

Electronic Supplementary Information (ESI)

Synthesis and Cytotoxicity against Tumor Cells of Pincer *N*-heterocyclic Ligands and their Transition Metal complexes

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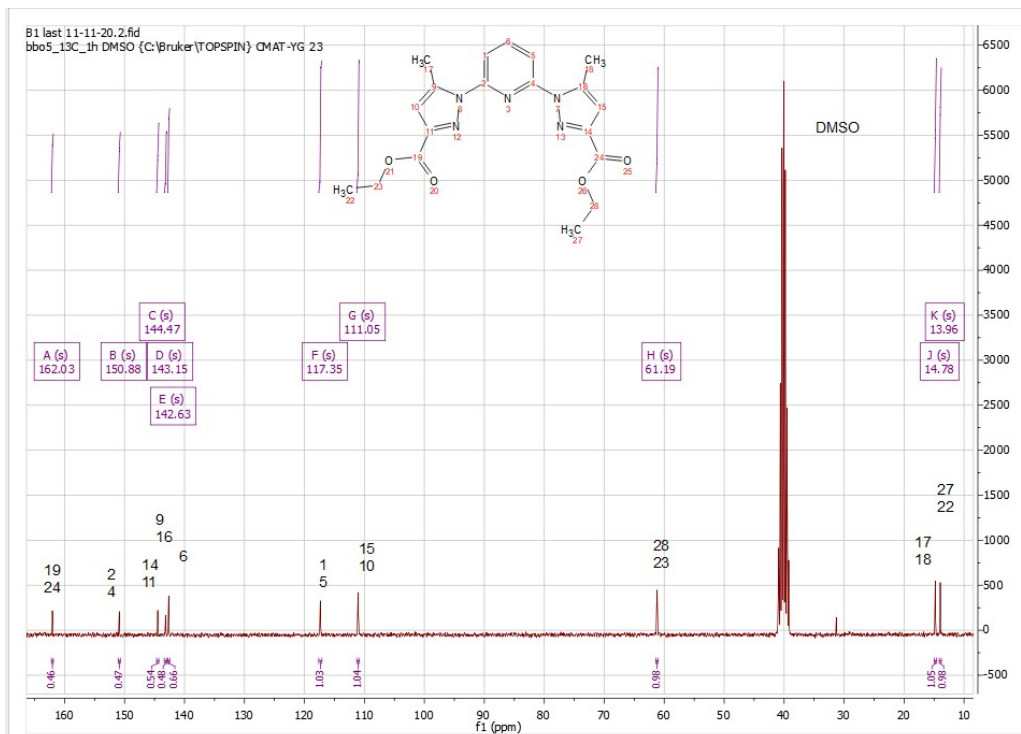
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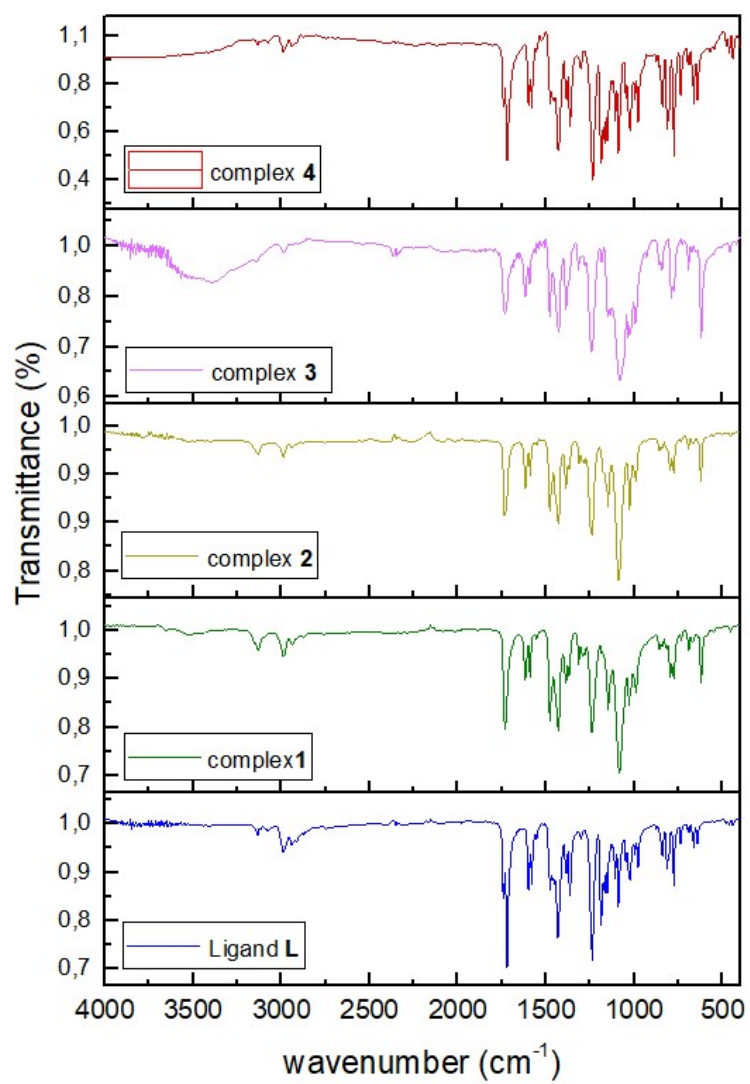
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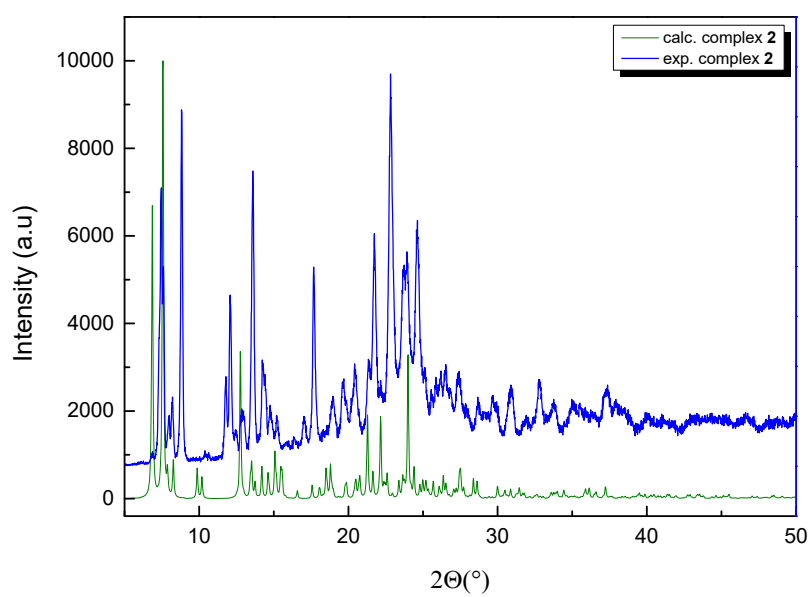
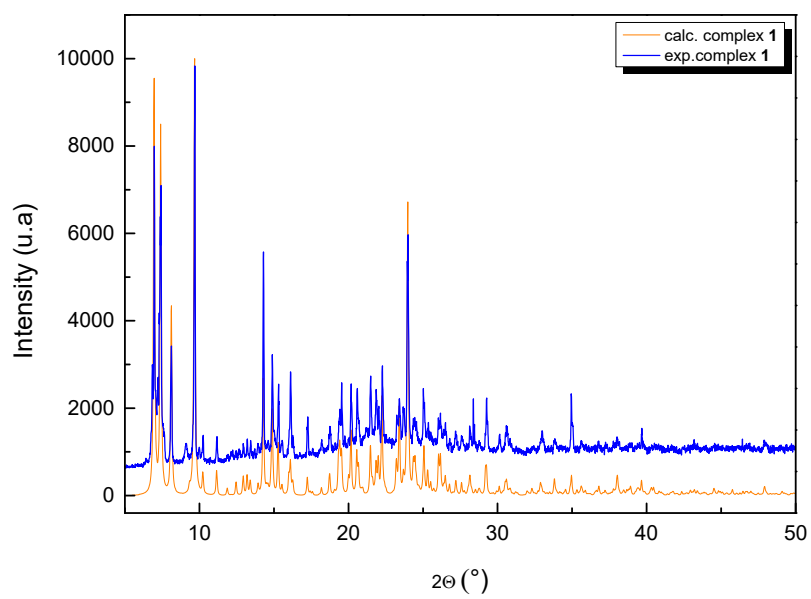
NMR ^1H and ^{13}C of ligand *L*

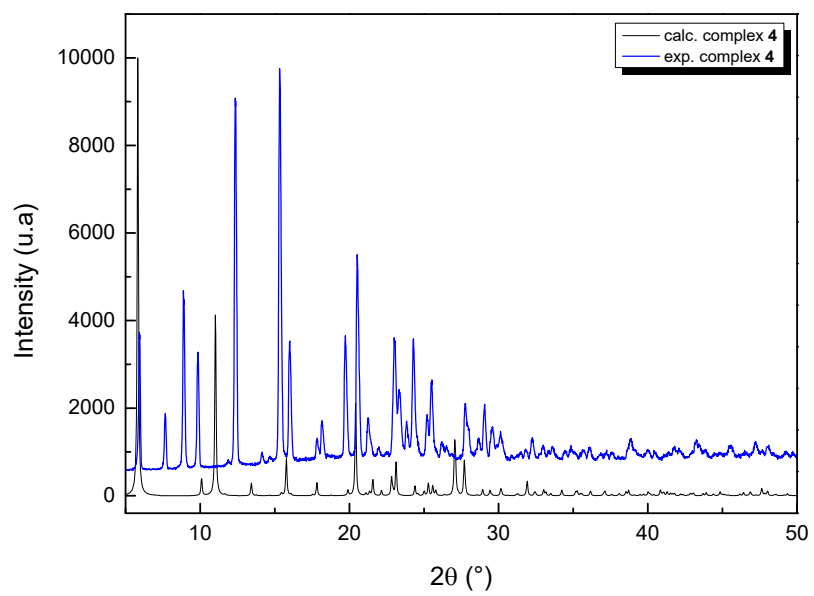
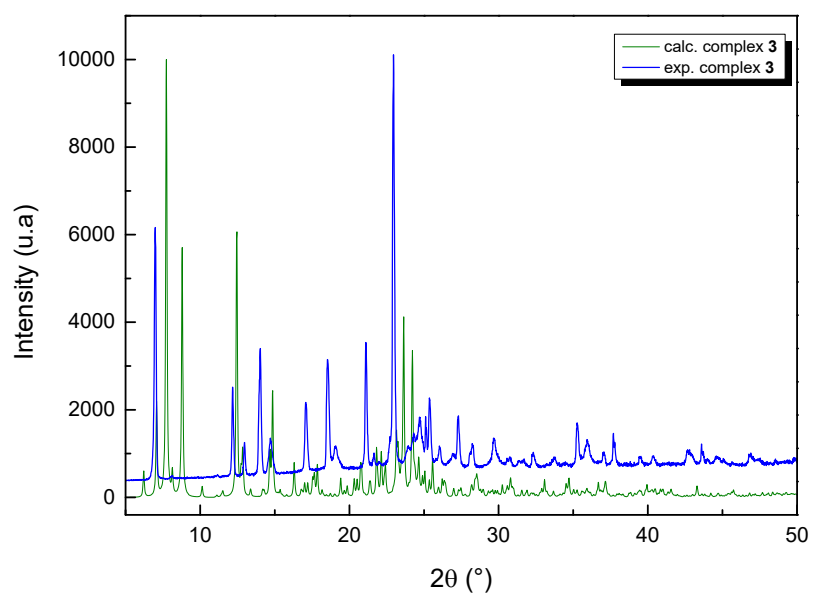


FT-IR spectroscopy comparison plot of *L* with 1-4.

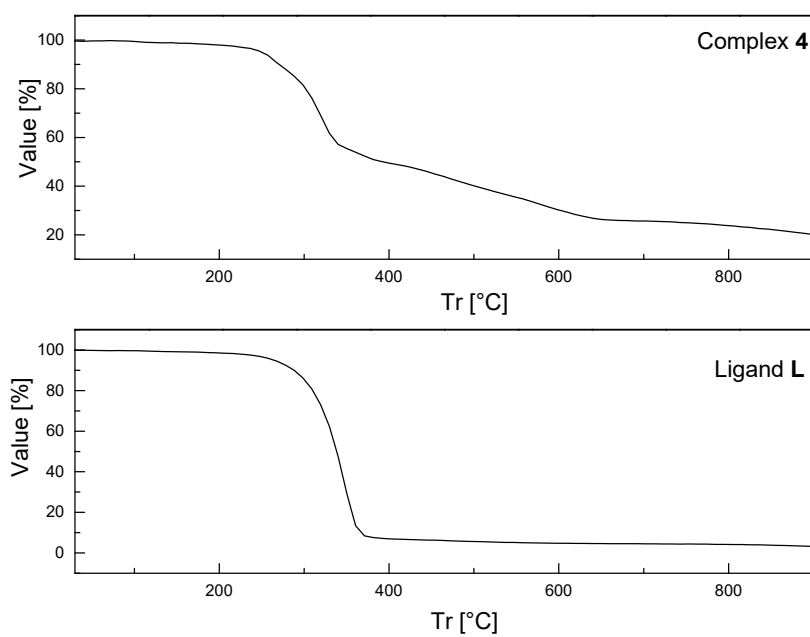


Powder X-ray diffraction of 1-4.



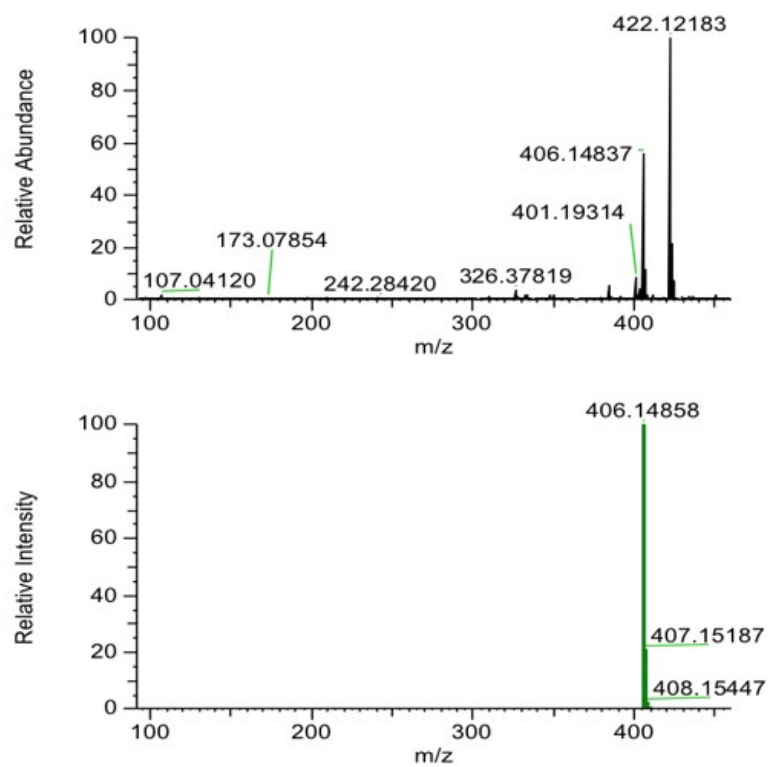


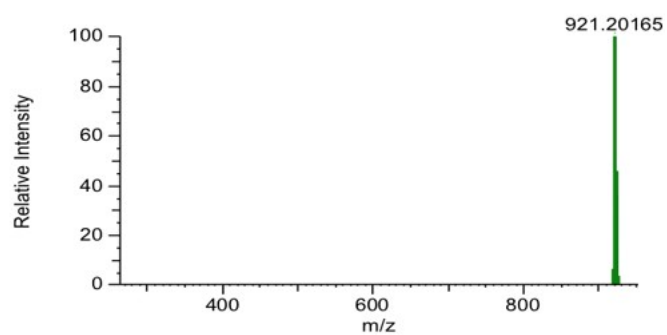
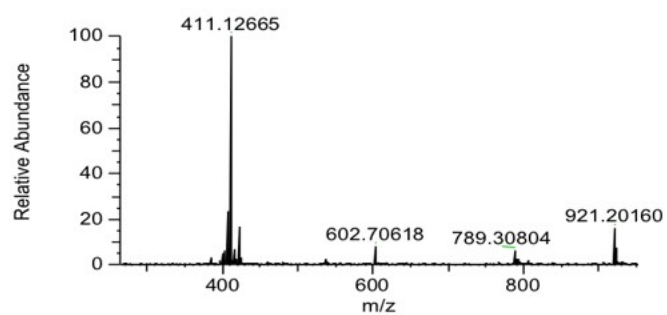
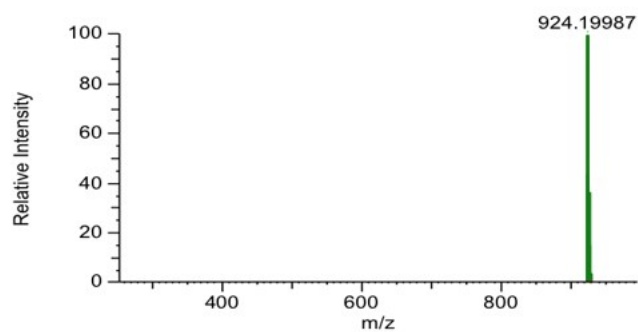
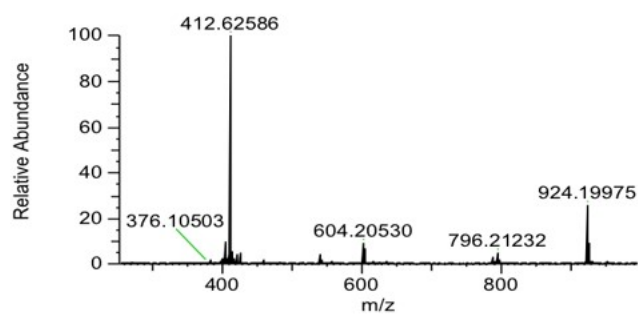
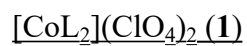
TGA of ligand *L* and 4.

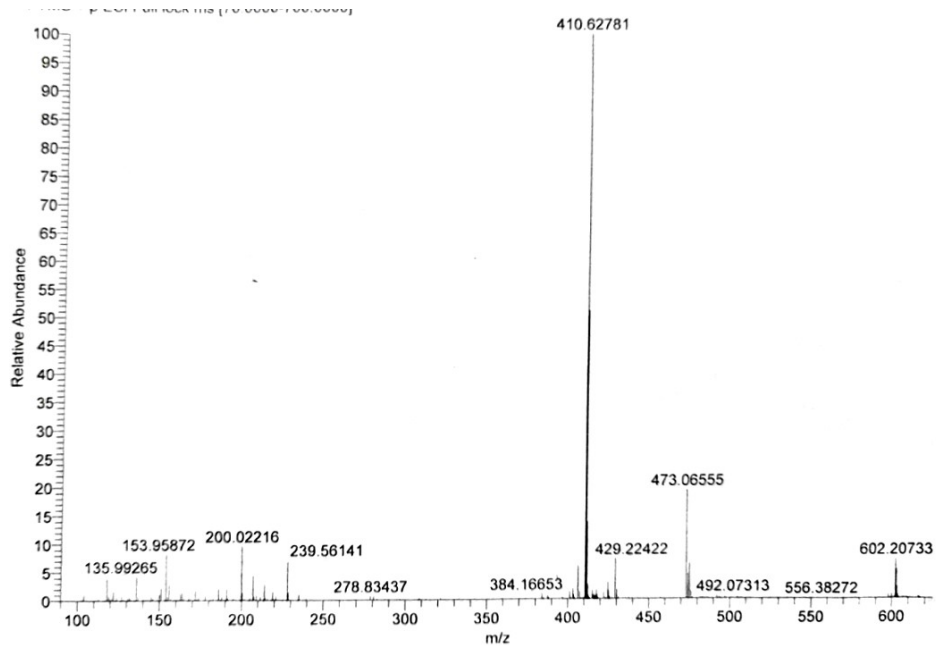
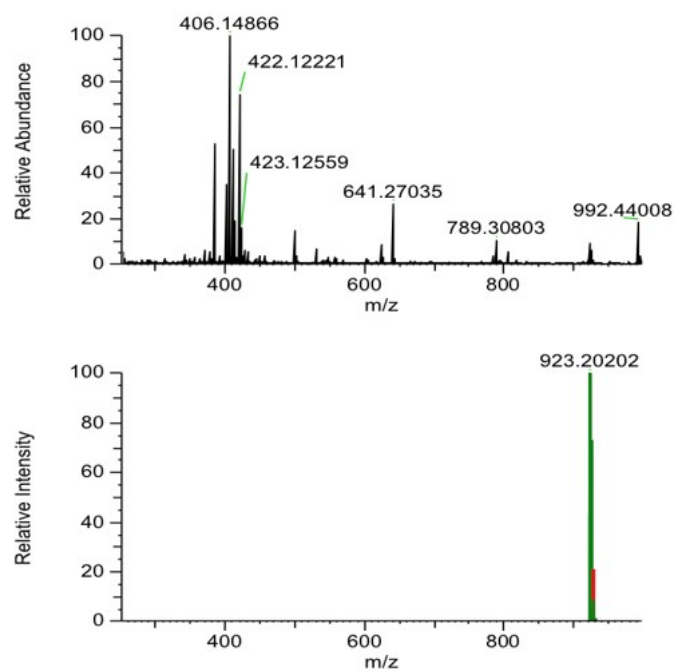
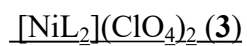


HRMS

Ligand *L*







XRD analysis

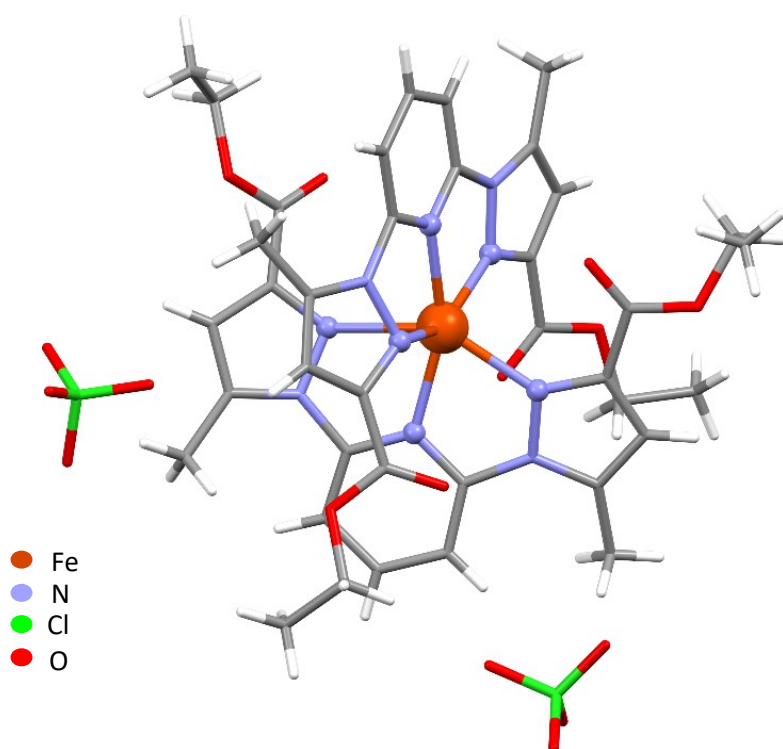


Figure S1: Perspective view of the molecular structure of **2**. Disorder was omitted for clarity.

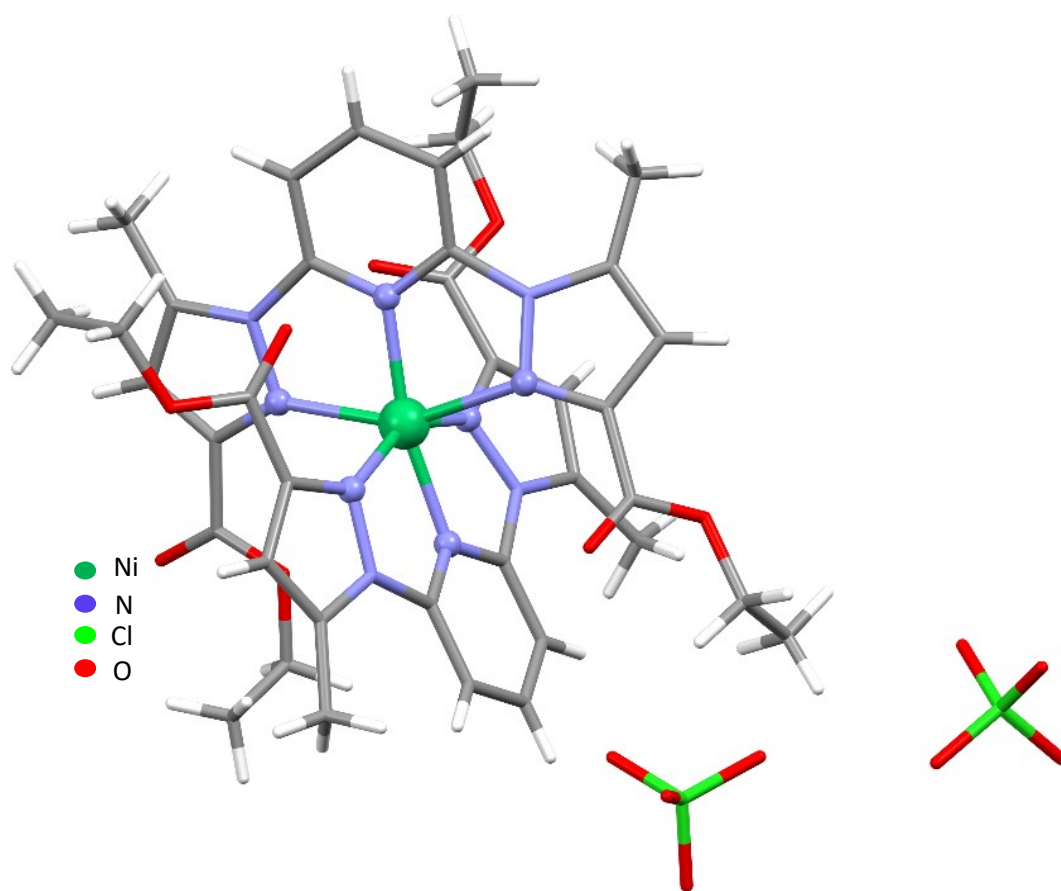


Figure S2: Perspective view of the molecular structure of **3**. Disorder was omitted for clarity.

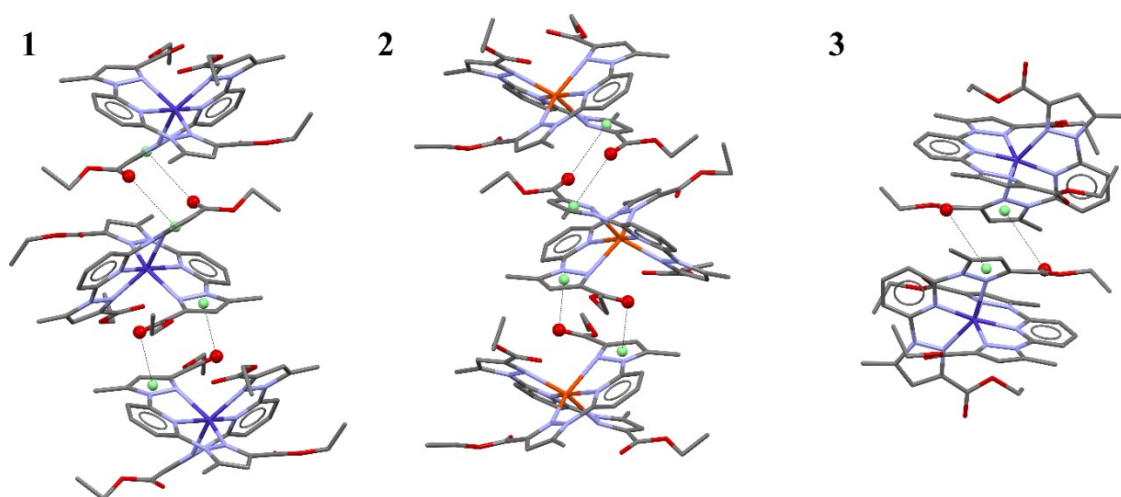


Figure S3: Packing motives found in complexes **1**, **2** and **3**, involving the twisted pyrazolyl ring centroids and the carbonyl oxygen.

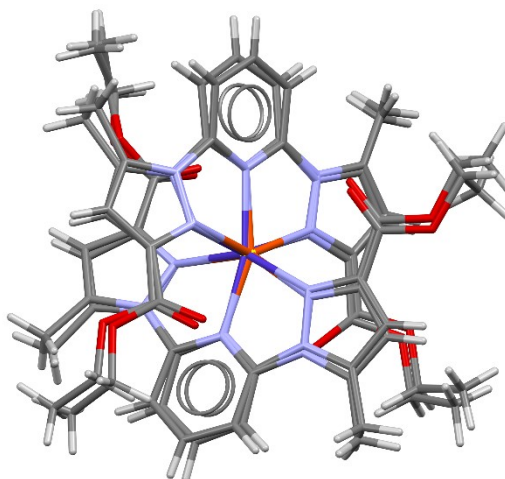


Figure S4: superposition of the Fe complex **2** on the Co complex **1**.

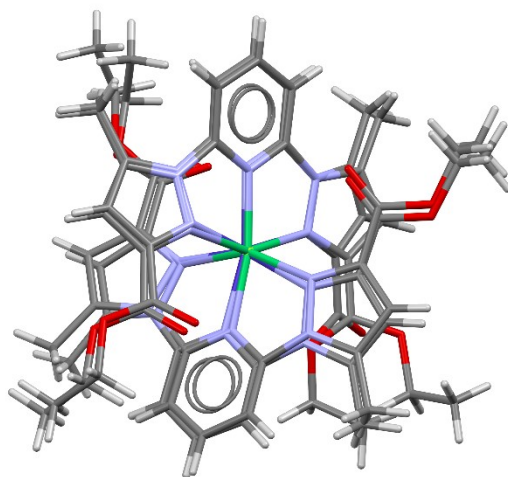


Figure S5: superposition of the Ni complex **3** on the Co complex **1**.

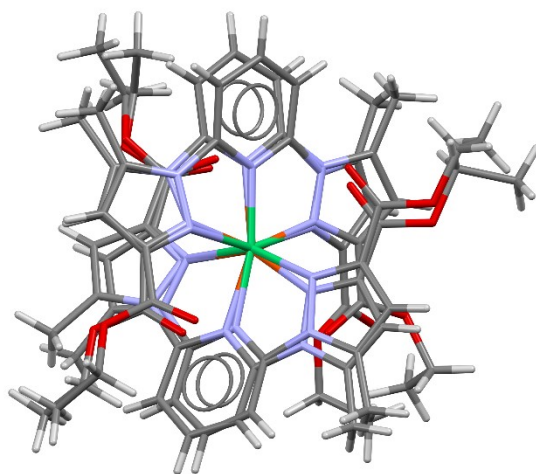


Figure S6: superposition of the Ni complex **3** on the Fe complex **2**.

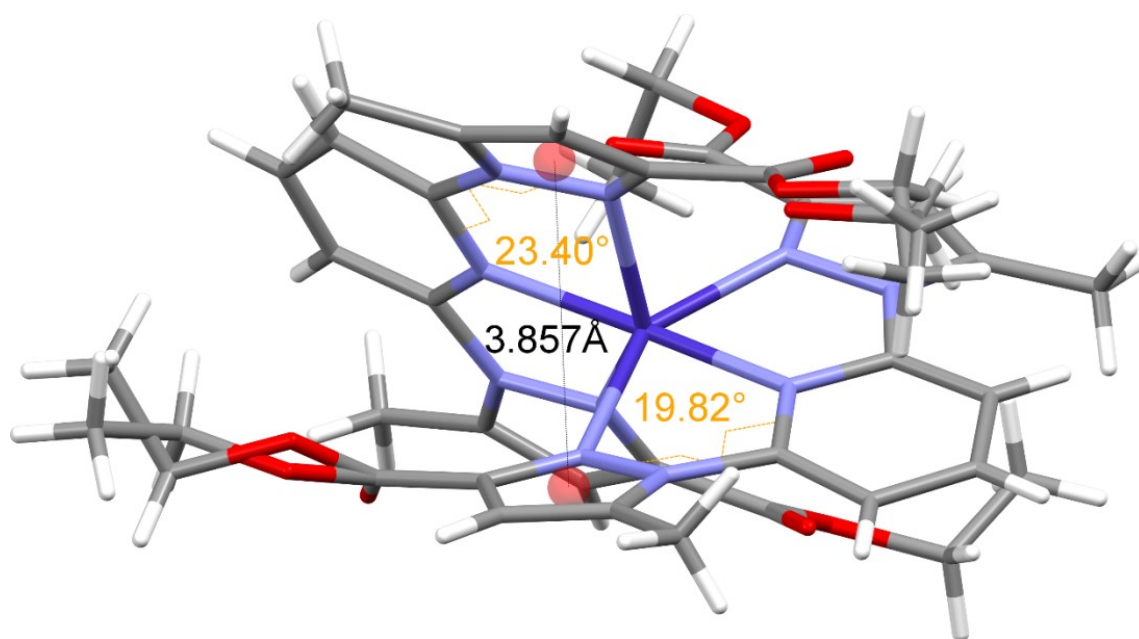


Figure S7: Torsion angles between the pyrazolyl and pyridyl rings, result in a short centroid-centroid distance between the pyrazolyl rings for **1**. Similar features are present in **2** and **3**.

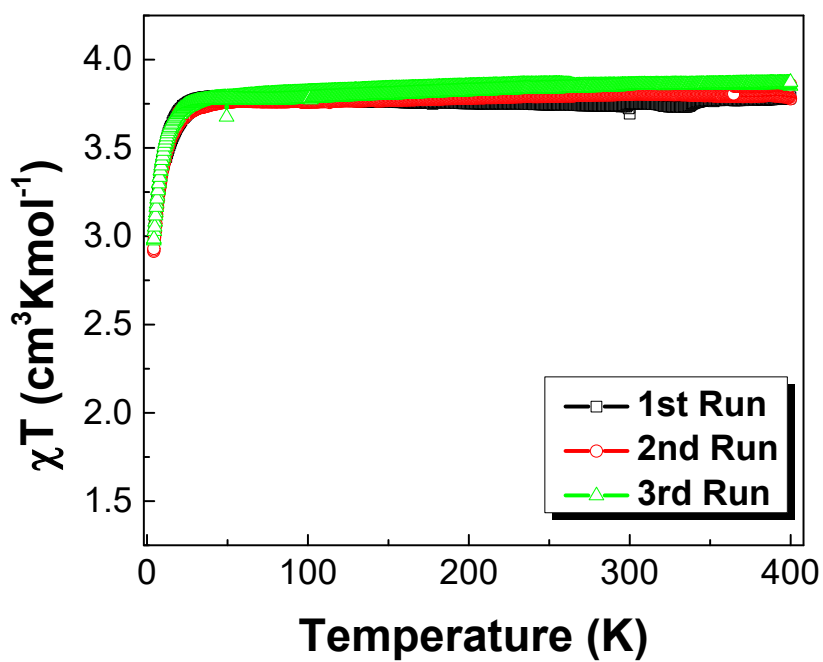


Figure S8 : Temperature-dependent $\chi_M T$ plot for **2** over three temperature runs, demonstrating the absence of any spin state crossover, over the whole temperature range investigated.

Table S1: Crystal data and structure refinements details for **1-4**.

Identification code	1	2	3	4
Empirical formula	C ₃₈ H ₄₂ C ₁₂ CoN ₁₀ O ₁₆	C ₃₈ H ₄₂ C ₁₂ FeN ₁₀ O ₁₆	C ₃₈ H ₄₂ C ₁₂ N ₁₀ NiO ₁₆	C ₁₉ H ₂₁ Cl ₂ MnN ₅ O ₄
Formula weight	1024.64	1034.24	1024.42	509.25
T (K)	297(2)	297(2)	150(2)	297(2)
Wavelength (Å)	0.71073	0.71073	0.71073	0.71073
Crystal system	Monoclinic	Monoclinic	Monoclinic	Trigonal
Space group	<i>P</i> 2 ₁ / <i>n</i>	<i>P</i> 2 ₁ / <i>c</i>	<i>C</i> 2/ <i>c</i>	<i>R</i> -3 <i>c</i>
Unit cell dim. <i>a</i> (Å)	12.8107(4)	14.3475(3)	29.5243(11)	30.3293(6)
<i>b</i> (Å)	14.1995(4)	25.7510(6)	13.8841(4)	30.3293(6)
<i>c</i> (Å)	25.5626(6)	13.6655(4)	22.5685(9)	13.6082(4)
β (°)	96.566(3)	114.200(3)	105.786(4)	
Volume (Å ³)	4619.5(2)	4605.2(2)	8902.4(6)	10840.7(5)
<i>Z</i>	4	4	8	18
Density (calc.) (g/cm ³)	1.473	1.492	1.529	1.404
Abs. coeff. (mm ⁻¹)	0.567	0.526	0.639	0.803
F(000)	2116	2137	4240	4698
Crystal size (mm ³)	0.50 x 0.30 x 0.10	0.47 x 0.30 x 0.08	0.50 x 0.40 x 0.20	0.50 x 0.30 x 0.25
θ range for data collection	3.201 to 25.700°.	2.794 to 25.243°.	2.846 to 25.689°.	2.686 to 25.690°.
Reflections collected	36203	30251	39648	30303
Independent reflections	8729 [<i>R</i> _(int) = 0.0350]	8298 [<i>R</i> _(int) = 0.0412]	8411 [<i>R</i> _(int) = 0.0353]	2279 [<i>R</i> _(int) = 0.0268]
Completeness to θ = 25.242°	99.4 %	99.5 %	99.4 %	99.2 %
Absorption correction	Semi-empirical from equivalents			
Max. and min. transm.	1.00000 and 0.74762	1.00000 and 0.37734	1.00000 and 0.71206	1.00000 and 0.83245
Refinement method	Full-matrix least-squares on <i>F</i> ²			
Data / restr. / param.	8729 / 136 / 762	8298 / 710 / 792	8411 / 10 / 628	2279 / 2 / 178
Goodness-of-fit on <i>F</i> ²	1.052	1.094	1.039	1.077
Final <i>R</i> indices [<i>I</i> > 2σ(<i>I</i>)]	<i>R</i> ₁ = 0.0453, <i>wR</i> ₂ = 0.1225	<i>R</i> ₁ = 0.0652, <i>wR</i> ₂ = 0.1625	<i>R</i> ₁ = 0.0369, <i>wR</i> ₂ = 0.0931	<i>R</i> ₁ = 0.0313, <i>wR</i> ₂ = 0.0830
<i>R</i> indices (all data)	<i>R</i> ₁ = 0.0542, <i>wR</i> ₂ = 0.1294	<i>R</i> ₁ = 0.0881, <i>wR</i> ₂ = 0.1780	<i>R</i> ₁ = 0.0400, <i>wR</i> ₂ = 0.0953	<i>R</i> ₁ = 0.0334, <i>wR</i> ₂ = 0.0844
$\Delta\rho$ max/min (e.Å ⁻³)	0.611 and -0.239	1.151 and -0.266	0.515 and -0.452	0.203 and -0.179

Table S2: Bond lengths and bond angles around the metal centre for **1-3**.

Bond lengths	1	2*	3*
M-N _{ax} 1			
M-N _{ax} 2			
M-N _{eq} 1			
M-N _{eq} 2			
M-N _{eq} 3			
M-N _{eq} 4			
Bond angles			
N _{Ax} 1-M-N _{Ax} 2	165.39(8)	157.61(11)	168.21(6)
N _{Ax} 1-M-N _{eq} 1	75.09(7)	72.54(10)	77.29(6)
N _{Ax} 1-M-N _{eq} 2	94.60(7)	90.42(9)	97.96(6)
N _{Ax} 1-M-N _{eq} 3	74.44(7)	72.48(11)	76.19(6)
N _{Ax} 1-M-N _{eq} 4	116.90(7)	125.27(11)	109.03(6)
N _{Ax} 2-M-N _{eq} 1	94.37(7)	92.25(10)	91.74(6)
N _{Ax} 2-M-N _{eq} 2	75.12(7)	71.67(10)	76.99(6)
N _{Ax} 2-M-N _{eq} 3	117.28(7)	123.74(11)	114.78(6)
N _{Ax} 2-M-N _{eq} 4	74.29(7)	72.89(11)	76.51(6)
N _{eq} 1-M- N _{eq} 2	90.63(8)	85.50(10)	87.54(6)
N _{eq} 1-M- N _{eq} 3	148.04(7)	143.97(10)	153.46(6)
N _{eq} 1-M- N _{eq} 4	99.36(8)	101.08(11)	98.30(6)
N _{eq} 2-M- N _{eq} 3	101.18(8)	103.08(10)	97.05(6)
N _{eq} 2-M- N _{eq} 4	148.40(7)	144.17(10)	153.01(6)
N _{eq} 3-M- N _{eq} 4	86.01(8)	92.11(12)	89.41(6)

* Values for bond lengths and angles were obtained after superposition of **2** and **3** on **1** for better comparison between the complexes

Table S3. Bond lengths and bond angles for **4**.

Bond lengths	4
Mn Cl ₂ _{Ax}	2.3359(5)
Mn N ₃ _{eq}	2.3120(15)

Mn N9 _{eq}	2.268(2)
Mn N3 _{eq}	2.3120(15)
Mn Cl2 _{eq}	2.3359(5)
Bond angles	
Cl _{Ax} Mn Cl _{eq}	133.83(4)
Cl _{Ax} Mn N _{eq1}	105.20(4)
Cl _{Ax} Mn N _{eq2}	113.2(5)
Cl _{Ax} Mn N _{eq3}	90.94(4)
Cl _{eq} Mn N _{eq1}	105.19(4)
Cl _{eq} Mn N _{eq2}	112.9(5)
Cl _{eq} Mn N _{eq3}	90.95(4)
N _{eq1} Mn N _{eq2} (*)	72.6(5)
N _{eq1} Mn N _{eq3}	138.37(8)
N _{eq2} Mn N _{eq3} (*)	65.8(5)

(*) Neq2 is disordered around the 2-fold axis,
hence the difference between both values

Table S4 RMSD and maximal displacement between complexes **1**, **2**, and **3**.

	RMSD (Å)	Max Deviation (Å)
Superposition of 2 on 1	0.5216	2.8204
Superposition of 3 on 1	0.7219	2.3544
Superposition of 3 on 2	0.8904	2.8630