

**Cyclometalated and NNN Terpyridine Ruthenium Photocatalysts and their Cytotoxic Activity**

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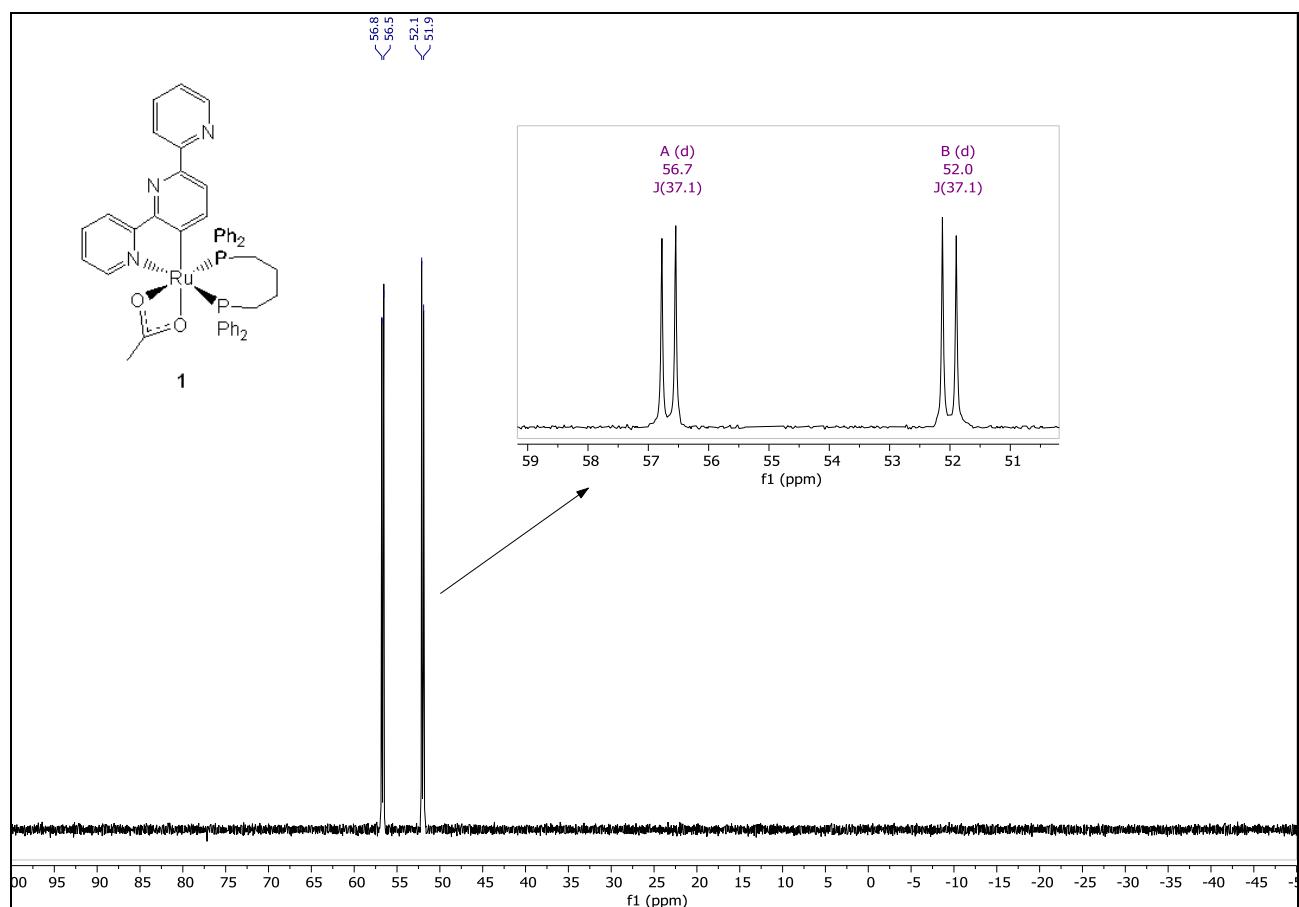
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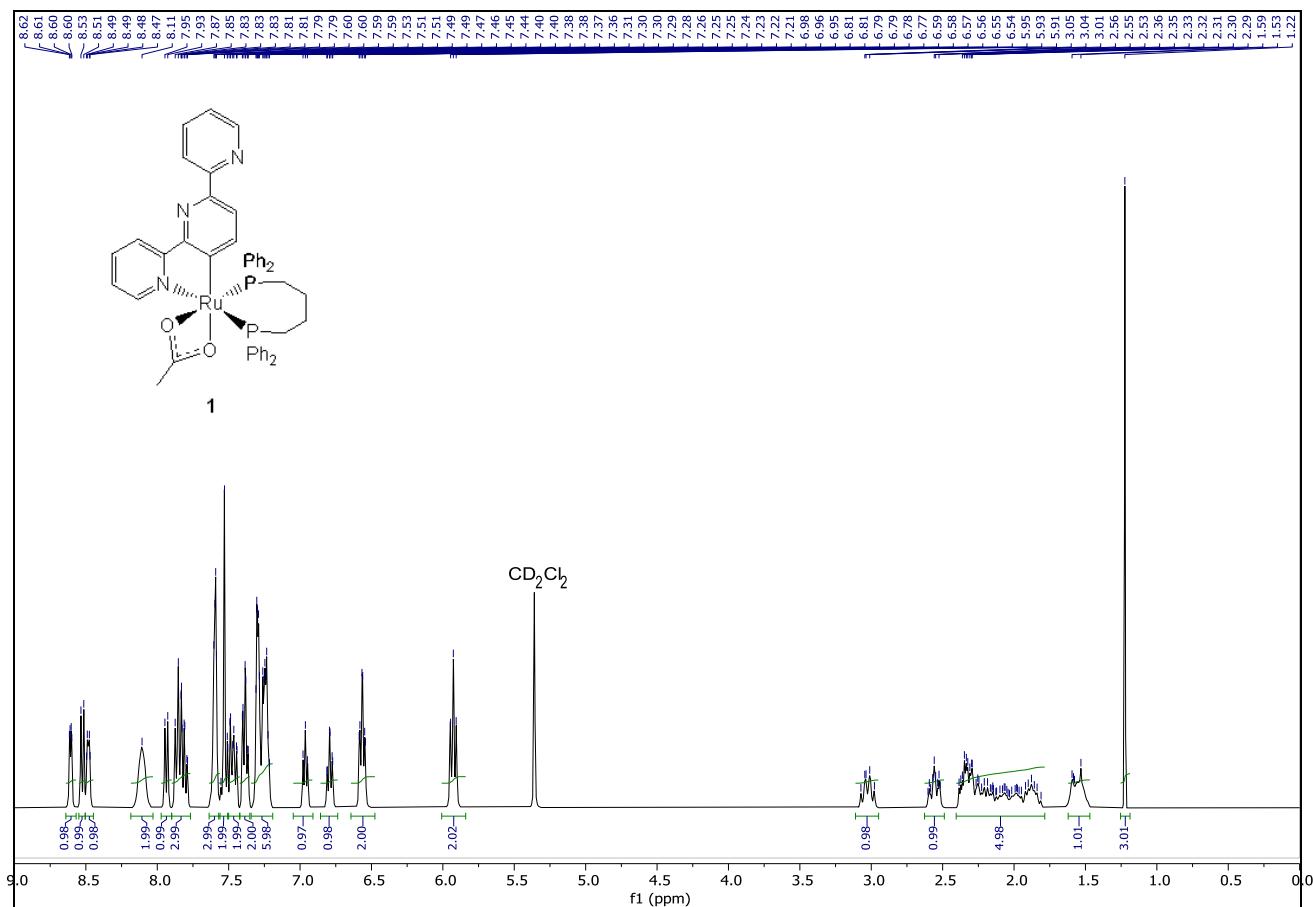
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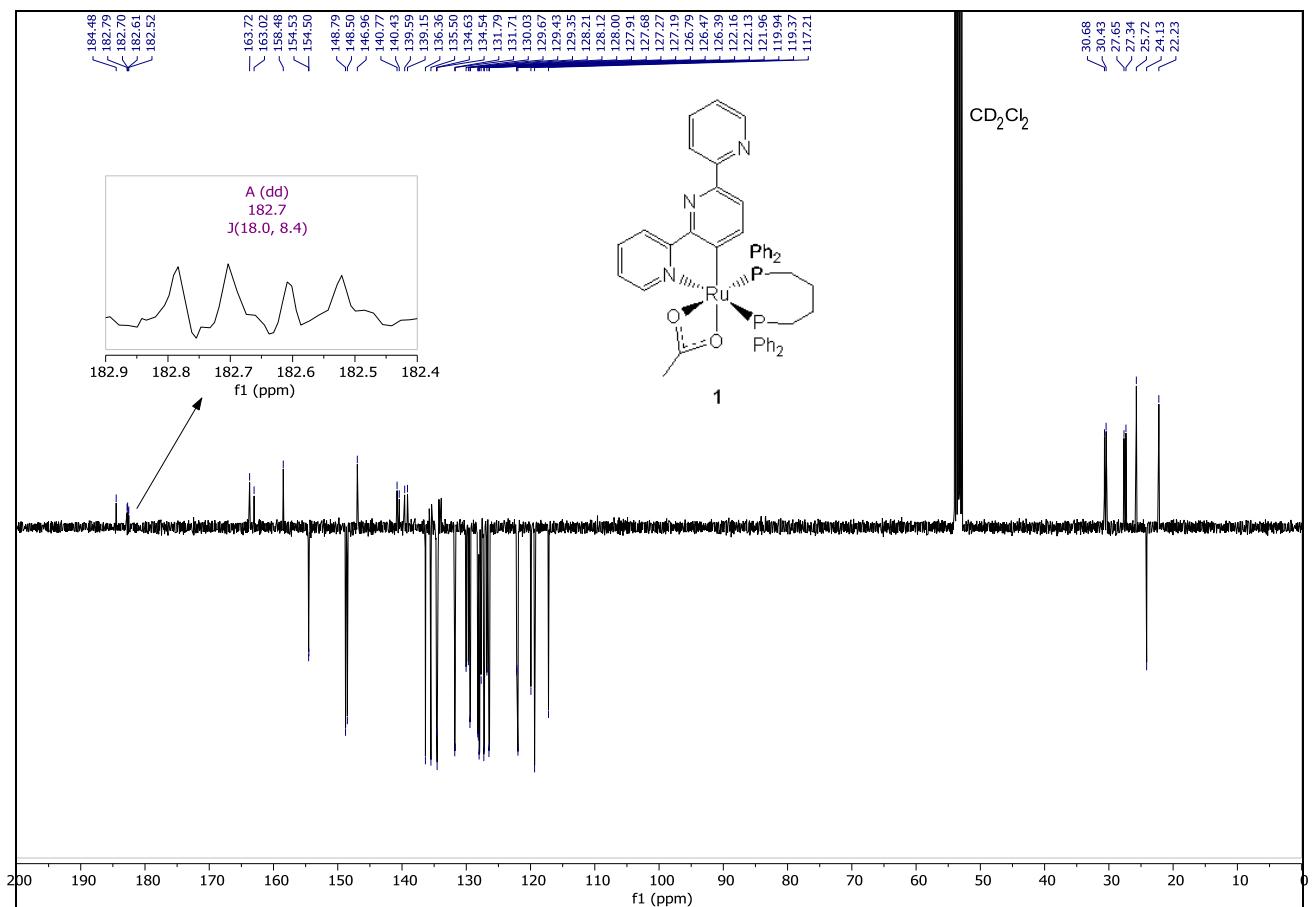
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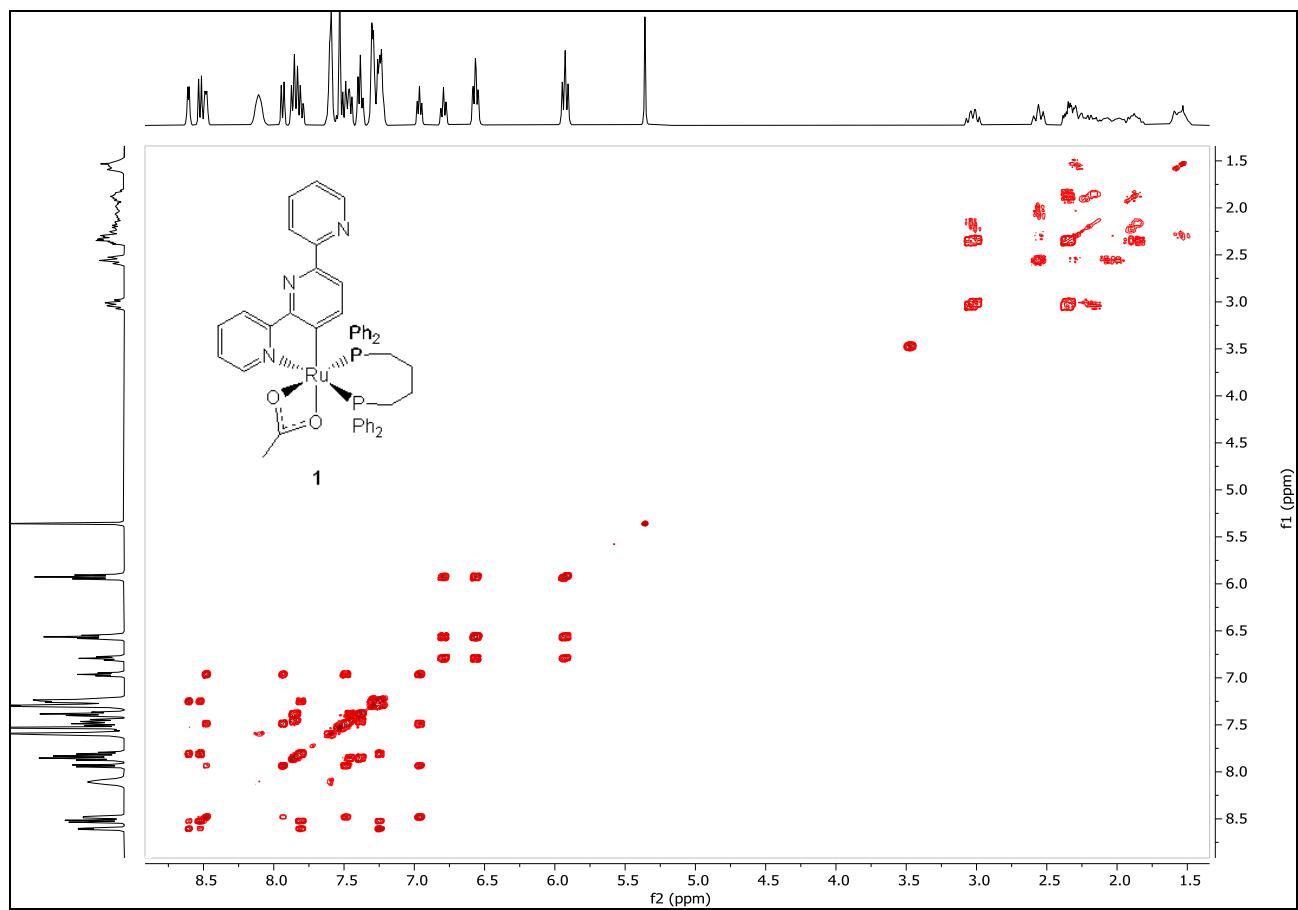
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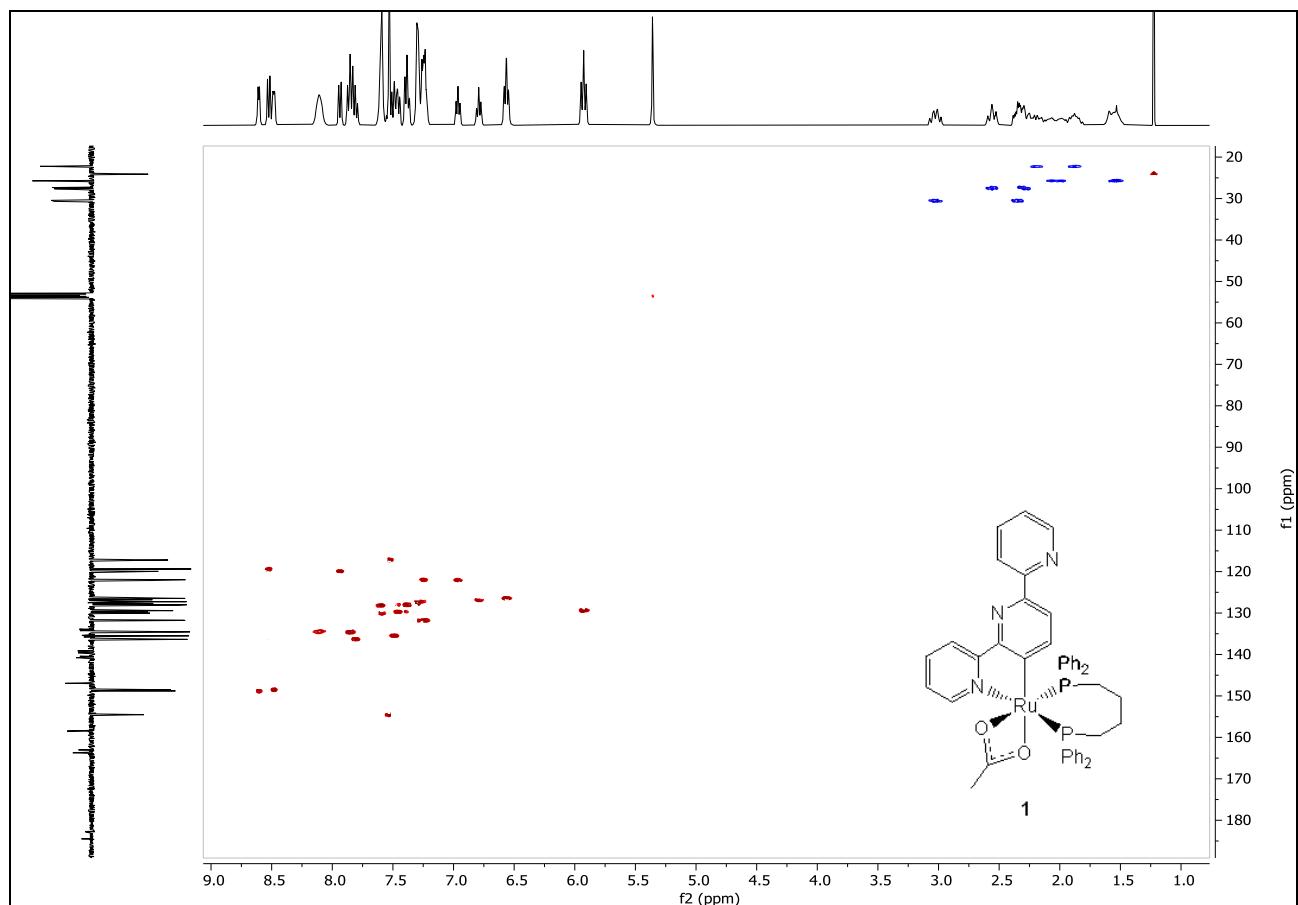
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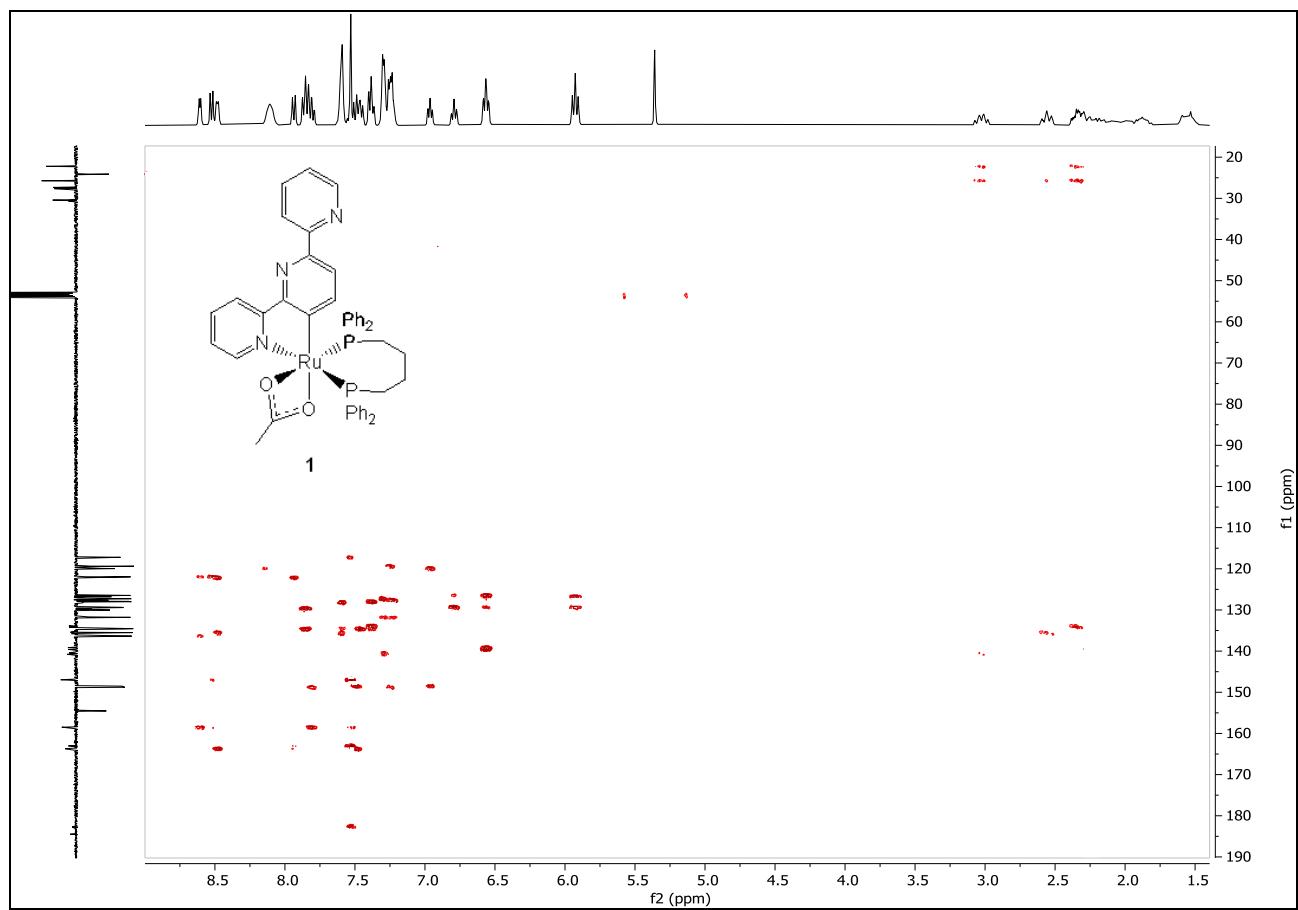
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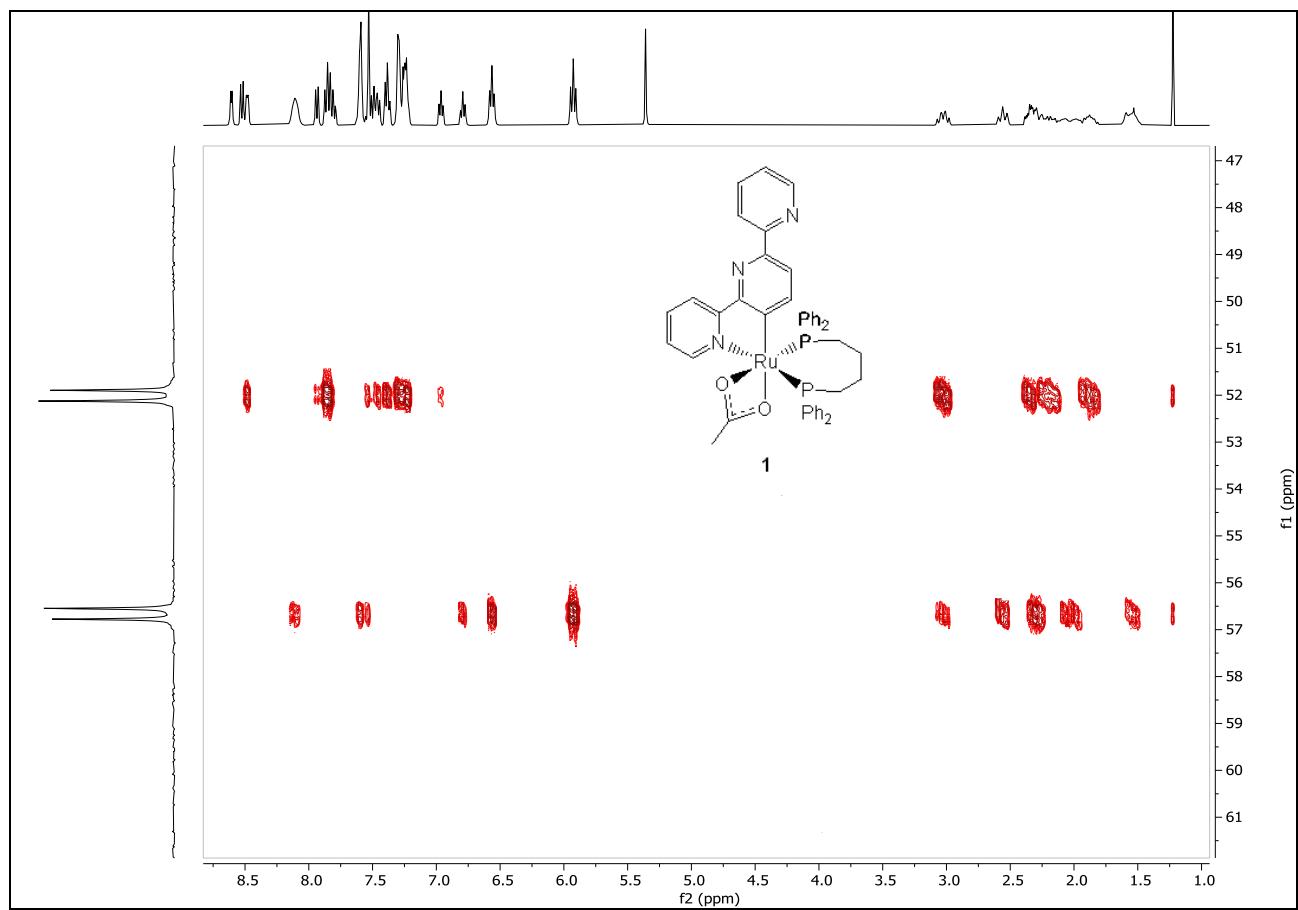
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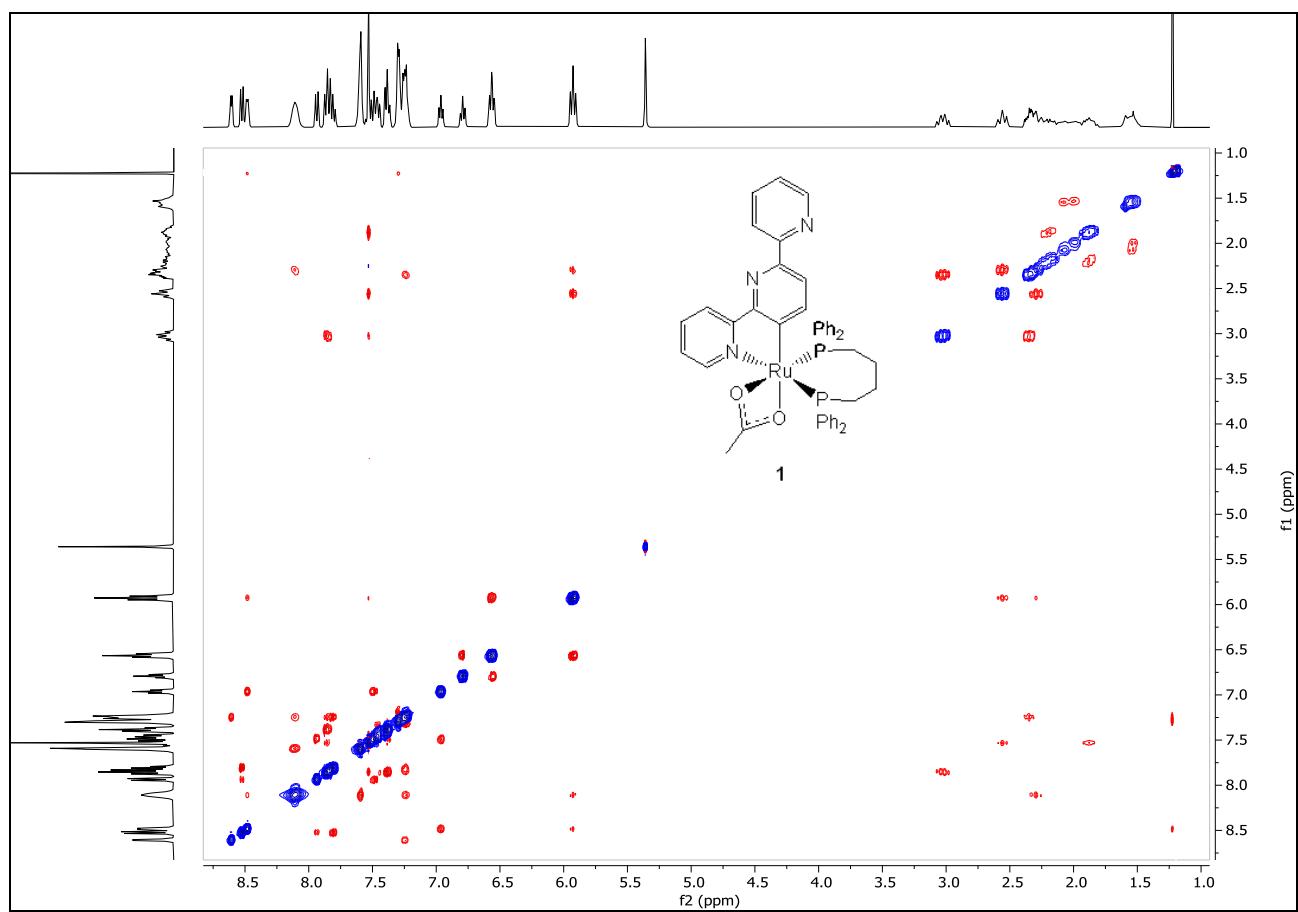
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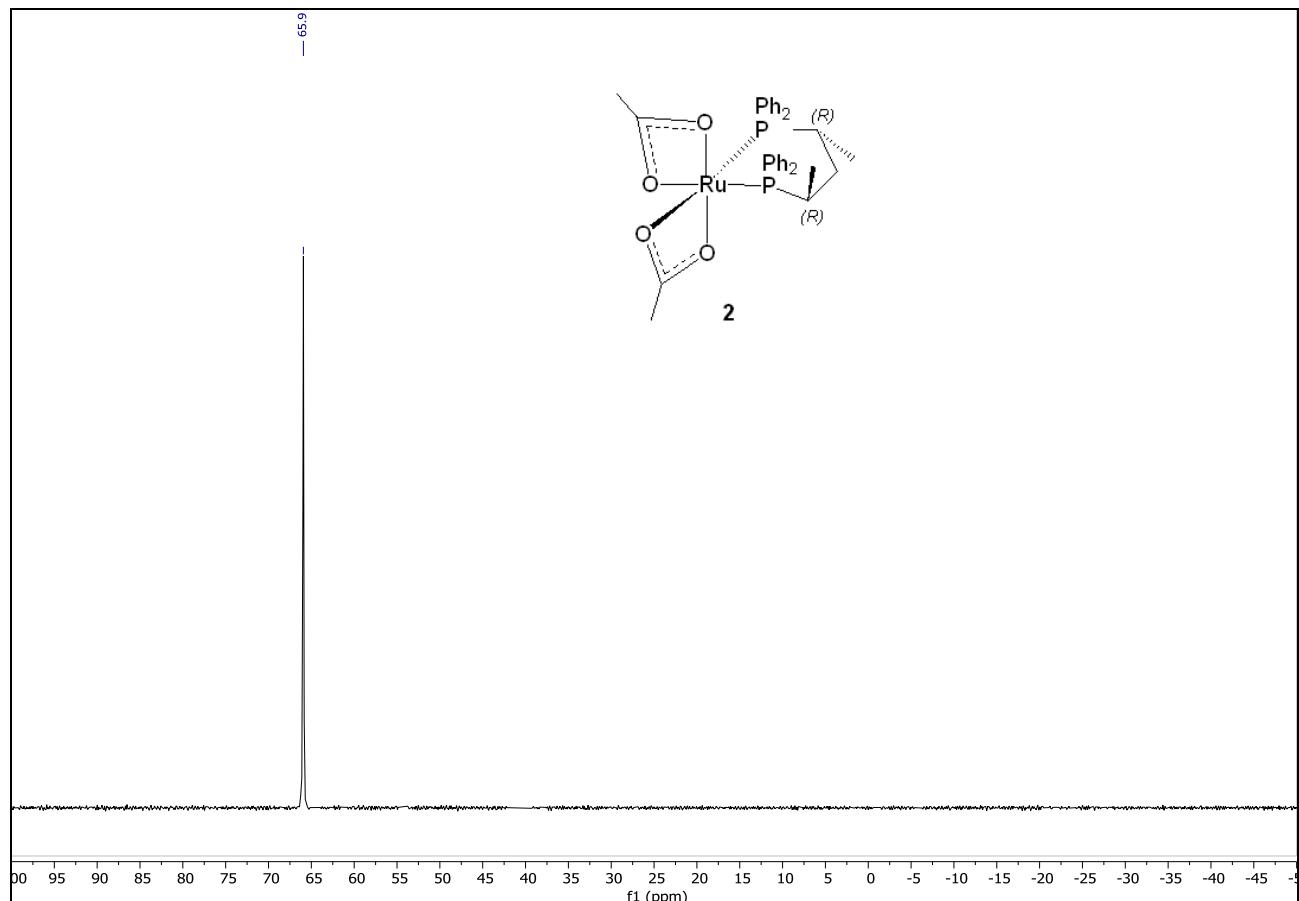
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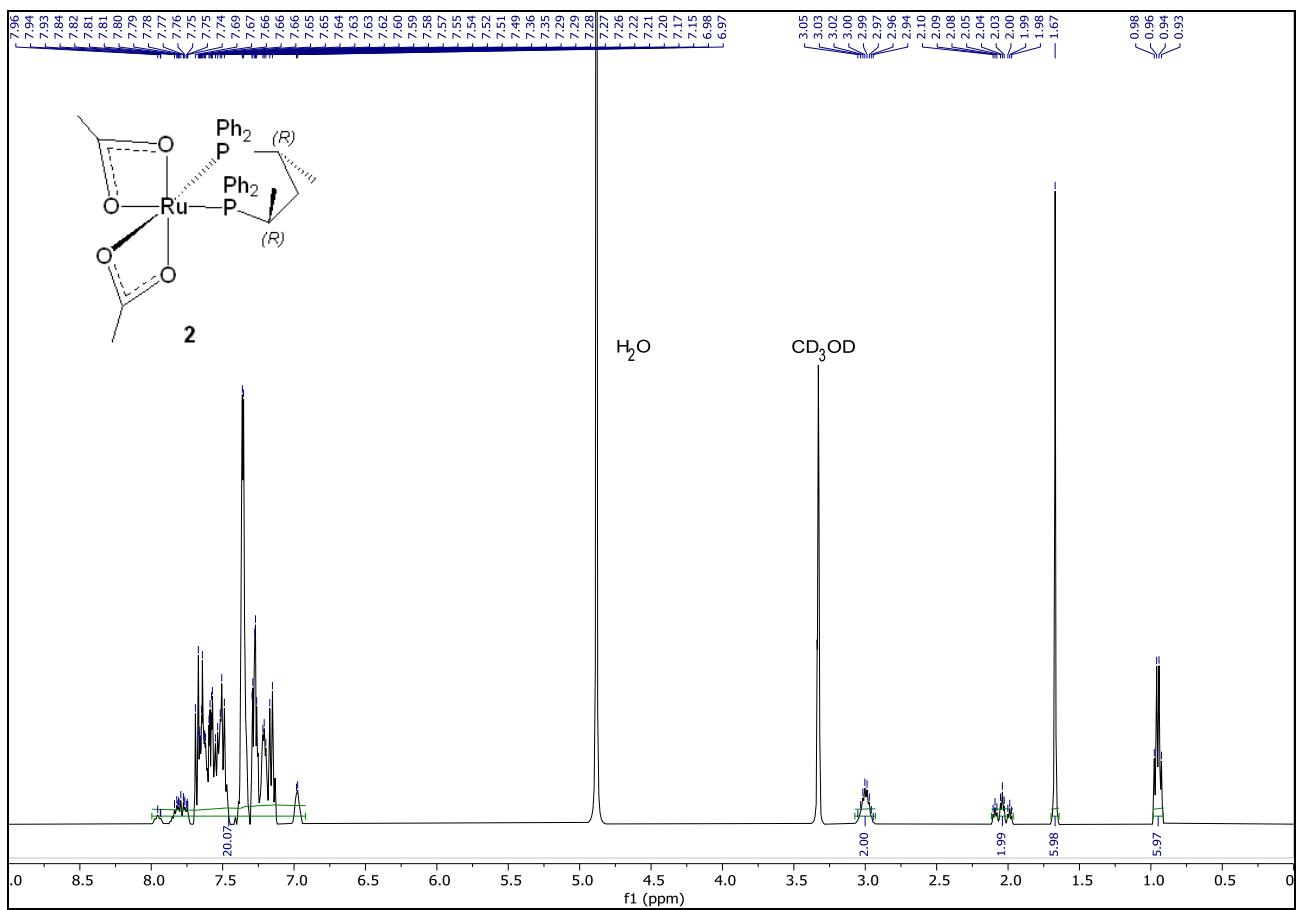
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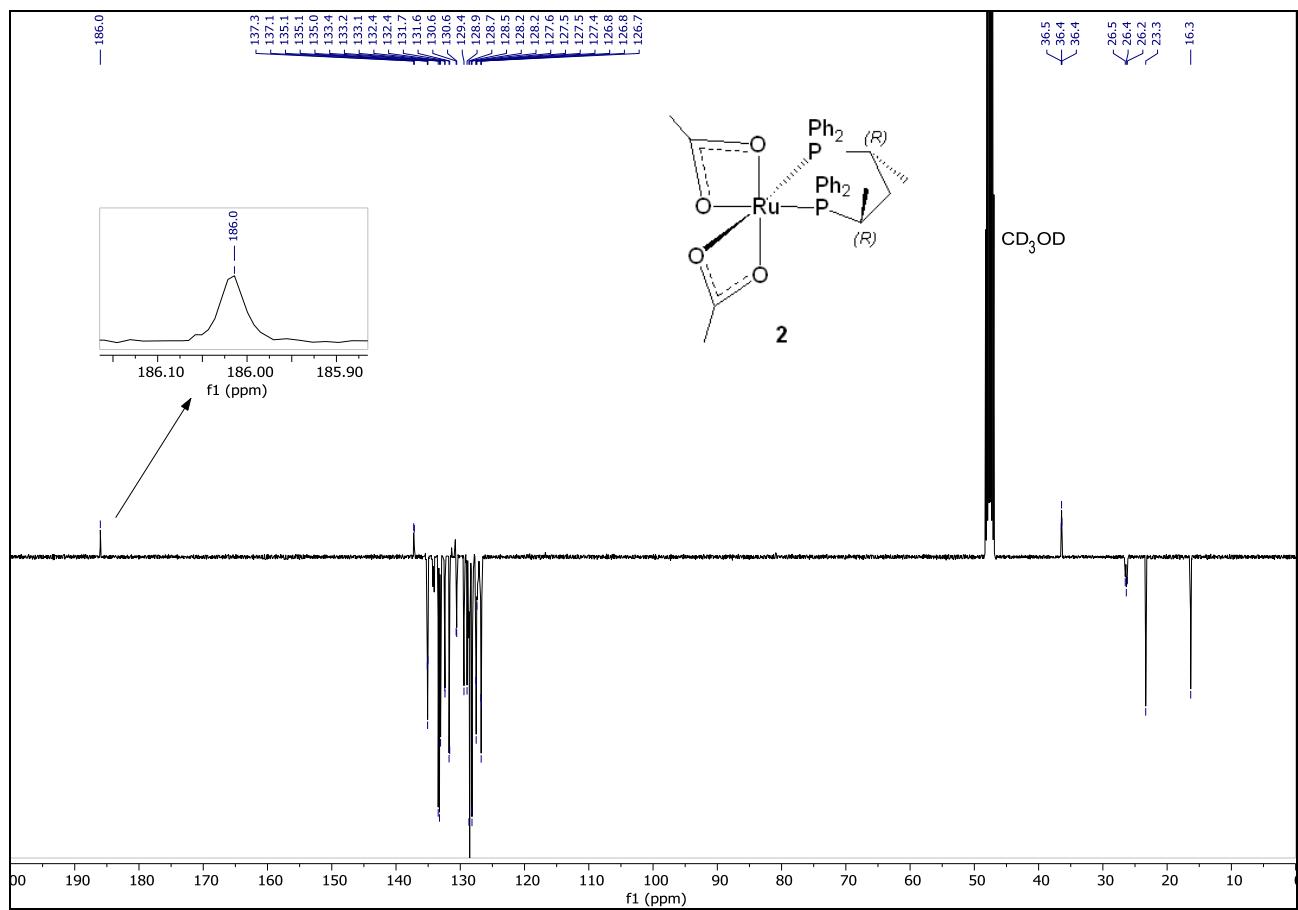
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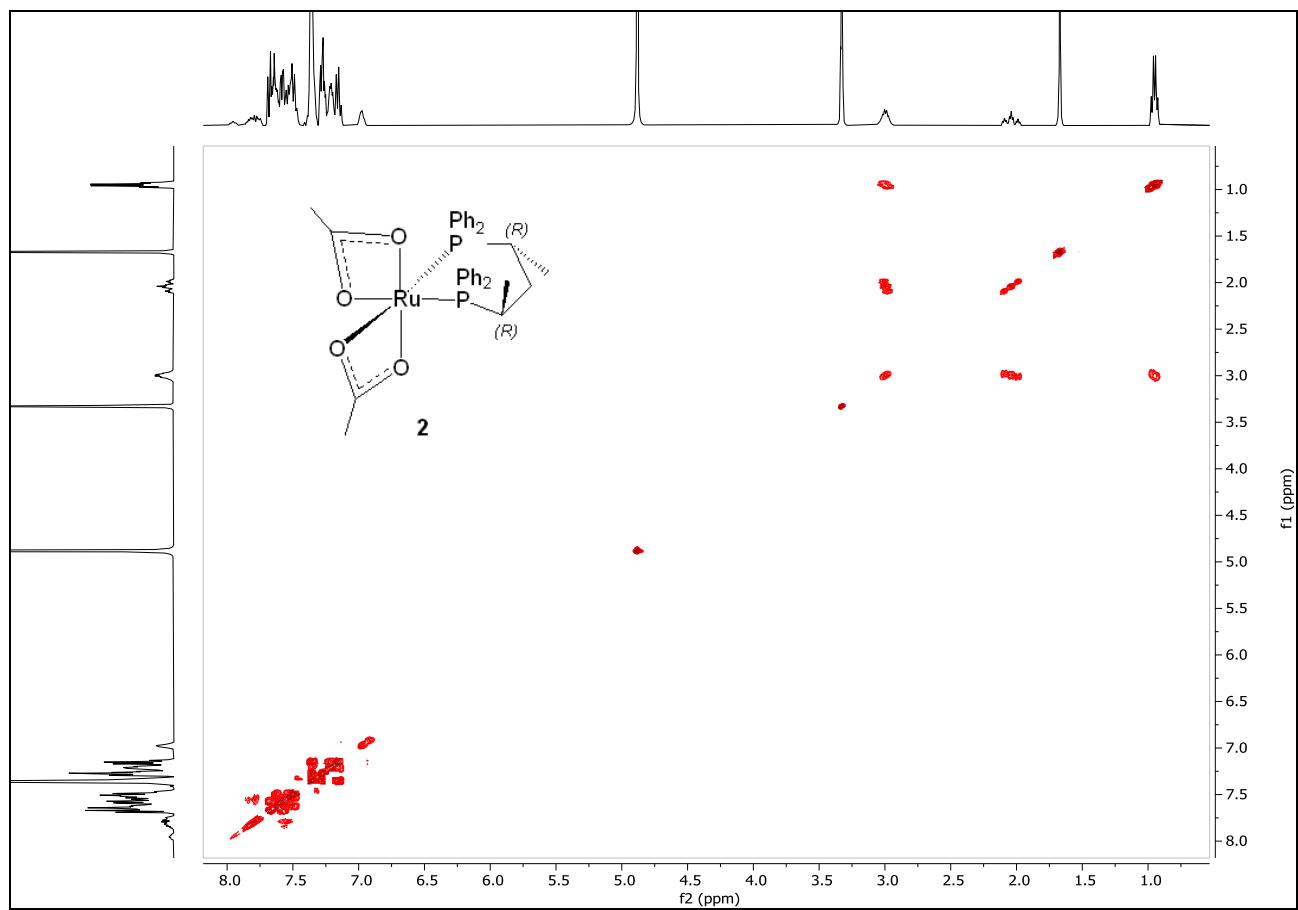
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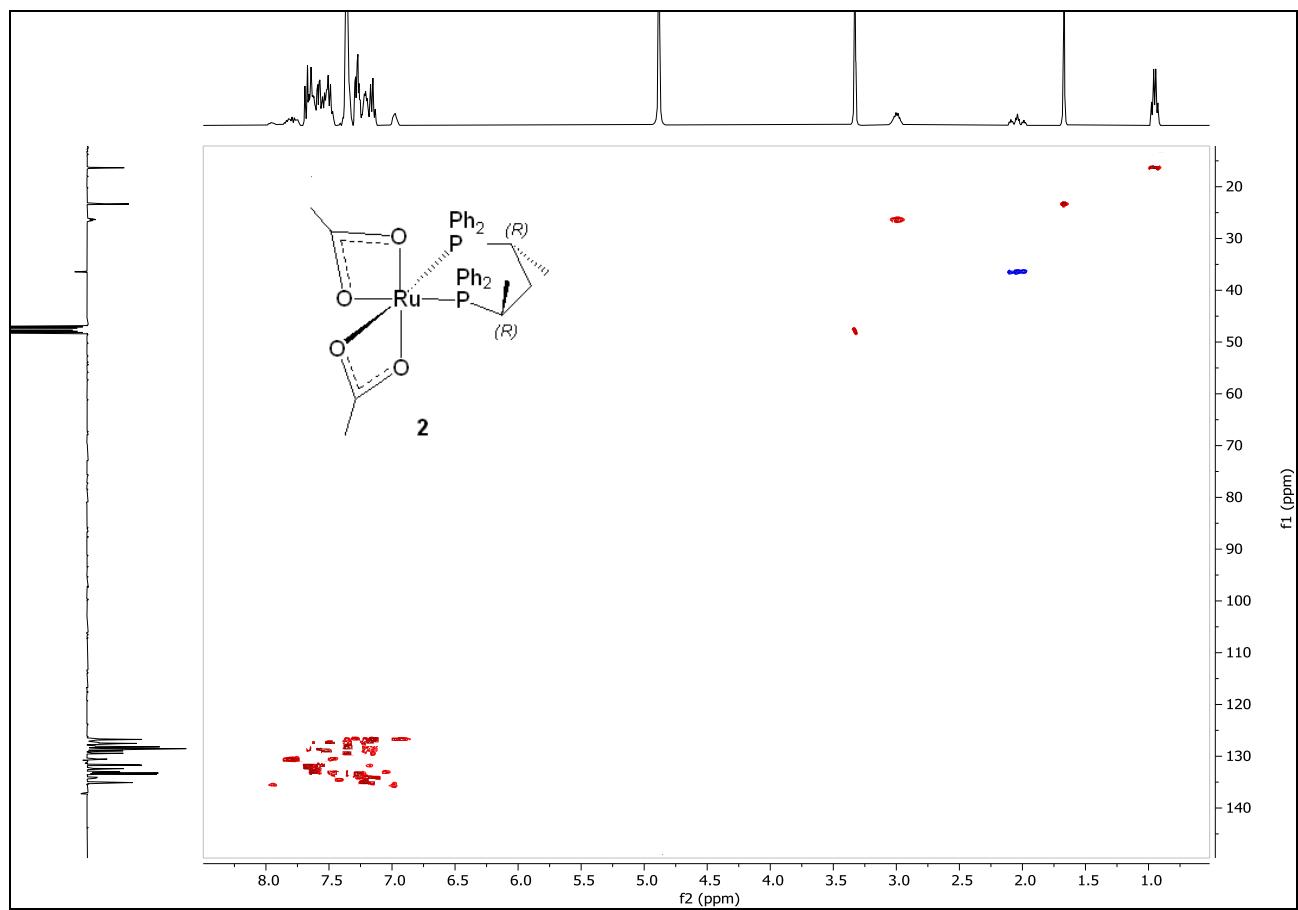
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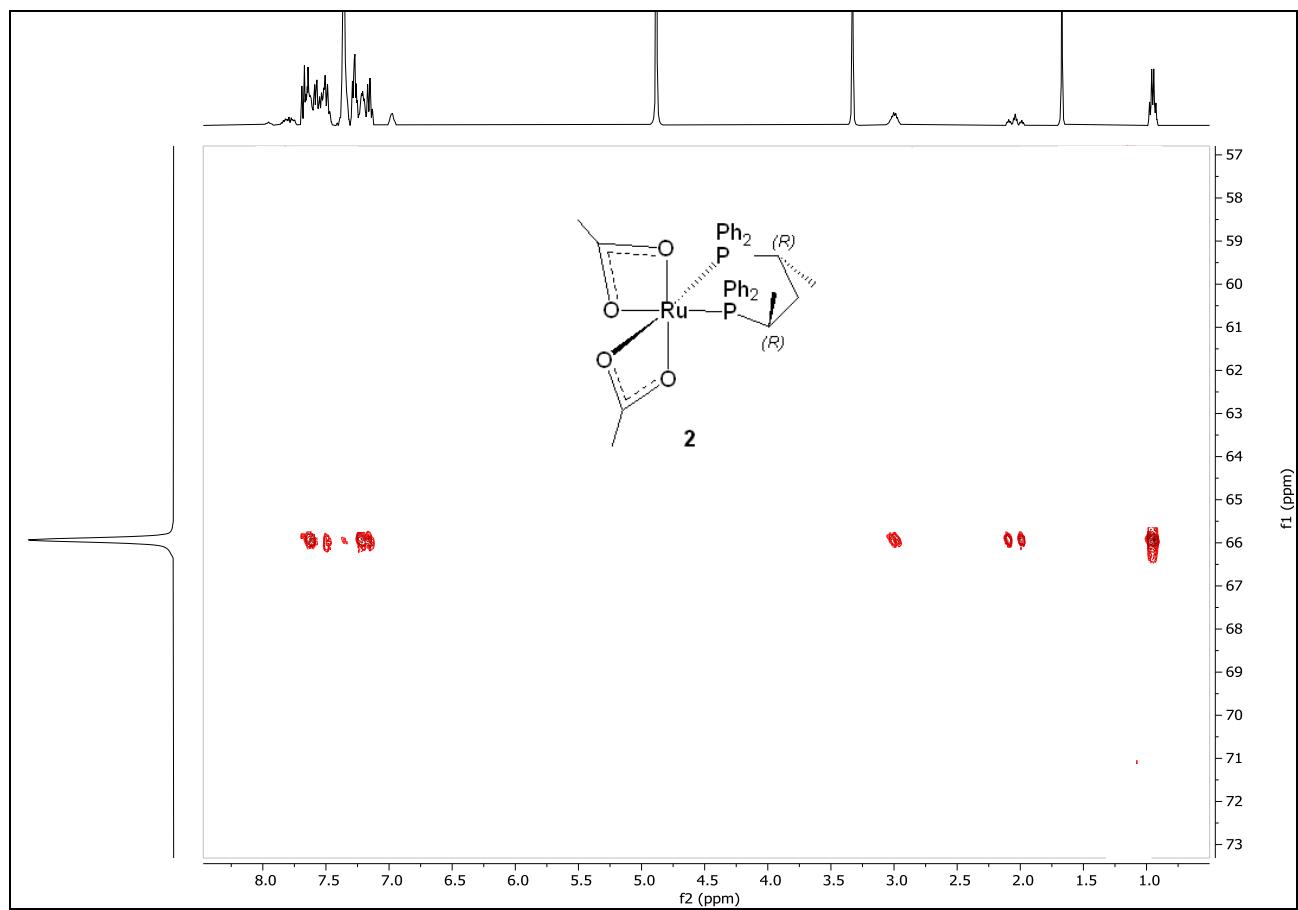
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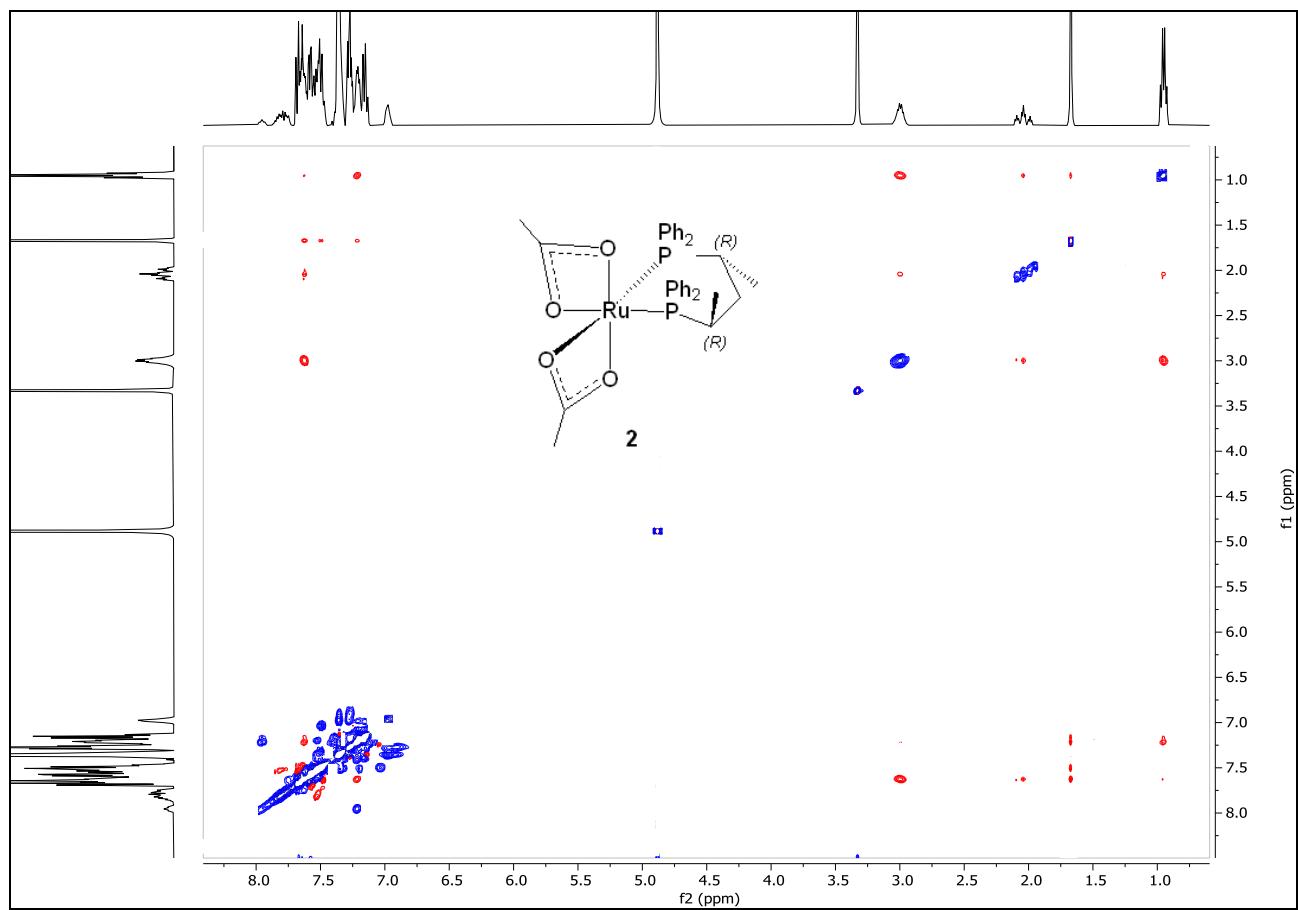
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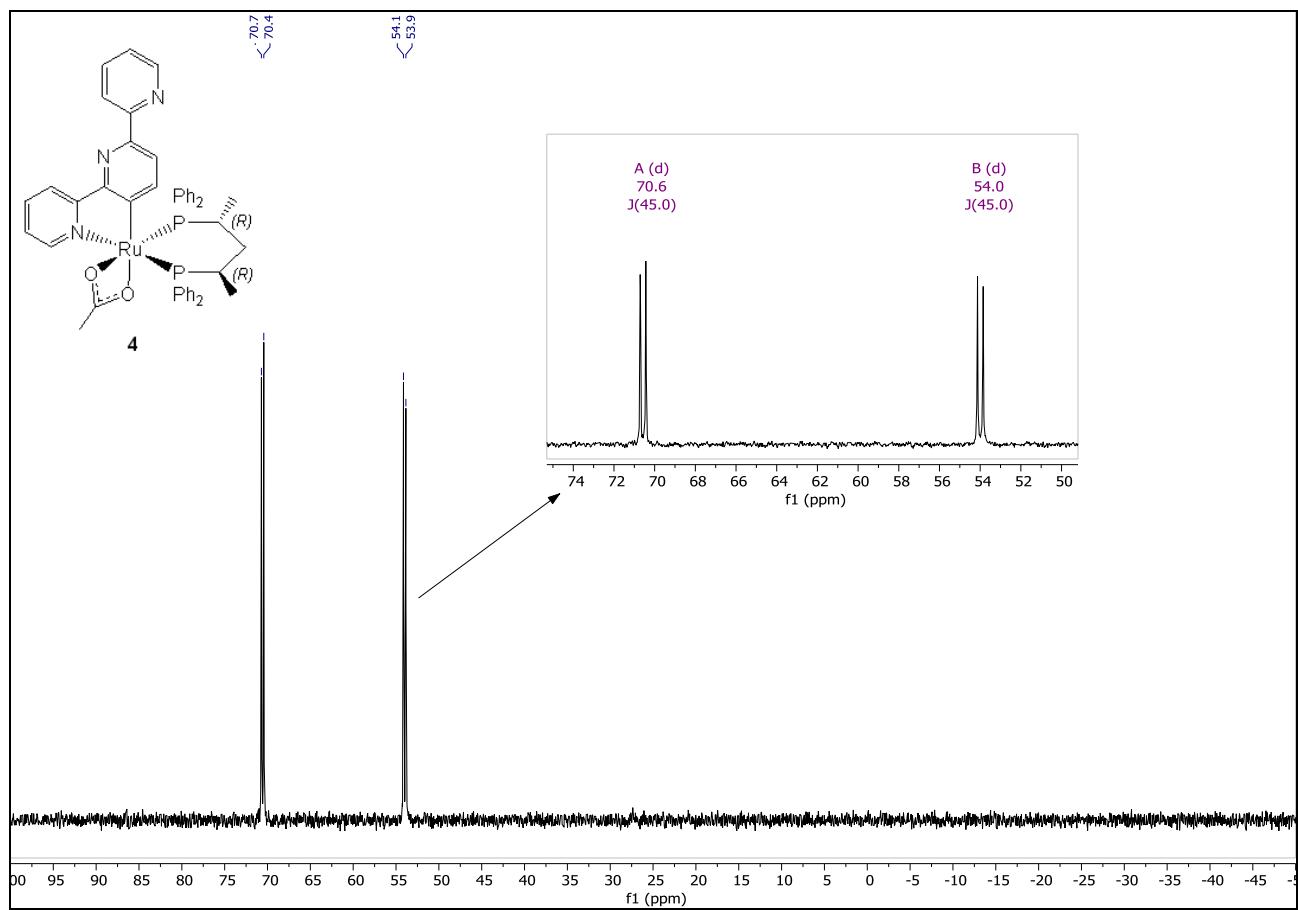
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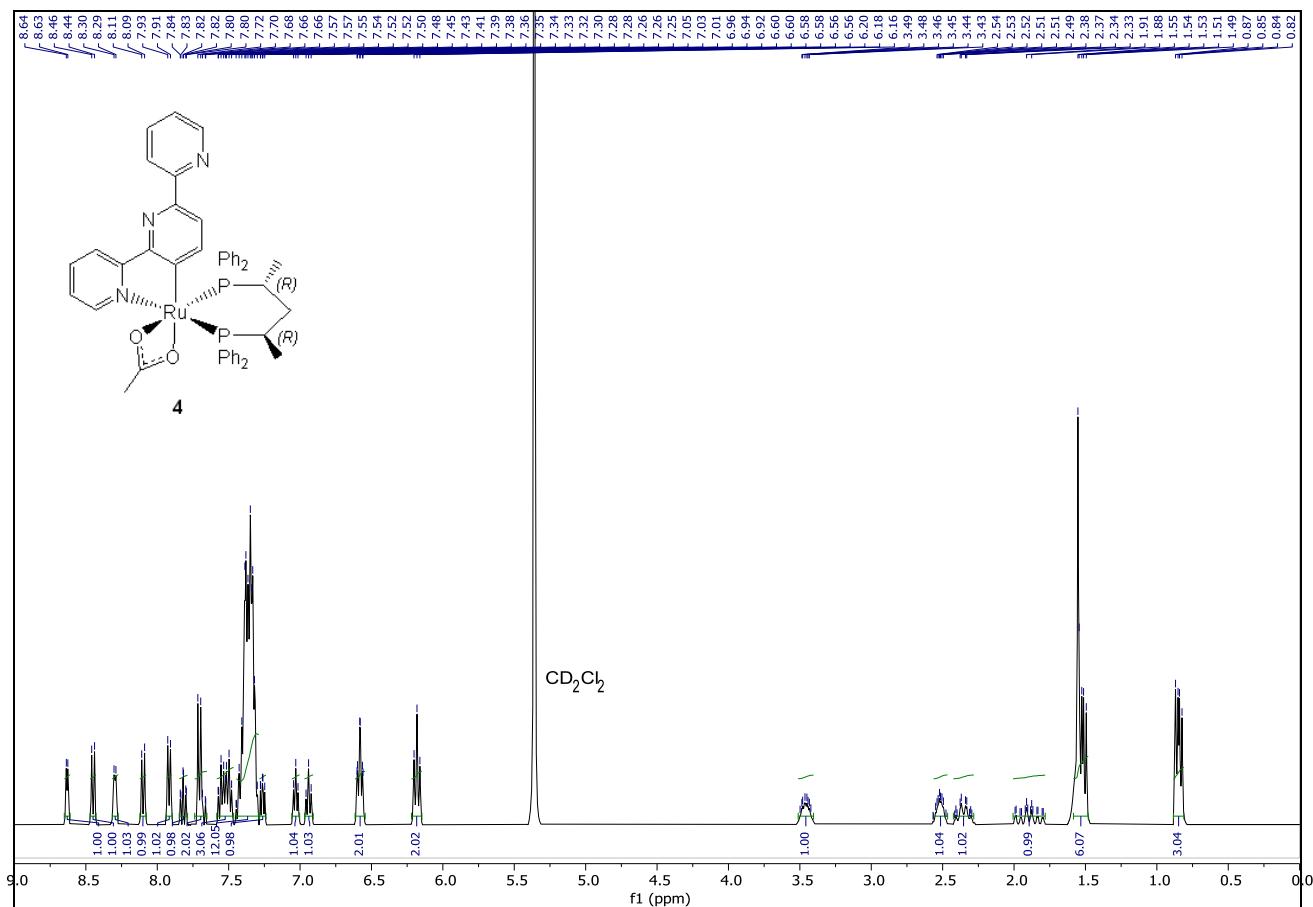
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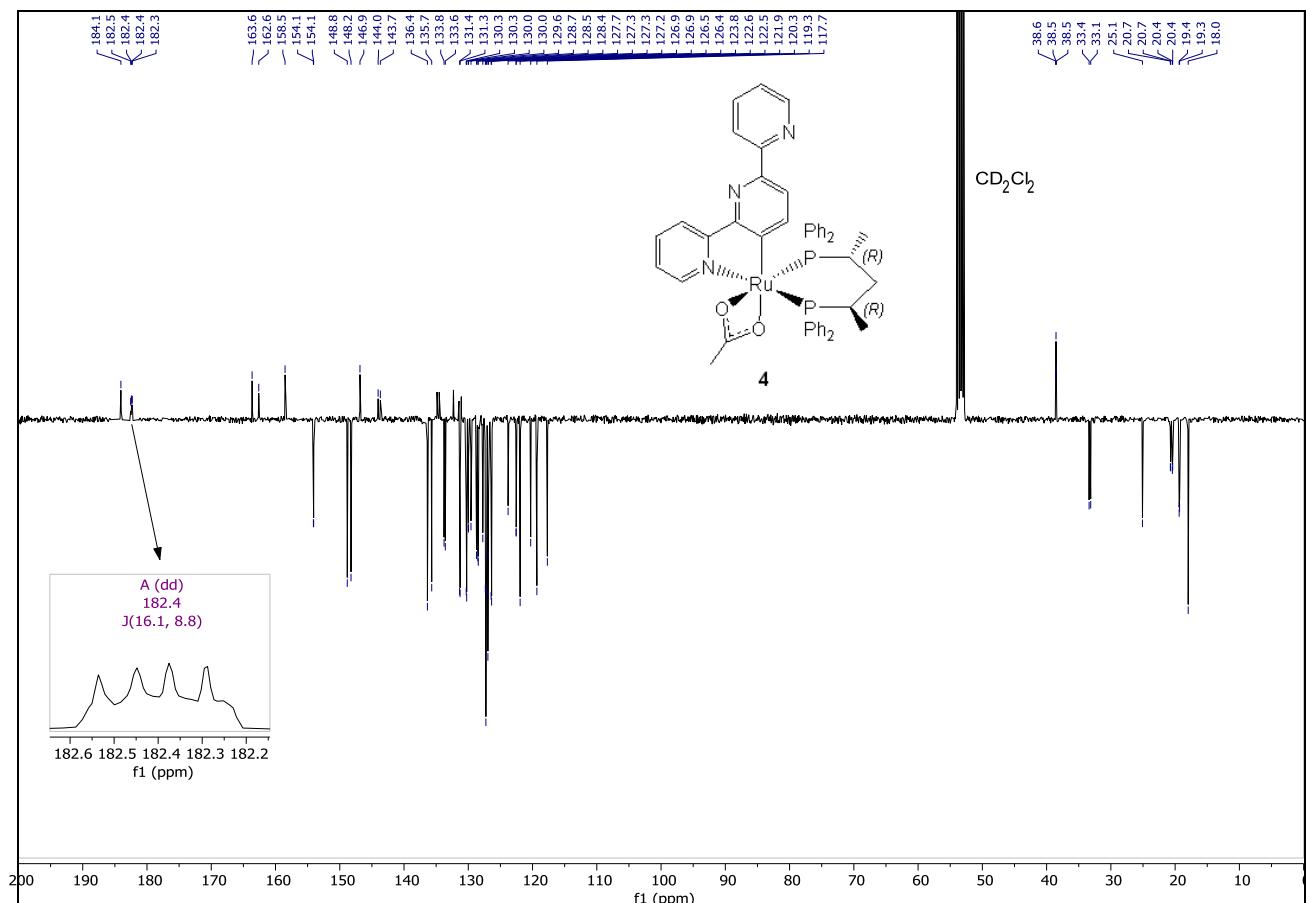
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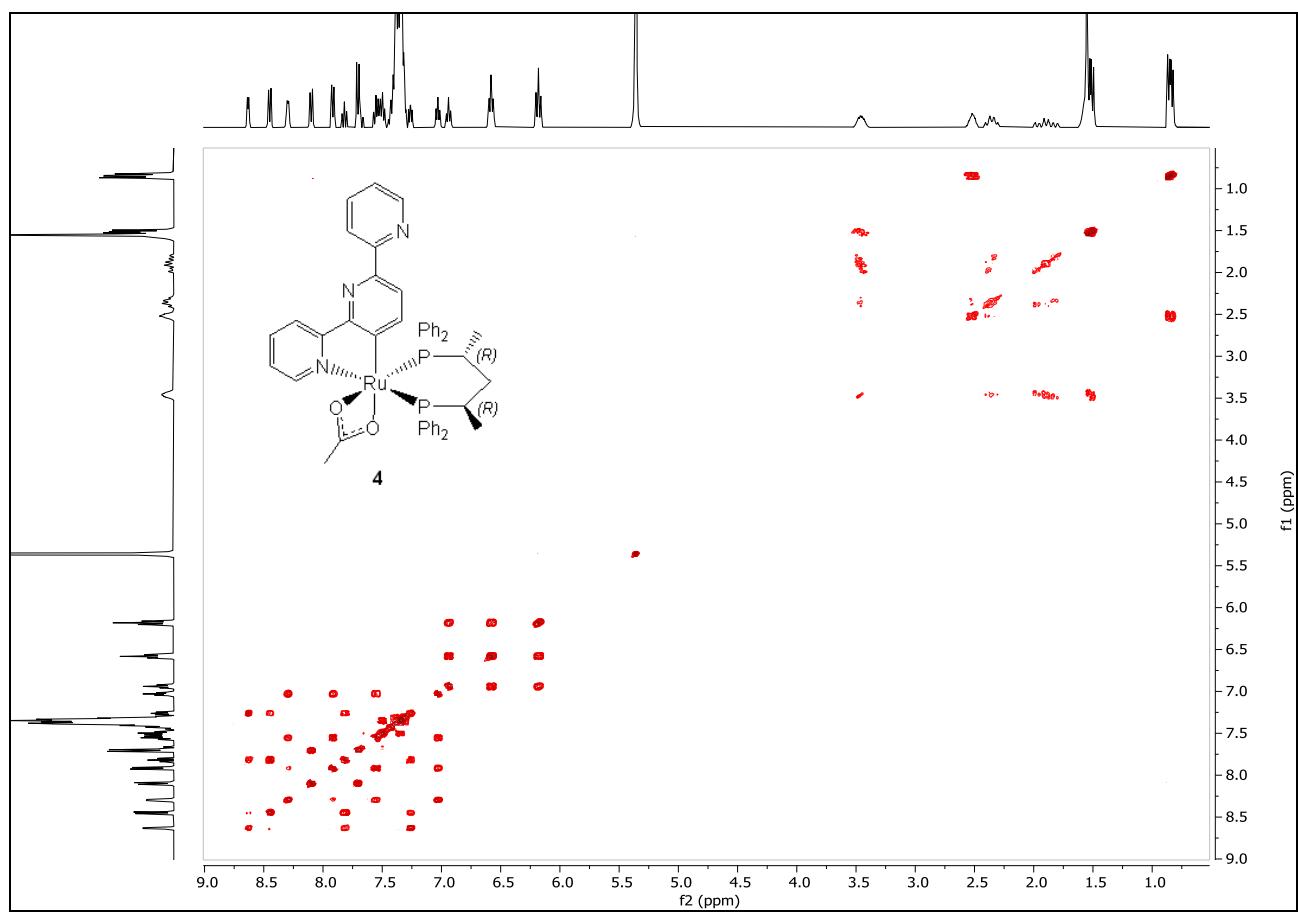
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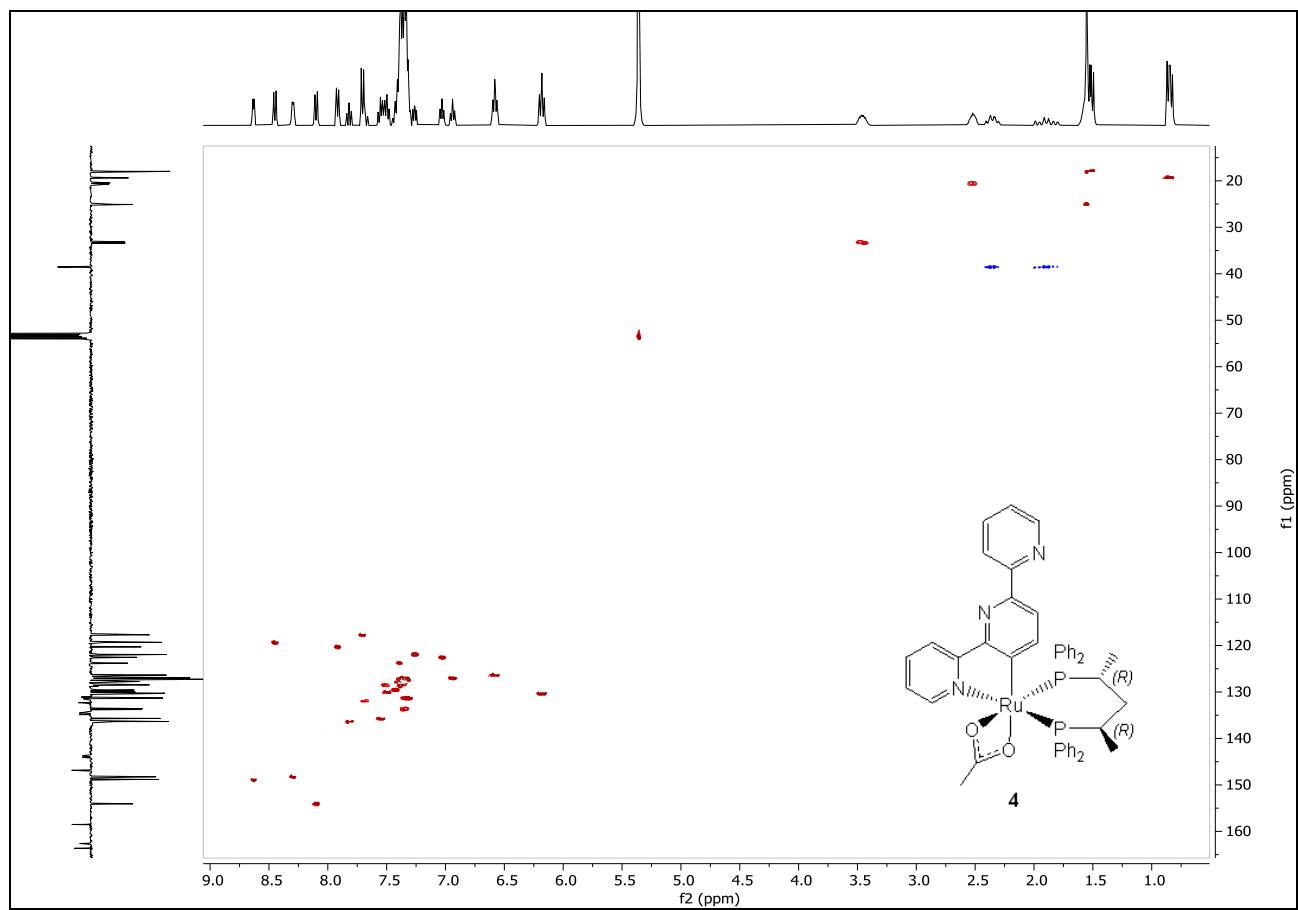
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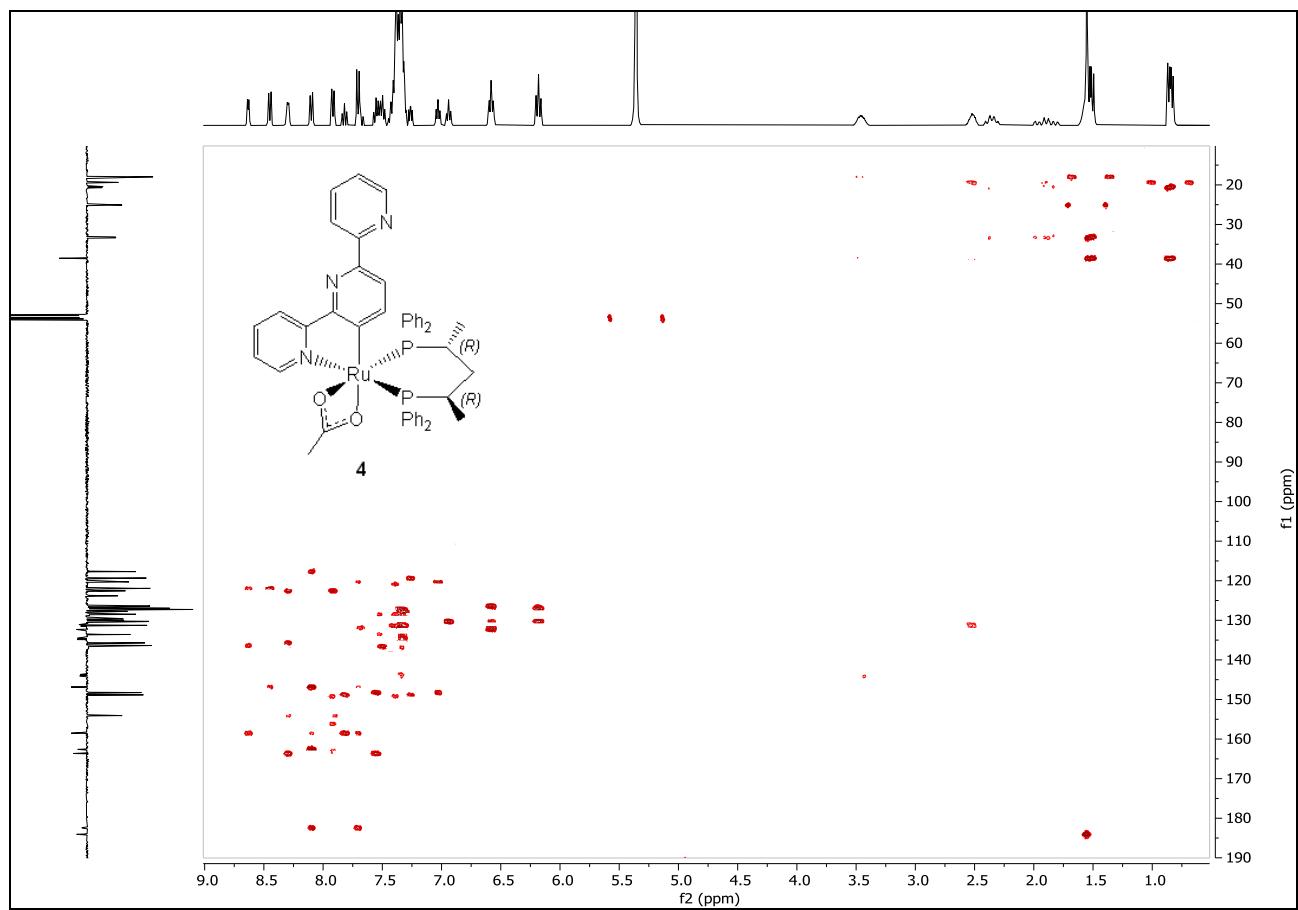
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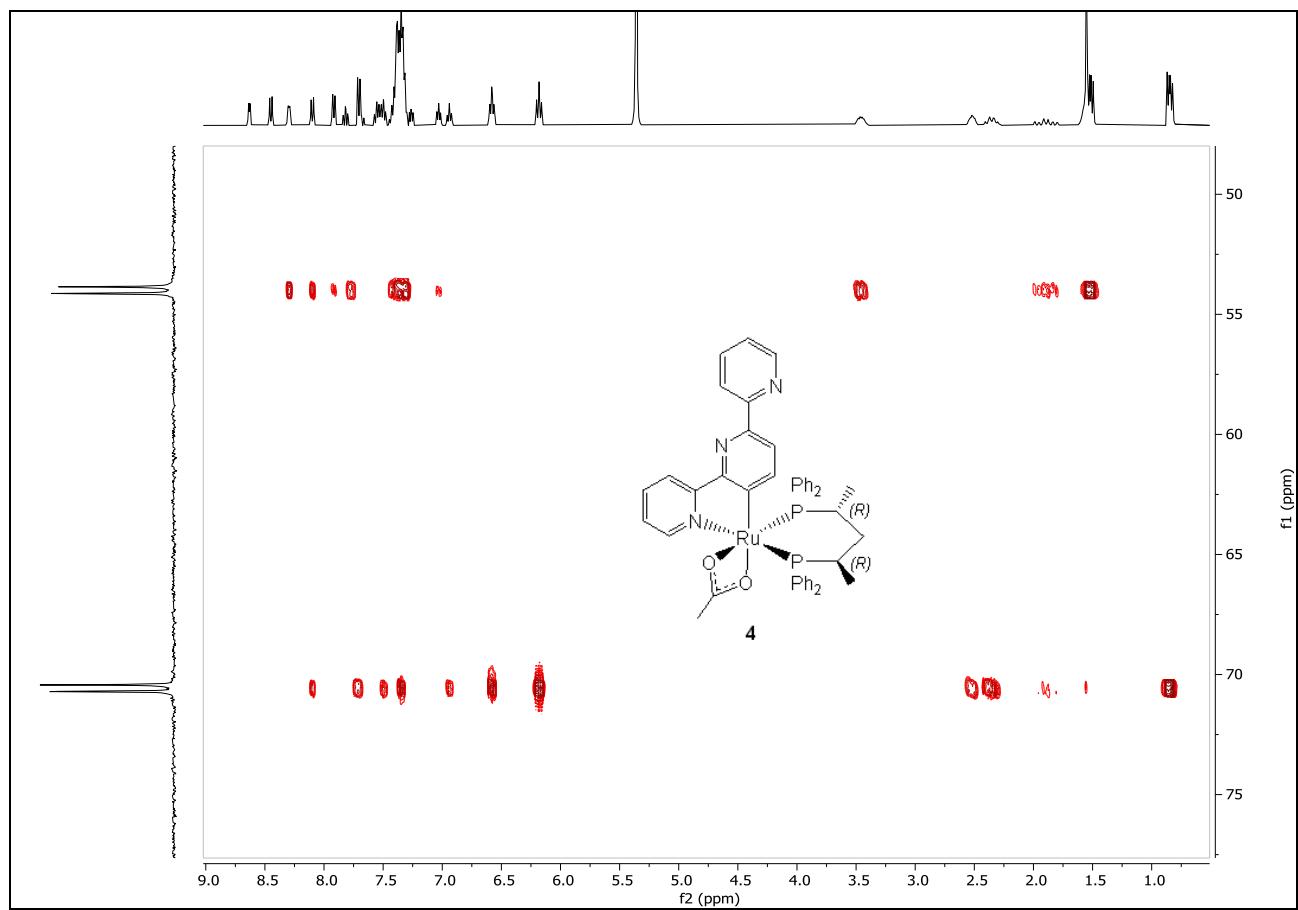
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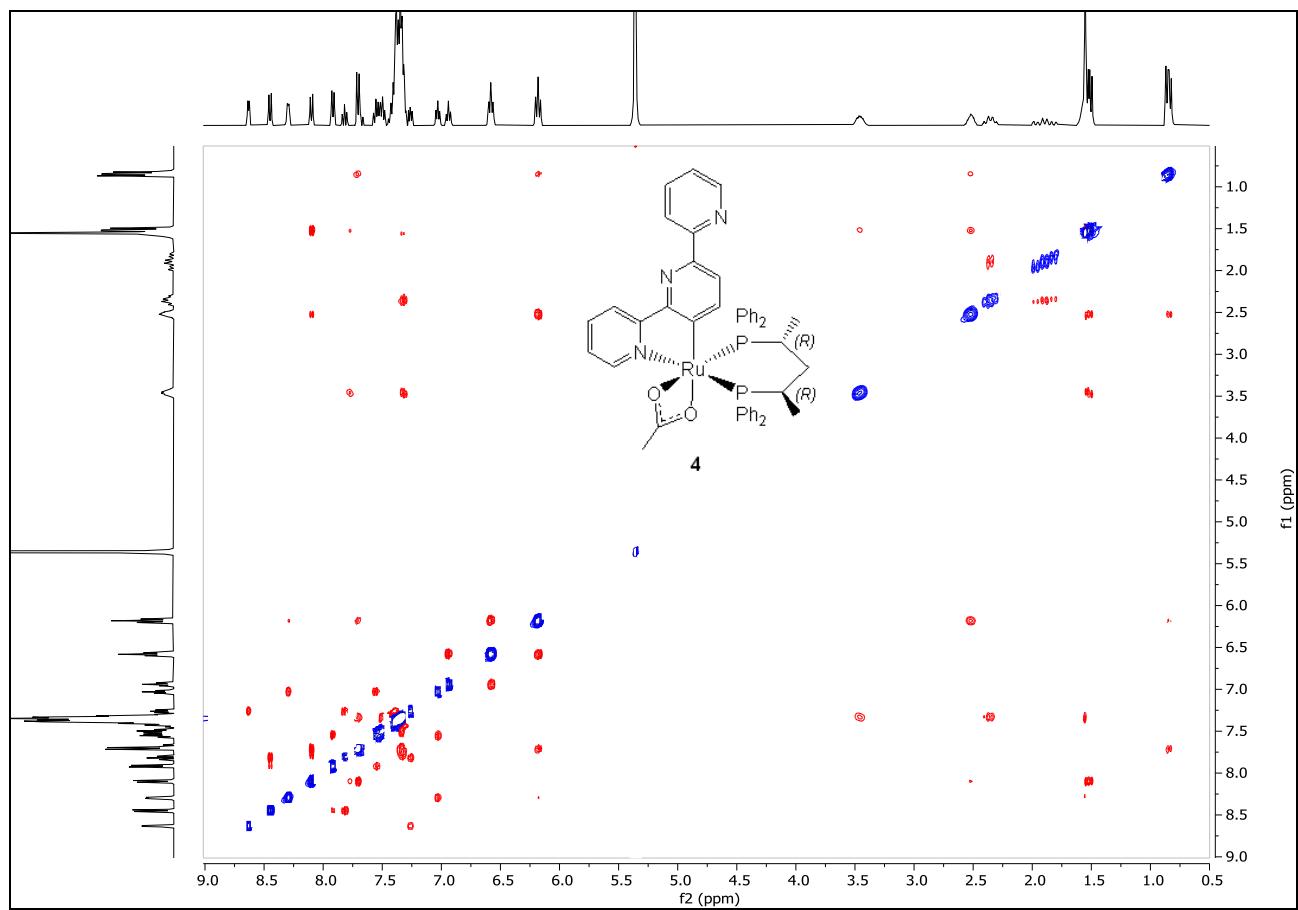
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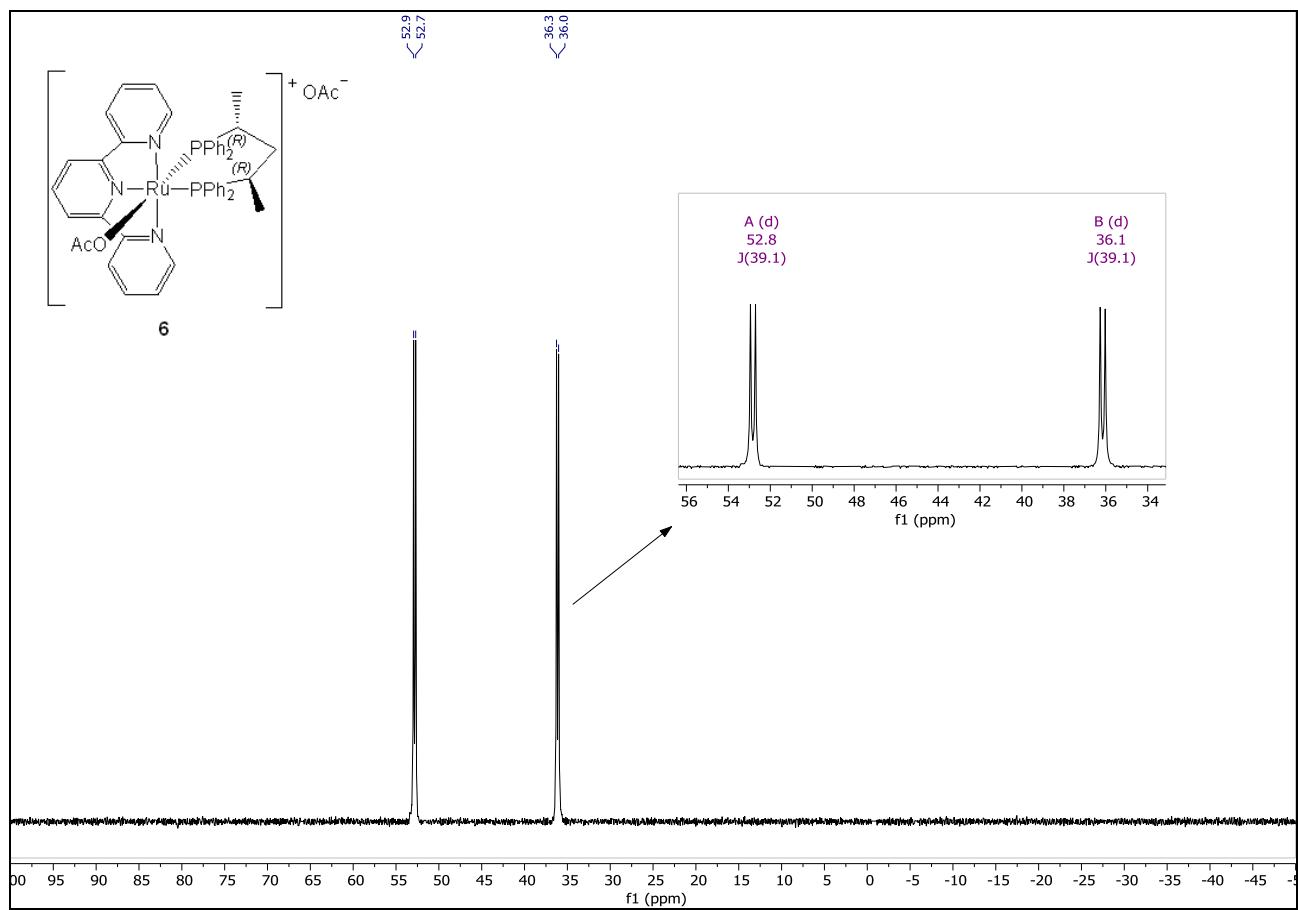
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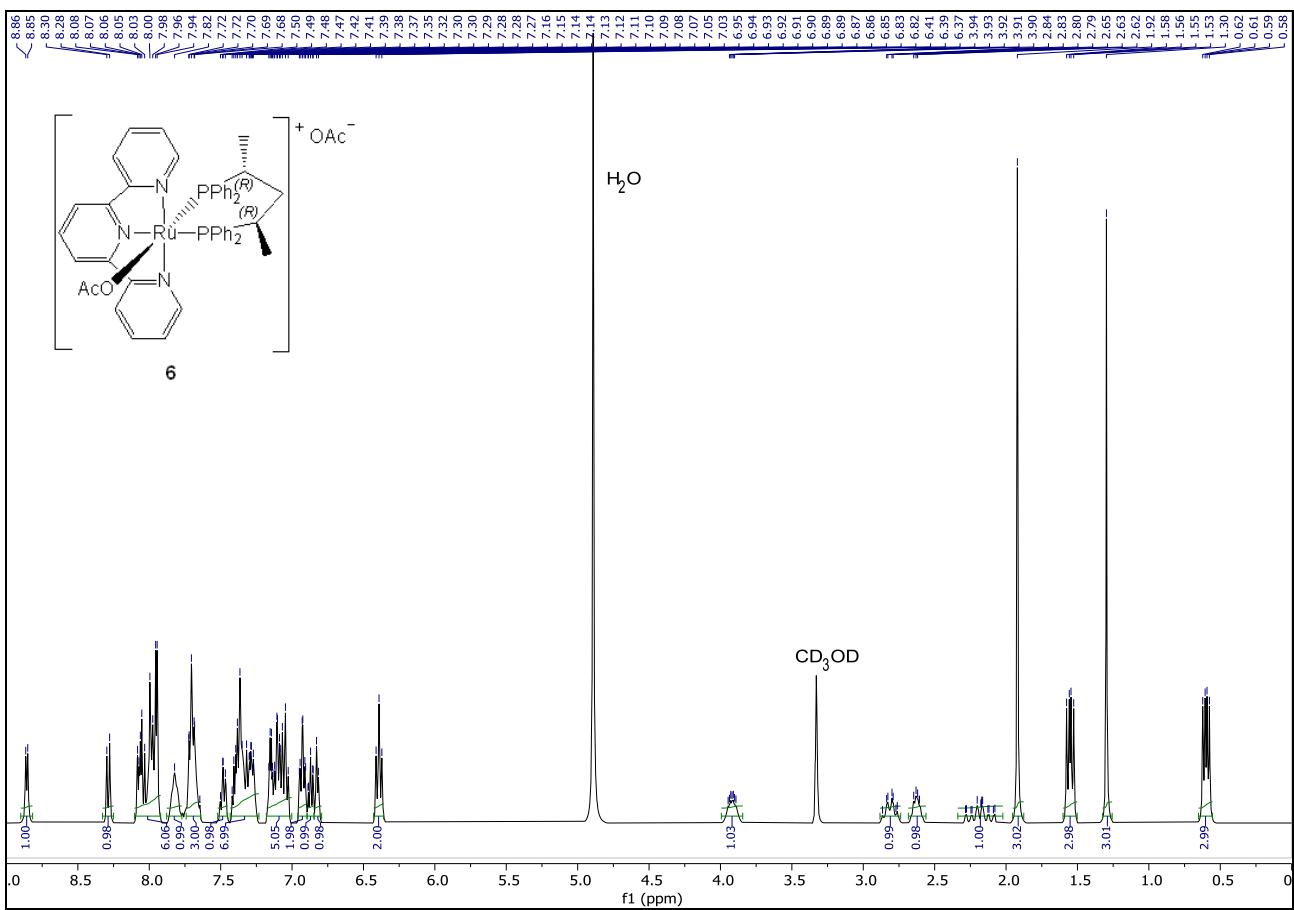
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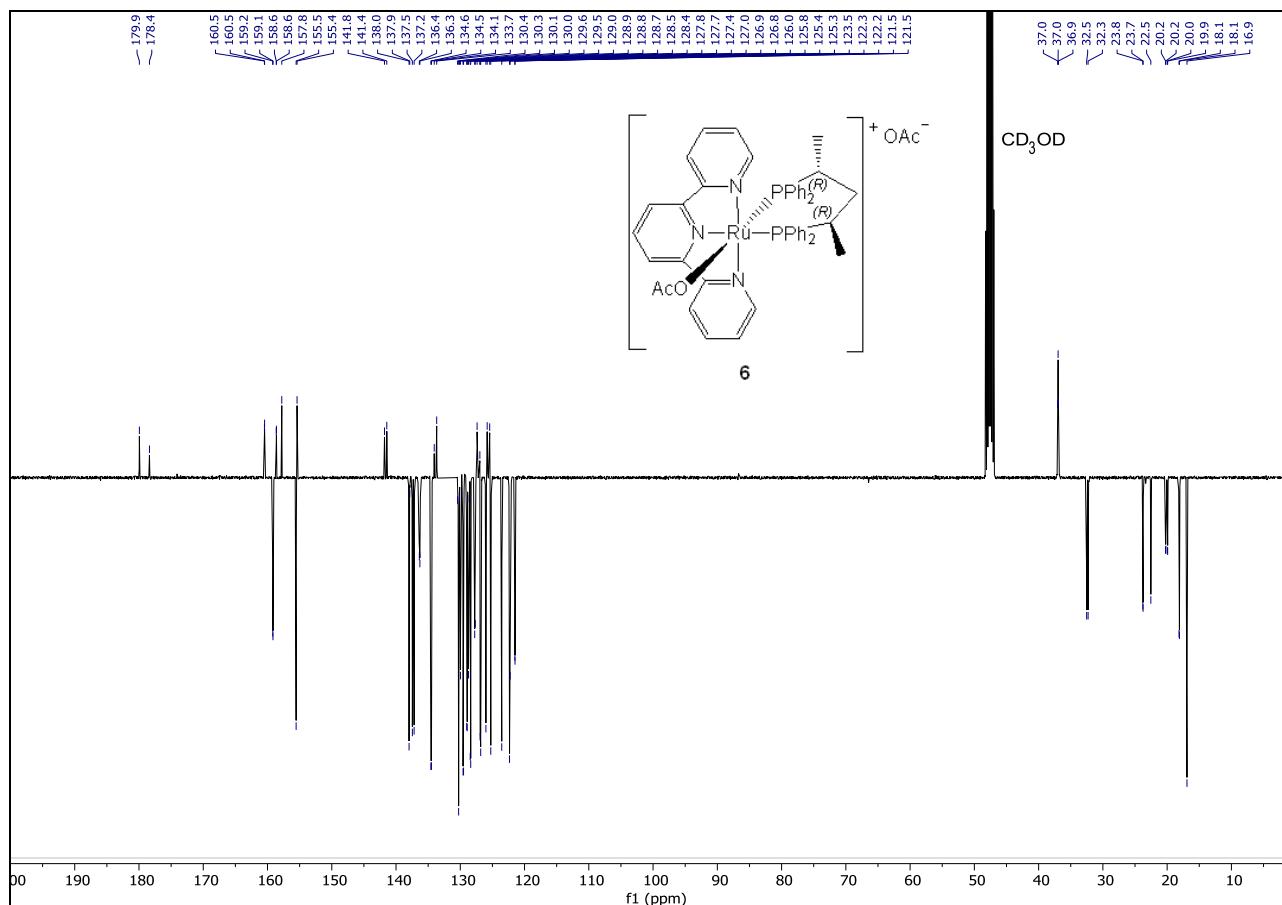
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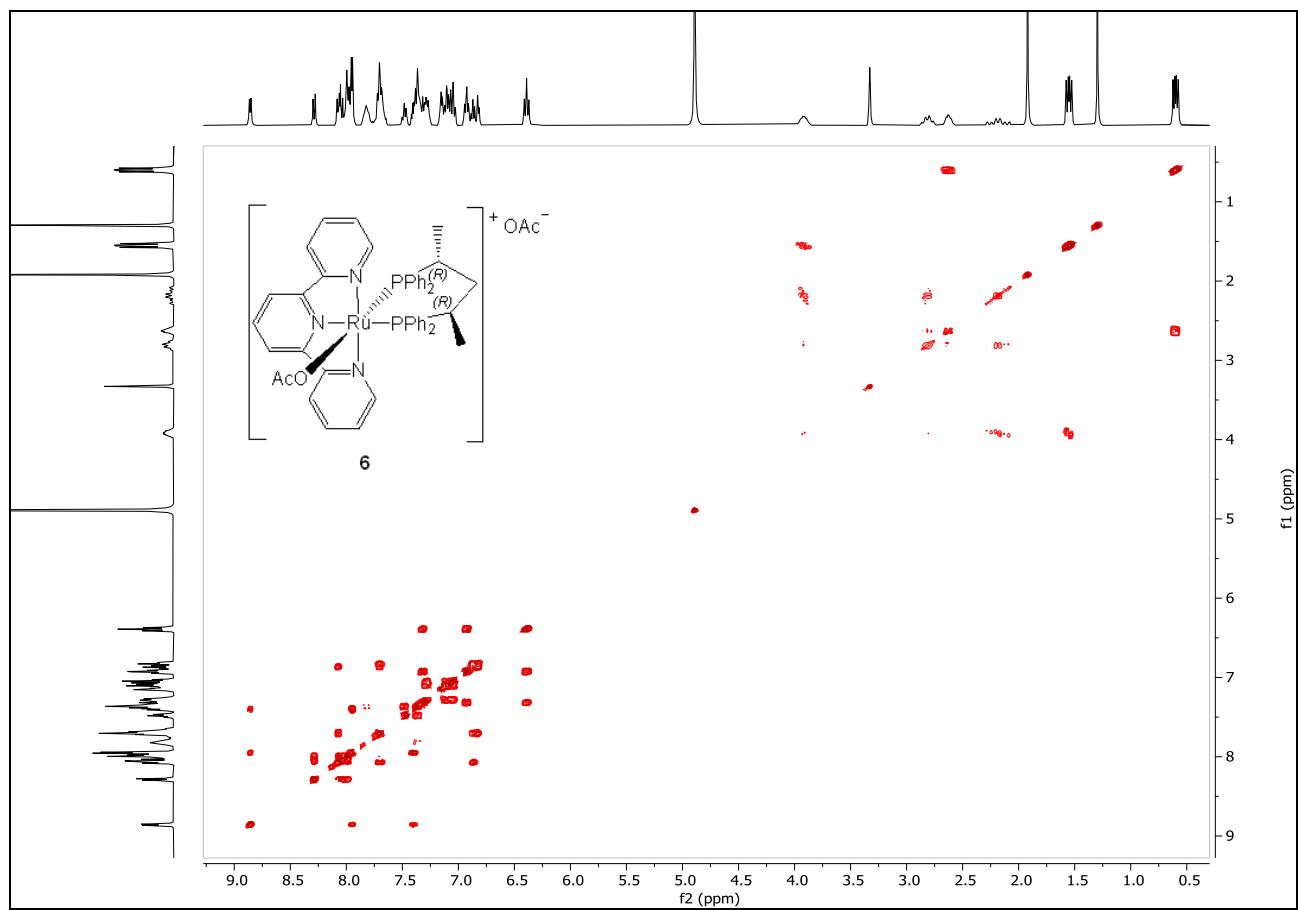
**Figure S24.**  $^{31}\text{P}\{\text{H}\}$  NMR spectrum (162.0 MHz) of  $[\text{Ru}(\eta^1\text{-OAc})(\text{NNN-tpy})((R,R)\text{-Skewphos})]\text{OAc}$  (**6**) in  $\text{CD}_3\text{OD}$  at 25 °C.



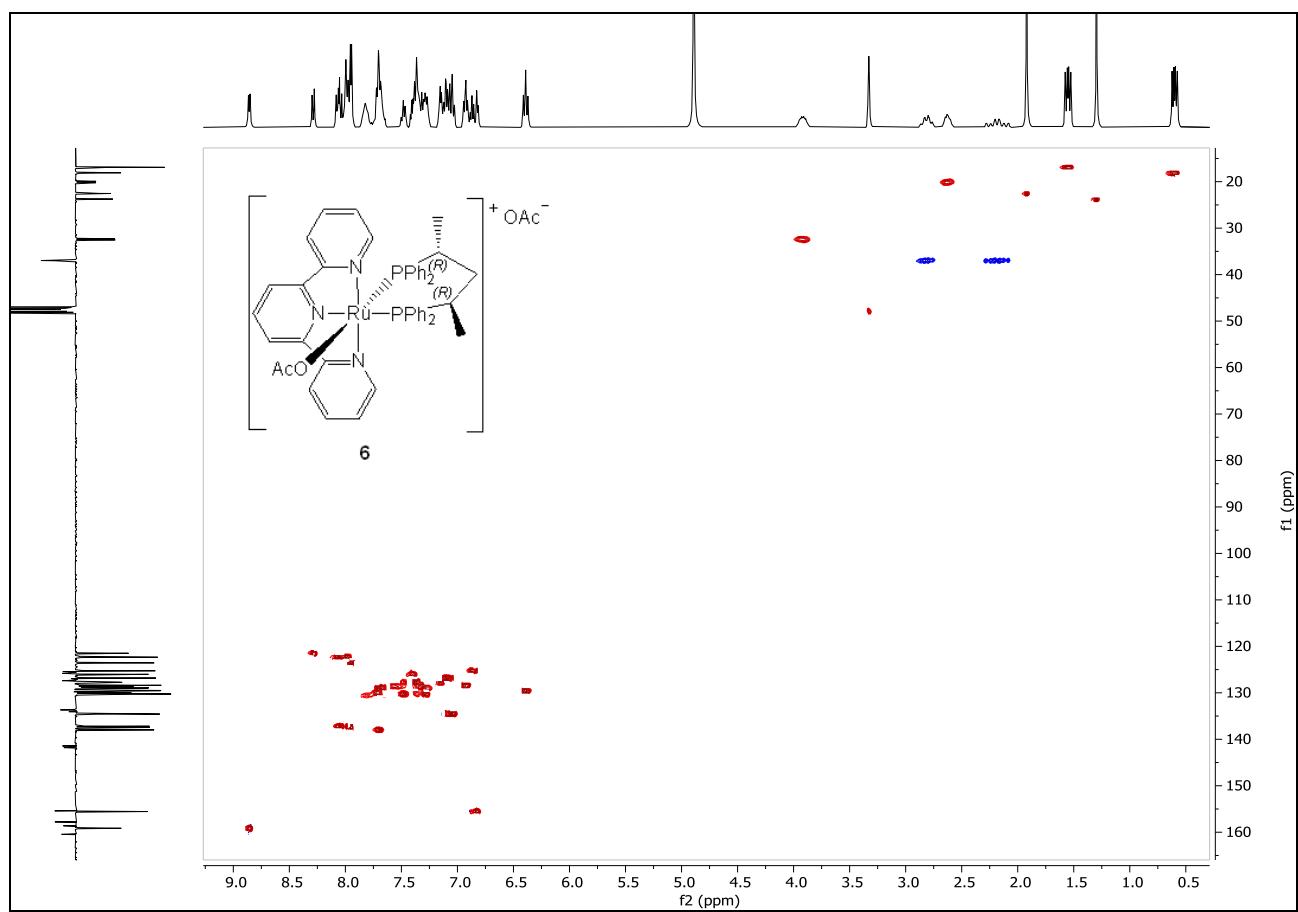
**Figure S25.**  $^1\text{H}$  NMR spectrum (400.1 MHz) of  $[\text{Ru}(\eta^1\text{-OAc})(\text{NNN-tpy})((R,R)\text{-Skewphos})]\text{OAc}$  (**6**) in  $\text{CD}_3\text{OD}$  at 25 °C.



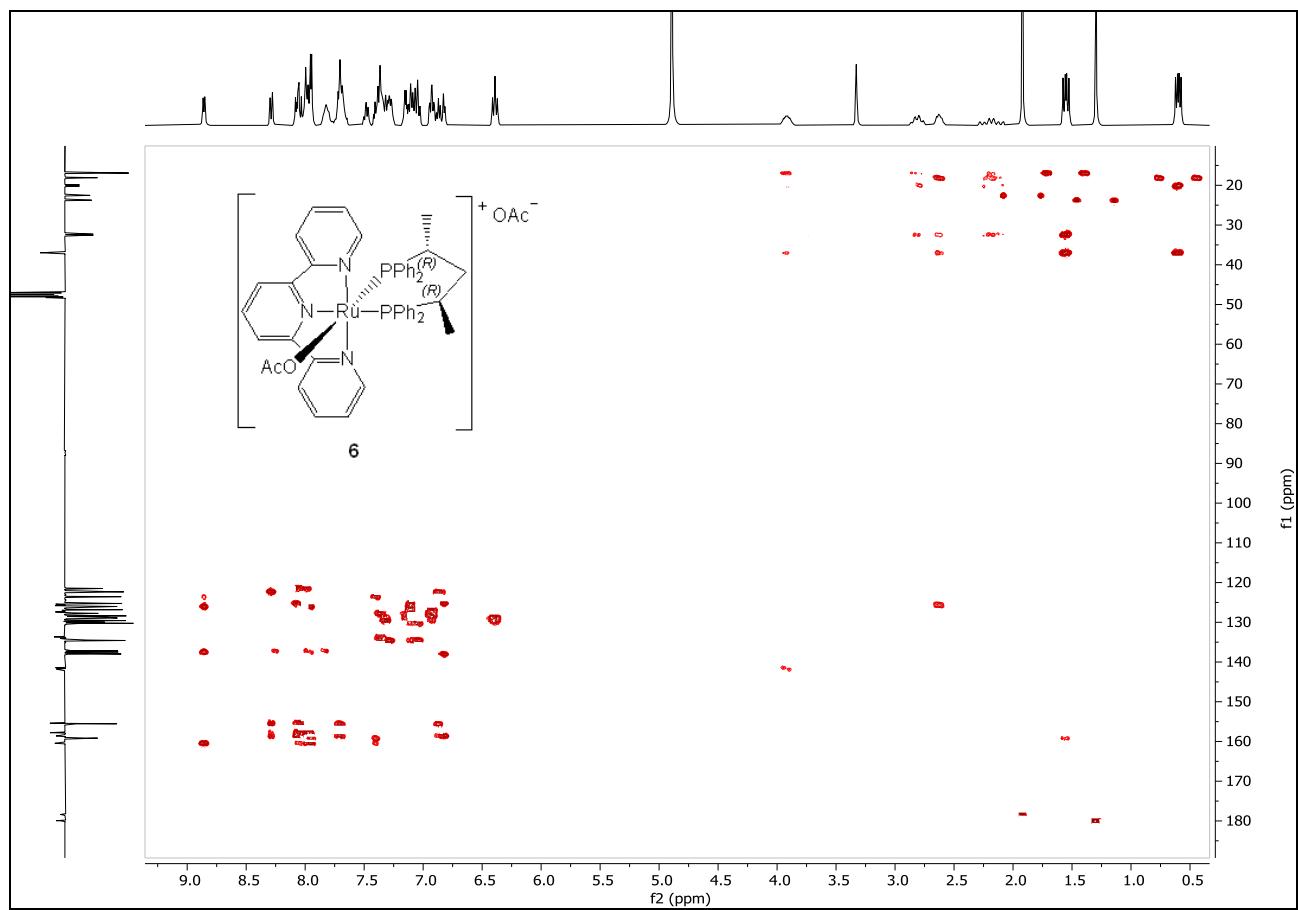
**Figure S26.**  $^{13}\text{C}\{^1\text{H}\}$  DEPTQ NMR spectrum (100.6 MHz) of  $[\text{Ru}(\eta^1\text{-OAc})(\text{NNN-tpy})((R,R)\text{-Skewphos})]\text{OAc}$  (**6**) in  $\text{CD}_3\text{OD}$  at 25 °C.



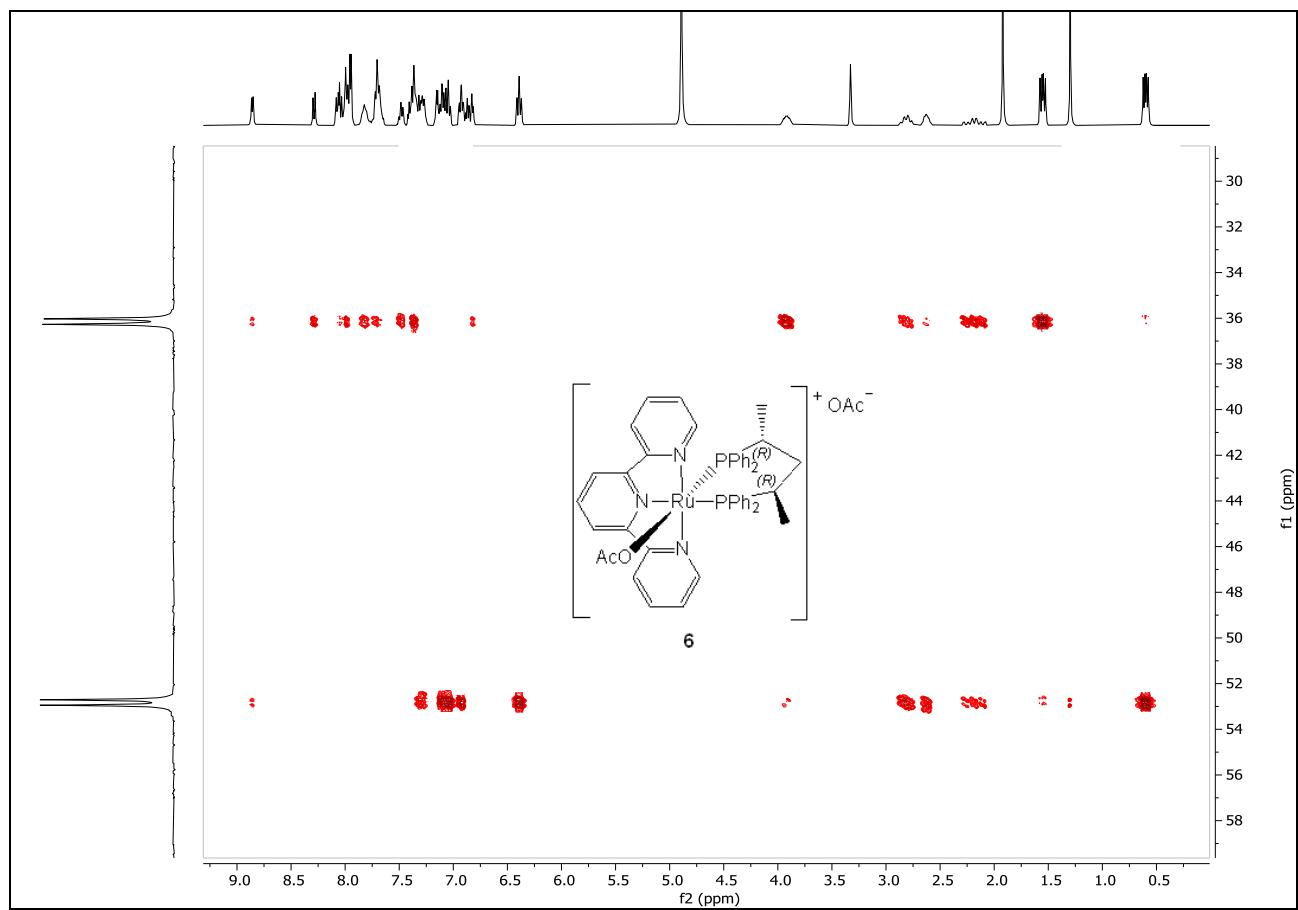
**Figure S27.**  $^1\text{H}$ - $^1\text{H}$  COSY 2D NMR spectrum of  $[\text{Ru}(\eta^1\text{-OAc})(\text{NNN-tpy})((R,R)\text{-Skewphos})]\text{OAc}$  (**6**) in  $\text{CD}_3\text{OD}$  at 25 °C.



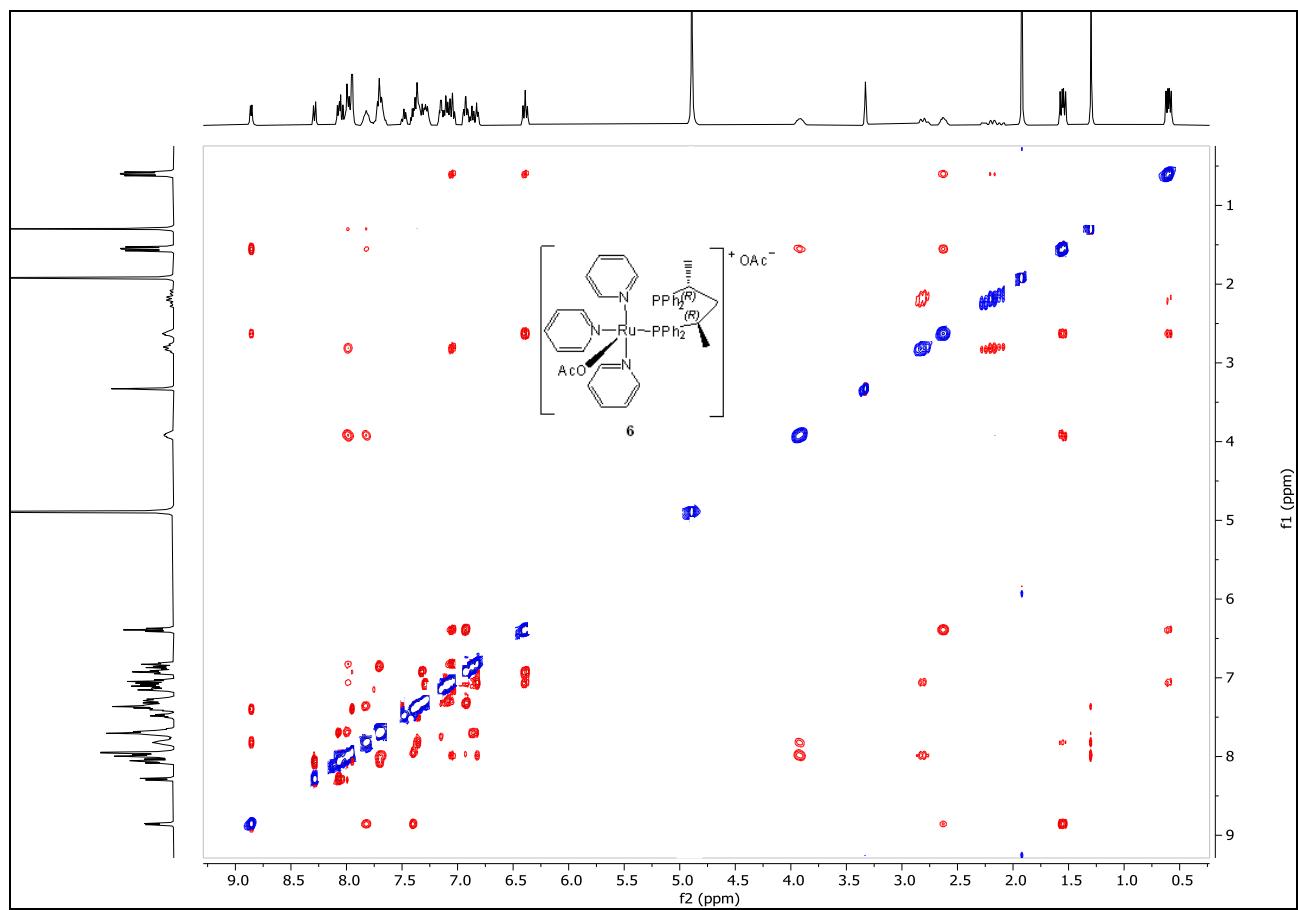
**Figure S28.**  $^1\text{H}$ - $^{13}\text{C}$  HSQC 2D NMR spectrum of  $[\text{Ru}(\eta^1\text{-OAc})(\text{NNN-tpy})((R,R)\text{-Skewphos})]\text{OAc}$  (**6**) in  $\text{CD}_3\text{OD}$  at  $25^\circ\text{C}$ .



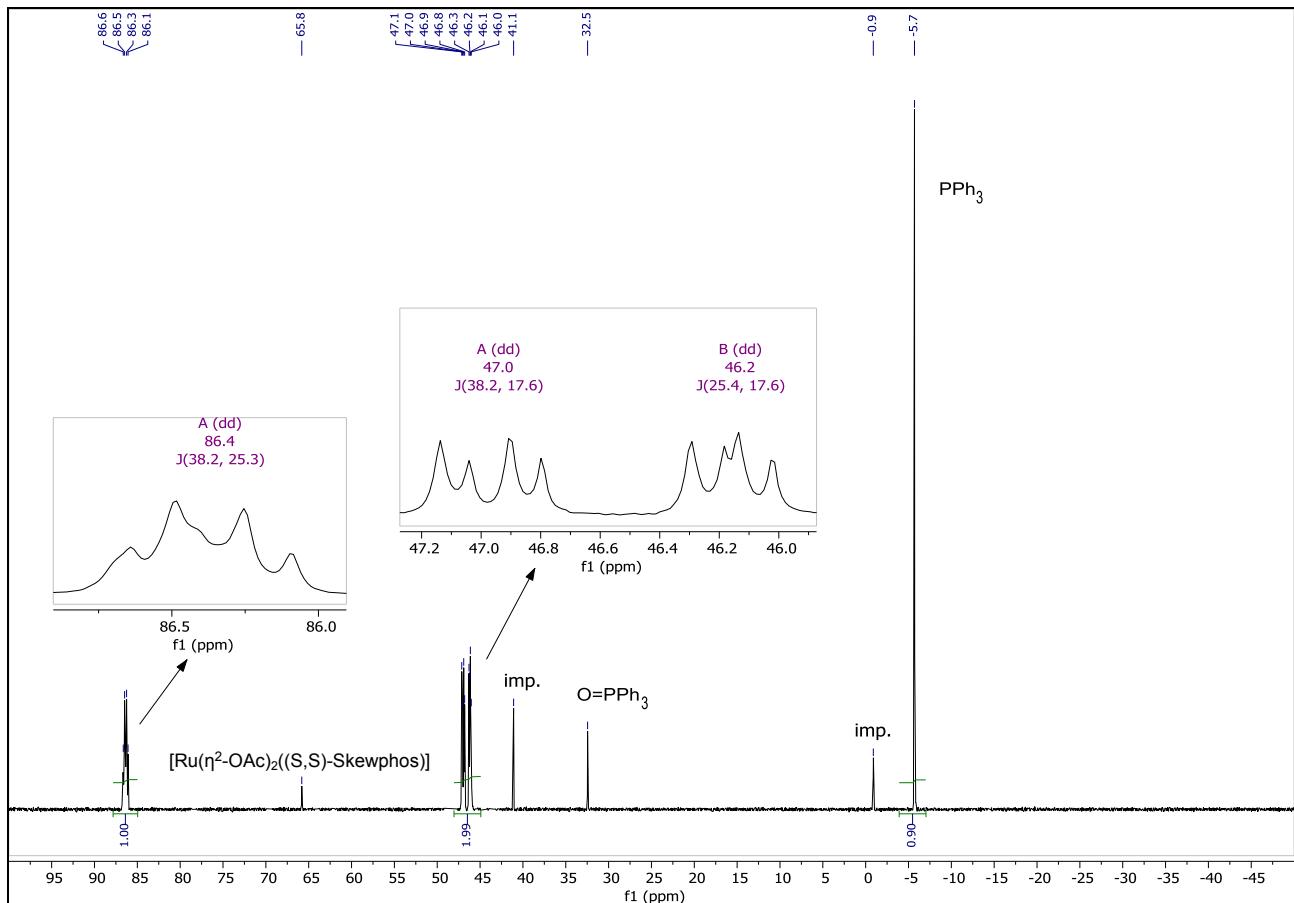
**Figure S29.**  $^1\text{H}$ - $^{13}\text{C}$  HMBC 2D NMR spectrum of  $[\text{Ru}(\eta^1\text{-OAc})(\text{NNN-tpy})((R,R)\text{-Skewphos})]\text{OAc}$  (**6**) in  $\text{CD}_3\text{OD}$  at 25 °C.



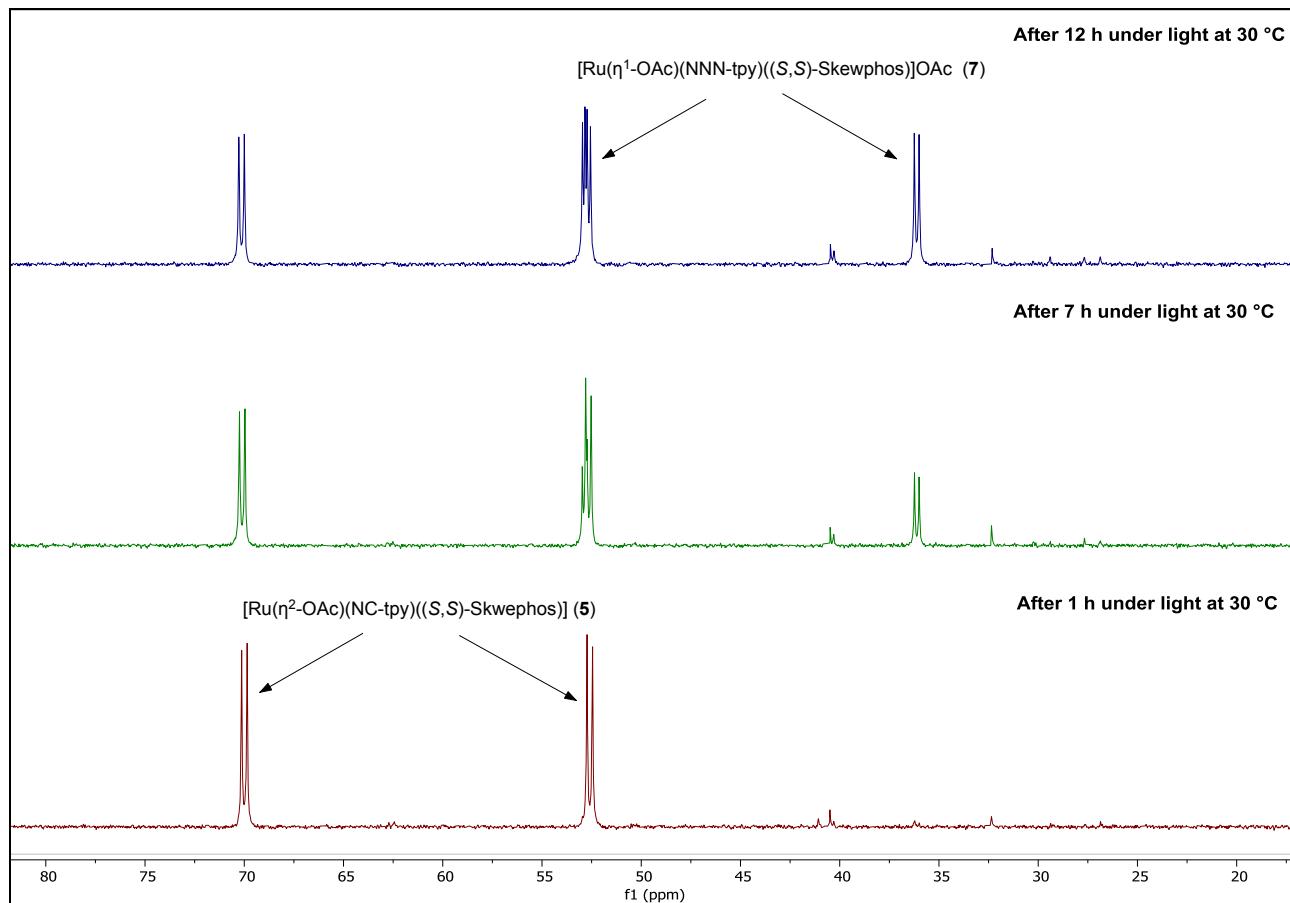
**Figure S30.**  $^1\text{H}$ - $^{31}\text{P}$  HMBC 2D NMR spectrum of  $[\text{Ru}(\eta^1\text{-OAc})(\text{NNN-tpy})((R,R)\text{-Skewphos})]\text{OAc}$  (**6**) in  $\text{CD}_3\text{OD}$  at 25 °C.



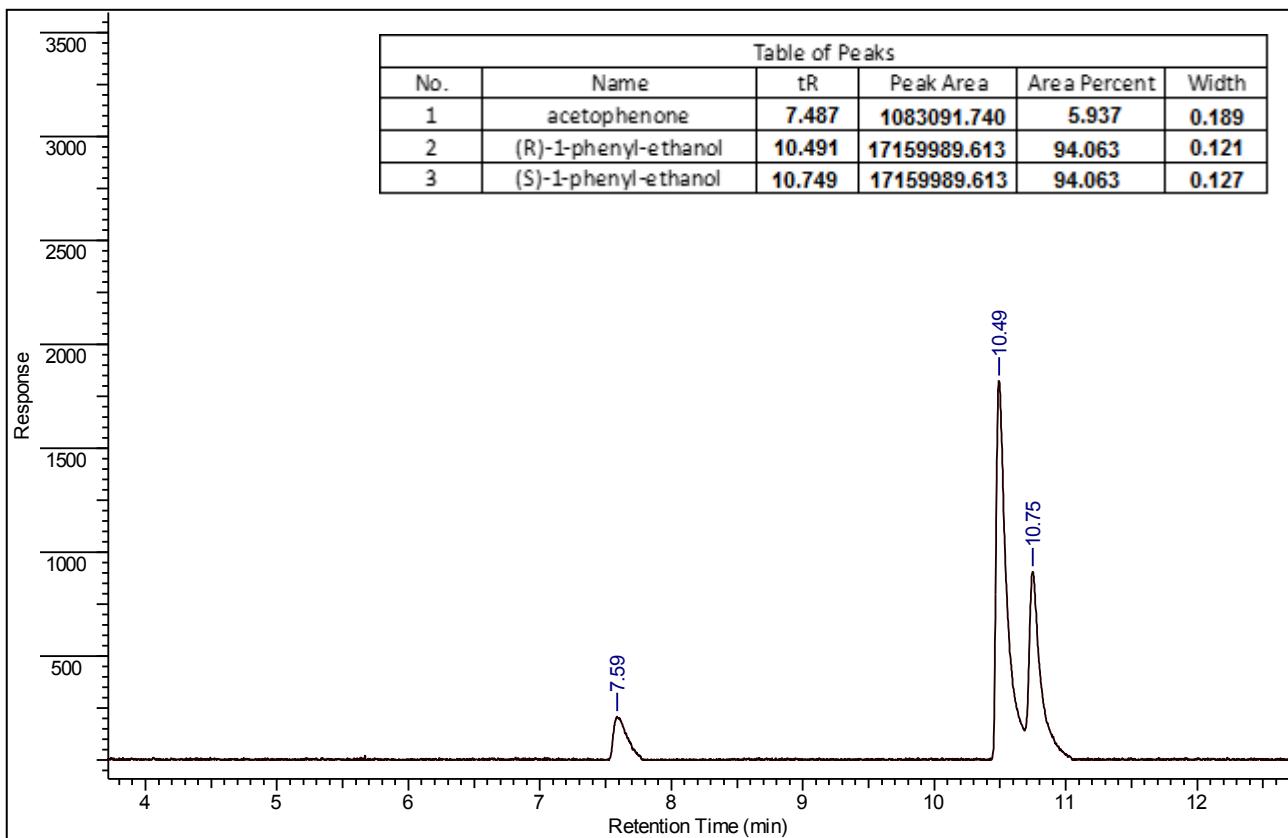
**Figure S31.**  $^1\text{H}$ - $^1\text{H}$  NOESY 2D NMR spectrum of  $[\text{Ru}(\eta^1\text{-OAc})(\text{NNN-tpy})((R,R)\text{-Skewphos})]\text{OAc}$  (**6**) in  $\text{CD}_3\text{OD}$  at  $25^\circ\text{C}$ .



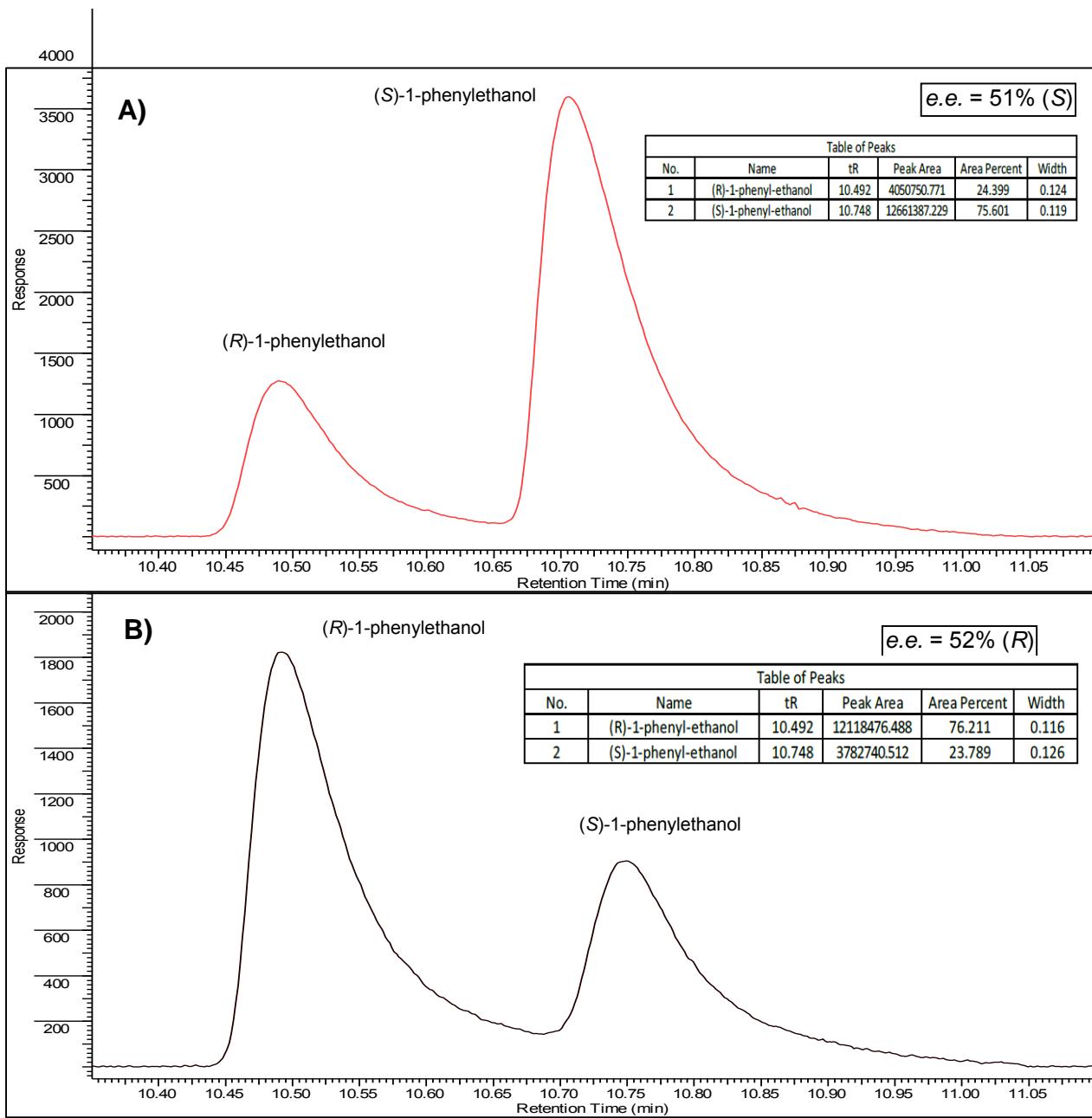
**Figure S32.** Evidence of the formation of the  $[\text{Ru}(\text{OAc})_2((S,S)\text{-Skewphos})(\text{PPh}_3)]$  intermediate in the reaction of (S,S)-Skewphos with  $[\text{Ru}(\eta^2\text{-OAc})_2(\text{PPh}_3)_2]$  in methanol at reflux in the  $^{31}\text{P}\{^1\text{H}\}$  NMR spectrum (162.0 MHz) with CD<sub>3</sub>OD as lock at 25 °C.



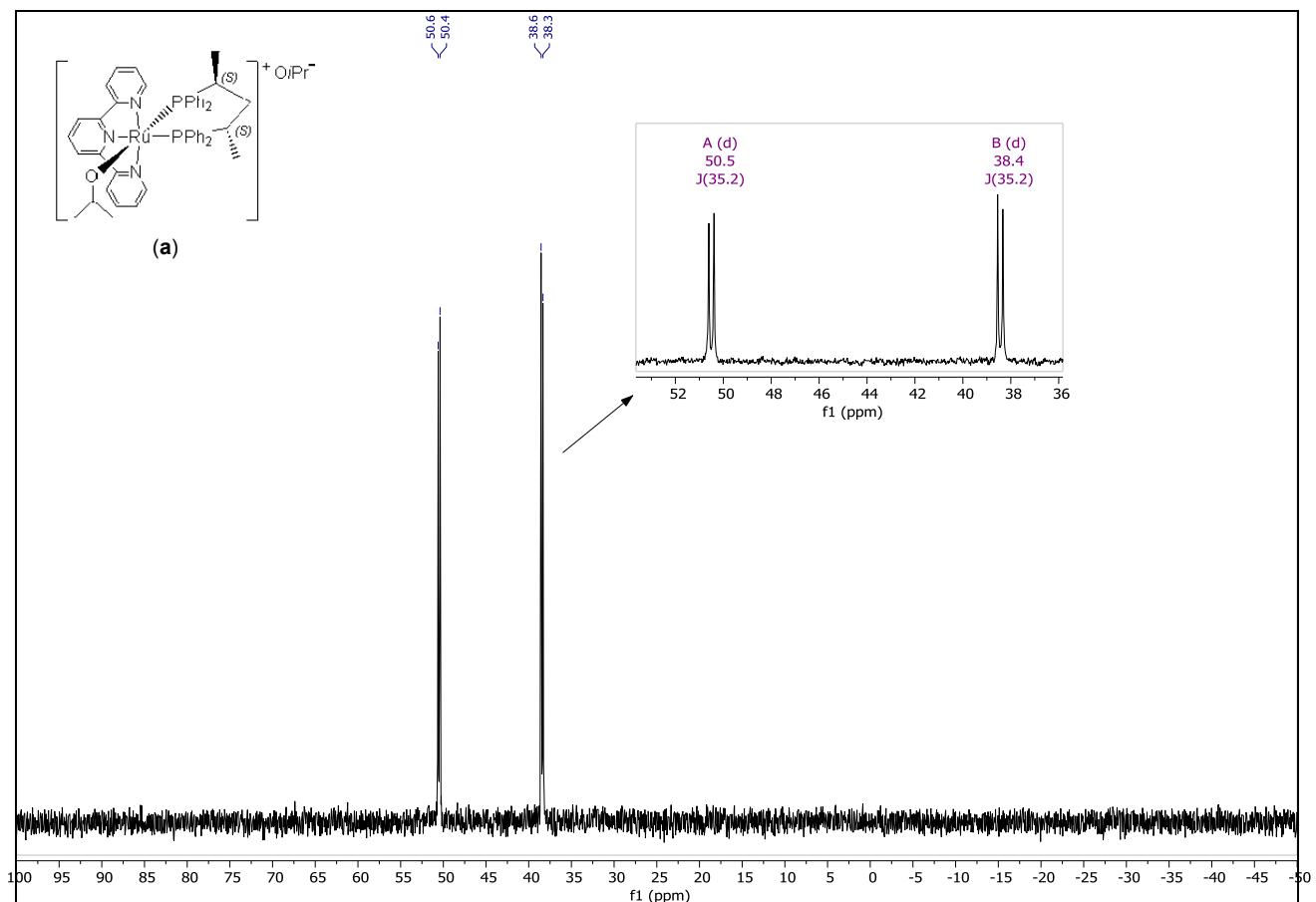
**Figure S33.** Conversion of complex  $[\text{Ru}(\eta^2\text{-OAc})(\text{NC-tpy})((S,S)\text{-Skewphos})]$  (5) to the cationic specie  $[\text{Ru}(\eta^1\text{-OAc})(\text{NNN-tpy})((S,S)\text{-Skewphos})]\text{OAc}$  (7) promoted by light with acetic acid (3 equiv) at  $30^\circ\text{C}$  in the  $^{31}\text{P}\{\text{H}\}$  NMR spectra (162.0 MHz) in  $\text{CD}_3\text{OD}$  (recorded at  $25^\circ\text{C}$ ).



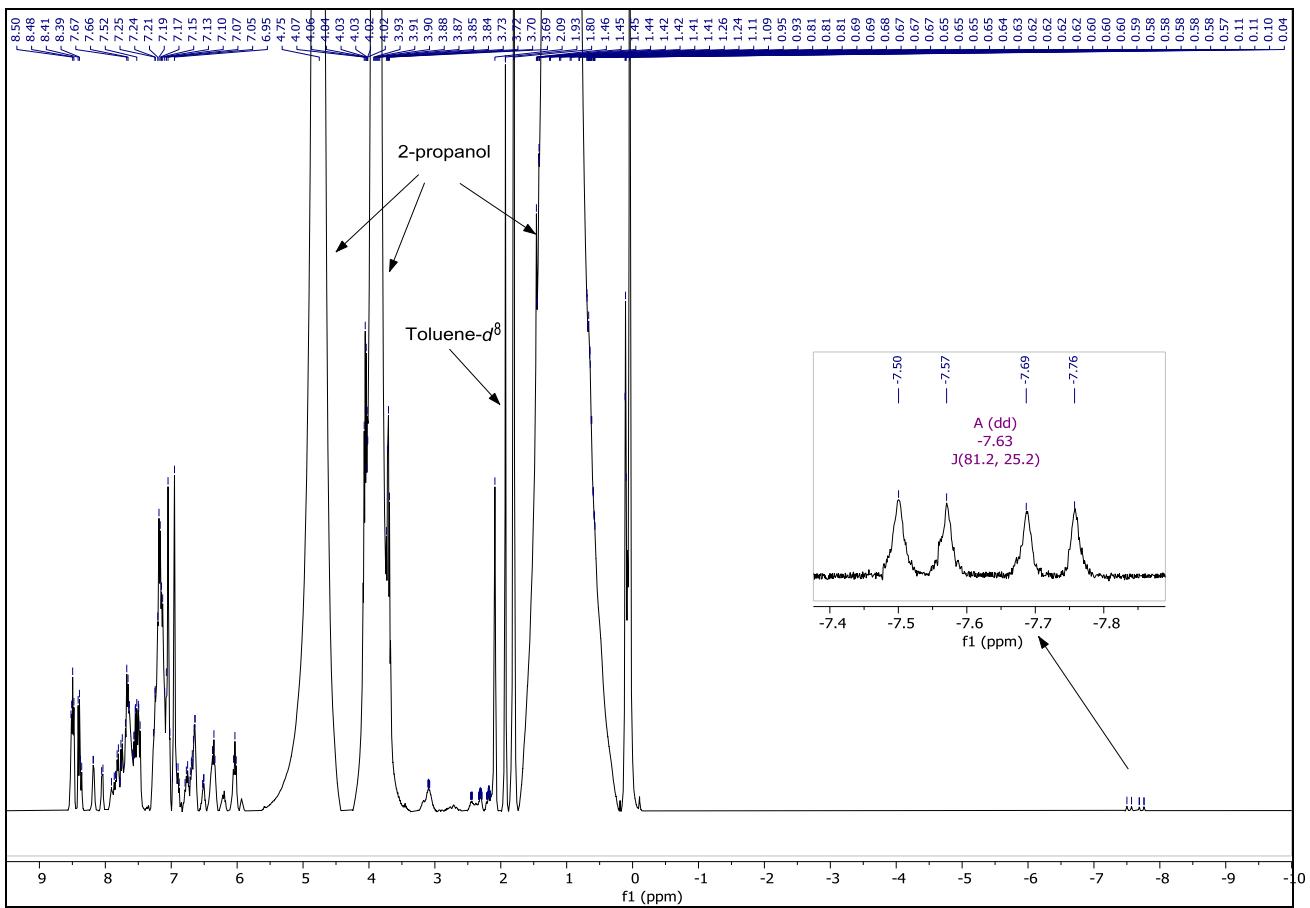
**Figure S34.** GC-FID chromatogram of the reaction mixture of the enantioselective photocatalytic TH of acetophenone in 2-propanol/MeOH 1:1 (v/v) at 30 °C and NaO*i*Pr 2 mol% promoted by complex **7** at S/C 1000 after 28 h of overall irradiation. GC analyses were performed with a Varian CP-3380 gas chromatograph equipped with a 25 m length MEGADEX-ETTBDMs- $\beta$  chiral column with hydrogen (5 psi) as the carrier gas and flame ionization detector (FID). The injector and detector temperature was 250 °C, with initial T = 95 °C ramped to 140 °C at 3 °C/min for a total of 15 min of analysis.



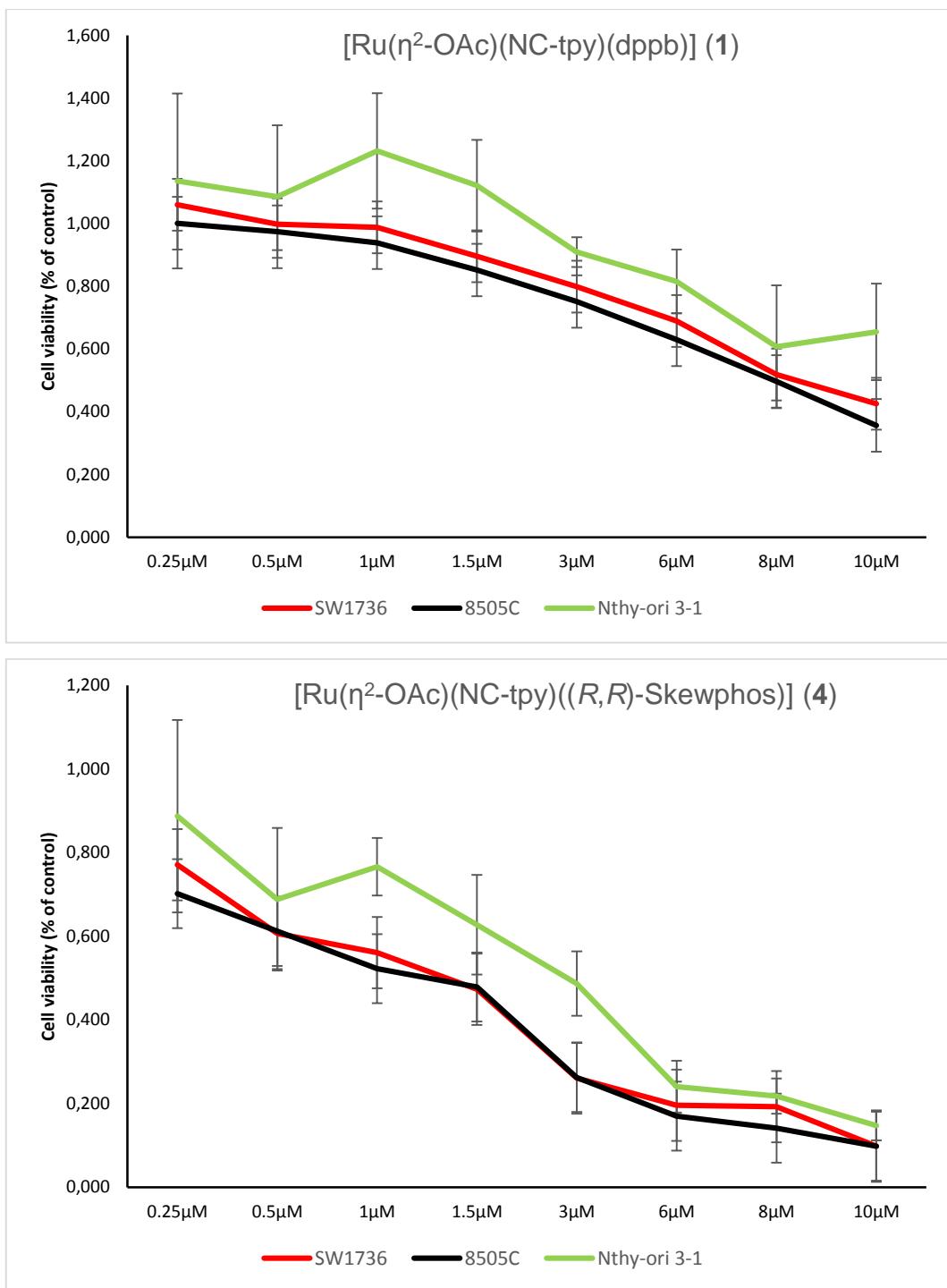
**Figure S35.** Comparison between the GC-FID chromatograms of the enantioselective photocatalytic TH of acetophenone in 2-propanol/MeOH 1:1 (v/v) with NaO*i*Pr 2 mol% at 30 °C promoted by complex **6** (A) and **7** (B) at S/C 1000.



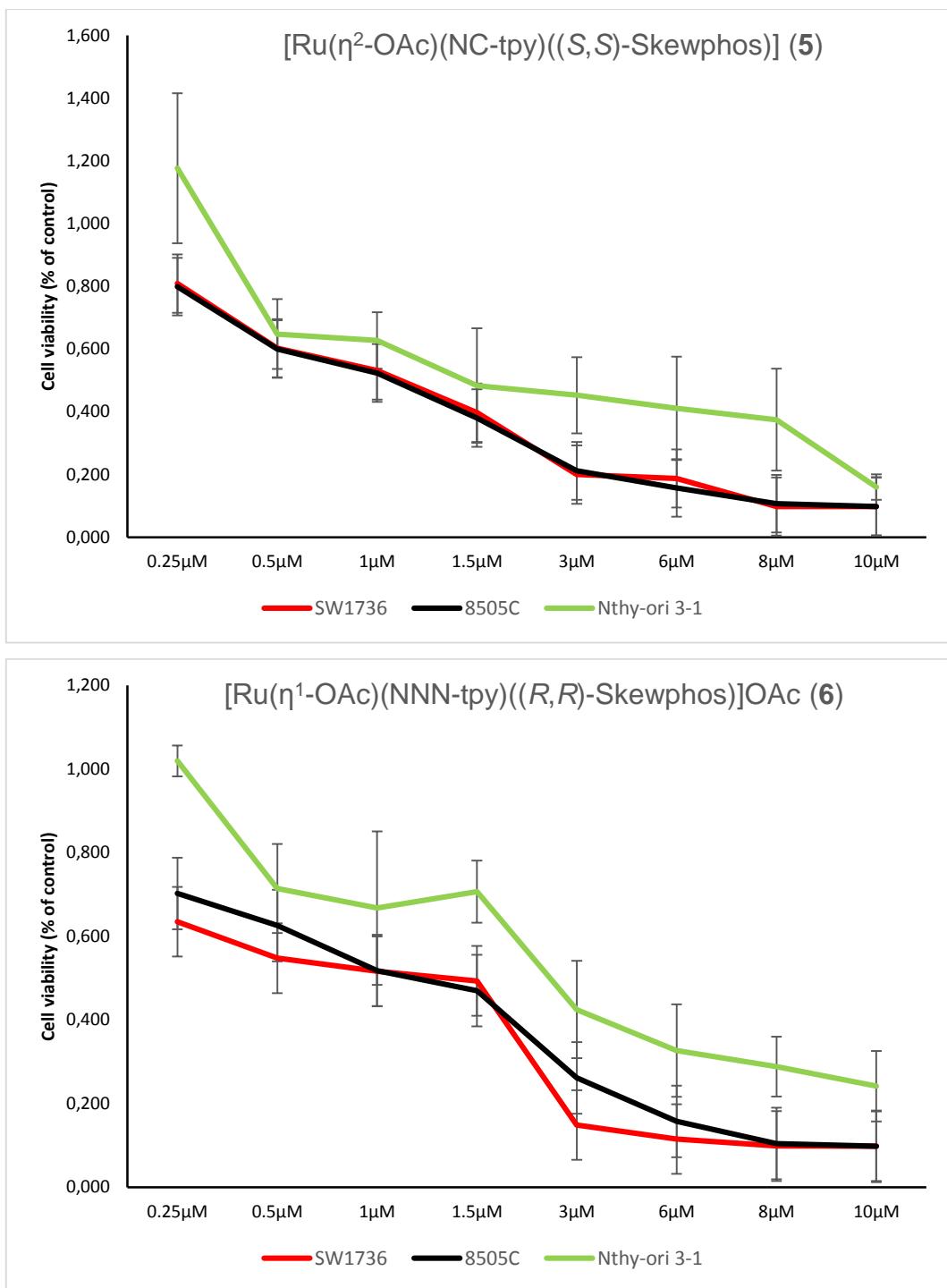
**Figure S36.**  $^{31}\text{P}\{\text{H}\}$  NMR spectrum (162.0 MHz) of  $[\text{Ru}(\text{O}i\text{Pr})(\text{NNN-tpy})((S,S)\text{-Skewphos})](\text{O}i\text{Pr})$  (a) in 2-propanol- $d^8$  at 25 °C, obtained after 1 h of visible light irradiation from complex 7 in presence of NaO*i*Pr (3 equiv).



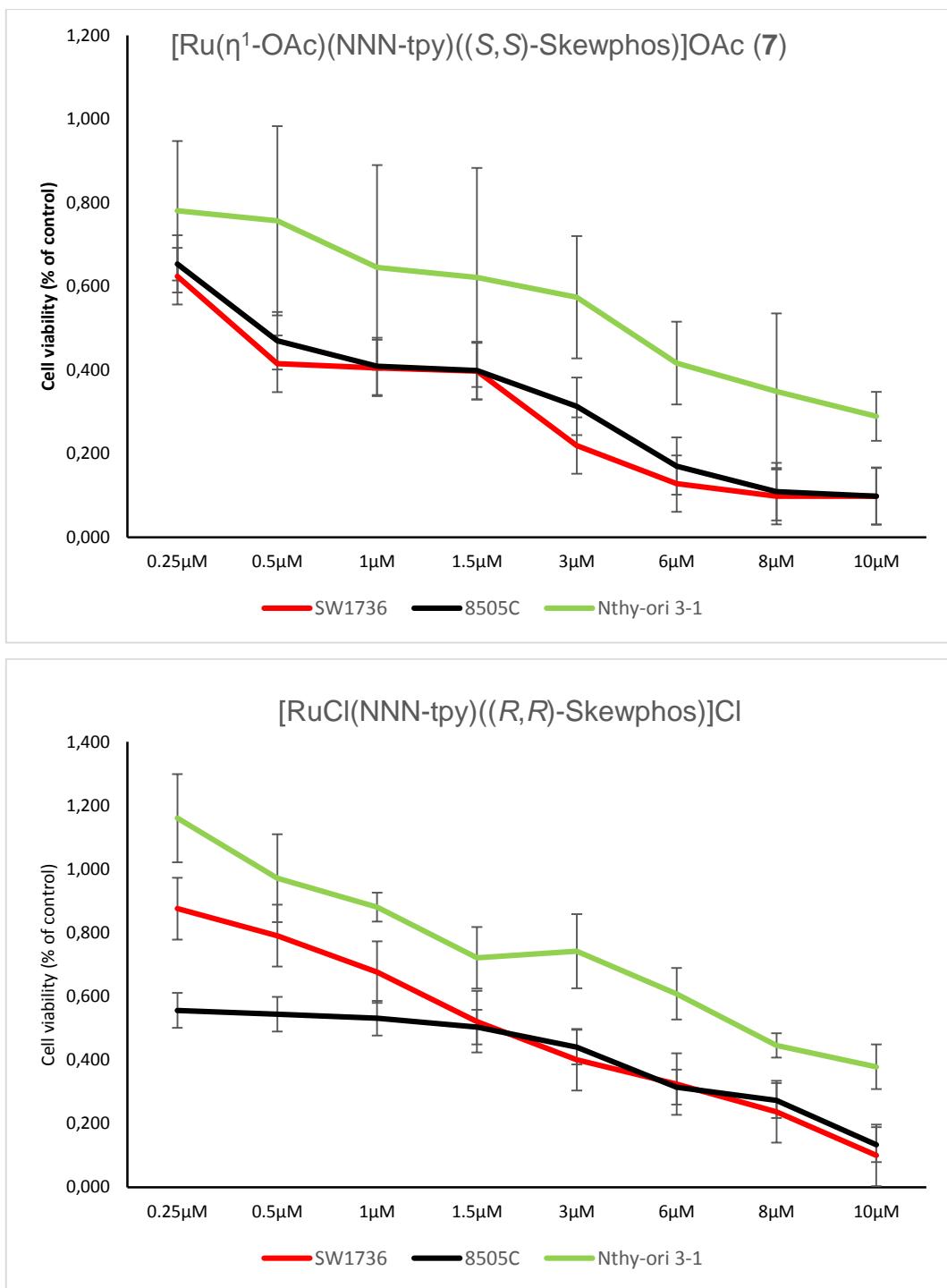
**Figure S37.**  ${}^1\text{H}$  NMR spectrum (400.1 MHz) with evidences of the formation of  $[\text{RuH}(\text{NNN-tpy})((S,S)\text{-Skewphos})](\text{OEt})$  (**b**) in 2-propanol/Toluene- $d^8$  1:1 (v/v) at 25 °C (in mixture with uncharacterized compounds), obtained after 6 h of visible light irradiation from complex **5** in presence of NaOEt (3 equiv).



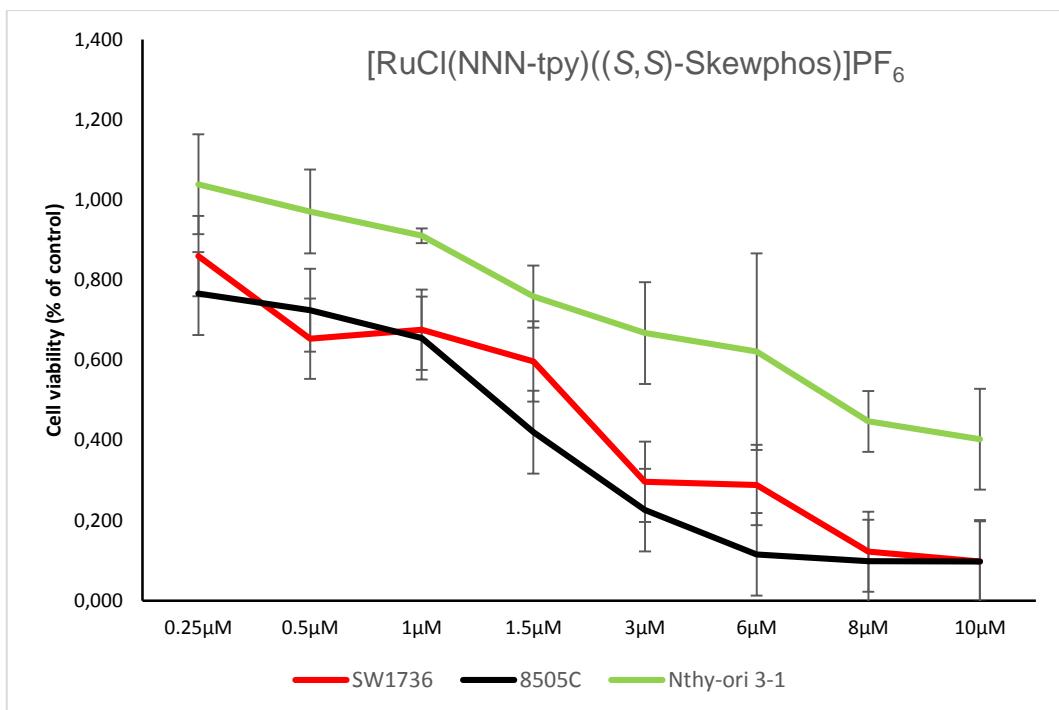
**Figure S38.** Effect of complexes **1** and **4** on ATC cell viability in SW1736 (red), 8505C (black) and Nthy-ori 3-1 (light green) cells. Cell viability was evaluated by using the MTT assay and expressed as the percentage of control (cells treated with DMSO alone). Each point represents the mean value of six-fold determinations. P-values obtained by statistical analysis (student's t test) are listed in the Table S1.



**Figure S39.** Effect of complexes **5** and **6** on ATC cell viability in SW1736 (red), 8505C (black) and Nthy-ori 3-1 (light green) cells. Cell viability was evaluated by using the MTT assay and expressed as the percentage of control (cells treated with DMSO alone). Each point represents the mean value of six-fold determinations. P-values obtained by statistical analysis (student's t test) are listed in the Table S1.



**Figure S40.** Effect of complexes **7** and [RuCl(NNN-tpy)((*R,R*)-Skewphos)]Cl on ATC cell viability in SW1736 (red), 8505C (black) and Nthy-ori 3-1 (light green) cells. Cell viability was evaluated by using the MTT assay and expressed as the percentage of control (cells treated with DMSO alone). Each point represents the mean value of six-fold determinations. P-values obtained by statistical analysis (student's t test) are listed in the Table S1.



**Figure S41.** Effect of complex [RuCl(NNN-tpy)((S,S)-Skewphos)]PF<sub>6</sub> on ATC cell viability in SW1736 (red), 8505C (black) and Nthy-ori 3-1 (light green) cells. Cell viability was evaluated by using the MTT assay and expressed as the percentage of control (cells treated with DMSO alone). Each point represents the mean value of six-fold determinations. P-values obtained by statistical analysis (student's t test) are listed in the Table S1.

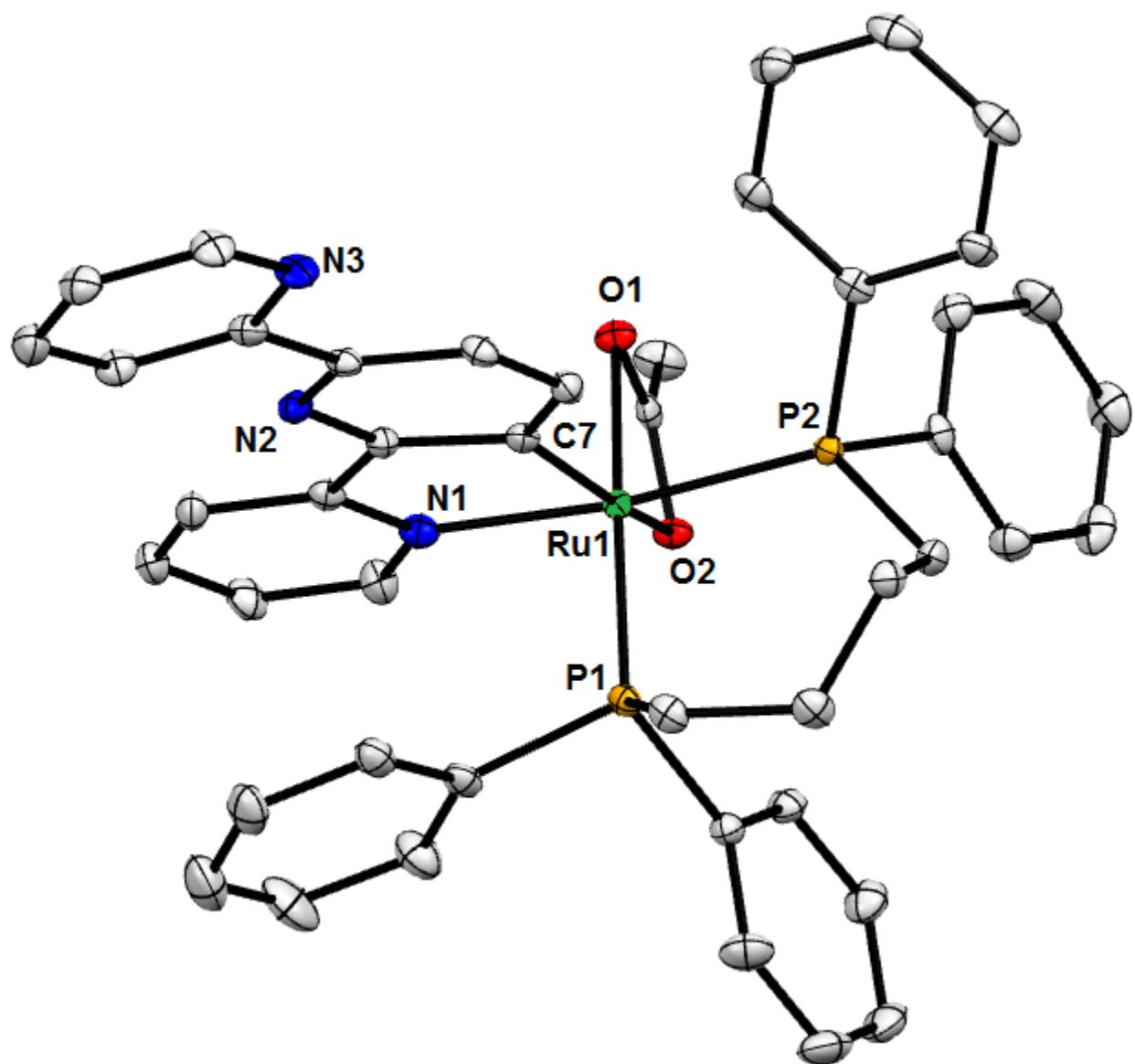
	[RuCl(NNN-tpy)( <i>R,R</i> -Skewphos)] <i>C</i>	[RuCl(NNN-tpy)( <i>S,S</i> -Skewphos)]PF <sub>6</sub>	[Ru(η <sup>2</sup> -OAc)(NNN-tpy)( <i>R,R</i> -Skewphos)]OAc <b>6</b>	[Ru(η <sup>2</sup> -OAc)(NC-tpy)(dppb)] <b>1</b>	[Ru(η <sup>2</sup> -OAc)(NC-tpy)( <i>R,R</i> -Skewphos)] <b>4</b>	[Ru(η <sup>2</sup> -OAc)(NNN-tpy)( <i>S,S</i> -Skewphos)]OAc <b>7</b>	[Ru(η <sup>2</sup> -OAc)(NC-tpy)( <i>S,S</i> -Skewphos)] <b>5</b>
complex dose	SW1736 8505C Nthy-ori 3:1	SW1736 8505C Nthy-ori 3:1	SW1736 8505C Nthy-ori 3:1	SW1736 8505C Nthy-ori 3:1	SW1736 8505C Nthy-ori 3:1	SW1736 8505C Nthy-ori 3:1	SW1736 8505C Nthy-ori 3:1
0.25µM	*	*	NS	***	*	NS	NS
0.5µM	NS	*	NS	*	NS	NS	NS
1µM	*	*	***	*	***	*	NS
1.5µM	***	*	***	***	*	NS	NS
3µM	***	*	***	***	*	NS	NS
6µM	***	*	****	****	*	NS	NS
8µM	***	*	****	****	***	NS	NS
10µM	****	****	****	****	***	NS	NS

**Table S1.** Statistical analysis for tested complexes. For each dose of each complex, the p-value in each of the two cell lines is listed. NS: not significant, \* p<0.05, \*\* p<0.01, \*\*\* p<0.001, \*\*\*\* p<0.0001.

## **Single Crystal X-Ray Structure Determination of Compound 1 (CCDC 2302606).**

### **General Data**

Data were collected on a Bruker D8 Venture single crystal x-ray diffractometer equipped with a CPAD detector (Bruker Photon II), an IMS microsource with MoK $\alpha$  radiation ( $\lambda = 0.71073 \text{ \AA}$ ) and a Helios optic using the APEX3 software package.<sup>83</sup> The measurement was performed on a single crystal coated with perfluorinated ether which was fixed on top of a kapton micro sampler and frozen under a stream of cold nitrogen. A matrix scan was used to determine the initial lattice parameters. Reflections were corrected for Lorentz and polarisation effects, scan speed, and background using SAINT.<sup>84</sup> Absorption correction, including odd and even ordered spherical harmonics was performed using SADABS.<sup>85</sup> Space group assignment was based upon systematic absences, E statistics, and successful refinement of the structure. The structure was solved using SHELXT with the aid of successive difference Fourier maps, and was refined against all data using SHELXL in conjunction with SHELXLE.<sup>86-88</sup> Hydrogen atoms were calculated in ideal positions as follows: Methyl hydrogen atoms were refined as part of rigid rotating groups, with a C–H distance of 0.98  $\text{\AA}$  and  $U_{\text{iso}}(\text{H}) = 1.5 \cdot U_{\text{eq}}(\text{C})$ . Non-methyl H atoms were placed in calculated positions and refined using a riding model with methylene, aromatic, and other C–H distances of 0.99  $\text{\AA}$ , 0.95  $\text{\AA}$  and 1.00  $\text{\AA}$ , respectively, and  $U_{\text{iso}}(\text{H}) = 1.2 \cdot U_{\text{eq}}(\text{C})$ . Non-hydrogen atoms were refined with anisotropic displacement parameters. Full-matrix least-squares refinements were carried out by minimizing  $\Sigma w(F_o^2 - F_c^2)^2$  with the SHELXL weighting scheme.<sup>86</sup> Neutral atom scattering factors for all atoms and anomalous dispersion corrections for the non-hydrogen atoms were taken from *International Tables for Crystallography*.<sup>89</sup> Images of the crystal structure were generated with Mercury.<sup>90</sup> CCDC 2302606 contains the supplementary crystallographic data for this paper. These data are provided free of charge by The Cambridge Crystallographic Data Centre.



**Figure S42.** ORTEP style plot of compound **1** (one out of two independent molecules) in the solid state (CCDC 2302606) as determined by single crystal X-ray diffraction. Ellipsoids are displayed at the 50% probability level. Hydrogen atoms and co-crystallized solvent molecules are omitted for clarity.

## Single Crystal X-Ray Structure Determination of Compound 1 (CCDC 2302606).

### Detailed Crystallographic Data.

Diffractometer operator:	C. Jandl
Scanspeed	2-20 s per frame
dx	90 mm
Frames:	2785 measured in 10 XYZ data sets
phi-scans with delta phi	0.5
omega-scans with delta omega	0.5
shutterless mode	

### Crystal Data:

Chemical formula: [C <sub>45</sub> H <sub>41</sub> N <sub>3</sub> O <sub>2</sub> P <sub>2</sub> Ru]·1/2(C <sub>7</sub> H <sub>16</sub> )	Density (calculated) = <u>1.396</u> g/cm <sup>3</sup>
Formula weight <u>868.91</u>	Absorption coefficient = <u>0.50</u> mm <sup>-1</sup>
Monoclinic, <u>P 1 21/n 1</u>	<u>Mo Kα</u> radiation, $\lambda = 0.71073$ Å
$a = 13.196(9)$ Å	$\alpha = 90^\circ$
$b = 44.86(3)$ Å	$\beta = 101.096(16)^\circ$
$c = 14.234(10)$ Å	$\gamma = 90^\circ$
$V = 8269.(10)$ Å <sup>3</sup>	<u>Yellow fragment</u>
$Z = 8$	<u>0.086 × 0.125 × 0.138</u> mm
$F(000) = 3608$	

### Data collection:

<u>Bruker D8 Venture Duo IMS</u> diffractometer	<u>15129</u> independent reflections
Radiation source: <u>IMS microsource, Mo</u>	<u>12749</u> reflections with $I > 2\sigma(F^2)$
<u>Helios optic</u> monochromator	$R_{\text{int}} = 0.1777$
Theta range for data collection	$\theta_{\max} = 25.38^\circ$ , $\theta_{\min} = 1.93^\circ$

Index ranges	<u>-15&lt;=h&lt;=15, -54&lt;=k&lt;=54, -17&lt;=l&lt;=17</u>
Absorption correction	Multi-Scan, <u>SADABS 2016/2, Bruker</u>
Max. and min. transmission	0.9580 and 0.9340
<u>189179</u> measured reflections	Coverage of independent reflections = 99.9%

Data refinement:

Refinement method	Full-matrix least-squares on $F^2$
Refinement program	SHELXL-2018/3 (Sheldrick, 2018)
Function minimized	$\Sigma w(Fo^2 - Fc^2)^2$
Data / restraints / parameters	<u>15129 / 0 / 1022</u>
Final R indices	<u>15129</u> data; $I > 2\sigma(I)$ $R_1 = 0.0472$ , $wR_2 = 0.0904$
Weighting scheme	all data $R_1 = 0.0614$ , $wR_2 = 0.0947$ $w = 1/[\sigma^2(Fo^2) + (0.0201P)^2 + 12.9123P]$ where $P = (Fo^2 + 2Fc^2)/3$
$\Delta/\sigma_{\text{max}}$	<u>0.003</u>
Goodness-of-fit on $F^2$	<u>1.109</u>
Largest diff. peak and hole	<u>0.524 and -0.739 eÅ<sup>-3</sup></u>
R.M.S. deviation from mean	<u>0.085 eÅ<sup>-3</sup></u>

## References

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- 86) G. M. Sheldrick, Crystal structure refinement with SHELXL. *Acta Crystallogr. Sect. A* **2015**, *71*, 3-8.
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- 88) C. B. Hübschle, G. M. Sheldrick, B. Dittrich, ShelXle: a Qt graphical user interface for SHELXL. *J. Appl. Cryst.* **2011**, *44*, 1281-1284.
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