

SUPPLEMENTARY MATERIAL

Half-Sandwich Rhodium Complexes with Releasable N-Donor Monodentate Ligands: Solution Chemical Properties and Possibility for Acidosis Activation

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FIGURE S1.

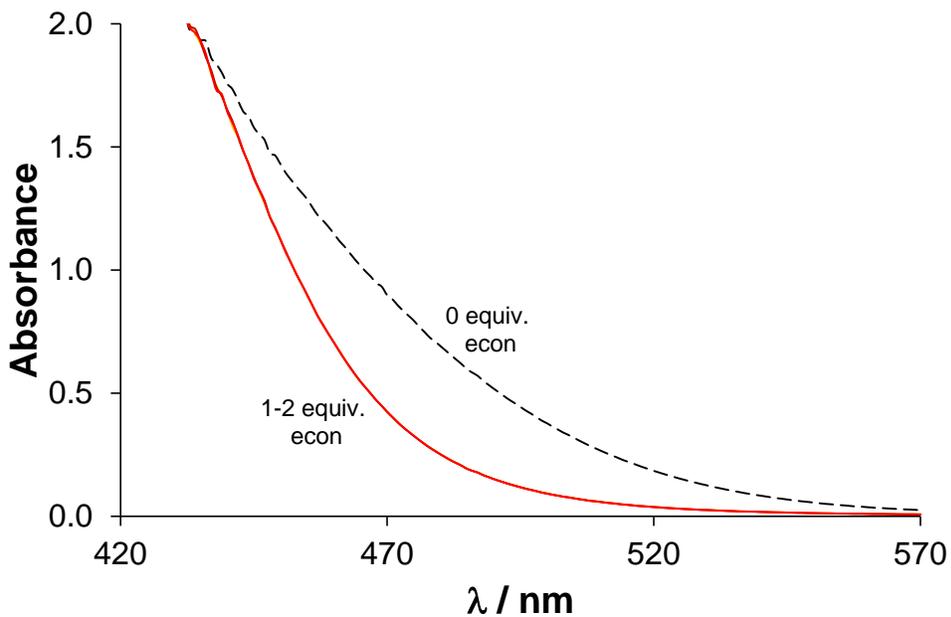


Figure S1. UV-vis spectra of the [RhCp*(8HQH₋₁)(H₂O)]⁺ complex at different concentrations of econazole. Conditions are the same as in the stock solutions prepared for anticancer and antibacterial experiments (pH = 7.4 (phosphate buffer) with 50% (v/v) ethanol). { c ([RhCp*(8HQH₋₁)(H₂O)]⁺) = 5 mM; c (econazole) = 0, 5 7.5 or 10 mM; t = 25 °C; ℓ = 0.2 cm}

CHARACTERIZATION OF THE ISOLATED COMPLEXES: ^1H NMR
AND ^{13}C NMR SPECTROSCOPIC DATA

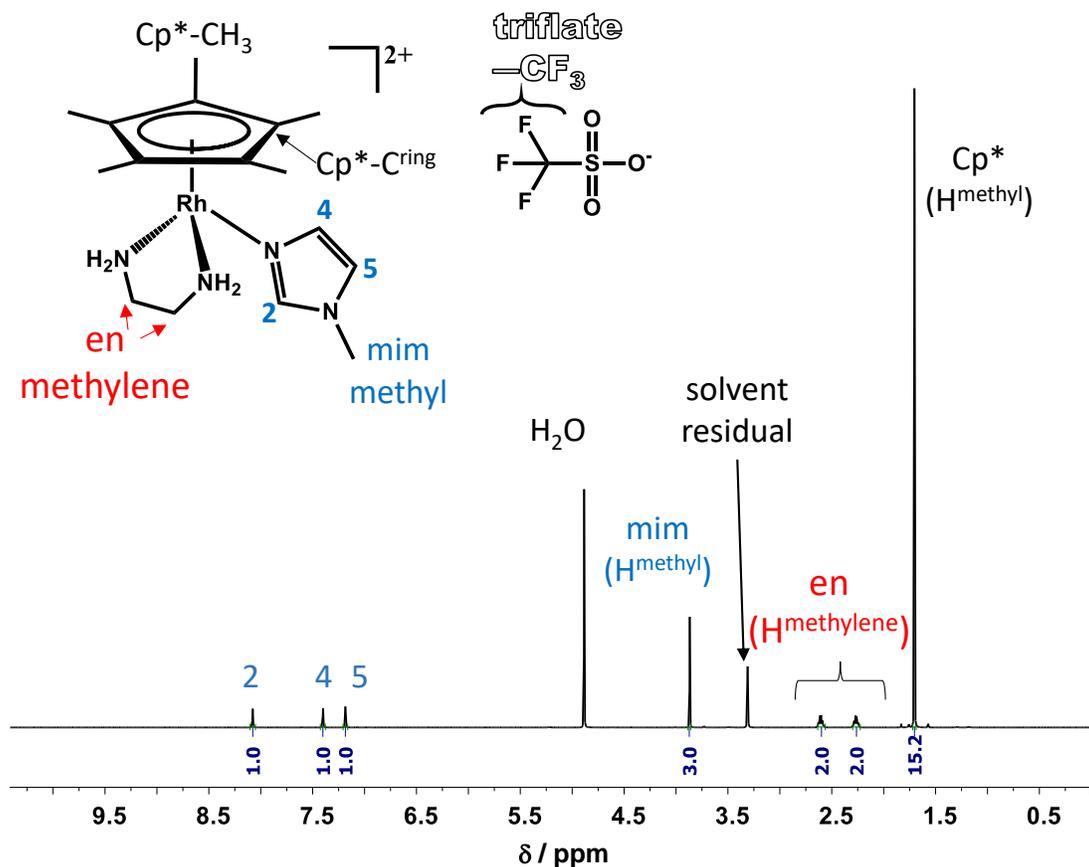


Figure S2. ^1H NMR spectrum of $[\text{RhCp}^*(\text{en})(\text{mim})](\text{CF}_3\text{SO}_3)_2$ in CD_3OD . Inserted structure shows numbering of peaks. $\{c(\text{complex}) = 10 \text{ mM}, t = 25.0^\circ\text{C}\}$

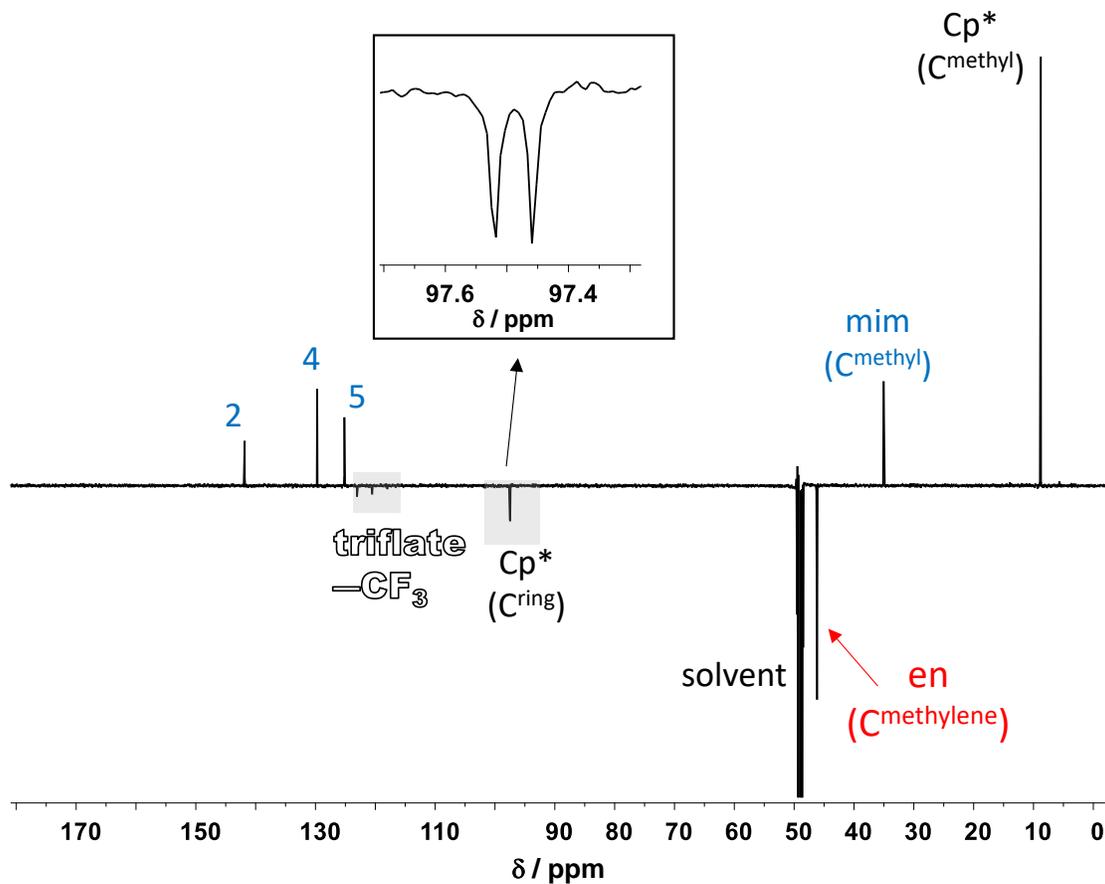


Figure S3. ^{13}C APT NMR spectrum of $[\text{RhCp}^*(\text{en})(\text{mim})](\text{CF}_3\text{SO}_3)_2$ in CD_3OD . For numbering see **Figure S2**. Inset figure shows the doublet of Cp^* ring carbon atoms as a result of coupling with ^{103}Rh . $\{c(\text{complex}) = 10 \text{ mM}, t = 25.0^\circ\text{C}\}$

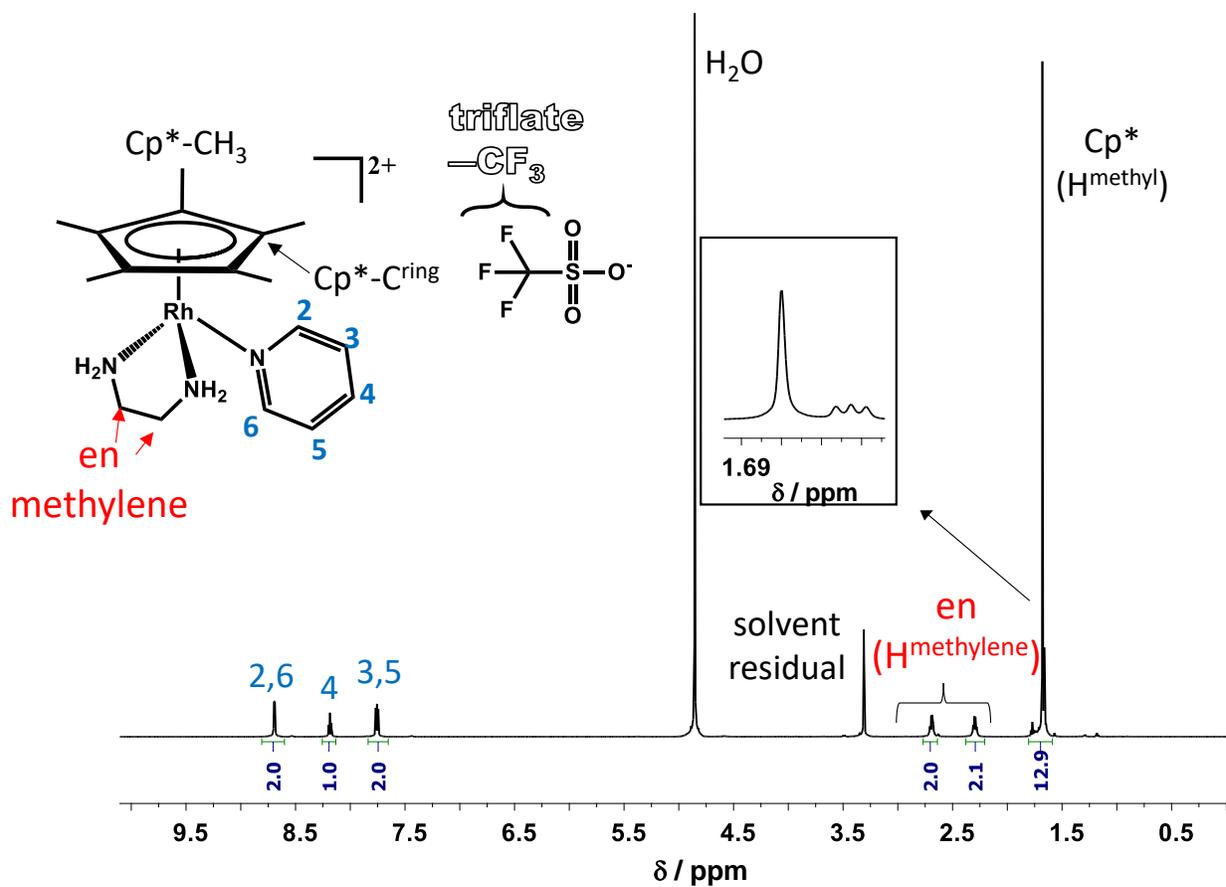


Figure S4. ^1H NMR spectrum of $[\text{RhCp}^*(\text{en})(\text{Py})](\text{CF}_3\text{SO}_3)_2$ in CD_3OD . Inserted structure shows numbering of peaks. Inset shows a triplet indicating the presence of $-\text{CH}_2\text{D}$ group. $\{c(\text{complex}) = 10 \text{ mM}, t = 25.0^\circ\text{C}\}$

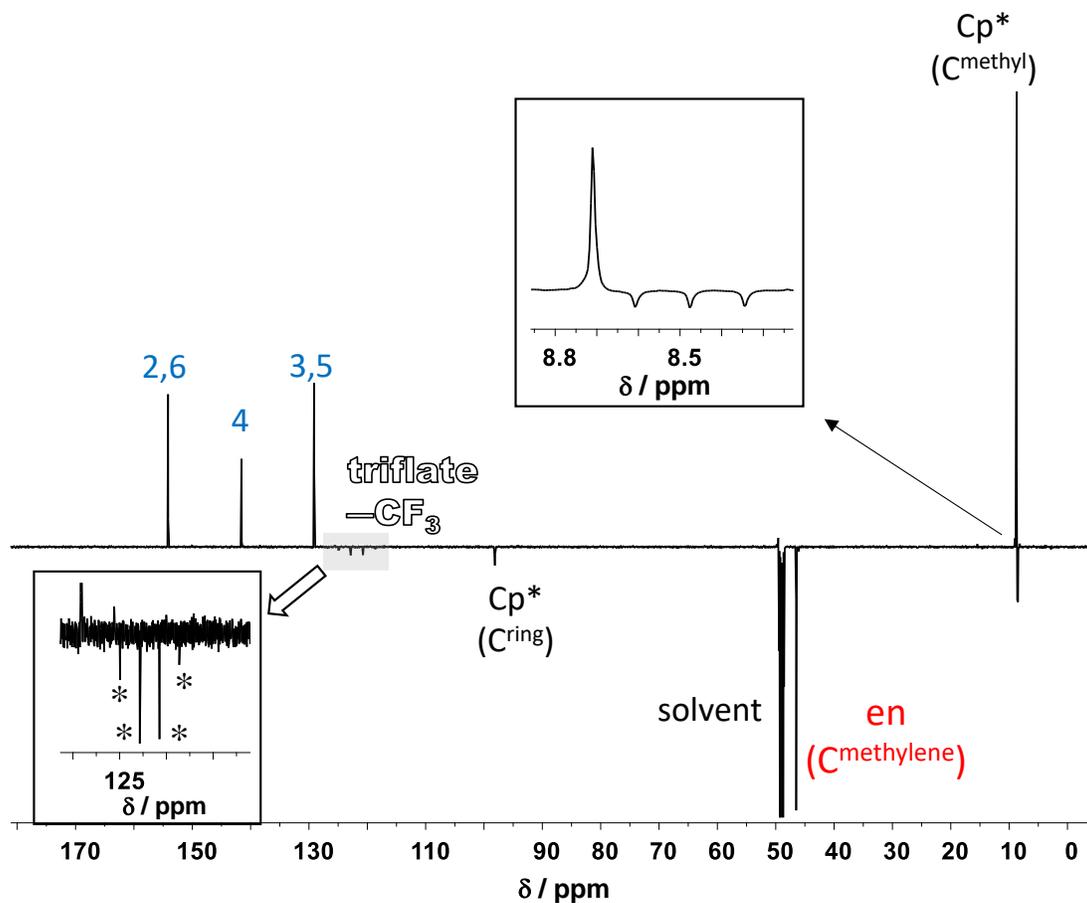


Figure S5. ^{13}C APT NMR spectrum of $[\text{RhCp}^*(\text{en})(\text{Py})](\text{CF}_3\text{SO}_3)_2$ in CD_3OD . For numbering see **Figure S4**. Inserted figure shows the quartet of $-\text{CF}_3$ carbon atom as a result of coupling with ^{19}F and a triplet indicating the presence of $-\text{CH}_2\text{D}$ group. $\{c(\text{complex}) = 10 \text{ mM}, t = 25.0^\circ\text{C}\}$

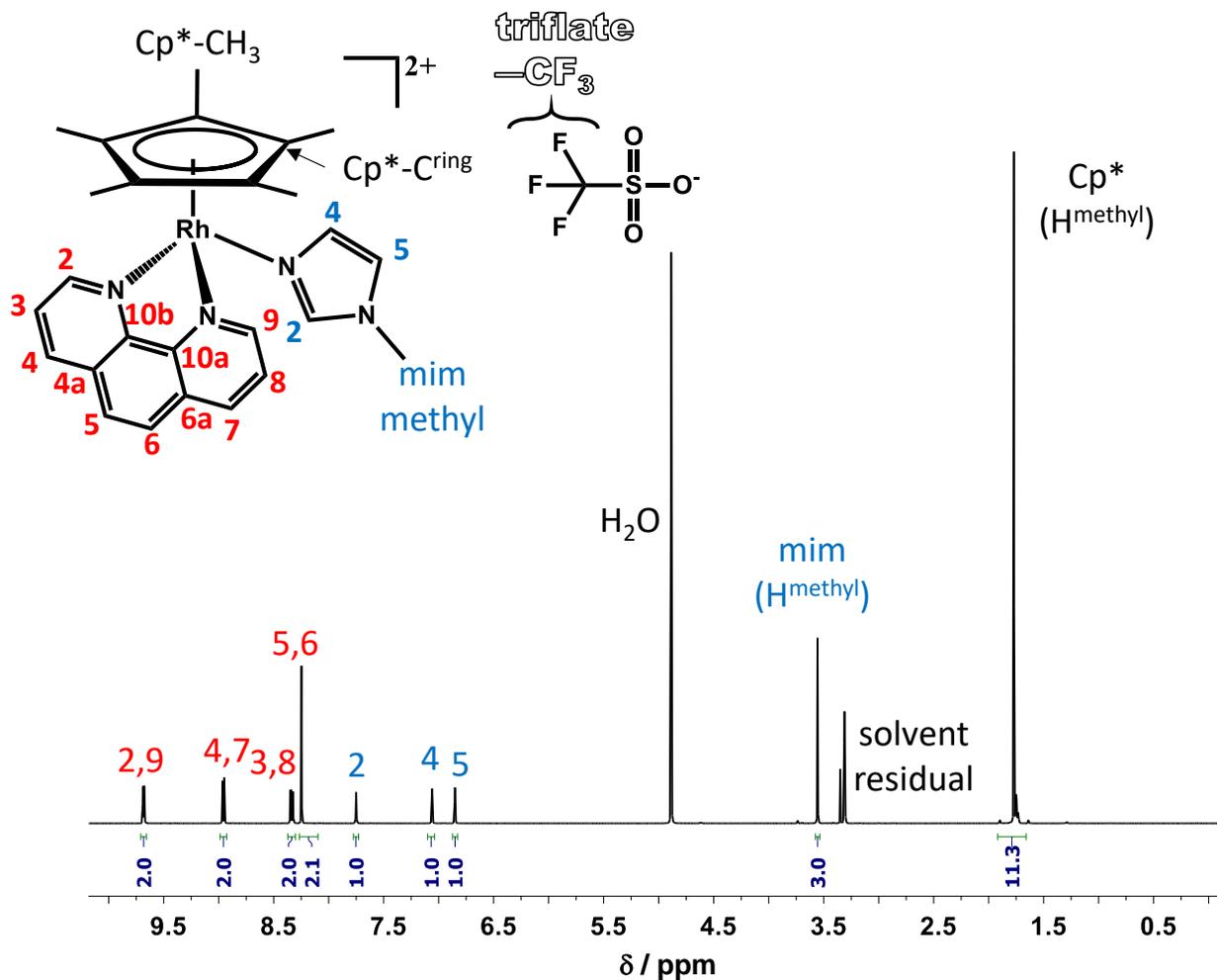


Figure S6. ^1H NMR spectrum of $[\text{RhCp}^*(\text{phen})(\text{mim})](\text{CF}_3\text{SO}_3)_2$ in CD_3OD . Inserted structure shows the numbering of peaks. $\{c(\text{complex}) = 10 \text{ mM}, t = 25.0^\circ\text{C}\}$

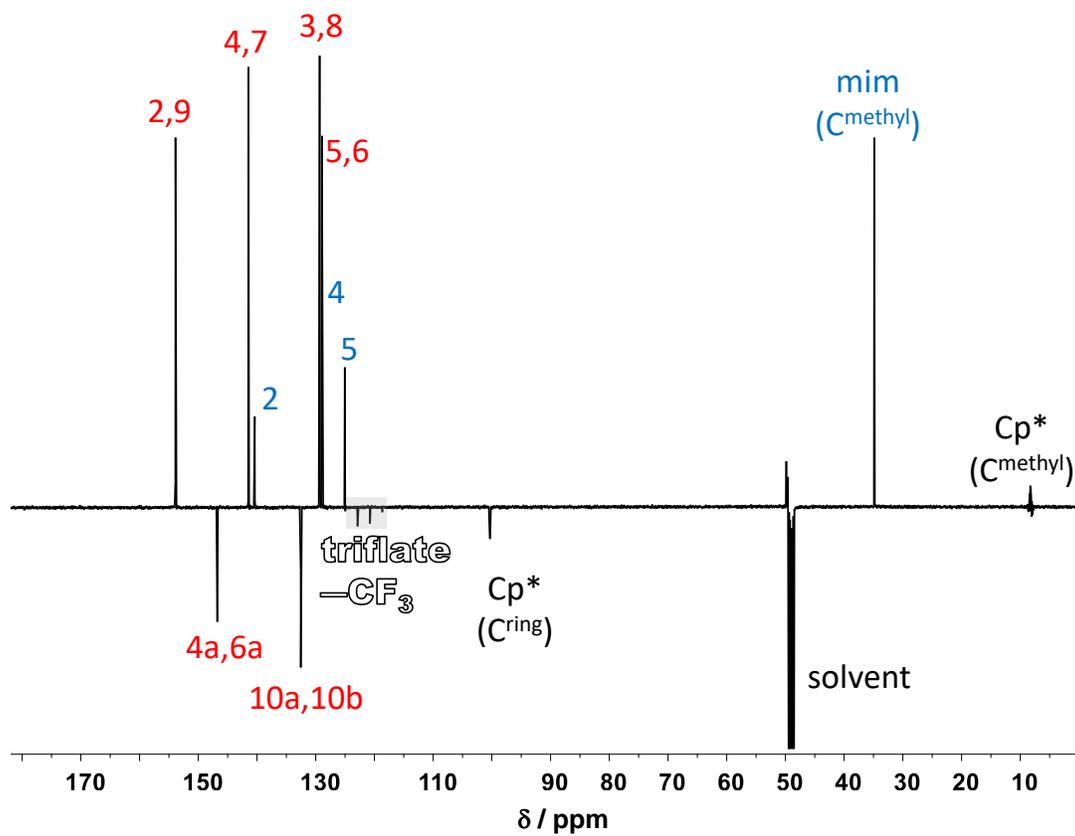


Figure S7. ^{13}C APT NMR spectrum of $[\text{RhCp}^*(\text{phen})(\text{mim})](\text{CF}_3\text{SO}_3)_2$ in CD_3OD . For numbering see **Figure S6**. $\{c(\text{complex}) = 10 \text{ mM}, t = 25.0^\circ\text{C}\}$

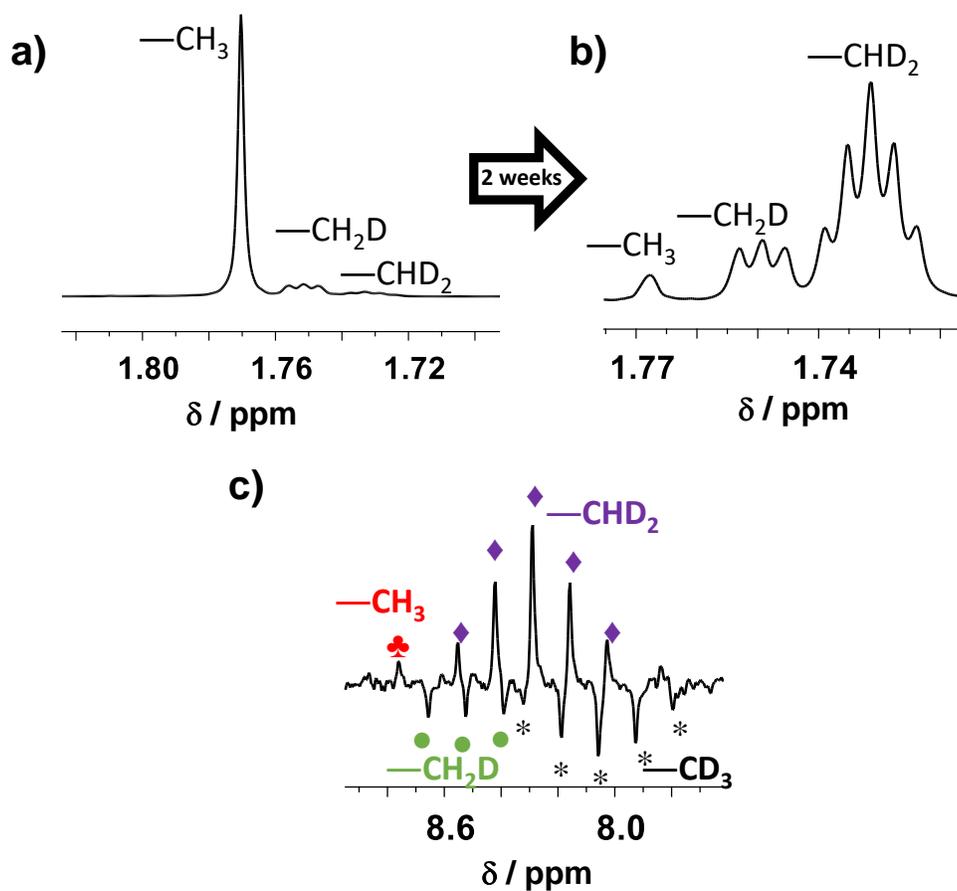


Figure S8. Upfield region of the ^1H NMR spectrum of $[\text{RhCp}^*(\text{phen})(\text{mim})](\text{CF}_3\text{SO}_3)_2$ in CD_3OD . a) Freshly prepared sample (waiting time < 1 day prior to measurement) and b) after ~2 weeks. c) Upfield region of the compound's ^{13}C NMR spectrum shows the presence of the Cp^* methyl groups at different deuterated stages measured 2-week-old sample. $\{c(\text{complex}) = 10 \text{ mM}, t = 25.0^\circ\text{C}\}$

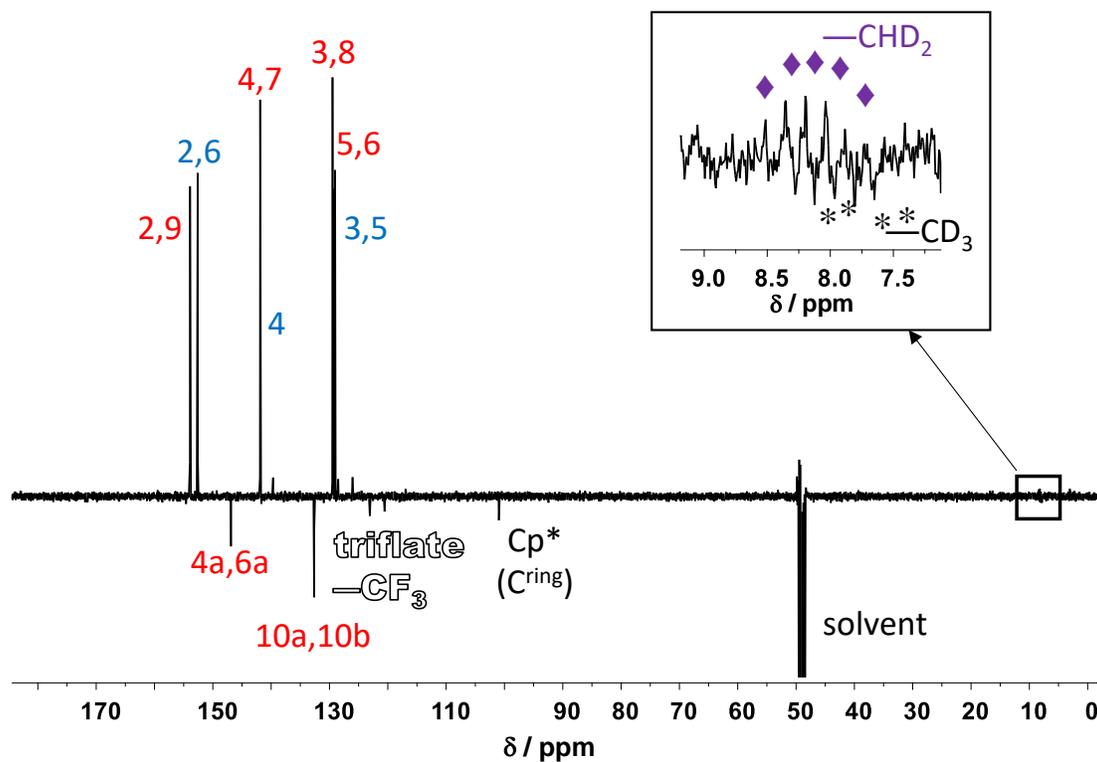


Figure S10. ^{13}C APT NMR spectrum of $[\text{RhCp}^*(\text{phen})(\text{Py})](\text{CF}_3\text{SO}_3)_2$ in CD_3OD . For numbering see **Figure S9**. Inset shows the presence of the Cp^* methyl groups at different deuterated stages. $\{c(\text{complex}) = 10 \text{ mM}, t = 25.0^\circ\text{C}\}$

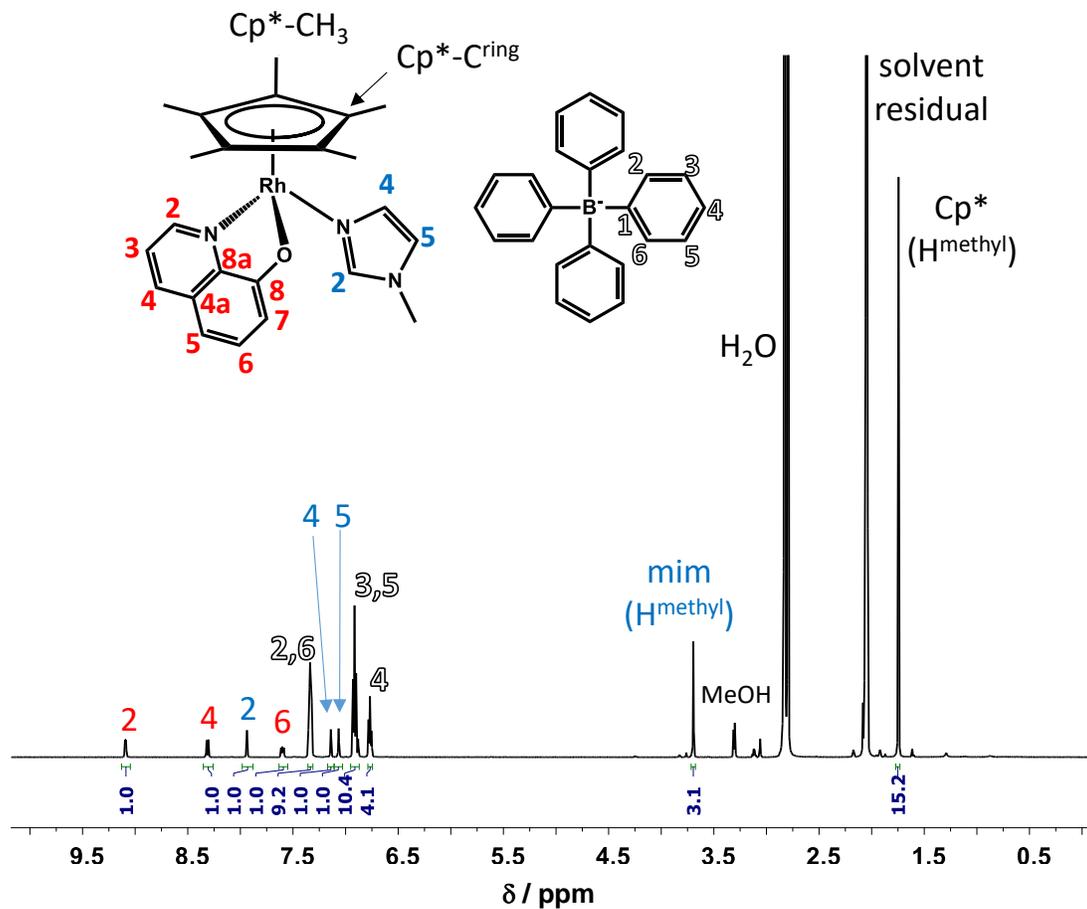


Figure S11. ^1H NMR spectrum of $[\text{RhCp}^*(8\text{HQH-1})(\text{mim})](\text{BPh}_4)$ in acetone- d_6 . Inserted structure shows numbering of peaks. $\{c(\text{complex}) = 2 \text{ mM}, t = 25.0^\circ\text{C}\}$

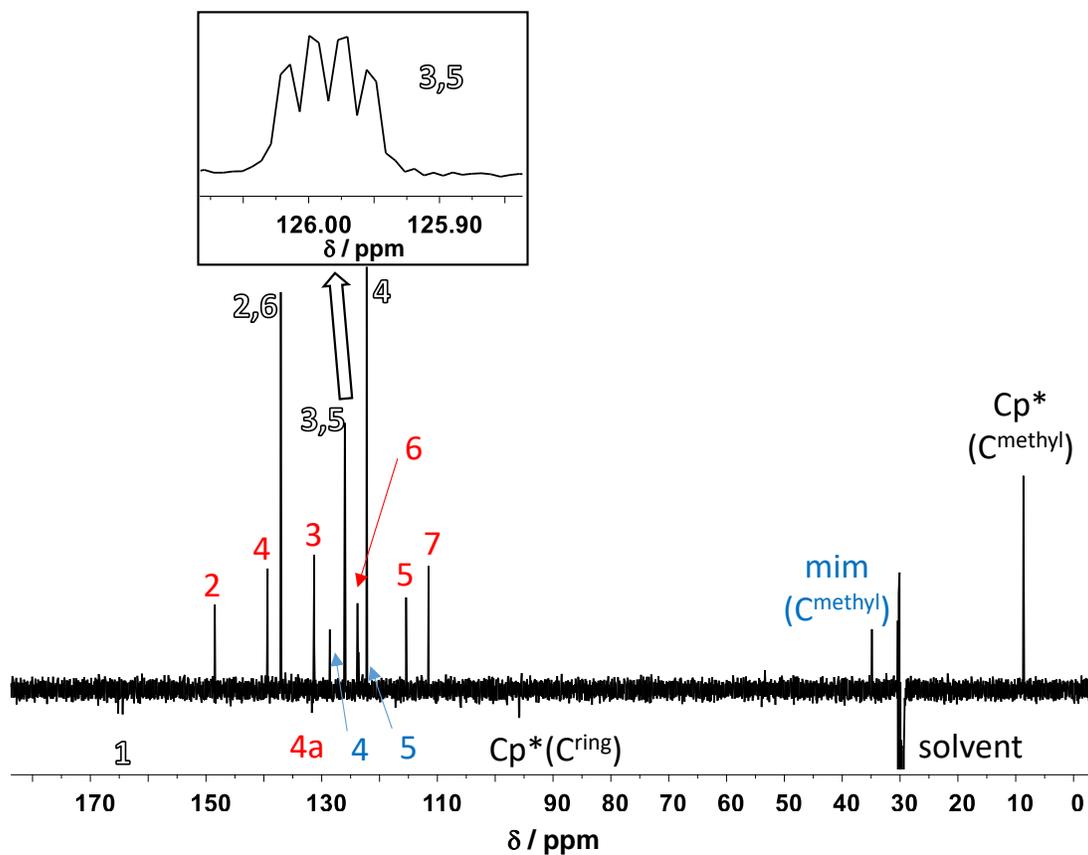


Figure S12. ^{13}C APT NMR spectrum of $[\text{RhCp}^*(8\text{HQH}_{-1})(\text{mim})](\text{BPh}_4)$ in acetone- d_6 . For numbering see **Figure S11**. Inset shows the coupling with ^{11}B in BPh_4 . $\{c(\text{complex}) = 2 \text{ mM}, t = 25.0^\circ\text{C}\}$

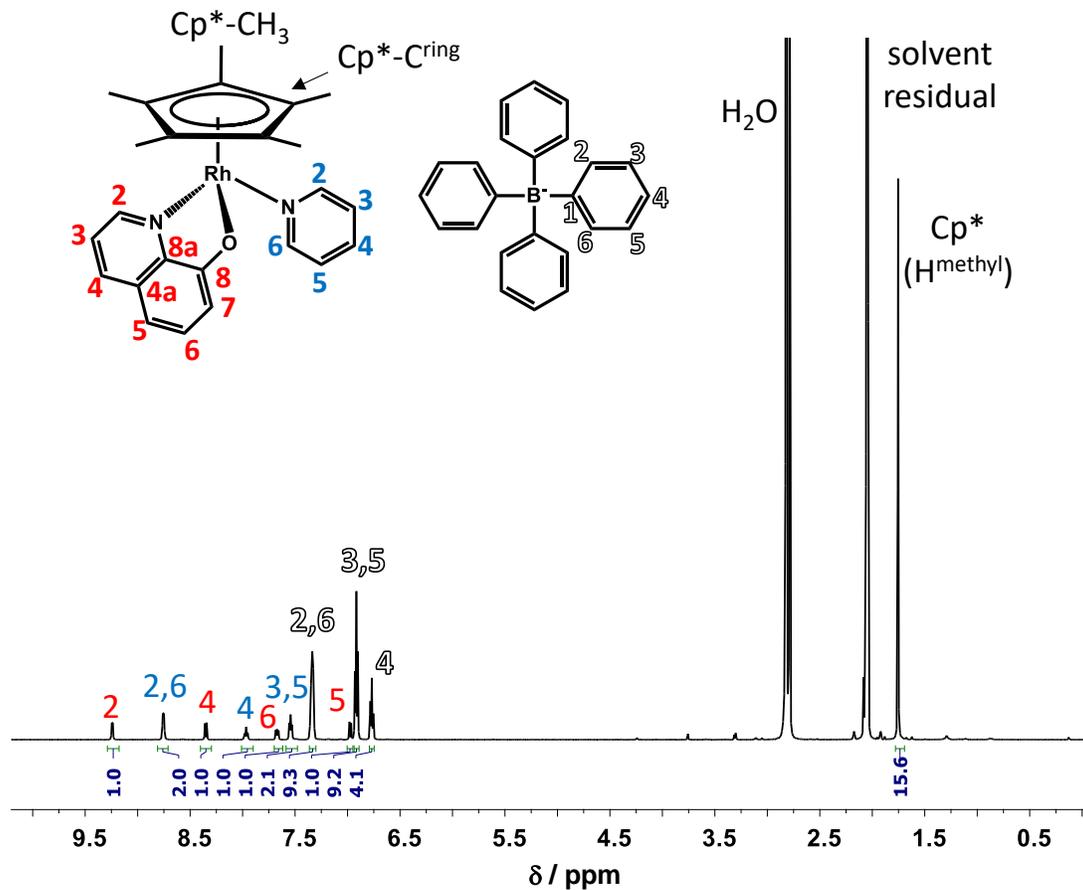


Figure S13. ^1H NMR spectrum of $[\text{RhCp}^*(8\text{HQH-1})(\text{Py})](\text{BPh}_4)$ in acetone- d_6 . Inserted structure shows numbering of peaks. $\{c(\text{complex}) = 2 \text{ mM}, t = 25.0^\circ\text{C}\}$

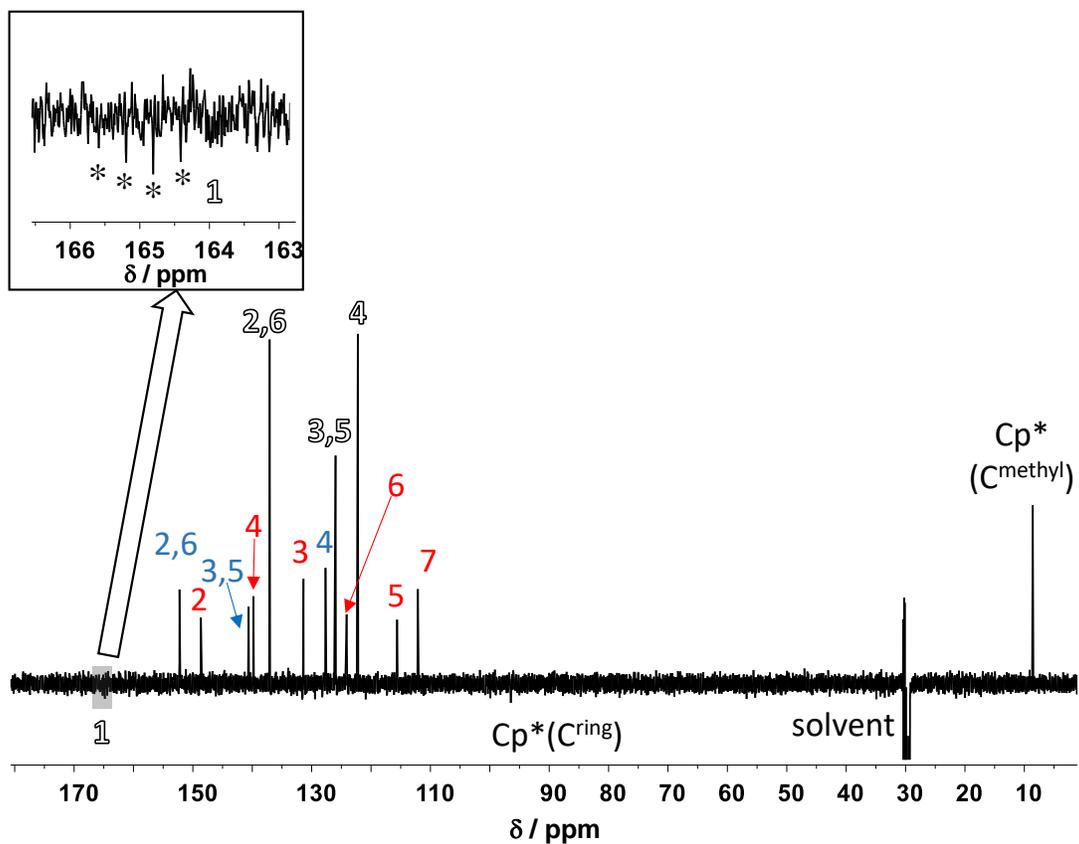


Figure S14. ^{13}C APT NMR spectrum of $[\text{RhCp}^*(8\text{HQH-1})(\text{Py})](\text{BPh}_4)$ in acetone- d_6 . For numbering see **Figure S13**. Inset shows the coupling with ^{11}B in BPh_4^- . $\{c(\text{complex}) = 2 \text{ mM}, t = 25.0^\circ\text{C}\}$

DATA QUALITY OF STRUCTURE DETERMINATION, COMPARISON OF M–N BOND LENGTHS

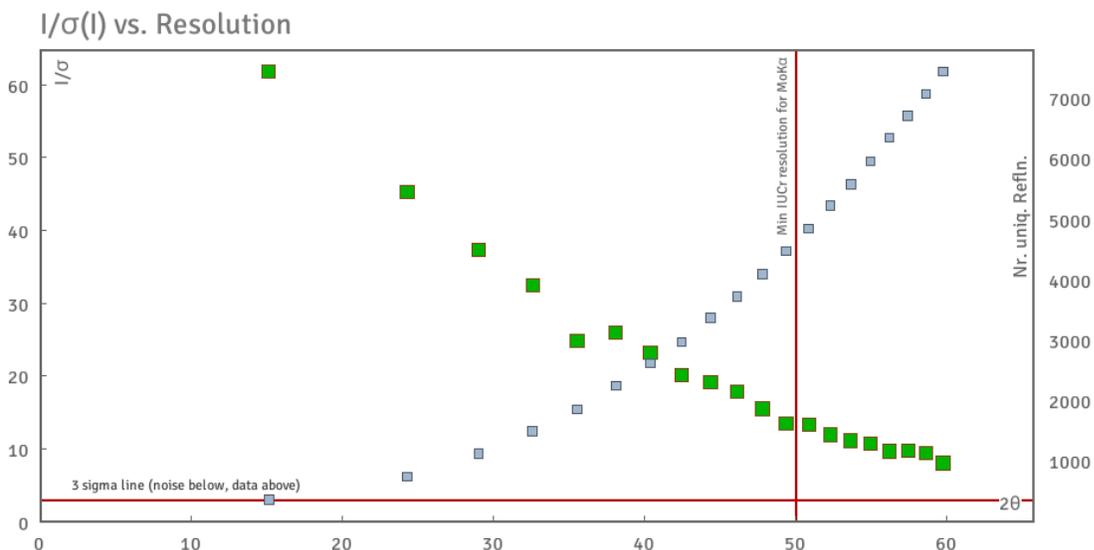


Figure S15. Data quality for complex 1': all data are above the three sigma line (along the minimum defined by the International Union of Crystallography (IUCR)).

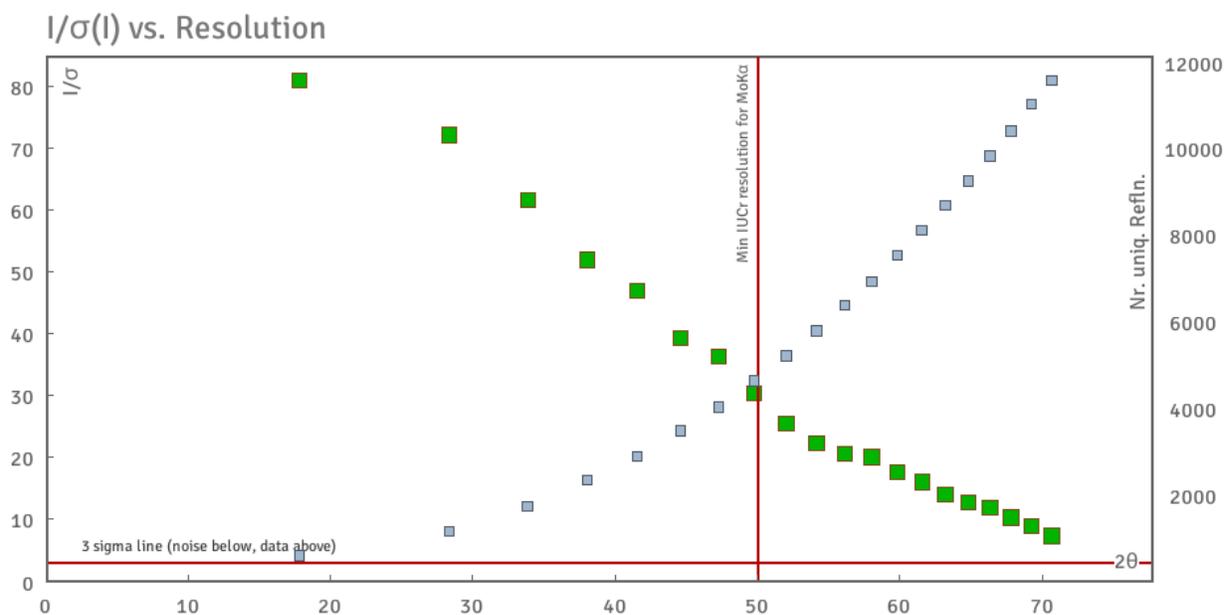


Figure S16. Data quality for complex 2': all data are above the three sigma line (along the minimum defined by IUCR).

I/σ(I) vs. Resolution

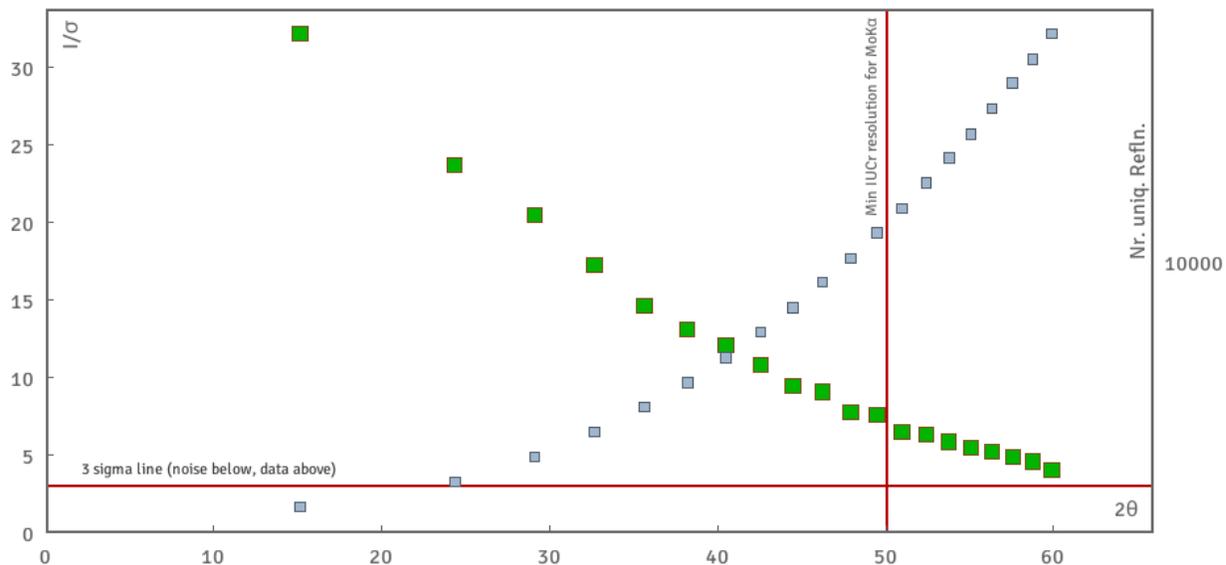


Figure S17. Data quality for complex 3': all data are above the three sigma line (along the minimum defined by IUCR).

I/σ(I) vs. Resolution

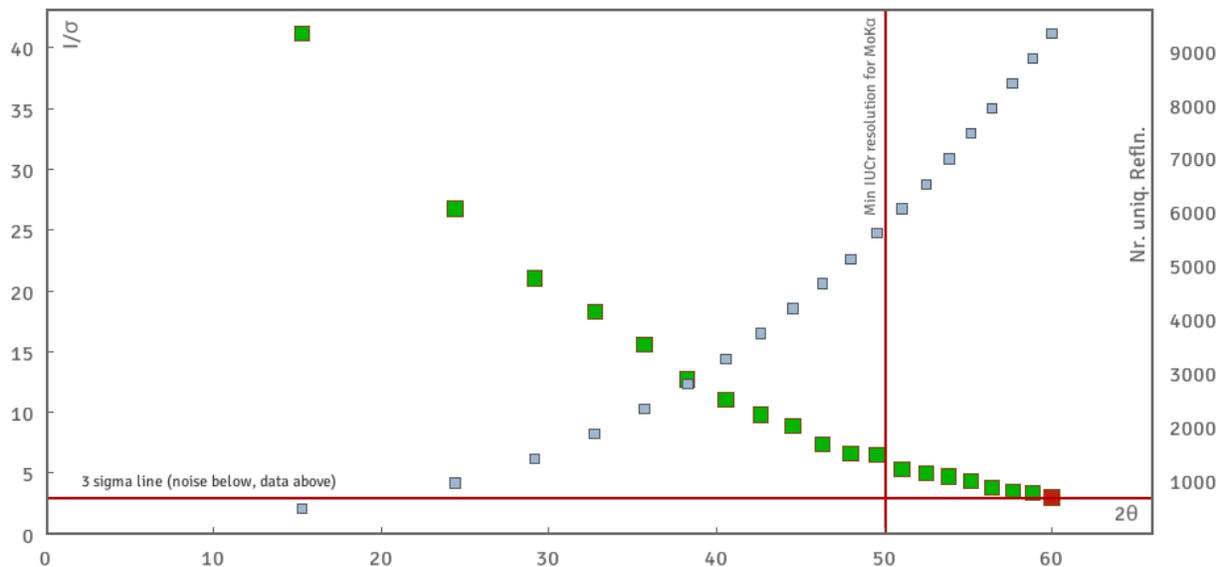


Figure S18. Data quality for complex 4': all data are above the three sigma line (along the minimum defined by IUCR).

$I/\sigma(I)$ vs. Resolution

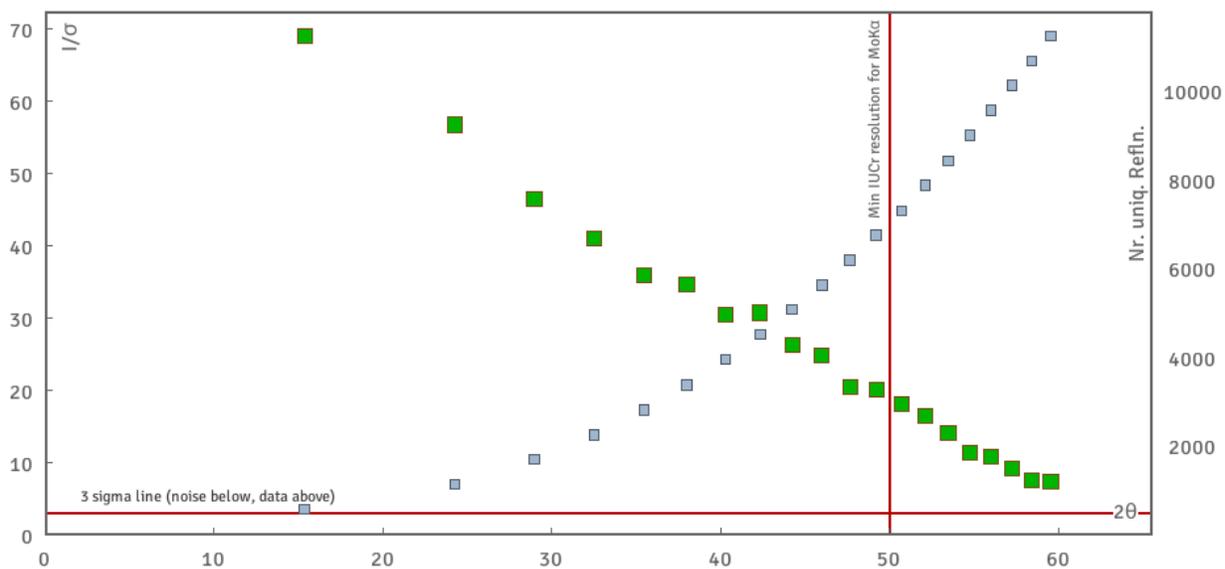


Figure S19. Data quality for complex 5': all data are above the three sigma line (along the minimum defined by IUCR).

$I/\sigma(I)$ vs. Resolution

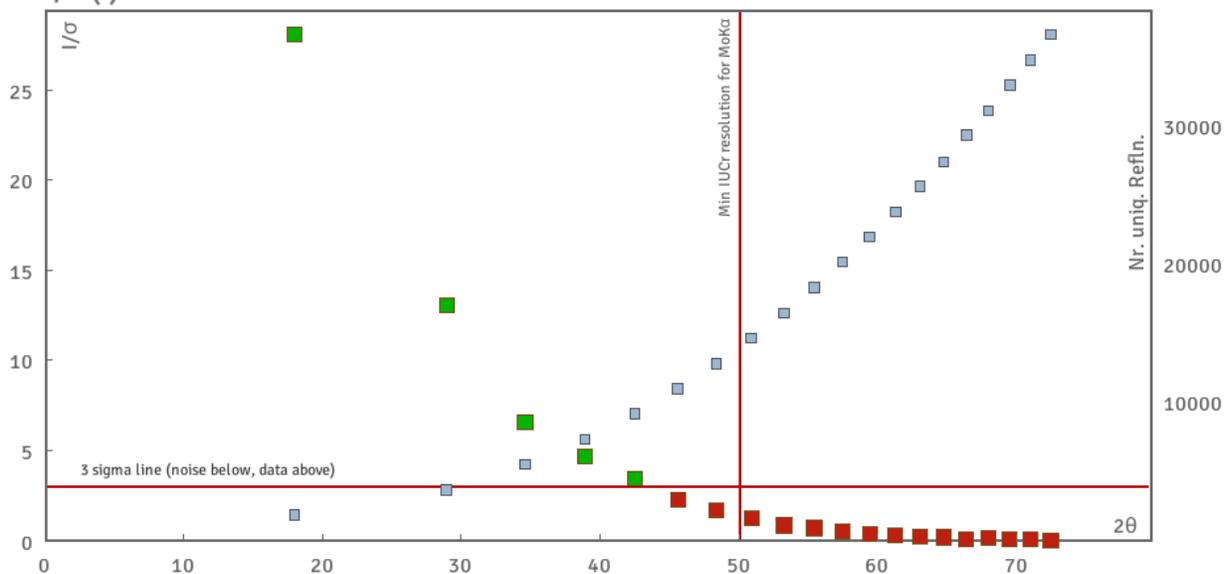


Figure S20. Data quality for complex 6': almost all data are above the three sigma line (along the minimum defined by IUCR). Data close to 50 degree (0.82, Mo) are still used in the refinement.

UV-Vis SPECTRA, FRACTIONS OF MIXED-LIGAND COMPLEXES AS THE FUNCTION OF pH, COMPARISON OF M-N BOND LENGTHS

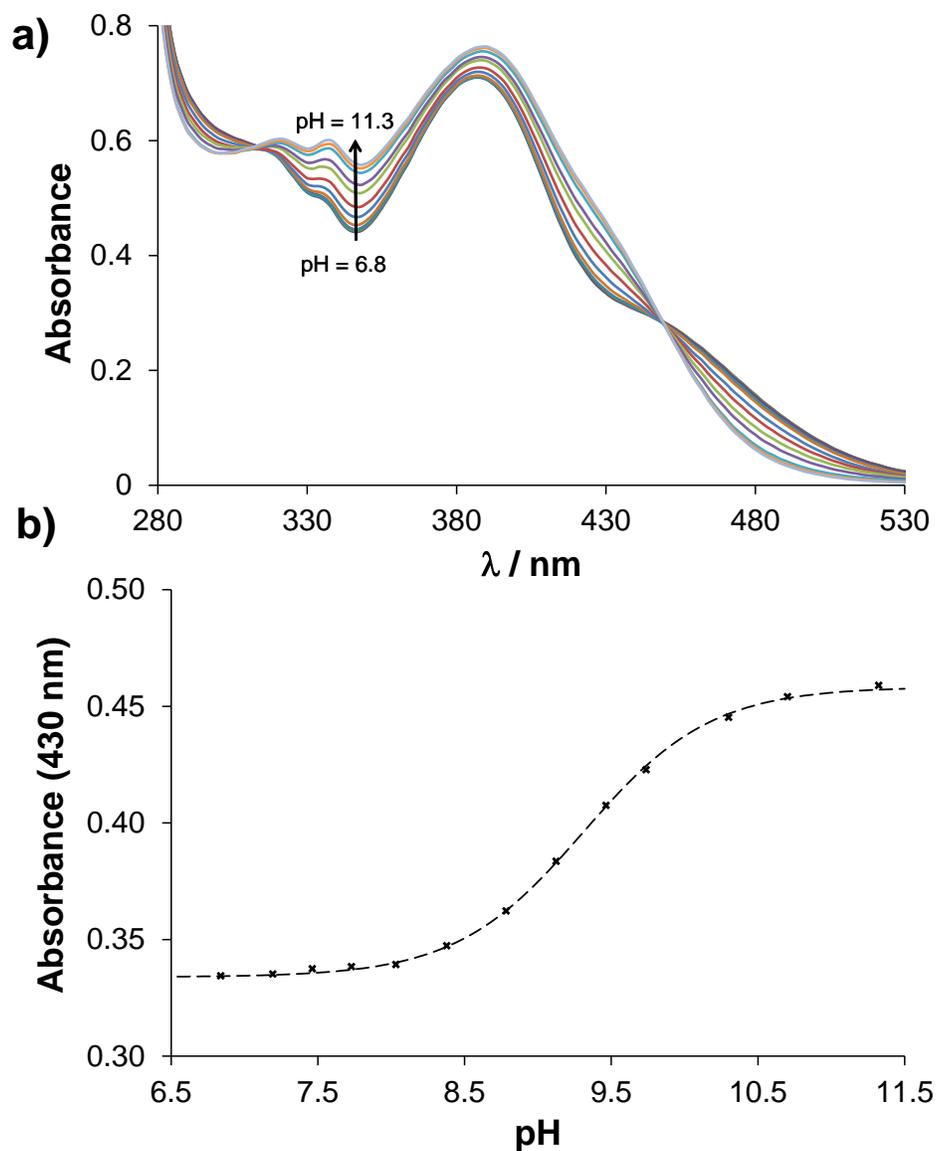


Figure S21. a) pH-dependent UV-vis spectra of [Ru(η^6 -*p*-cym)(8HQH₋₁)(H₂O)]⁺ in the pH-range 2.0–11.5. b) Absorbance values at 430 nm in the function of pH. {*c*([Ru(η^6 -*p*-cym)(8HQH₋₁)(H₂O)]⁺) = 200 μ M; *I* = 0.10 M (KCl), *t* = 25 $^{\circ}$ C, *l* = 1 cm}

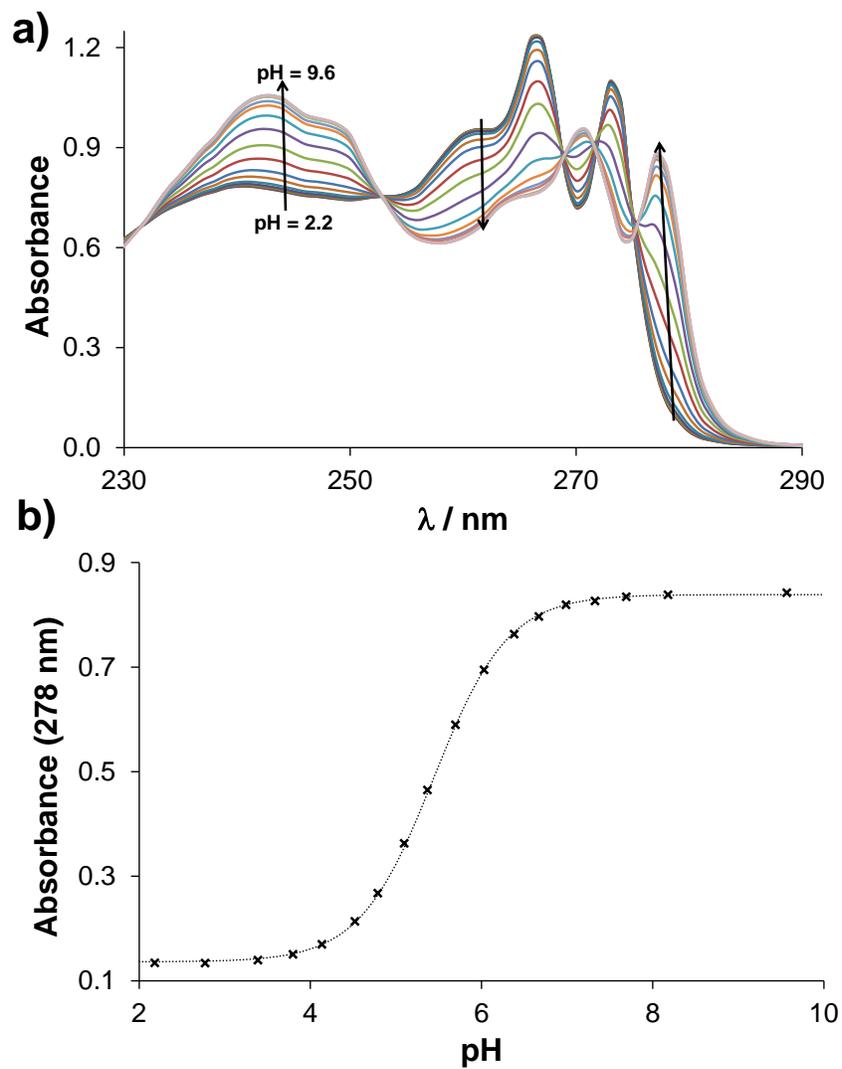


Figure S22. a) pH-dependent UV-vis spectra of benzimidazole in the pH-range 2.0–10.5. b) Absorbance values at 278 nm in the function of pH together with the fitted line (dotted line). $\{c(\text{bim}) = 200 \mu\text{M}; I = 0.10 \text{ M (KCl)}, t = 25 \text{ }^\circ\text{C}, \ell = 0.2 \text{ cm}\}$.

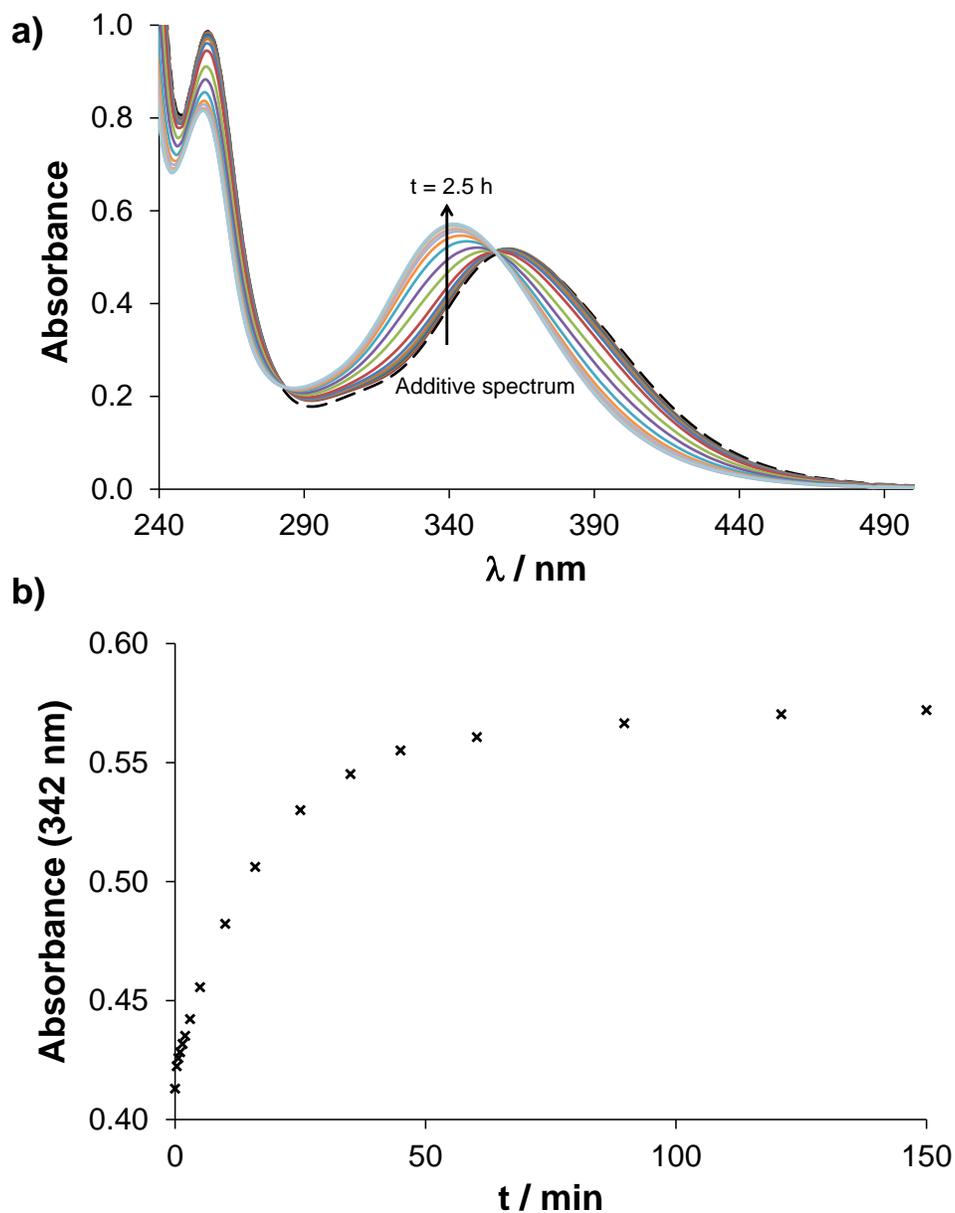


Figure S23. a) Time-dependent UV-vis spectra of the sample containing a mixture with $c([\text{RhCp}^*(\text{en})(\text{H}_2\text{O})]^{2+}):c(1\text{-methylimidazole}) = 1:2$ composition at $\text{pH} = 5.94$. b) Time dependence of absorbance values at 342 nm. $\{c([\text{RhCp}^*(\text{en})(\text{H}_2\text{O})]^{2+}) = 200 \mu\text{M}; c(1\text{-methylimidazole}) = 400 \mu\text{M}; \text{pH} = 5.94 (\text{PBS}'); t = 25 \text{ }^\circ\text{C}; \ell = 1 \text{ cm}\}$

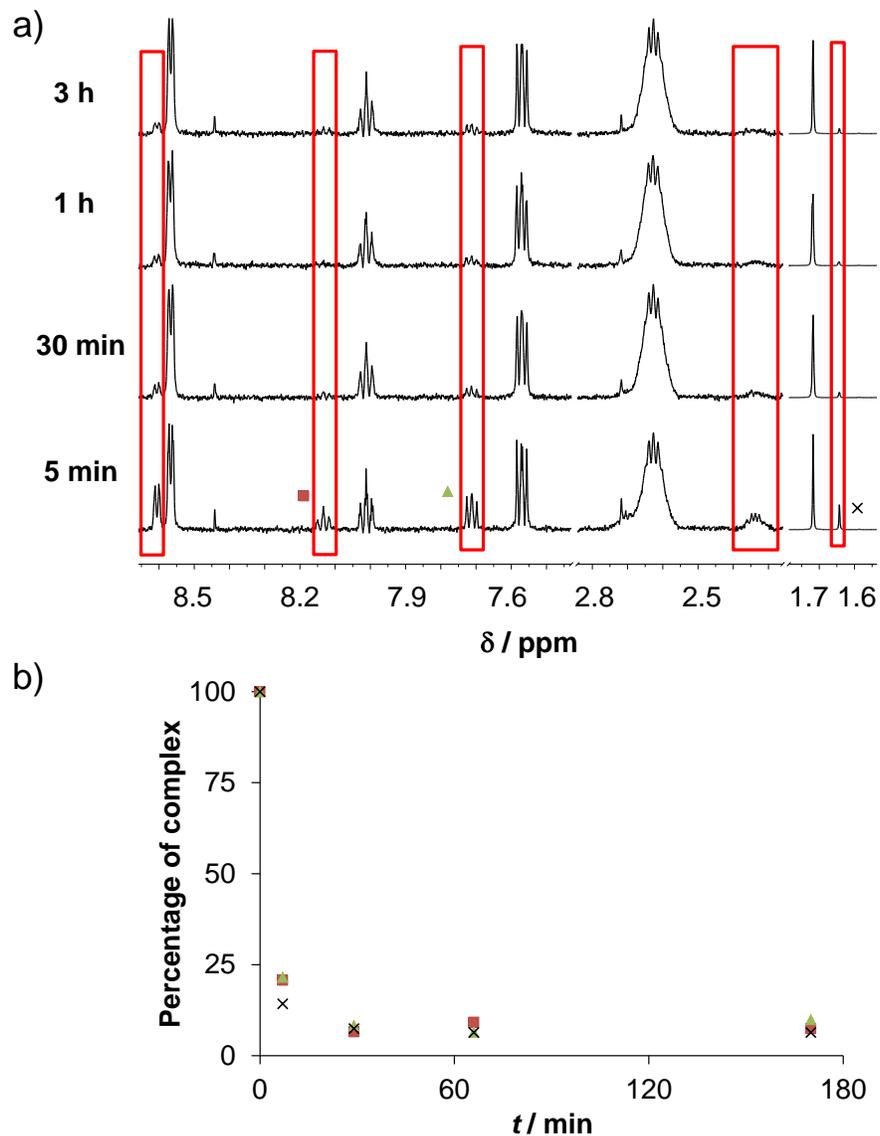


Figure S24. a) Time-dependent ^1H NMR spectra of the $[\text{RhCp}^*(\text{en})(\text{Py})](\text{CF}_3\text{SO}_3)_2$ synthesized complex at pH = 6.0. Red rectangles are highlighting peaks of the remaining mixed-ligand complex. b) Time-dependence of the presence of mixed-ligand complex (in percentage) based on integrals of the peaks of pyridine and Cp^* . Assignment: \times = Cp^* ; \blacksquare = $\text{pyr-H}^{3,5}$; \blacktriangle = pyr-H^4 . $\{c([\text{RhCp}^*(\text{en})(\text{Py})](\text{CF}_3\text{SO}_3)_2) = 200 \mu\text{M}; \text{pH} = 6.0 (\text{PBS}'); 10\% \text{D}_2\text{O}; t = 25 \text{ }^\circ\text{C}; \}$

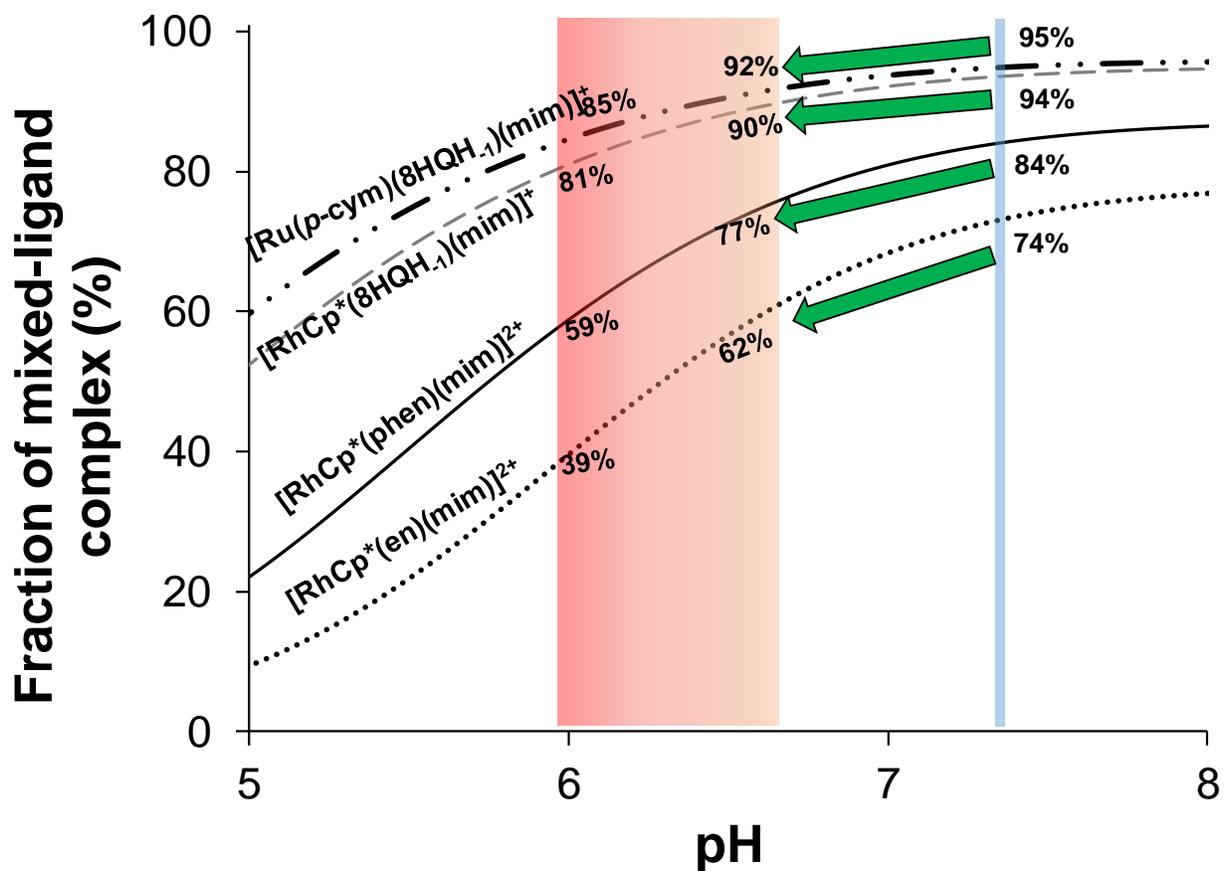


Figure S25. Fractions of the formed mixed-ligand complexes of 1-methylimidazole in the pH-range 5.0–8.0. Calculations based on the stability constants of the mixed-ligand complexes (Table S2) and proton dissociation constants in Table 1. $\{c([\text{M}(\text{arene})(\text{N},\text{N}/\text{O})(\text{H}_2\text{O})) = c(\text{mim}) = 100 \mu\text{M}; I = 0.10 \text{ M (KCl)}, t = 25 \text{ }^\circ\text{C}\}$.

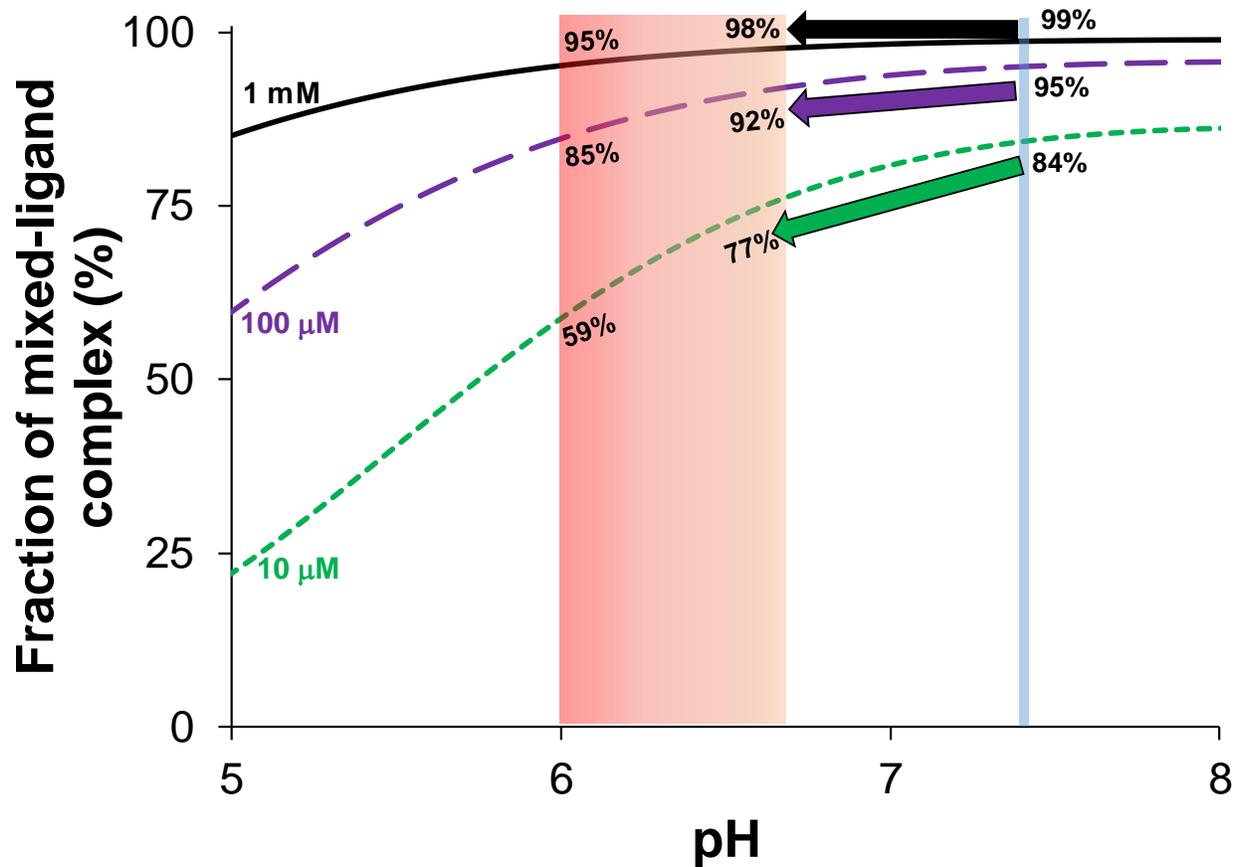


Figure S26. Formation (%) of $[\text{Ru}(\eta^6\text{-}p\text{-cym})(8\text{HQH-1})(\text{mim})]^+$ in the pH-range 5.0–8.0 at various equimolar concentrations of monodentate ligand and aqua complex. $\{c([\text{Ru}(\eta^6\text{-}p\text{-cym})(8\text{HQH-1})(\text{H}_2\text{O})]^+) = c(\text{mim}) = 10 \mu\text{M}, 100 \mu\text{M} \text{ or } 1 \text{ mM}; I = 0.10 \text{ M (KCl)}, t = 25 \text{ }^\circ\text{C}\}$.

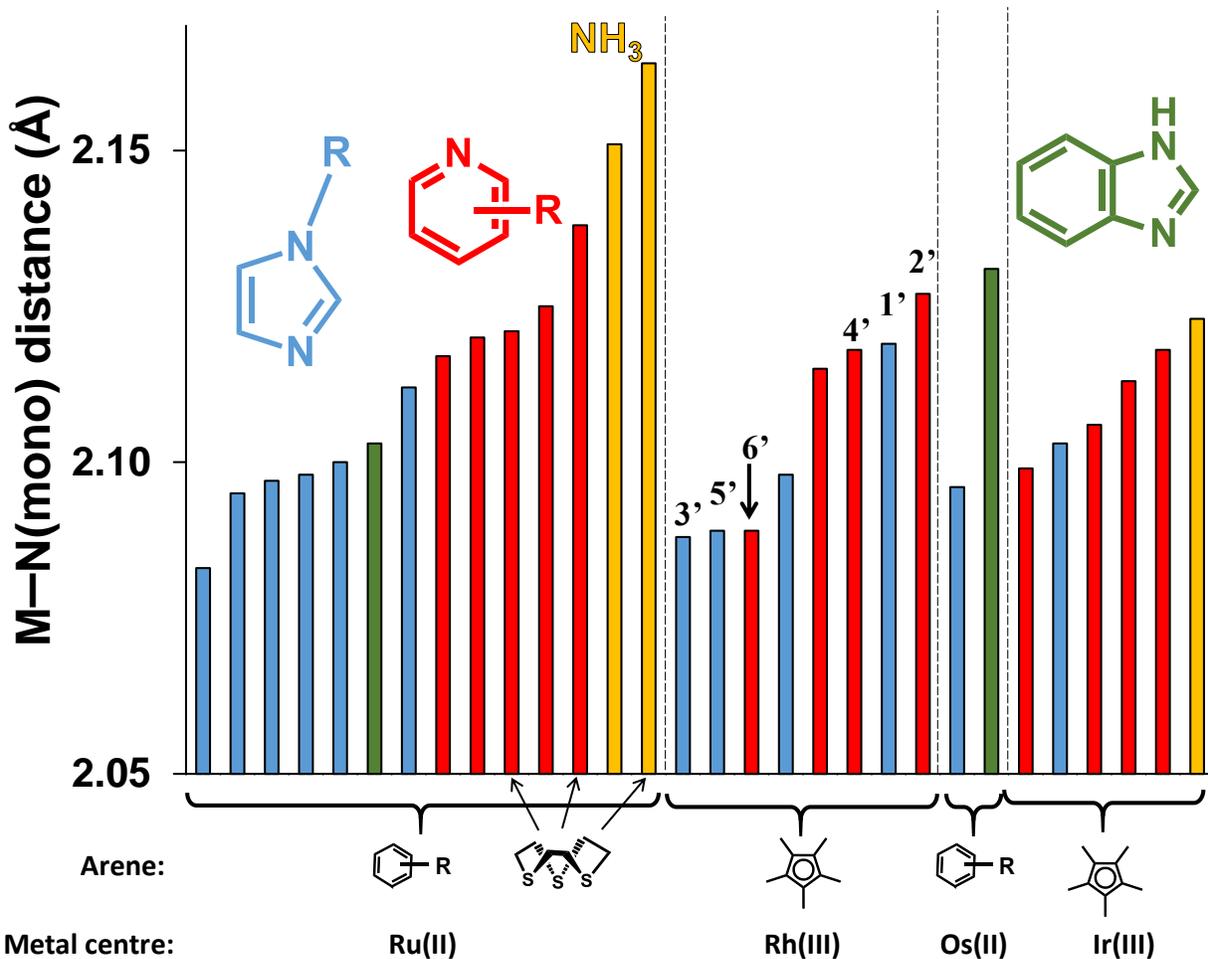


Figure S27. Comparison of the M–N(monodentate) distances in different [M(arene)(N,N/O)(N)]-type complexes reported in references [1-12]. Colour code for the monodentate ligand: **blue**: imidazole derivatives; **red**: pyridine derivatives; **green**: benzimidazole; **yellow**: NH₃. Roman numerals indicate the values derived from complexes 1'–6'.

TABLES: PROTON DISSOCIATION CONSTANTS AND FORMATION CONSTANTS

Table S1. Proton dissociation constants of the coordinated water molecules in $[M(\text{arene})(\text{N},\text{N}/\text{O})(\text{H}_2\text{O})]^{2+/+}$ complexes $\{I = 0.10 \text{ M (KCl)}; t = 25.0 \text{ }^\circ\text{C}\}$.

Compound	pK _a	Method	Literature data
[RhCp*(en)(H ₂ O)] ²⁺	10.6±0.1	UV-vis	9.58 ¹ [13] 11.05 ² [13]
[RhCp*(8HQH ₋₁)(H ₂ O)] ⁺	10.6±0.1	UV-vis	10.27 ² [14]
[RhCp*(phen)(H ₂ O)] ²⁺	10.05±0.01	UV-vis	8.58 ² [15]
[Ru(<i>p</i> -cym)(en)(H ₂ O)] ²⁺	9.03±0.01	UV-vis	8.14 ² [15]
[Ru(<i>p</i> -cym)(8HQH ₋₁)(H ₂ O)] ⁺	9.31±0.01	UV-vis	9.19 ² [14]
[Ru(<i>p</i> -cym)(phen)(H ₂ O)] ²⁺	8.91±0.01	UV-vis	7.59 ² [15]

¹ *I* = 0.20 M (KNO₃); ² *I* = 0.20 M (KCl)

Table S2. Formation constants (logK) determined for mixed-ligand half-sandwich complexes of Rh(III) and Ru(II) by UV-vis and ¹H NMR spectroscopy $\{I = 0.10 \text{ M (KCl)}; t = 25.0 \text{ }^\circ\text{C}\}$.

$[M(\text{arene})(\text{N},\text{N}/\text{O})(\text{H}_2\text{O})]^{2+/+} + \text{N}_{\text{mono}} \rightleftharpoons [M(\text{arene})(\text{N},\text{N}/\text{O})(\text{N}_{\text{mono}})]^{2+/+} + \text{H}_2\text{O}$		
Aqua complex	Monodentate ligand	logK
[RhCp*(en)(H ₂ O)] ²⁺	bim	3.3 ± 0.2
[RhCp*(en)(H ₂ O)] ²⁺	mim	4.2 ± 0.1
[RhCp*(en)(H ₂ O)] ²⁺	Py	2.6 ± 0.1
[RhCp*(phen)(H ₂ O)] ²⁺	bim	4.9 ± 0.2
[RhCp*(phen)(H ₂ O)] ²⁺	mim	5.5 ± 0.1
[RhCp*(phen)(H ₂ O)] ²⁺	Py	4.1 ± 0.1
[RhCp*(8HQH ₋₁)(H ₂ O)] ⁺	bim	3.4 ± 0.1
[RhCp*(8HQH ₋₁)(H ₂ O)] ⁺	mim	4.7 ± 0.2
[RhCp*(8HQH ₋₁)(H ₂ O)] ⁺	Py	3.0 ± 0.1
[RuCym(en)(H ₂ O)] ²⁺	bim	3.5 ± 0.1
[RuCym(en)(H ₂ O)] ²⁺	mim	< 2.6
[RuCym(en)(H ₂ O)] ²⁺	Py	< 1.5
[RuCym(phen)(H ₂ O)] ²⁺	bim	5.6 ± 0.2
[RuCym(phen)(H ₂ O)] ²⁺	mim	6.7 ± 0.2
[RuCym(phen)(H ₂ O)] ²⁺	Py	4.3 ± 0.2
[RuCym(8HQH ₋₁)(H ₂ O)] ⁺	bim	2.9 ± 0.1
[RuCym(8HQH ₋₁)(H ₂ O)] ⁺	mim	< 2.6
[RuCym(8HQH ₋₁)(H ₂ O)] ⁺	Py	1.6 ± 0.2

EXPERIMENTAL DATA TABLES FOR SINGLE-CRYSTAL XRD

Table S3. Experimental parameters for the single crystal X-ray diffraction data collection and CCDC-code of the synthesized complexes.

CCDC-code	Sample	Machine	Source	Temp.	Detector Distance	Time/Frame	Frames	Frame width
				[K]	[mm]	[s]		[°]
2209815	6'	Stoe Stadivari	Mo	100	40	10	1804	0.5
2209816	4'	Bruker X8	Mo	100	36	6	1347	0.5
2209817	2'	Bruker D8	Mo	100	34	6	1805	0.5
2209818	3'	Bruker X8	Mo	100	40	8.1	5595	0.5
2209819	1'	Bruker X8	Mo	100	35	7	833	0.6
2209820	5'	Bruker D8	Mo	100	35	3	1390	0.6

Table S4. Crystal data and structure refinement for complexes **1'-3'**.

	1'	2'	3'
Empirical formula	C ₁₈ H ₂₉ F ₆ N ₄ O ₆ RhS ₂	C ₁₉ H ₂₈ F ₆ N ₃ O ₆ RhS ₂	C ₇₆ H ₇₃ B ₂ Cl ₄ N ₄ Rh
Formula weight	678.48	675.47	1308.71
Temperature (K)	100.0	100.0	100.0
Crystal system, Space group	triclinic	triclinic	triclinic
Space group	P-1	P-1	P-1
Unit cell dimensions			
a (Å)	7.7751(4)	7.8309(4)	12.9421(9)
b (Å)	11.5350(7)	10.8399(5)	12.9452(9)
c (Å)	15.4437(9)	15.7969(7)	20.0369(13)
α (°)	100.4691(17)	89.5353(16)	76.964(3)
β (°)	91.3163(17)	85.0745(16)	85.322(3)
γ (°)	103.4897(17)	79.0821(16)	86.772(4)
Volume (Å ³)	1321.35(13)	1311.76(11)	3255.8(4)
Z	2	2	2
ρ _{calc} (g/cm ³)	1.705	1.710	1.335
μ (mm ⁻¹)	0.887	0.892	0.474
F(000)	688.0	684.0	1360.0
Crystal size (mm)	0.13×0.1×0.07	0.423×0.227×0.08	0.3×0.11×0.08
Radiation	Mo-K _α (λ=0.71073 Å)		
2θ range for data collection (°)	5.378-54.206	4.602-71.378	4.432-50.7
Index ranges	-9≤h≤9, -14≤k≤14, -19≤l≤19	-12≤h≤12, -17≤k≤17, -25≤l≤23	-15≤h≤15, -15≤k≤15, -23≤l≤24
Reflections collected	14736	58577	97481
Independent reflections	5816 [R _{int} =0.0269, R _{sigma} =0.0308]	12115 [R _{int} =0.0303, R _{sigma} =0.0271]	11943 [R _{int} =0.0804, R _{sigma} =0.0482]
Data / restraints / parameters	5816 / 36 / 364	12115 / 0 / 339	11943 / 0 / 790
Goodness-of-fit on F ²	1.030	1.073	1.089
Final R indices [I>2σ(I)]	R ₁ =0.0331, wR ₂ =0.0788	R ₁ =0.0265, wR ₂ =0.0571	R ₁ =0.0452, wR ₂ =0.1179
R indices (all data)	R ₁ =0.0361, wR ₂ =0.0809	R ₁ =0.0328, wR ₂ =0.0592	R ₁ =0.0501, wR ₂ =0.1220
Largest diff. peak and hole (e ⁻ Å ⁻³)	1.64 / -1.04	0.95 / -1.08	1.50 / -1.67

Table S5. Crystal data and structure refinement for complexes **4'**-**6'**.

	4'	5'	6'
Empirical formula	C ₃₀ H ₃₂ F ₆ N ₃ O ₇ RhS ₂	C ₄₈ H ₅₁ BN ₃ O ₂ Rh	C ₄₈ H ₄₆ BN ₂ ORh
Formula weight	827.61	815.64	780.59
Temperature (K)	100.0	100.0	100.0
Crystal system, Space group	monoclinic	monoclinic	monoclinic
Space group	P2 ₁ /c	P2 ₁ /n	P2 ₁ /c
Unit cell dimensions			
a (Å)	13.0591(11)	11.8020(5)	29.1880(10)
b (Å)	13.5115(9)	27.8595(9)	11.8065(3)
c (Å)	18.9926(15)	12.2552(9)	24.8781(10)
α (°)	90	90	90
β (°)	102.252(3)	90.8912(11)	114.956(3)
γ (°)	90	90	90
Volume (Å ³)	3274.9(4)	4029.0(4)	7772.7(5)
Z	4	4	8
ρ _{calc} (g/cm ³)	1.679	1.345	1.334
μ (mm ⁻¹)	0.735	0.467	0.478
F(000)	1680.0	1704.0	3248.0
Crystal size (mm)	0.22×0.2×0.03	0.3×0.07×0.07	0.28×0.2×0.06
Radiation	Mo-Kα (λ=0.71073 Å)		
2Θ range for data collection (°)	3.728-50.69	4.524-60.092	3.274-51.364
Index ranges	-14≤h≤15, -15≤k≤16, -22≤l≤22	-16≤h≤16, -39≤k≤37, -17≤l≤17	-35≤h≤35, -14≤k≤14, -30≤l≤25
Reflections collected	28186	144915	83725
Independent reflections	5887 [R _{int} =0.0506, R _{sigma} =0.0447]	11780 [R _{int} =0.0628, R _{sigma} =0.0249]	14760 [R _{int} =0.1697, R _{sigma} =0.1700]
Data / restraints / parameters	5887 / 4 / 469	11780 / 0 / 504	14760 / 0 / 966
Goodness-of-fit on F ²	1.046	1.074	0.953
Final R indices [I>2σ(I)]	R ₁ =0.0442, wR ₂ =0.1065	R ₁ =0.0316, wR ₂ =0.0641	R ₁ =0.0661, wR ₂ =0.1492
R indices (all data)	R ₁ =0.0534, wR ₂ =0.1133	R ₁ =0.0404, wR ₂ =0.0697	R ₁ =0.1304, wR ₂ =0.1676
Largest diff. peak and hole (e ⁻ Å ⁻³)	0.90 / -0.80	0.67 / -0.75	2.26 / -1.31

MIC VALUES FOR ALL COMPOUNDS, pH = 5–8

Table S6. Minimal inhibition concentration (MIC) (μM) of premixed complexes, bidentate and monodentate ligands on Gram-positive *S. aureus* bacterial strains after one day.

MIC (μM)	<i>S. aureus</i> , sensitive				<i>S. aureus</i> , resistant				
	pH =	5	6	7	8	5	6	7	8
mim	Bacterial cell proliferation was inhibited at that pH.	>100	>100	>100	>100	>100	>100	>100	>100
Py		>100	>100	>100	>100	>100	>100	>100	>100
econ		0.78	0.19	25	12.5	6.25	3.125		
zol		>100	>100	>100	>100	>100	>100	>100	>100
phen		>100	>100	50	>100	>100	>100	>100	>100
8HQ		25	25	100	6.25	6.25	6.25		
[RhCp*(phen)Cl]Cl		>100	>100	>100	>100	>100	>100	>100	>100
+mim		>100	>100	>100	>100	>100	>100	>100	>100
+Py		>100	>100	>100	>100	>100	>100	>100	>100
+econ		0.78	1.56	25	12.5	3.125	3.125		
[RhCp*(8HQH ₋₁)Cl]		12.5	12.5	50	12.5	12.5	12.5		
+mim		12.5	12.5	50	25	12.5	12.5		
+econ		0.39	0.39	12.5	6.25	3.125	3.125		
+zol		12.5	25	50	50	12.5	12.5		

Table S7. Minimal inhibition concentration (MIC) (μM) of premixed complexes, bidentate and monodentate ligands on Gram-negative *E. coli* bacterial strains after one day.

MIC (μM)	<i>E. coli</i> , sensitive				<i>E. coli</i> , resistant				
	pH =	5	6	7	8	5	6	7	8
mim	>100	>100	>100	>100	>100	>100	>100	>100	>100
Py	>100	>100	>100	>100	>100	>100	>100	>100	>100
econ	>100	>100	>100	>100	>100	>100	>100	>100	>100
zol	>100	>100	>100	>100	>100	>100	>100	>100	>100
phen	50	50	50	25	50	50	50	50	50
8HQ	>100	>100	>100	>100	>100	>100	>100	>100	>100
[RhCp*(phen)Cl]Cl	>100	>100	>100	>100	>100	>100	>100	>100	>100
+mim	>100	>100	>100	>100	>100	>100	>100	>100	>100
+Py	>100	>100	>100	>100	>100	>100	>100	>100	>100
+econ	>100	>100	>100	>100	>100	>100	>100	>100	>100
[RhCp*(8HQH ₋₁)Cl]	>100	>100	>100	>100	>100	>100	>100	>100	>100
+mim	>100	>100	>100	>100	>100	>100	>100	>100	>100
+econ	25	25	25	25	50	100	100	50	
+zol	>100	>100	>100	>100	>100	>100	>100	>100	>100

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