 1.9267584 sec
RG 7.85
DM 29.460 usec
DE 10.00 usec
TE 298.0 K
D1 1.00000000 sec
TD0 1

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SFO1 850.1552500 MHz
NUC1 1H
P1 8.00 usec
PLM1 15.30000019 H

F2 - Processing parameters
SI 65536
SF 850.1500185 MHz
WDW EM
SSB 0
LB 0.30 Hz
GB 0
PC 2.00

Dr.Hossam
Sample : PARADOL CDCL3



Current Data Parameters
NAME HOSSAM PARADOL 27-01-2015
EXPNO 40
PROCNO 1

F2 - Acquisition Parameters
Date_ 20150127
Time 18:52
INSTRUM spect
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PULPROG zgpg30
TD 65536
SOLVENT CDCL3
NS 4096
DS 4
SFO1 51020.406 Hz
FIDRES 0.778810 Hz
AQ 0.642528 sec
RG 186.93
DM 2.800 usec
DE 18.00 usec
TE 298.0 K
D1 2.00000000 sec
D11 0.03000000 sec
TD0 1

----- CHANNEL f1 -----
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NUC1 13C
P1 12.00 usec
PLM1 130.00000000 H

----- CHANNEL f2 -----
SFO2 850.1534906 MHz
NUC2 1H
CPDPRG2 waltz16
PCPD2 80.00 usec
PLM2 13.80000019 H
PLM12 0.13800000 H
PLM13 0.98832000 H

F2 - Processing parameters
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WDW EM
SSB 0
LB 1.50 Hz
GB 0
PC 2.00

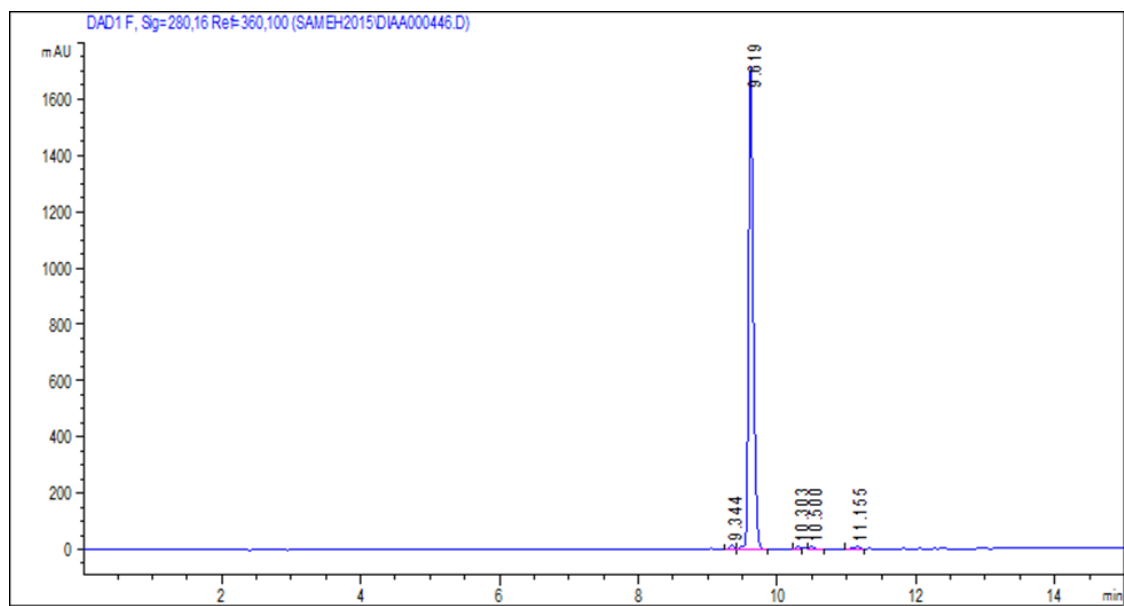
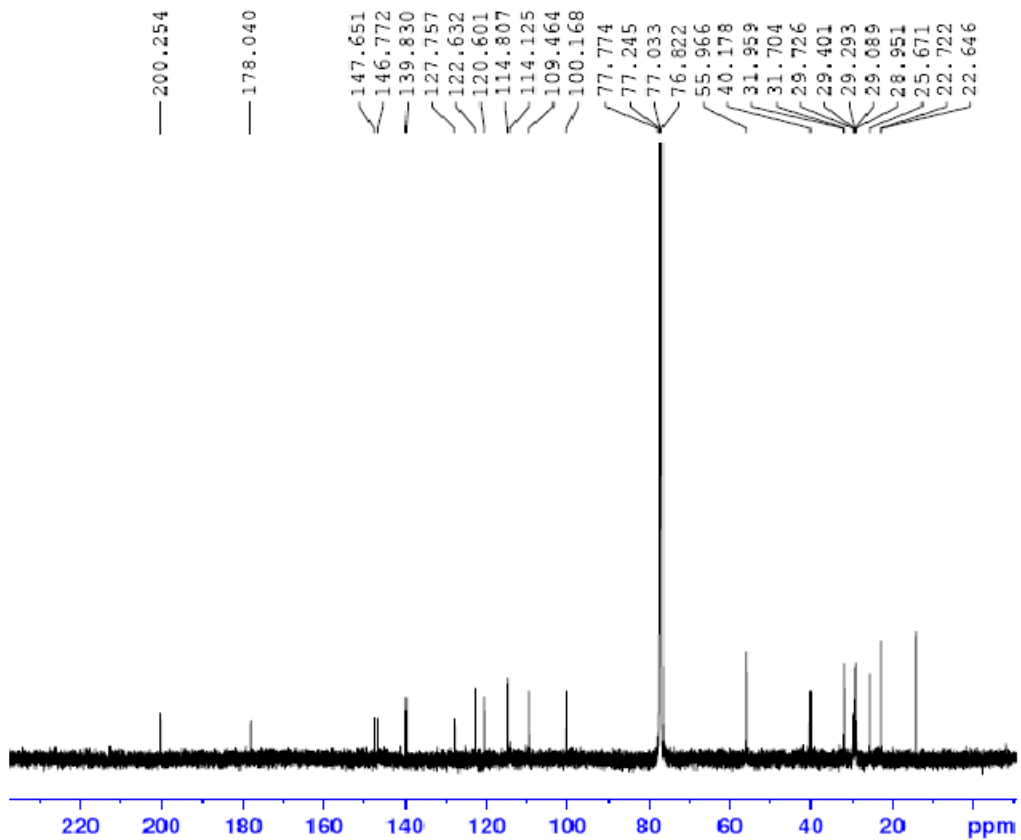
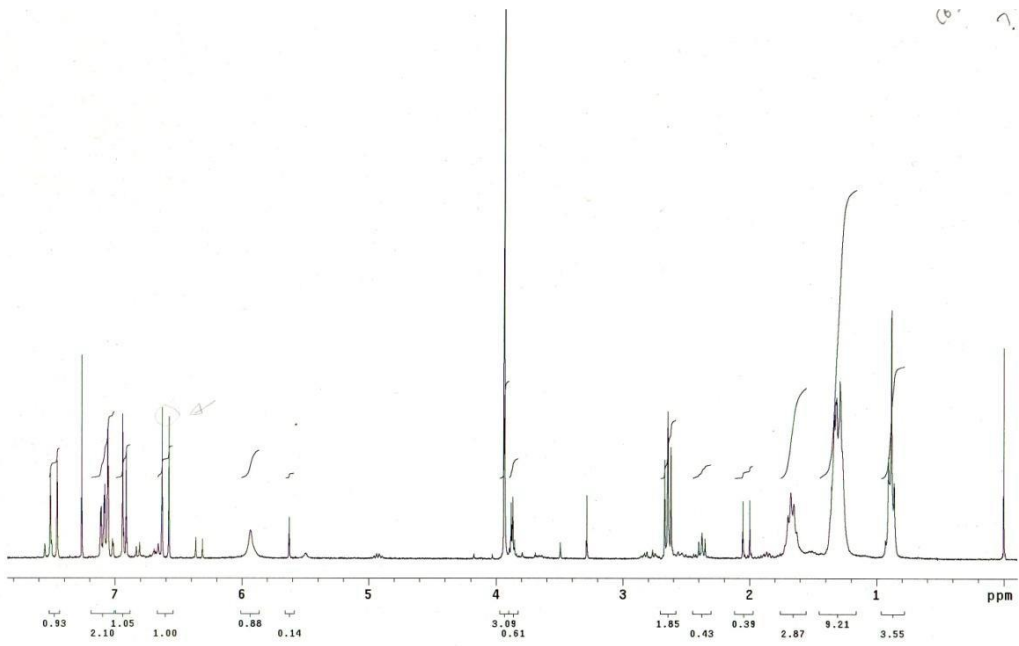


Figure S2. ^1H and ^{13}C NMR charts and HPLC chromatogram of **6-paradol** (Samples were recorded on DRX-850 spectrometer (Bruker BioSpin, Billerica, MA, USA)).



Scan: 84
Base: m/z 177; 16.2%FS TIC: 6148640

R.T.: 2.1

#Ions: 460

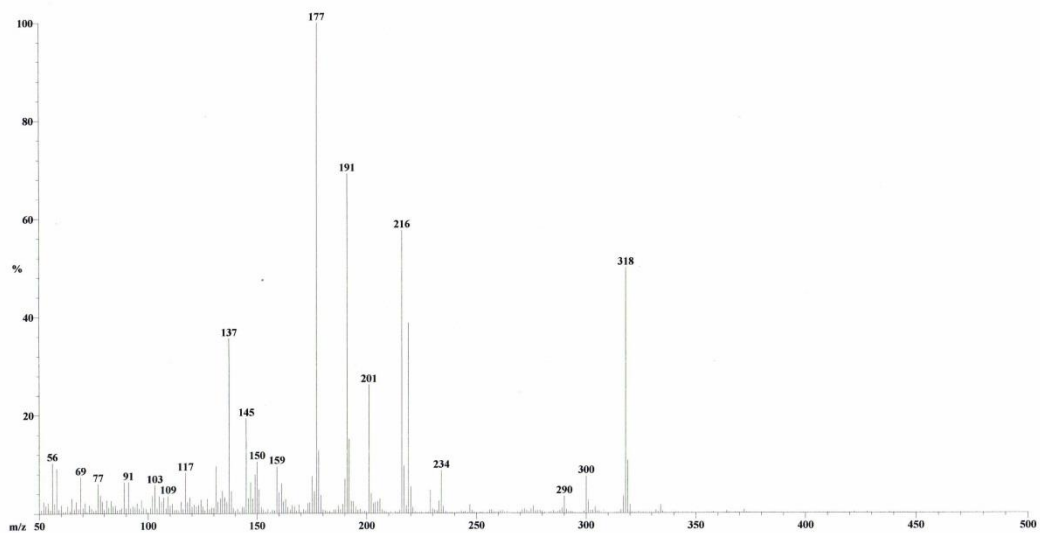
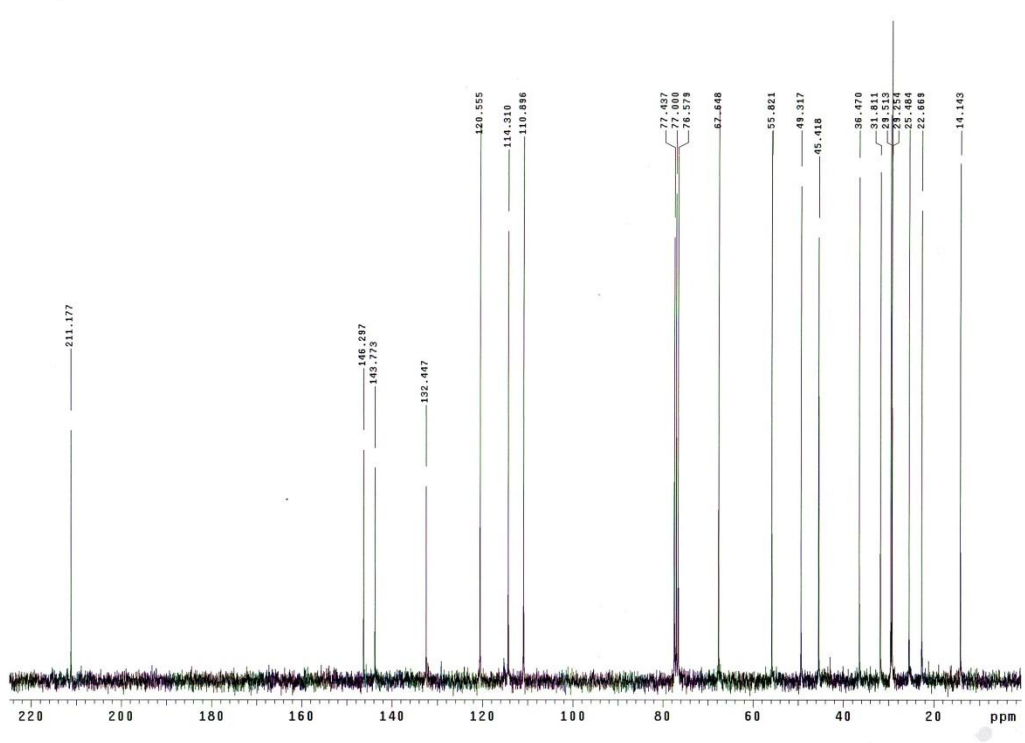
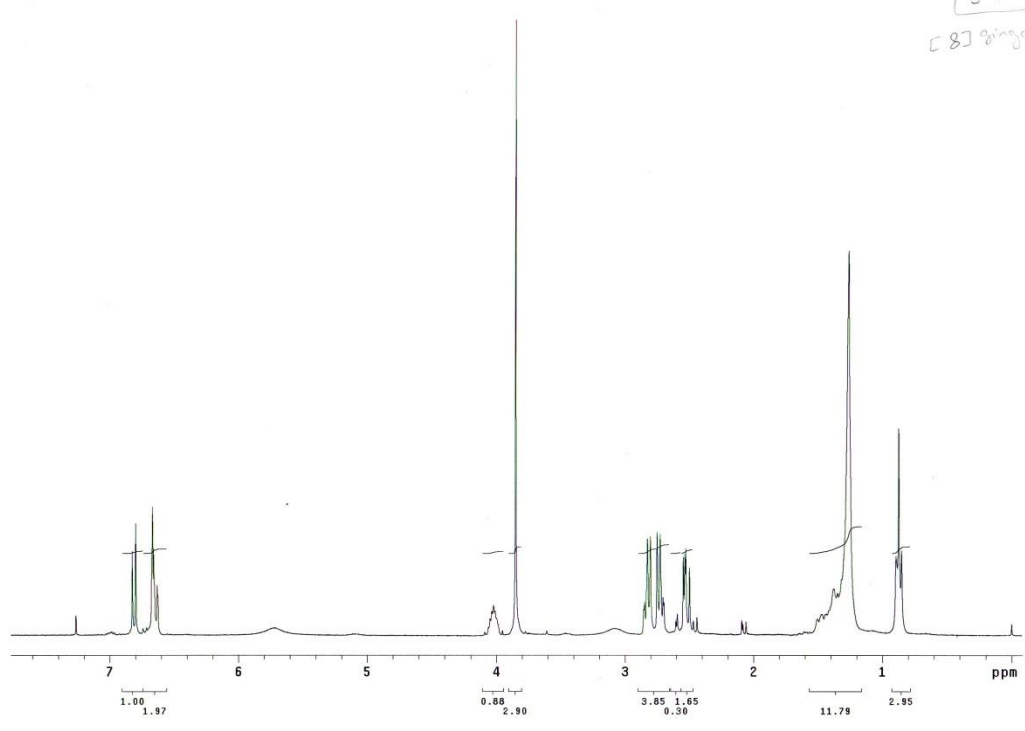


Figure S3. ^1H and ^{13}C NMR and Ms Charts of 8-dehydrogingerdione (3)

3-7-3
[8] gingerol



2007/12/19

Page 1

File: LR00
Sample: 3-7-3
Instrument: JEOL GCmate
Inlet: Direct Probe

Date Run: 2007-12-19 (Time Run: 20:44:07)

Ionization mode: EI+

8-gingerol

Scan: 65
Base: m/z 137; 12.2%FS TIC: 2905988

R.T.: 1.62

#Ions: 2

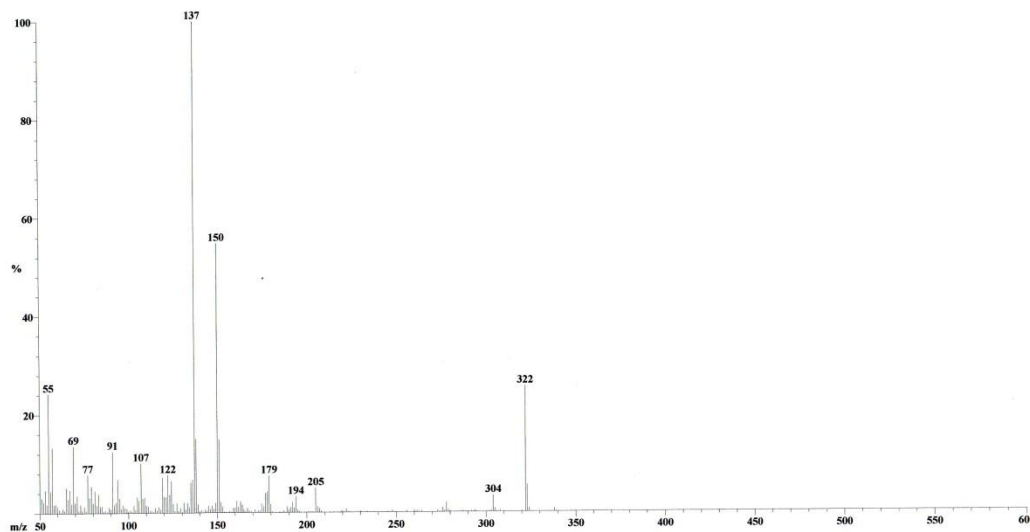


Figure S4. ^1H and ^{13}C NMR and MS charts of 8-gingerol (4)

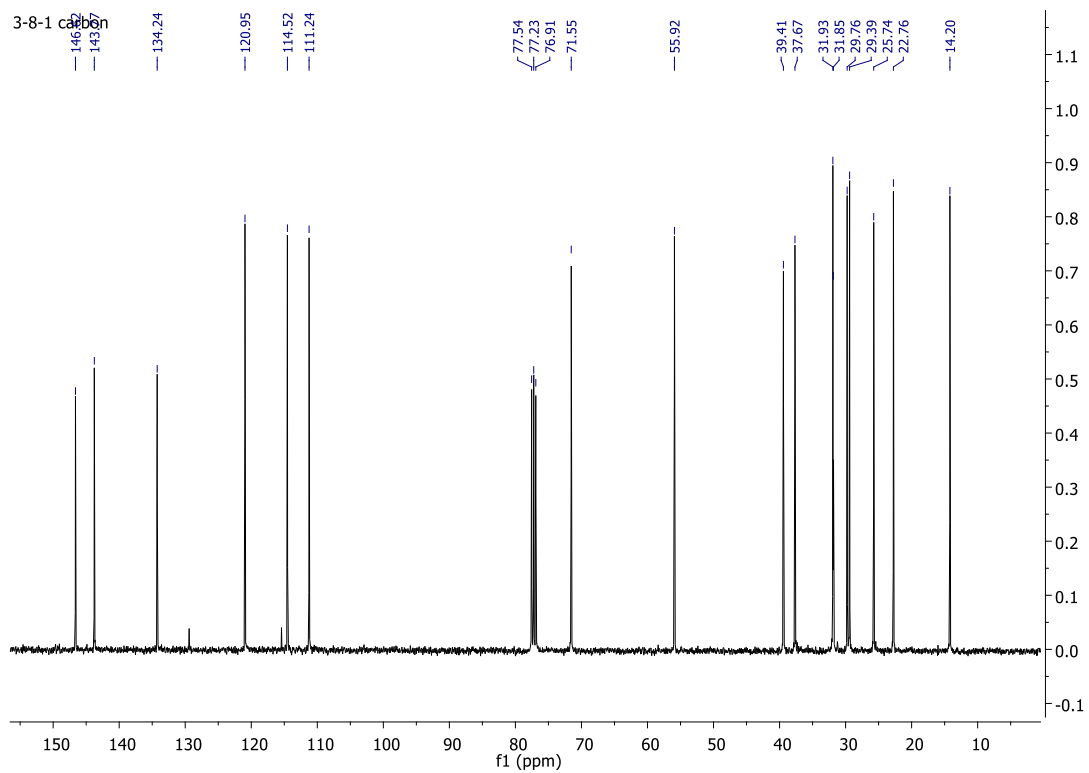
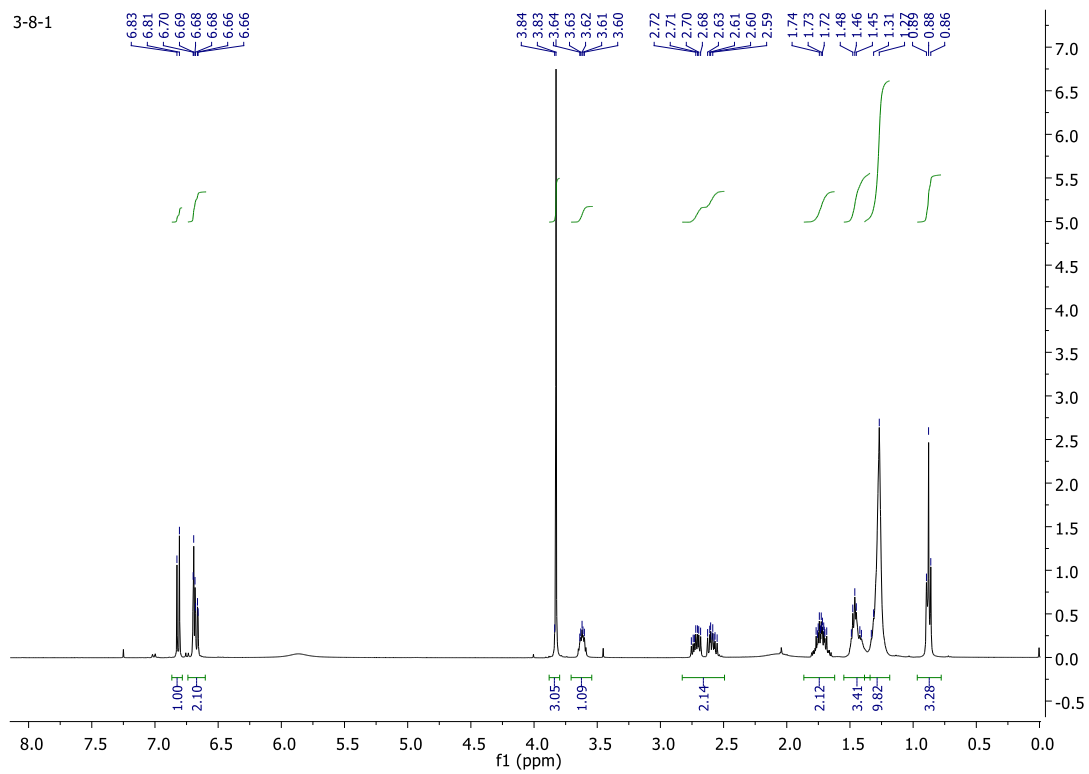
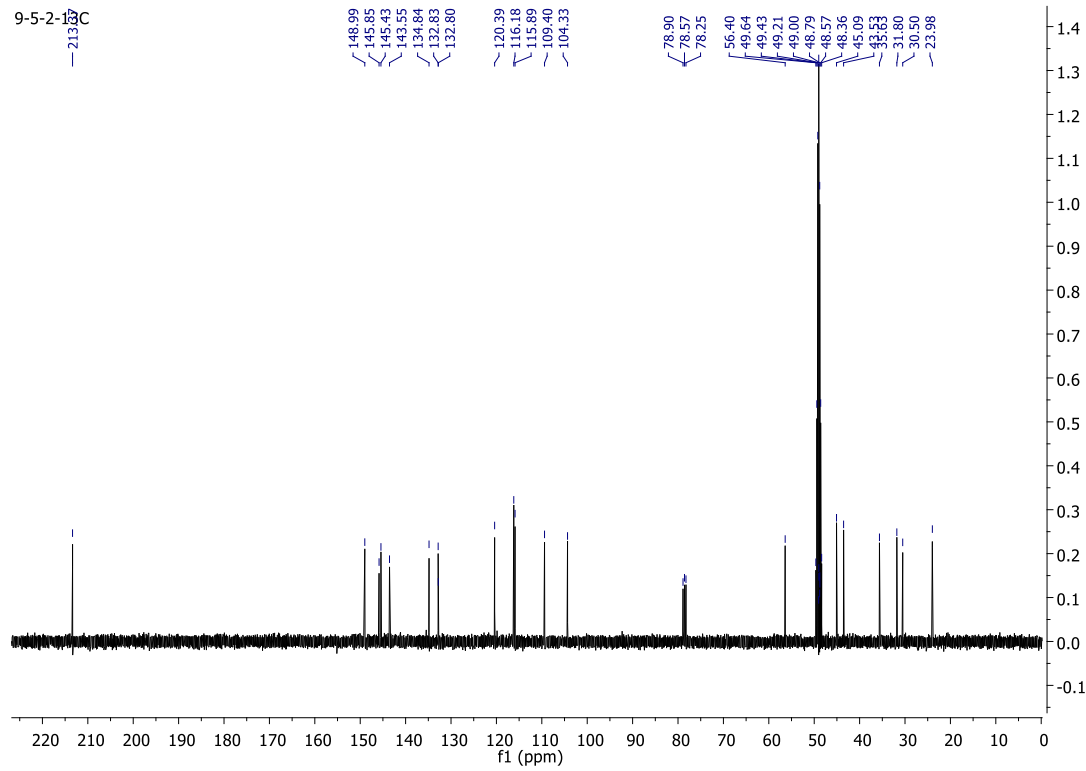
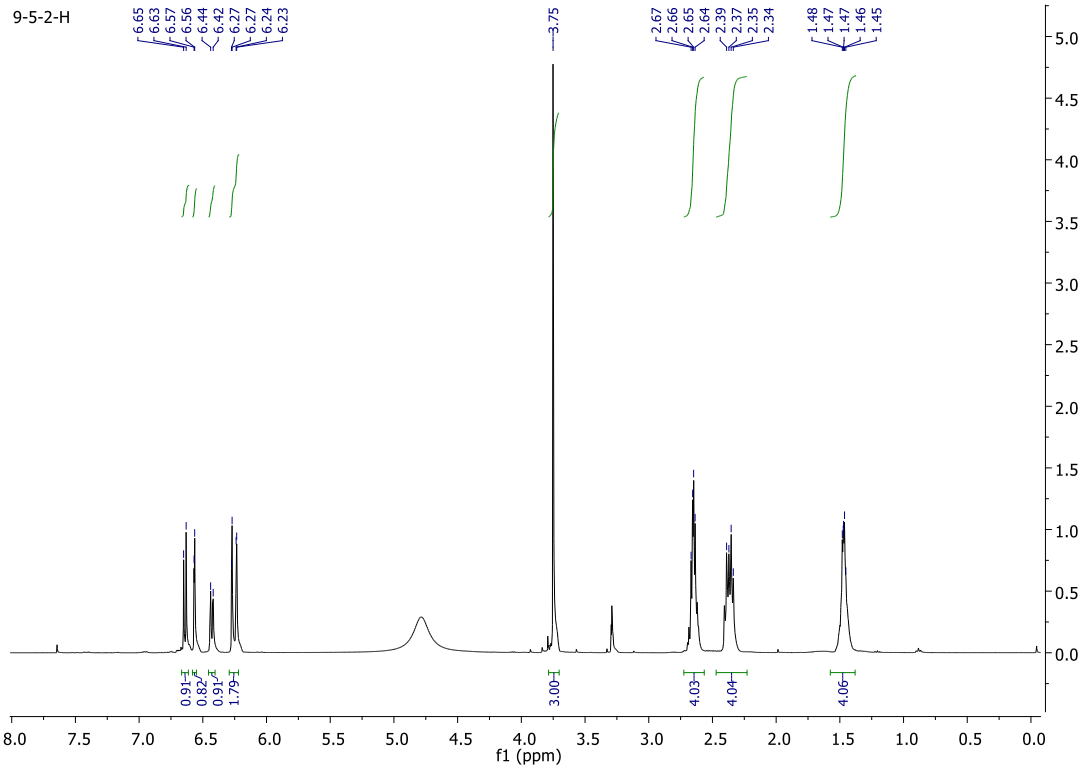


Figure S5. ^1H and ^{13}C NMR charts of dihydro-6-paradol (**5**)



Scan: 160
Base: m/z 153; 2.5%FS TIC: 1670832

R.T.: 4.02

#1or

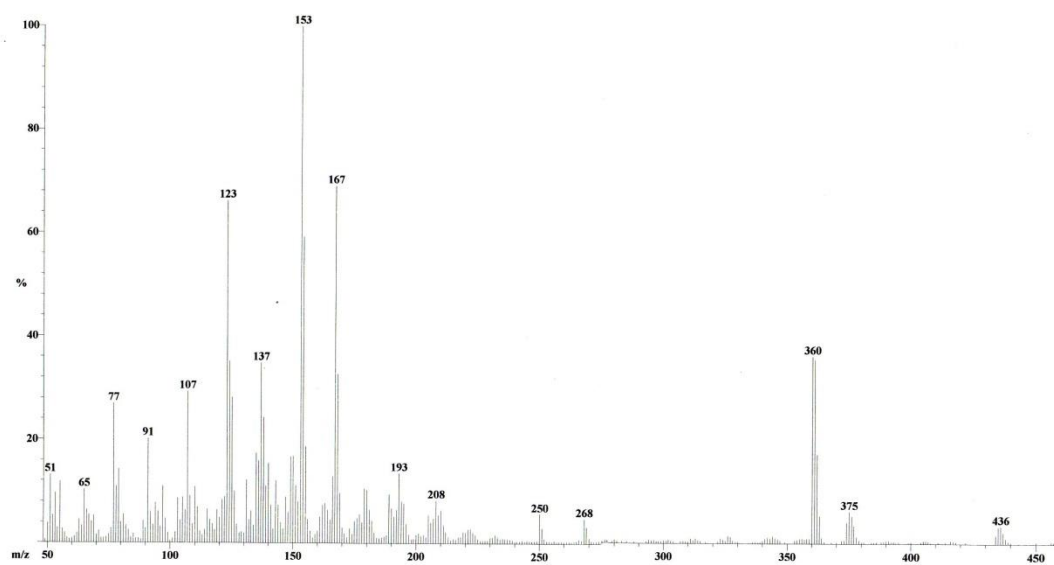


Figure S6. ^1H and ^{13}C NMR and Ms charts of dihydrogingerone-A (6)

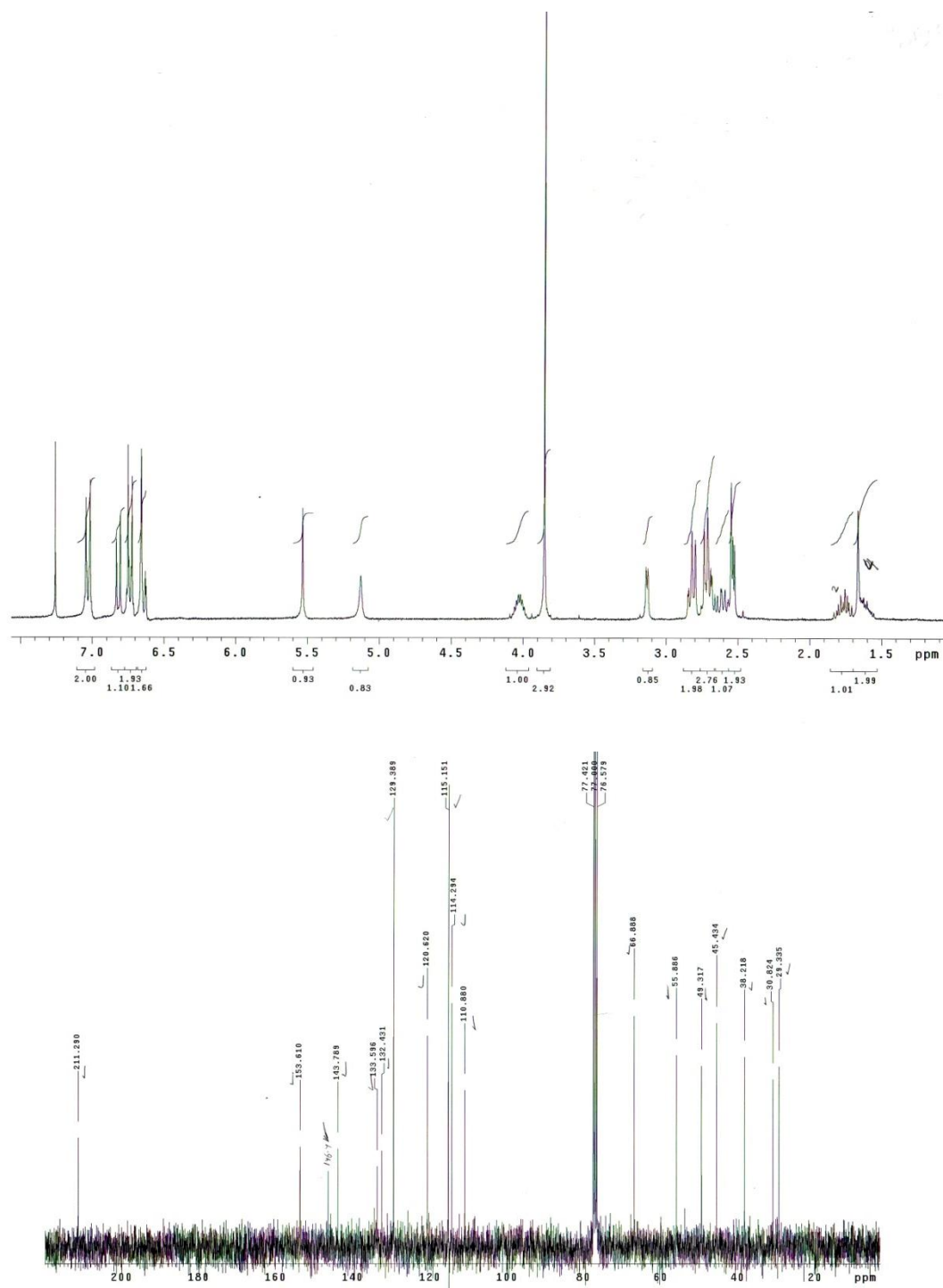


Figure S7. ^1H and ^{13}C NMR charts of dihydrogingerone-C (7)

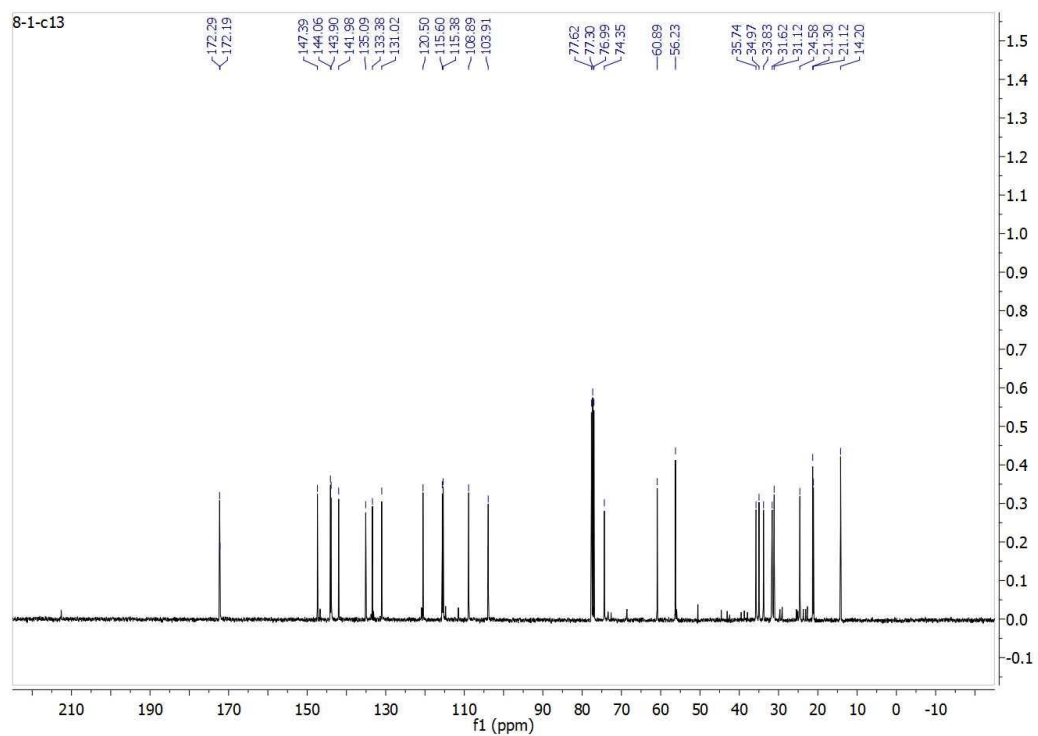
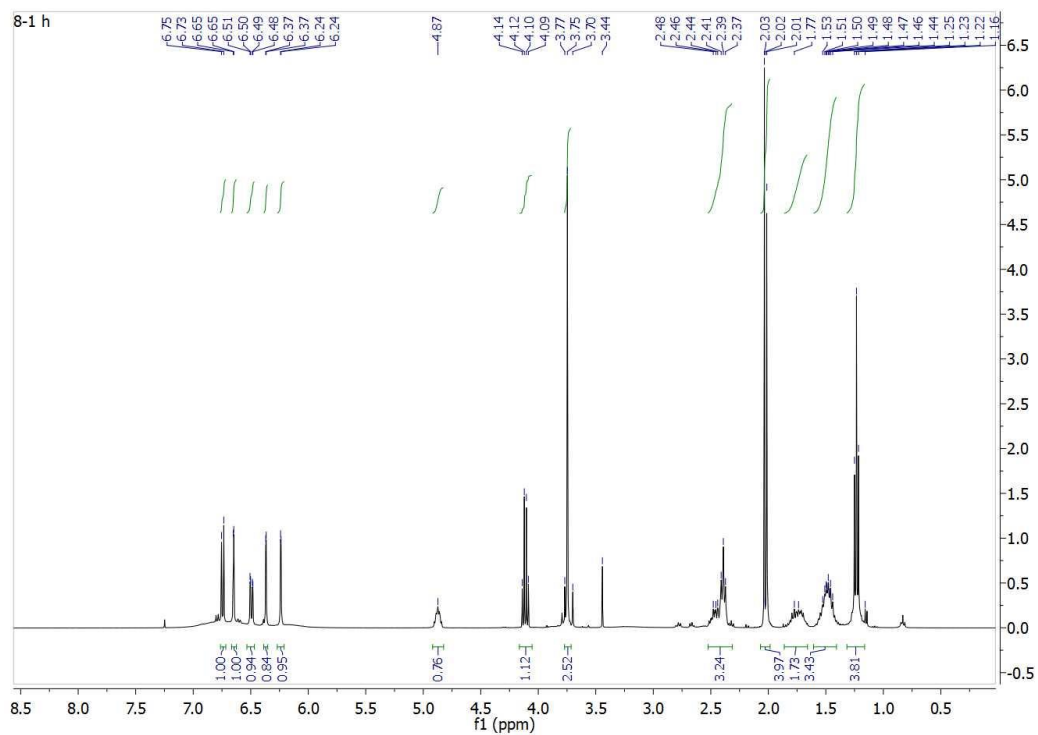


Figure S8. ^1H and ^{13}C NMR charts of compound **8**

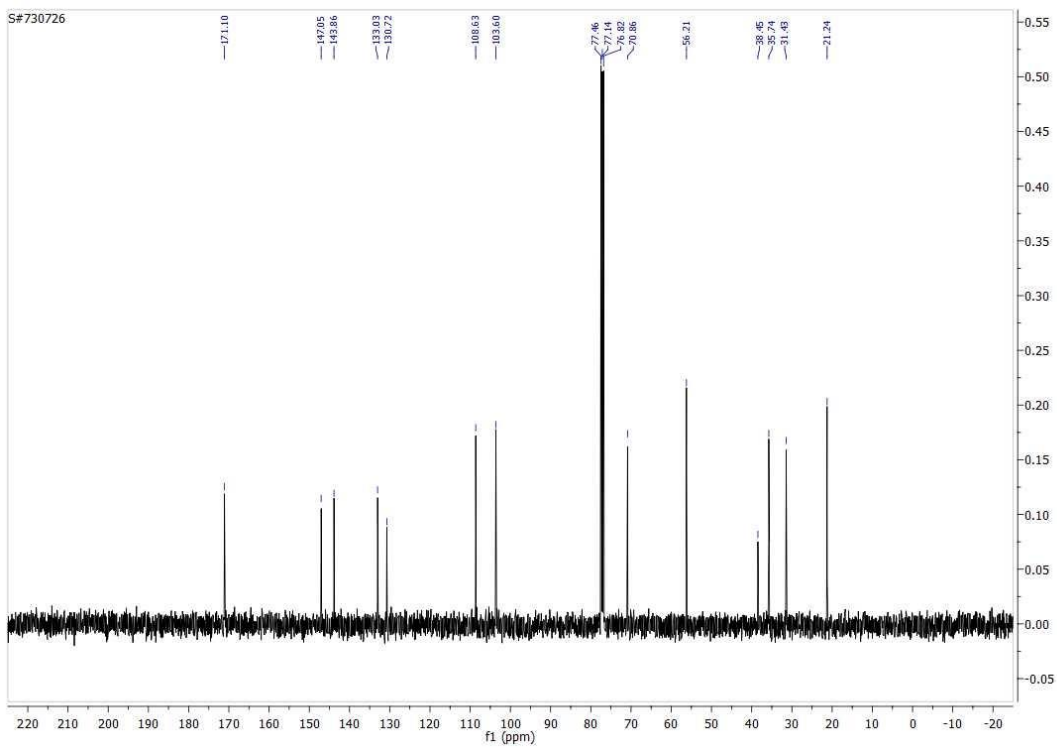
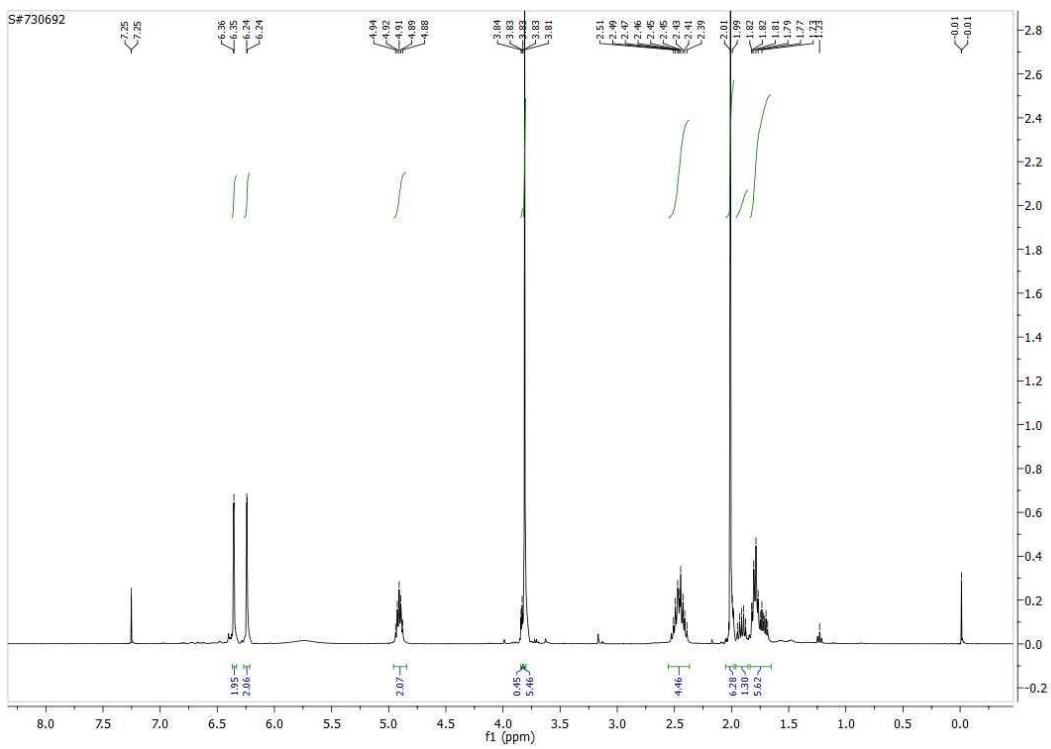


Figure S9. ^1H and ^{13}C NMR charts of **compound 9**

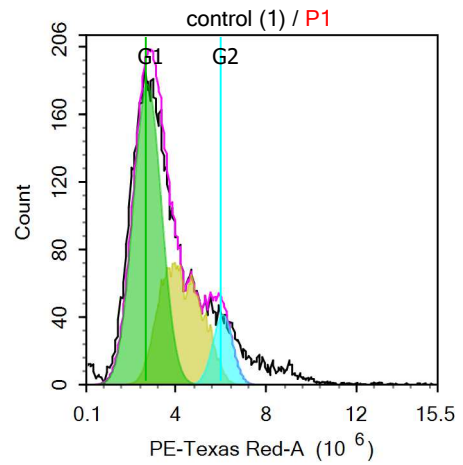
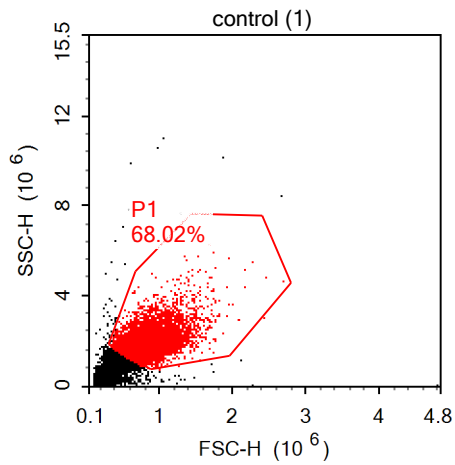
Report of Specimen 1

Specimen Name: Specimen1

Run Time: 11/8/2016 2:22 PM

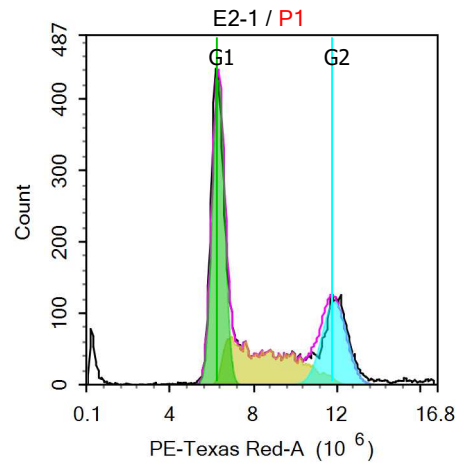
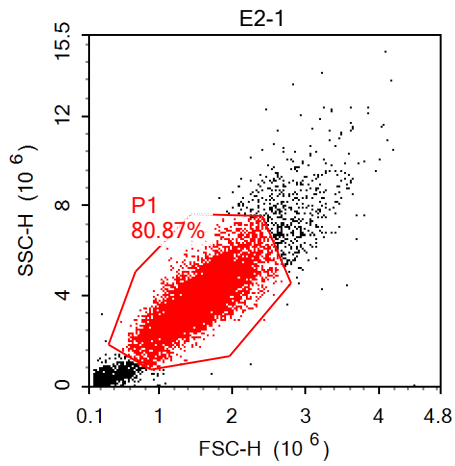
Cytometer: NovoCyte 451141011592

Software: NovoExpress 1.1.0



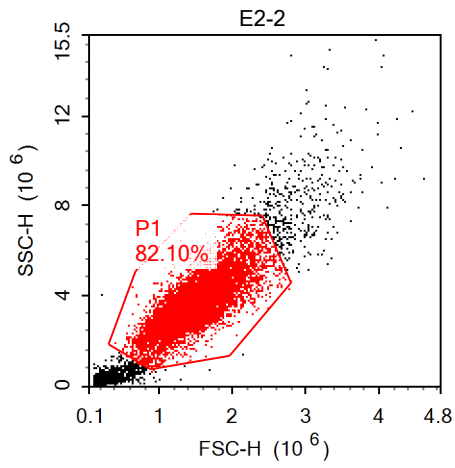
Gate	Count	% All	Mean X	Mean Y
All	12,000	100.00 %	661,934	1,522,012
P1	8,162	68.02 %	814,598	1,906,709

RMS	Freq G1	Freq S	Freq G2	G2/G1	CV G1
11.50	52.54	31.35	8.67	2.21	22.48 %

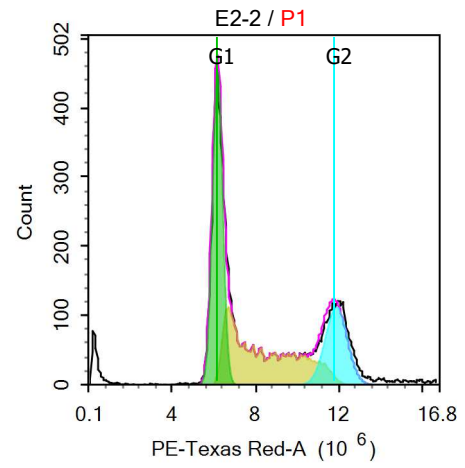


Gate	Count	% All	Mean X	Mean Y
All	12,000	100.00 %	1,356,756	3,379,520
P1	9,704	80.87 %	1,478,921	3,699,377

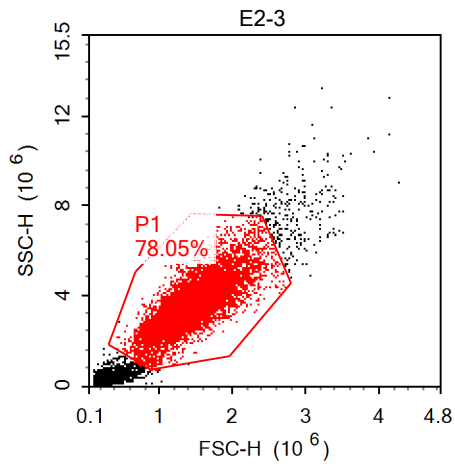
RMS	Freq G1	Freq S	Freq G2	G2/G1	CV G1
15.72	41.83	29.44	23.25	1.88	4.56 %



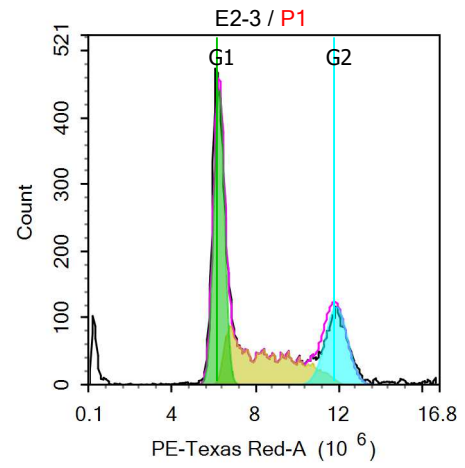
Gate	Count	% All	Mean X	Mean Y
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P1	9,852	82.10 %	1,484,045	3,647,778



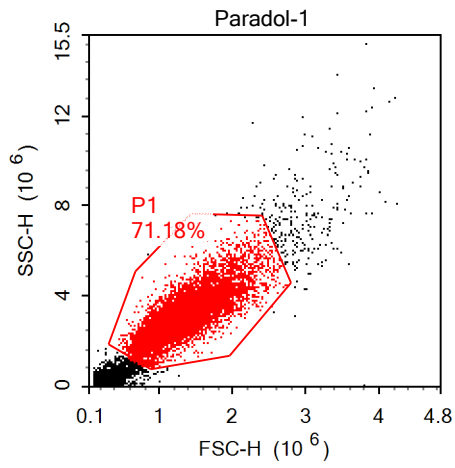
RMS	Freq G1	Freq S	Freq G2	G2/G1	CV G1
11.04	36.81	36.63	19.89	1.92	3.92 %



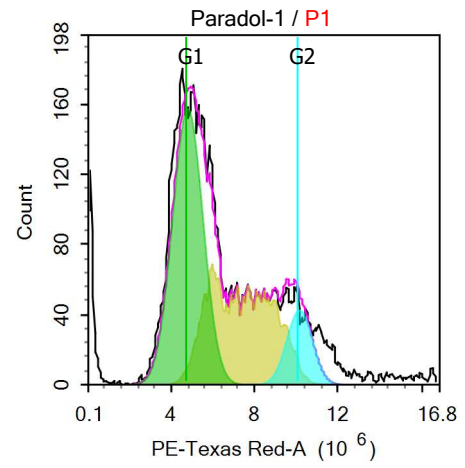
Gate	Count	% All	Mean X	Mean Y
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P1	9,366	78.05 %	1,441,499	3,563,016



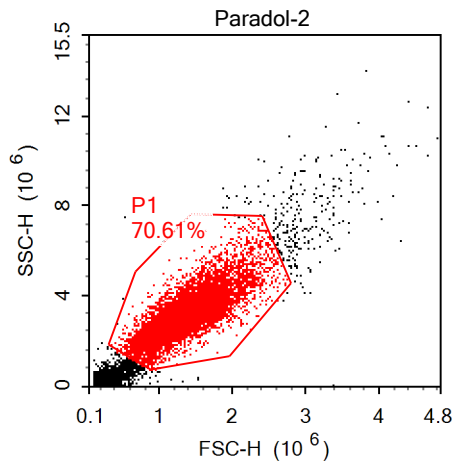
RMS	Freq G1	Freq S	Freq G2	G2/G1	CV G1
16.50	39.07	32.04	21.73	1.90	4.02 %



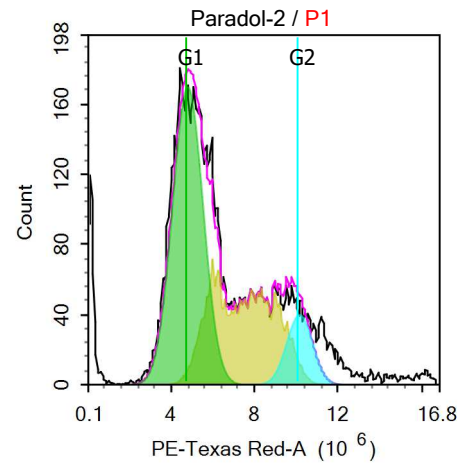
Gate	Count	% All	Mean X	Mean Y
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P1	8,542	71.18 %	1,363,014	3,175,923



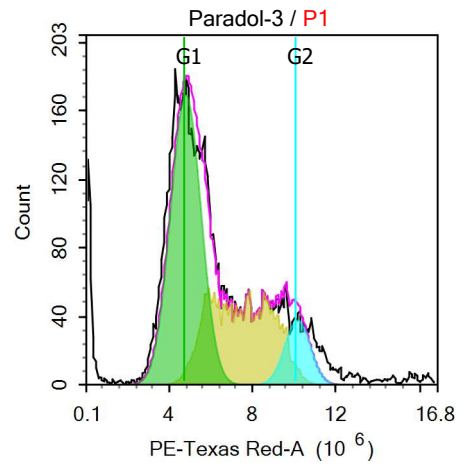
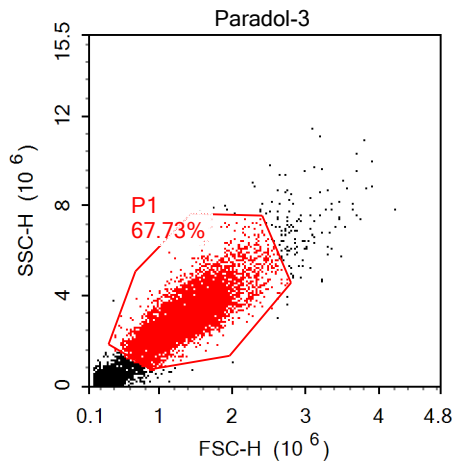
RMS	Freq G1	Freq S	Freq G2	G2/G1	CV G1
10.77	41.79	34.58	9.56	2.11	15.06 %



Gate	Count	% All	Mean X	Mean Y
All	12,000	100.00 %	1,094,594	2,489,568
P1	8,473	70.61 %	1,359,290	3,180,320



RMS	Freq G1	Freq S	Freq G2	G2/G1	CV G1
10.63	44.84	33.32	8.93	2.11	15.06 %



Gate	Count	% All	Mean X	Mean Y
All	12,000	100.00 %	1,007,418	2,248,819
P1	8,127	67.73 %	1,324,895	3,076,992

RMS	Freq G1	Freq S	Freq G2	G2/G1	CV G1
11.48	46.88	32.19	8.85	2.11	15.06 %