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**Supporting information for article:**

**First heterobimetallic Ag<sup>I</sup>-Co<sup>III</sup> coordination compound with both bridging and terminal -NO<sub>2</sub> coordination modes: synthesis, characterization, structural and computational studies of (PPh<sub>3</sub>)<sub>2</sub>Ag<sup>I</sup>-(μ-κ<sup>2</sup>O,O';κN-NO<sub>2</sub>-Co<sup>III</sup>(DMGH)<sub>2</sub>(κN-NO<sub>2</sub>)**

**Reza Kia, Shiva Batmanghelich and Paul R. Raithby**

## Supporting Information

### First Heterobimetallic Ag<sup>I</sup>-Co<sup>III</sup> Coordination Compound with both Bridging and Terminal -NO<sub>2</sub> Coordination Modes: Synthesis, Characterization, Structural and Computational Studies of (PPh<sub>3</sub>)<sub>2</sub>Ag<sup>I</sup>-(κ<sup>2</sup>-O,O'-κ<sup>1</sup>-N)NO<sub>2</sub>-Co<sup>III</sup>(DMGH)<sub>2</sub>(κ<sup>1</sup>-NO<sub>2</sub>)

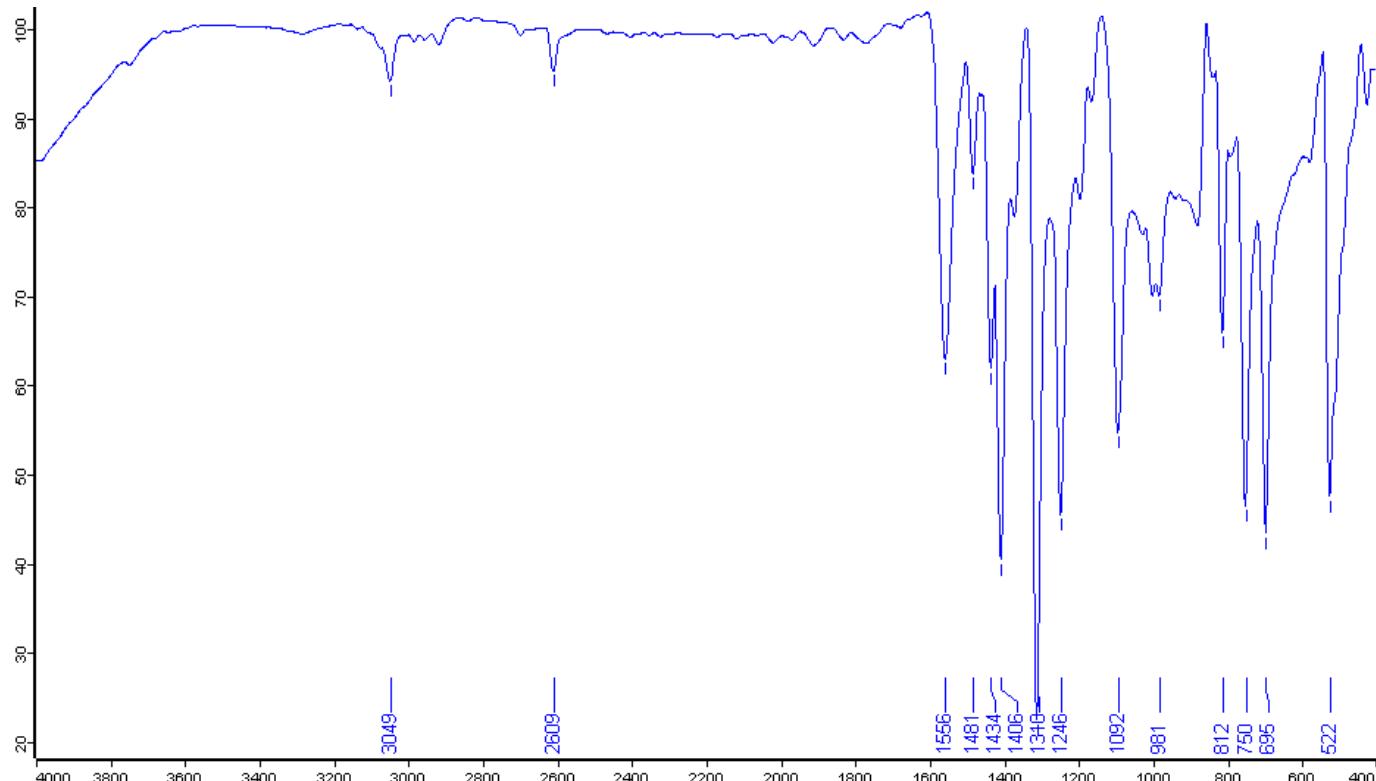
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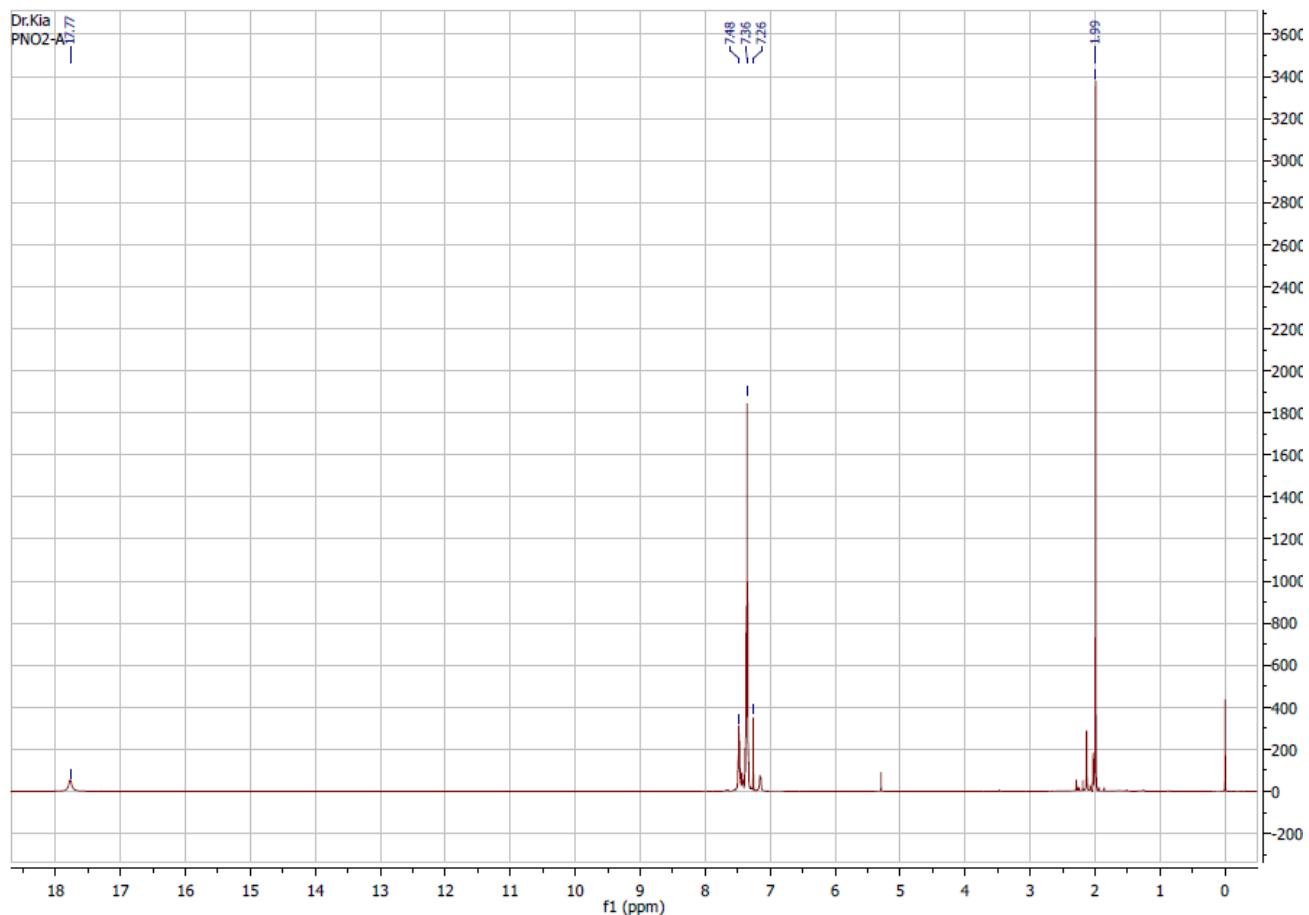
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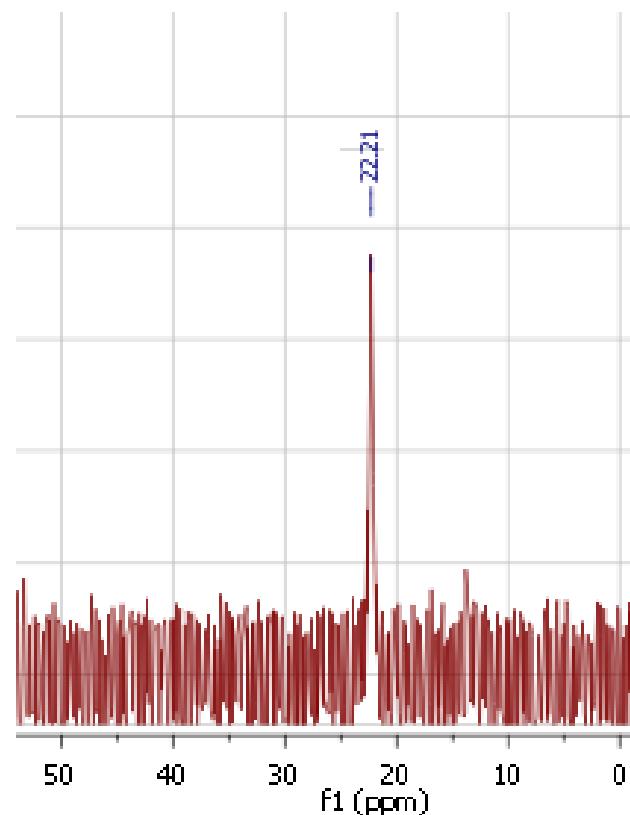
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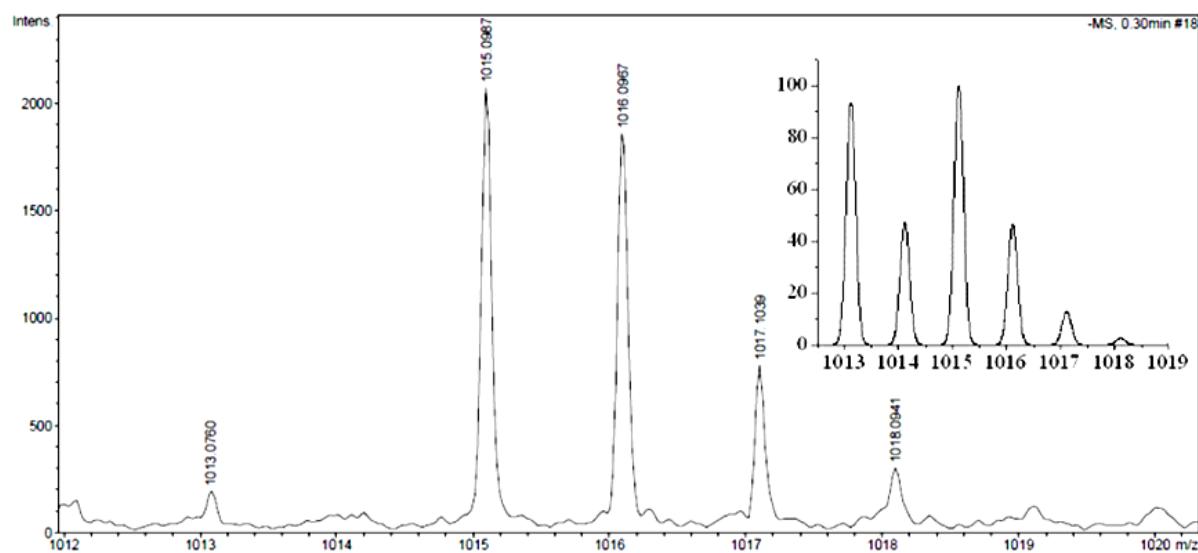
**Fig. S1.** The FT-IR spectrum of **1**.



**Figure S2.**  $^1\text{H}$  NMR (500 MHz), spectrum of complex **2** in  $\text{CDCl}_3$  at room temperature.



**Fig. S3.**  $^{31}\text{P} \{^1\text{H}\}$  NMR spectrum of **1** in  $\text{CDCl}_3$  at room temperature.



**Fig. S4.** Expanded ESI-MS spectrum of **1** in base peak position. Inset shows the simulated isotope pattern.

**Table S1.** G03/B3LYP calculated one-electron energy and percentage composition of selected frontier MOs of **1** expressed in terms of component fragments

<b>MO</b>	<b>Name</b>	<b>E (eV)</b>	<b>Ag</b>	<b>PPh<sub>3</sub></b>	<b>NO<sub>2</sub><sup>b</sup></b>	<b>Co</b>	<b>DMGH<sup>c</sup></b>	<b>NO<sub>2</sub><sup>t</sup></b>
246	LUMO + 5	-1	4.1	90.8	4.6	0.2	0.4	0
245	LUMO + 4	-1.1	7	54.6	4.8	5.1	28.2	0.3
244	LUMO + 3	-1.1	6.6	79.2	1.7	6.7	5.2	0.7
243	LUMO + 2	-1.2	6.8	50.9	5.1	15.8	17.8	3.6
242	LUMO + 1	-1.2	4.5	95	0.4	0.1	0	0
<b>241</b>	<b>LUMO</b>	<b>-1.3</b>	<b>5.7</b>	<b>11.8</b>	<b>76</b>	<b>2.9</b>	<b>3.2</b>	<b>0.3</b>
<b>240</b>	<b>HOMO</b>	<b>-5</b>	<b>0.1</b>	<b>2.6</b>	<b>0.5</b>	<b>0.3</b>	<b>94.8</b>	<b>1.7</b>
239	HOMO - 1	-5.1	0.2	2	0.1	6.1	91.2	0.3
238	HOMO - 2	-5.7	3.9	3.8	15.4	13.1	3.6	60
237	HOMO - 3	-5.8	0.1	2	0.5	3.4	88.4	5.4
236	HOMO - 4	-6.2	0.4	0.4	3.1	2.5	25.2	68.5
235	HOMO - 5	-6.4	15.5	83.8	0.4	0.1	0	0

<sup>b</sup> Bridged NO<sub>2</sub>; <sup>c</sup> Dimethylglyoximate; <sup>t</sup> Terminal