

ESI for:

## Conformational and Electronic Variability of *N,N',O*-Ligand Documented on its Coordination to Main Group Halides

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**Table S1.** Crystallographic data for **2**

Crystal data	
Chemical formula	C <sub>25</sub> H <sub>37</sub> ClN <sub>2</sub> O <sub>3</sub> PGe·Cl <sub>3</sub> Ge
M <sub>r</sub>	731.51
Crystal system, space group	Triclinic, <i>P</i> -1
Temperature (K)	150
<i>a</i> , <i>b</i> , <i>c</i> (Å)	8.4329(9), 14.0969(13), 14.9495(14)
α, β, γ (°)	70.465(5), 87.024(5), 73.513(5)
<i>V</i> (Å <sup>3</sup> )	1604.2(3)
<i>Z</i>	2
Radiation type	Mo <i>K</i> α
μ (mm <sup>-1</sup> )	2.285
Crystal size (mm)	0.290 × 0.123 × 0.059
Data collection	
Diffractometer	Bruker D8 - Venture
Absorption correction	Multi-scan <i>SADABS2016/2</i> - Bruker AXS area detector scaling and absorption correction
<i>T</i> <sub>min</sub> , <i>T</i> <sub>max</sub>	0.4879, 0.7456
No. of measured, independent and observed [ <i>I</i> > 2σ( <i>I</i> )] reflections	38102, 6290, 4639
<i>R</i> <sub>int</sub>	0.1034
(sin θ/λ) <sub>max</sub> (Å <sup>-1</sup> )	0.596
Refinement	
<i>R</i> [ <i>F</i> <sup>2</sup> > 2σ( <i>F</i> <sup>2</sup> )], <i>wR</i> ( <i>F</i> <sup>2</sup> ), <i>S</i>	0.0652, 0.1580, 1.079
No. of reflections	6290
No. of parameters	362
No. of restraints	292
H-atom treatment	H-atom parameters constrained
Δρ <sub>max</sub> , Δρ <sub>min</sub> (e Å <sup>-3</sup> )	1.386, -0.783

Computer programs: Bruker Instrument Service vV6.2.3, *APEX3* v2016.5-0 (Bruker AXS), *SAINT* V8.37A (Bruker AXS Inc., 2015), *XT*, VERSION 2014/5, *SHELXL2014/7* (Sheldrick, 2014), *PLATON* (Spek, 2009).

**Table S2.** Crystallographic data for **3·0.5(C<sub>6</sub>H<sub>14</sub>)**

Crystal data	
Chemical formula	C <sub>43</sub> H <sub>52</sub> ClN <sub>2</sub> O <sub>3</sub> PSn·0.5(C <sub>6</sub> H <sub>14</sub> )
$M_r$	873.06
Crystal system, space group	Triclinic, <i>P</i> -1
Temperature (K)	150
<i>a</i> , <i>b</i> , <i>c</i> (Å)	9.7555(3), 10.6048(3), 23.1622(7)
$\alpha$ , $\beta$ , $\gamma$ (°)	86.347(2), 81.2130(10), 69.5690(10)
<i>V</i> (Å <sup>3</sup> )	2219.01(12)
<i>Z</i>	2
Radiation type	Mo <i>K</i> α
$\mu$ (mm <sup>-1</sup> )	0.712
Crystal size (mm)	0.593 × 0.246 × 0.060
Data collection	
Diffractometer	Bruker D8 - Venture
Absorption correction	Multi-scan <i>SADABS2016/2</i> - Bruker AXS area detector scaling and absorption correction
$T_{\min}$ , $T_{\max}$	0.6487, 0.7469
No. of measured, independent and observed [ $I > 2\sigma(I)$ ] reflections	77217, 9196, 7779
$R_{\text{int}}$	0.0965
(sin $\theta/\lambda$ ) <sub>max</sub> (Å <sup>-1</sup> )	0.606
Refinement	
$R[F^2 > 2\sigma(F^2)]$ , $wR(F^2)$ , <i>S</i>	0.0460, 0.0850, 1.060
No. of reflections	9196
No. of parameters	493
No. of restraints	441
H-atom treatment	H-atom parameters constrained
$\Delta\rho_{\max}$ , $\Delta\rho_{\min}$ (e Å <sup>-3</sup> )	0.591, -0.986

Computer programs: Bruker Instrument Service vV6.2.3, *APEX3* v2016.5-0 (Bruker AXS), *SAINT* V8.37A (Bruker AXS Inc., 2015), *XT*, VERSION 2014/5, *SHELXL2014/7* (Sheldrick, 2014), *PLATON* (Spek, 2009).

**Table S3.** Crystallographic data for **4**

Crystal data	
Chemical formula	C <sub>37</sub> H <sub>47</sub> Cl <sub>2</sub> N <sub>2</sub> O <sub>3</sub> PSn
M <sub>r</sub>	788.32
Crystal system, space group	Triclinic, <i>P</i> -1
Temperature (K)	150
<i>a</i> , <i>b</i> , <i>c</i> (Å)	8.9582(5), 9.7607(6), 22.8368(14)
α, β, γ (°)	79.246(2), 86.851(3), 75.846(3)
<i>V</i> (Å <sup>3</sup> )	1902.1(2)
<i>Z</i>	2
Radiation type	Mo <i>K</i> α
μ (mm <sup>-1</sup> )	0.890
Crystal size (mm)	0.591 × 0.170 × 0.156
Data collection	
Diffractometer	Bruker D8 - Venture
Absorption correction	Multi-scan <i>SADABS2016/2</i> - Bruker AXS area detector scaling and absorption correction
<i>T</i> <sub>min</sub> , <i>T</i> <sub>max</sub>	0.5753, 0.7456
No. of measured, independent and observed [ <i>I</i> > 2σ( <i>I</i> )] reflections	78519, 8746, 8147
<i>R</i> <sub>int</sub>	0.0421
(sin θ/λ) <sub>max</sub> (Å <sup>-1</sup> )	0.627
Refinement	
<i>R</i> [ <i>F</i> <sup>2</sup> > 2σ( <i>F</i> <sup>2</sup> )], <i>wR</i> ( <i>F</i> <sup>2</sup> ), <i>S</i>	0.0295, 0.0662, 1.112
No. of reflections	8746
No. of parameters	424
No. of restraints	366
H-atom treatment	H-atom parameters constrained
Δρ <sub>max</sub> , Δρ <sub>min</sub> (e Å <sup>-3</sup> )	0.661, -0.872

Computer programs: Bruker Instrument Service vV6.2.3, *APEX3* v2016.5-0 (Bruker AXS), *SAINT* V8.37A (Bruker AXS Inc., 2015), *XT*, VERSION 2014/5, *SHELXL2014/7* (Sheldrick, 2014), *PLATON* (Spek, 2009).

**Table S4.** Crystallographic data for **5**·CH<sub>2</sub>Cl<sub>2</sub>

Crystal data	
Chemical formula	C <sub>25</sub> H <sub>37</sub> Cl <sub>3</sub> InN <sub>2</sub> O <sub>3</sub> P·CH <sub>2</sub> Cl <sub>2</sub>
$M_r$	750.63
Crystal system, space group	Monoclinic, <i>P2<sub>1</sub>/n</i>
Temperature (K)	150
<i>a</i> , <i>b</i> , <i>c</i> (Å)	10.0275(4), 24.3548(9), 14.0572(6)
$\alpha$ , $\beta$ , $\gamma$ (°)	90, 100.5700(10), 90
<i>V</i> (Å <sup>3</sup> )	3374.8(2)
<i>Z</i>	4
Radiation type	Mo $K\alpha$
$\mu$ (mm <sup>-1</sup> )	1.172
Crystal size (mm)	0.592 × 0.200 × 0.052
Data collection	
Diffractometer	Bruker D8 - Venture
Absorption correction	Multi-scan <i>SADABS2016/2</i> - Bruker AXS area detector scaling and absorption correction
$T_{\min}$ , $T_{\max}$	0.6529, 0.7456
No. of measured, independent and observed [ $I > 2\sigma(I)$ ] reflections	96027, 7772, 5822
$R_{\text{int}}$	0.1476
(sin $\theta/\lambda$ ) <sub>max</sub> (Å <sup>-1</sup> )	0.626
Refinement	
$R[F^2 > 2\sigma(F^2)]$ , $wR(F^2)$ , <i>S</i>	0.0591, 0.1125, 1.058
No. of reflections	7772
No. of parameters	338
No. of restraints	375
H-atom treatment	H-atom parameters constrained
$\Delta\rho_{\max}$ , $\Delta\rho_{\min}$ (e Å <sup>-3</sup> )	1.230, -1.475

Computer programs: Bruker Instrument Service vV6.2.3, *APEX3* v2016.5-0 (Bruker AXS), *SAINT* V8.37A (Bruker AXS Inc., 2015), *XT*, VERSION 2014/5, *SHELXL2014/7* (Sheldrick, 2014), *PLATON* (Spek, 2009).

**Table S5.** Crystallographic data for **7**

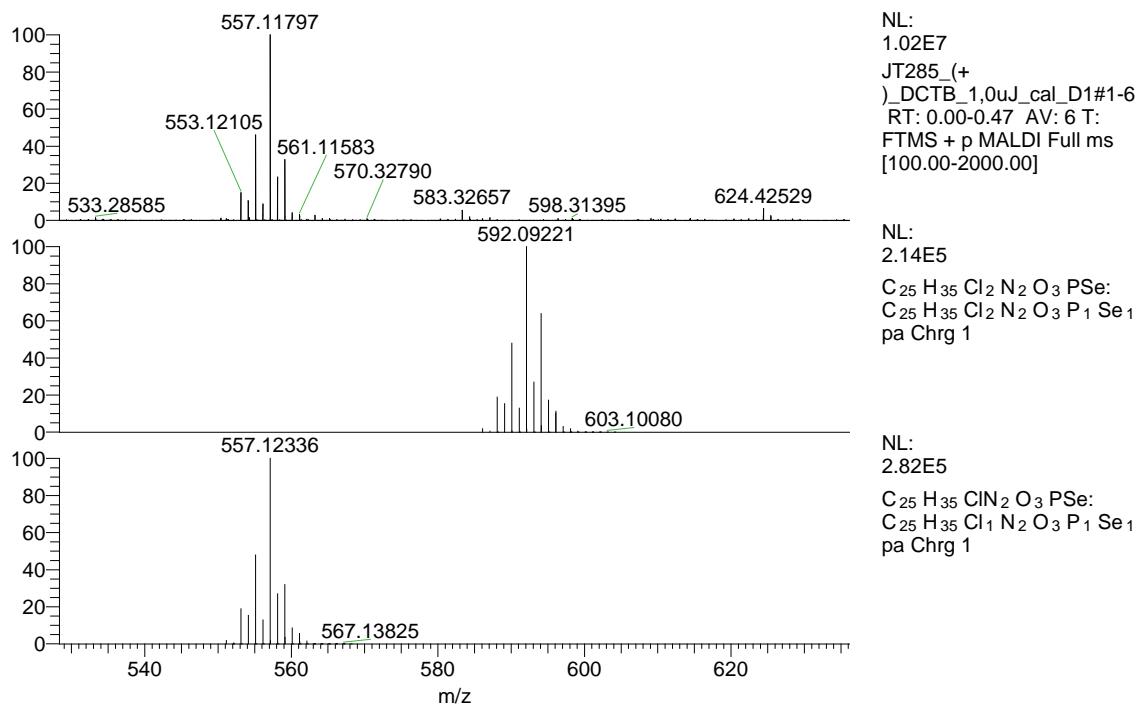
Crystal data	
Chemical formula	C <sub>25</sub> H <sub>36</sub> Cl <sub>3</sub> N <sub>2</sub> O <sub>3</sub> PTe·C <sub>6</sub> H <sub>14</sub>
M <sub>r</sub>	763.65
Crystal system, space group	Orthorhombic, P2 <sub>1</sub> 2 <sub>1</sub> 2 <sub>1</sub>
Temperature (K)	150
a, b, c (Å)	10.6757(4), 15.4741(6), 21.0949(7)
α, β, γ (°)	90, 90, 90
V (Å <sup>3</sup> )	3484.8(2)
Z	4
Radiation type	Mo Kα
μ (mm <sup>-1</sup> )	1.163
Crystal size (mm)	0.495 × 0.224 × 0.142
Data collection	
Diffractometer	Bruker D8 - Venture
Absorption correction	Multi-scan SADABS2016/2 - Bruker AXS area detector scaling and absorption correction
T <sub>min</sub> , T <sub>max</sub>	0.6035, 0.7464
No. of measured, independent and observed [I > 2σ(I)] reflections	33790, 10325, 8671
R <sub>int</sub>	0.0431
(sin θ/λ) <sub>max</sub> (Å <sup>-1</sup> )	0.713
Refinement	
R[F <sup>2</sup> > 2σ(F <sup>2</sup> )], wR(F <sup>2</sup> ), S	0.0403, 0.0720, 1.046
No. of reflections	10325
No. of parameters	324
No. of restraints	288
H-atom treatment	H-atom parameters constrained
Δρ <sub>max</sub> , Δρ <sub>min</sub> (e Å <sup>-3</sup> )	1.035, -0.913

Computer programs: Bruker Instrument Service vV6.2.3, APEX3 v2016.5-0 (Bruker AXS), SAINT V8.37A (Bruker AXS Inc., 2015), XT, VERSION 2014/5, SHELLXL2014/7 (Sheldrick, 2014), PLATON (Spek, 2009).

**Table S6.** Crystallographic data for **8**

Crystal data	
Chemical formula	C <sub>25</sub> H <sub>37</sub> GeN <sub>2</sub> O <sub>3</sub> P
M <sub>r</sub>	517.12
Crystal system, space group	Triclinic, <i>P</i> -1
Temperature (K)	150
<i>a</i> , <i>b</i> , <i>c</i> (Å)	7.9728(13), 11.803(2), 15.182(3)
α, β, γ (°)	80.580(9), 83.224(9), 71.301(8)
<i>V</i> (Å <sup>3</sup> )	1331.8(4)
<i>Z</i>	2
Radiation type	Mo <i>K</i> α
μ (mm <sup>-1</sup> )	1.237
Crystal size (mm)	0.371 × 0.353 × 0.174
Data collection	
Diffractometer	Bruker D8 - Venture
Absorption correction	Multi-scan <i>SADABS2016/2</i> - Bruker AXS area detector scaling and absorption correction
<i>T</i> <sub>min</sub> , <i>T</i> <sub>max</sub>	0.5384, 0.7456
No. of measured, independent and observed [ <i>I</i> > 2σ( <i>I</i> )] reflections	25989, 5460, 3990
<i>R</i> <sub>int</sub>	0.1175
(sin θ/λ) <sub>max</sub> (Å <sup>-1</sup> )	0.606
Refinement	
<i>R</i> [ <i>F</i> <sup>2</sup> > 2σ( <i>F</i> <sup>2</sup> )], <i>wR</i> ( <i>F</i> <sup>2</sup> ), <i>S</i>	0.1079, 0.2519, 1.114
No. of reflections	5460
No. of parameters	317
No. of restraints	264
H-atom treatment	H-atom parameters constrained
Δρ <sub>max</sub> , Δρ <sub>min</sub> (e Å <sup>-3</sup> )	1.563, -1.207

Computer programs: Bruker Instrument Service vV6.2.3, *APEX3* v2016.5-0 (Bruker AXS), *SAINT* V8.37A (Bruker AXS Inc., 2015), *XT, VERSION 2014/5*, *SHELXL2014/7* (Sheldrick, 2014), *PLATON* (Spek, 2009).



**Figure S1.** FTMS+ MALDI MS spectrum of **6**

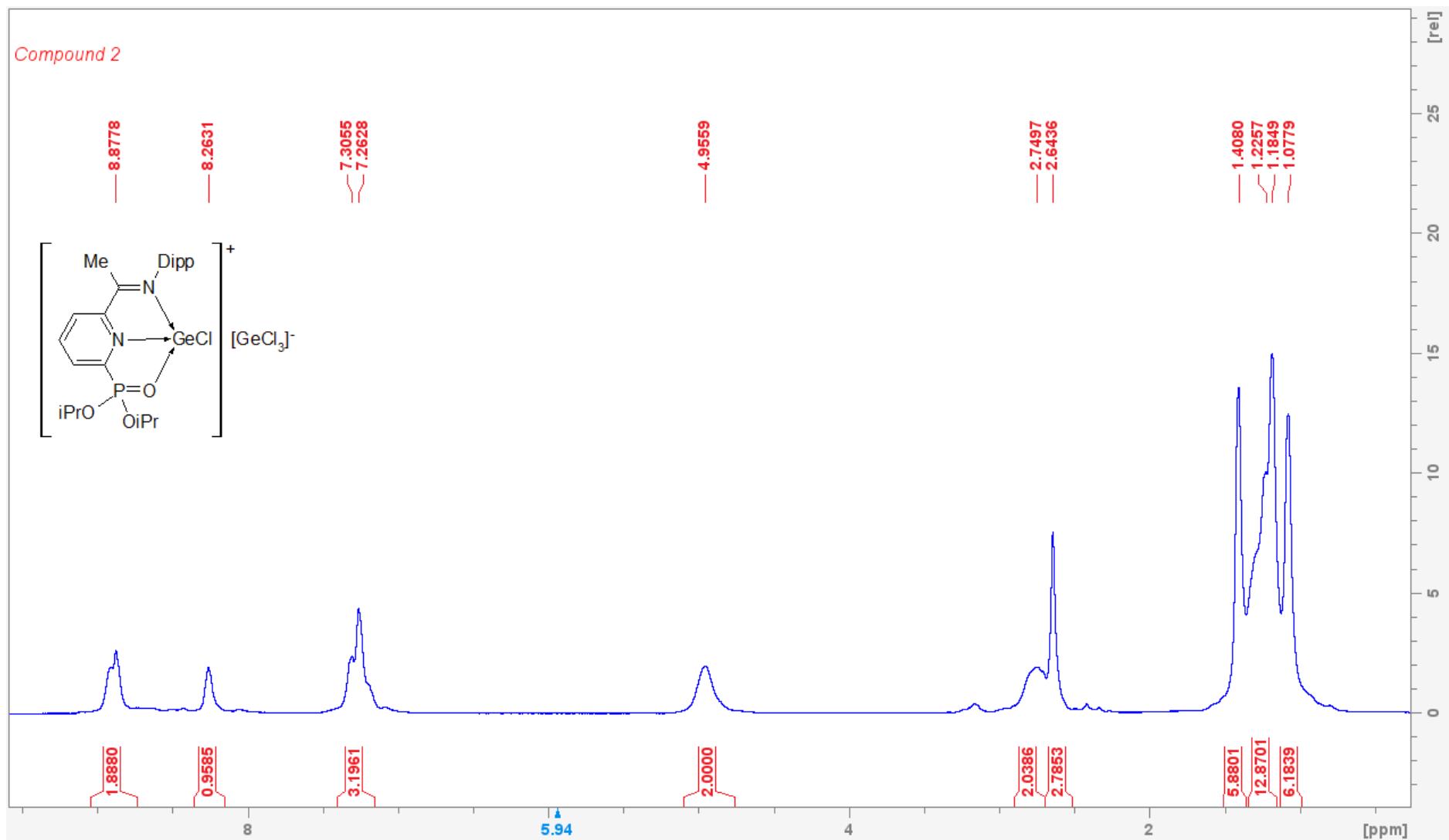
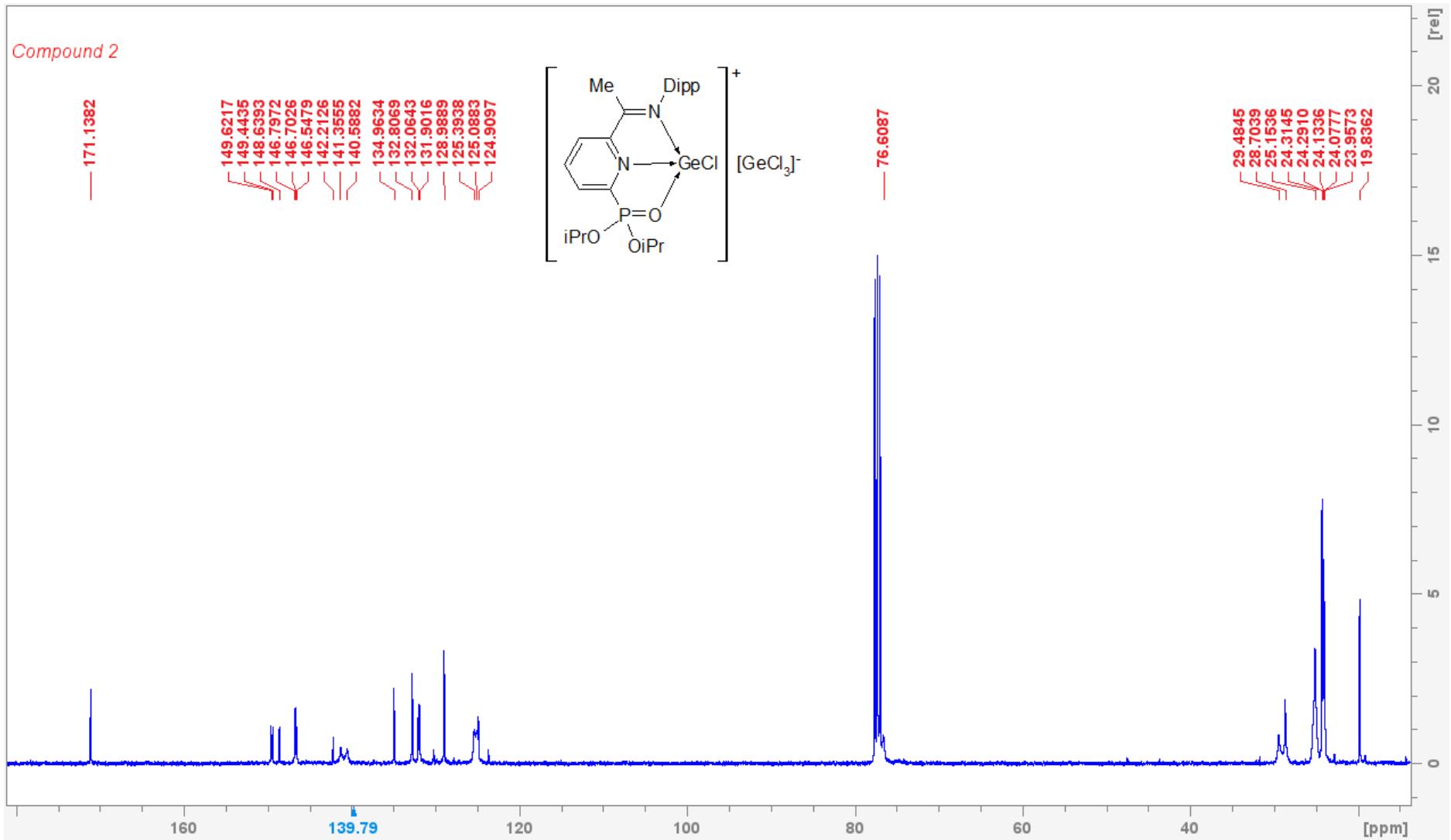
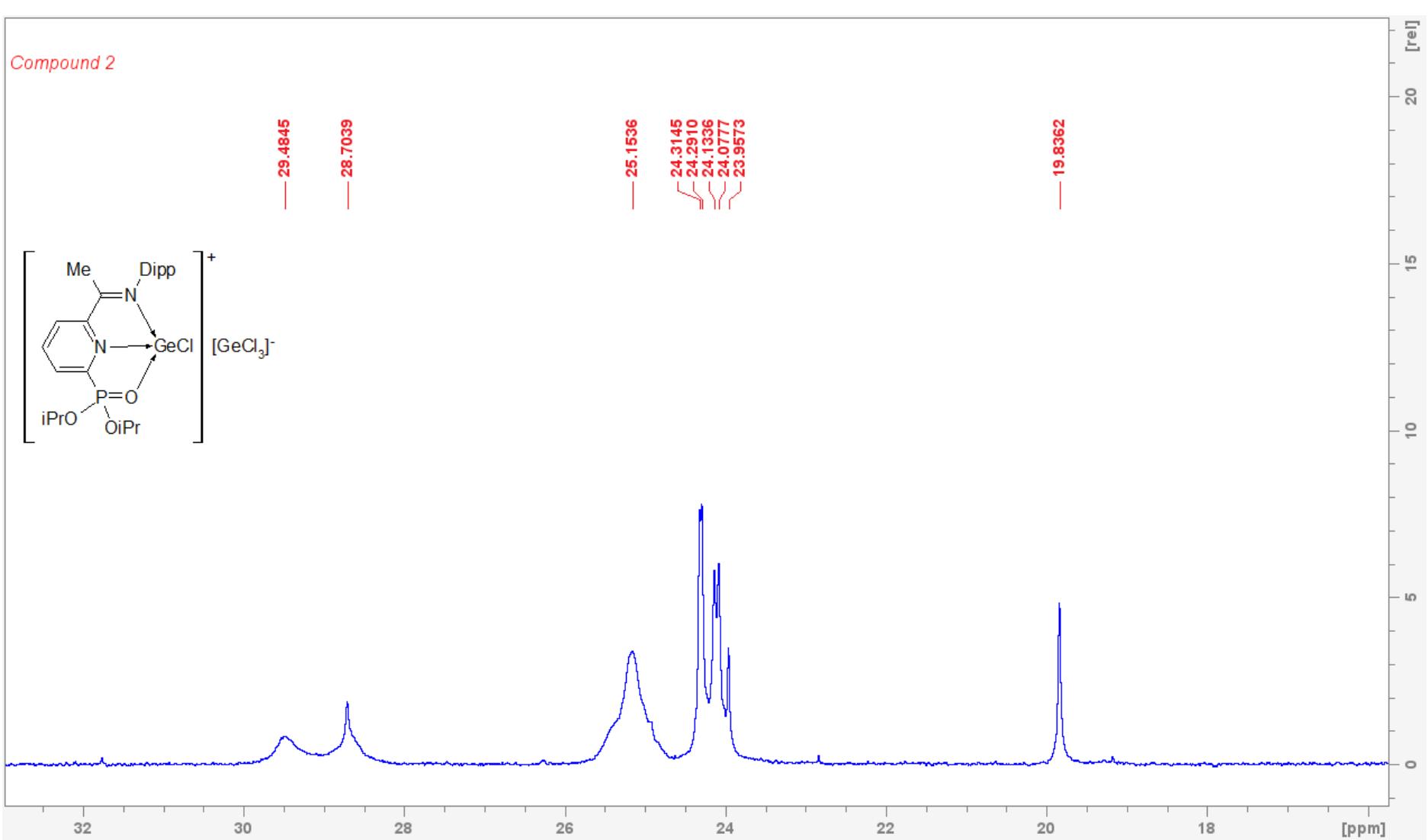


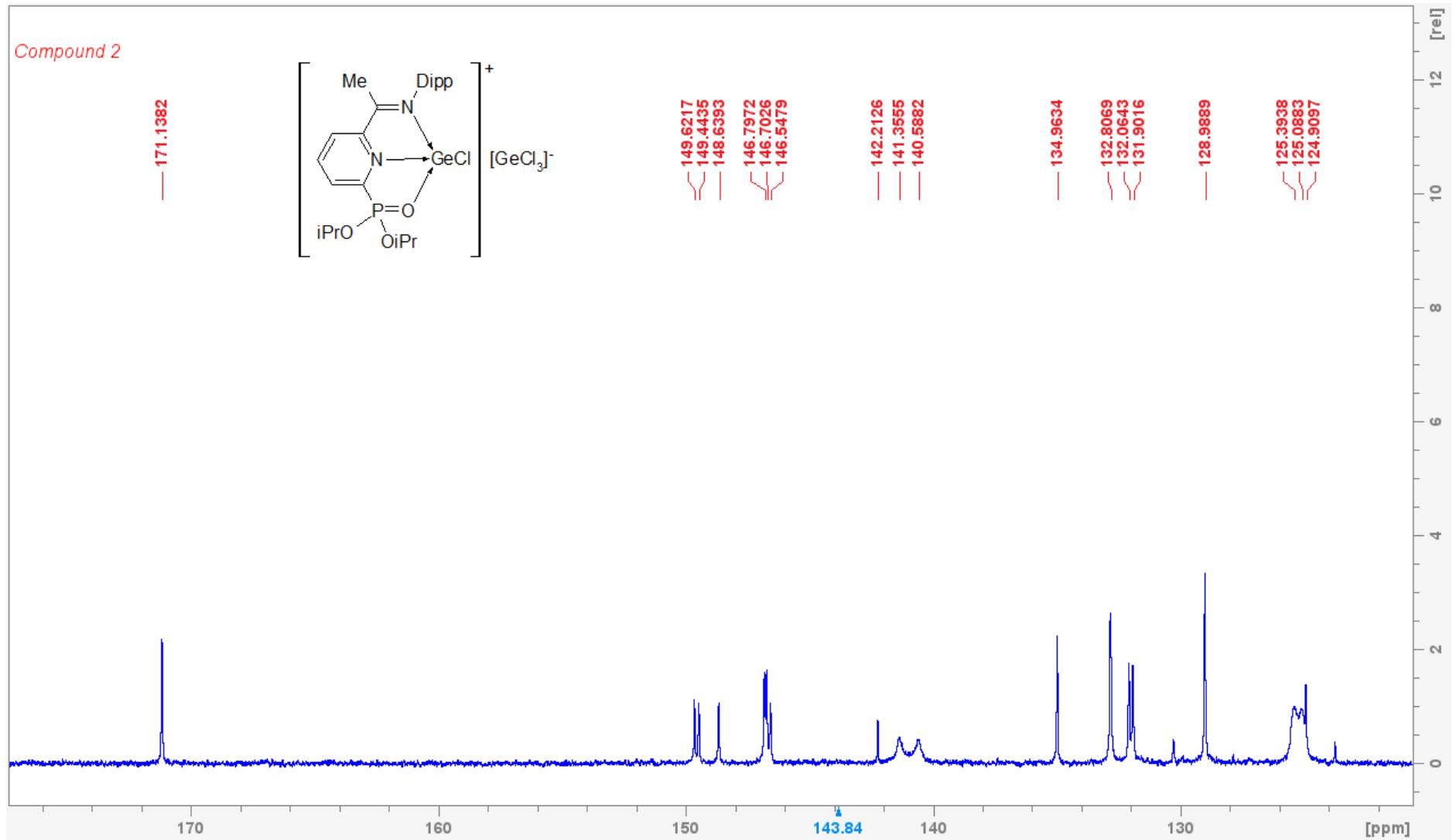
Figure S2.  $^1\text{H}$  NMR spectrum of 2 in  $\text{CDCl}_3$



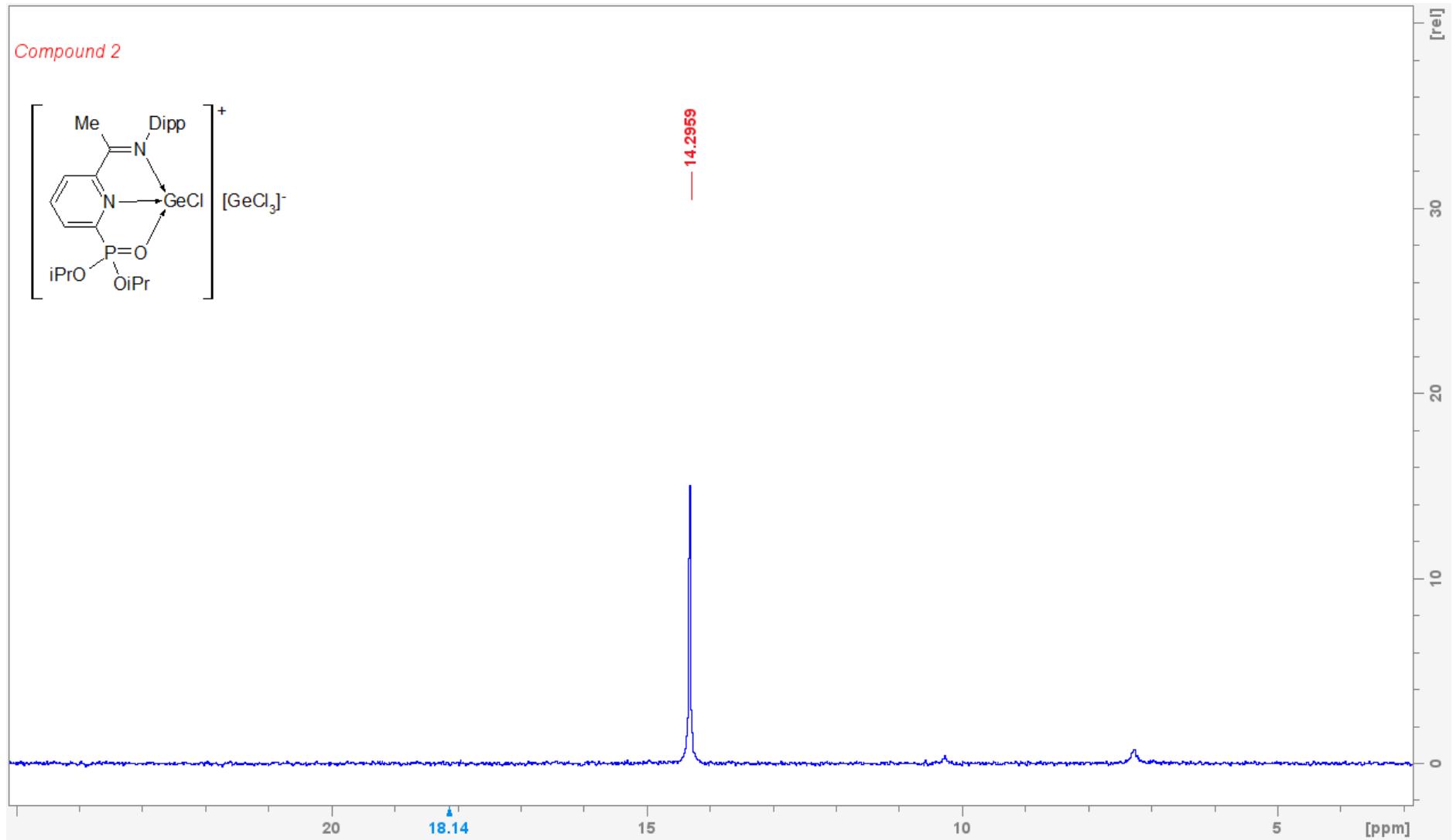
**Figure S3.**  $^{13}\text{C}\{\text{H}\}$  NMR spectrum of **2** in  $\text{CDCl}_3$



**Figure S4.**  $^{13}\text{C}\{\text{H}\}$  NMR spectrum of **2** in  $\text{CDCl}_3$  – aliphatic region



**Figure S5.**  $^{13}\text{C}\{\text{H}\}$  NMR spectrum of **2** in  $\text{CDCl}_3$  – aromatic region



**Figure S6.**  $^{31}\text{P}\{\text{H}\}$  NMR spectrum of **2** in  $\text{CDCl}_3$

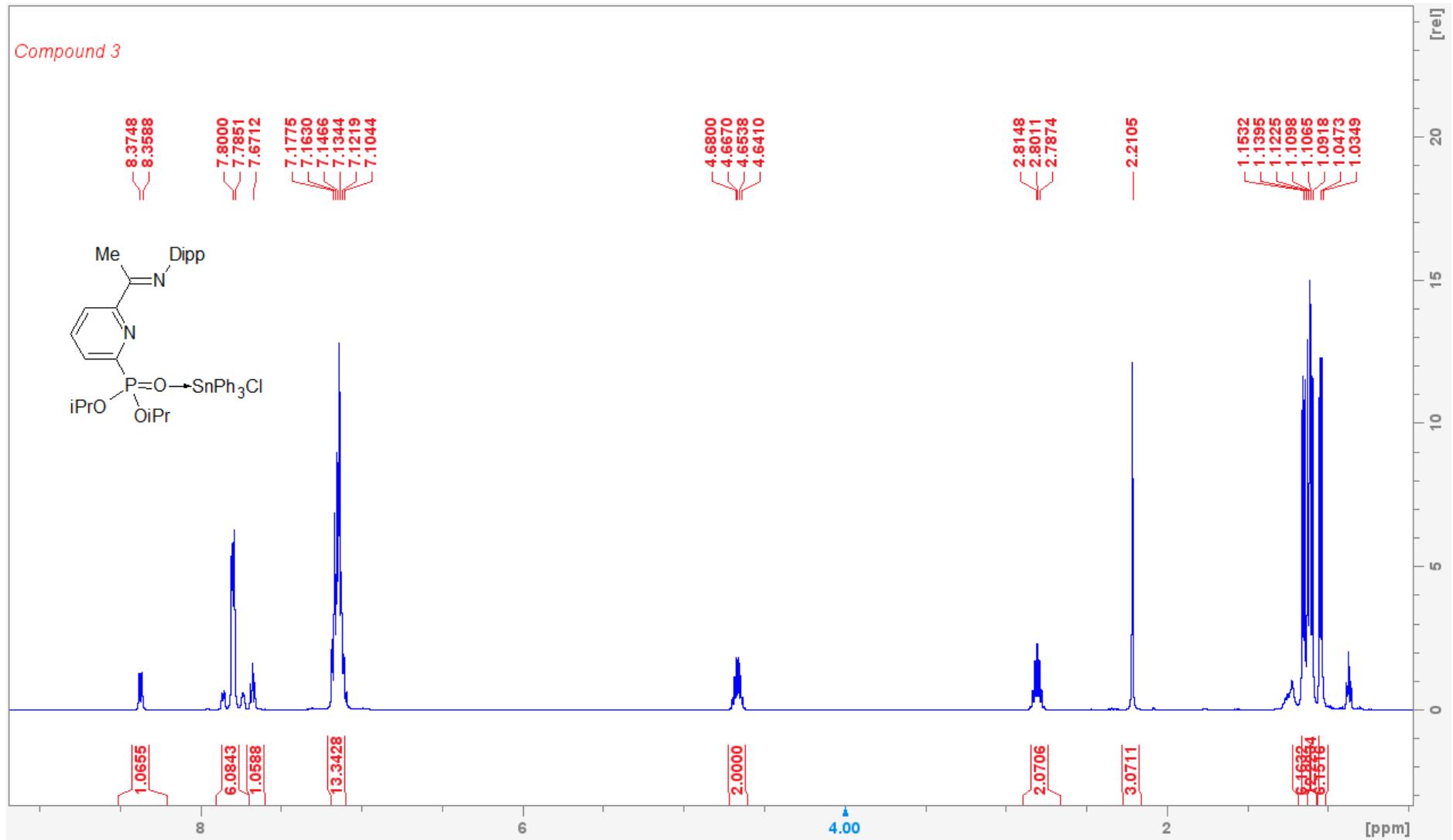
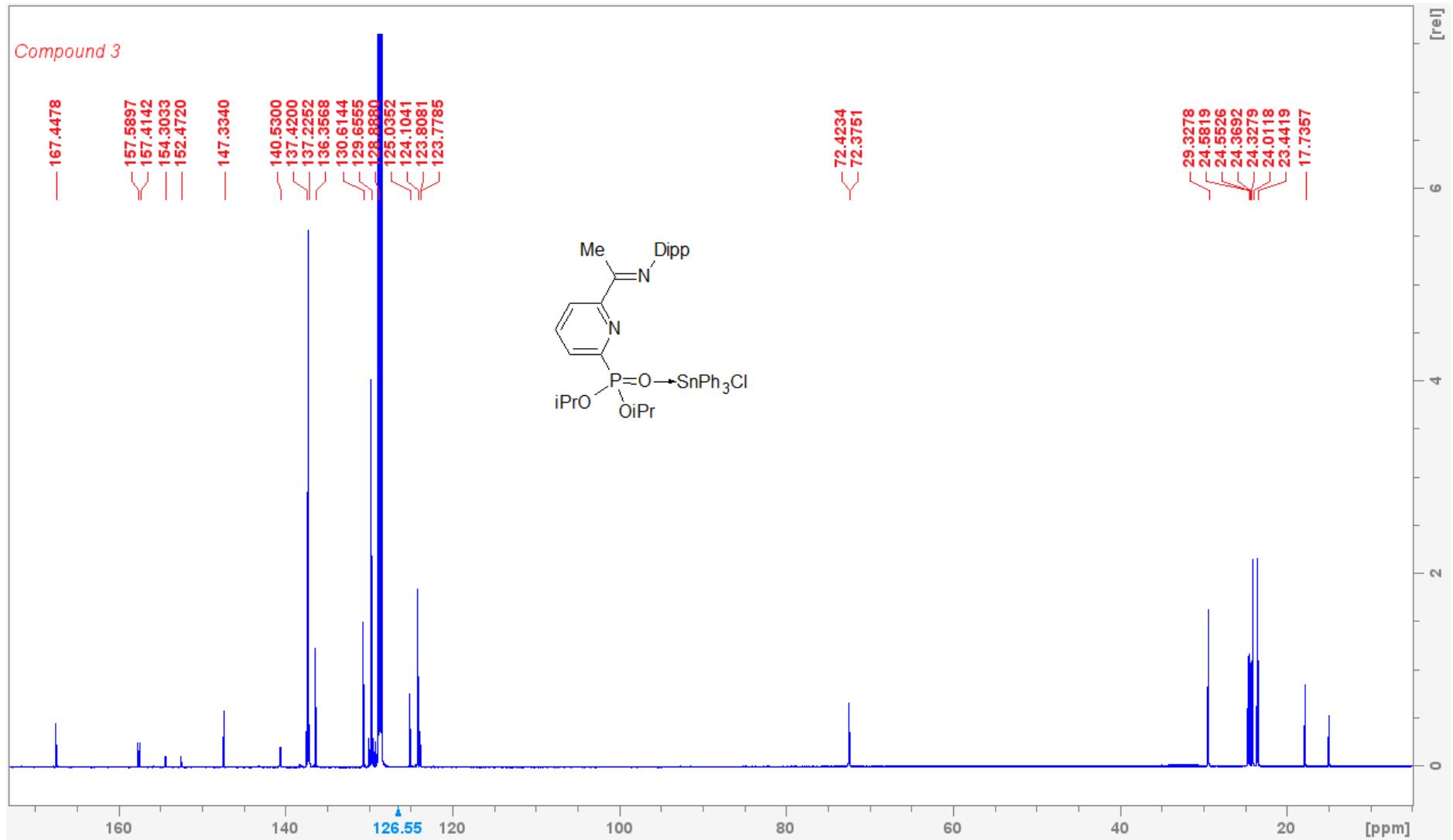
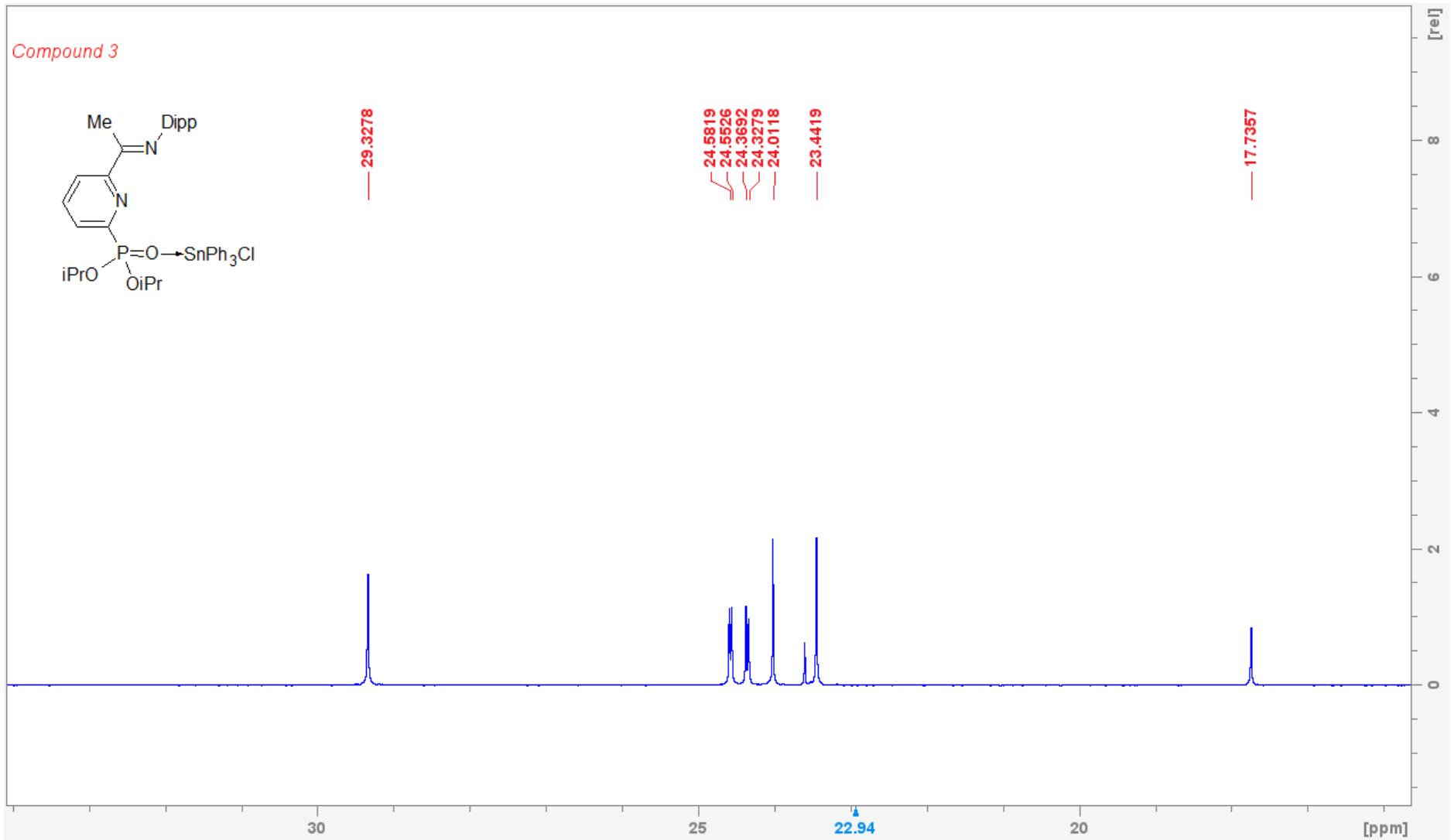


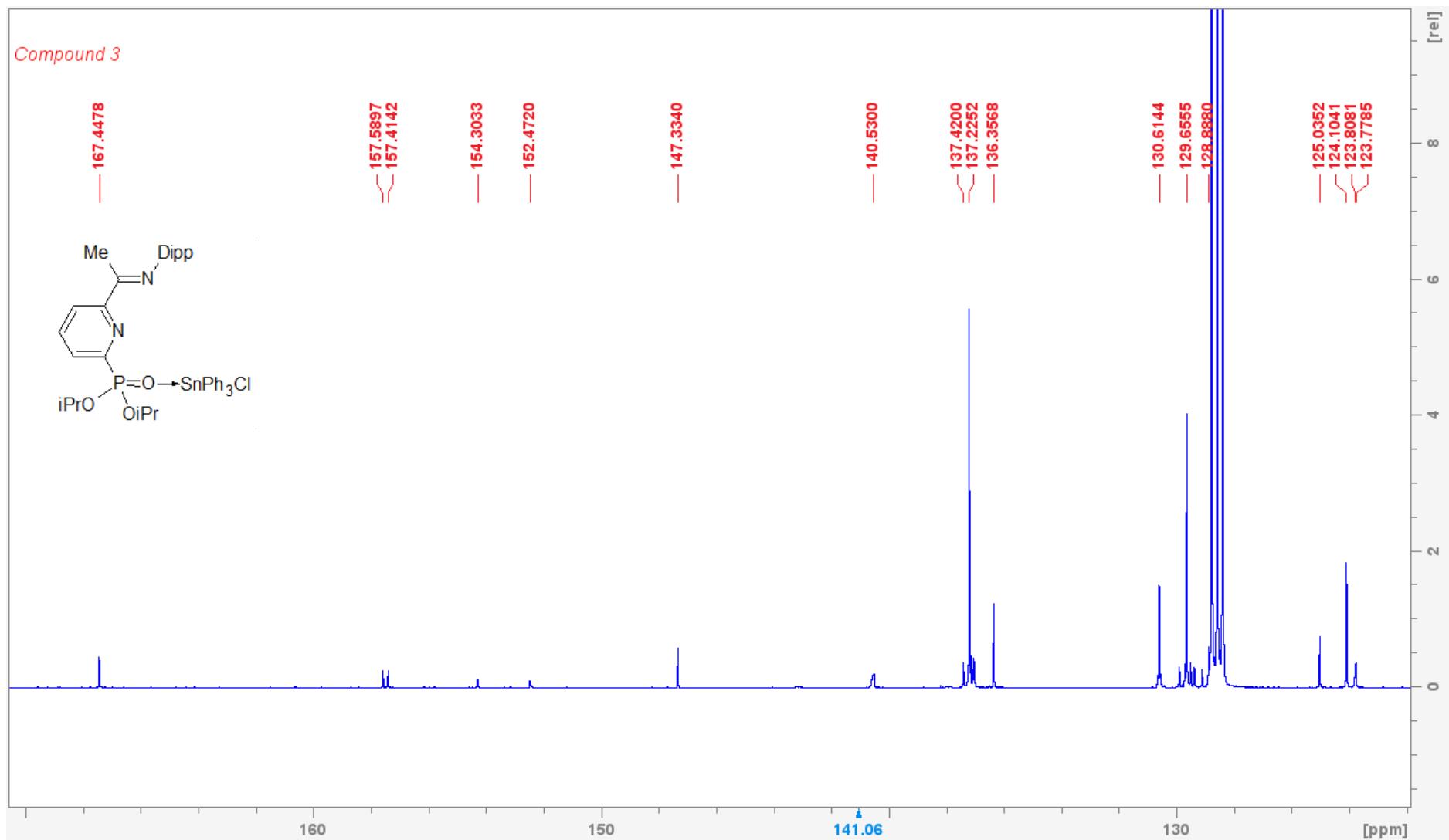
Figure S7.  $^1\text{H}$  NMR spectrum of **3** in  $\text{C}_6\text{D}_6$



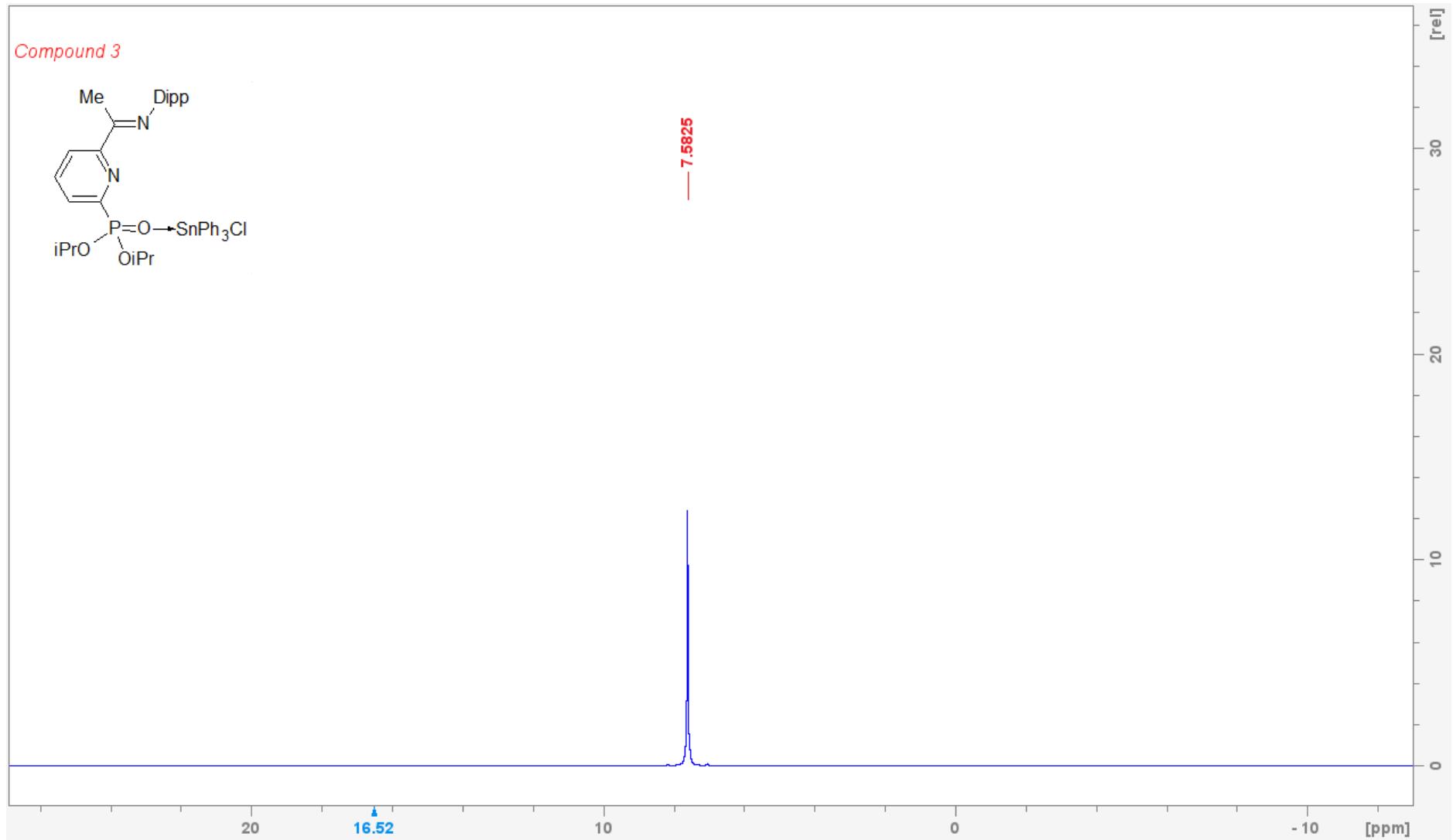
**Figure S8.**  $^{13}\text{C}\{^1\text{H}\}$  NMR spectrum of **3** in  $\text{C}_6\text{D}_6$



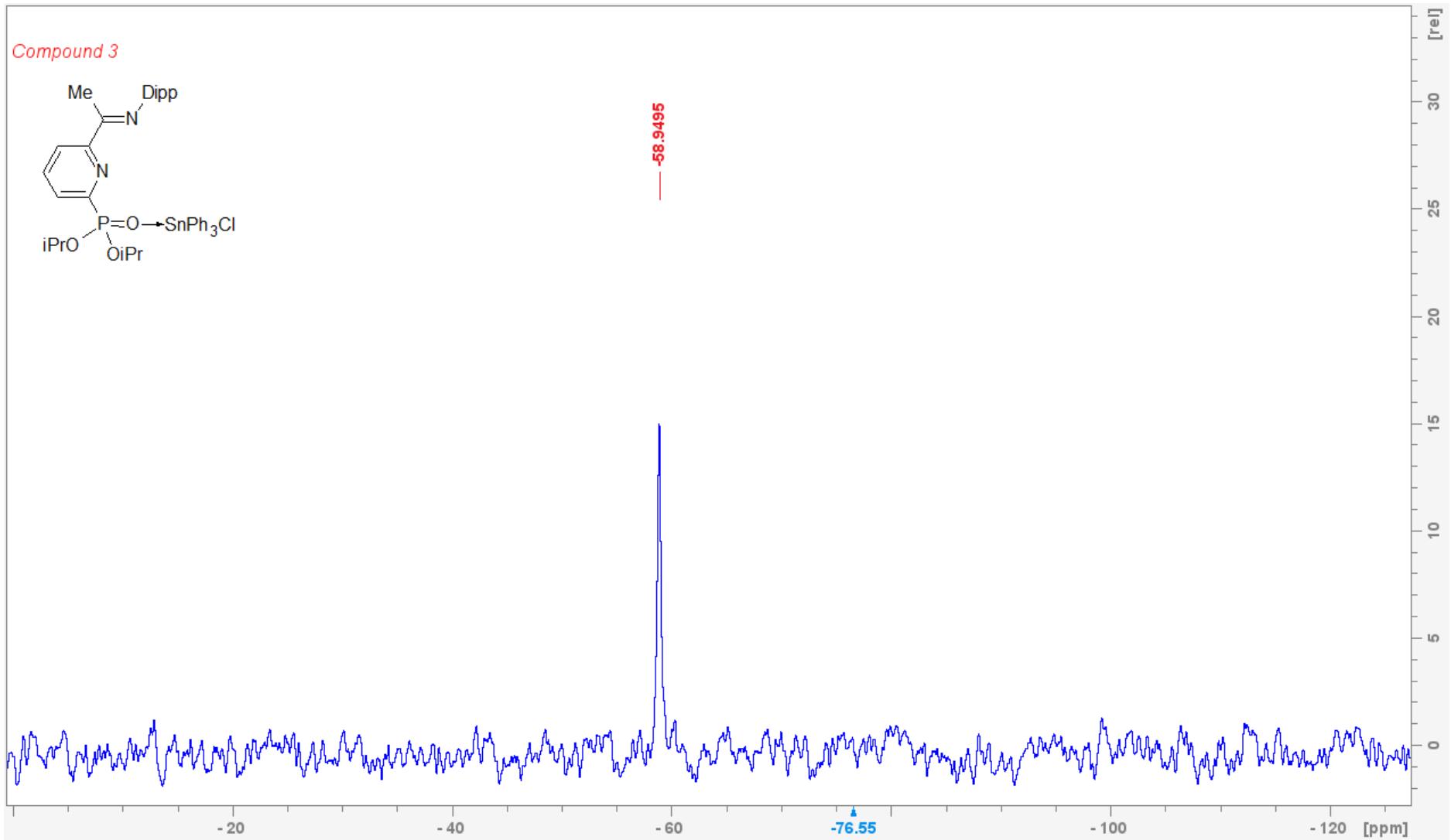
**Figure S9.**  $^{13}\text{C}\{^1\text{H}\}$  NMR spectrum of **3** in  $\text{C}_6\text{D}_6$  – aliphatic region



**Figure S10.**  $^{13}\text{C}\{^1\text{H}\}$  NMR spectrum of **3** in  $\text{C}_6\text{D}_6$  – aromatic region



**Figure S11.**  $^{31}\text{P}\{\text{H}\}$  NMR spectrum of **3** in C<sub>6</sub>D<sub>6</sub>



**Figure S12.**  $^{119}\text{Sn}\{\text{H}\}$  NMR spectrum of **3** in  $\text{C}_6\text{D}_6$

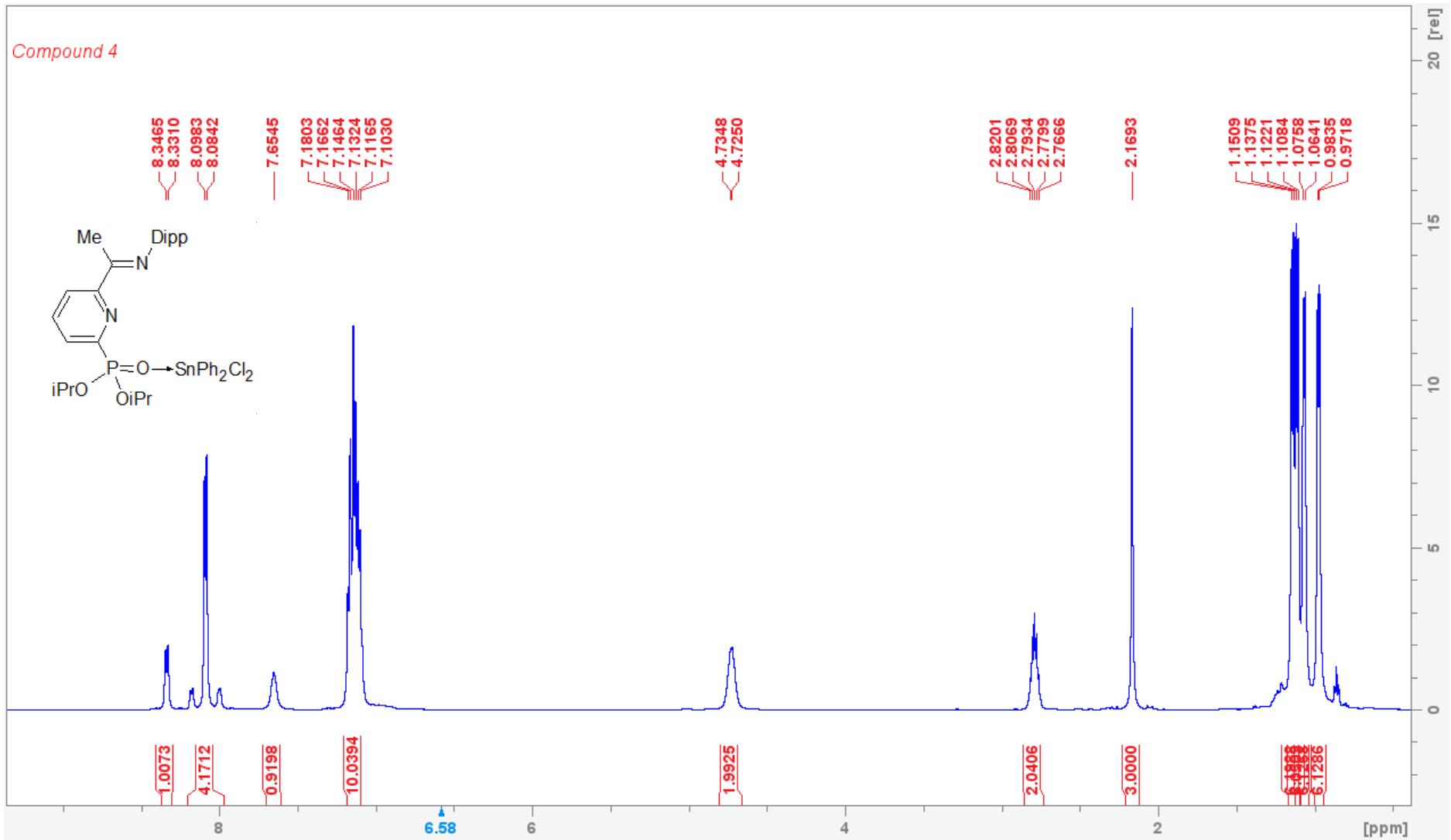


Figure S13.  $^1\text{H}$  NMR spectrum of **4** in  $\text{C}_6\text{D}_6$

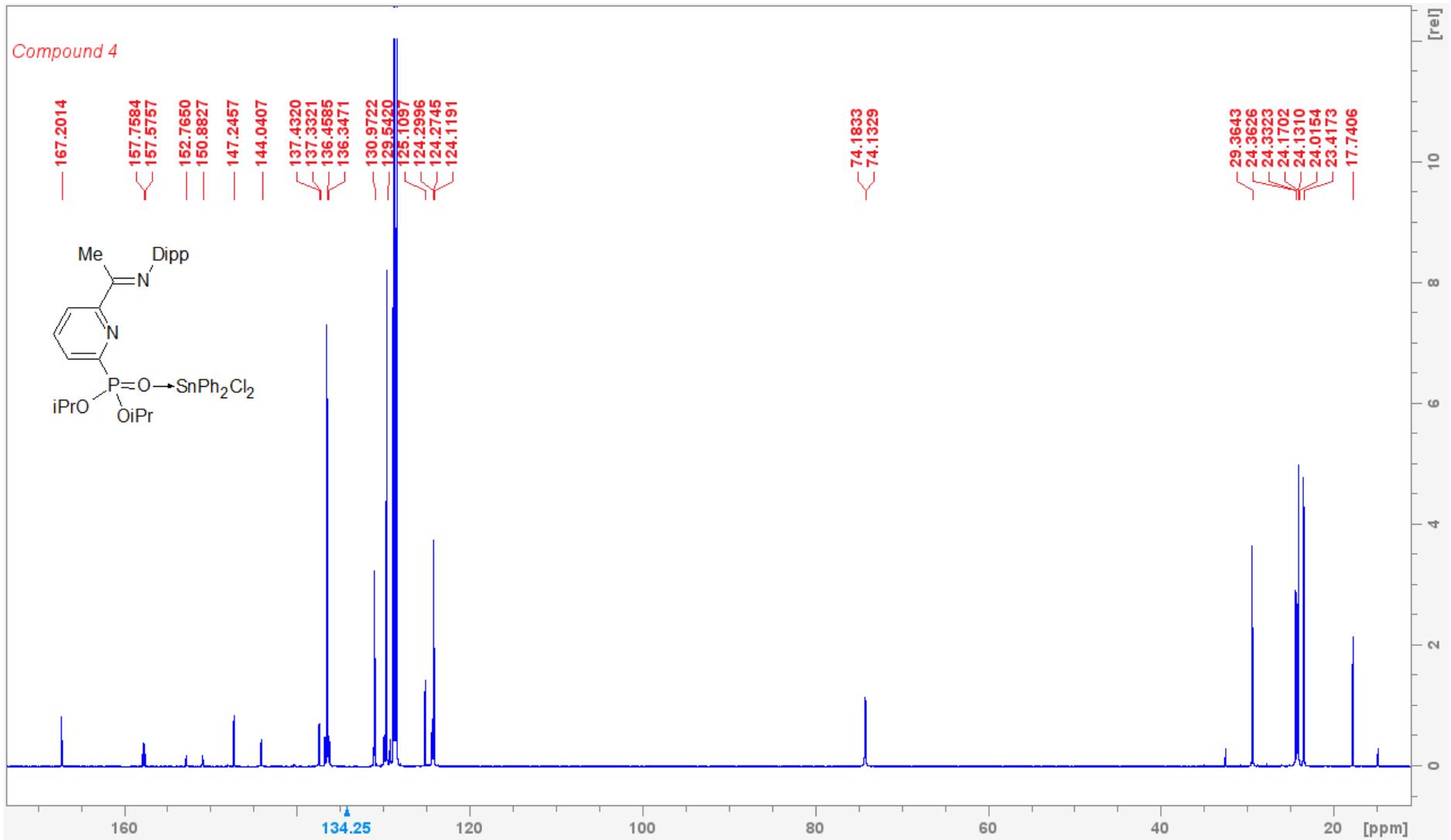
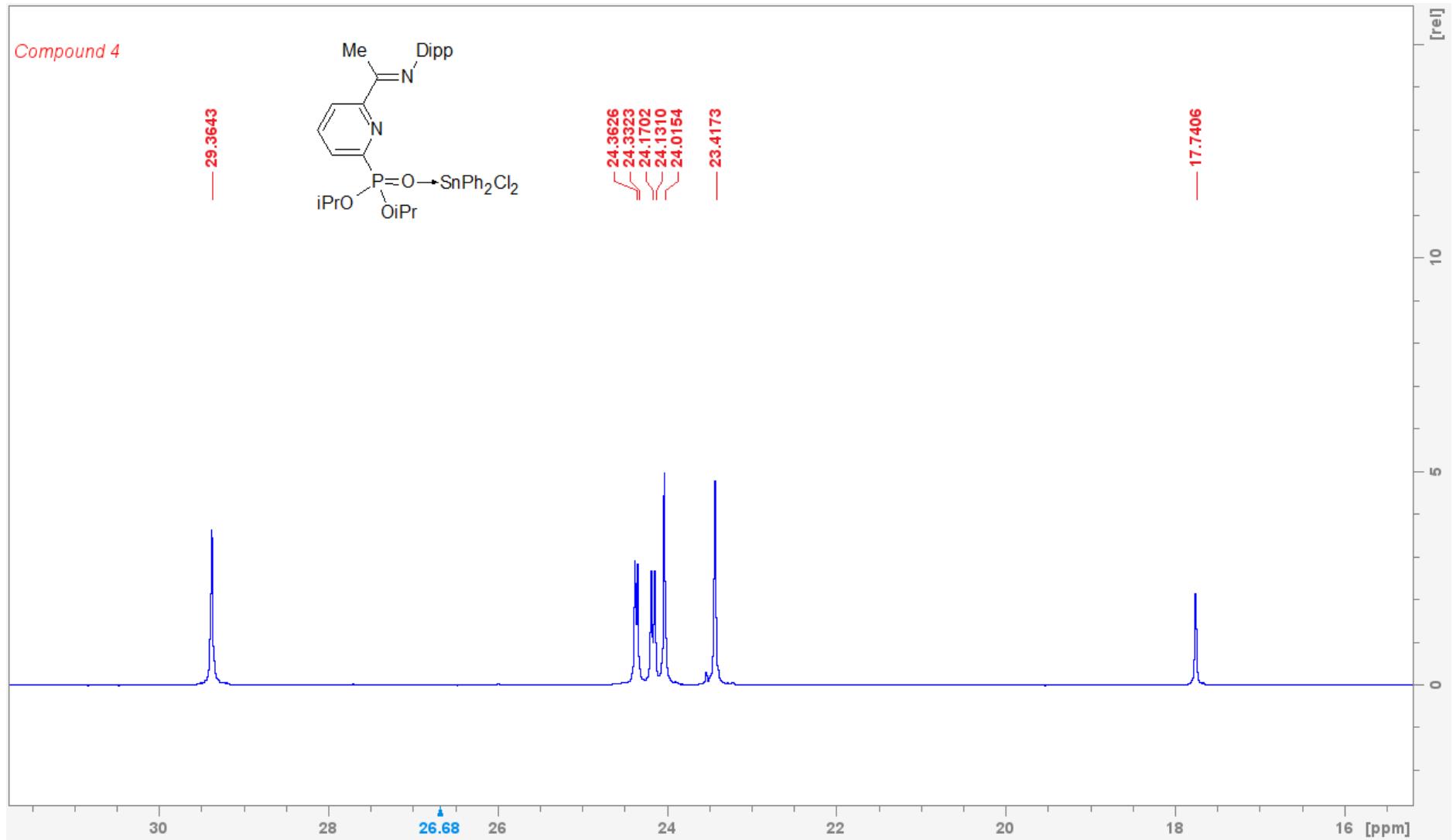
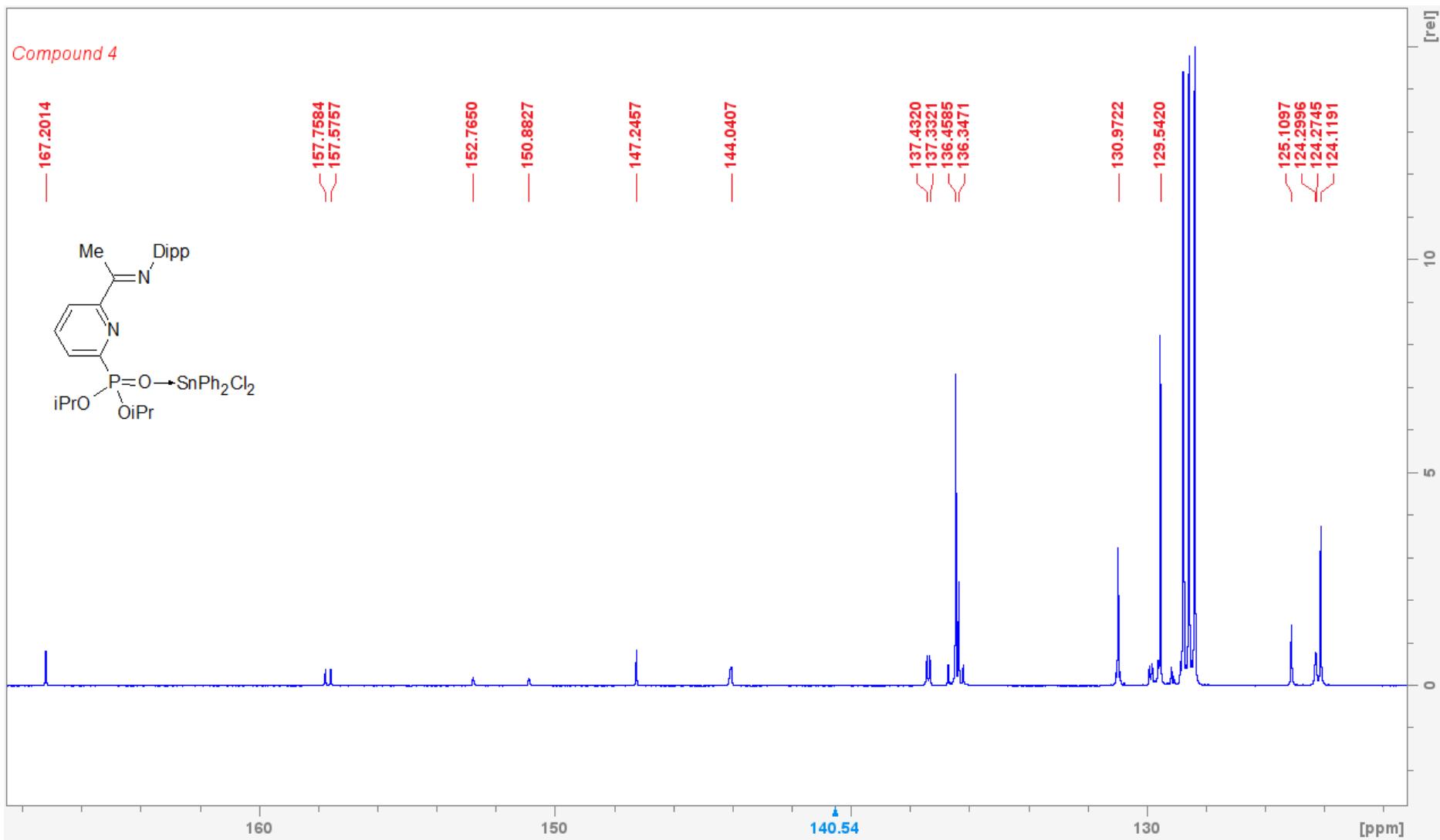


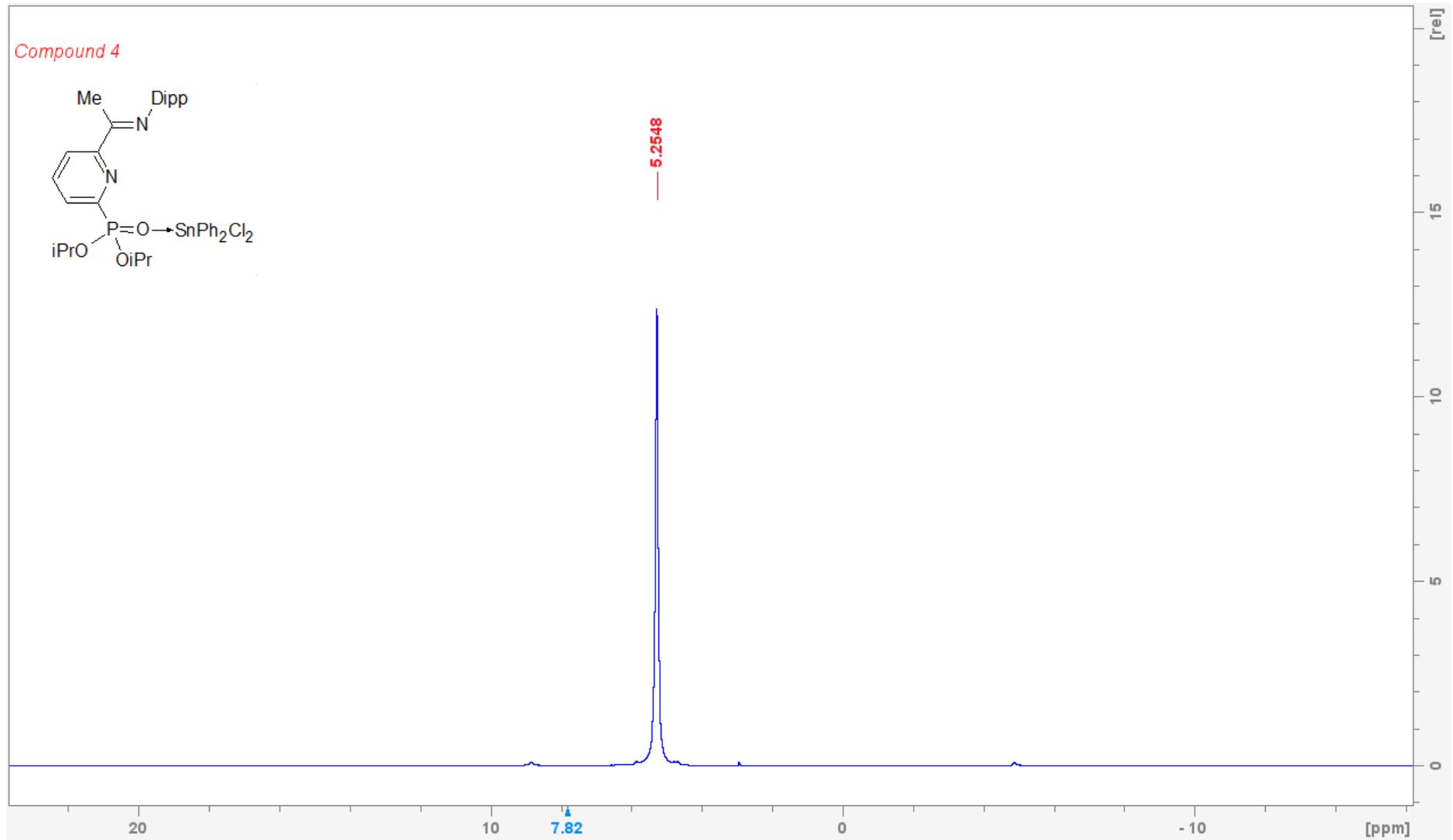
Figure S14.  $^{13}\text{C}\{\text{H}\}$  NMR spectrum of **4** in  $\text{C}_6\text{D}_6$



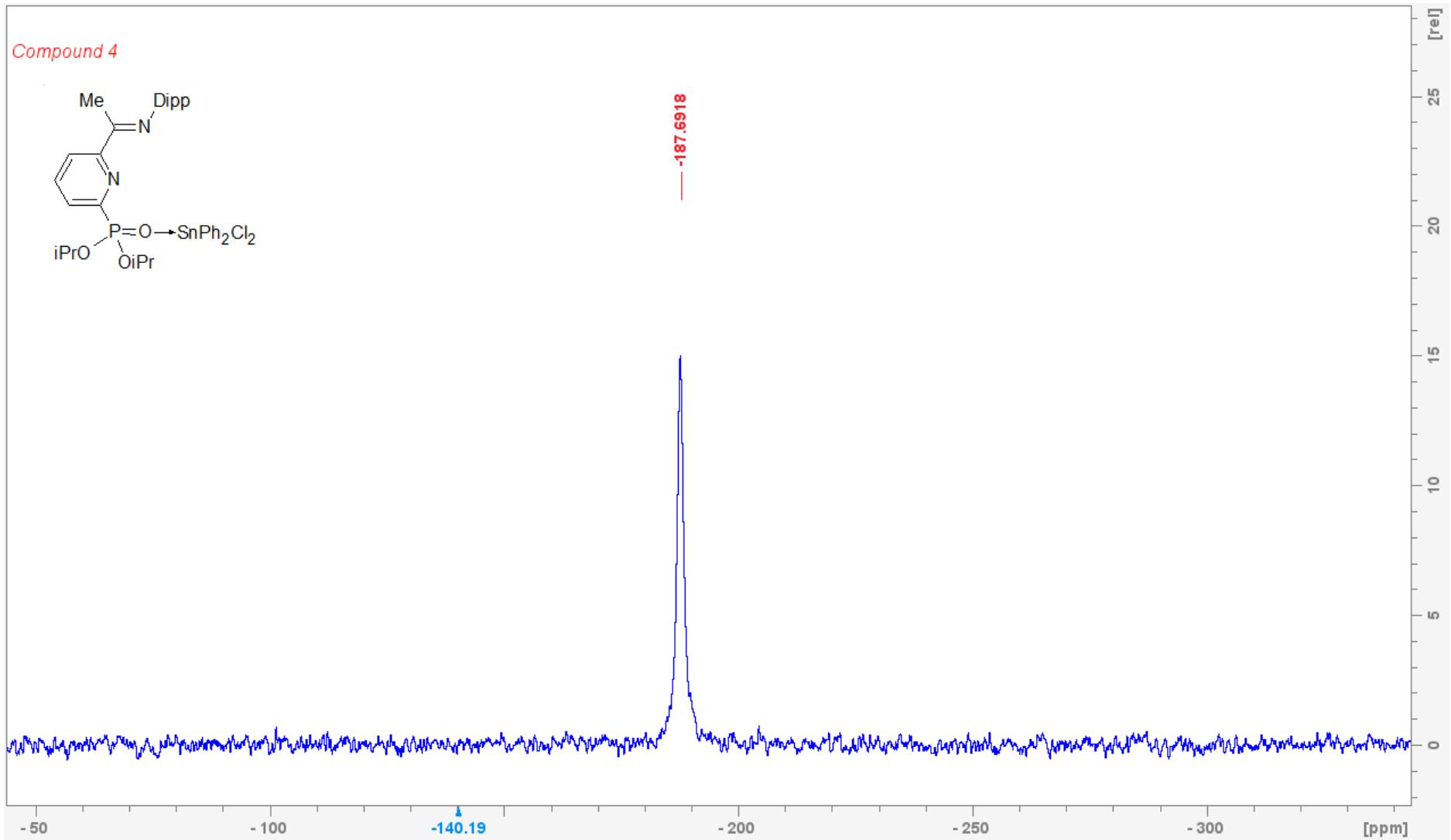
**Figure S15.**  $^{13}\text{C}\{\text{H}\}$  NMR spectrum of **4** in  $\text{C}_6\text{D}_6$  – aliphatic region



**Figure S16.**  $^{13}\text{C}\{\text{H}\}$  NMR spectrum of **4** in  $\text{C}_6\text{D}_6$  – aromatic region



**Figure S17.**  $^{31}\text{P}\{\text{H}\}$  NMR spectrum of **4** in  $\text{C}_6\text{D}_6$



**Figure S18.**  $^{119}\text{Sn}\{^1\text{H}\}$  NMR spectrum of **4** in C<sub>6</sub>D<sub>6</sub>

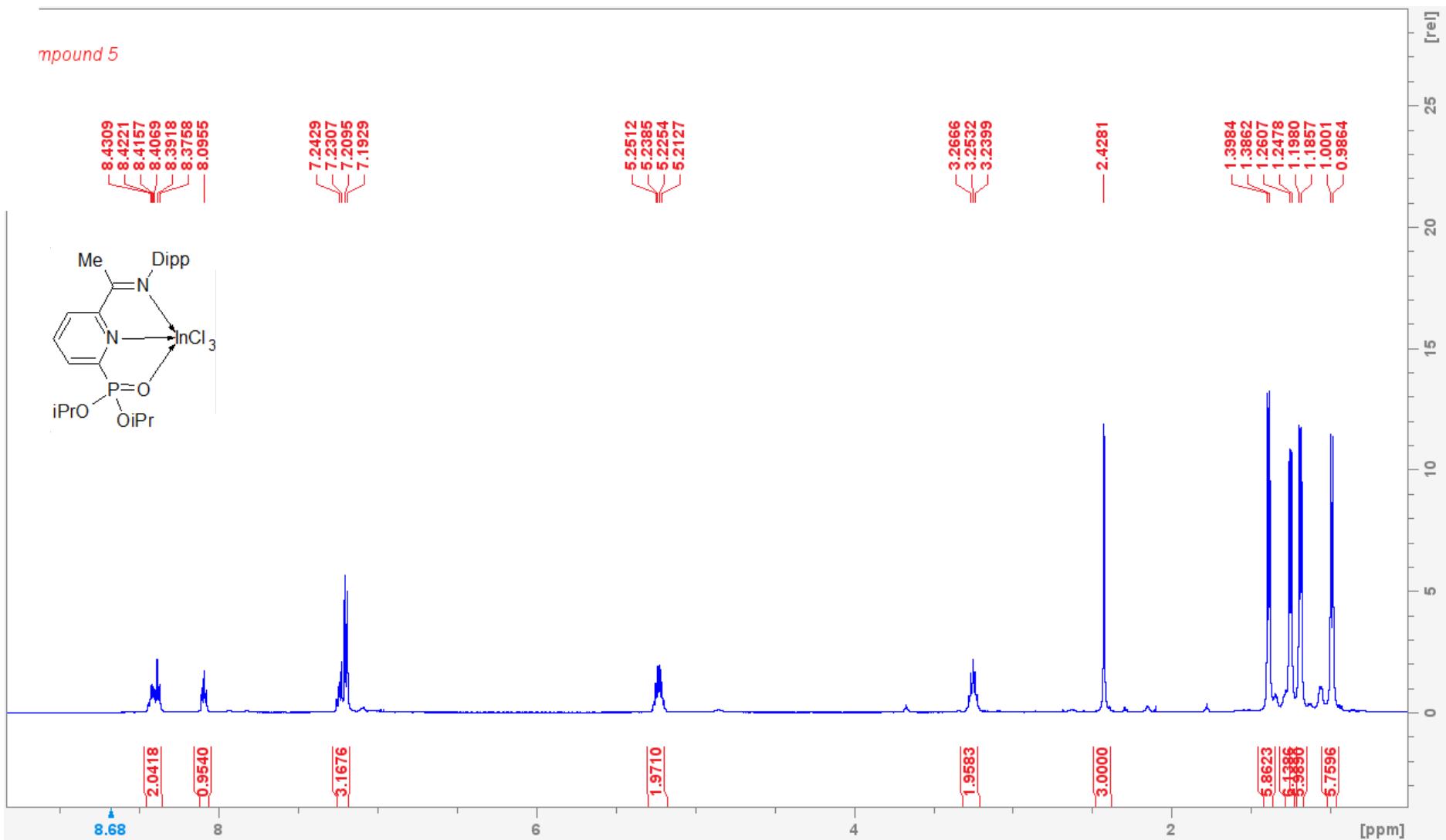
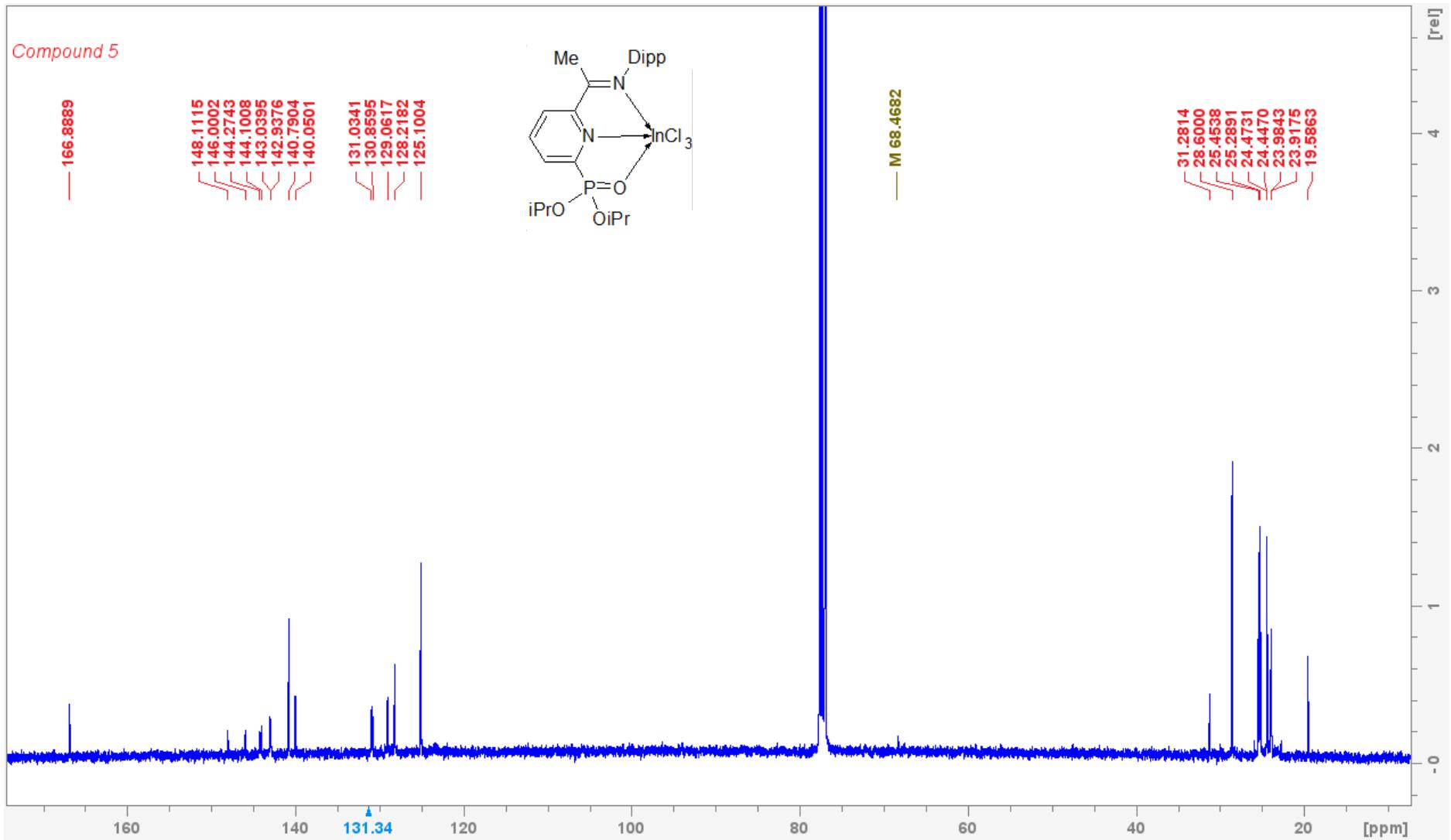
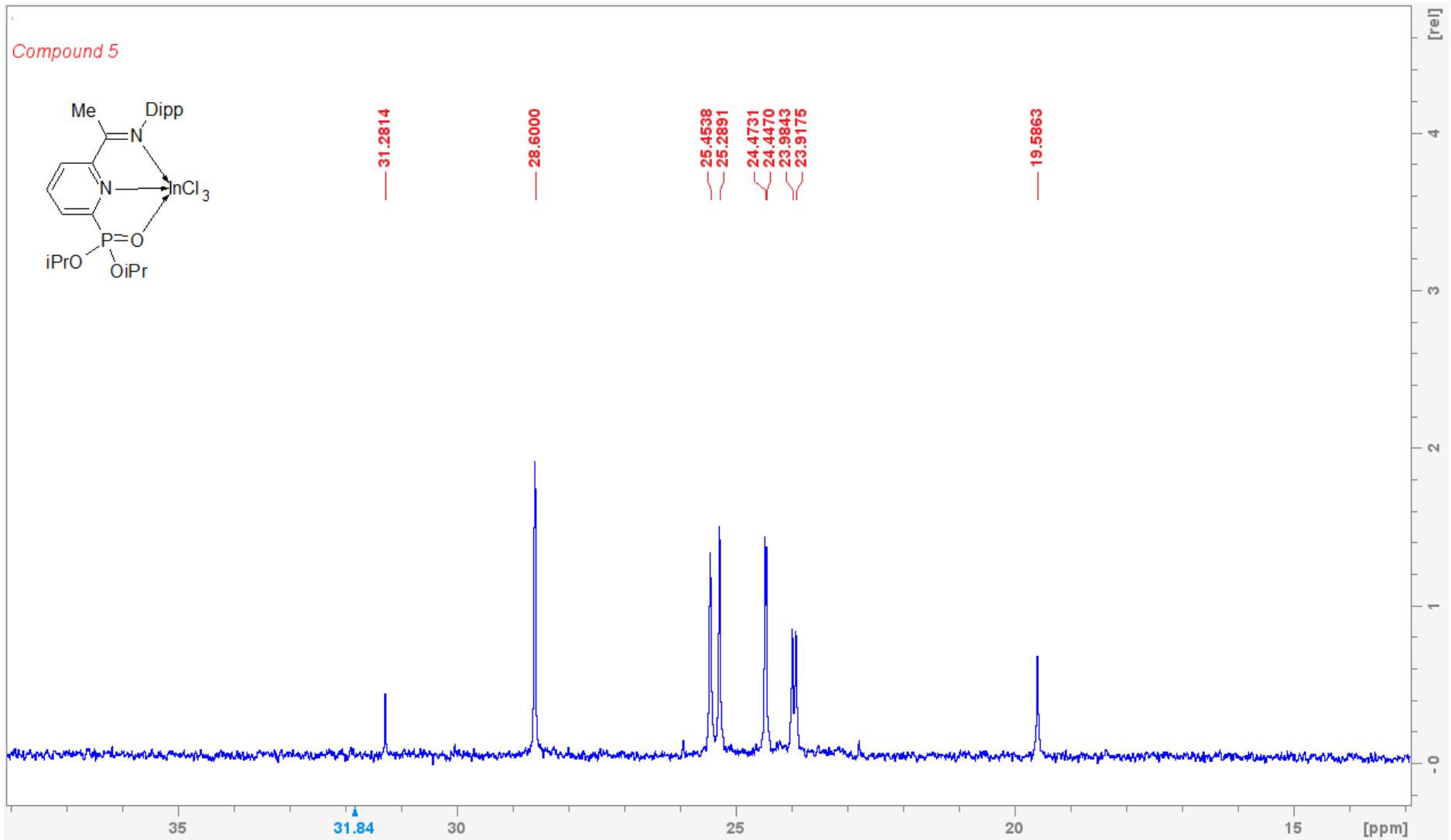


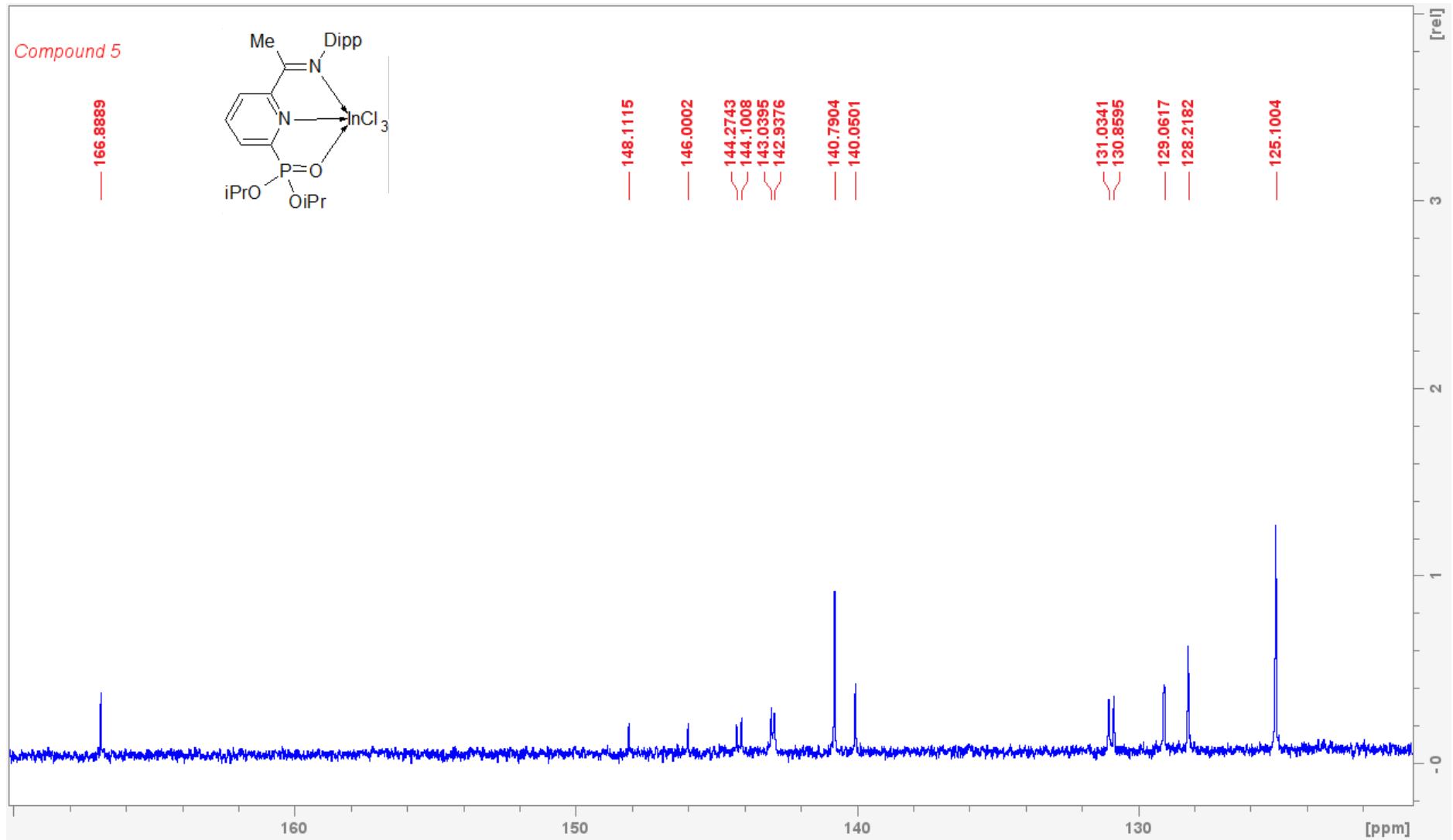
Figure S19.  $^1\text{H}$  NMR spectrum of 5 in  $\text{CDCl}_3$



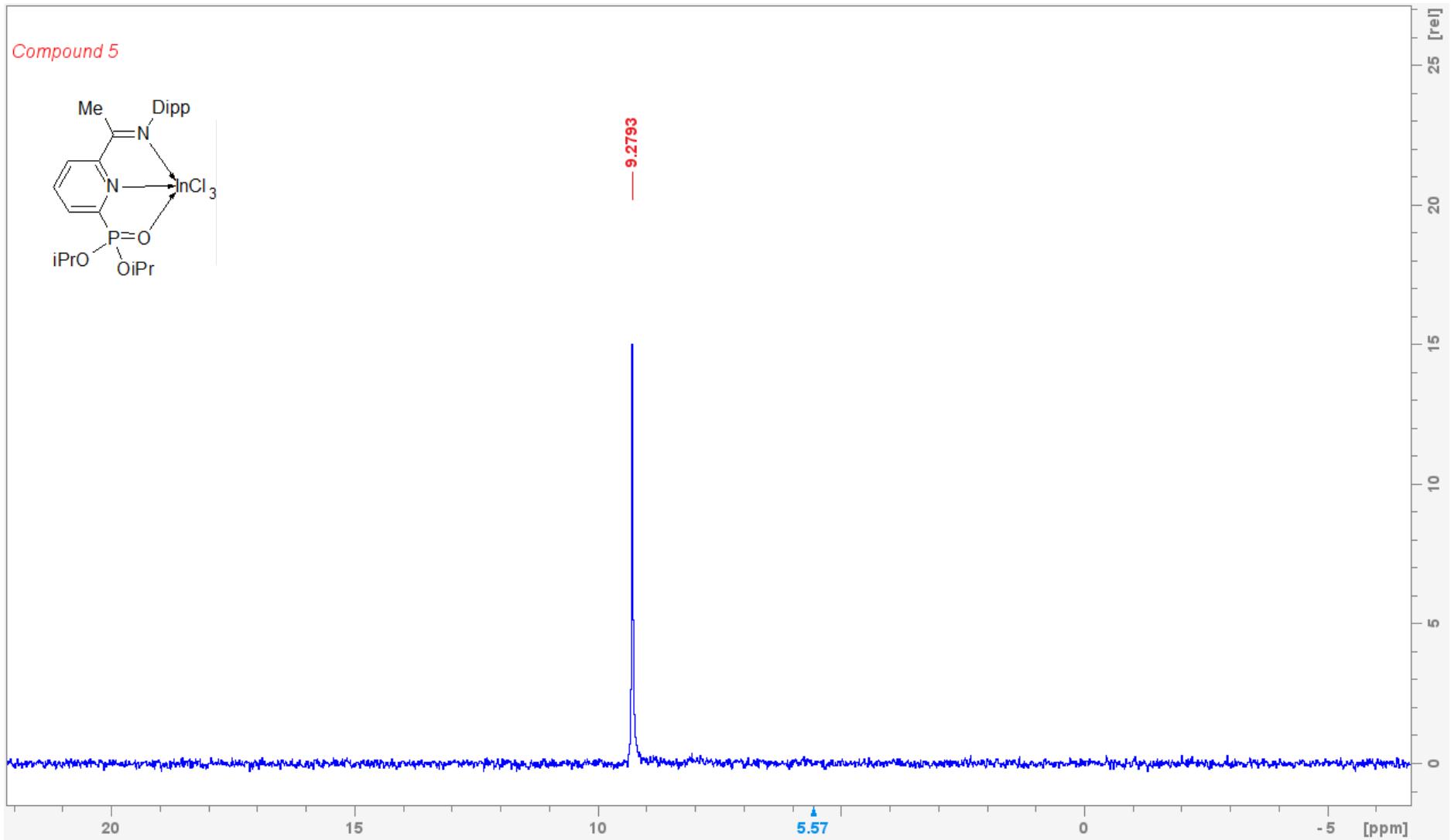
**Figure S20.**  $^{13}\text{C}\{\text{H}\}$  NMR spectrum of **5** in  $\text{CDCl}_3$



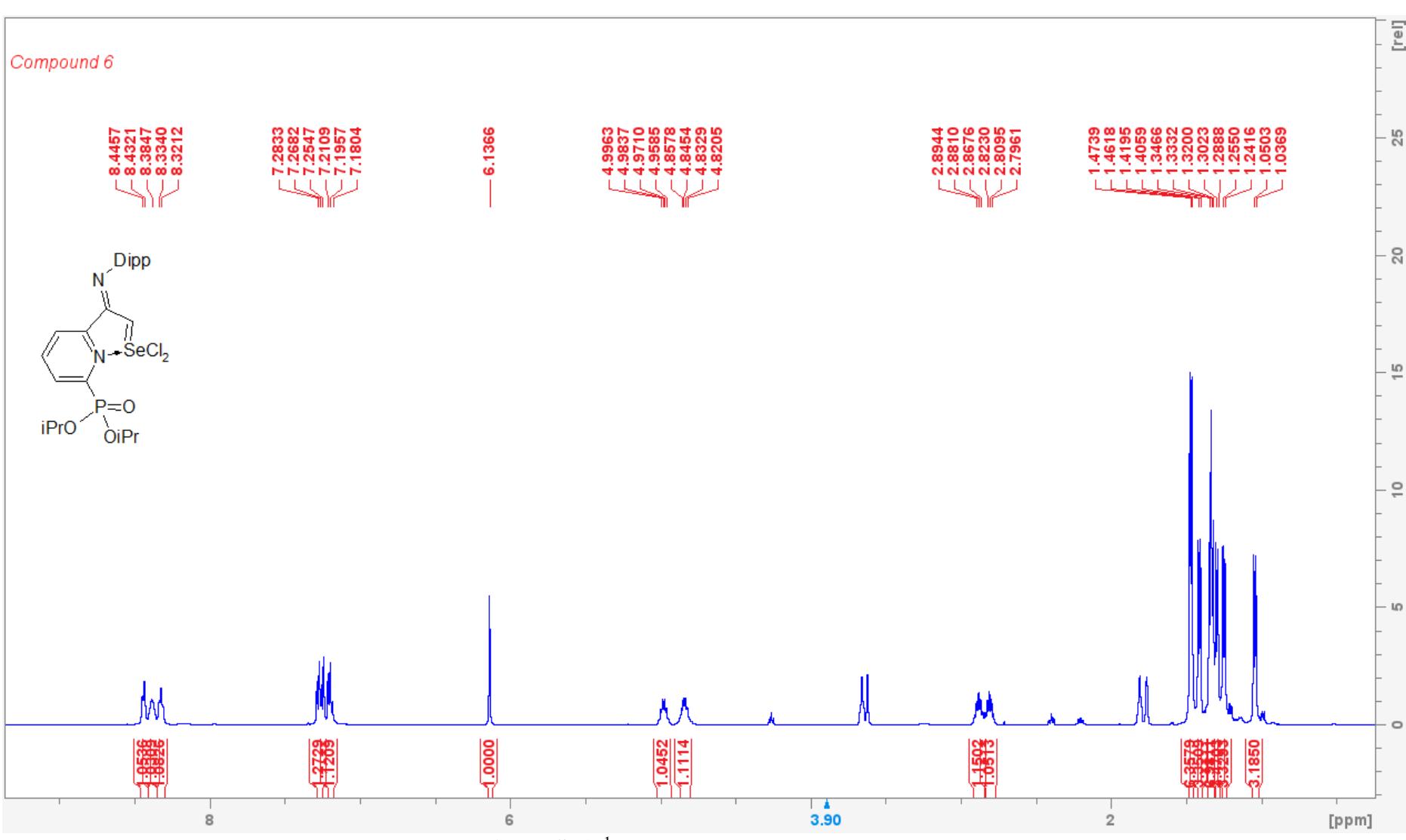
**Figure S21.**  $^{13}\text{C}\{\text{H}\}$  NMR spectrum of **5** in  $\text{CDCl}_3$  – aliphatic region



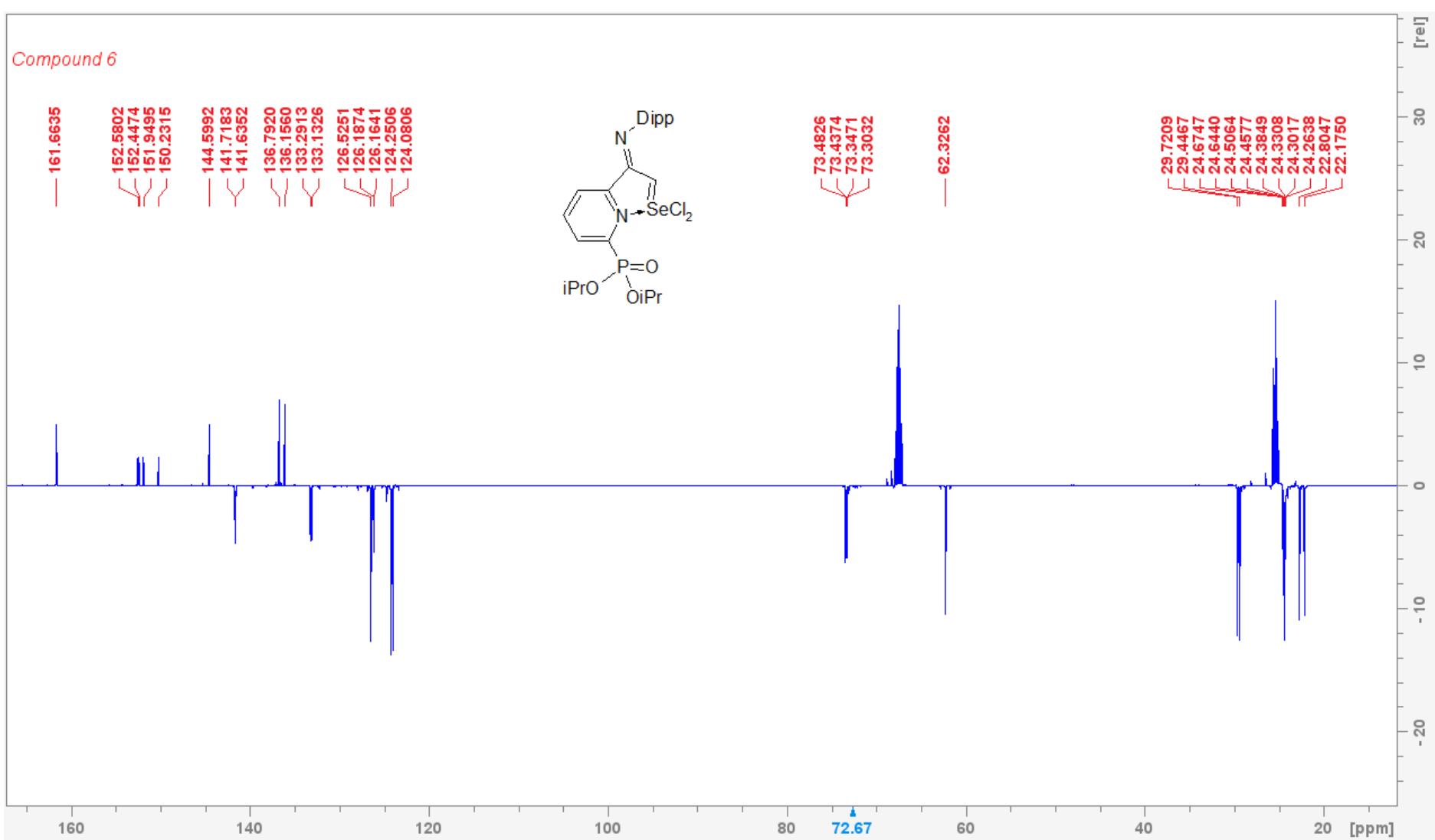
**Figure S22.**  $^{13}\text{C}\{\text{H}\}$  NMR spectrum of **5** in  $\text{CDCl}_3$  – aromatic region



**Figure S23.**  $^{31}\text{P}\{\text{H}\}$  NMR spectrum of **5** in CDCl<sub>3</sub>



**Figure S24.**  $^1\text{H}$  NMR spectrum of **6** in  $\text{THF-d}_8$



**Figure S25.**  $^{13}\text{C}\{\text{H}\}$  APT NMR spectrum of **6** in THF-d8

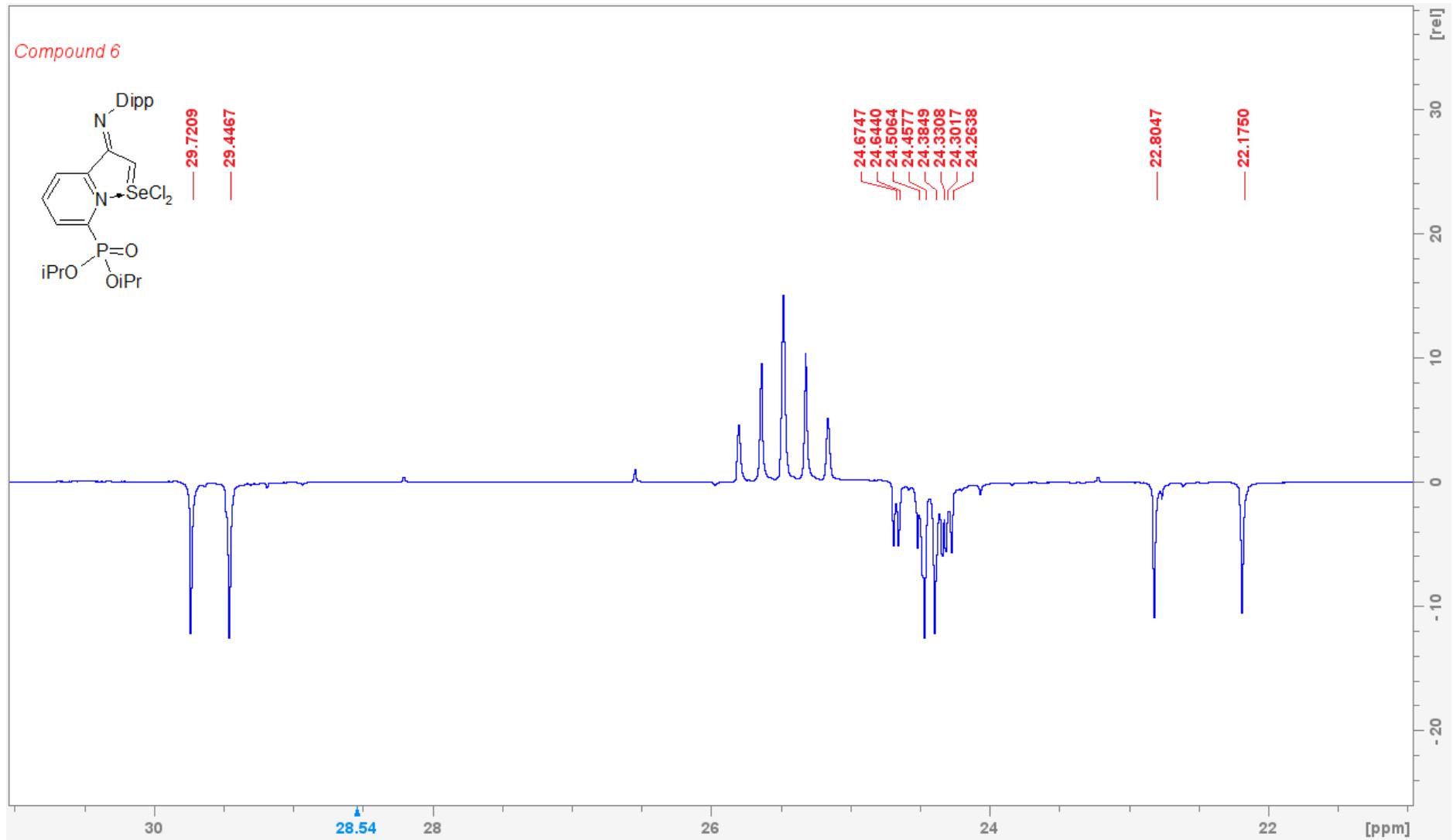
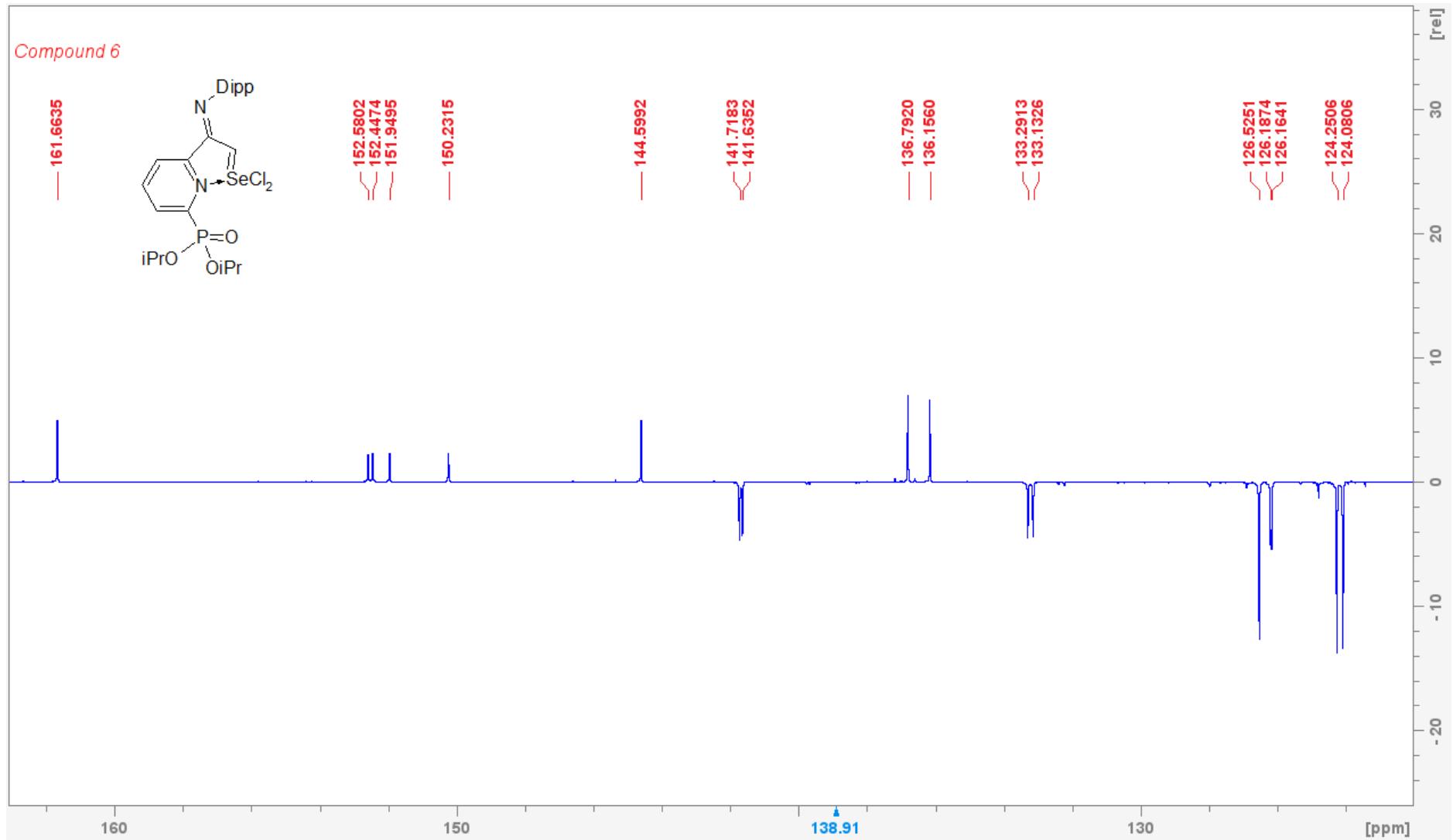
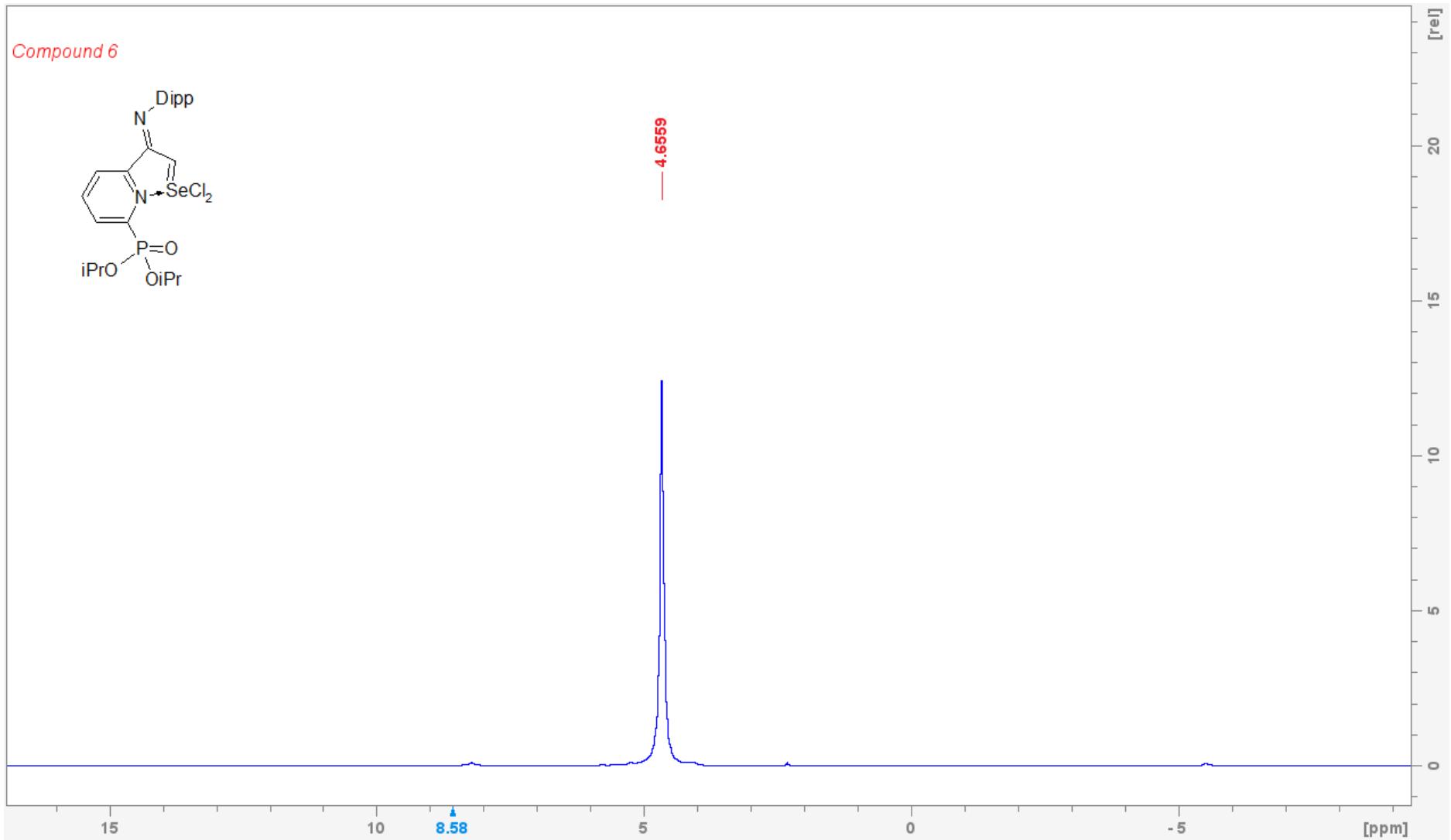


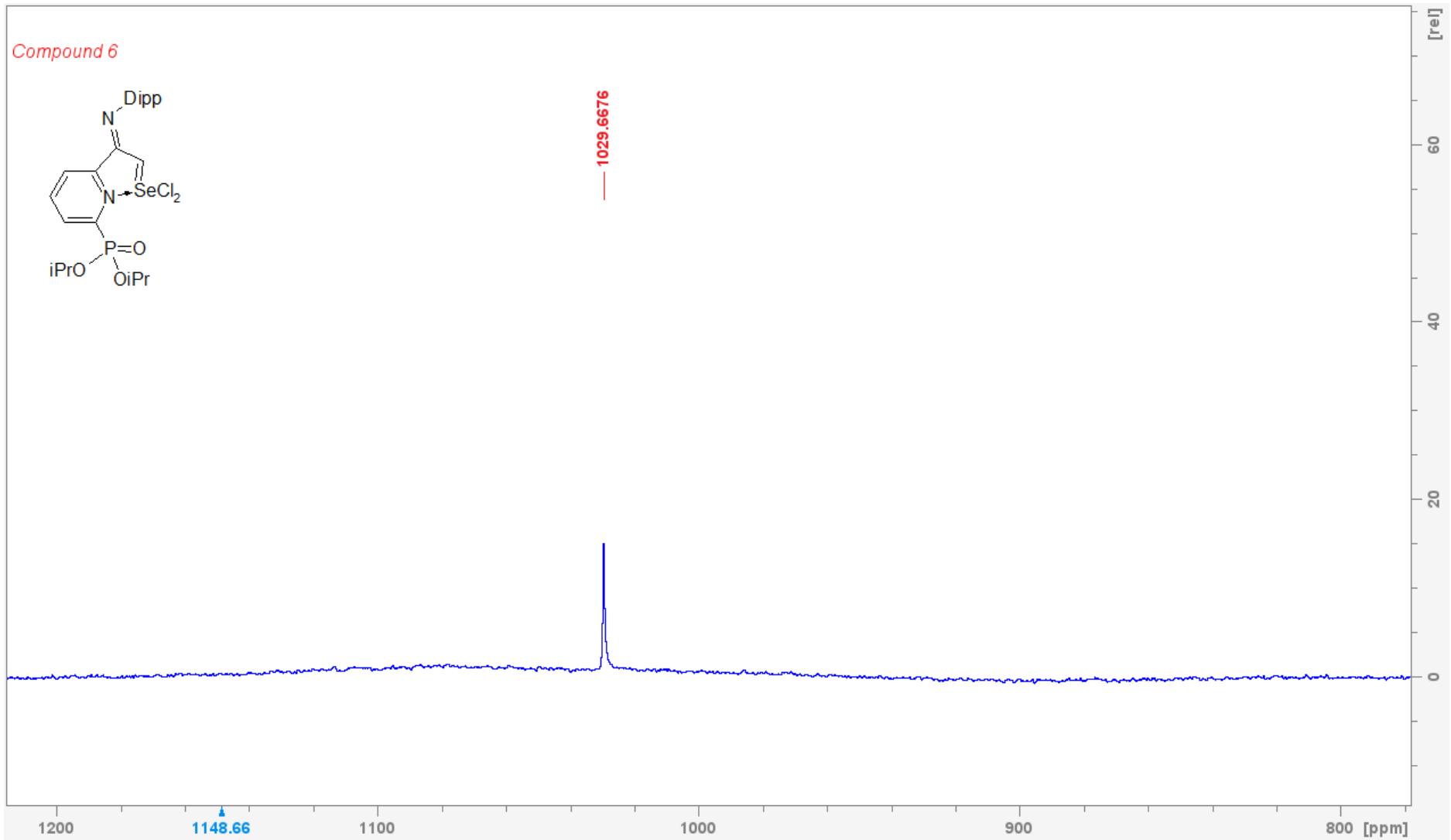
Figure S26.  $^{13}\text{C}\{\text{H}\}$  APT NMR spectrum of **6** in THF-d8 – aliphatic region



**Figure S27.**  $^{13}\text{C}\{\text{H}\}$  APT NMR spectrum of **6** in  $\text{THF-d}_8$  – aromatic region



**Figure S28.**  $^{31}\text{P}\{\text{H}\}$  NMR spectrum of **6** in THF-d8



**Figure S29.**  $^{77}\text{Se}\{\text{H}\}$  NMR spectrum of **6** in THF-d8

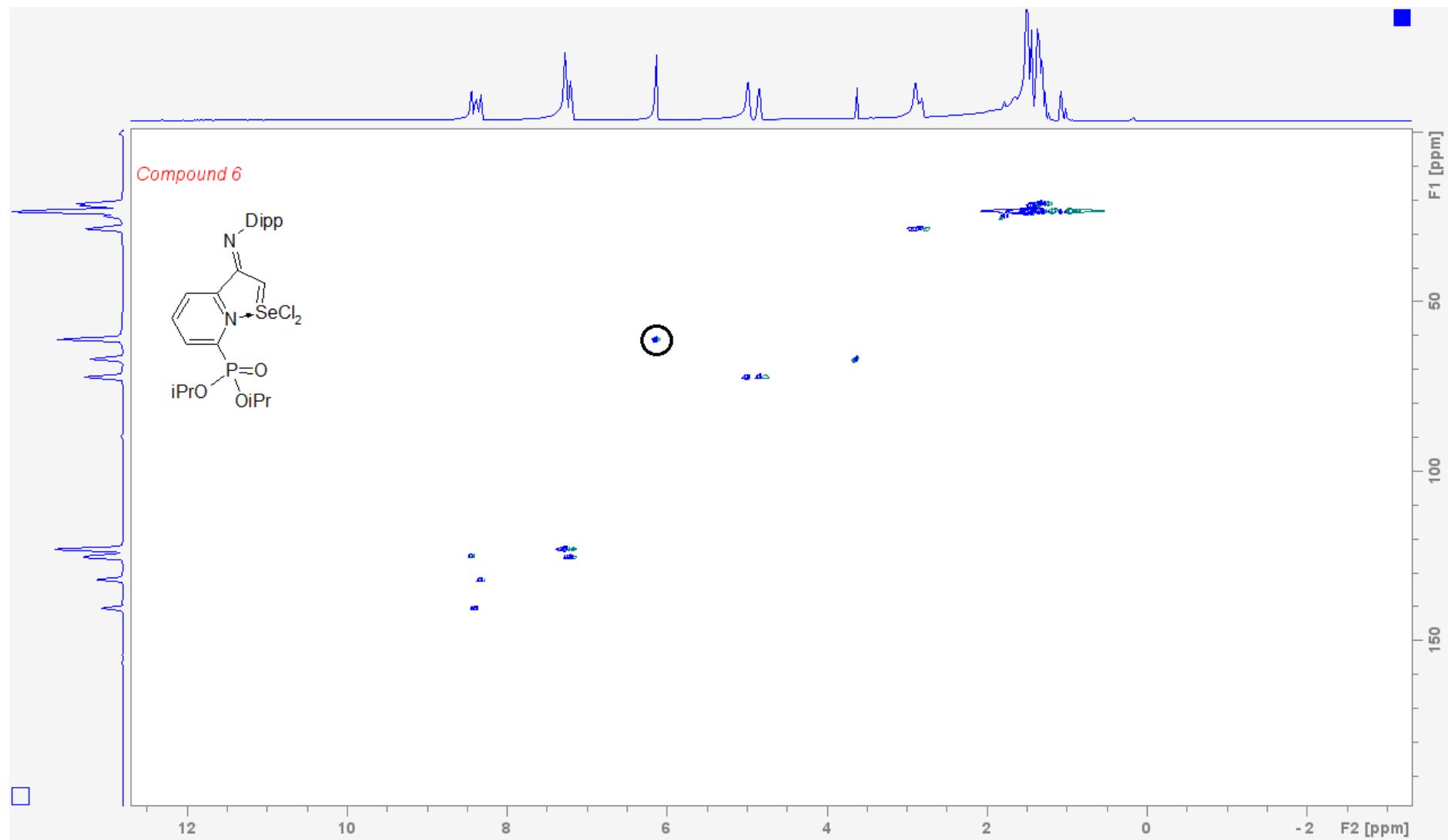
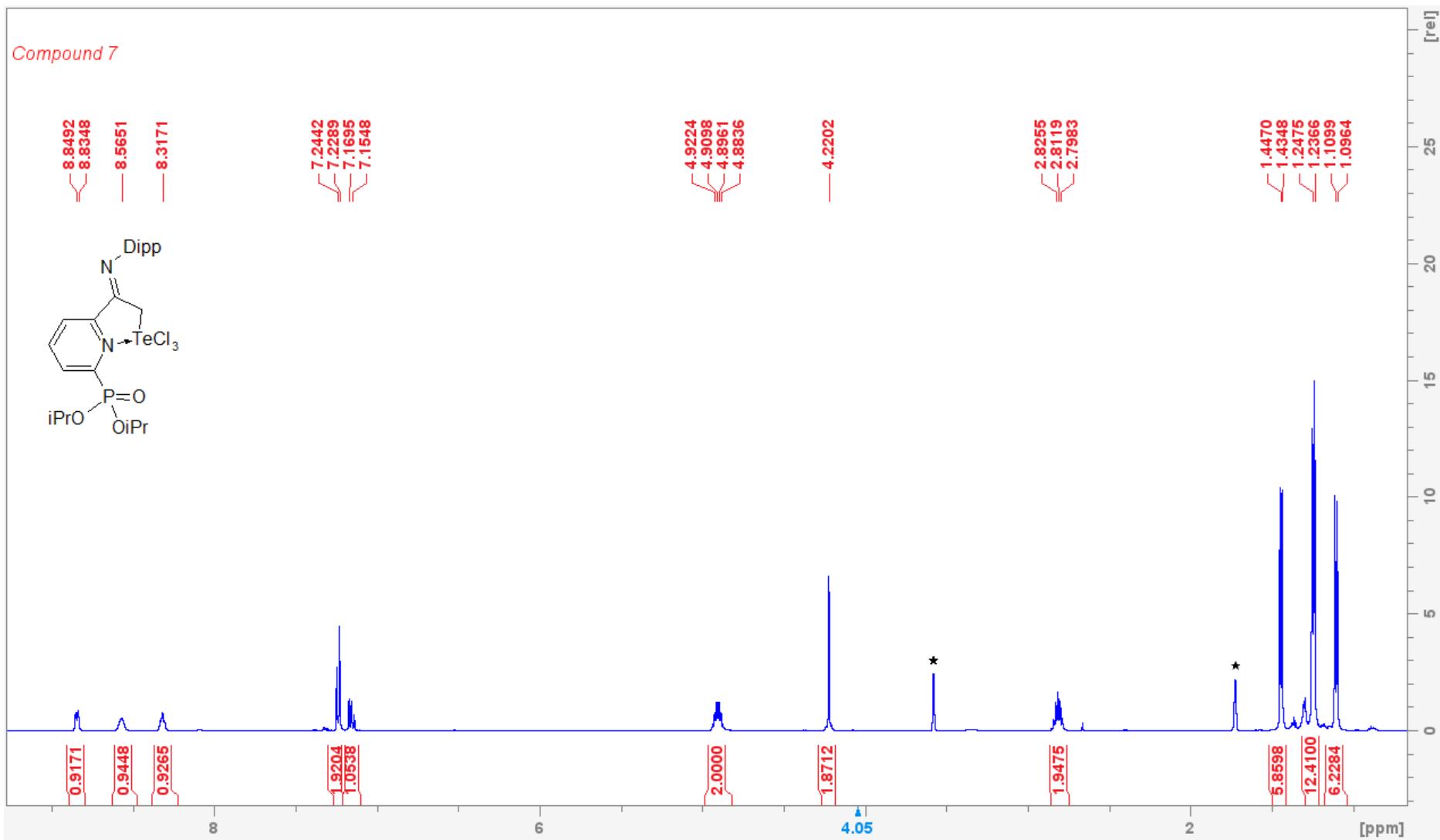


Figure S30.  $^1\text{H}$ - $^{13}\text{C}$  HSQC NMR spectrum of **6** in  $\text{THF-d}_8$



**Figure S31.**  $^1\text{H}$  NMR spectrum of **7** in THF-d8 (\* residual signal of THF)

Compound 7

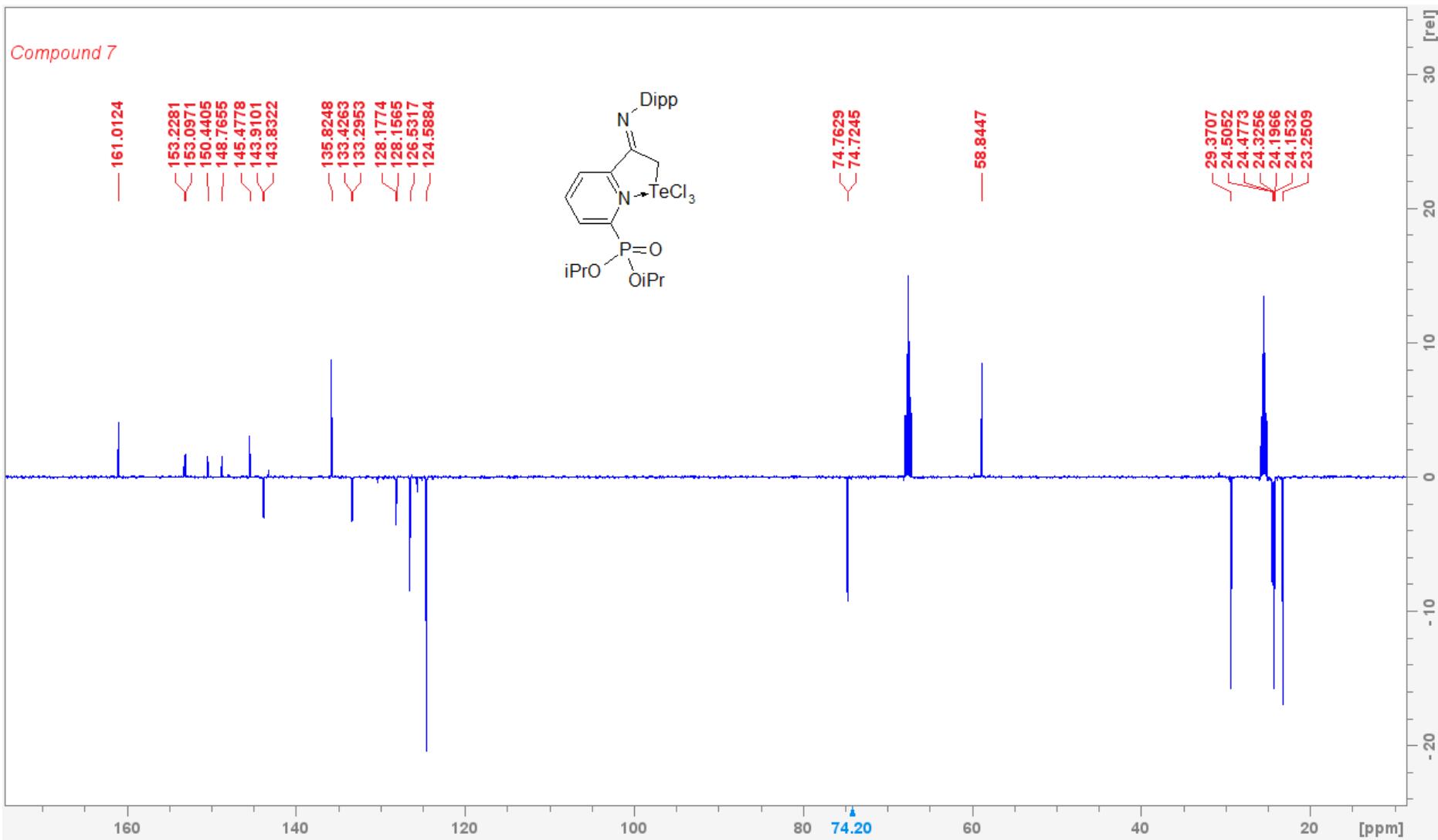
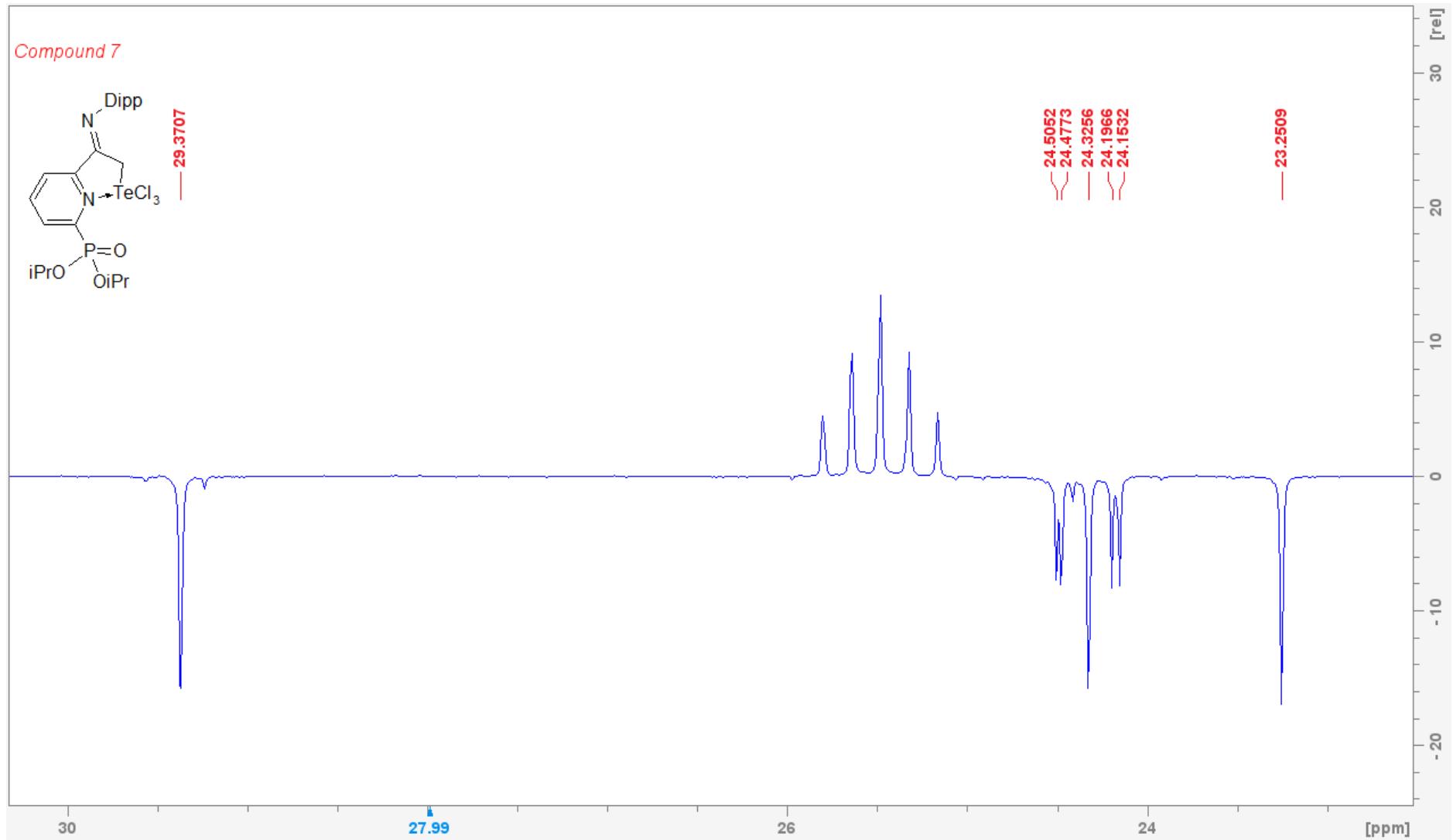
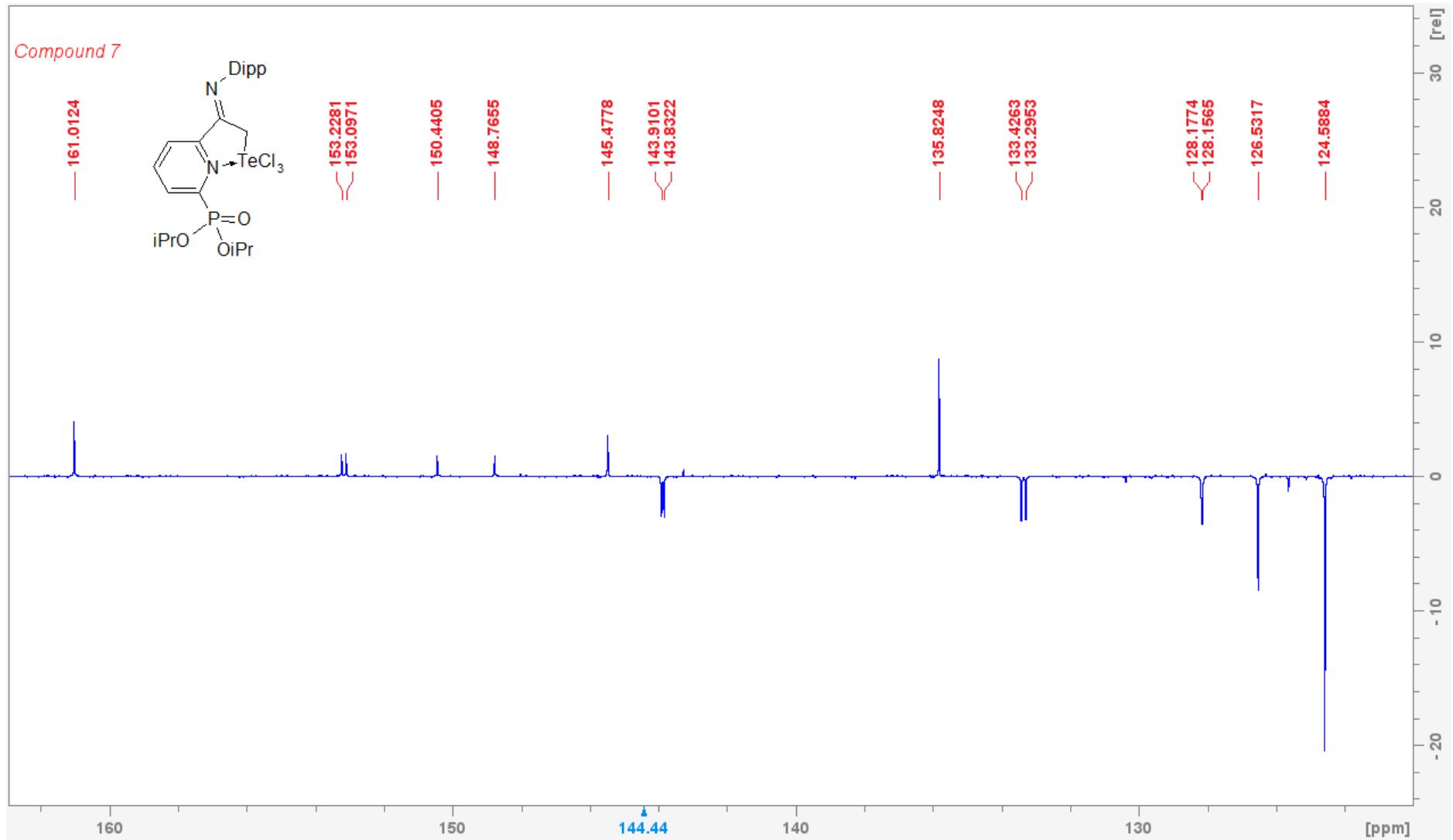


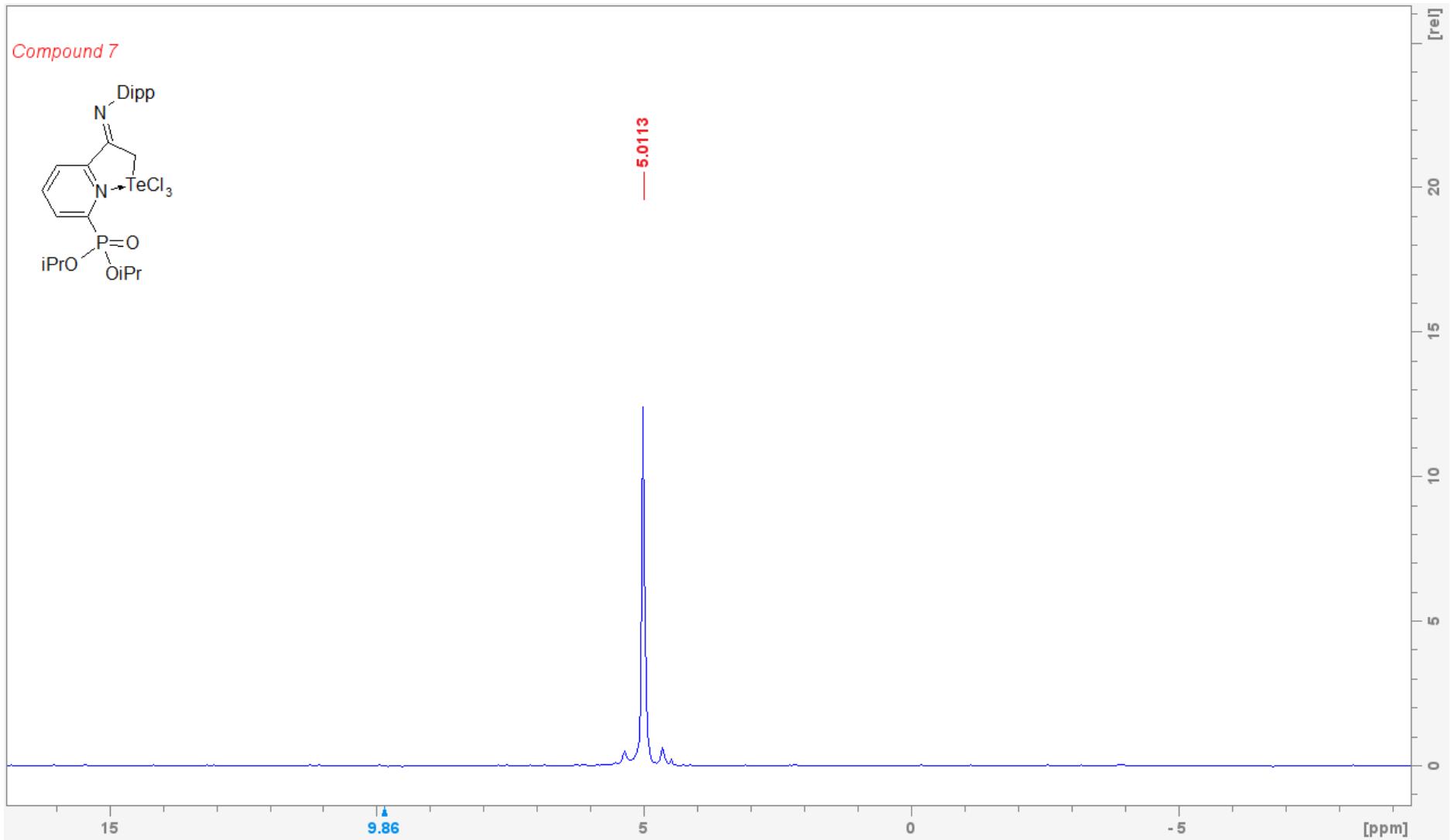
Figure S32.  $^{13}\text{C}\{\text{H}\}$  APT NMR spectrum of 7 in  $\text{THF-d}_8$



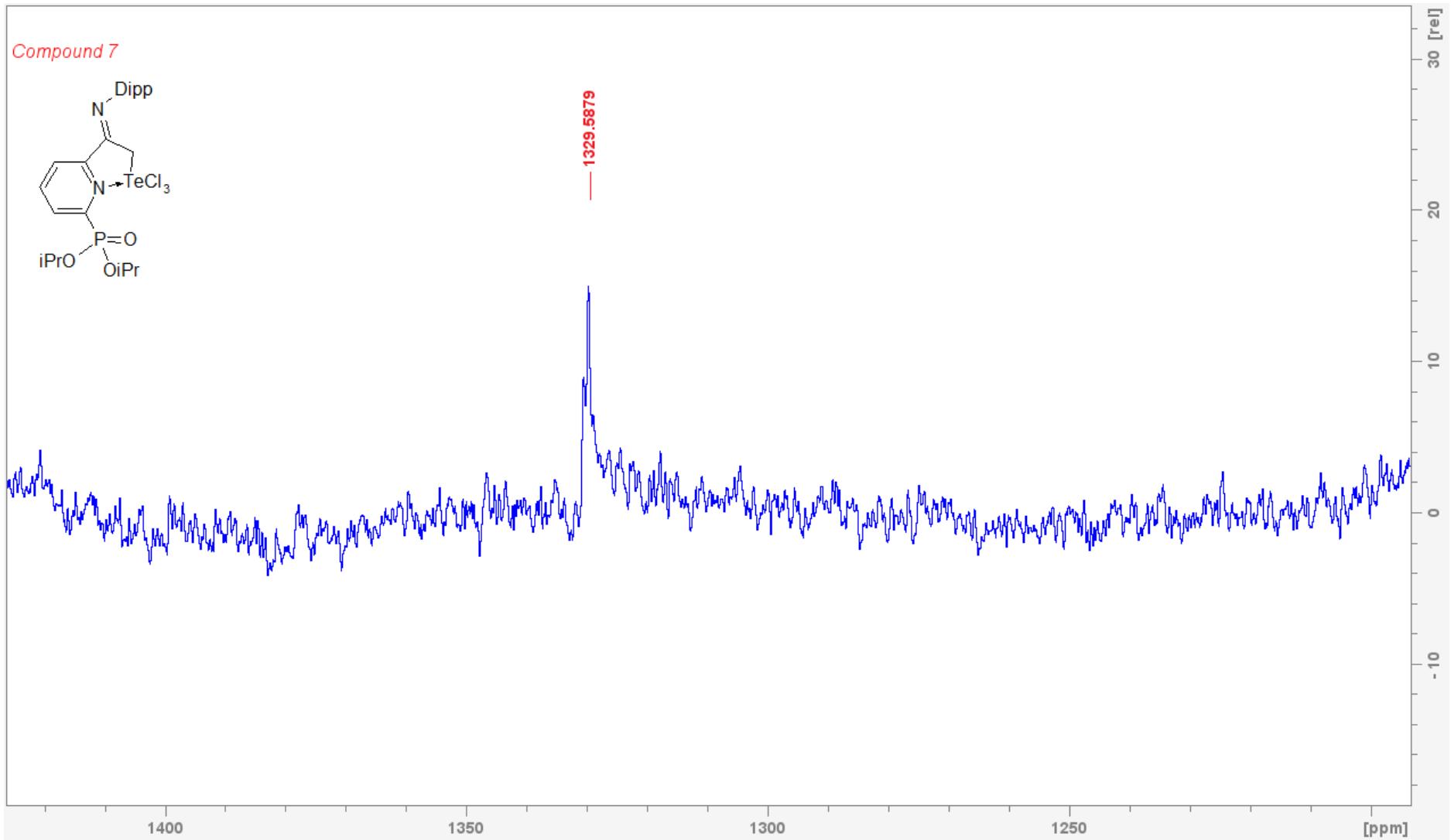
**Figure S33.**  $^{13}\text{C}\{\text{H}\}$  APT NMR spectrum of **7** in THF-d8 – aliphatic region



**Figure S34.**  $^{13}\text{C}\{\text{H}\}$  APT NMR spectrum of **7** in THF-d8 – aromatic region



**Figure S35.**  $^{31}\text{P}\{\text{H}\}$  NMR spectrum of **7** in THF-d<sub>8</sub>



**Figure S36.**  $^{125}\text{Te}\{\text{H}\}$  NMR spectrum of **7** in THF-d8

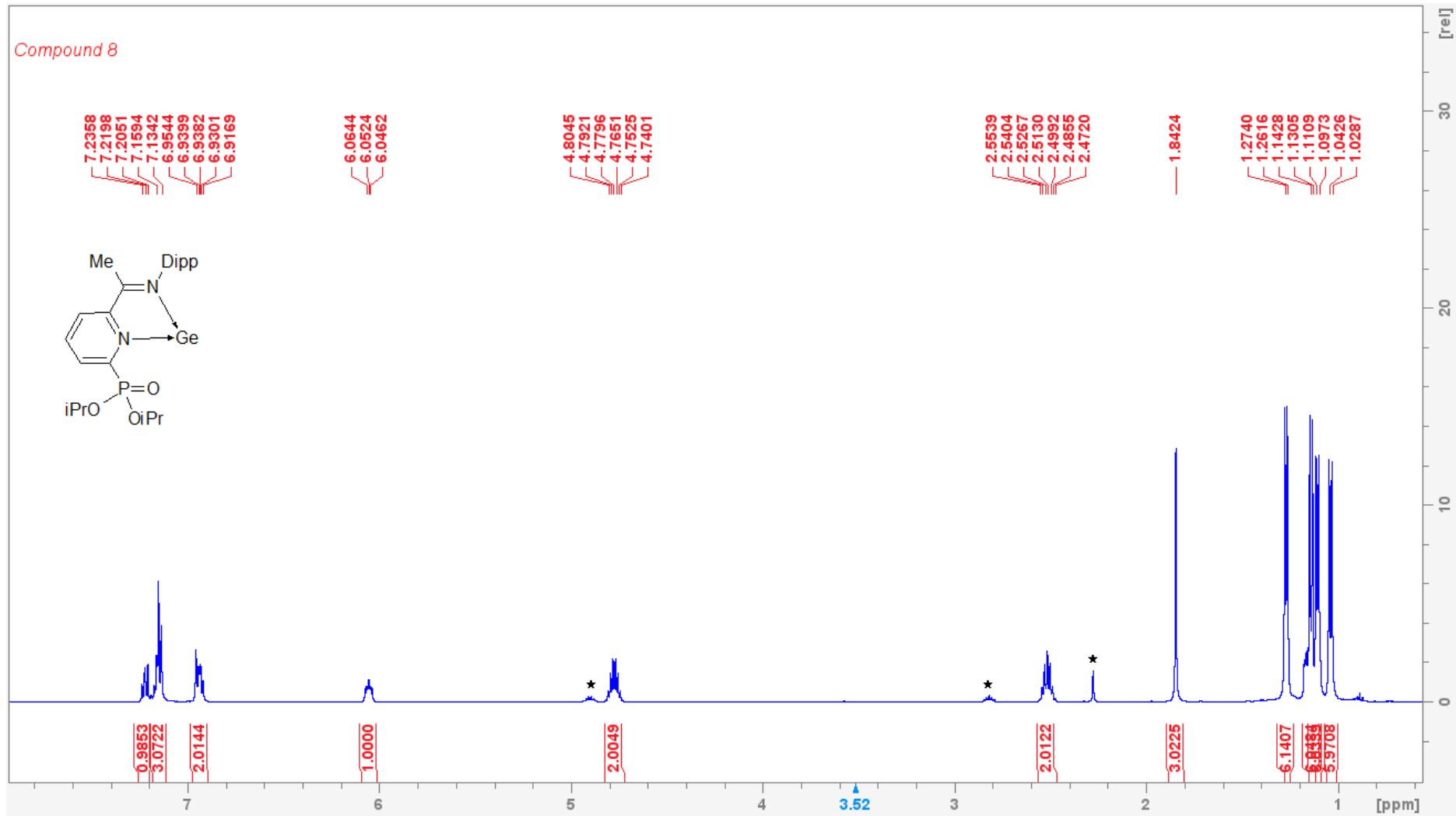
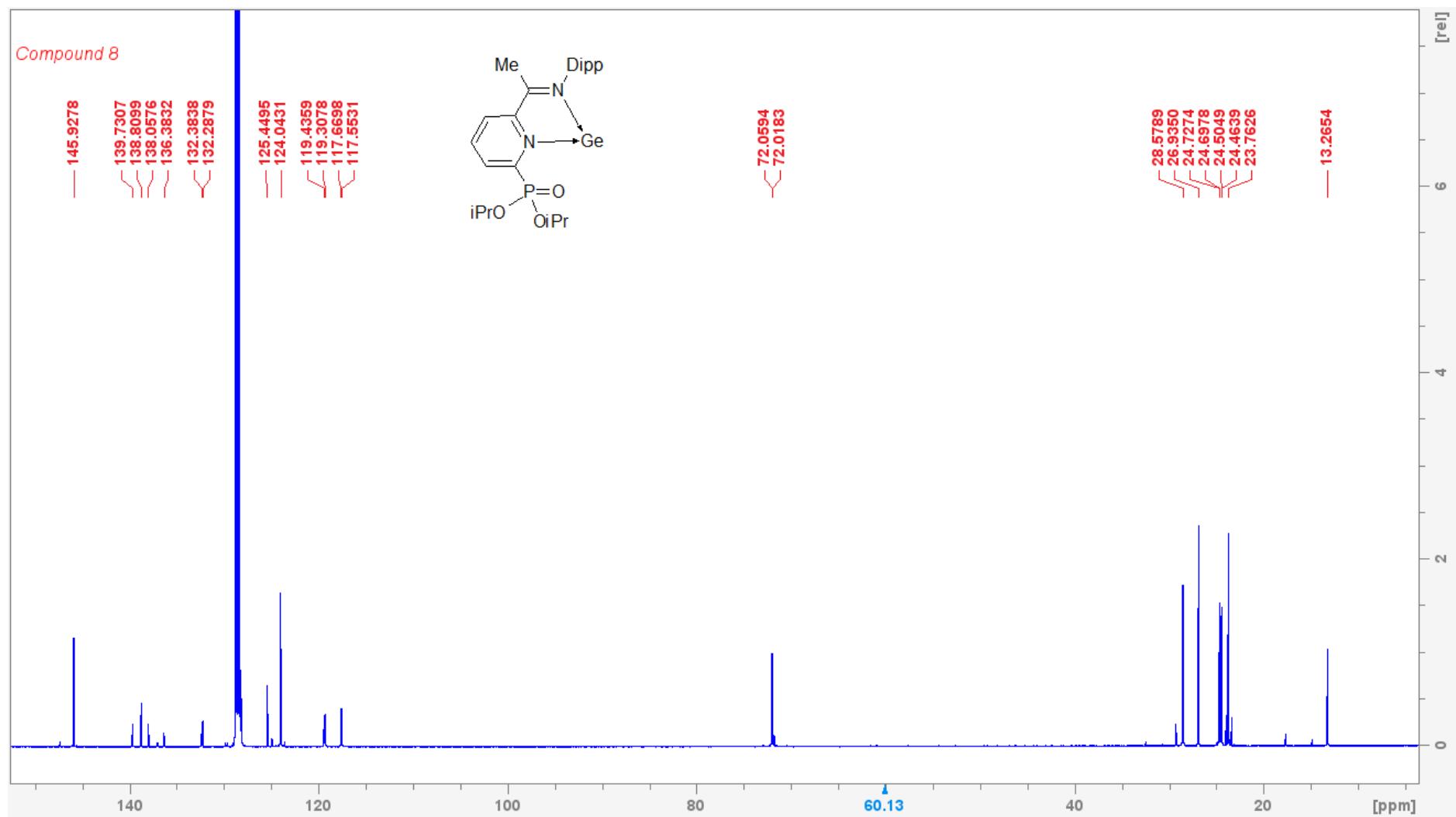
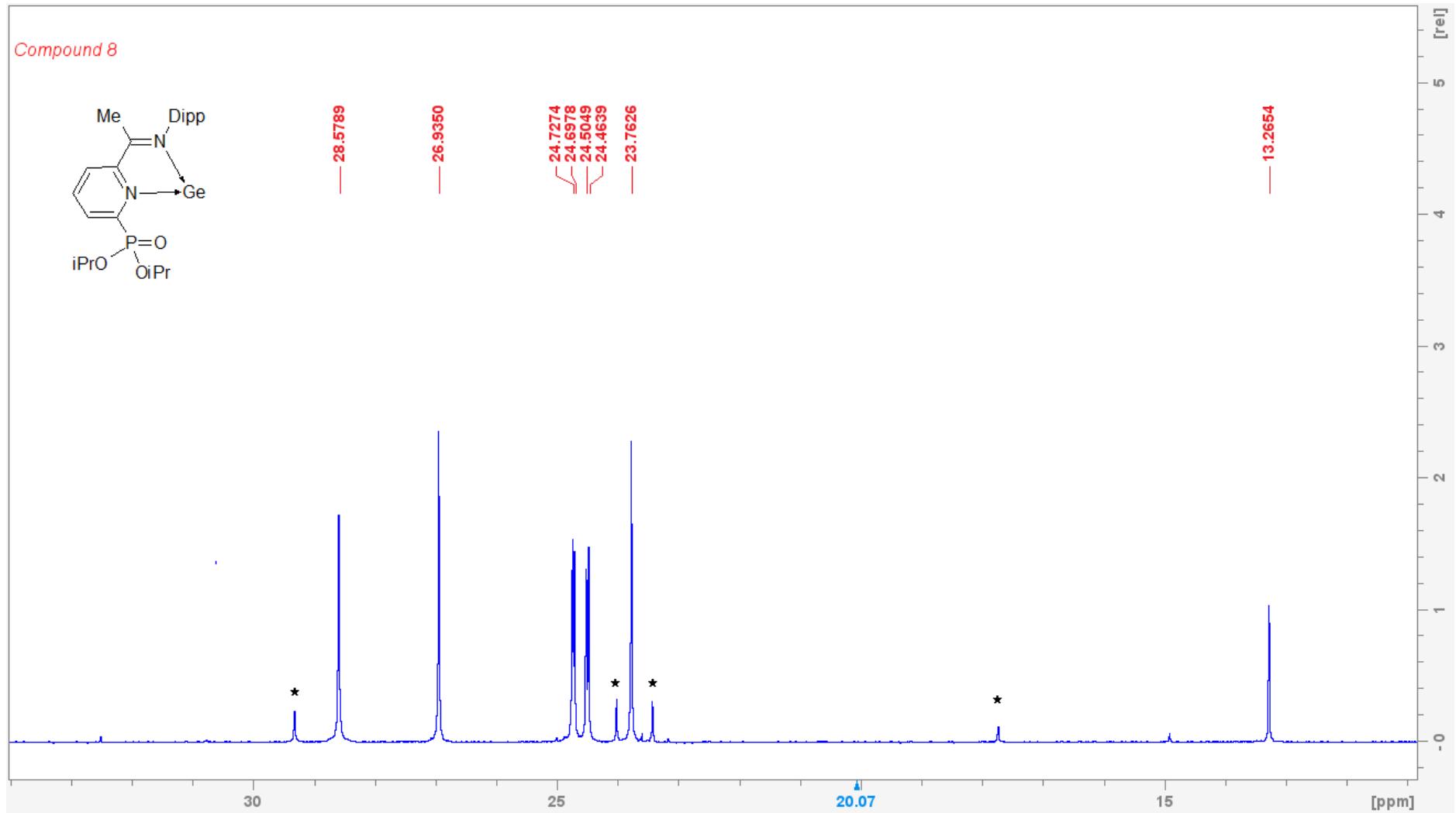


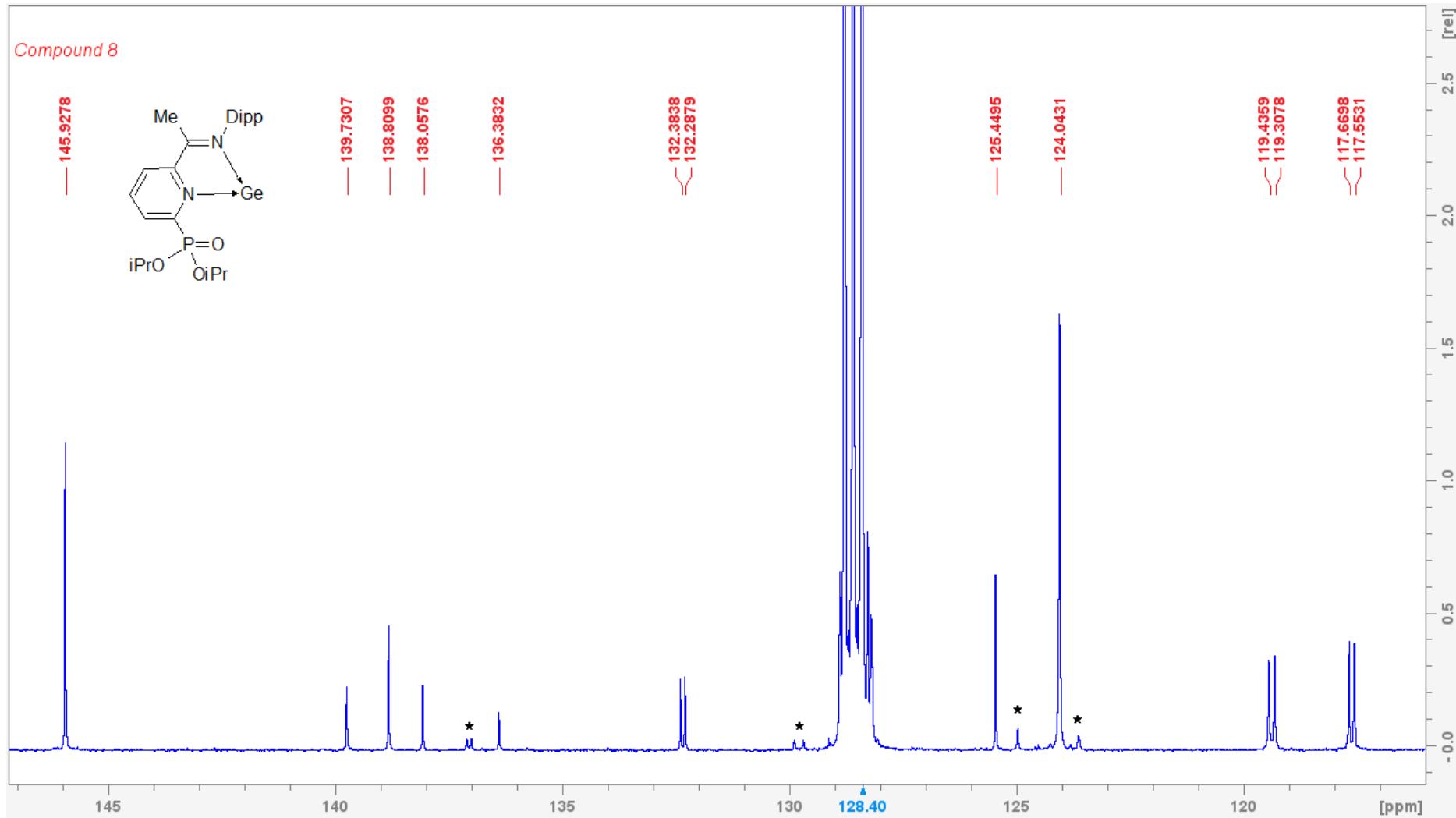
Figure S37.  $^1\text{H}$  NMR spectrum of **8** in  $\text{C}_6\text{D}_6$  (\* signal of **L**)



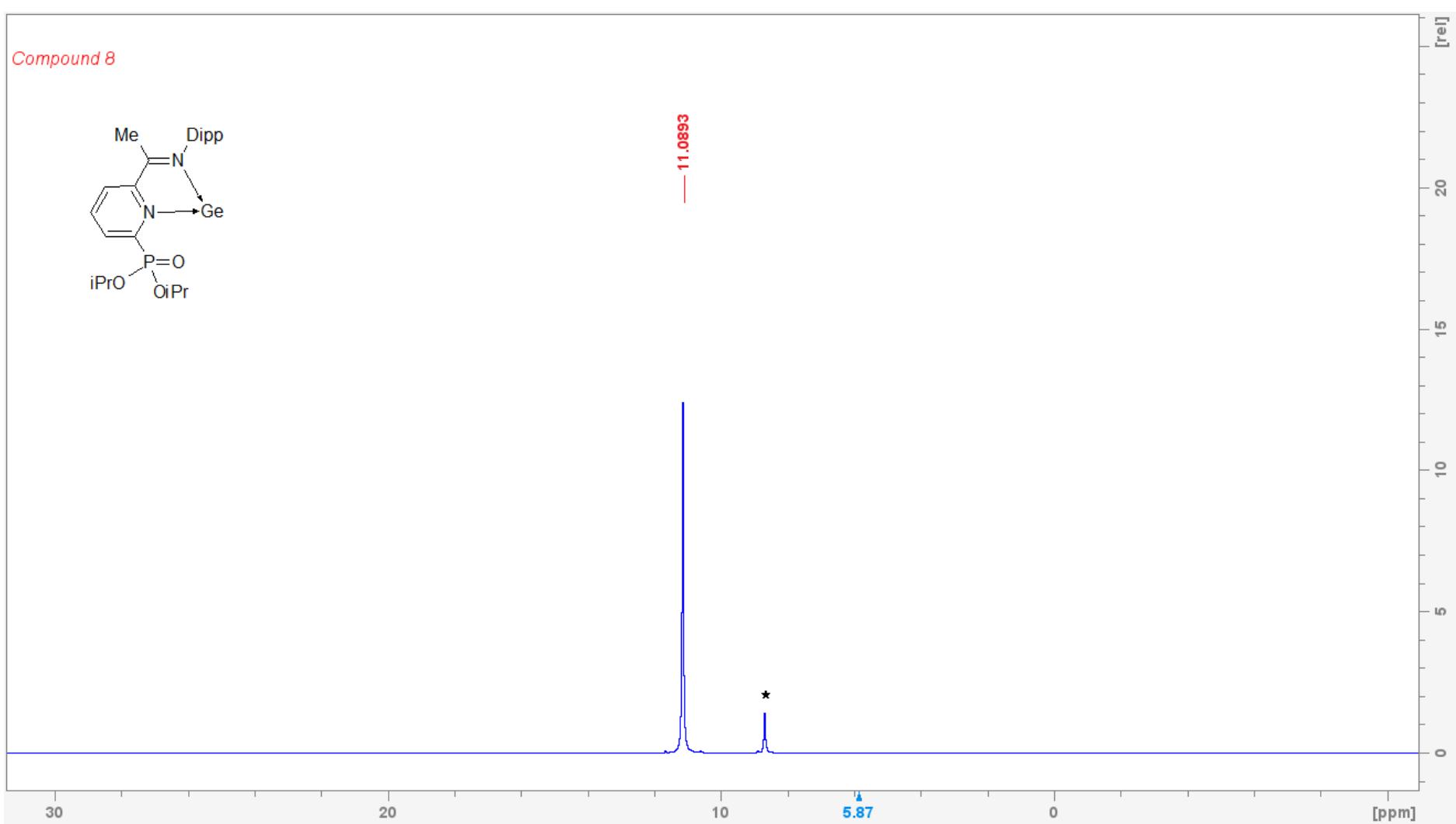
**Figure S38.**  $^{13}\text{C}\{\text{H}\}$  NMR spectrum of **8** in  $\text{C}_6\text{D}_6$



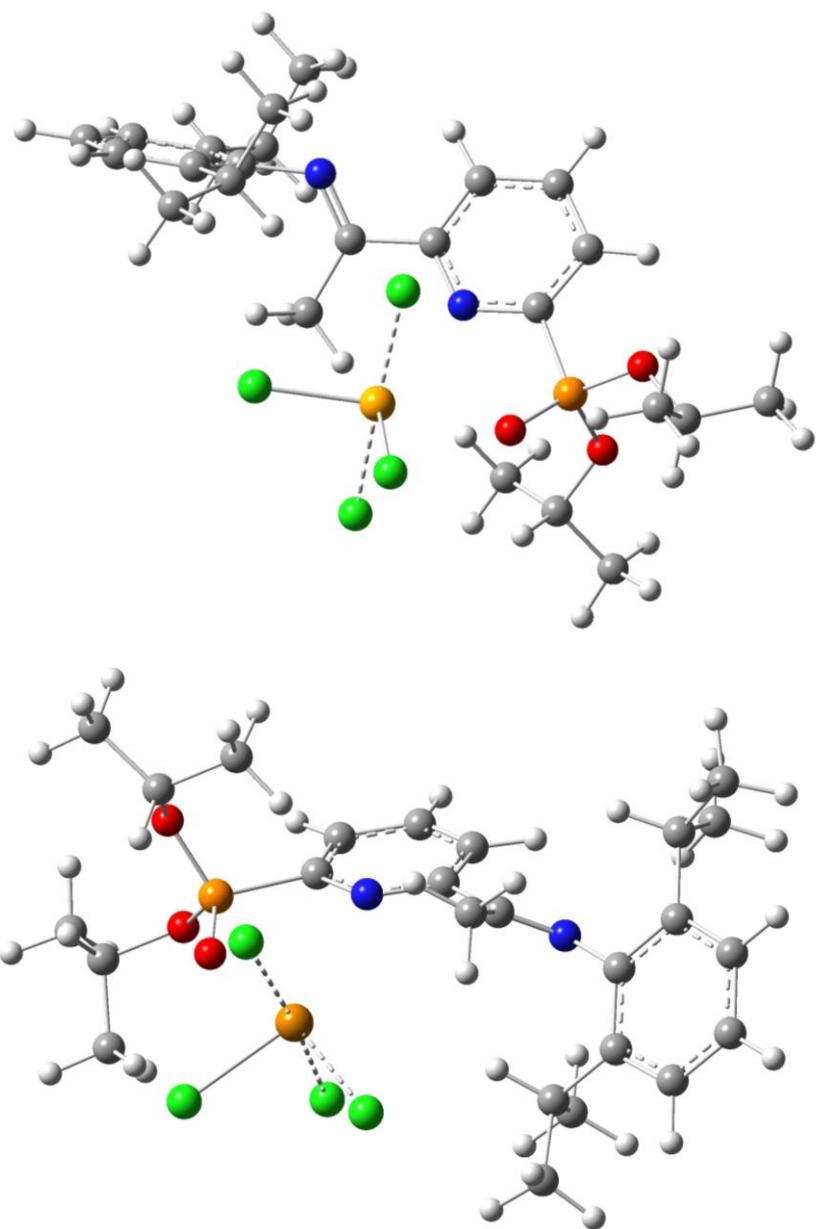
**Figure S39.**  $^{13}\text{C}\{^1\text{H}\}$  NMR spectrum of **8** in  $\text{C}_6\text{D}_6$  – aliphatic region (\* signal of **L**)



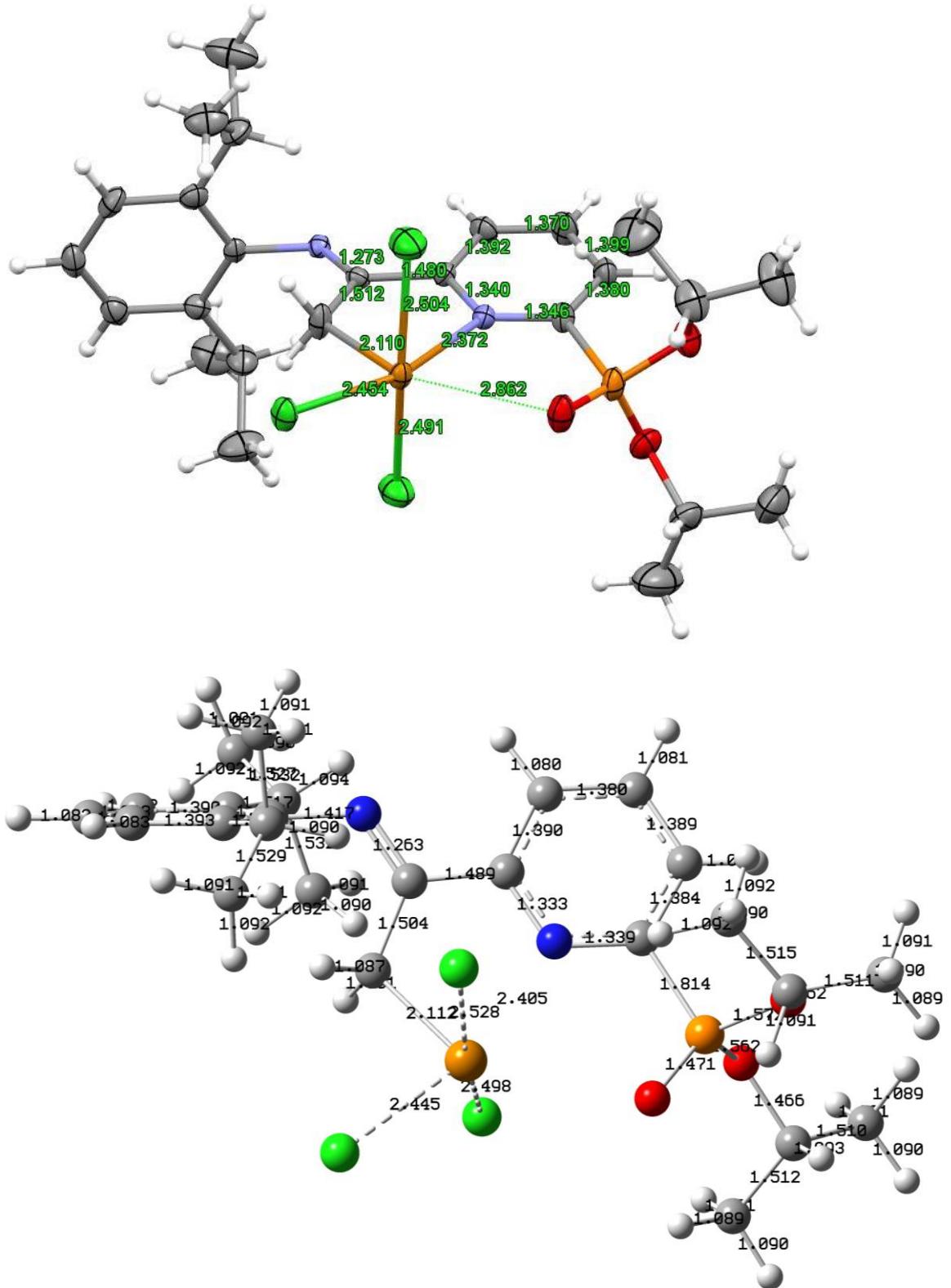
**Figure S40.**  $^{13}\text{C}\{\text{H}\}$  NMR spectrum of **8** in  $\text{C}_6\text{D}_6$  – aromatic region (\* signal of **L**)



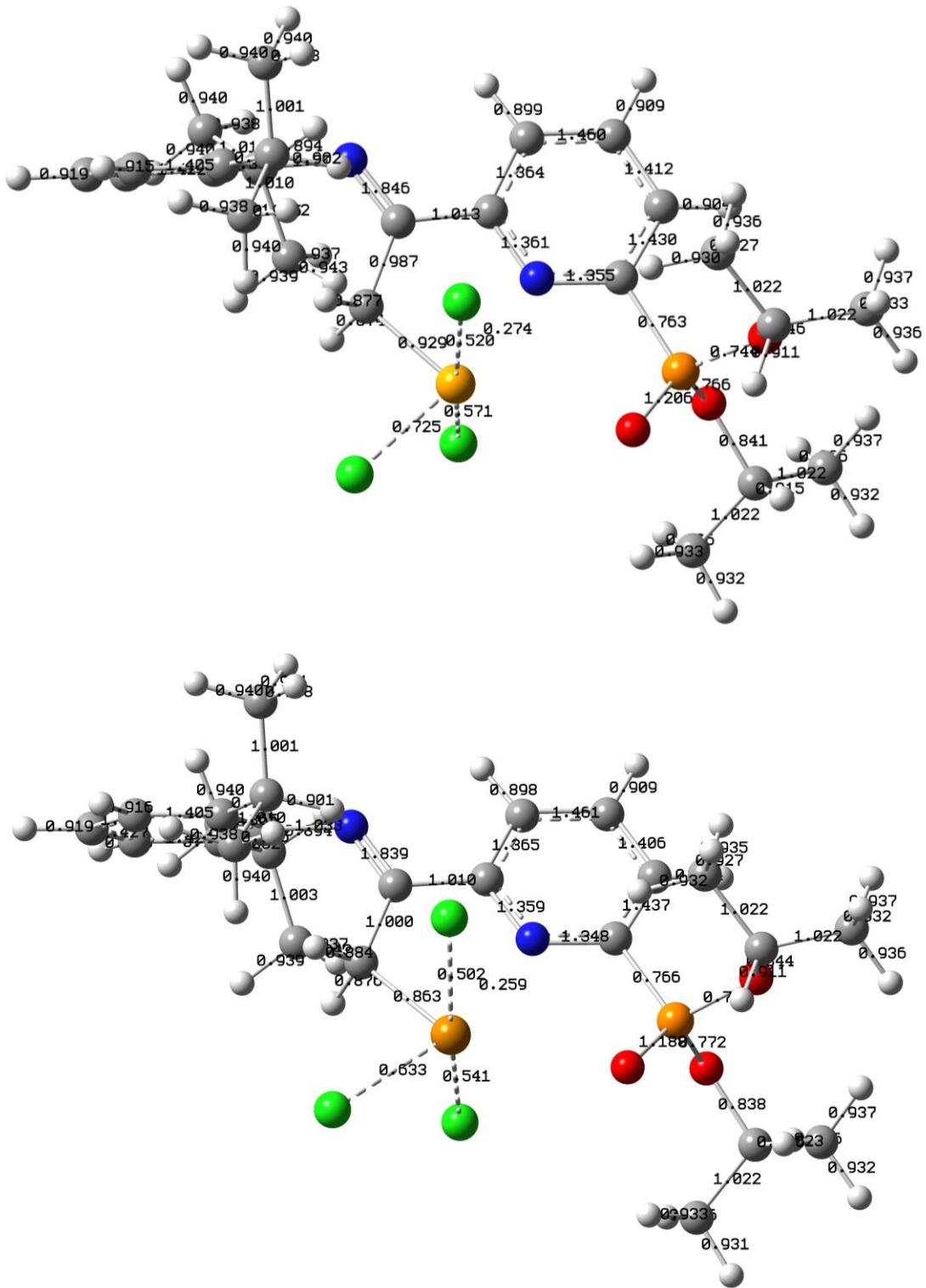
**Figure S41.**  $^{31}\text{P}\{\text{H}\}$  NMR spectrum of **8** in  $\text{C}_6\text{D}_6$  (\* signal of **L**).



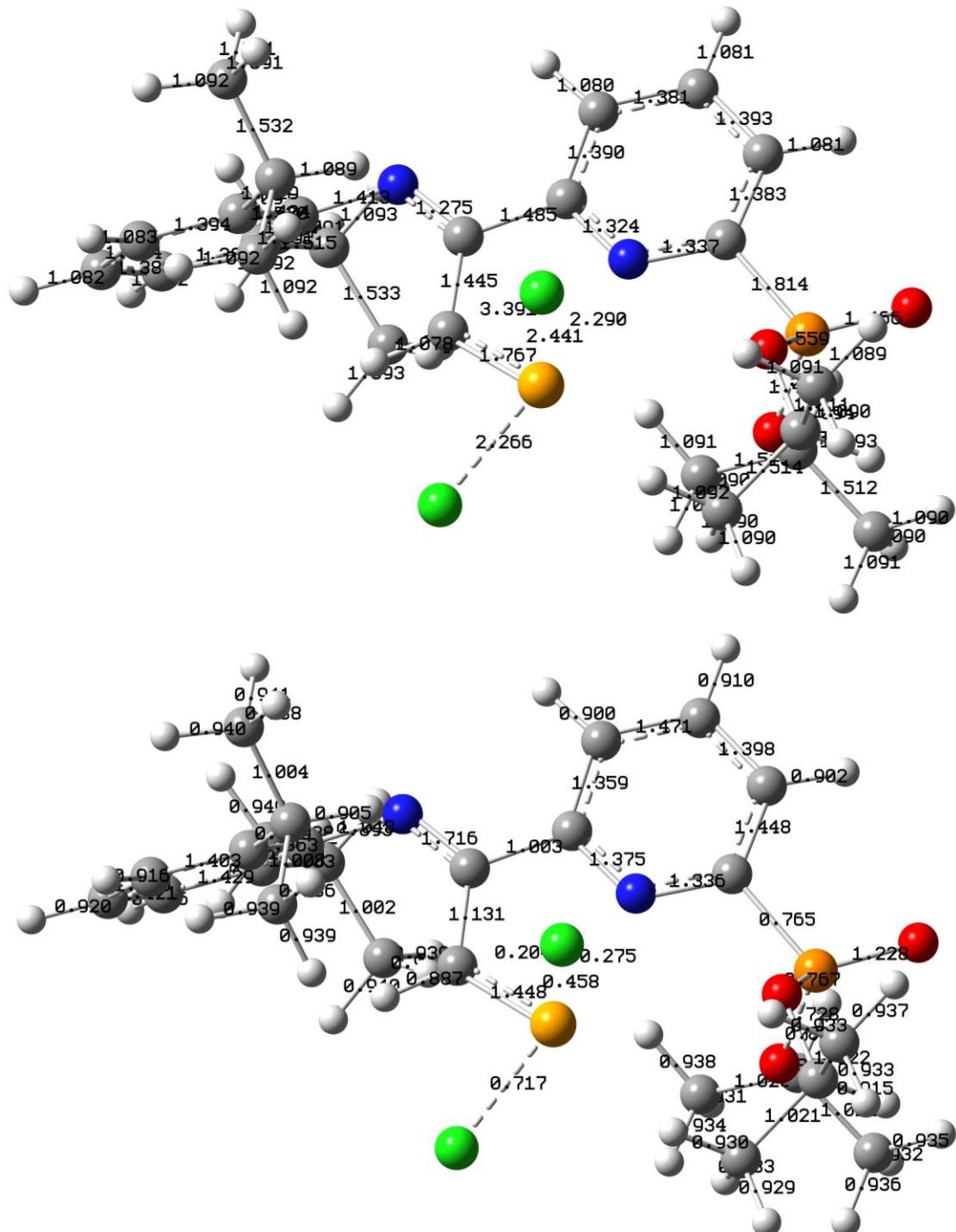
**Figure S42.** Comparison of optimized structures of **L**→SeCl<sub>4</sub> (top) and **L**→TeCl<sub>4</sub> (bottom).



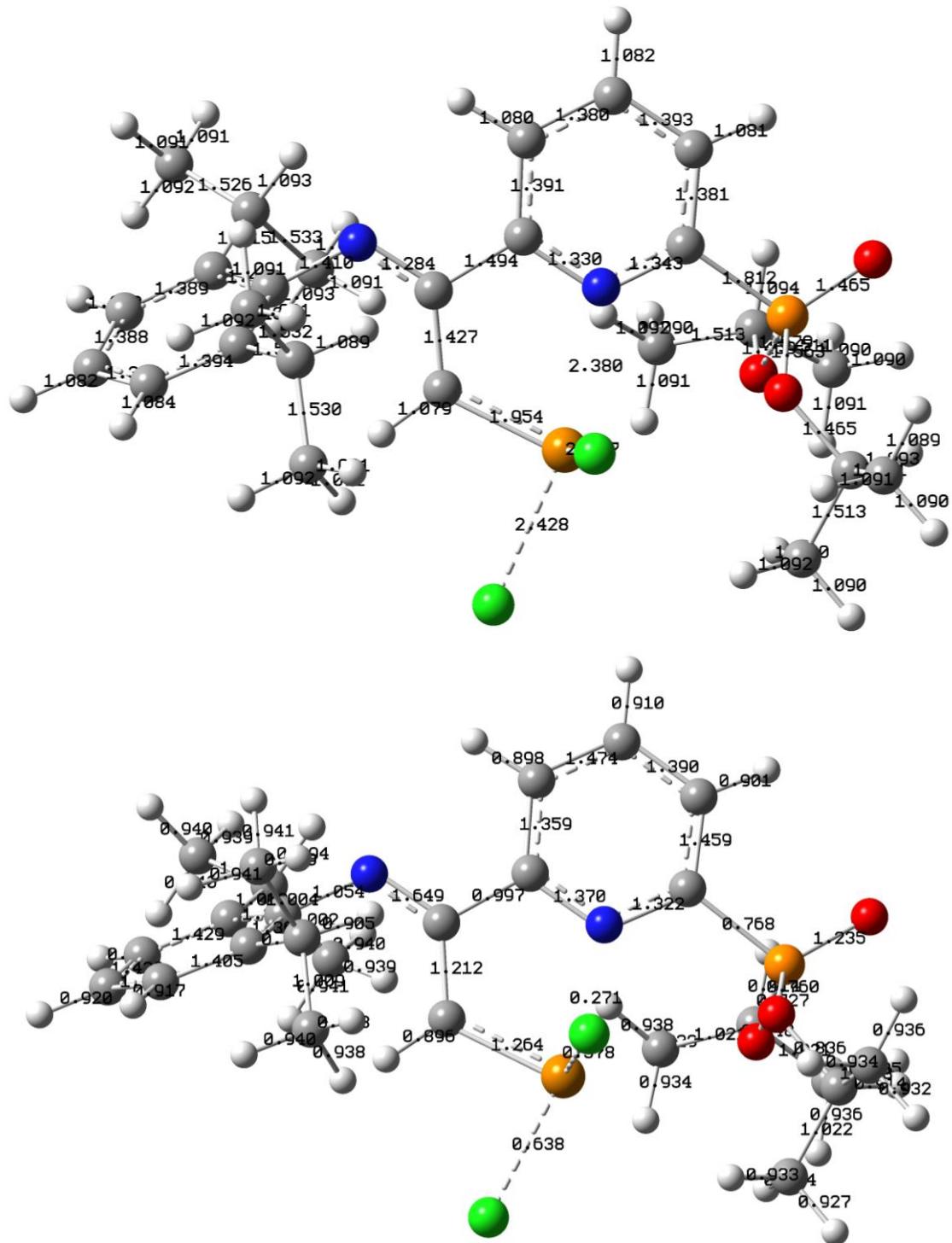
**Figure S43.** Comparison of molecular structures of **7**, XRD (top) and calculated (bottom, with Te...O 2.913) – bond lengths are given in Å.



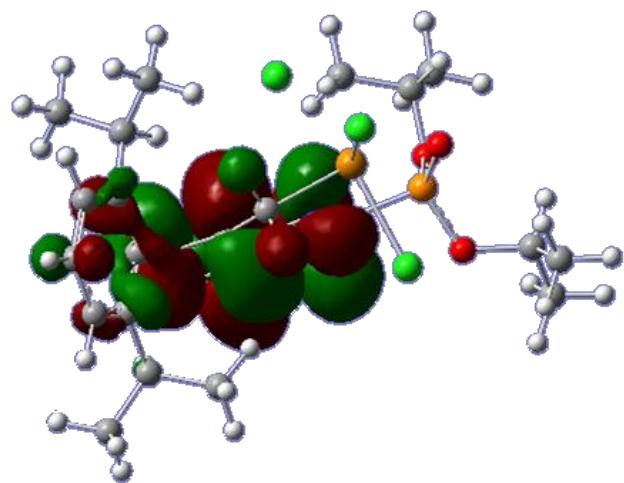
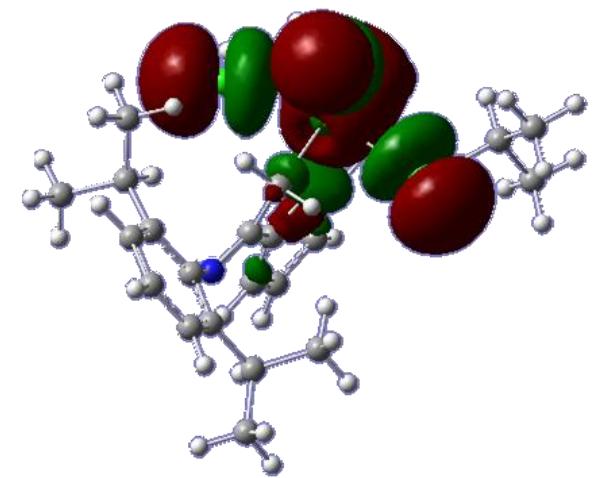
**Figure S44.** Comparison of optimized structures of **7'**(Se) (top) and **7** (bottom) including WBIs.



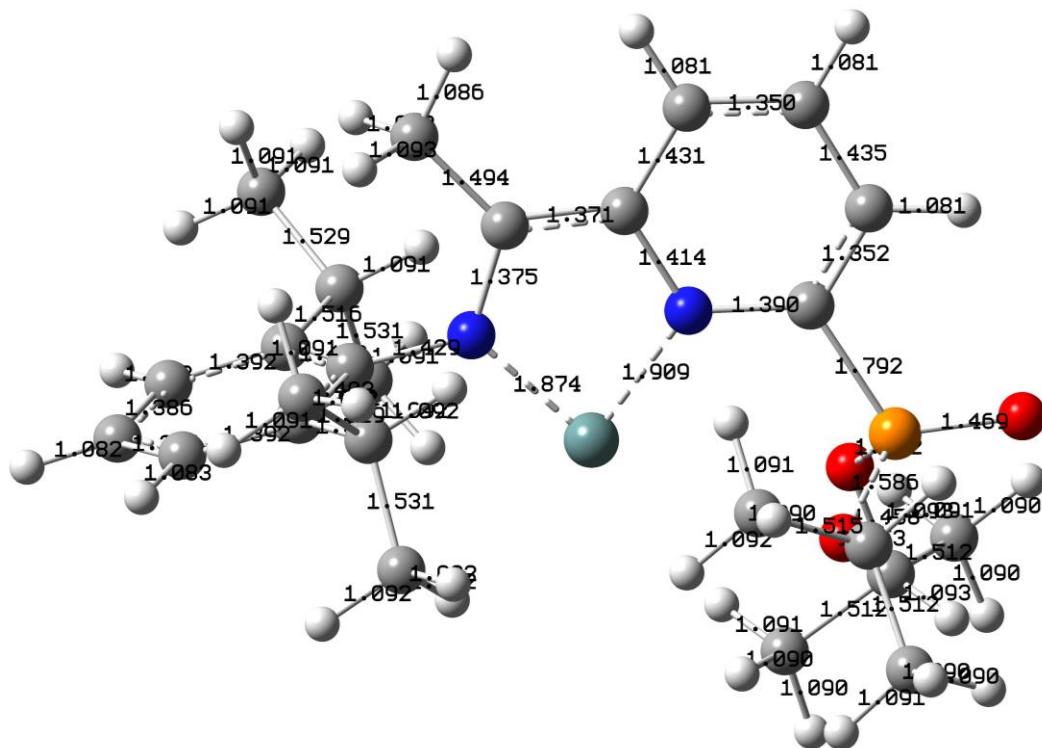
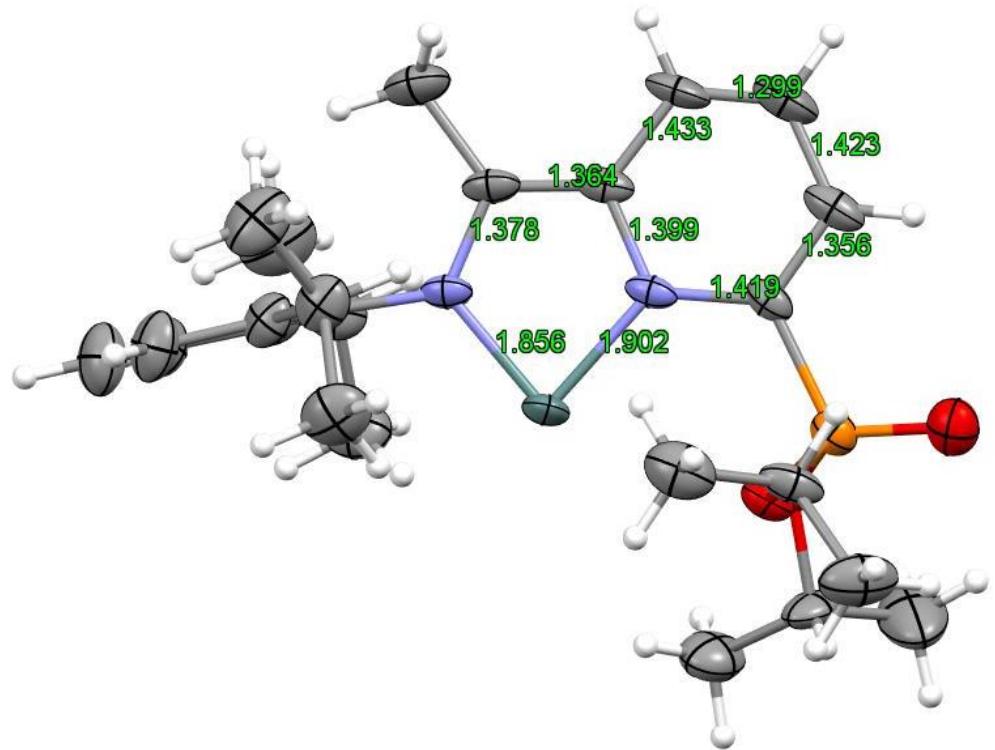
**Figure S45.** Optimized structure of **6** (top: distances in Å, bottom: WBIs).



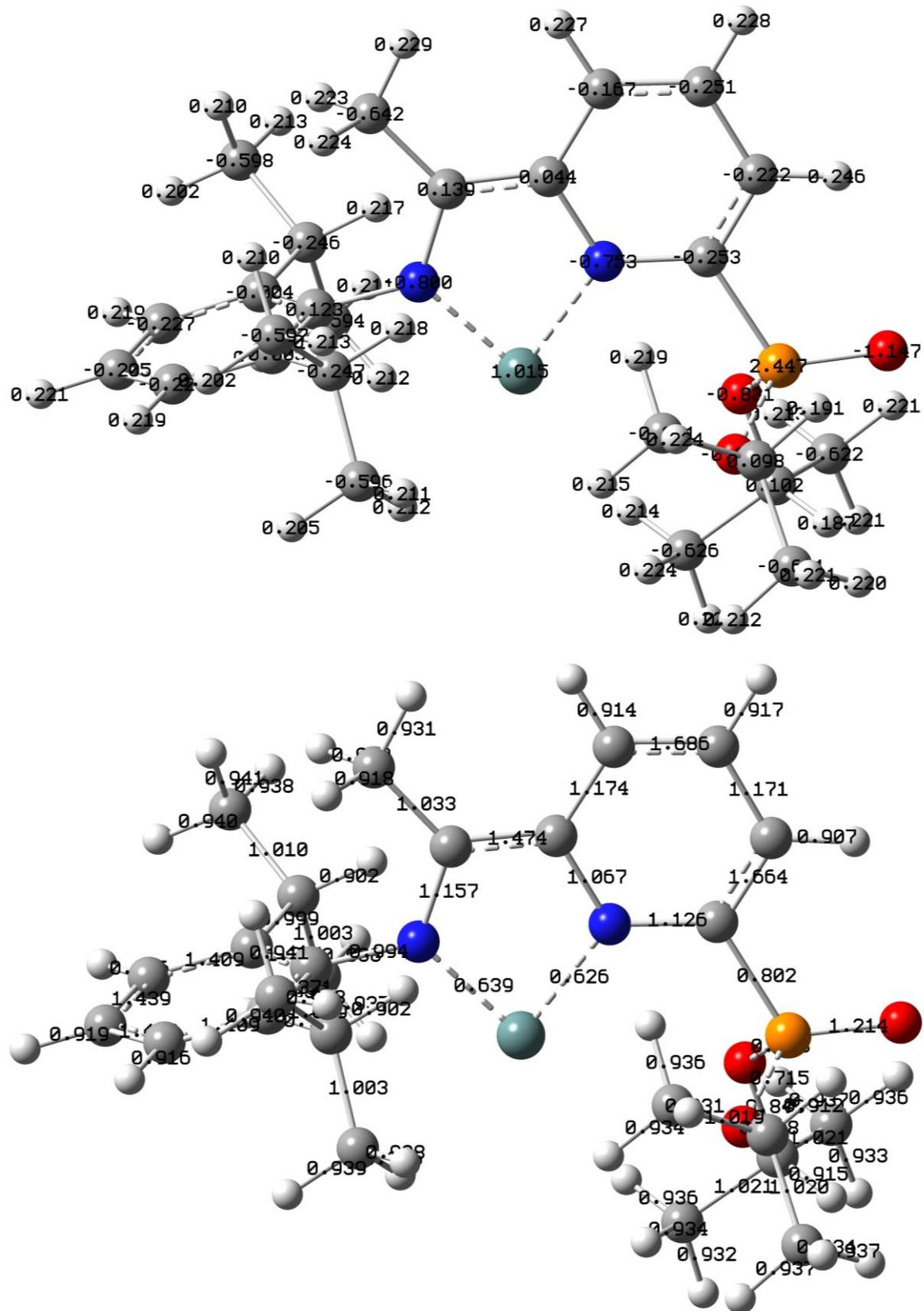
**Figure S46.** Optimized structure of **6'(Te)** (top: distances in Å, bottom: WBIs).

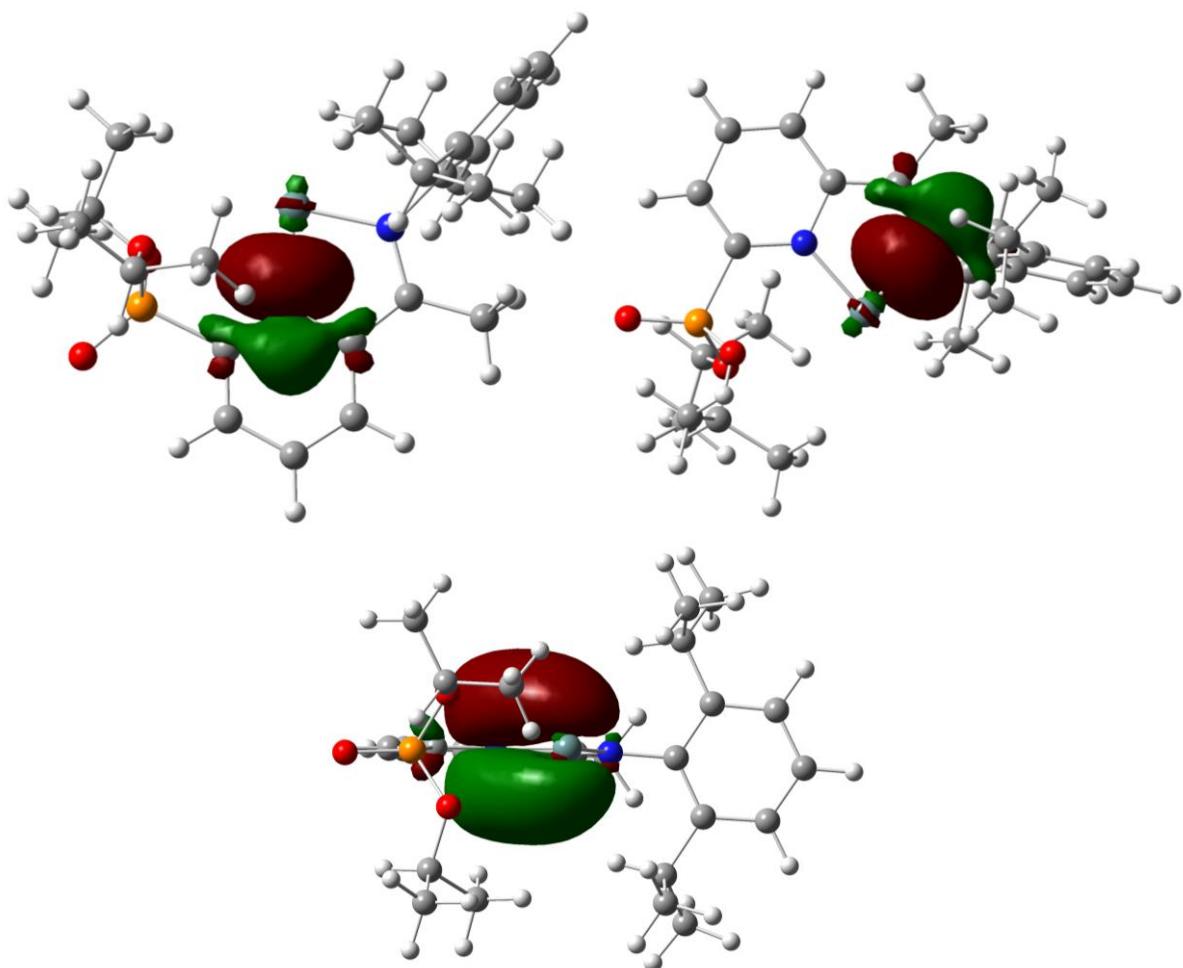


**Figure S47.** Visualization of HOMO (top) and LUMO (bottom) in **7** and **7'**(Se).

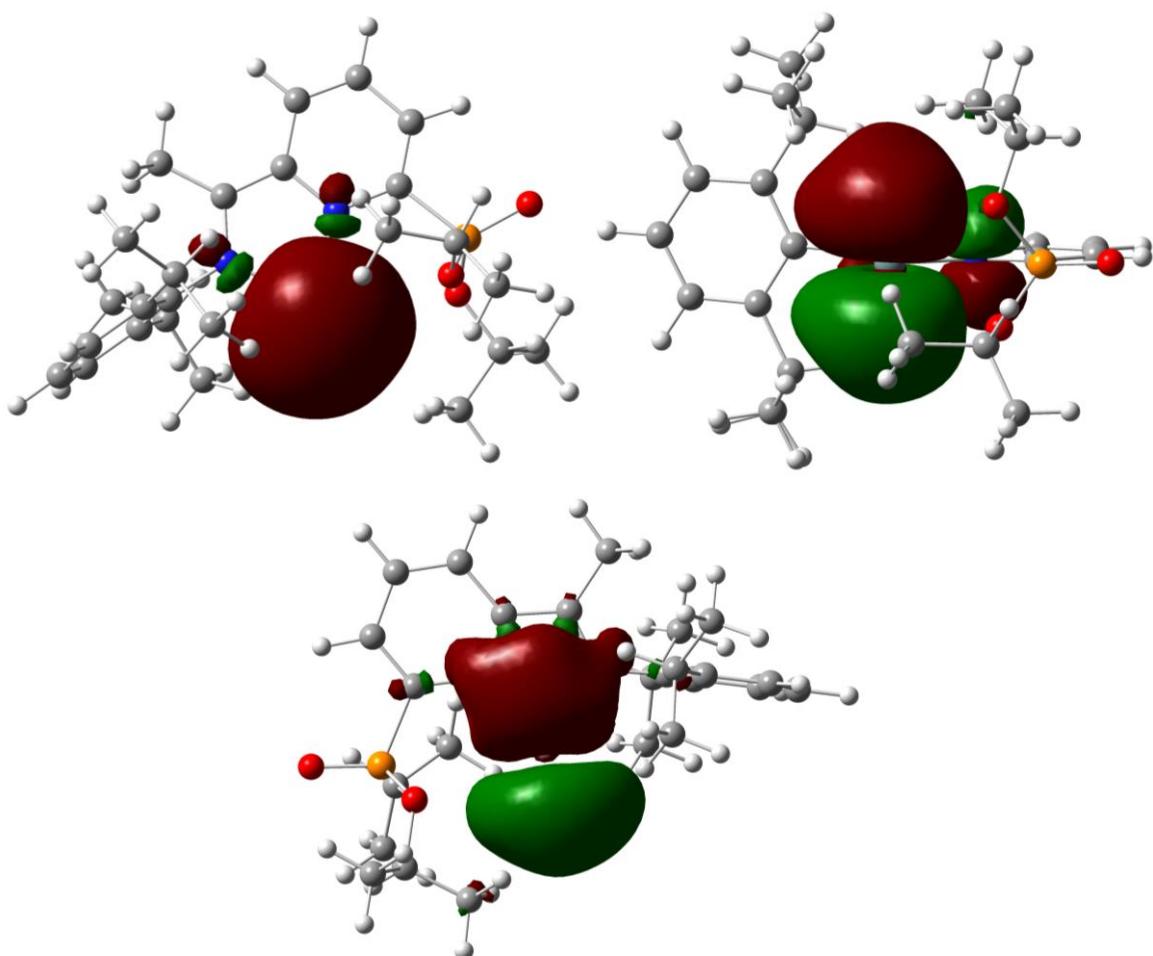


**Figure S48.** Comparison of molecular structures of **8**, XRD (top) and calculated (bottom) – bond lengths are given in Å.





**Figure S50.** Visualization of HOMO-14 (top-left), HOMO-13 (top-right) and HOMO-5 (bottom) orbitals involved in connection of Ge atom and ligand in **8**.



**Figure S51.** Visualization of HOMO-10 (top-left, lone electron pair on Ge), LUMO (top-right, gap 7.38 eV) and LUMO+8 (bottom) orbitals situated close to Ge atom in **8**.

**Table S7.** Gibbs' free energy in Hartrees

<b>compound</b>	<b>Gibbs' free energy</b>	<b>complex</b>	<b>Gibbs' free energy</b>
HCl	-460.811943	L <sup>NO</sup> SeCl <sub>4</sub>	-5893.407581
SeCl <sub>4</sub>	-4242.388935	L <sup>CO</sup> SeCl <sub>4</sub>	-5893.404968
TeCl <sub>4</sub>	-2108.83124	L <sup>CO</sup> SeCl <sub>3</sub>	-5432.620862
L <sup>CO(ROT)</sup>	-1651.011969	L <sup>CO</sup> SeCl <sub>2</sub>	-4971.767615
L <sup>CO</sup>	-1651.010611	L <sup>CO(ROT)</sup> SeCl <sub>2</sub>	-4971.789719
L <sup>NO(ROT)</sup>	-1651.005528	L <sup>NO</sup> TeCl <sub>4</sub>	-3759.862807
L <sup>NO</sup>	-1651.00387	L <sup>CO</sup> TeCl <sub>4</sub>	-3759.858043
		L <sup>CO</sup> TeCl <sub>3</sub>	-3299.048927
		L <sup>CO</sup> TeCl <sub>2</sub>	-2838.178768
		L <sup>CO(ROT)</sup> TeCl <sub>2</sub>	-2838.194775