

## Supporting Information

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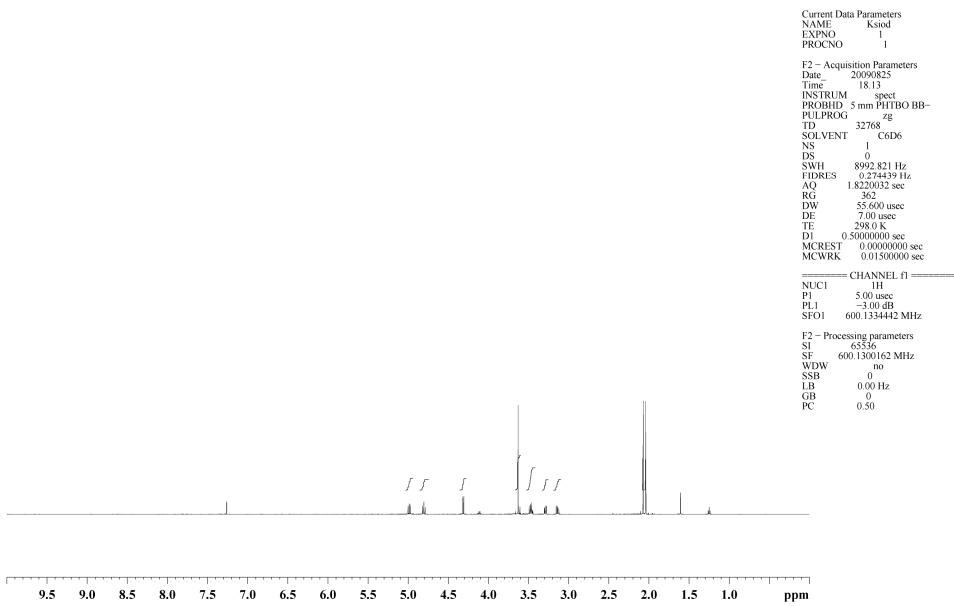
### **The Development of Selective Inhibitors of NagZ: Increased Susceptibility of Gram-Negative Bacteria to $\beta$ -Lactams**

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Tracey M. Gloster,<sup>[c, f]</sup> David J. Vocadlo,<sup>[c]</sup> and Brian L. Mark<sup>\*[b]</sup>

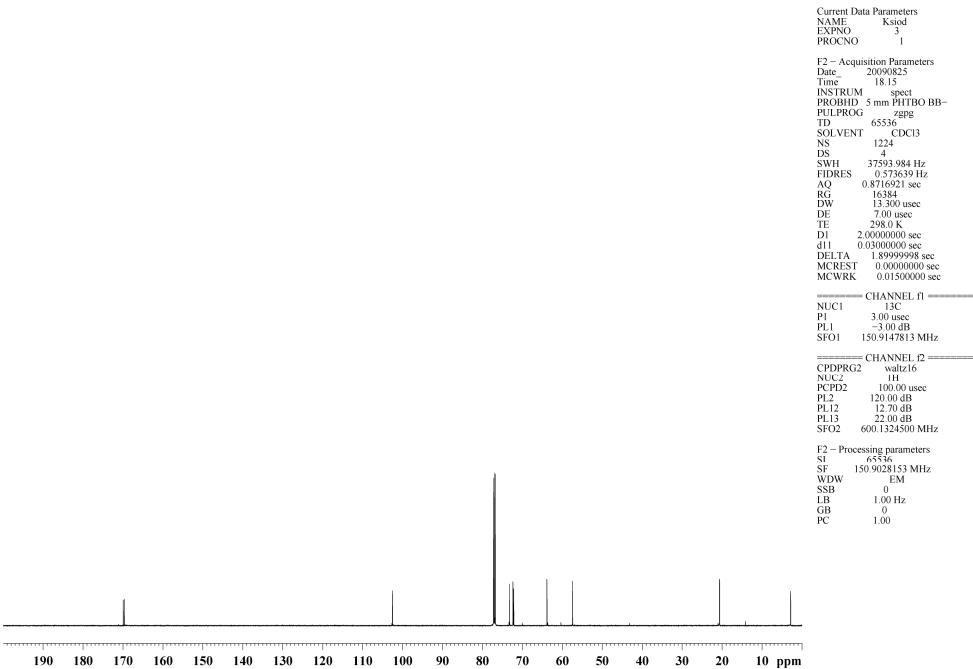
cbic\_201300395\_sm\_miscellaneous\_information.pdf

**Table S1: Crystallographic data and model refinement**

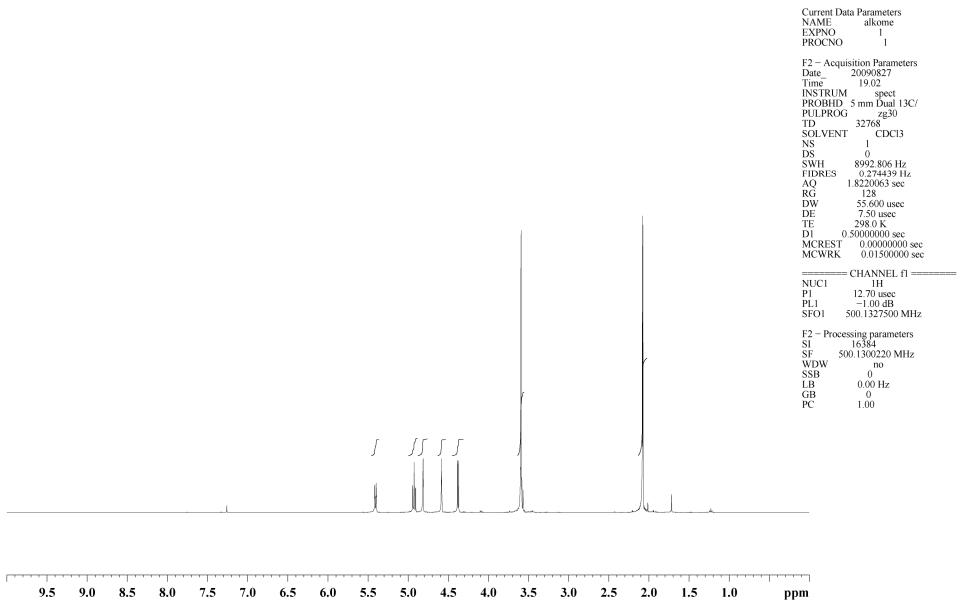
<b><i>StNagZ-21</i> (PDB:4HZM)</b>	
<b>Data Collection<sup>a</sup></b>	
<hr/>	
Space group	P1 2 <sub>1</sub> 1
Unit cell (Å)	a=49.77, b=66.19, c=95.07 $\alpha = 90, \beta = 99.21, \gamma = 90^\circ$
Wavelength (Å)	0.9795
Resolution range (Å)	49.12 – 1.45 (1.53 – 1.45)
Total observations	378412
Total unique observations	105765
I/ $\sigma_I$	13.0 (3.1)
Completeness (%)	98.0 (87.5)
R <sub>merge</sub>	0.065 (0.365)
Redundancy	3.6 (2.5)
<hr/>	
<b>Refinement Statistics</b>	
<hr/>	
Resolution (Å)	49.12 – 1.45
Reflections (total)	105697
Reflections (test)	3160
Total atoms refined	6089
Solvent	920
R <sub>work</sub> (R <sub>free</sub> )	0.169 (0.193)
Mean B values (Å <sup>2</sup> )	14.5
Overall	
Bound ligand (A/B subunit)	7.9/7.9
RMSDs	0.008/1.172
Bond lengths (Å) / angles (°)	
Ramachandran plot	97.1/2.9
Favored/allowed (%)	



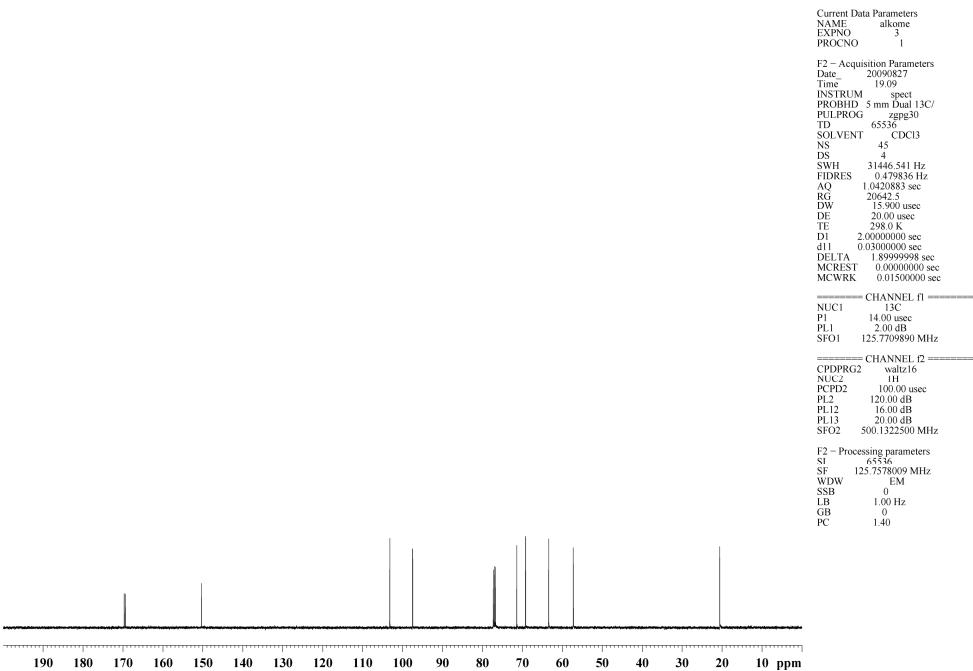
<sup>1</sup>H NMR of **5**.



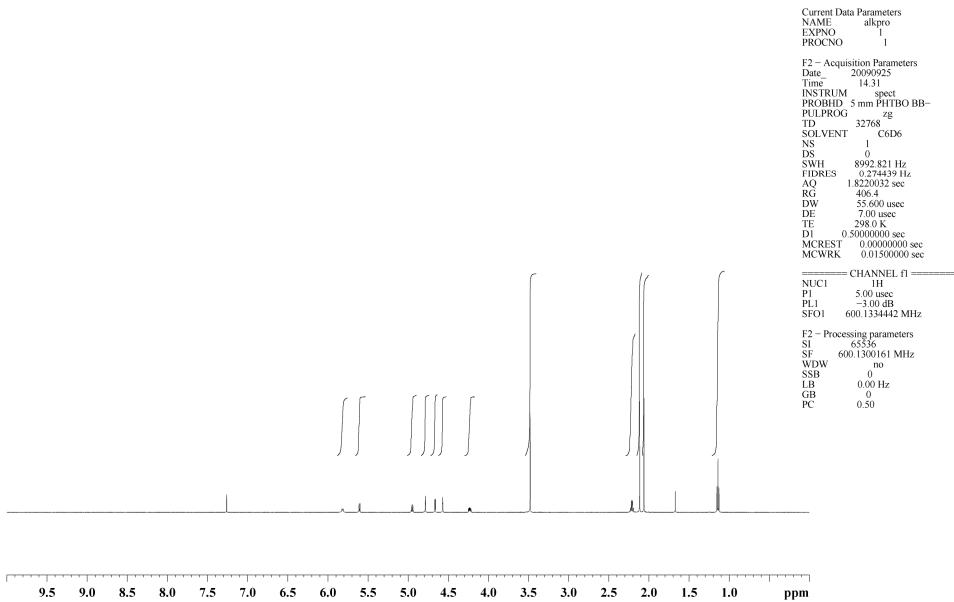
<sup>13</sup>C NMR of **5**.



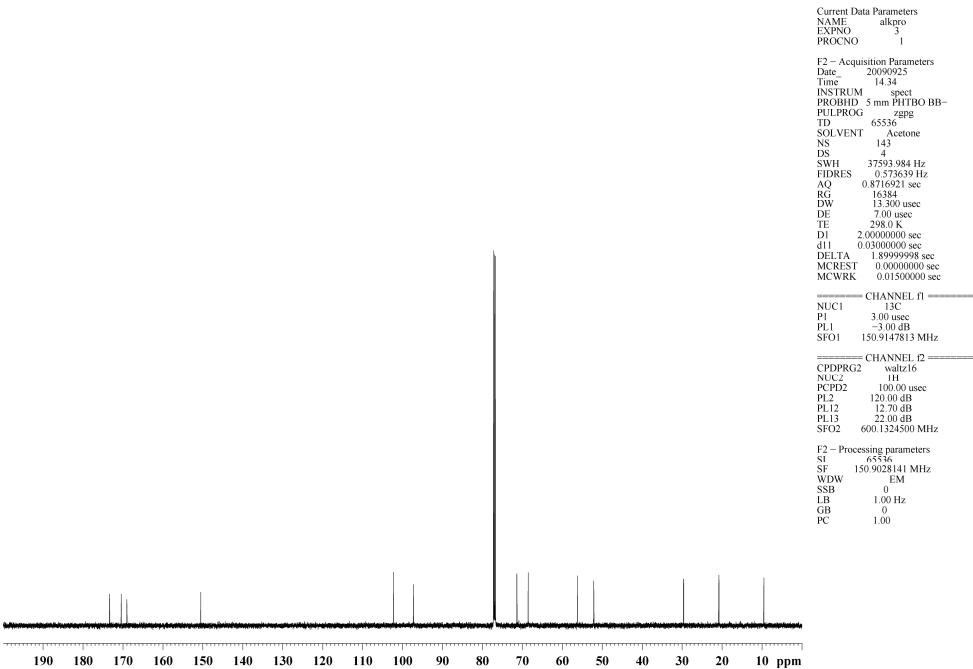
<sup>1</sup>H NMR of **2**.



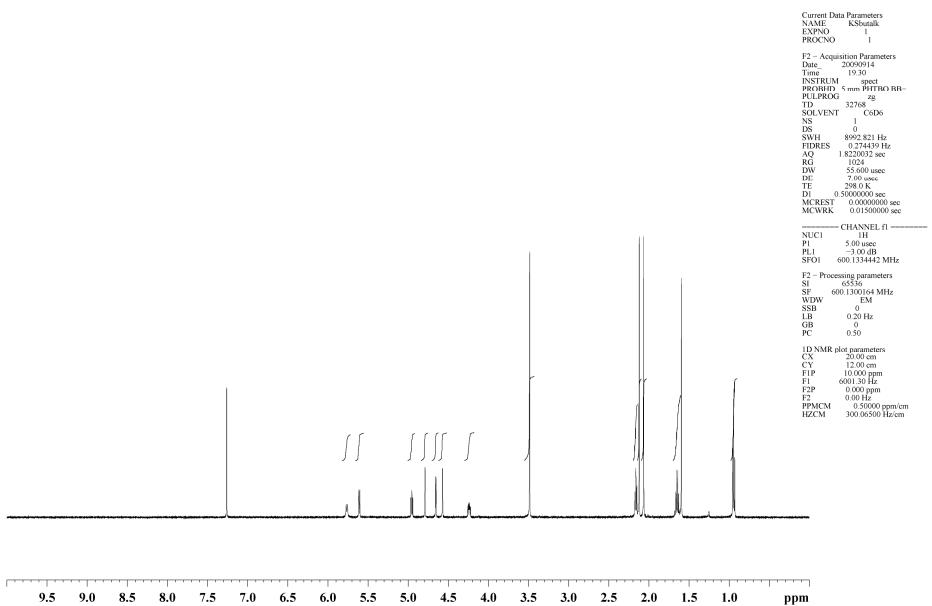
<sup>13</sup>C NMR of **2**.



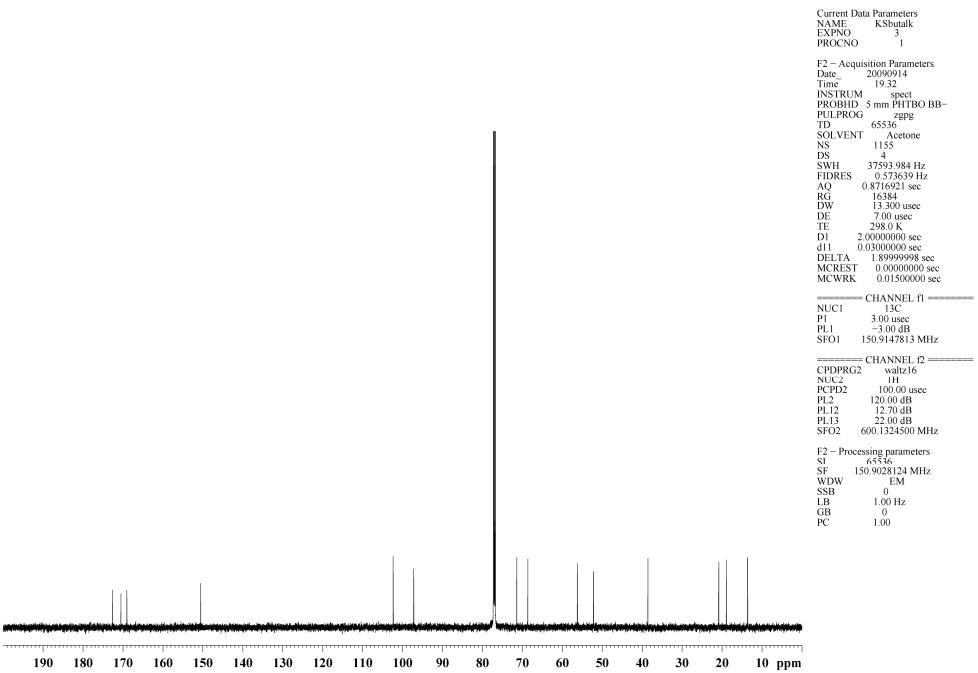
<sup>1</sup>H NMR of **7**.



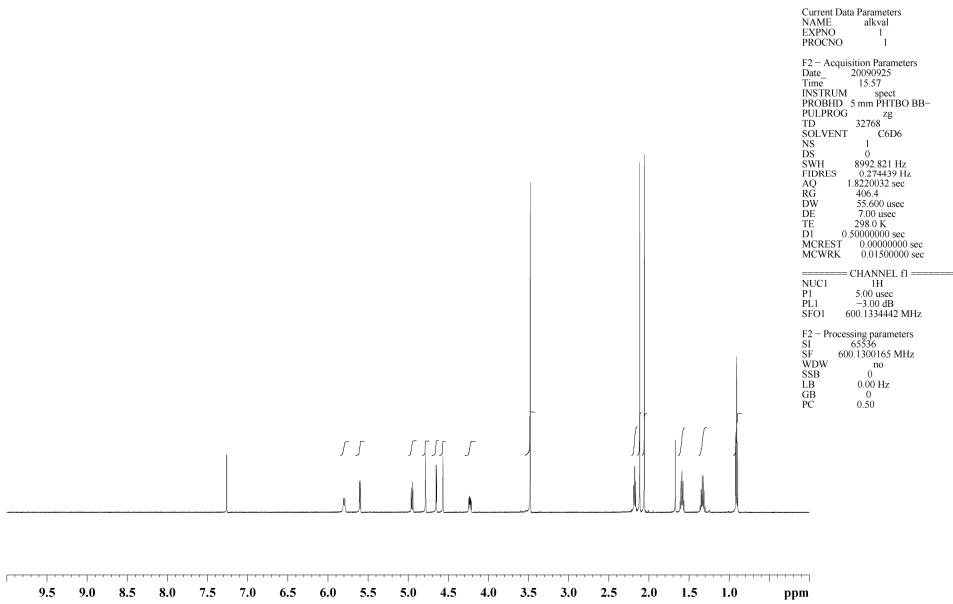
<sup>13</sup>C NMR of **7**.



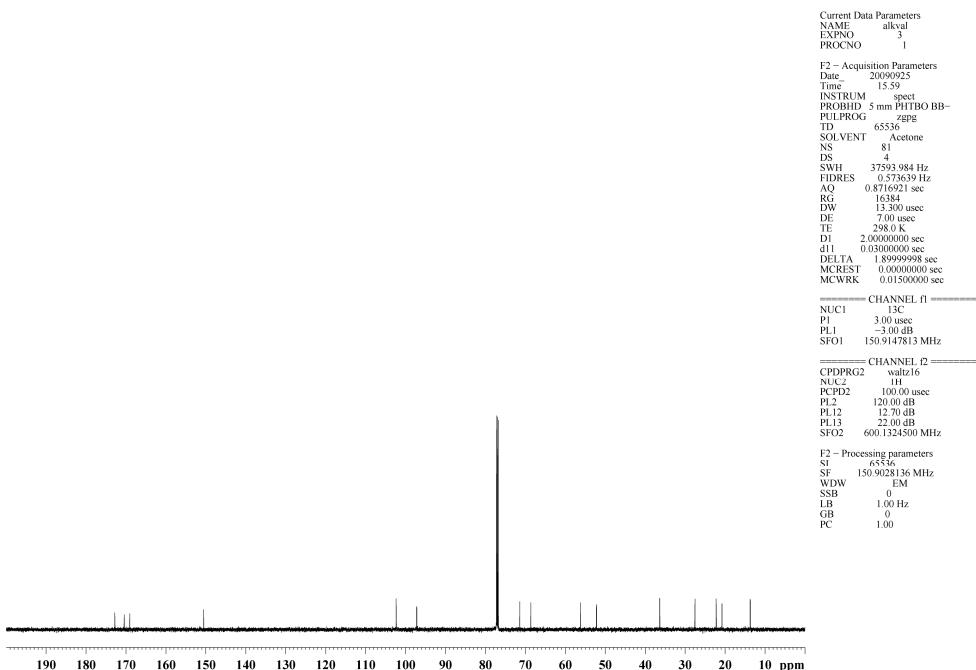
<sup>1</sup>H NMR of **8**.



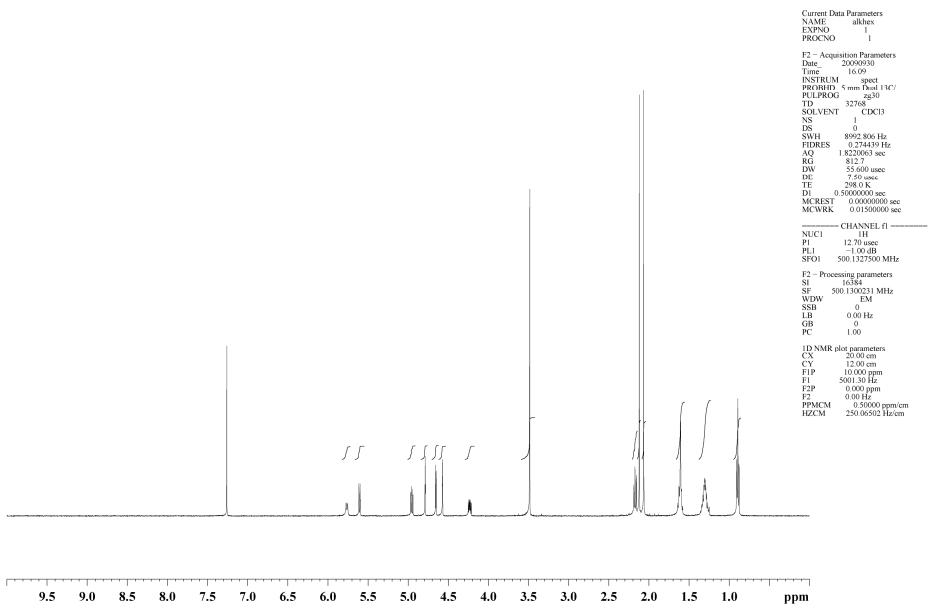
<sup>13</sup>C NMR of **8**.



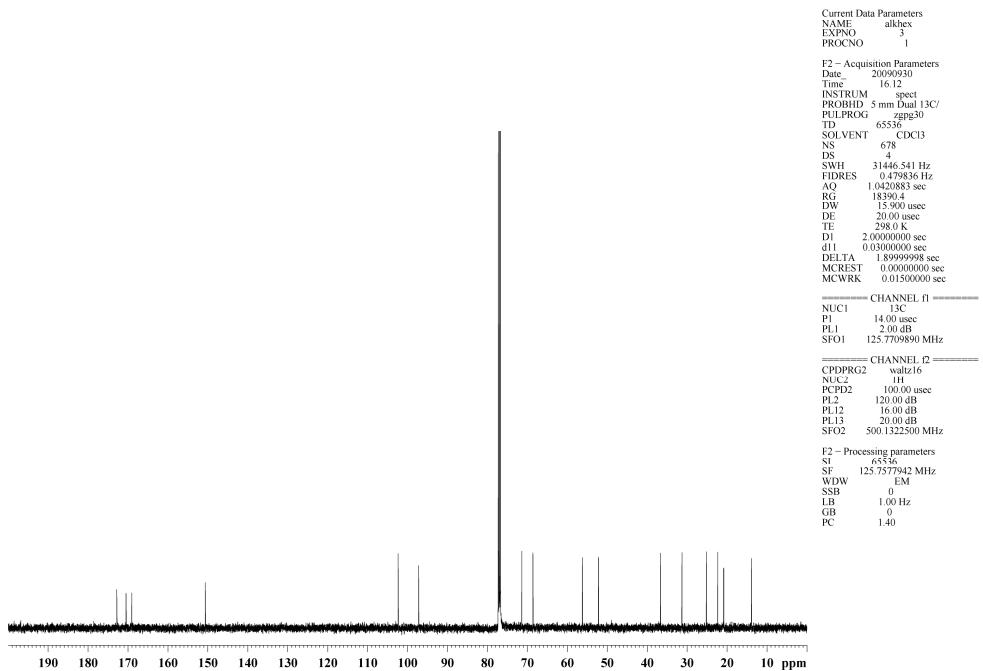
<sup>1</sup>H NMR of **9**.



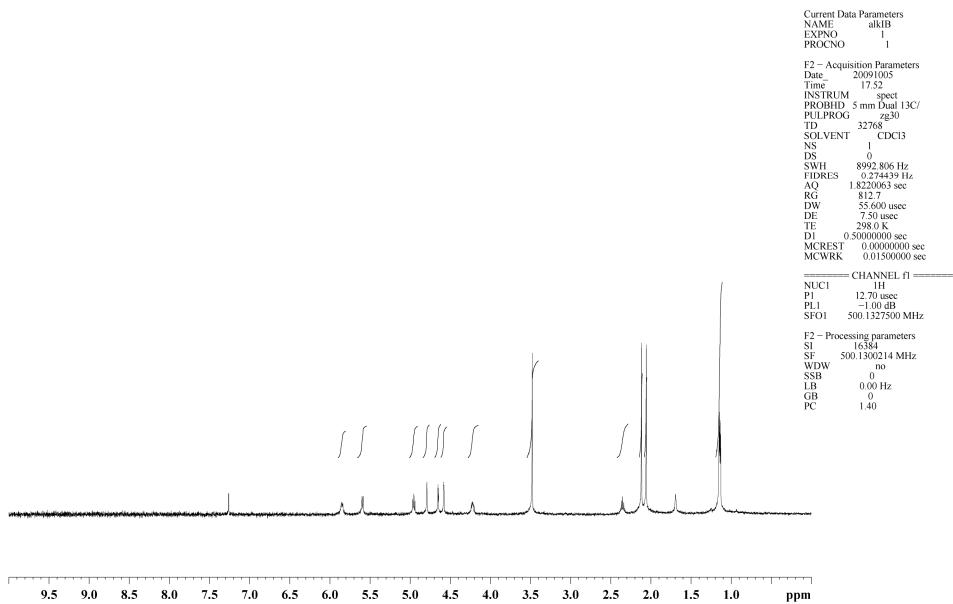
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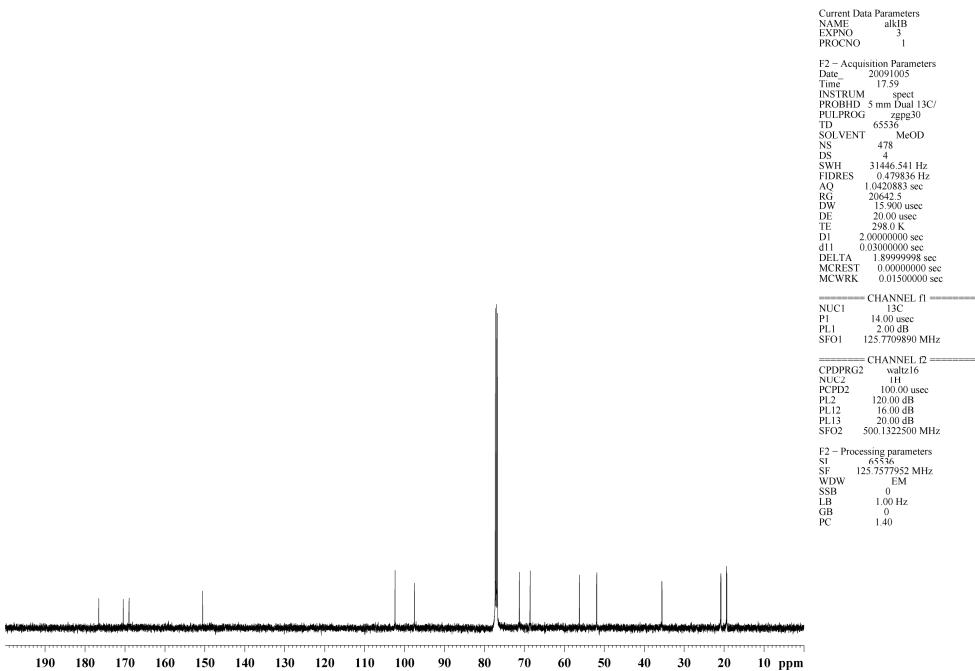
<sup>1</sup>H NMR of **10**.



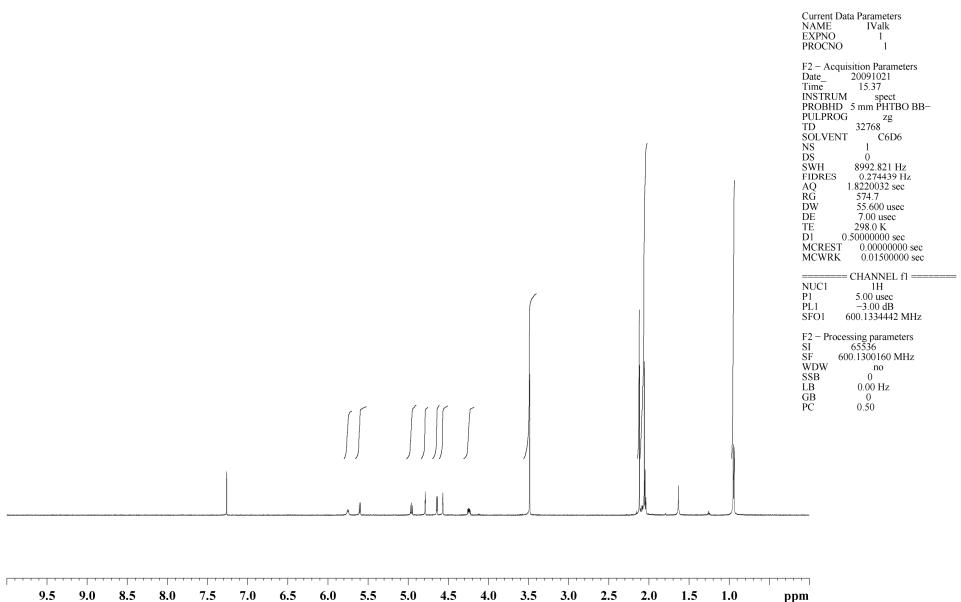
<sup>13</sup>C NMR of **10**.



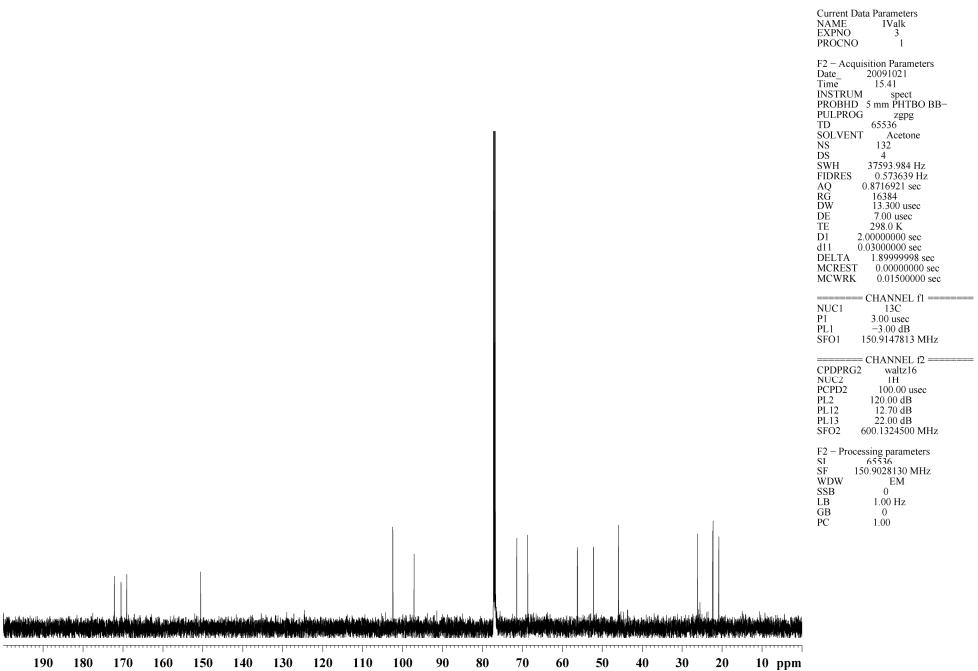
<sup>1</sup>H NMR of **11**.



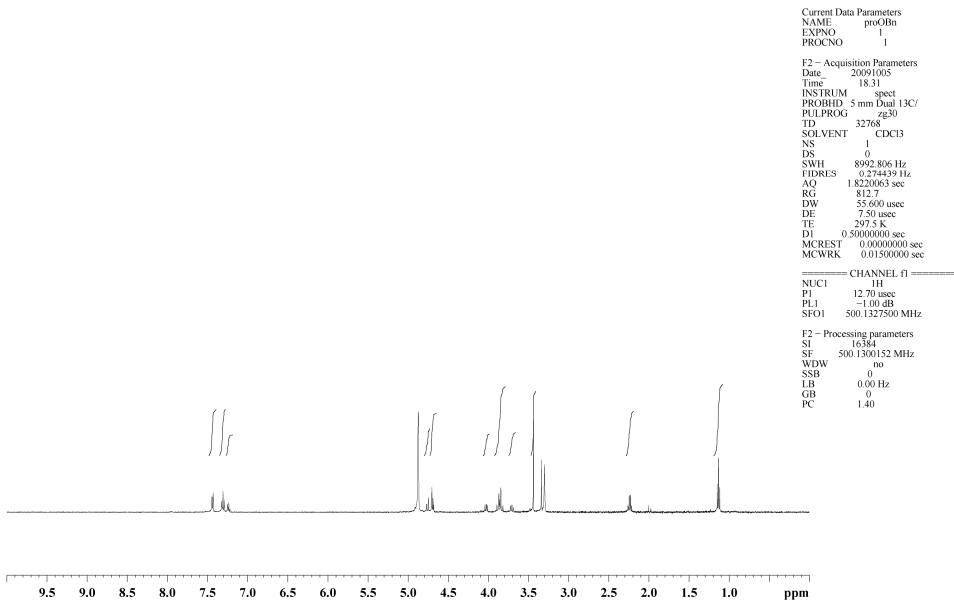
<sup>13</sup>C NMR of **11**.



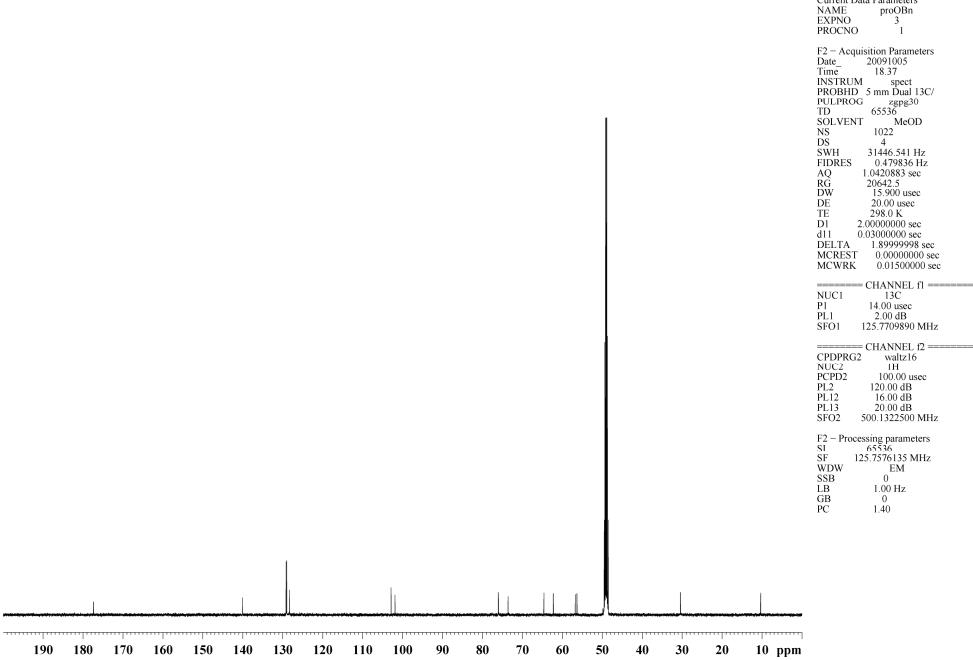
<sup>1</sup>H NMR of **12**.



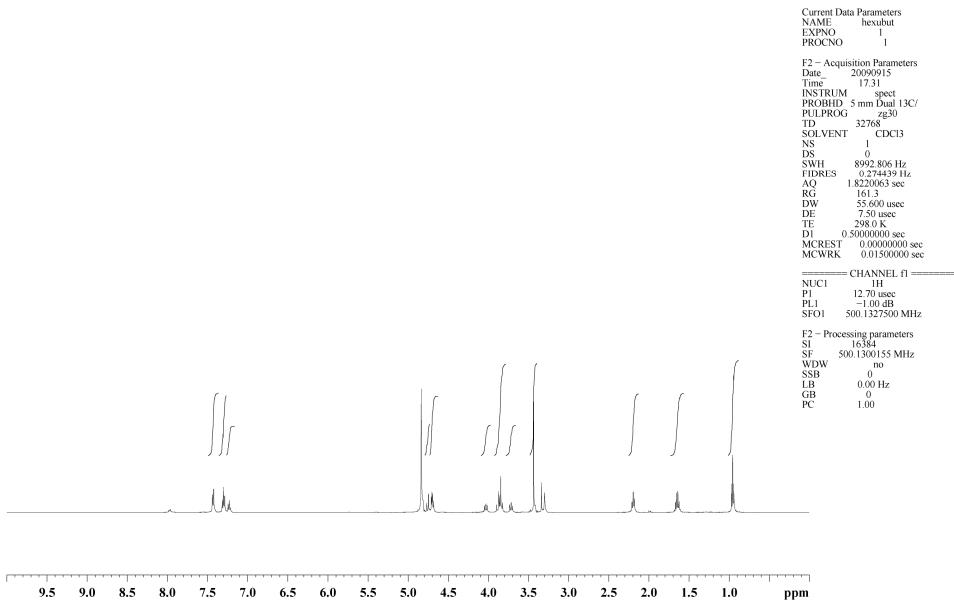
<sup>13</sup>C NMR of **12**.



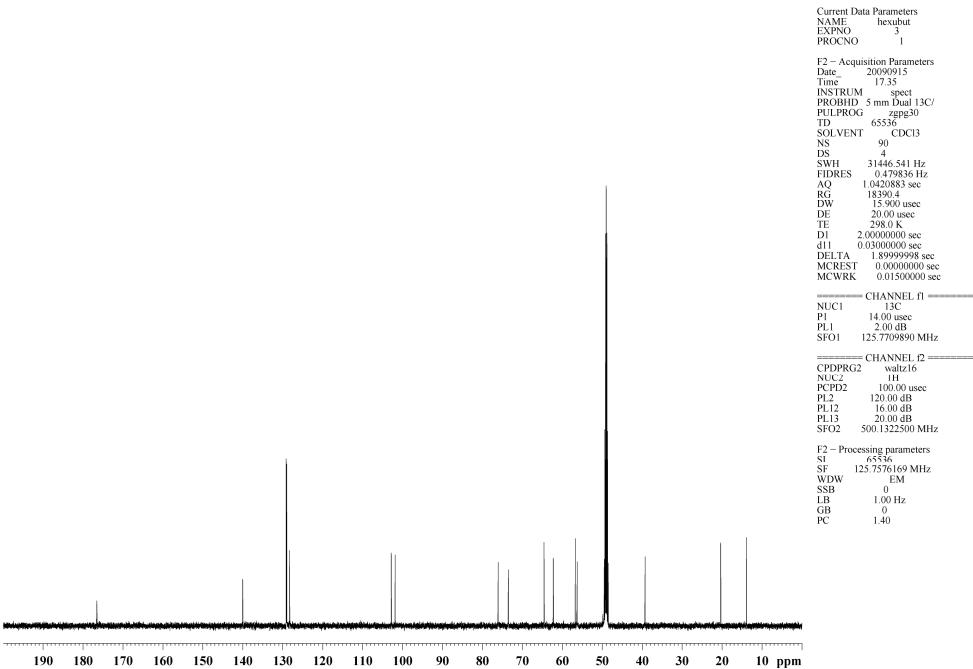
<sup>1</sup>H NMR of **14**.



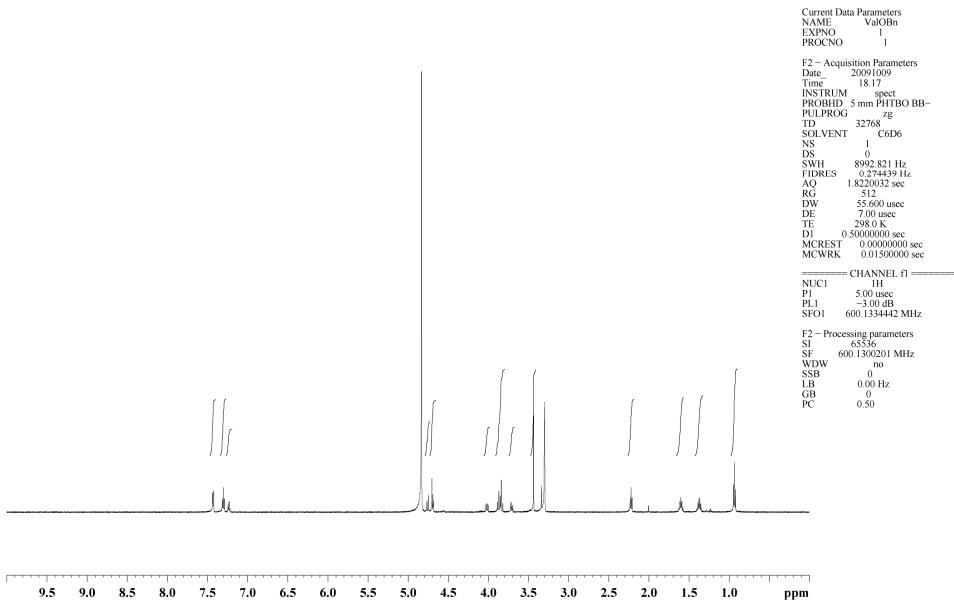
<sup>13</sup>C NMR of **14**.



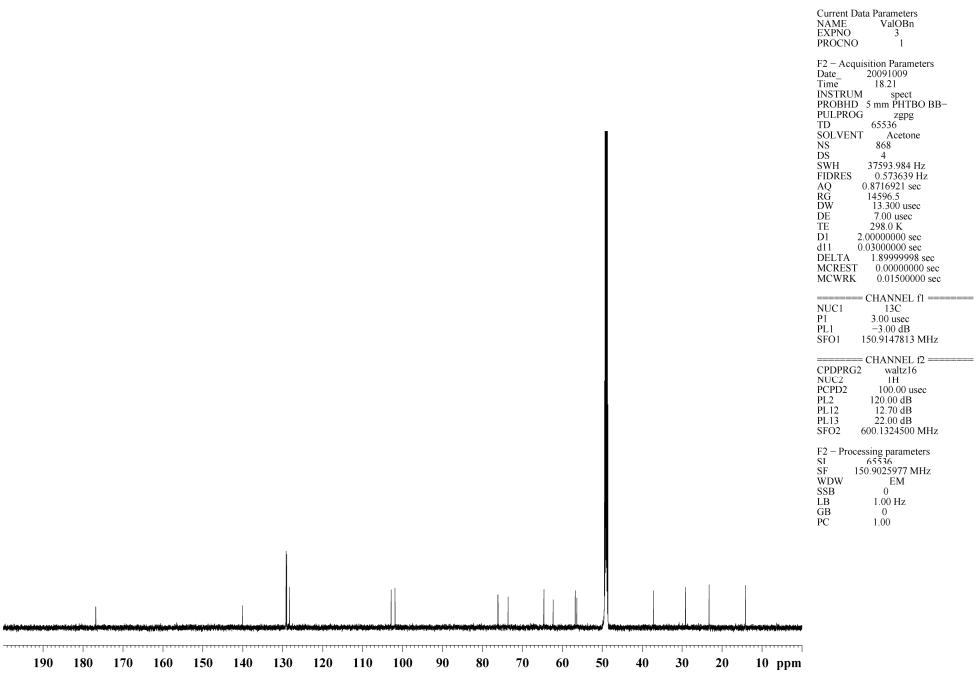
<sup>1</sup>H NMR of **15**.

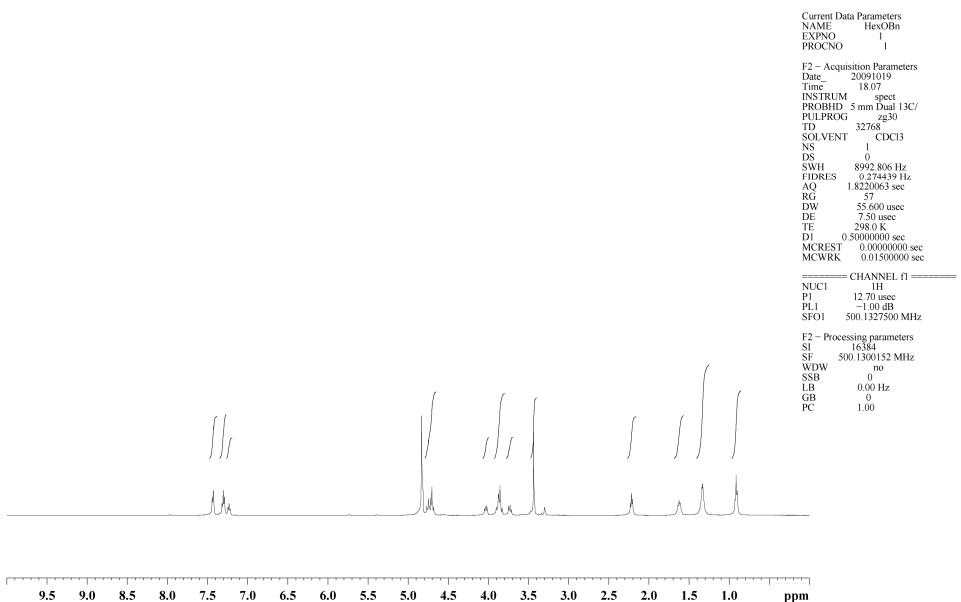


<sup>13</sup>C NMR of **15**.

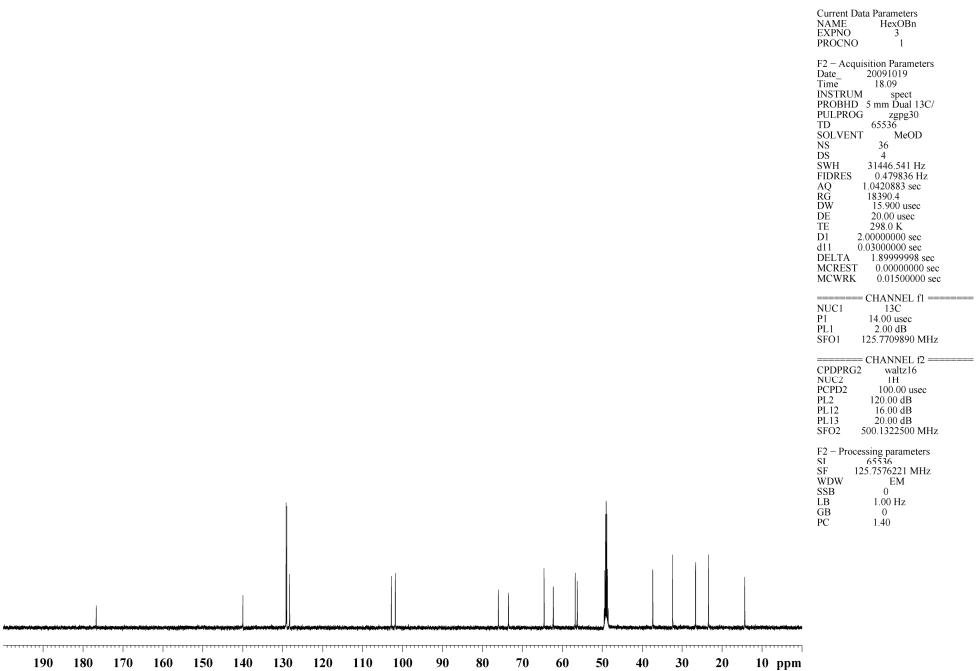


<sup>1</sup>H NMR of **16**.

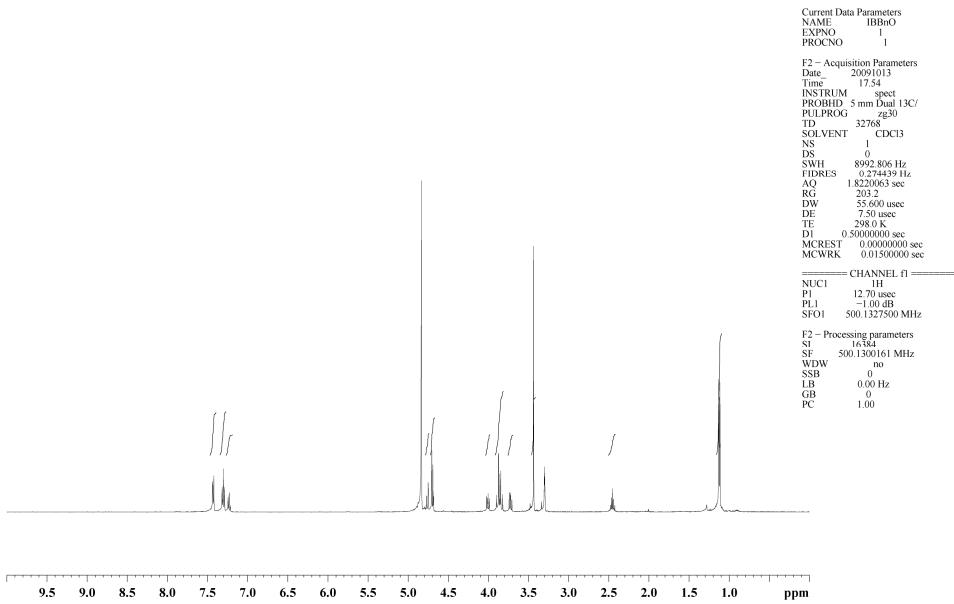




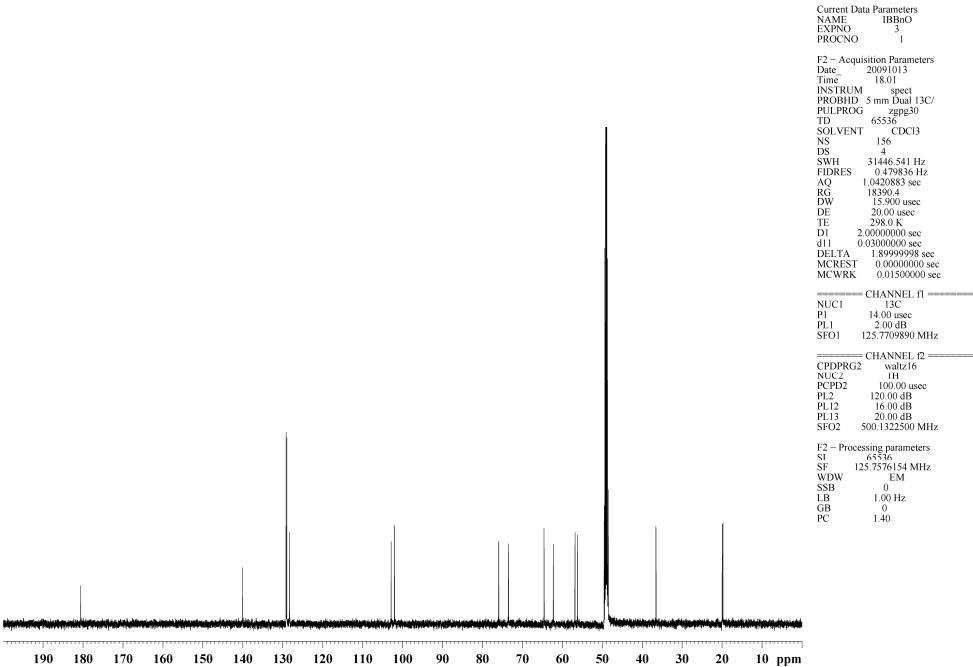
<sup>1</sup>H NMR of **17**.



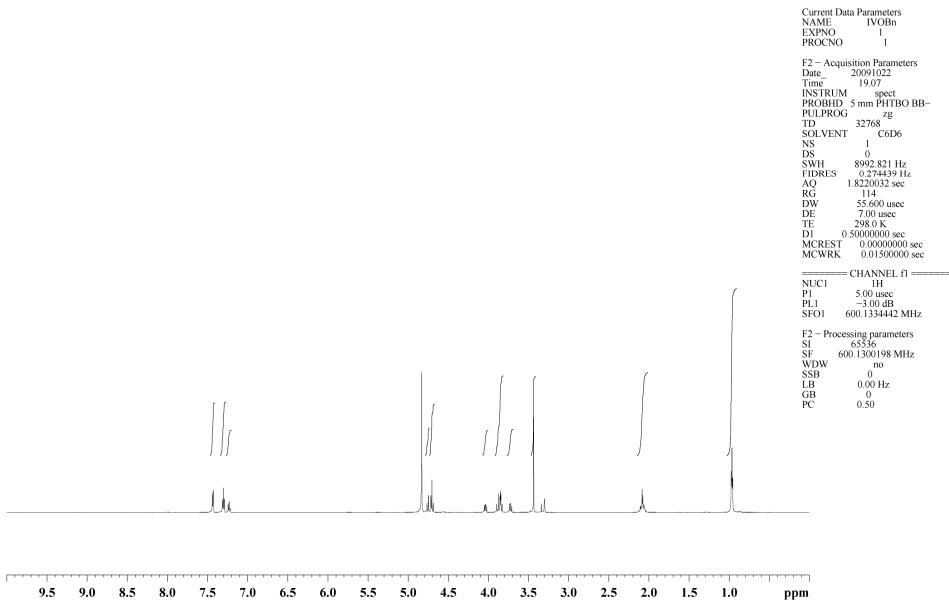
<sup>13</sup>C NMR of **17**.



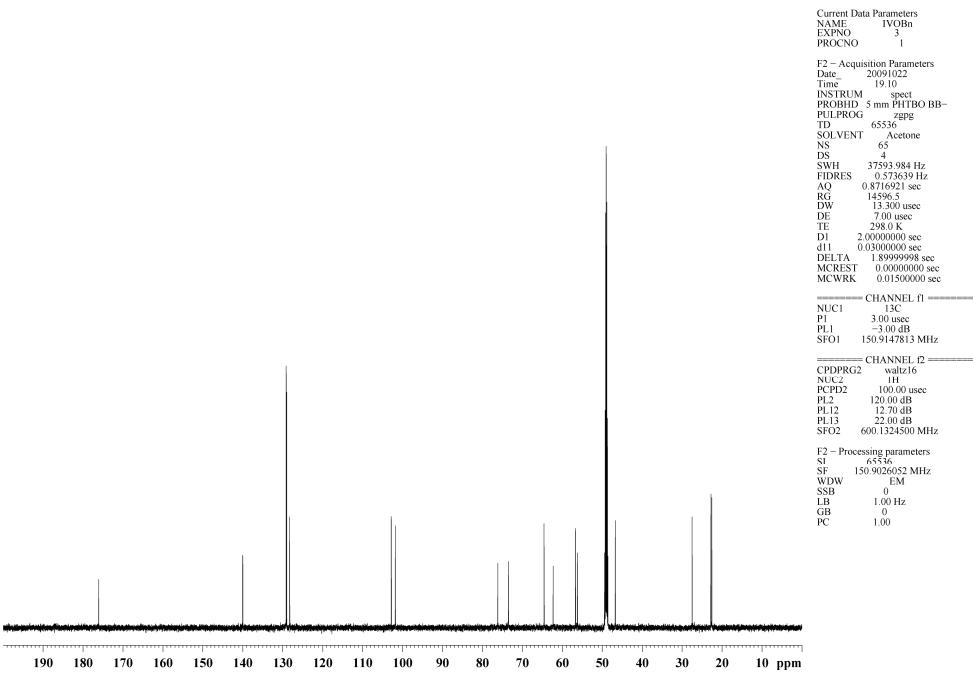
<sup>1</sup>H NMR of **18**.



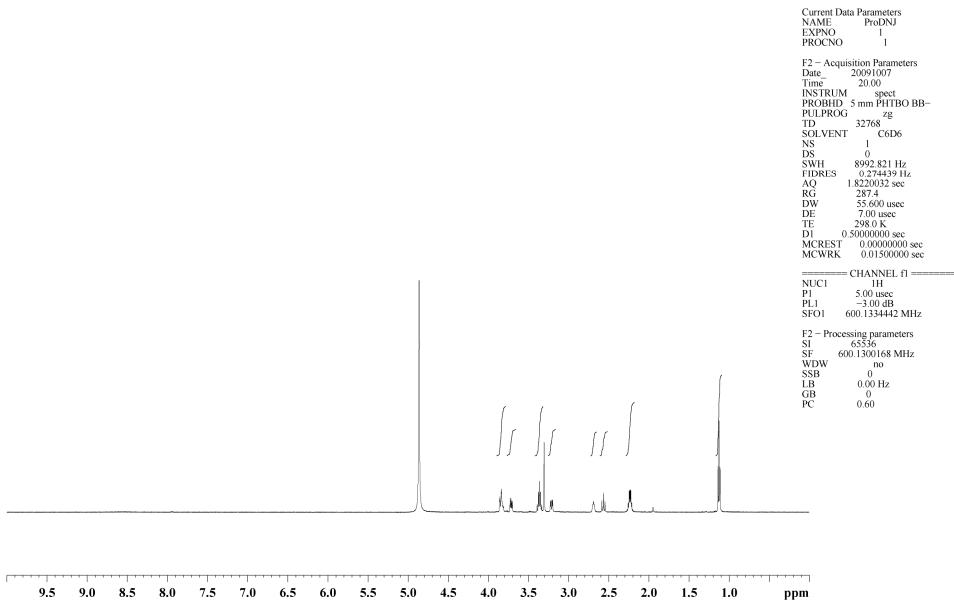
<sup>13</sup>C NMR of **18**.



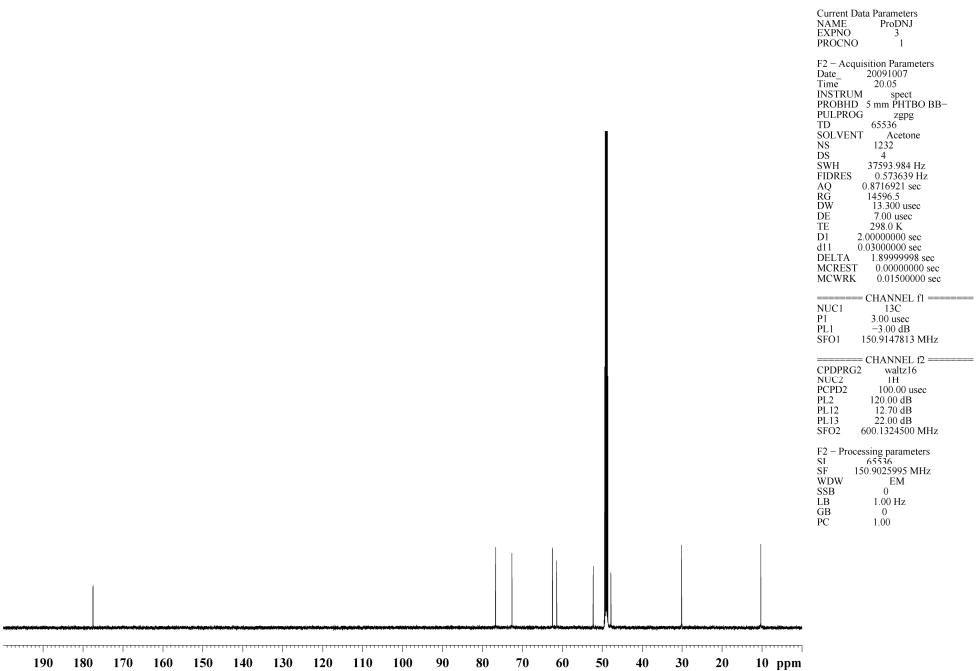
<sup>1</sup>H NMR of **19**.



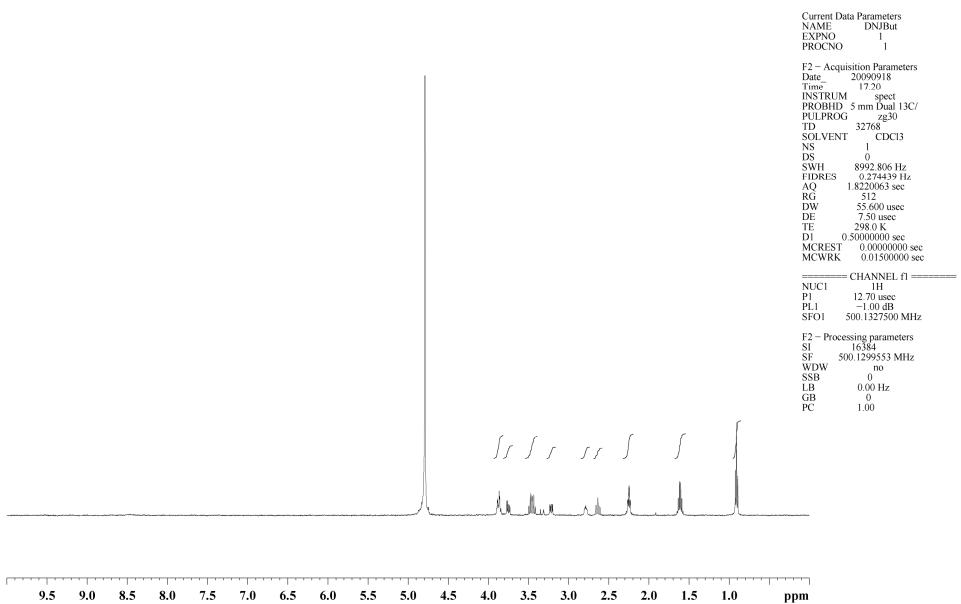
<sup>13</sup>C NMR of **19**.



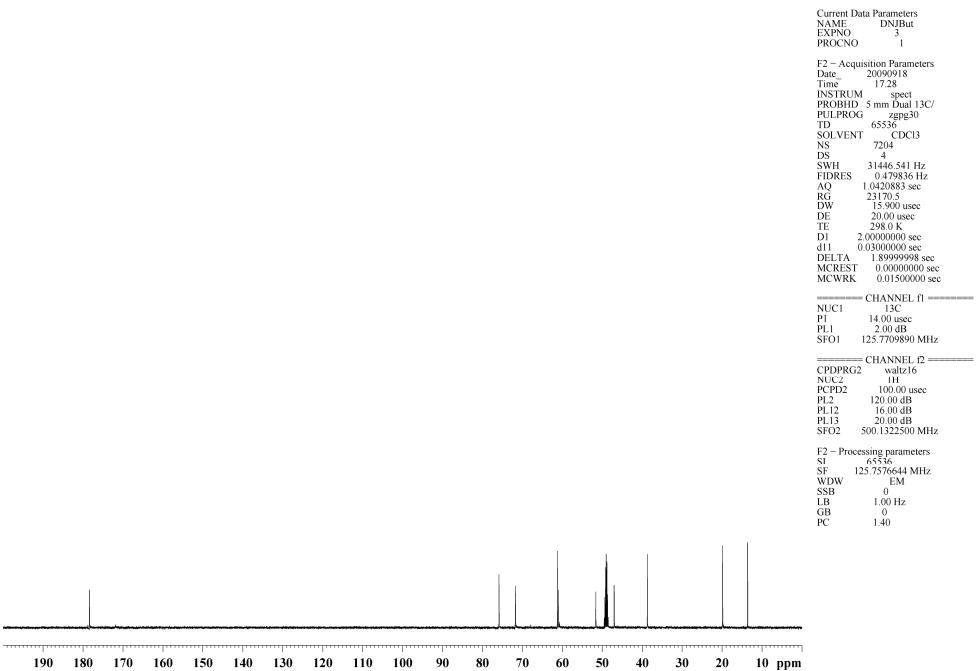
<sup>1</sup>H NMR of **20**.



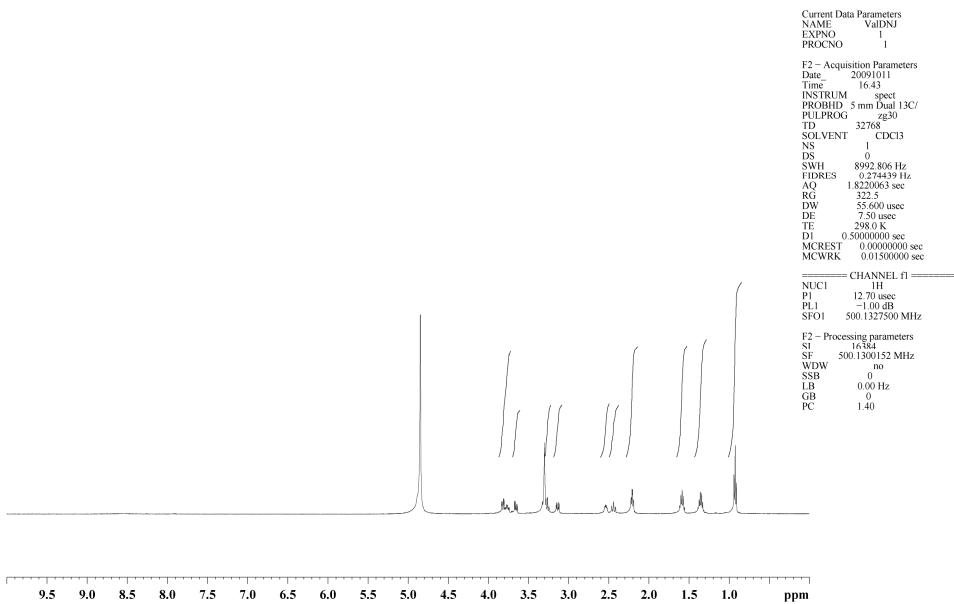
<sup>13</sup>C NMR of **20**.



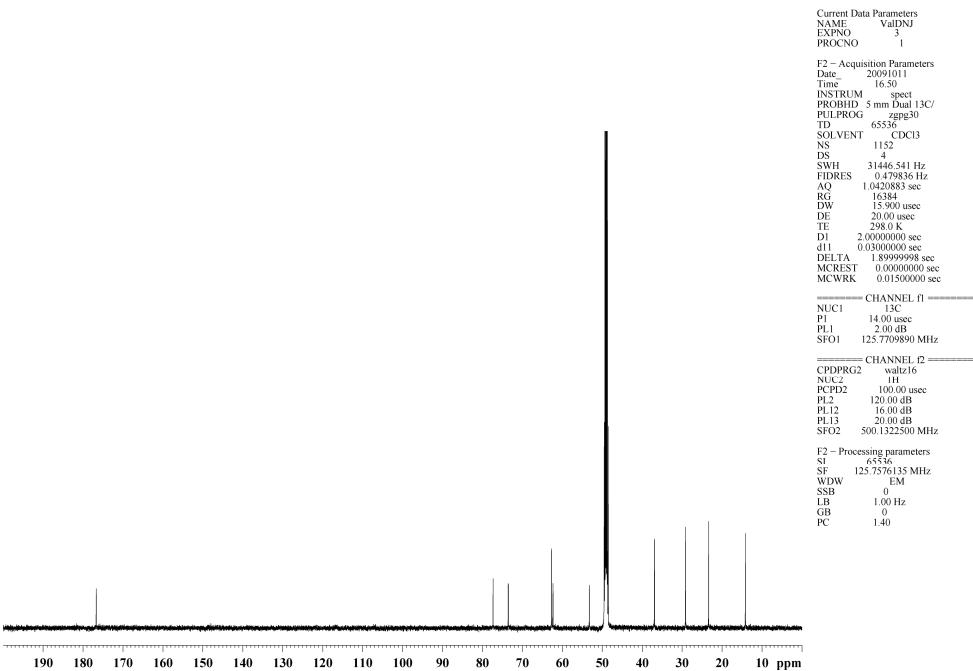
<sup>1</sup>H NMR of **21**.



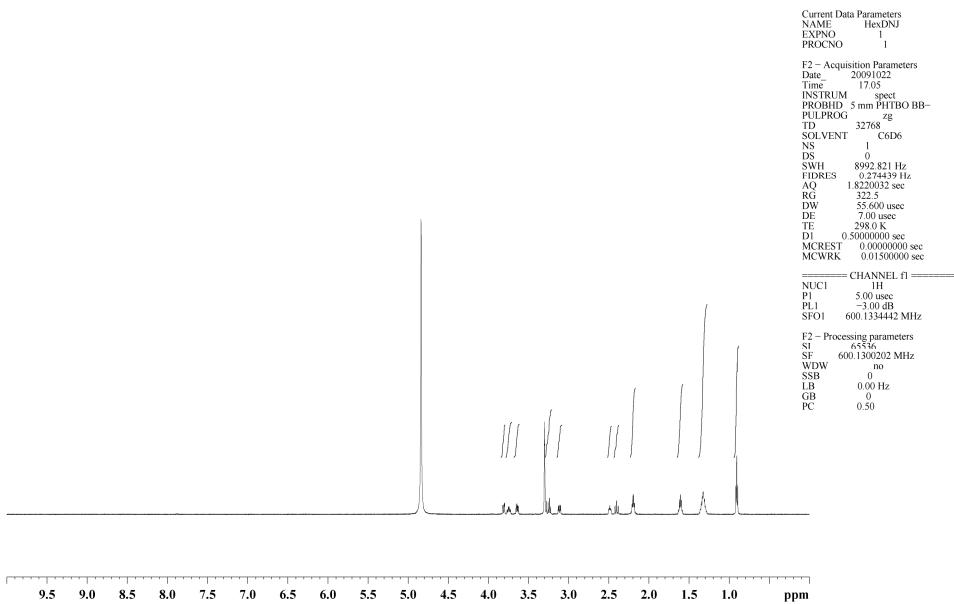
<sup>13</sup>C NMR of **21**.



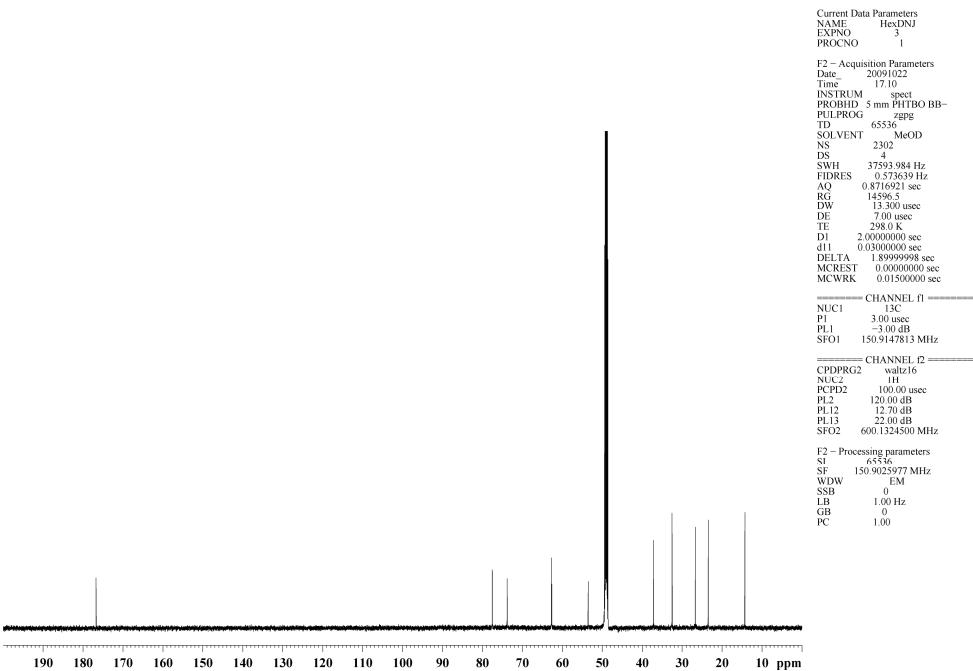
<sup>1</sup>H NMR of **22**.



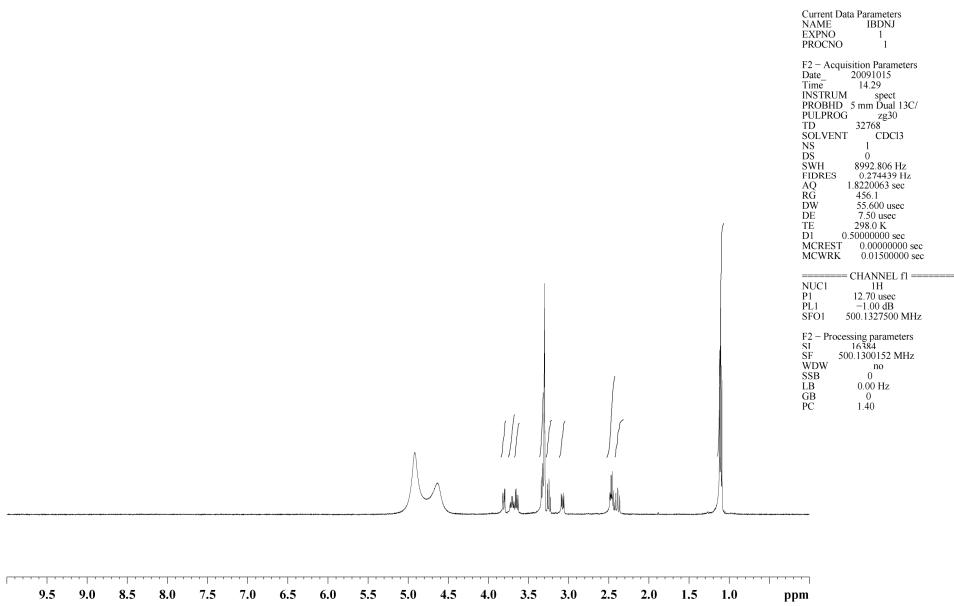
<sup>13</sup>C NMR of **22**.



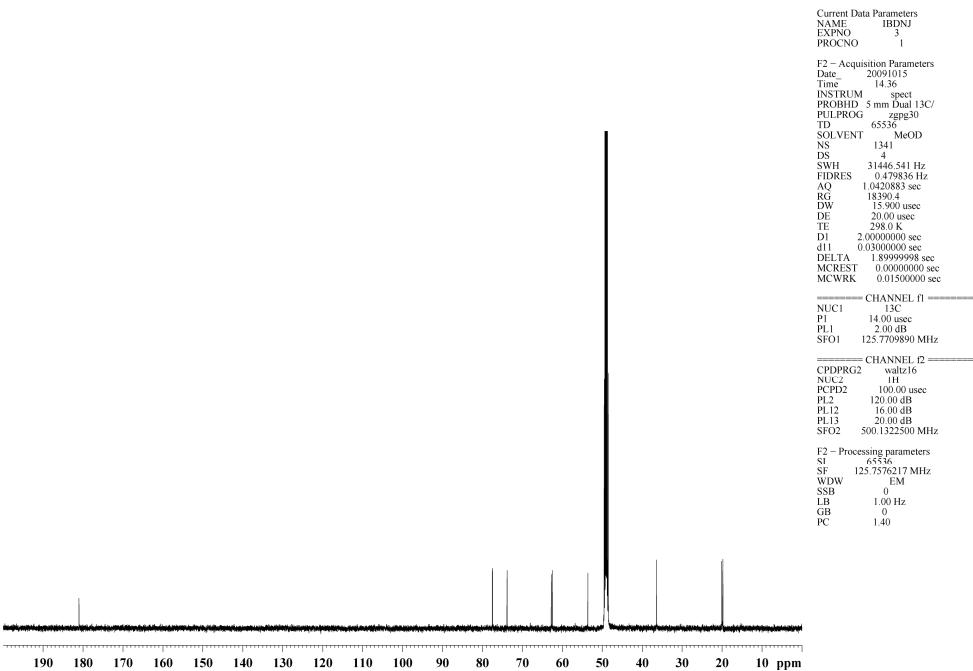
<sup>1</sup>H NMR of **23**.



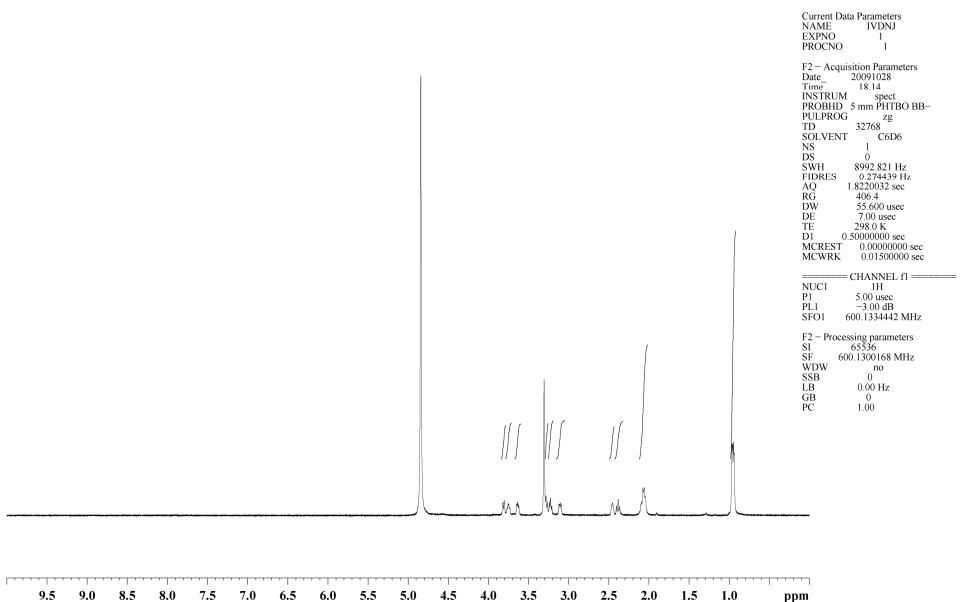
<sup>13</sup>C NMR of **23**.



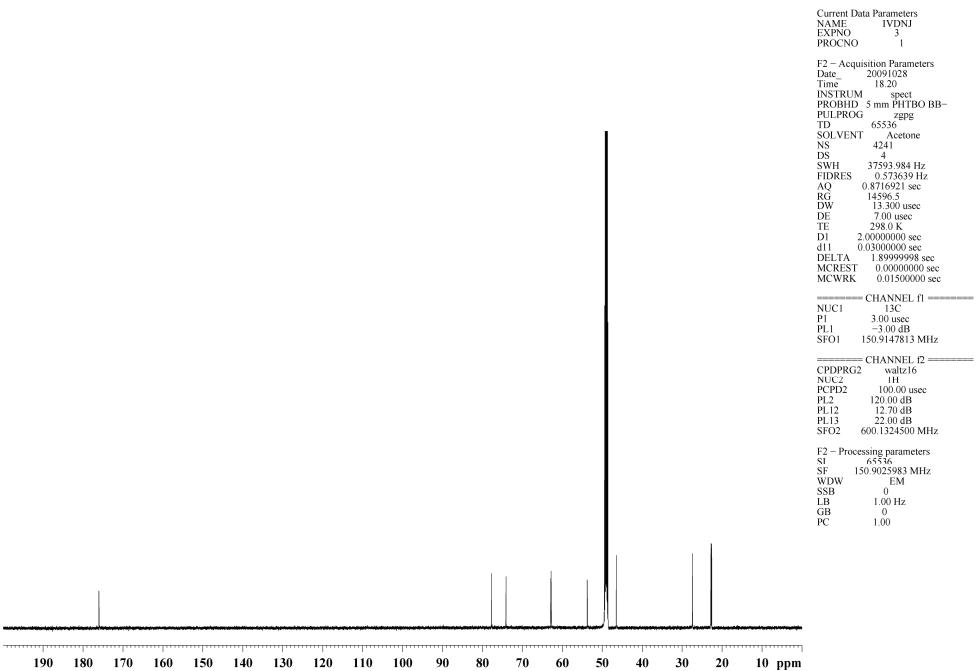
<sup>1</sup>H NMR of **24**.



<sup>13</sup>C NMR of **24**.



<sup>1</sup>H NMR of **25**.



<sup>13</sup>C NMR of **25**.