

Supporting information

Synthesis and Biological Evaluation of Heterocyclic Ring-substituted Betulinic Acid Derivatives as Novel Inhibitors of Osteoclast Differentiation and Bone Resorption

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1. HPLC Analysis of BA analogues

Purity of all final analogues for biological testing was confirmed to be > 95% (Table 2) as determined by HPLC analysis. HPLC analysis was conducted according to the following method with the retention time expressed in min at UV detection of 210, 230 and 254 nm. For HPLC method, an Agilent 1200 series HPLC instrument was used, with chromatography performed on a ZORBAX 150x4.6 mm, 5 μ m C18 column with mobile phase gradient of 0-10% H₂O in MeOH (Table 1), with a follow rate of 1.0 mL/min.

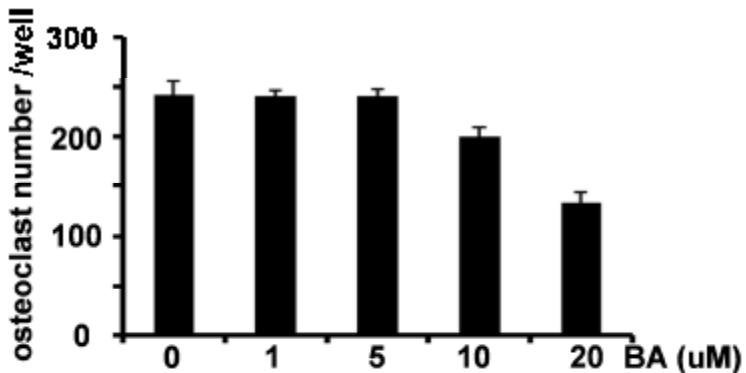
Table 1. HPLC conditions

Time (min)	H ₂ O (V%)	MeOH (V%)
0	10	90
5	0	100
20	0	100
21	10	90
30	10	90

Table 2. HPLC purity data for BA derivatives

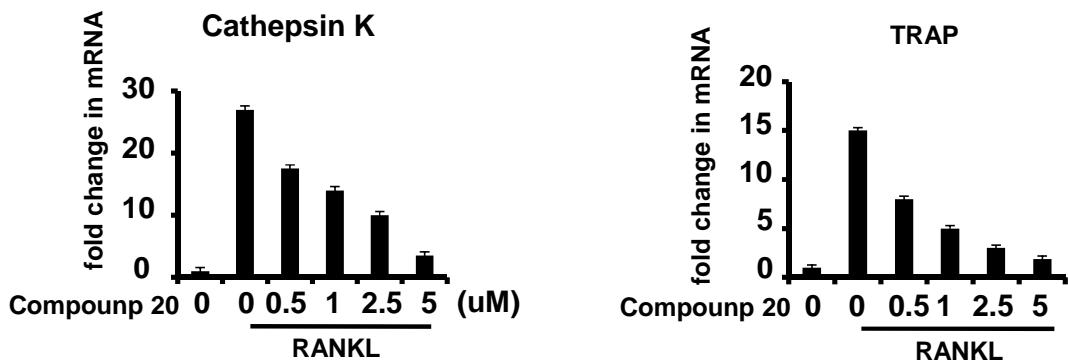
Compound	UV(nm)	RT(min)	Purity(%)	Compound	UV(nm)	RT (min)	Purity(%)
2	254	6. 636	96. 9	20	230	5. 387	98. 2
4	254	6. 318	95. 4	21	230	6. 265	97. 6
5	210	6. 164	97. 8	22	230	5. 579	98. 5
12	254	4. 902	96. 0	23	230	4. 755	98. 7
13	230	5. 738	96. 1	27	230	6. 636	95. 2
14	230	6. 562	95. 2	29	230	7. 861	96. 2
15	230	6. 003	95. 1	30	230	8. 250	97. 7
16	230	5. 166	96. 5	32	230	10. 507	97. 8
17	230	5. 814	96. 2	33	230	8. 743	95. 1
18	254	4. 708	96. 3	34	230	7. 009	96. 7
19	230	6. 497	96. 2	35	230	6. 307	98. 3

2. The effect of BA on osteoclast differentiation



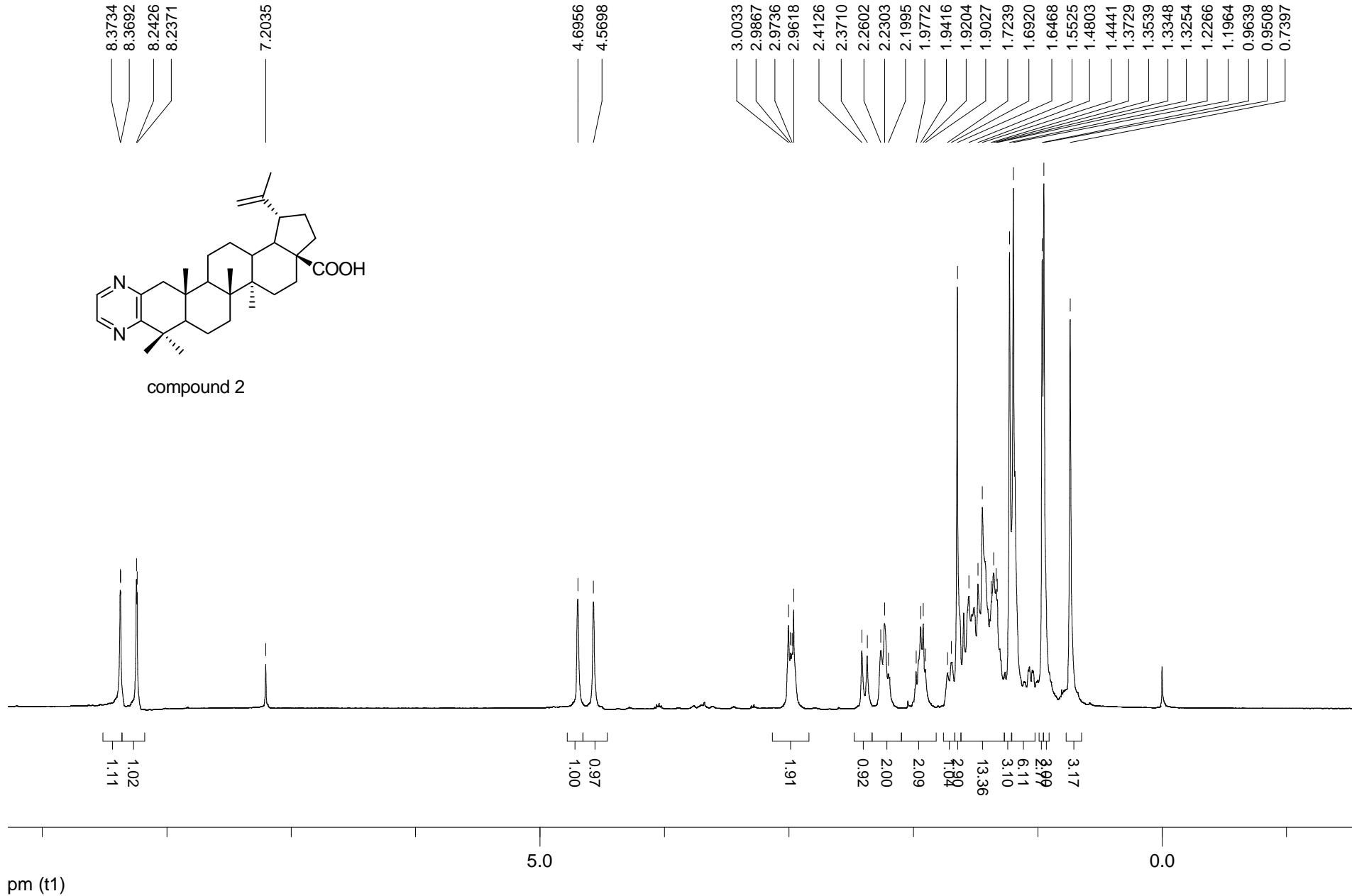
SFigure 1. The effect of BA on osteoclast differentiation. RAW 264.7 cells (3×10^3 cells/well) were treated with or without RANKL (50 ng/mL), followed by addition of the indicated concentrations of BA for 3-5 days and stained for TRAP expression. TRAP positive multinucleated (>3 nuclei) osteoclasts were counted. Column, means of three experiments carried out in triplicates. SD, bar.

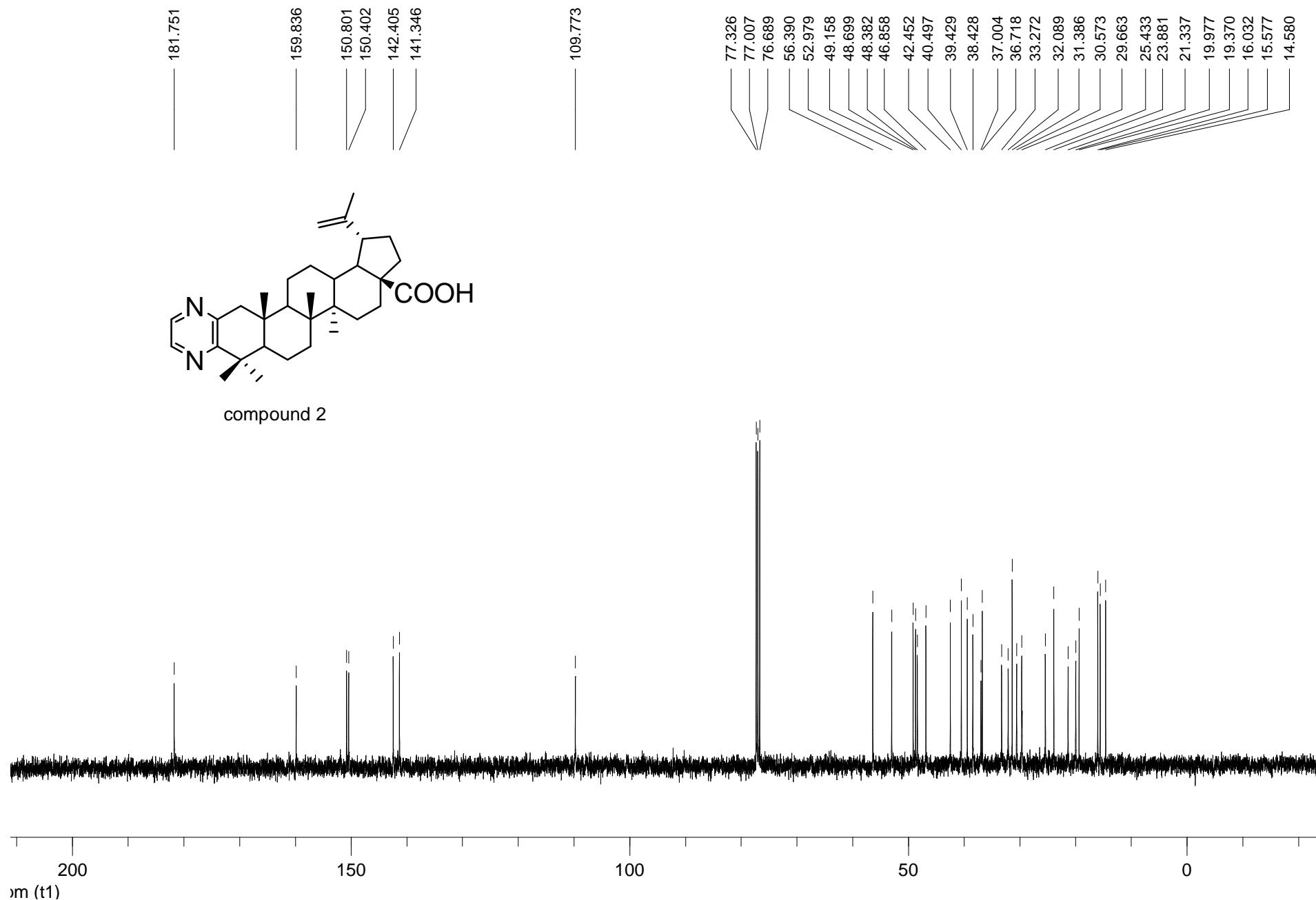
3. Compound 20 inhibits the expression of osteoclast mark genes in a dose-dependent manner

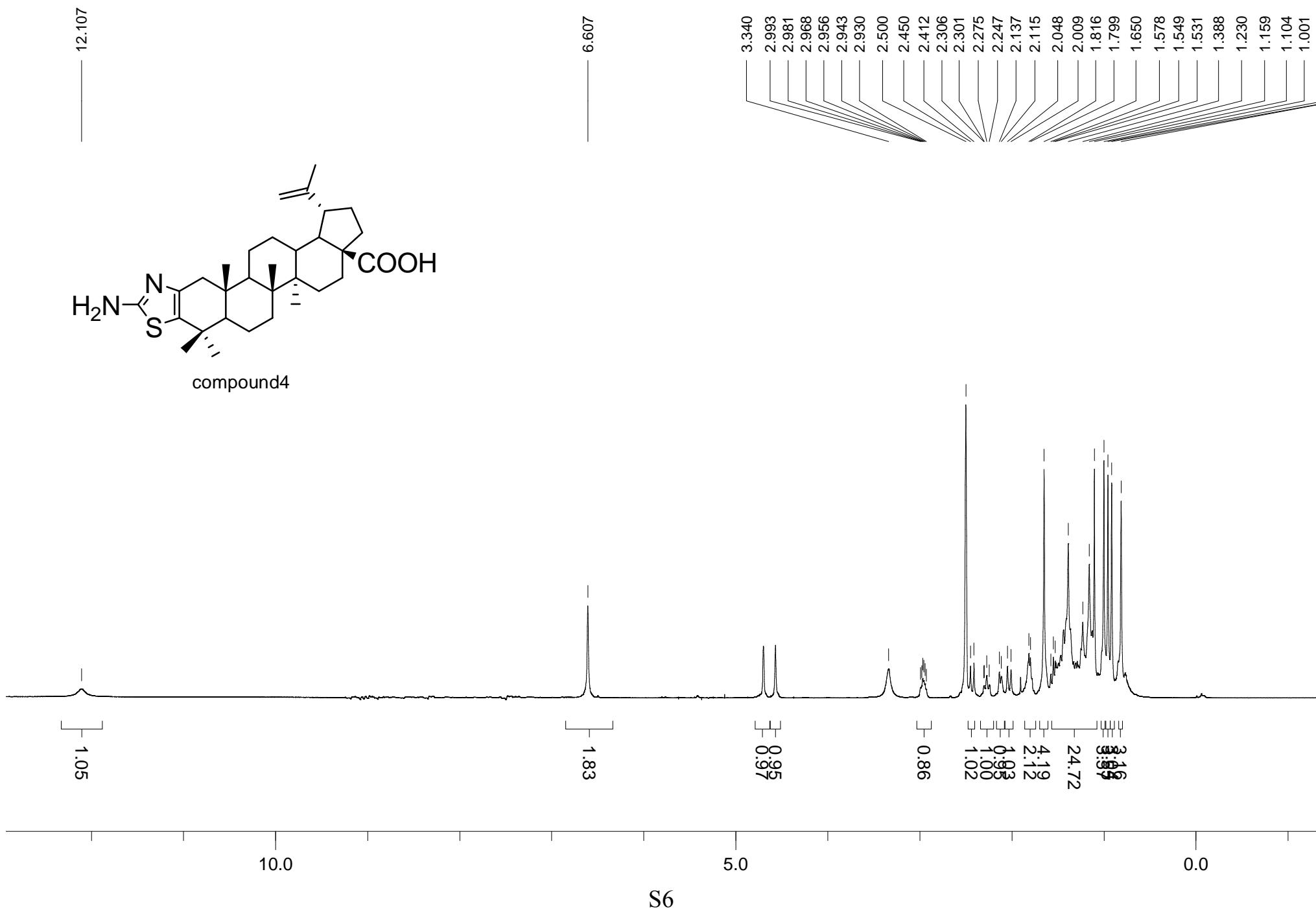


SFigure 2. Compound 20 inhibits the expression of osteoclast mark genes in a dose-dependent manner. RAW264.7 cells were pretreated with indicated concentration of 20 for 6 hours, followed by stimulation of RANKL (50ng/mL) for 3 days. Total RNA was isolated and subjected to quantitative real-time PCR for the expression levels of TRAP and cathepsin K.

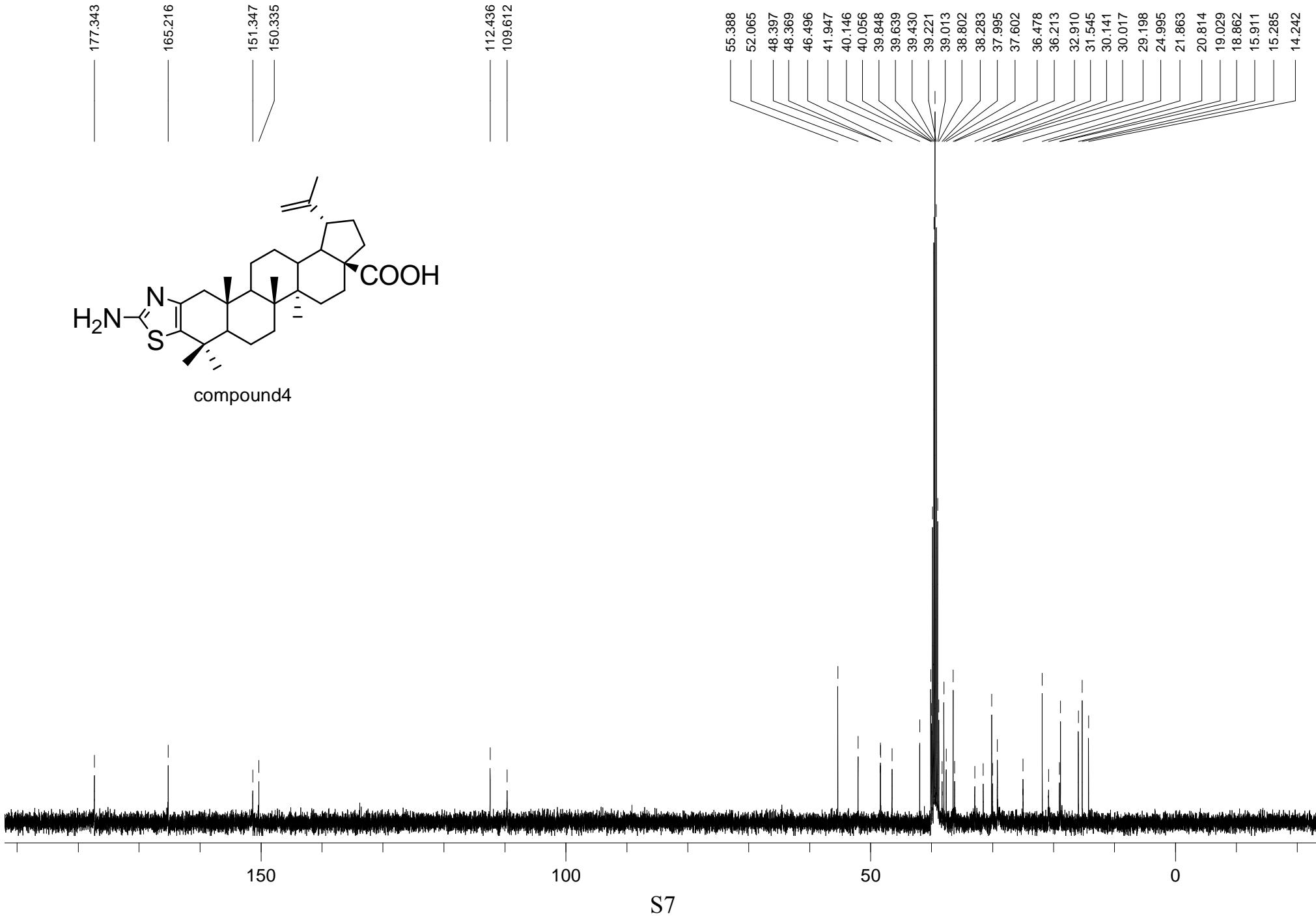
4. Copies of Selected ^1H and ^{13}C NMR Spectra

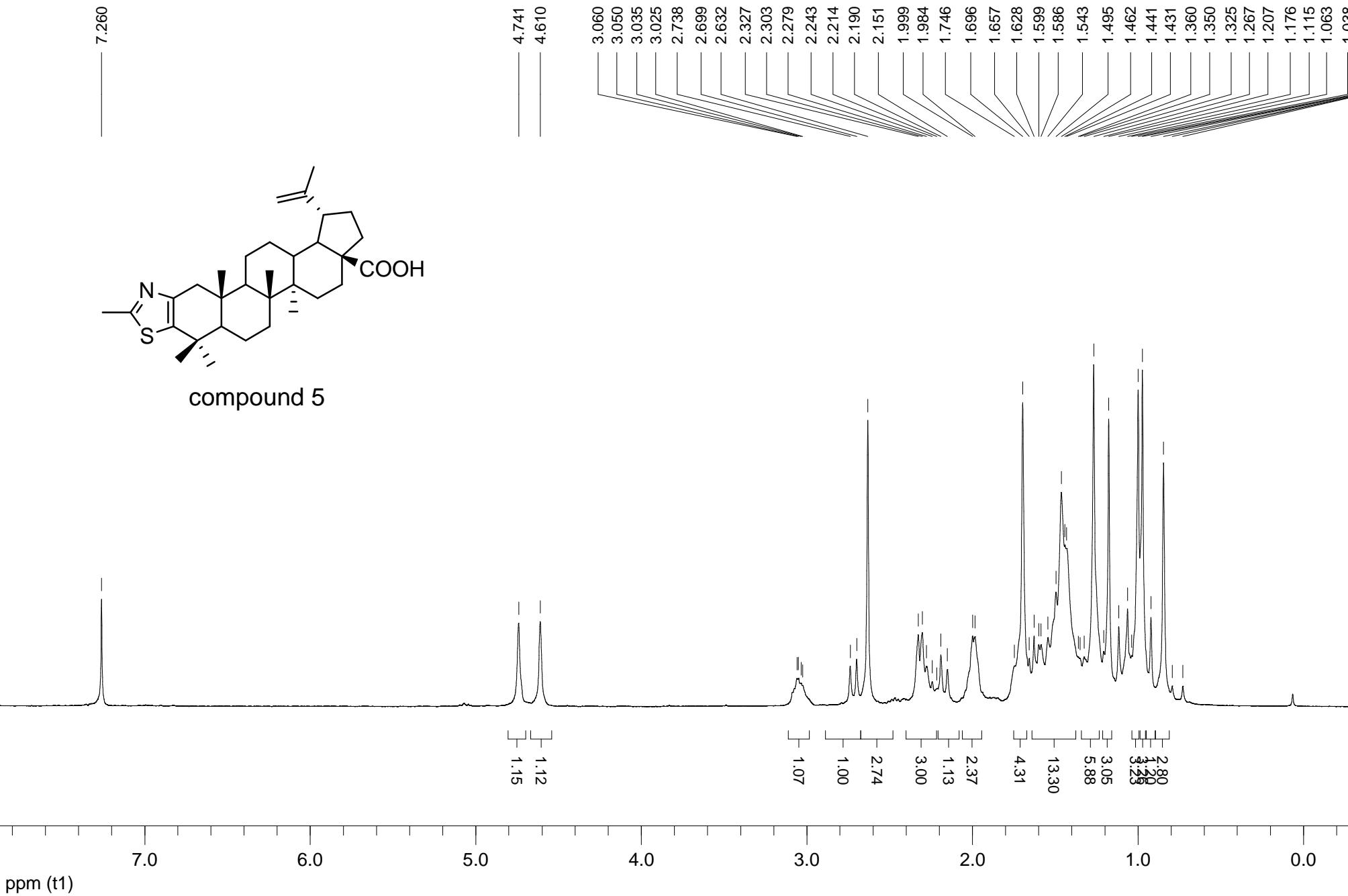


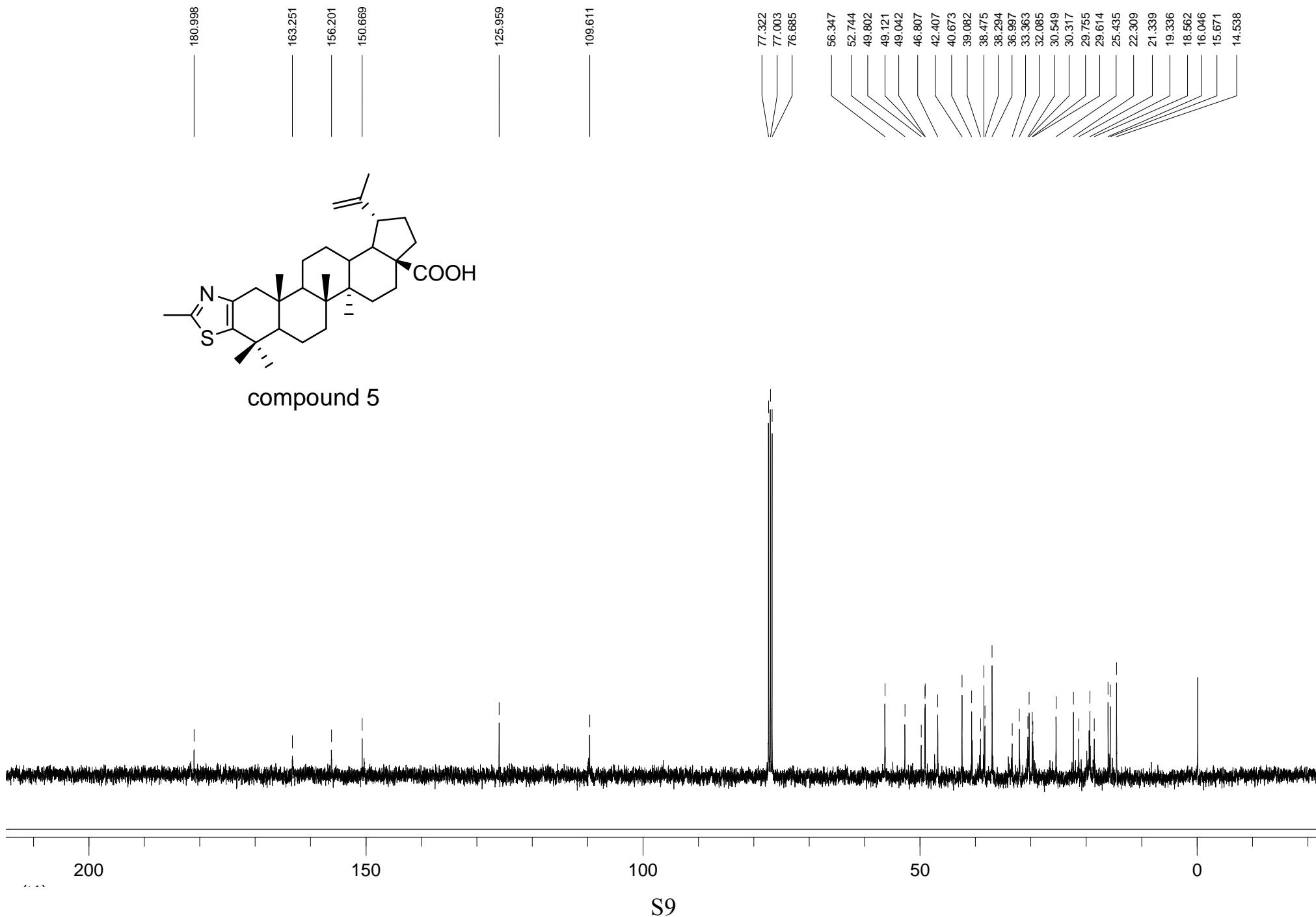


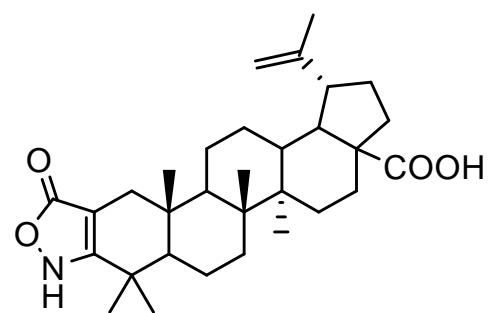


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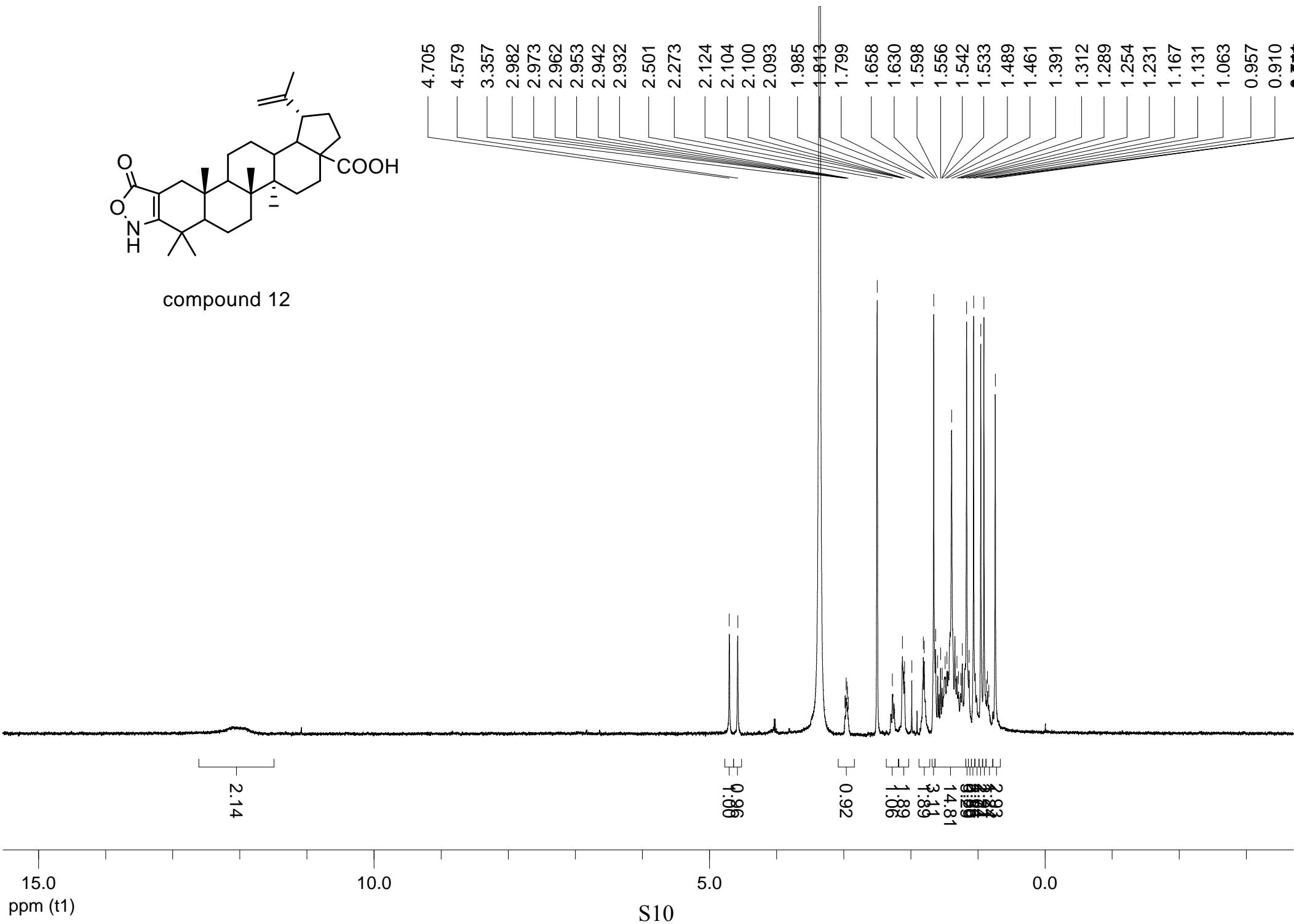


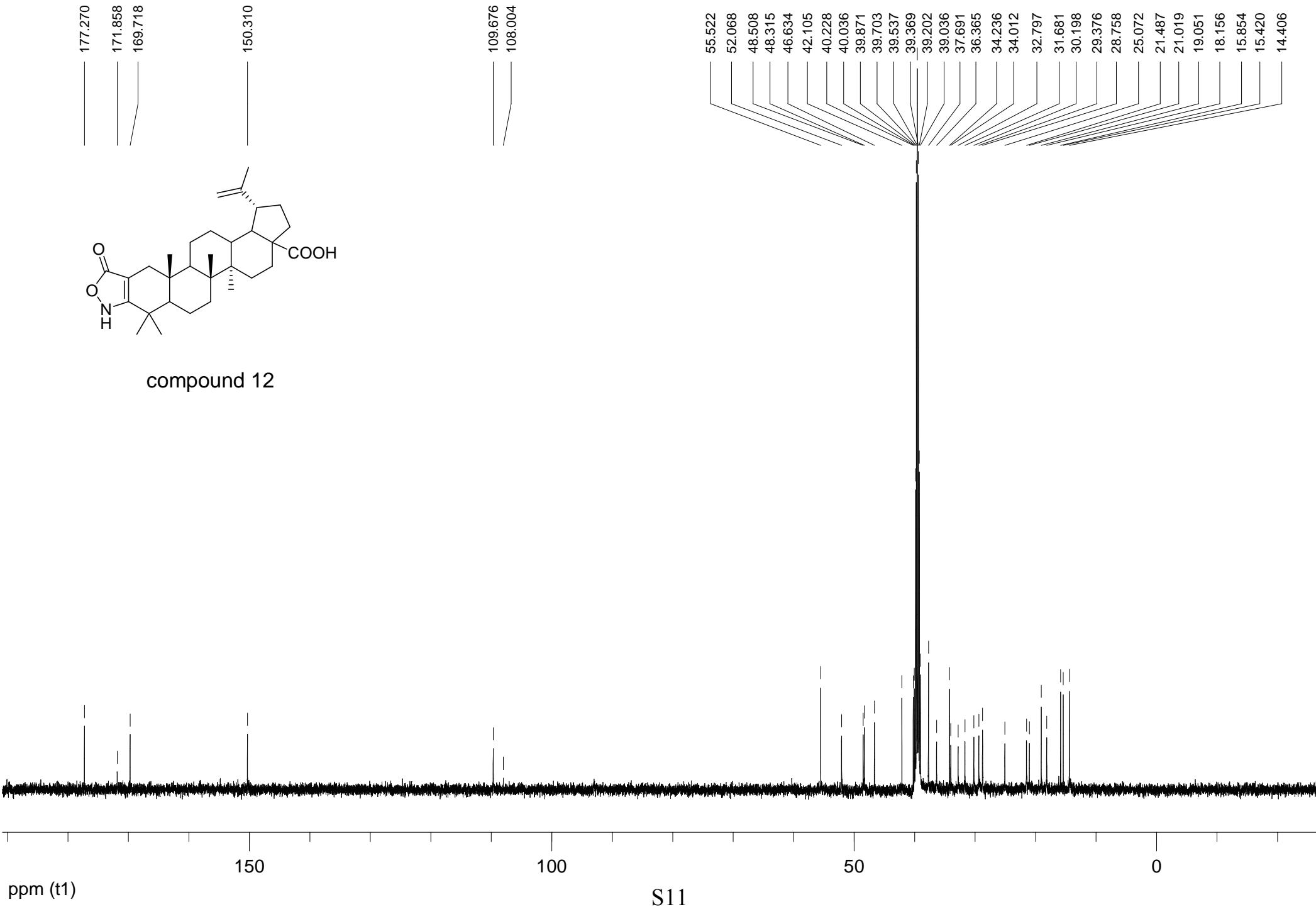


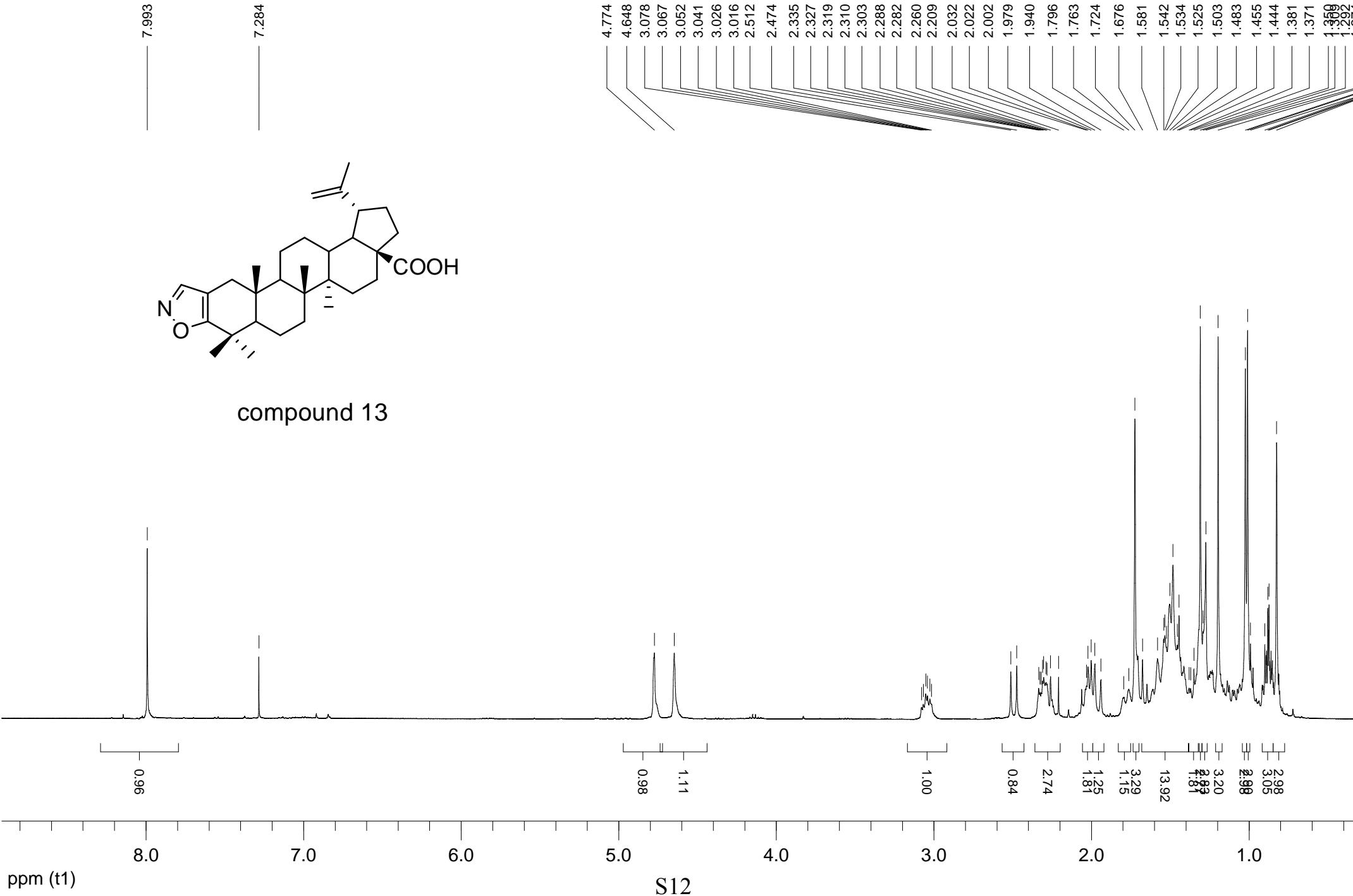


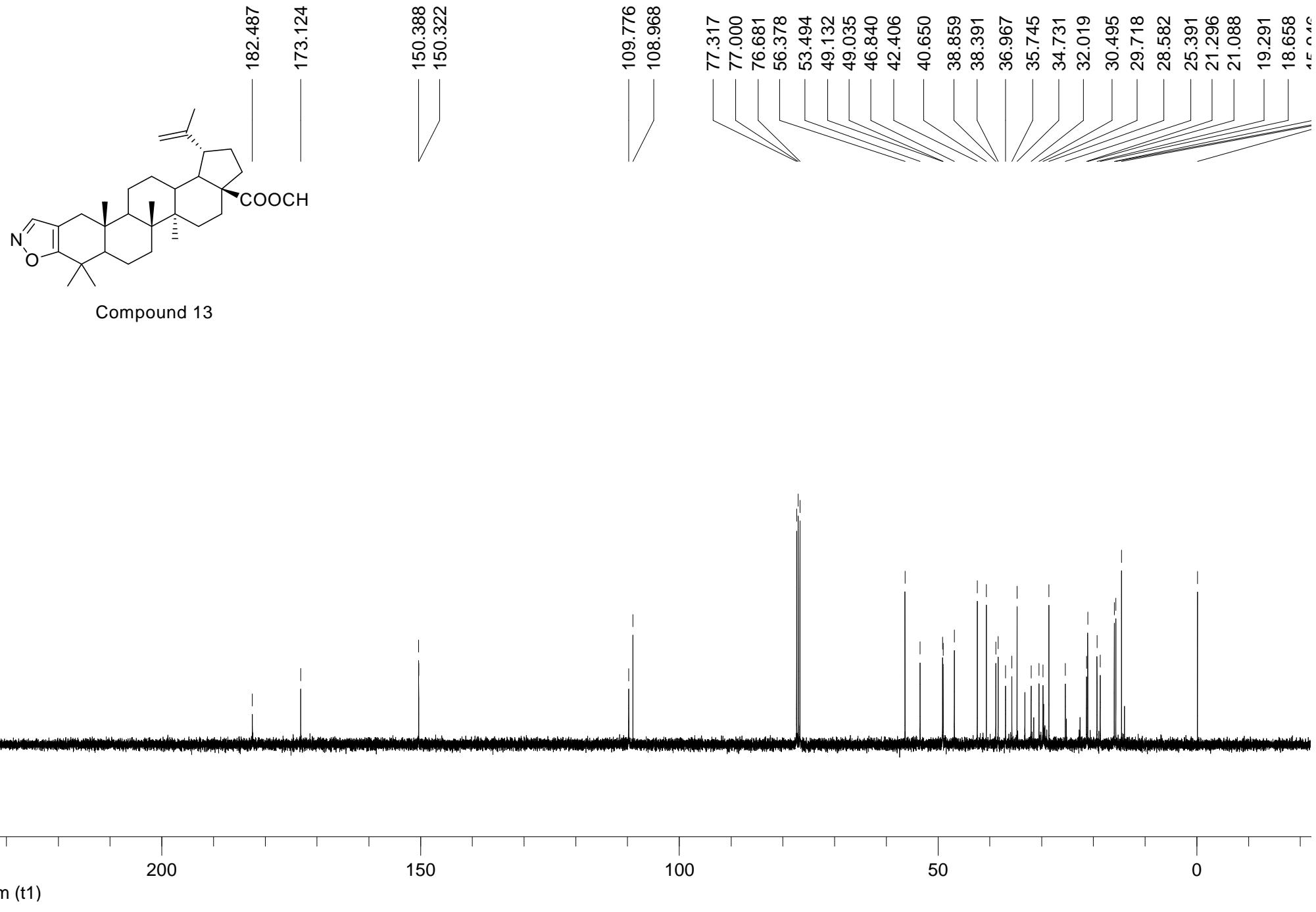


compound 12





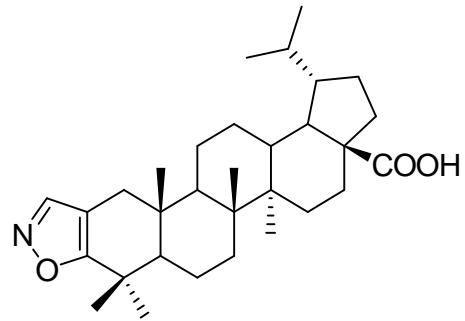




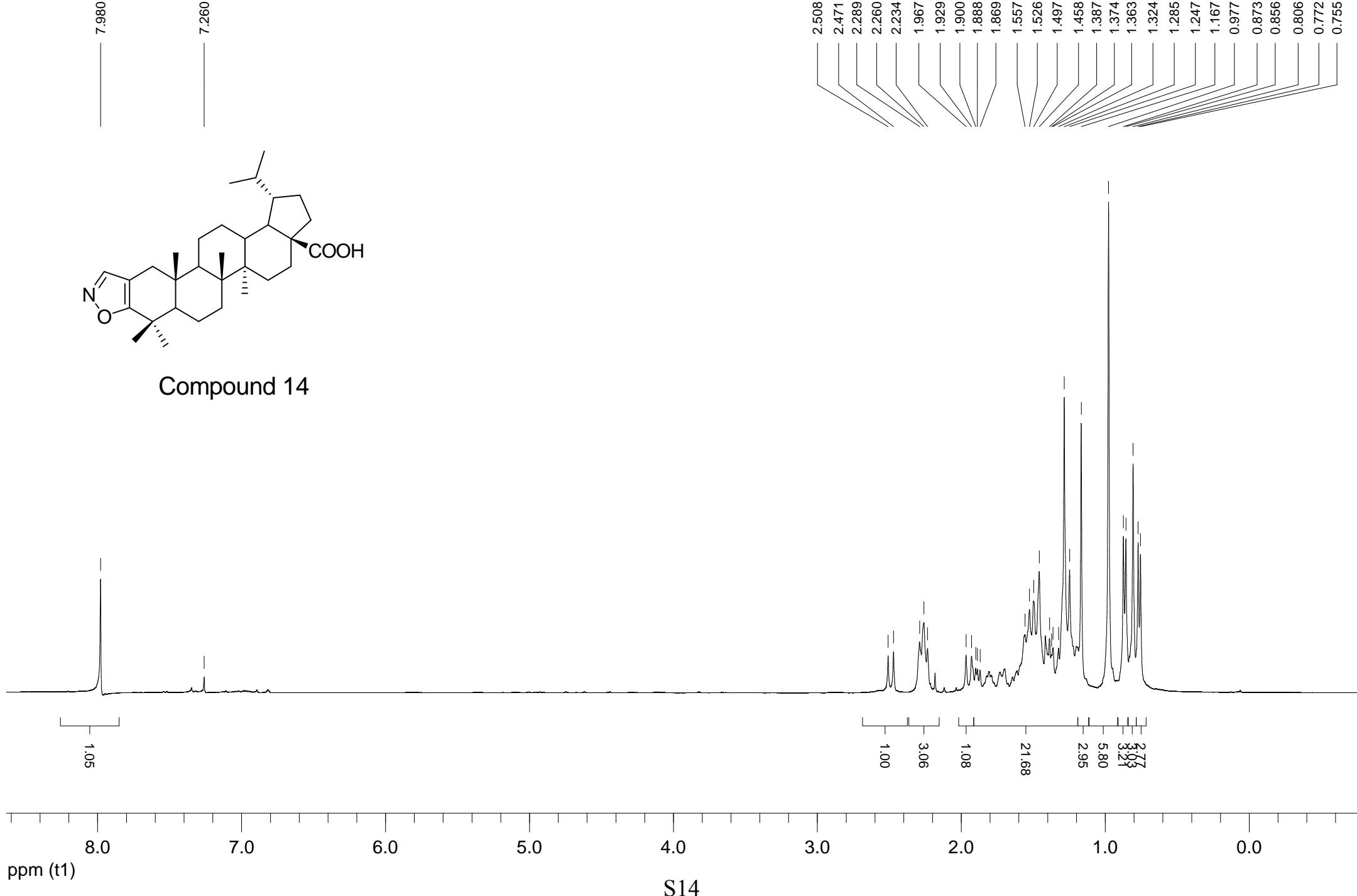
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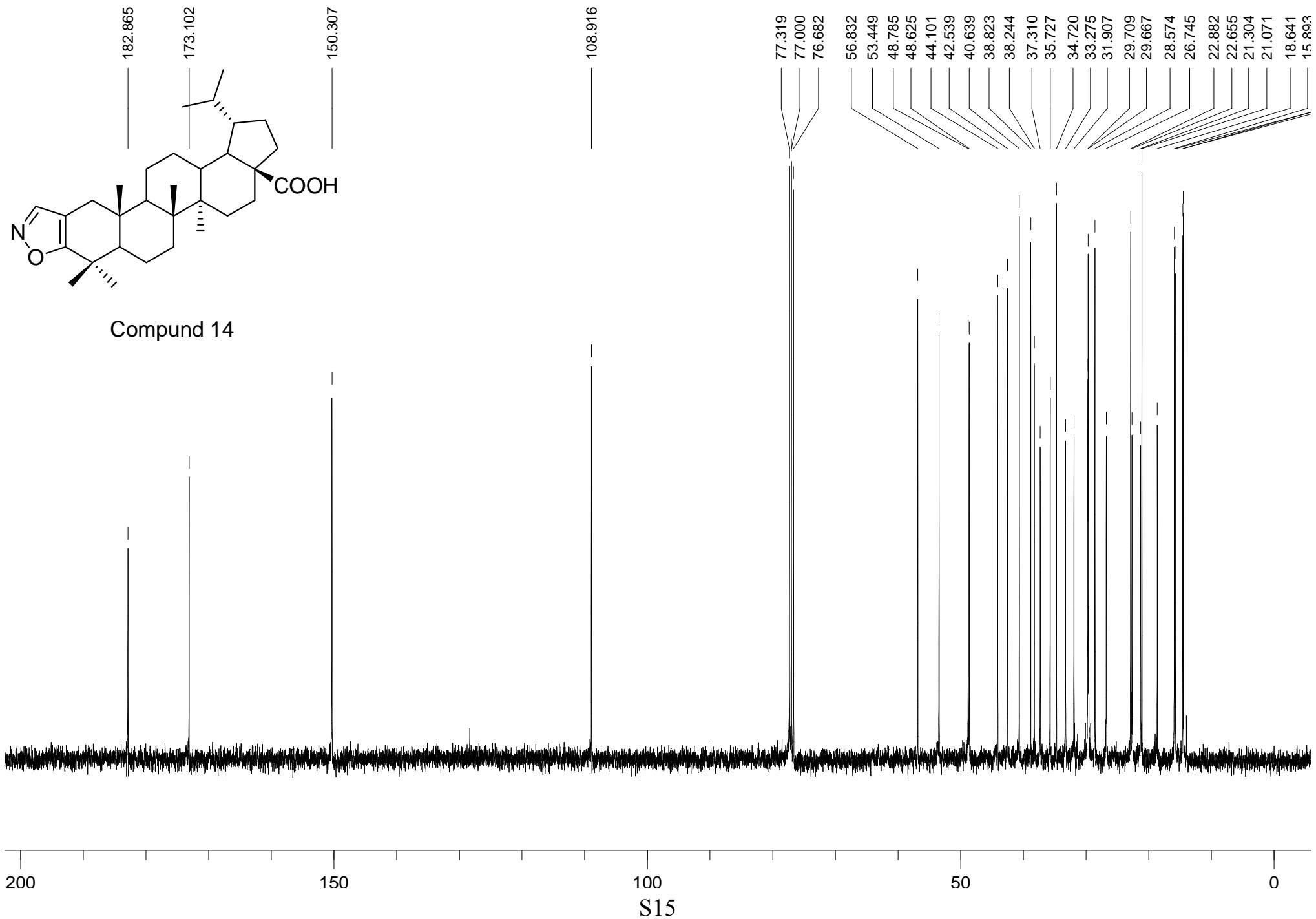
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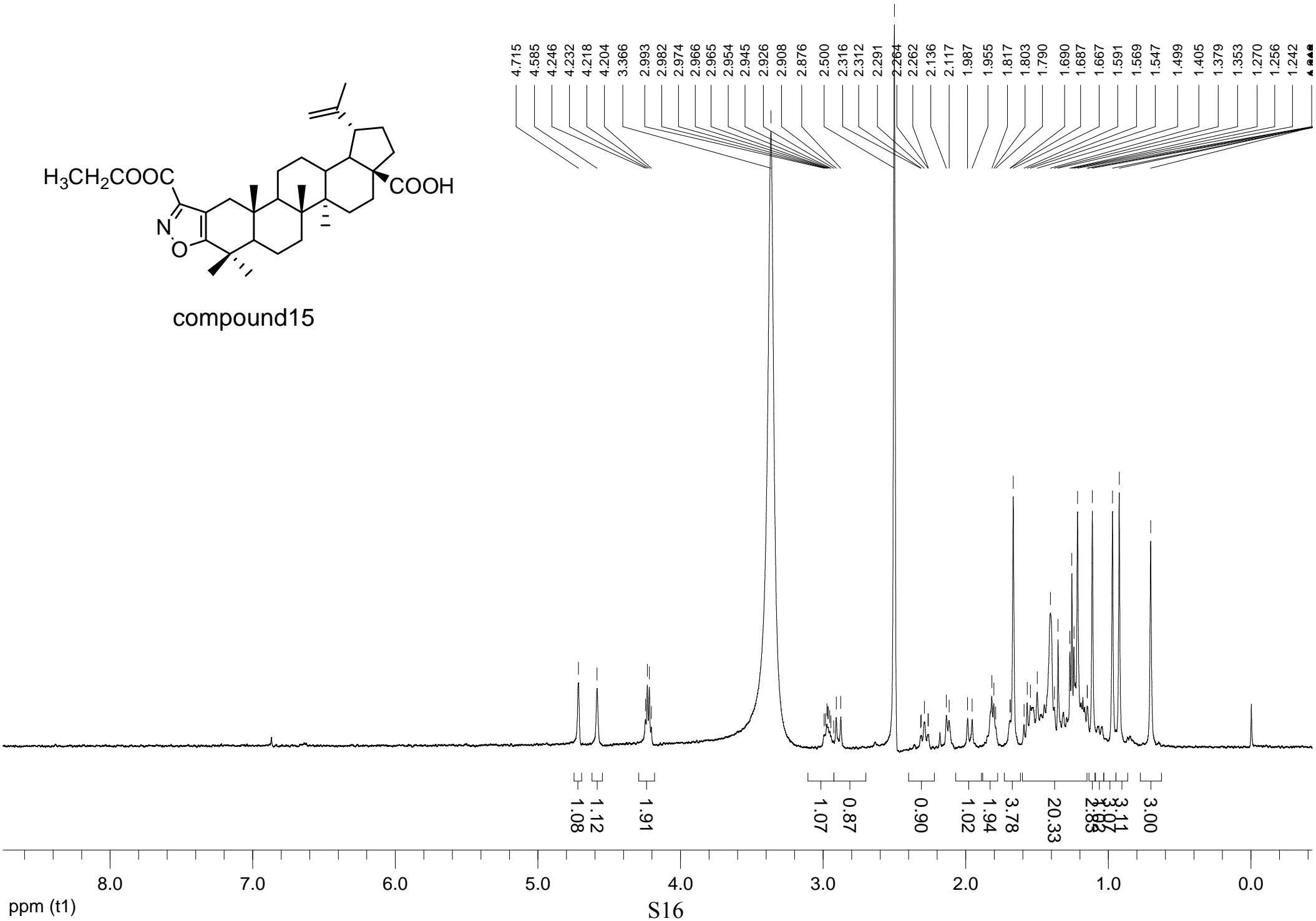
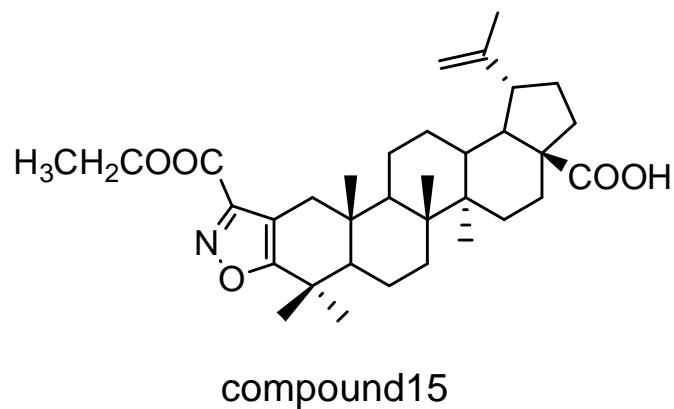
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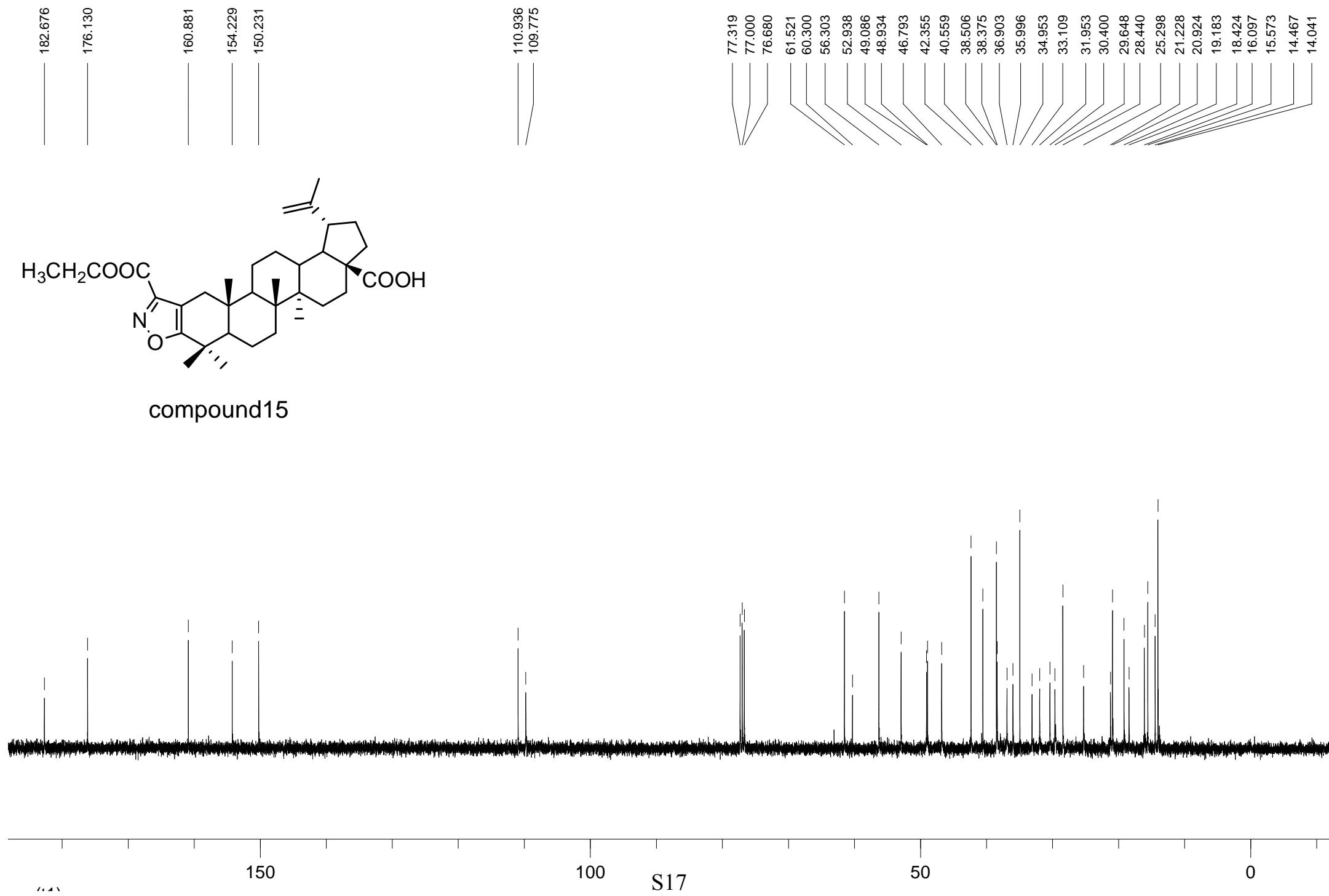


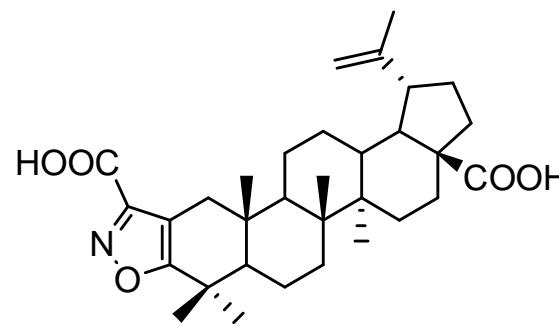
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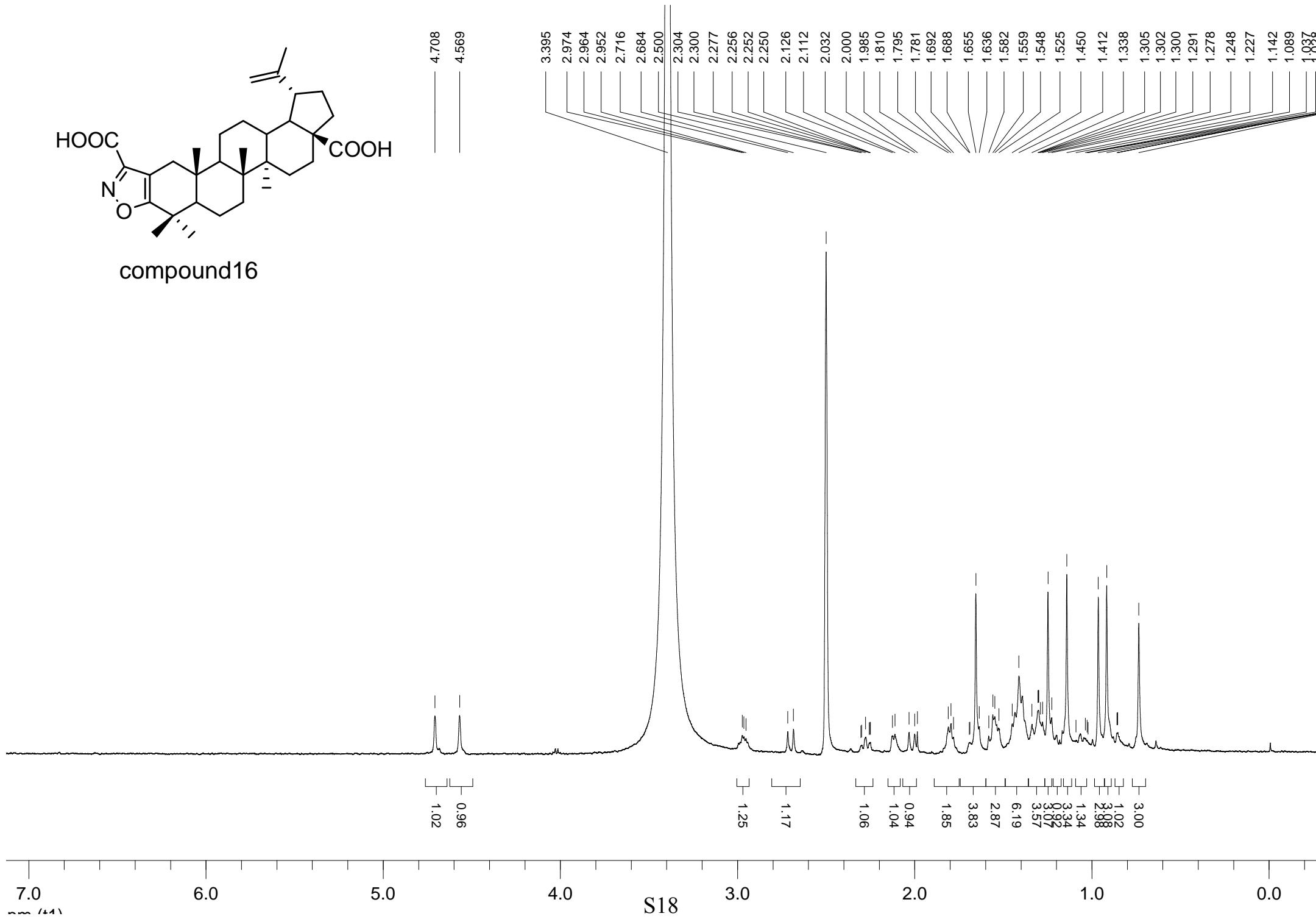


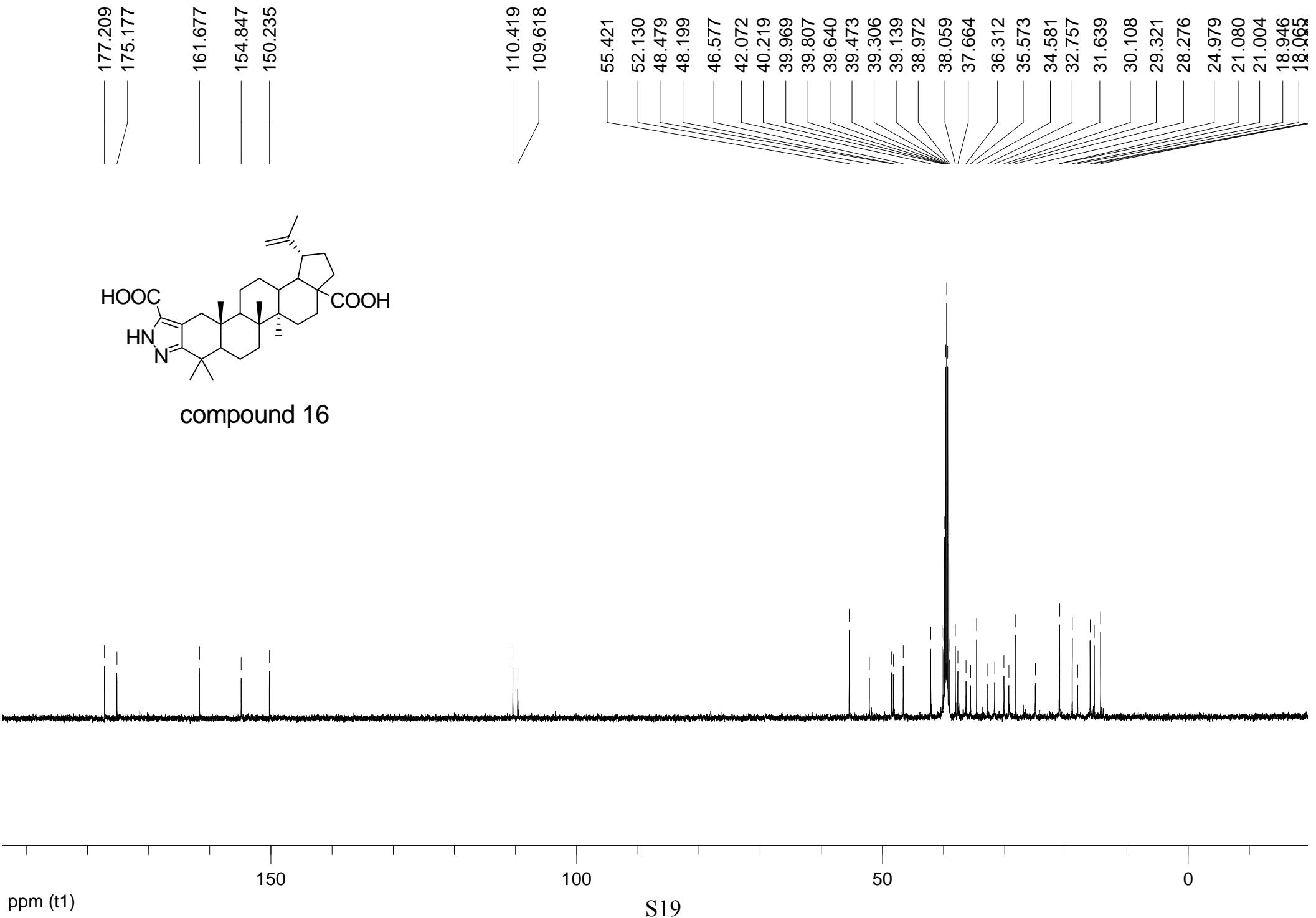


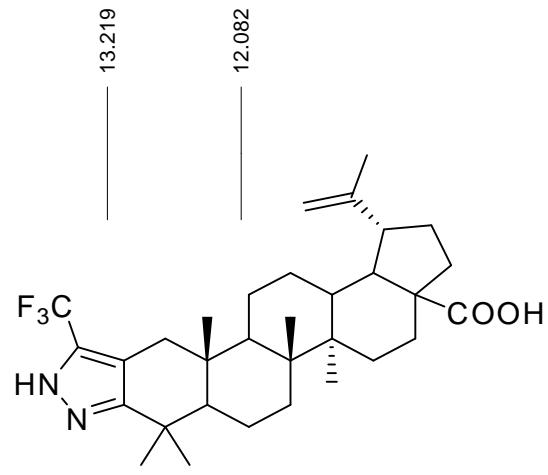




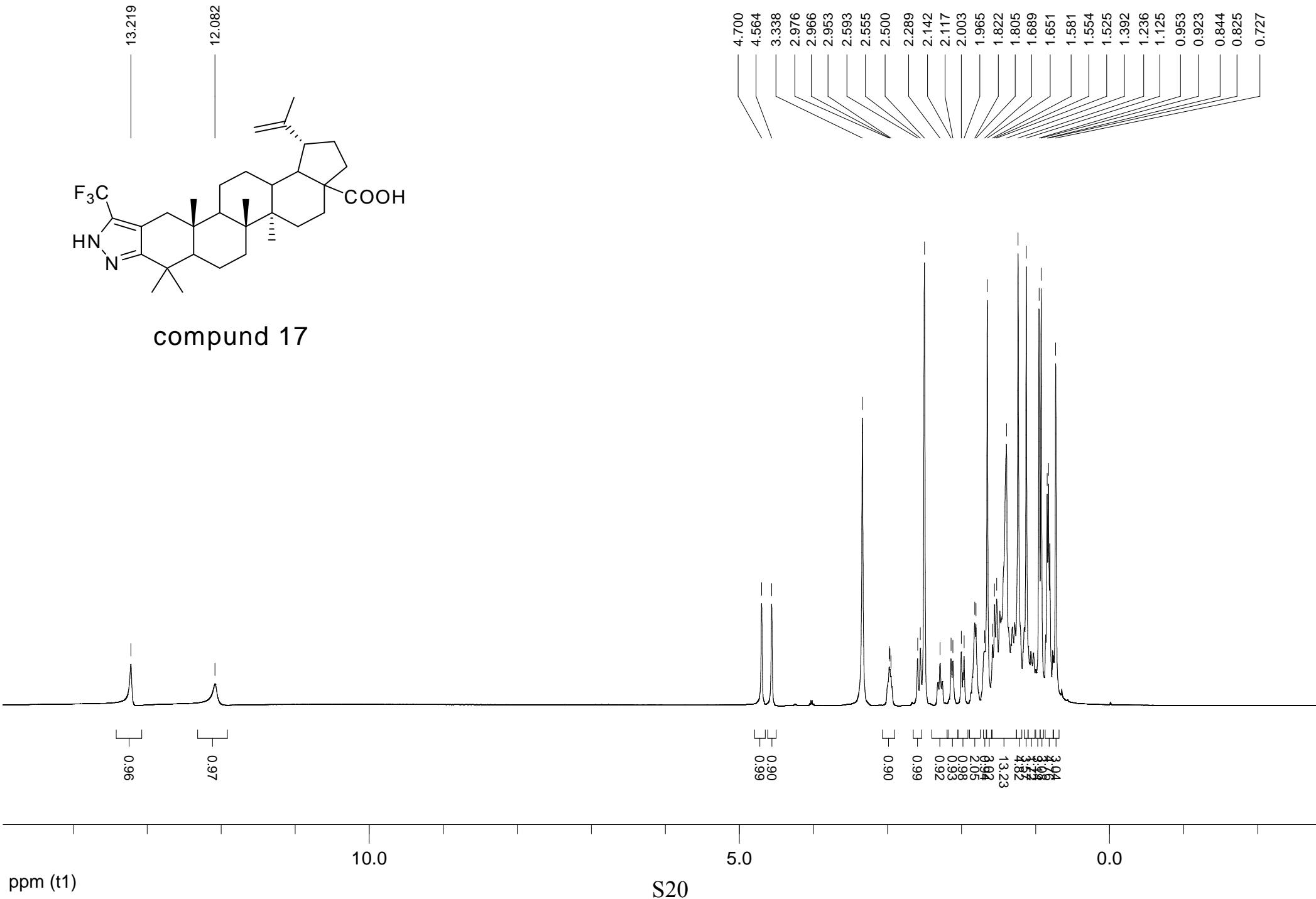
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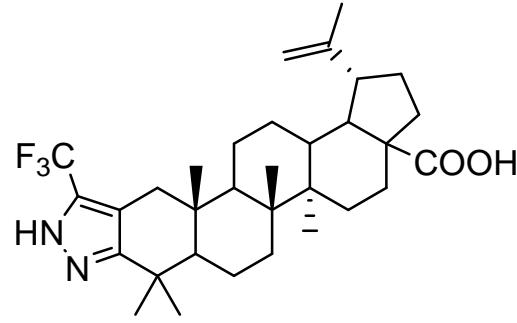
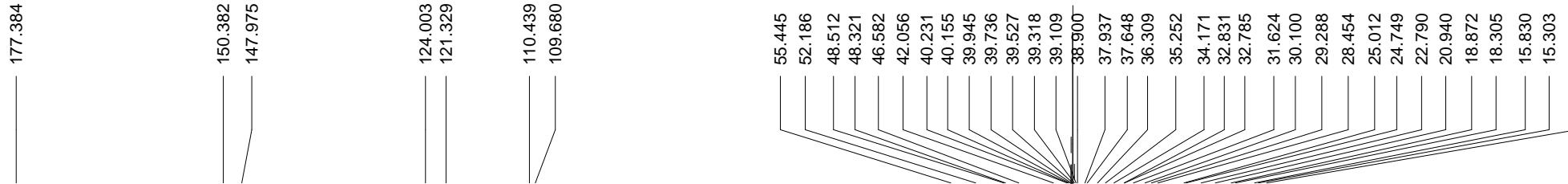




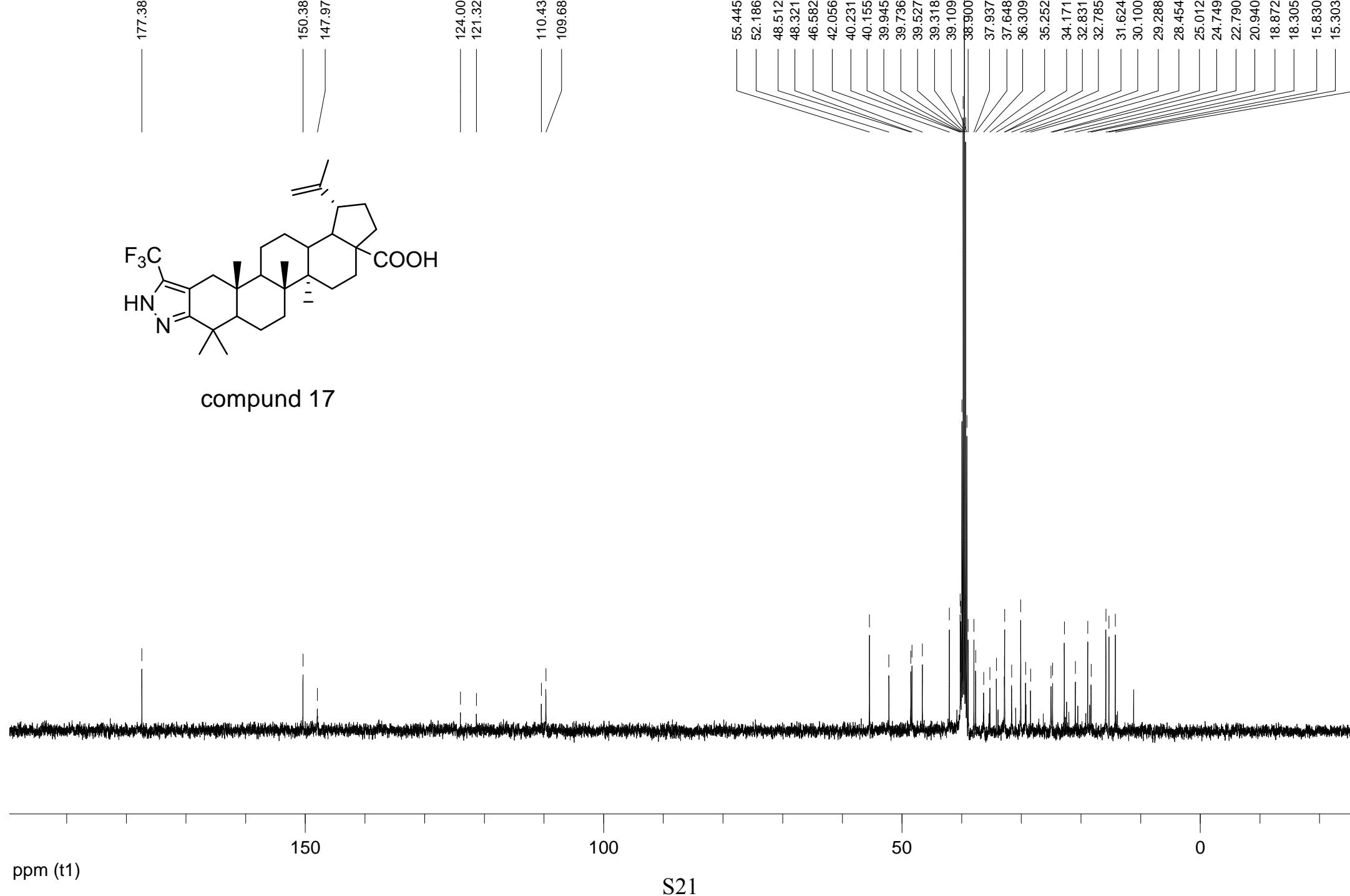


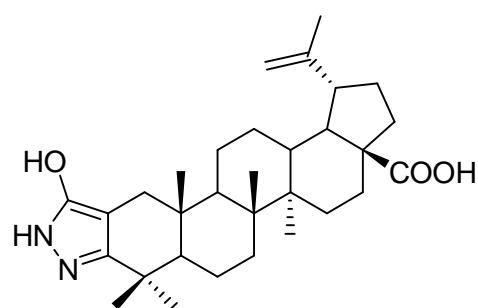
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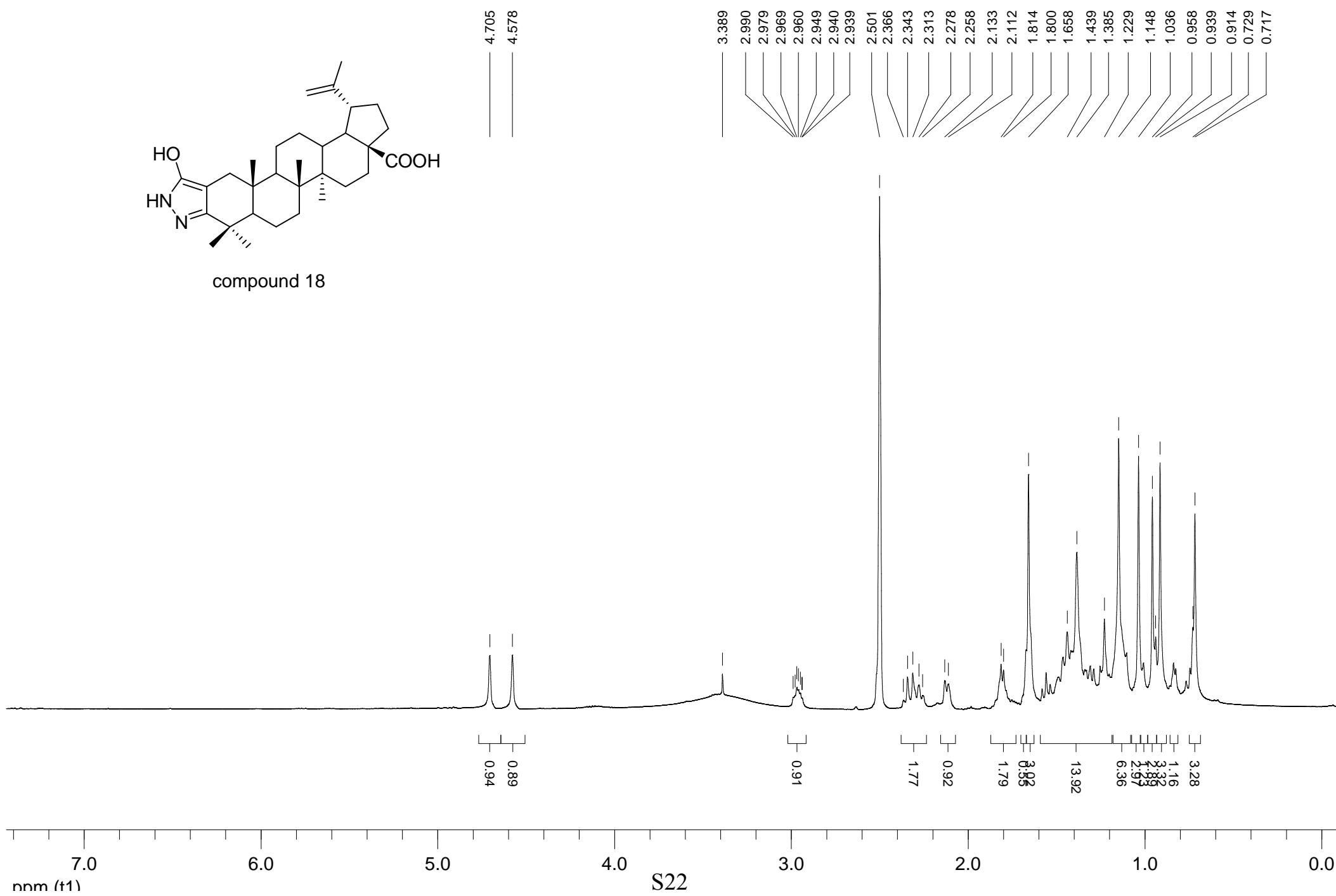


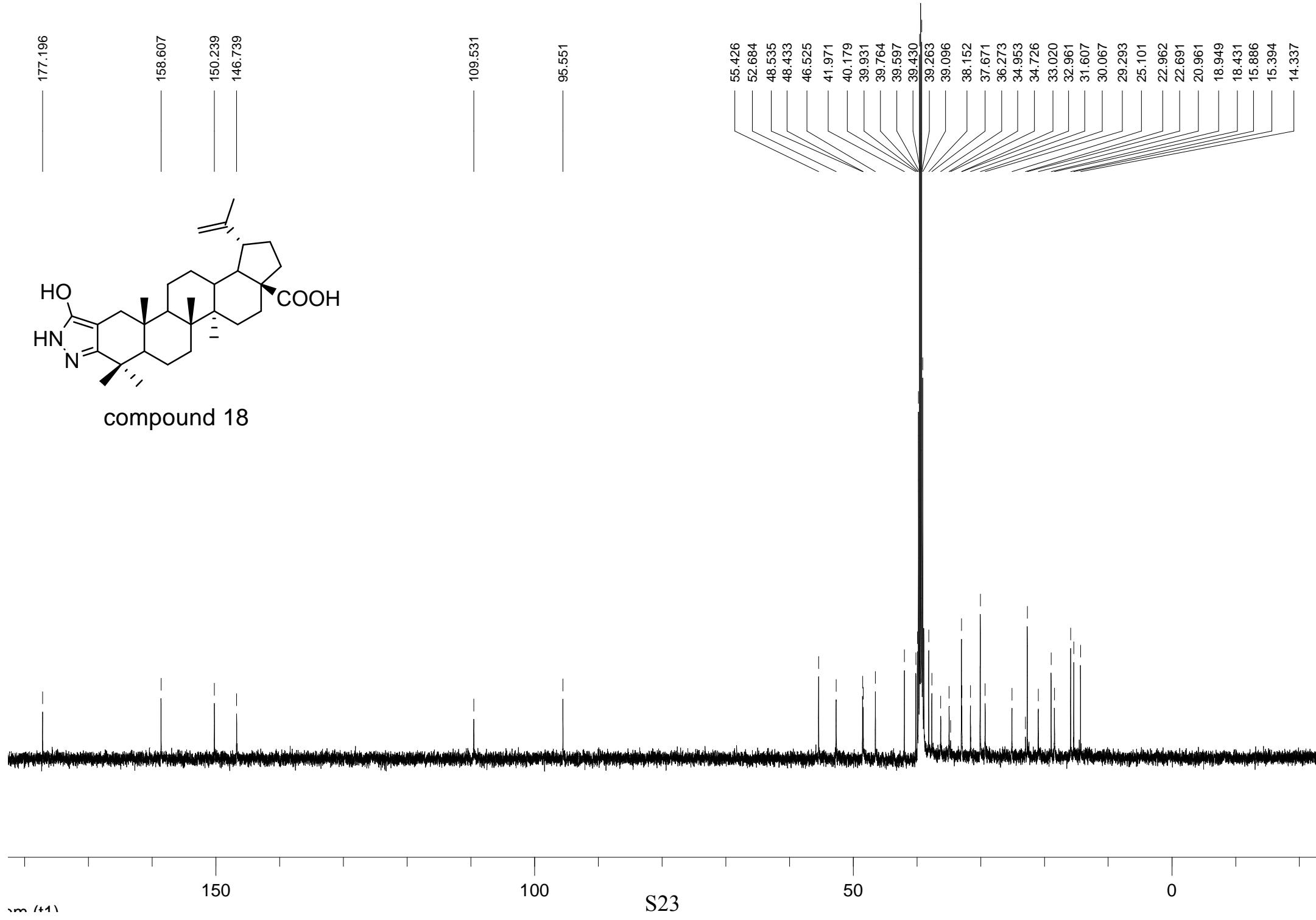
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compound 18





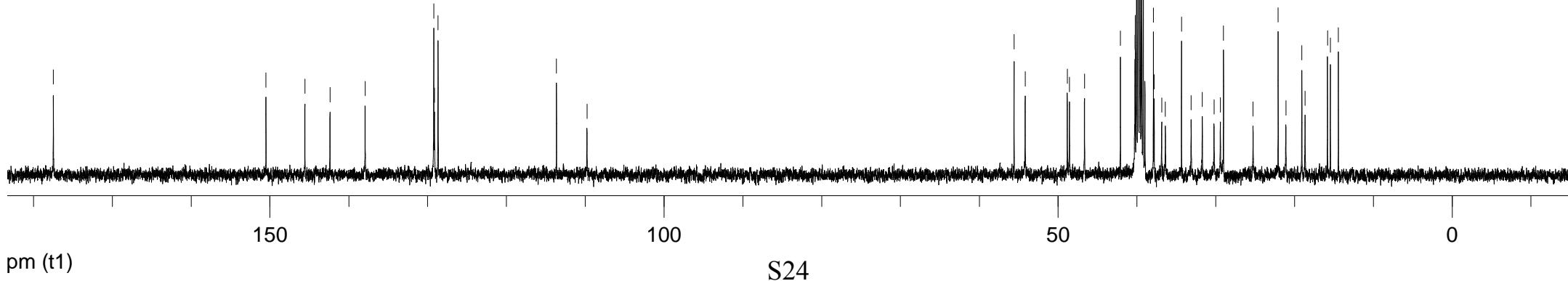
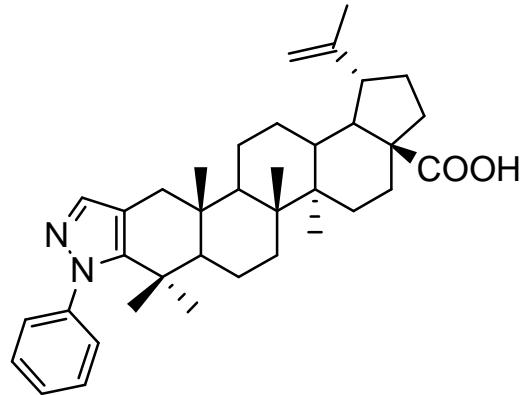
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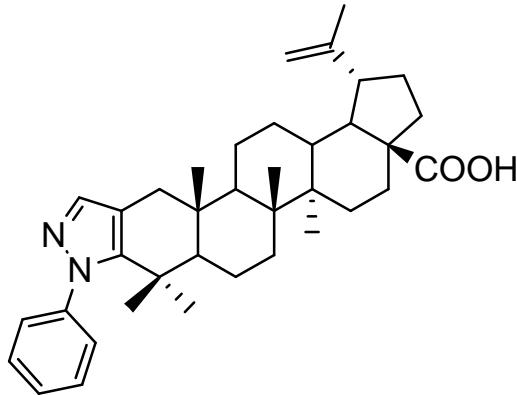
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12.093



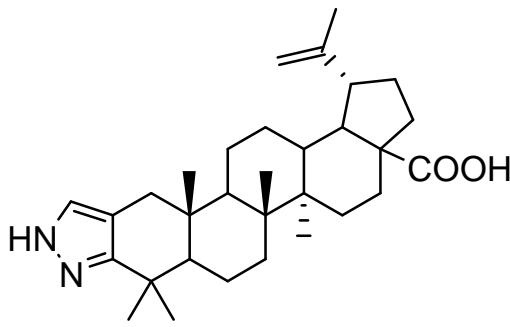
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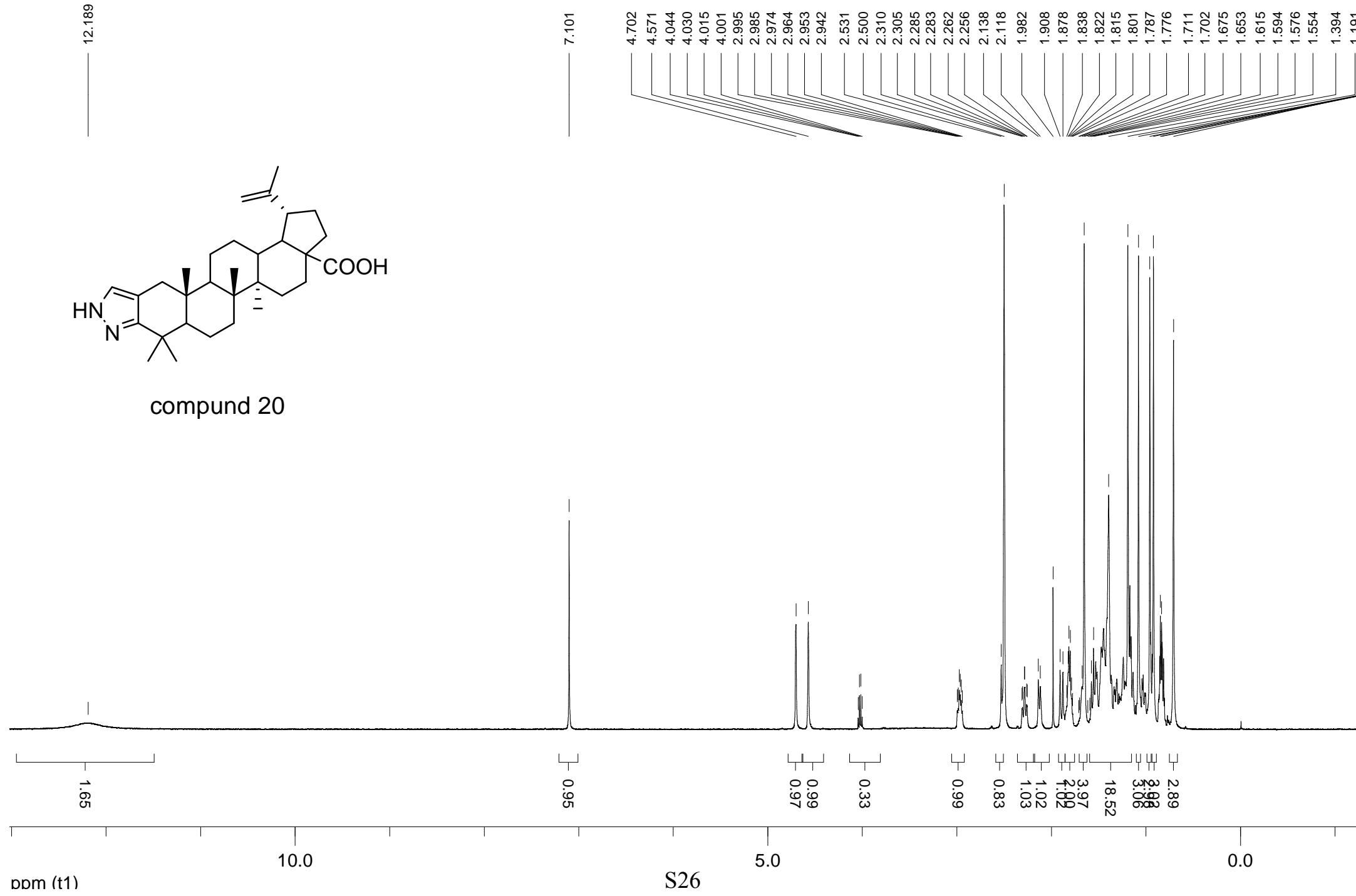
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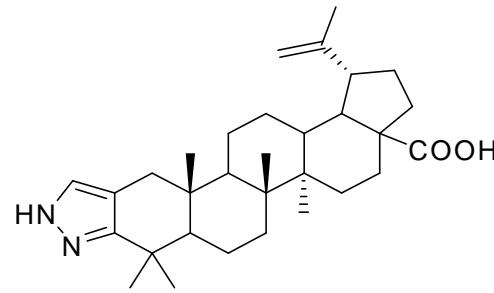
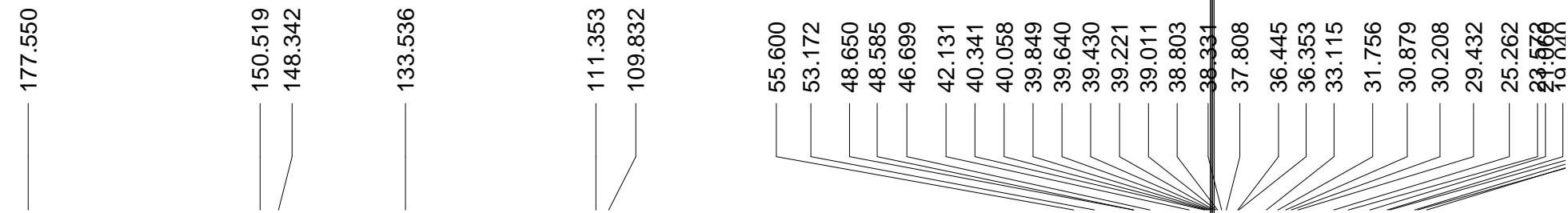
S25^{5.0}

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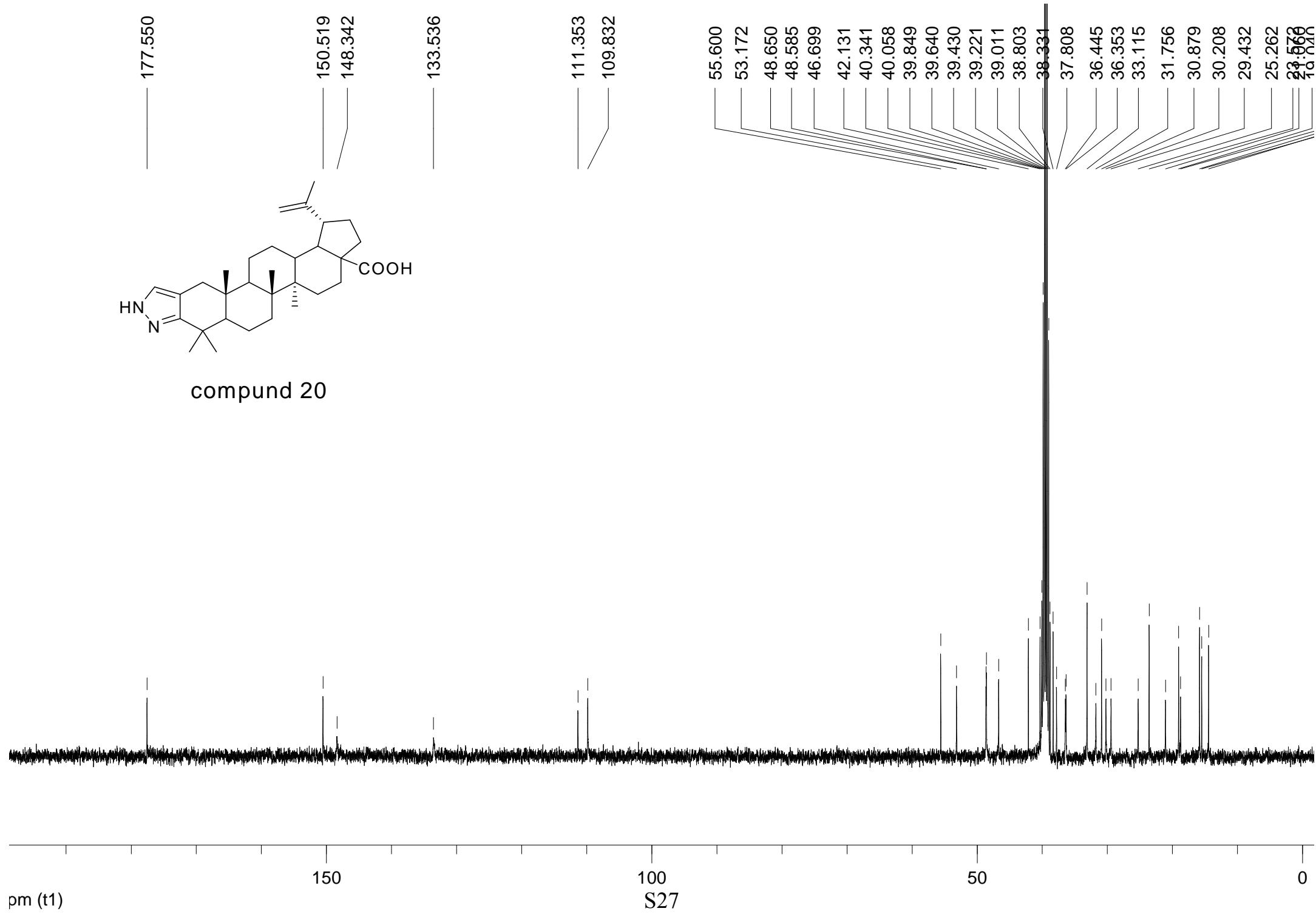


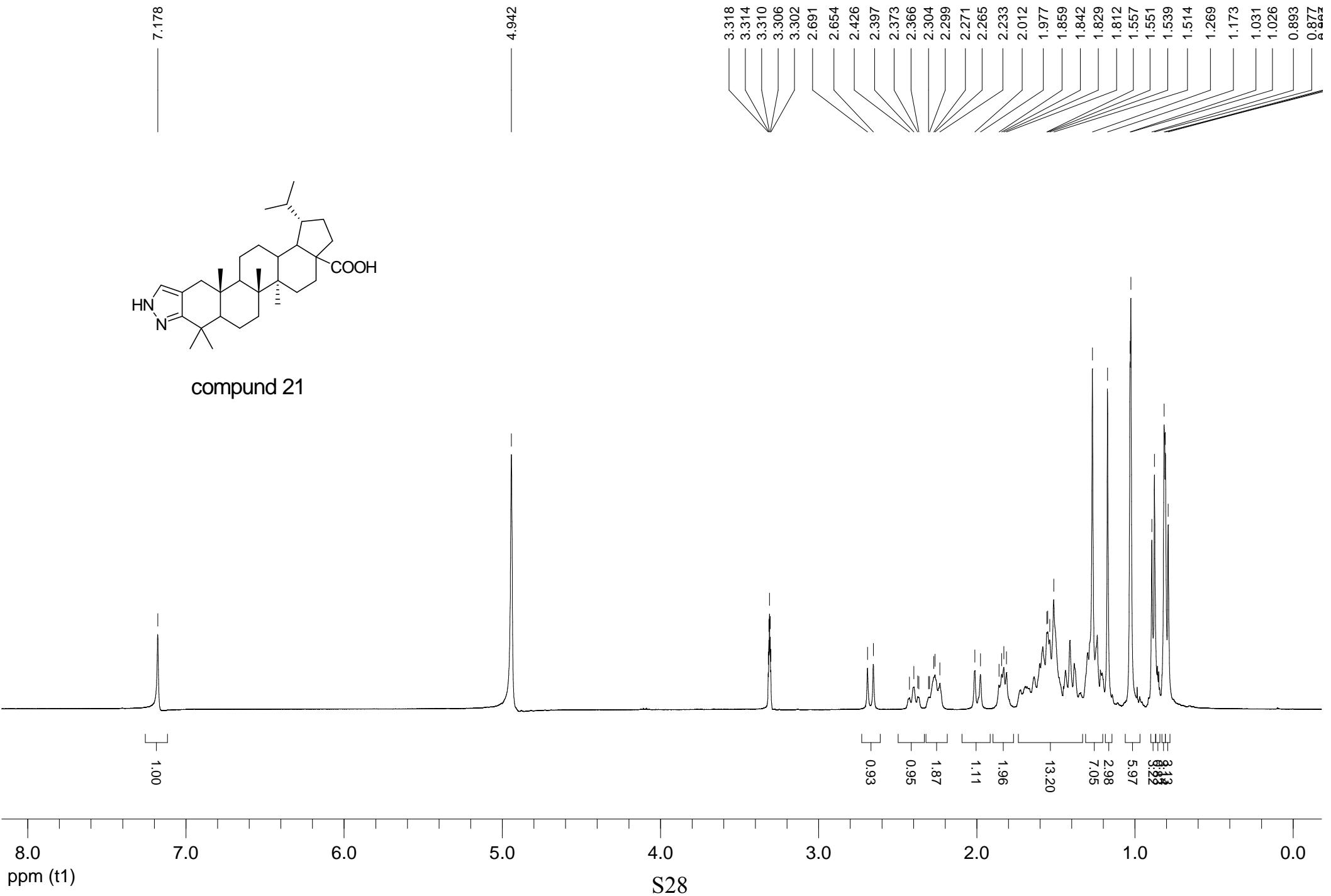
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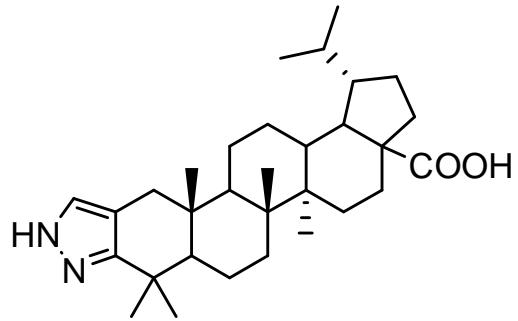




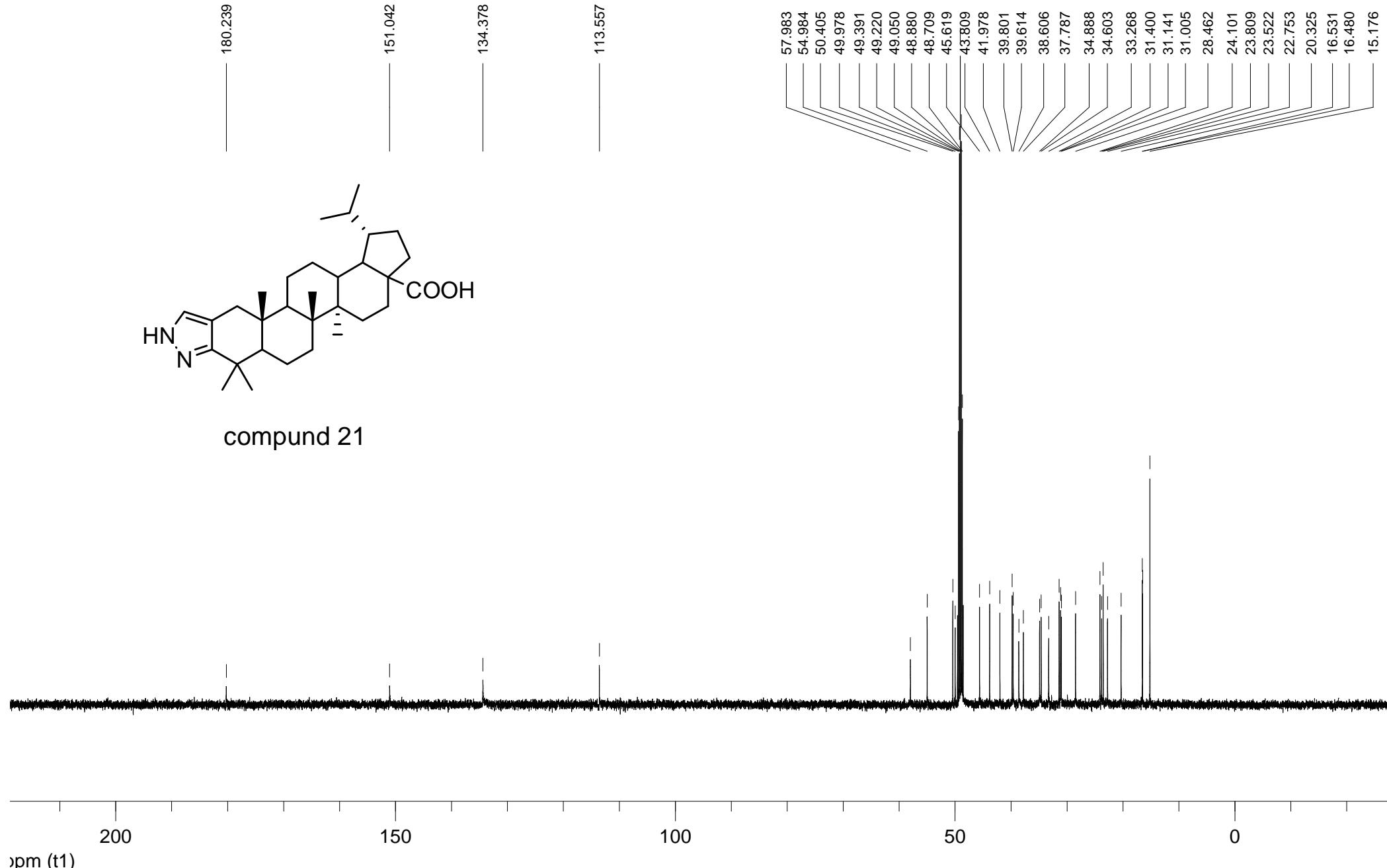
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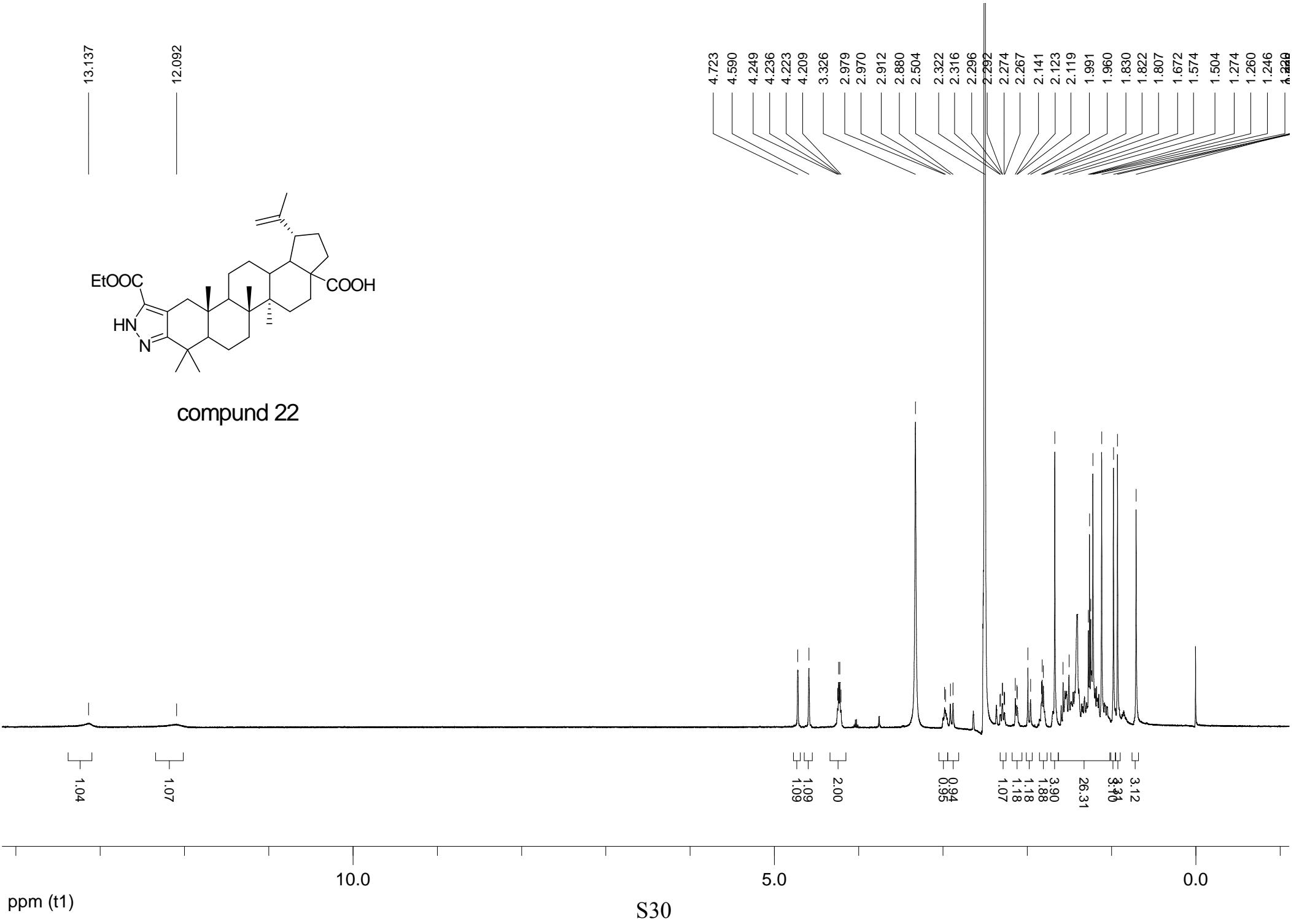


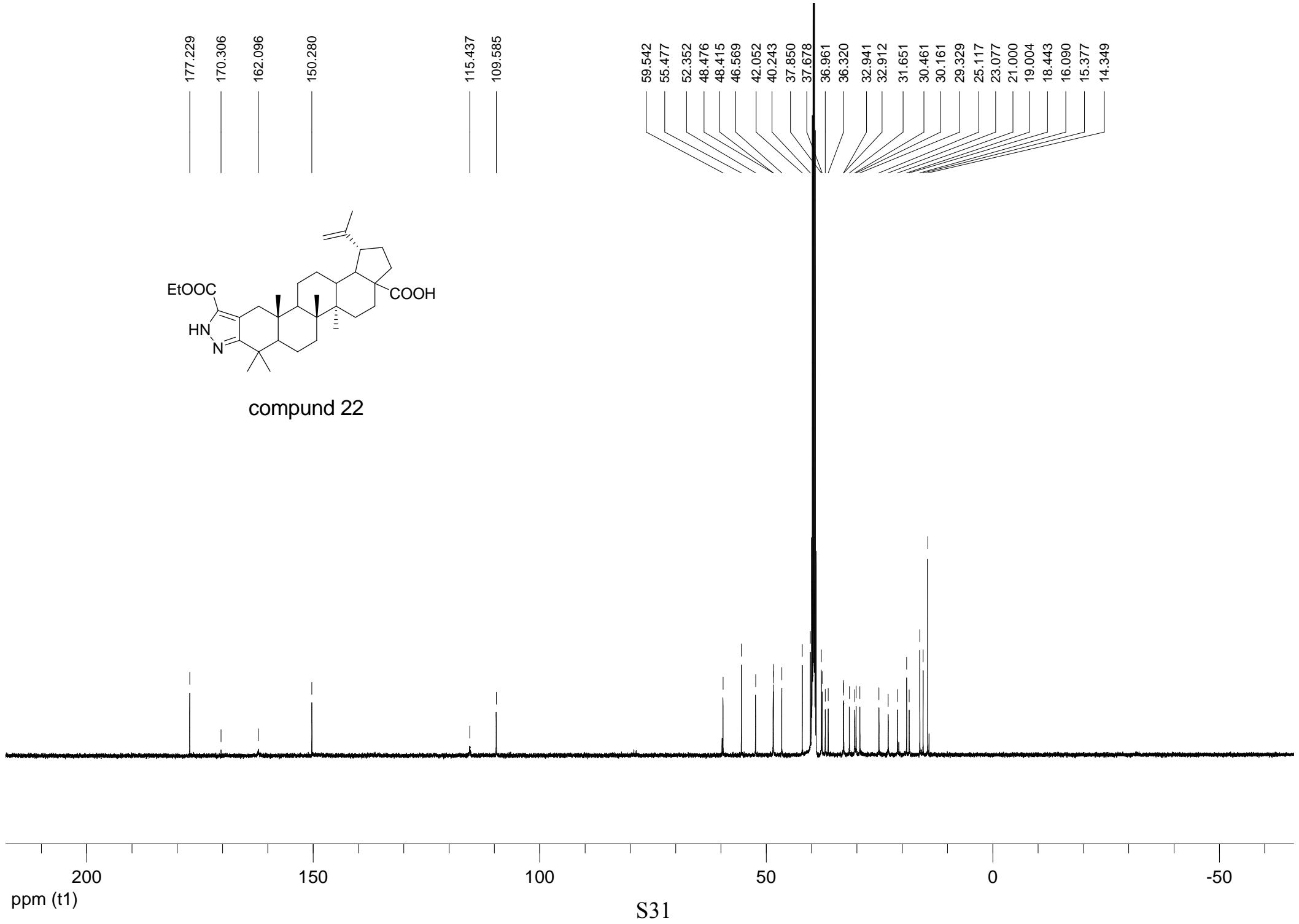


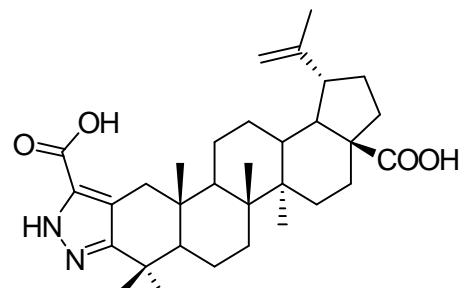


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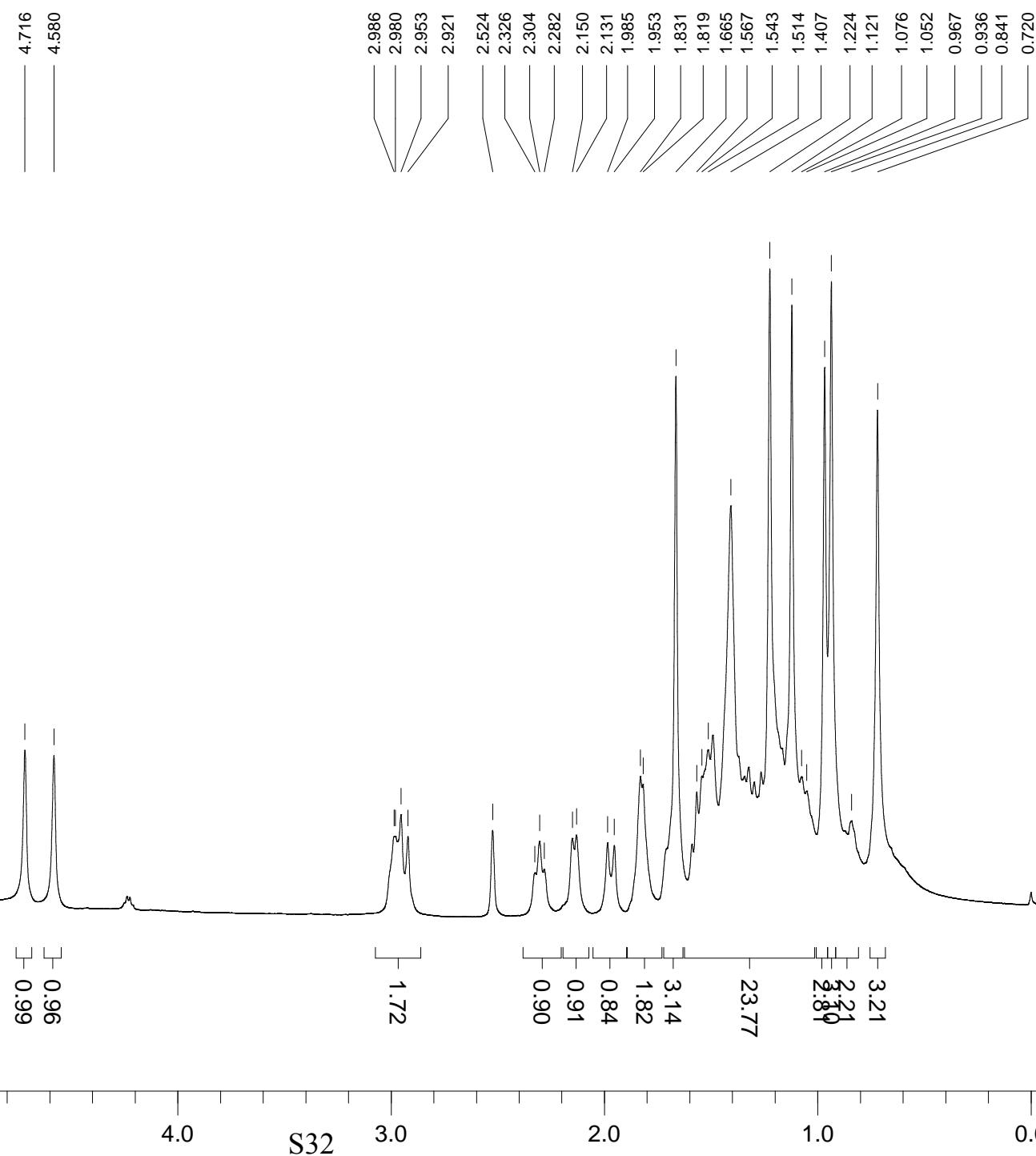


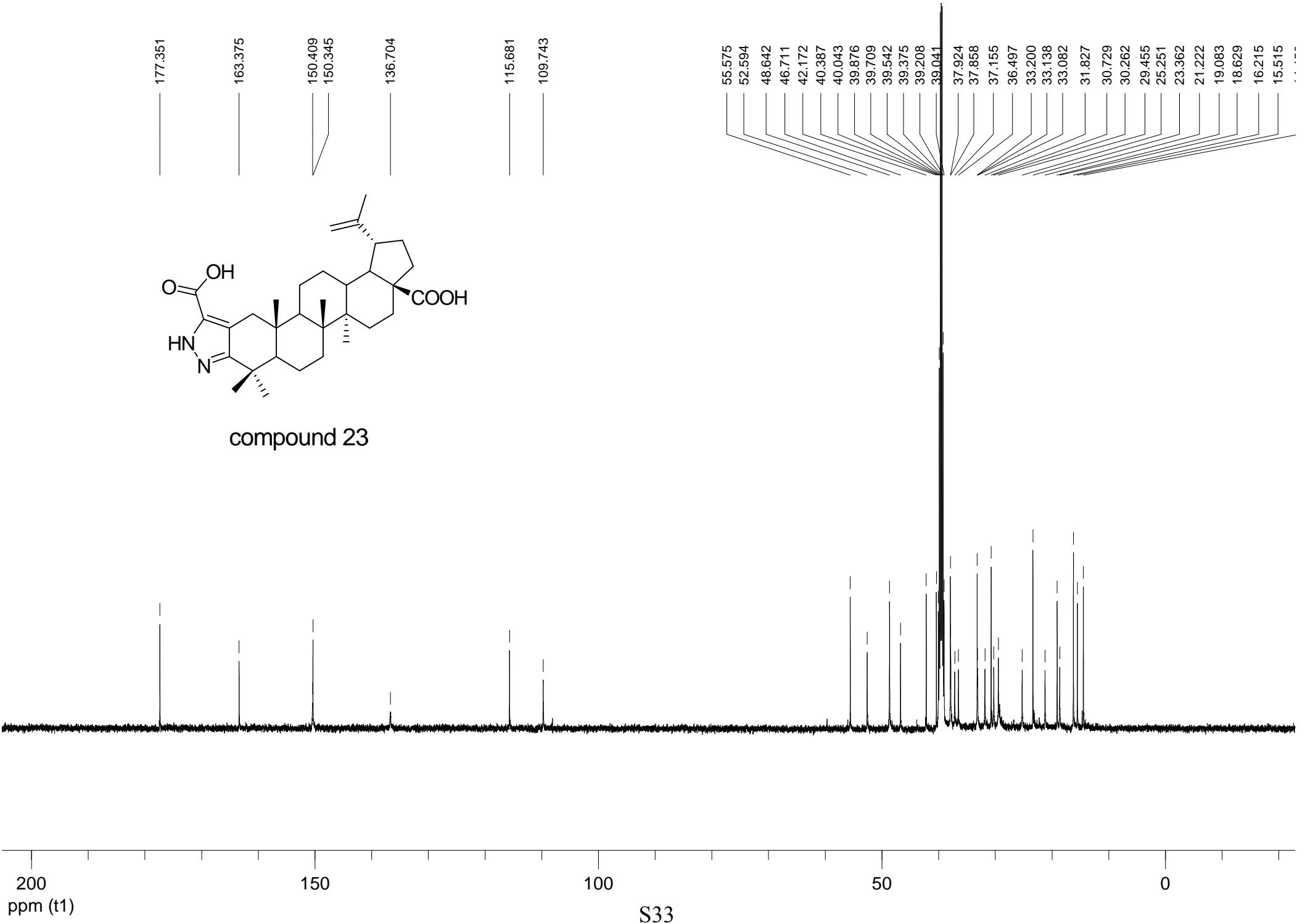




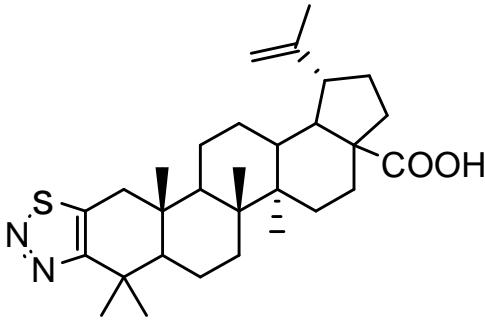


compound 23





7.260



compund 27

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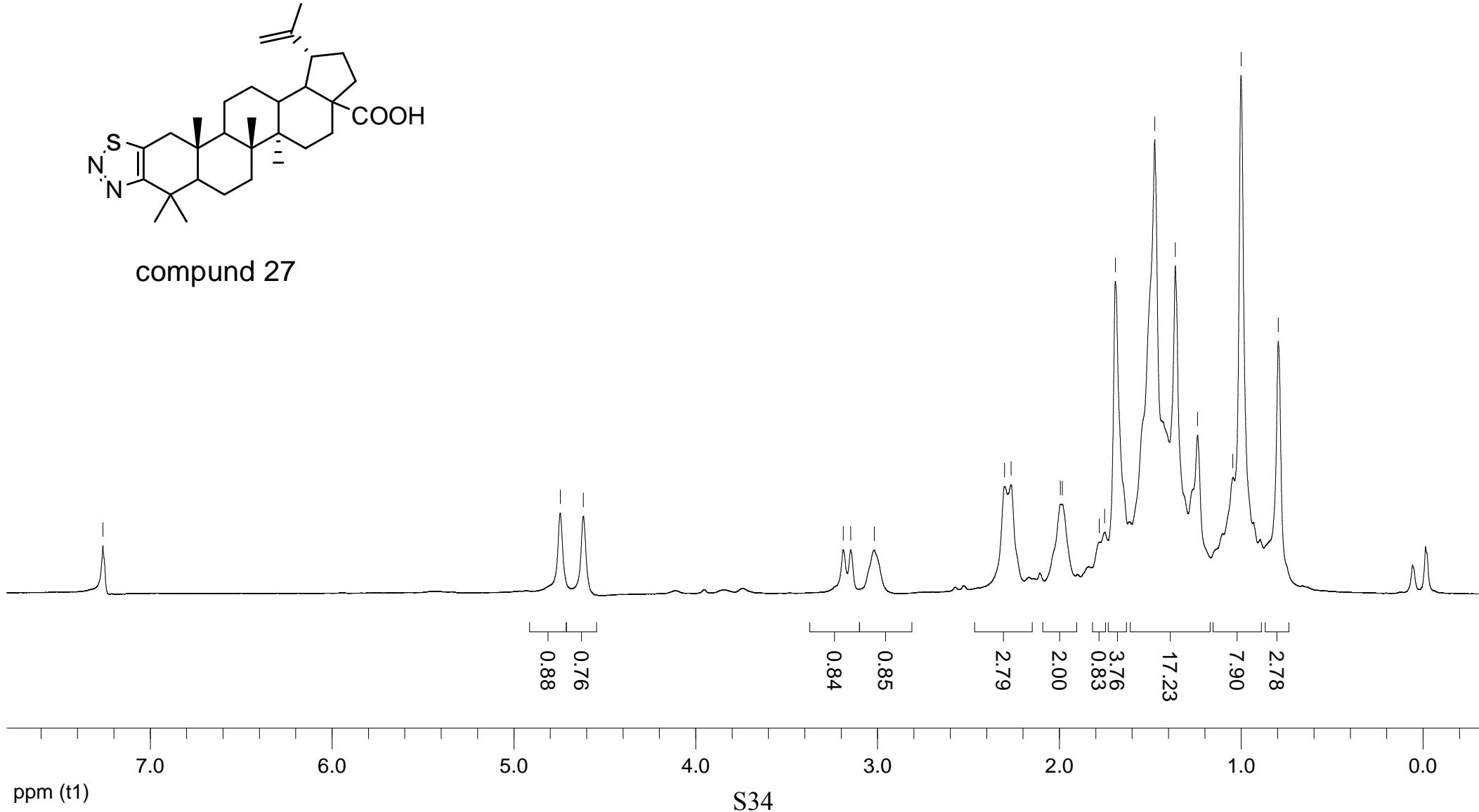
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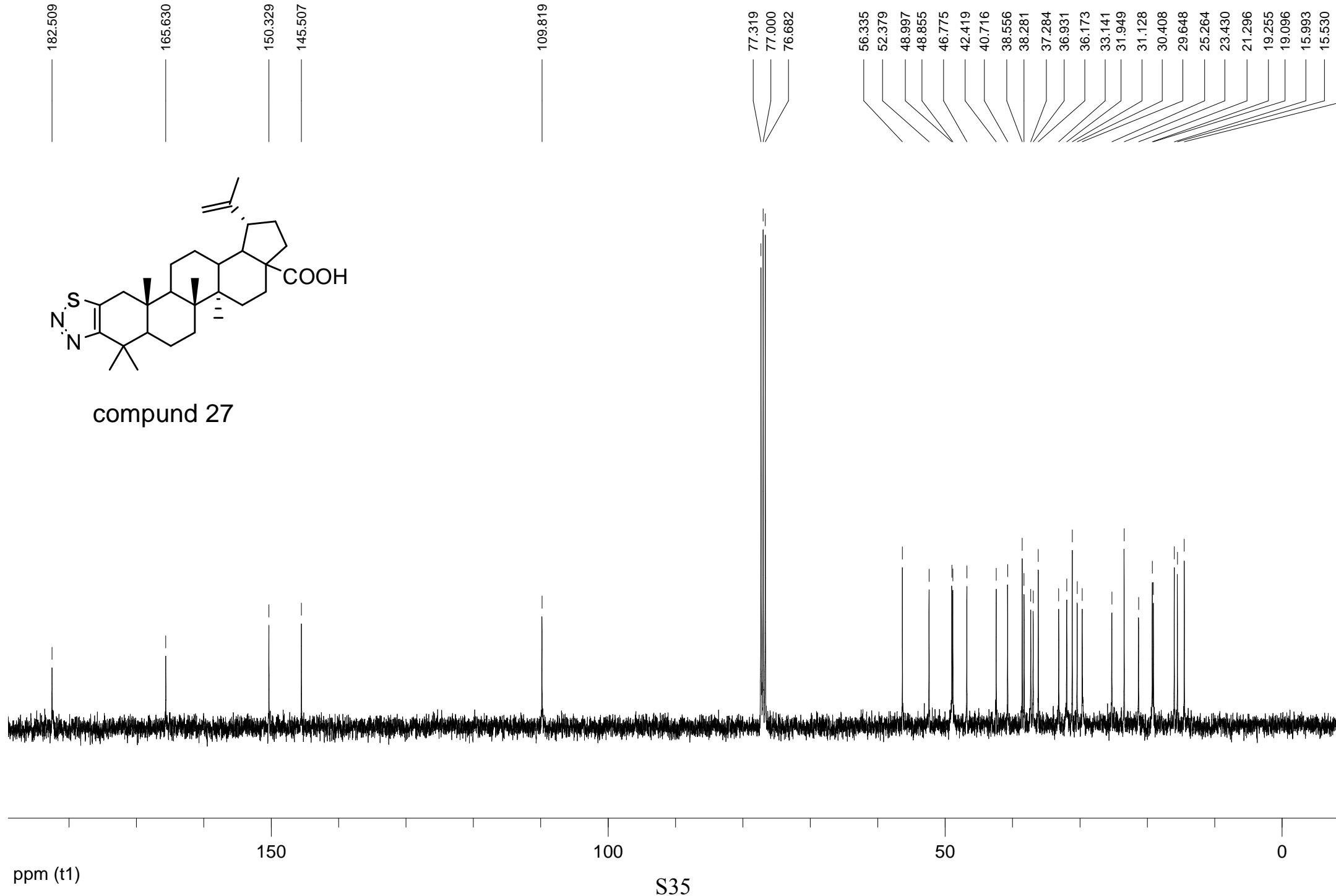
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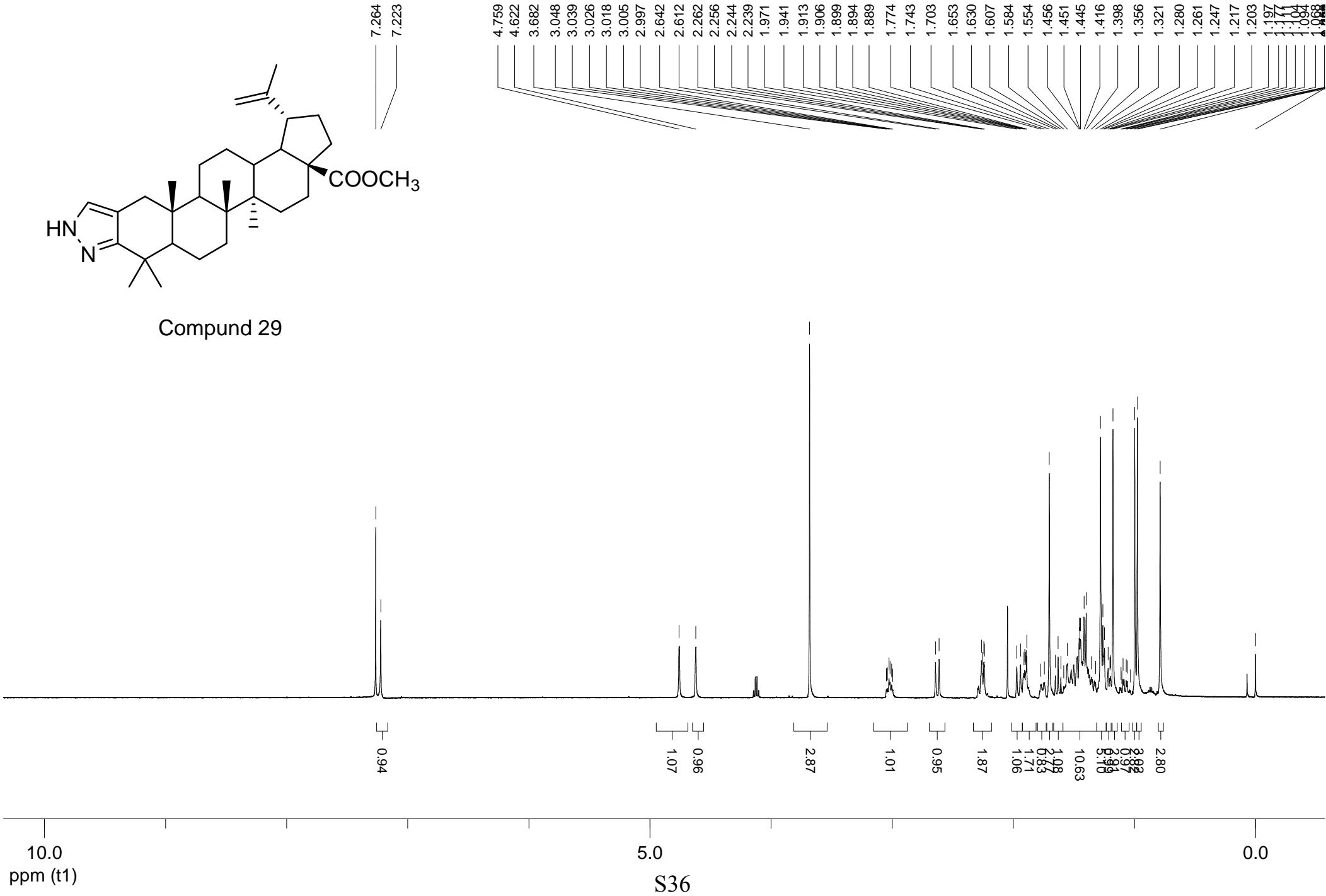
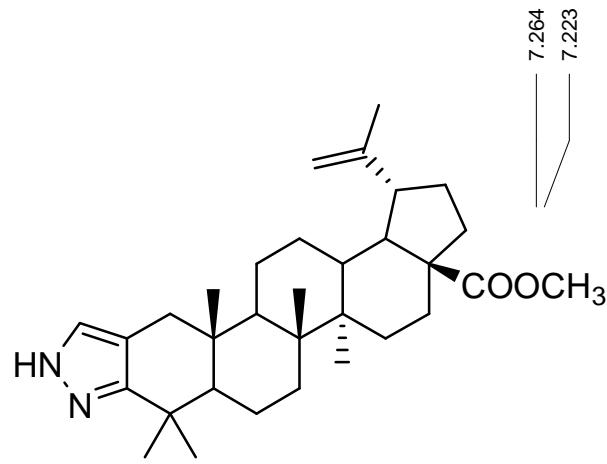
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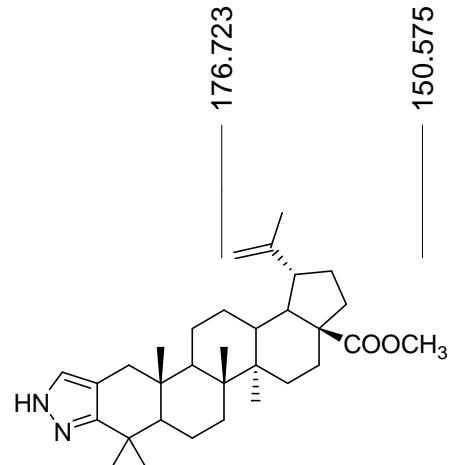


ppm (t1)

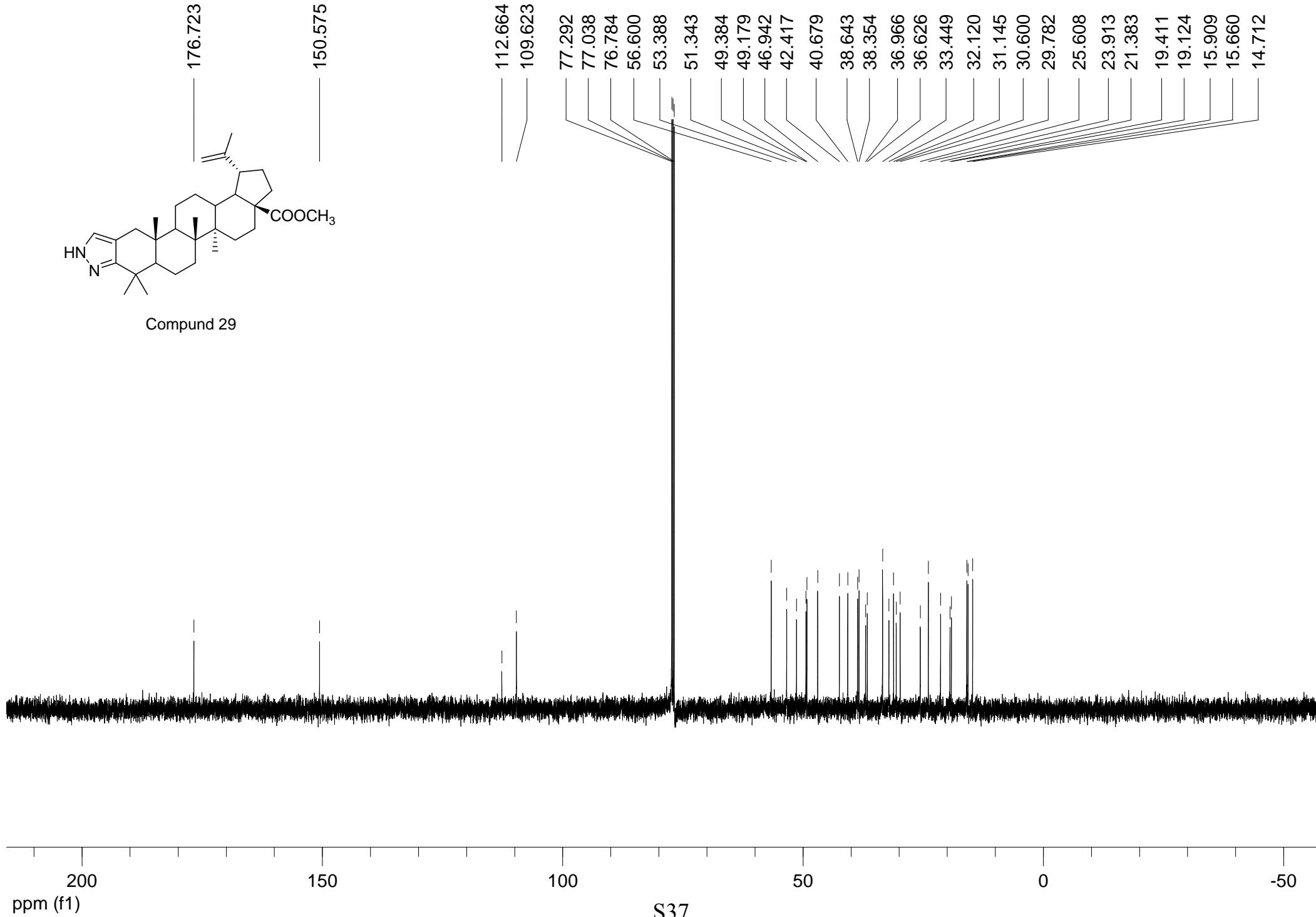
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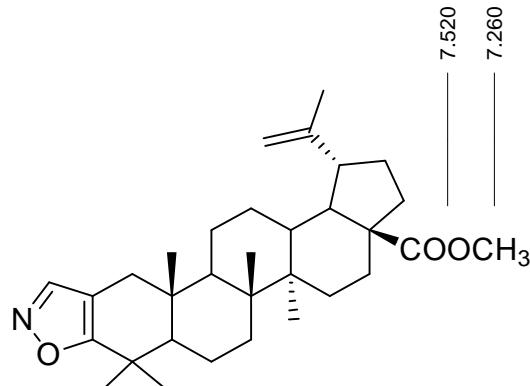




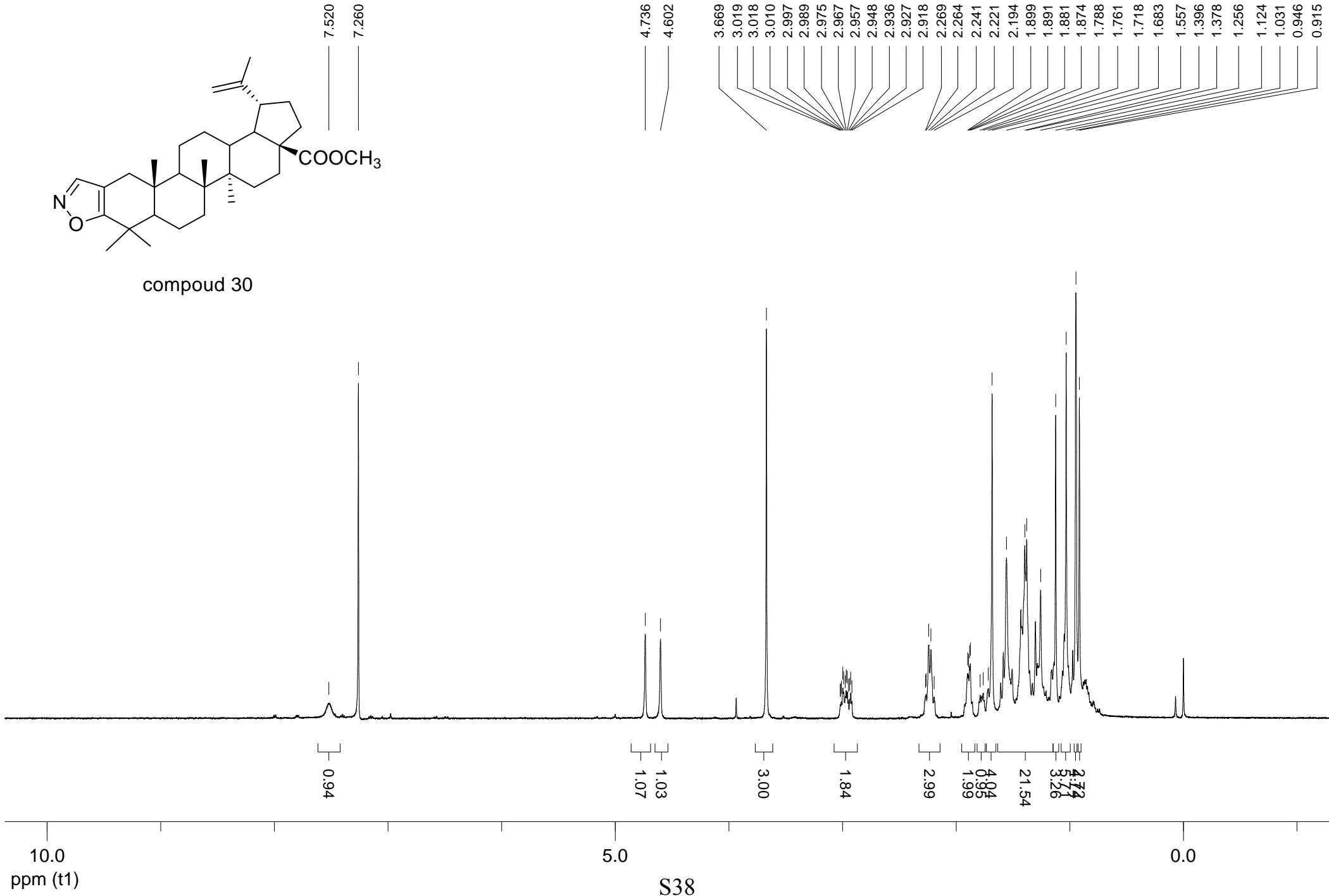


Compound 29





compoud 30



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S38

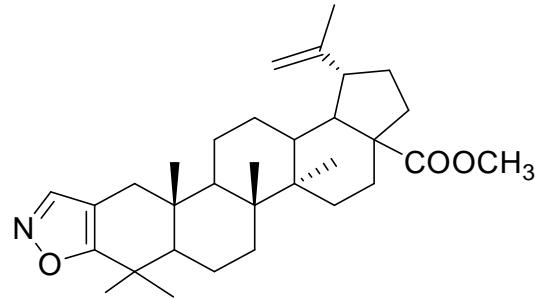
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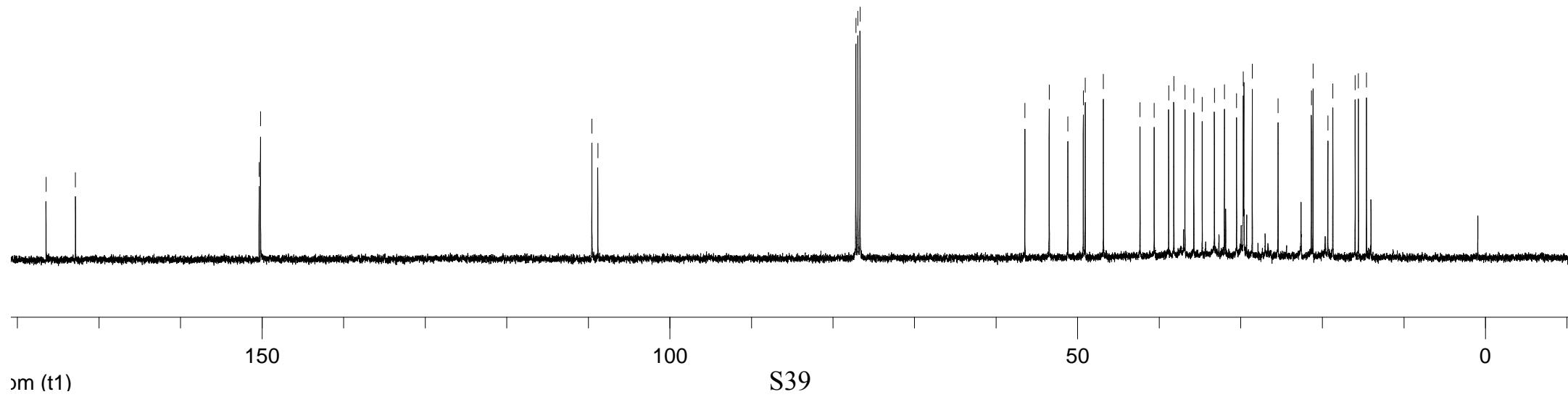
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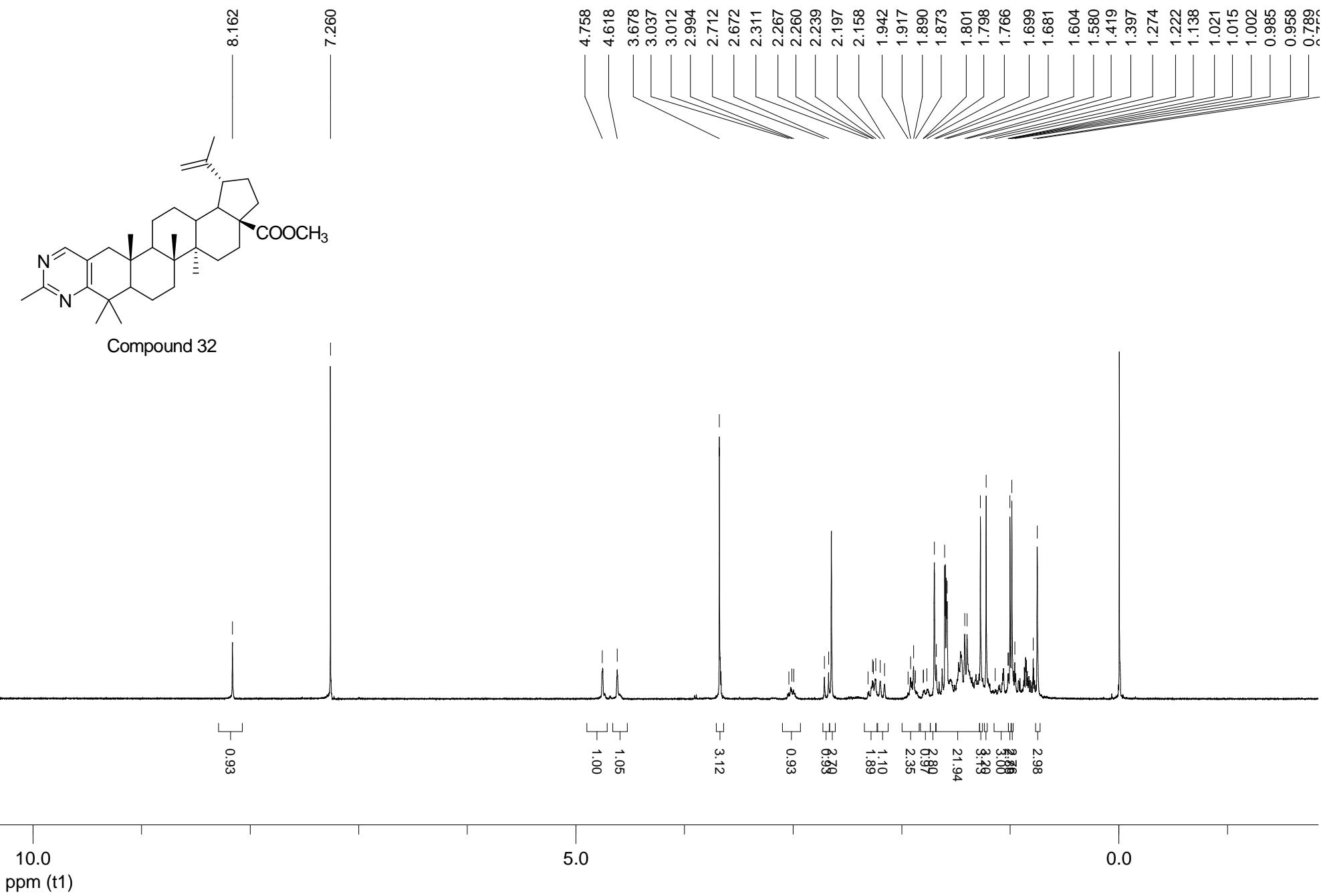
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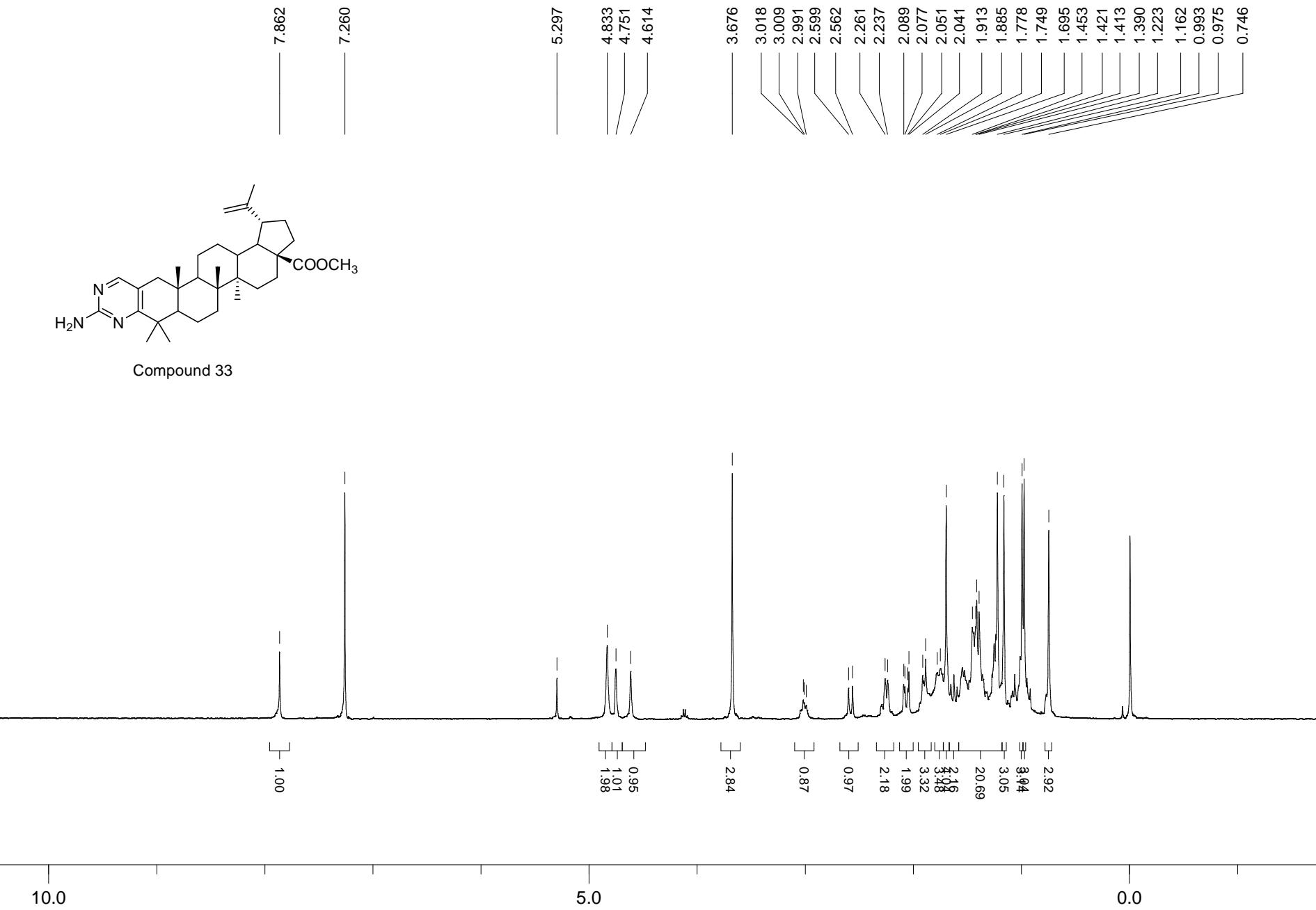
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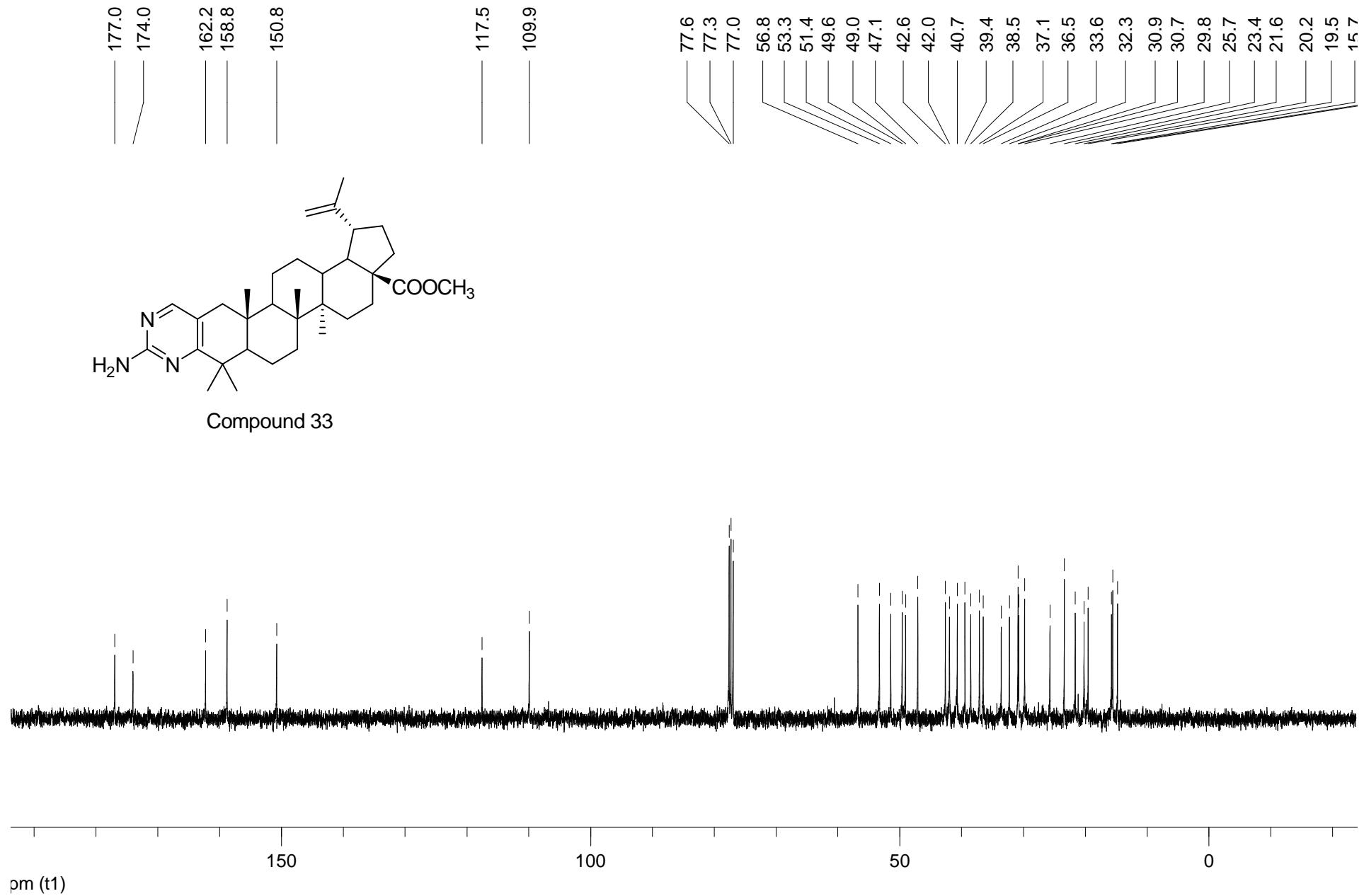


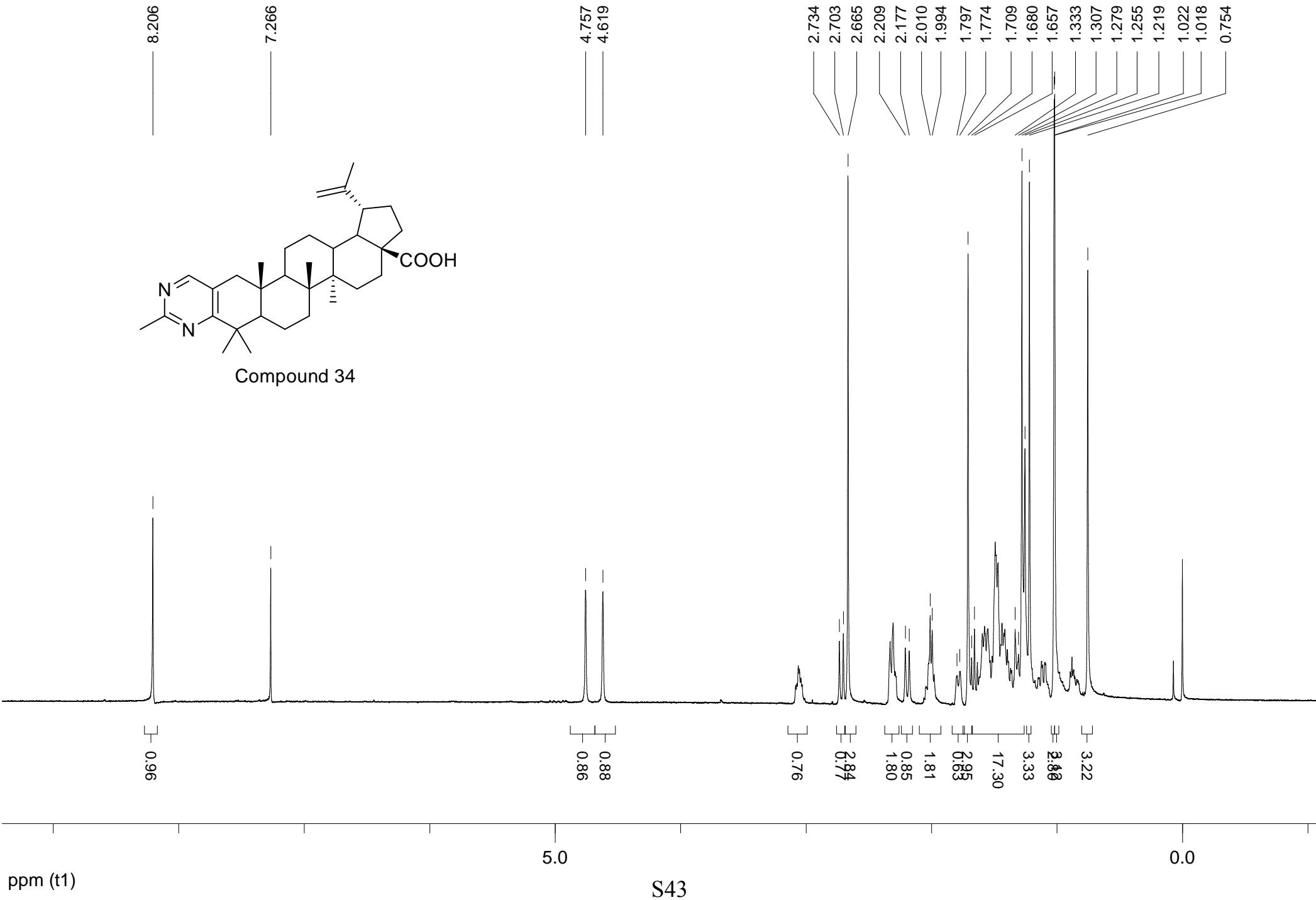
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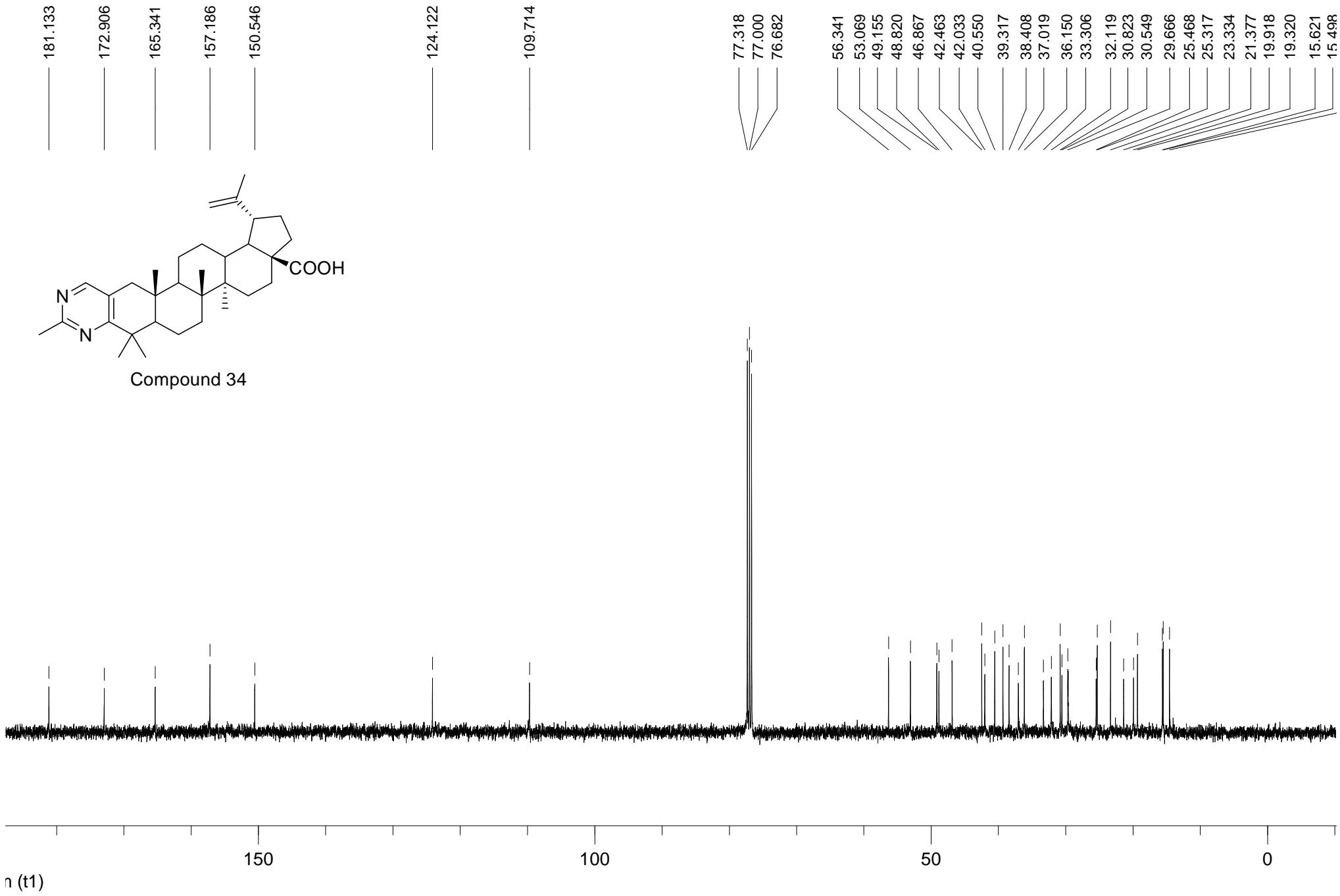


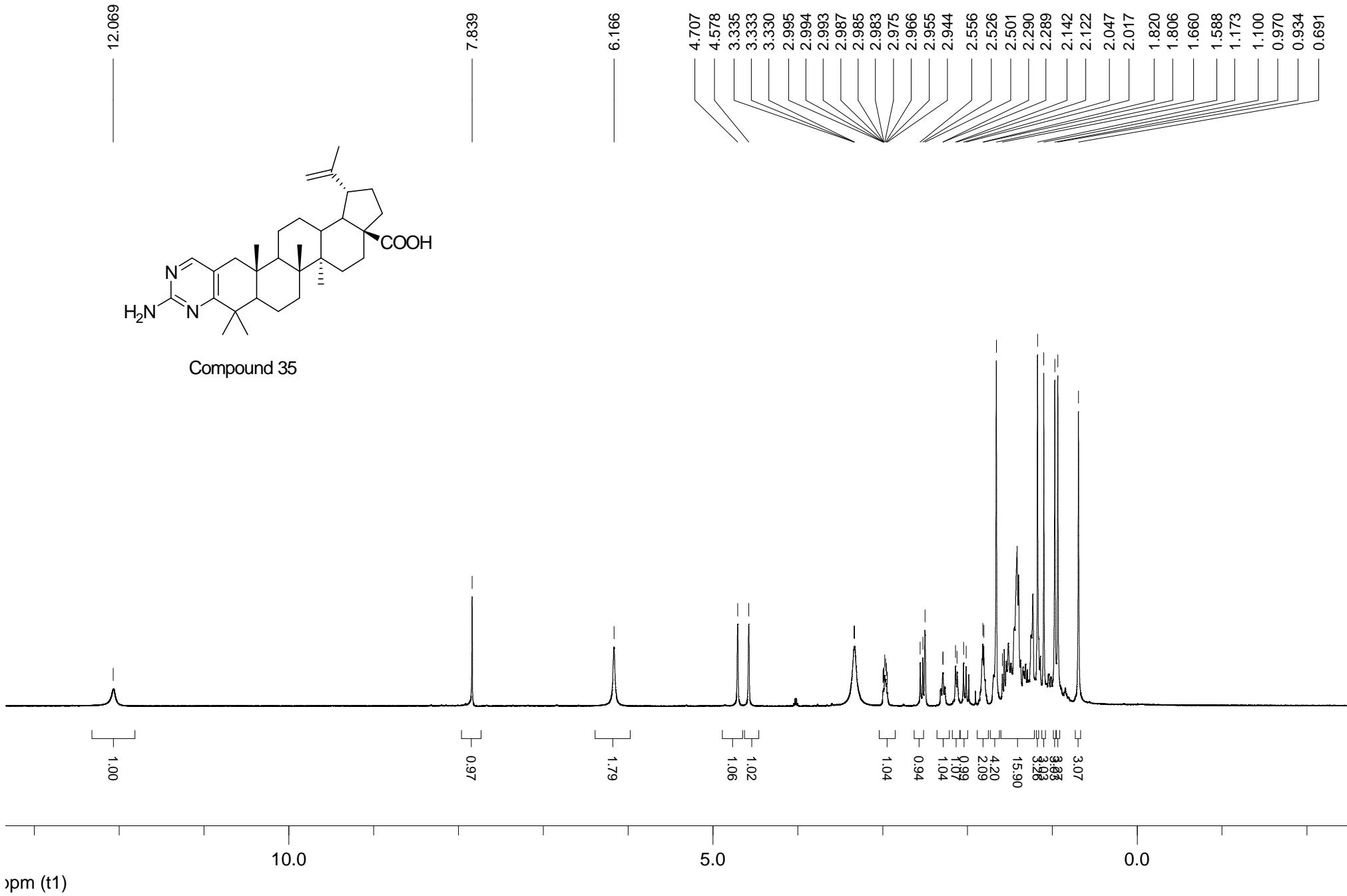


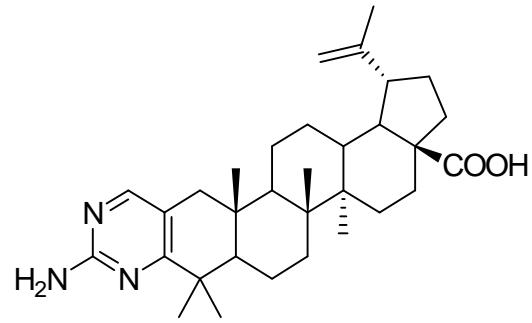
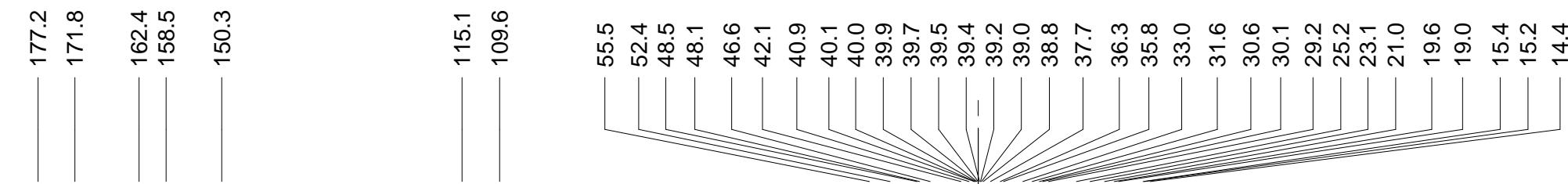




S43







Compound 35

