

Supplementary information for

Mononuclear complexes of Fe^{II}, Co^{II} and Co^{III} containing imine-based ligands of 8-aminoquinoline and 7-aminoindazole: spin state tuning of Fe^{II} complexes in solution

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1. NMR spectroscopy

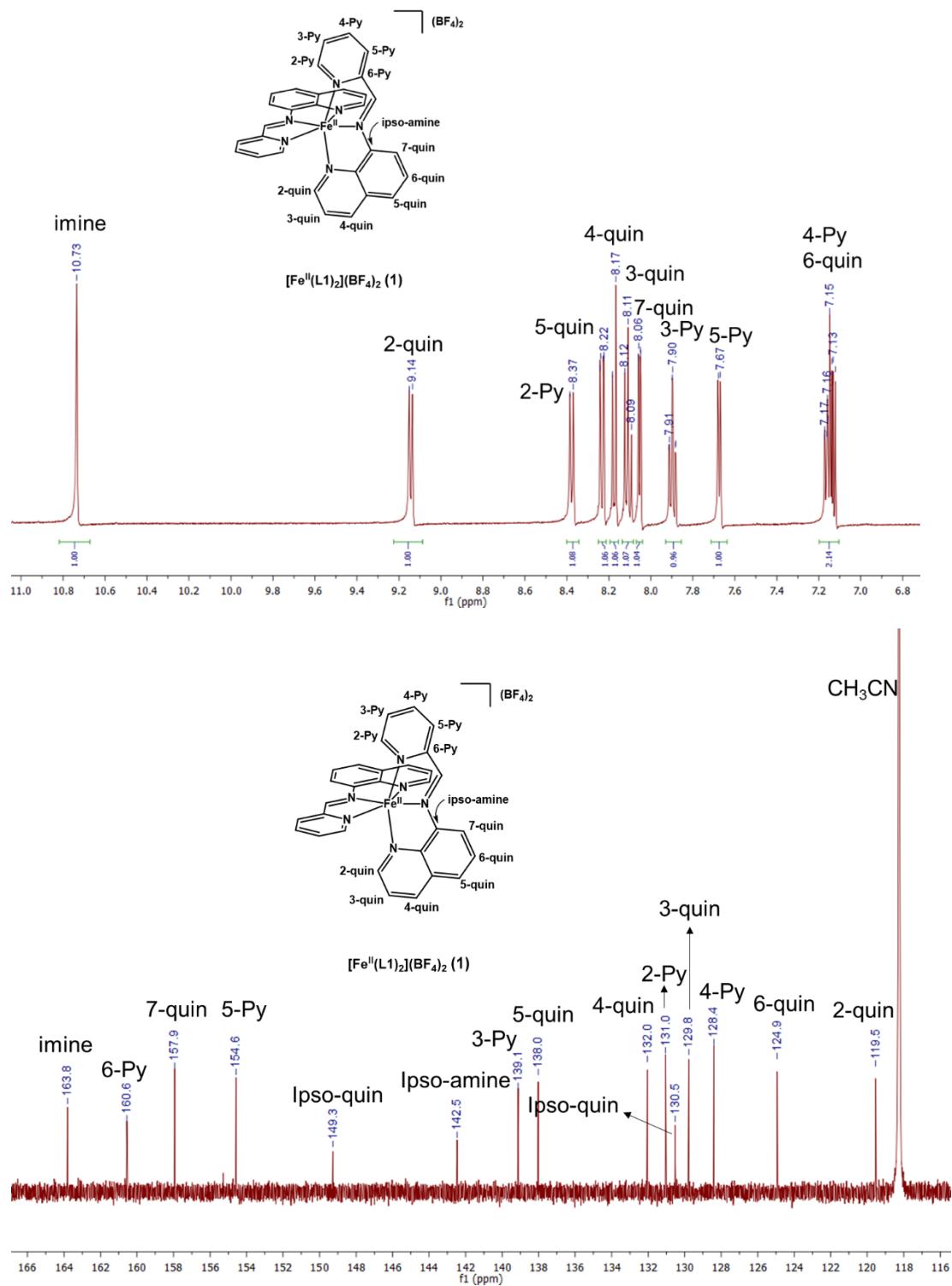


Fig S1. ^1H NMR and ^{13}C NMR of complex **1** in CD_3CN solution.

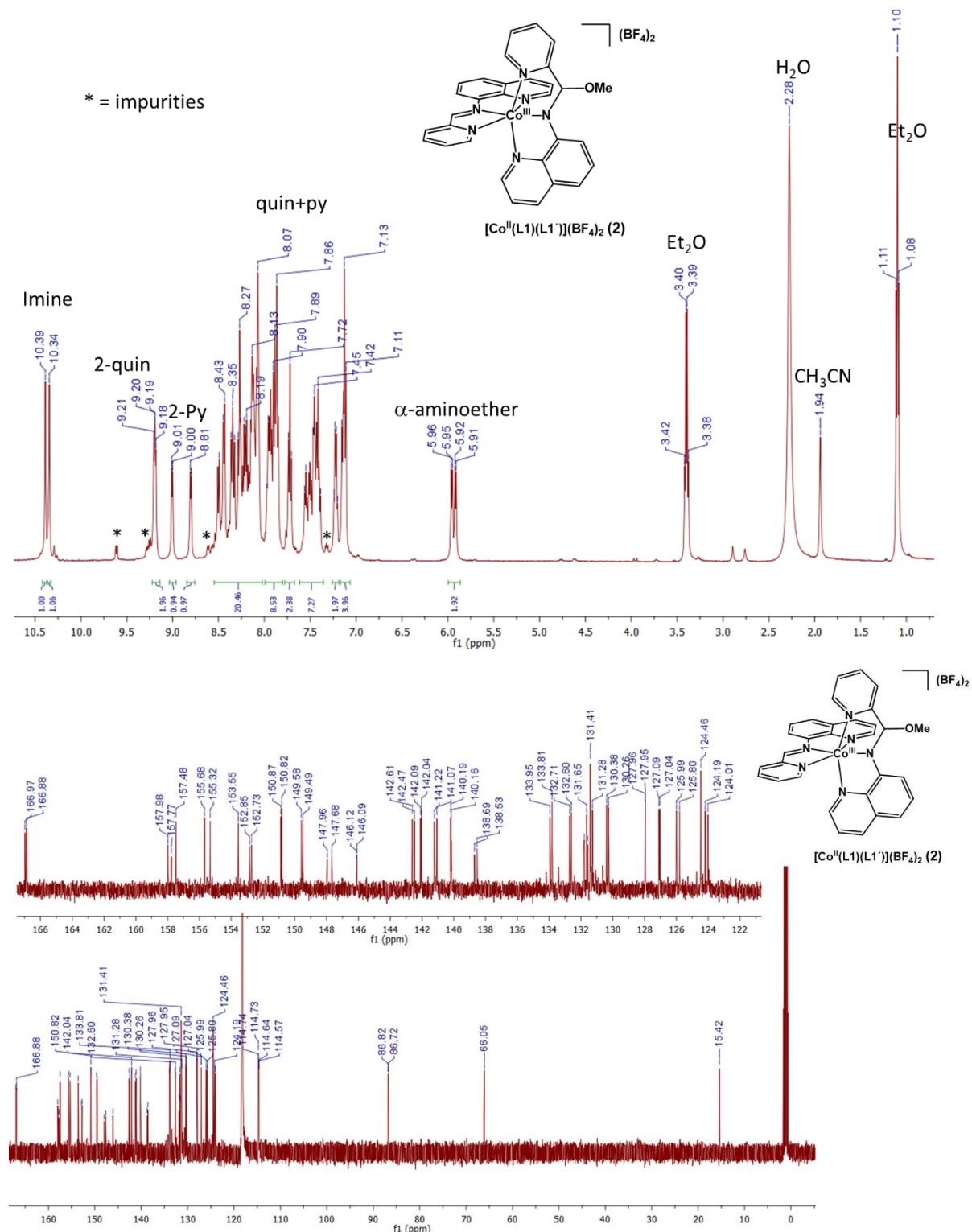


Fig S2. ^1H NMR and ^{13}C NMR of complex **2** in CD_3CN solution.

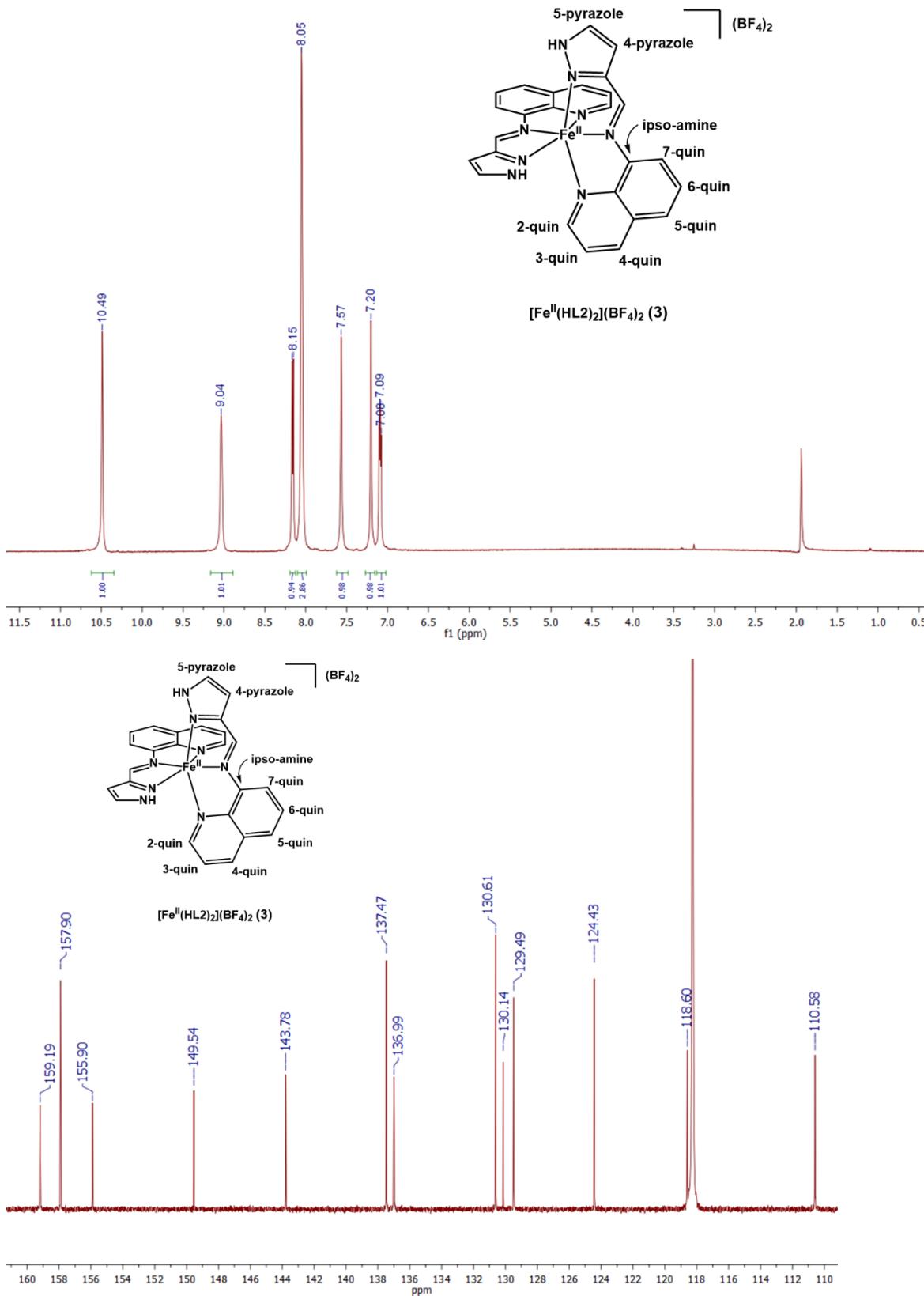


Fig S3. ^1H NMR and ^{13}C NMR of complex **3** in CD_3CN solution.

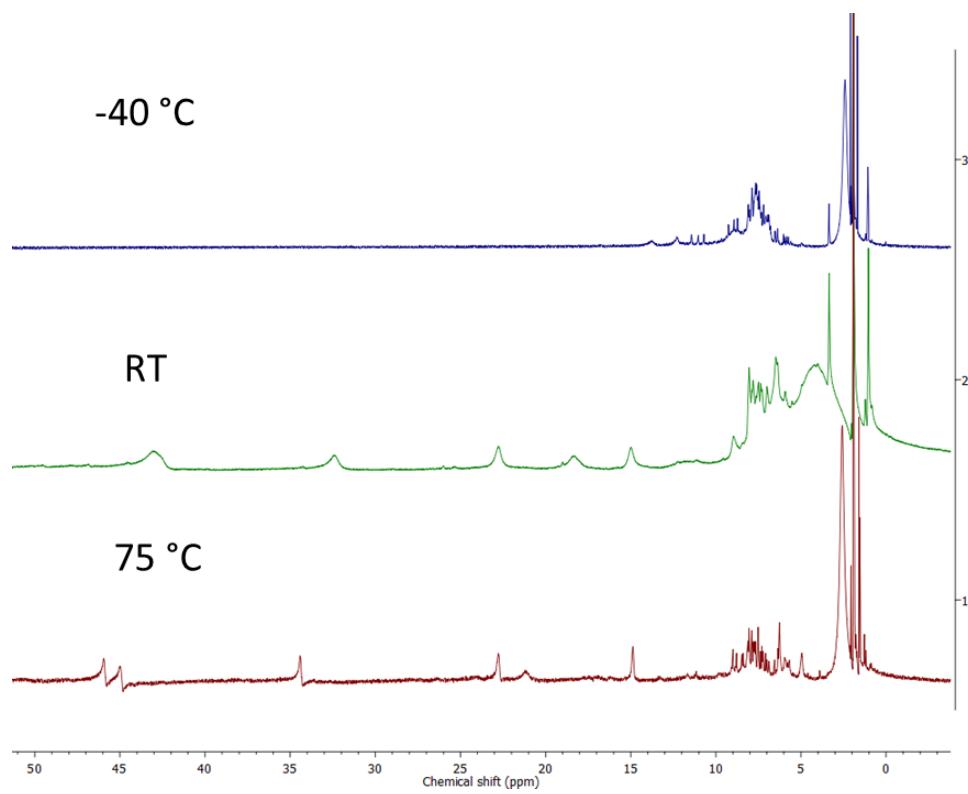


Figure S4. Paramagnetic VT-¹H NMR for complex 4 in CD_3CN solution.

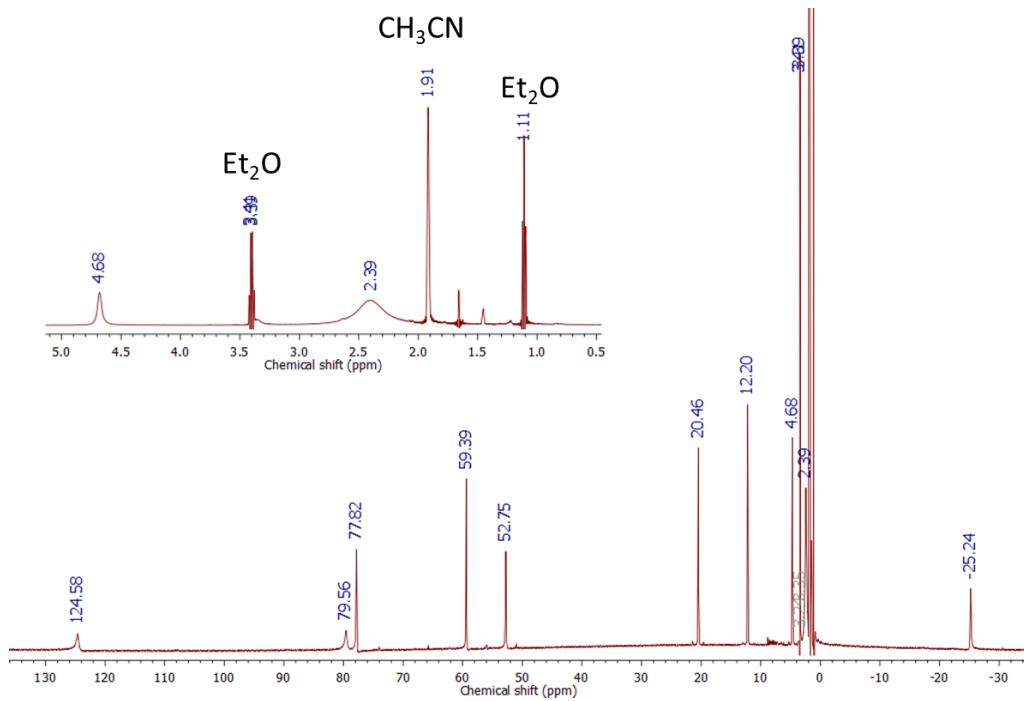


Figure S5. Paramagnetic ¹H NMR for complex 5-(MeOH)₂(Et₂O)_{0.5} in CD_3CN solution. Inset shows an expansion of the 0.5-5.0 ppm range.

2. IR Spectroscopy

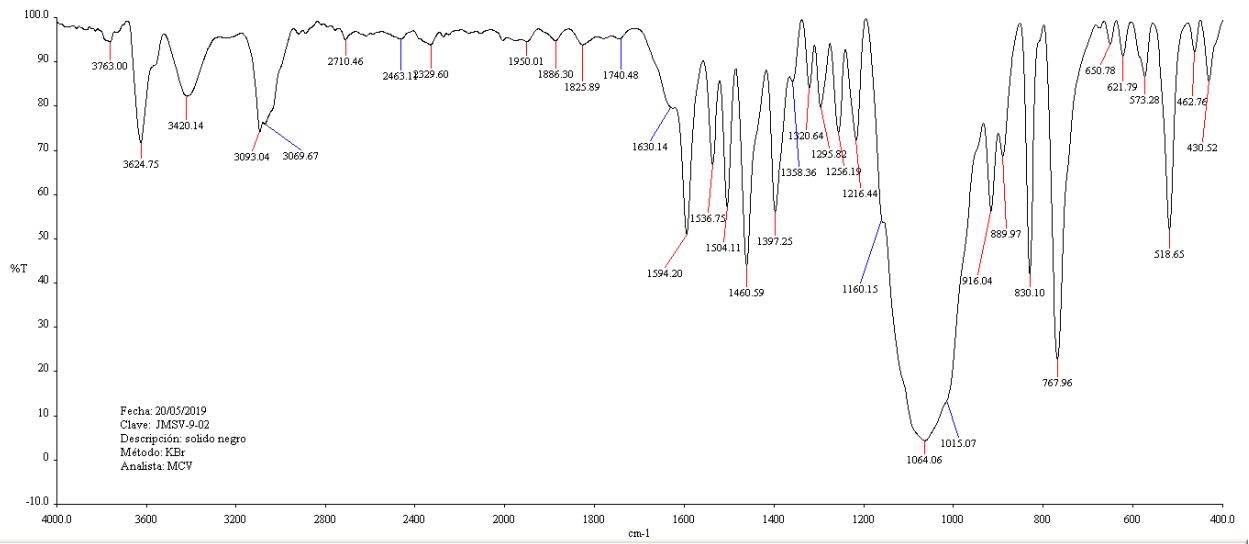


Figure S6. IR spectrum for complex 1.

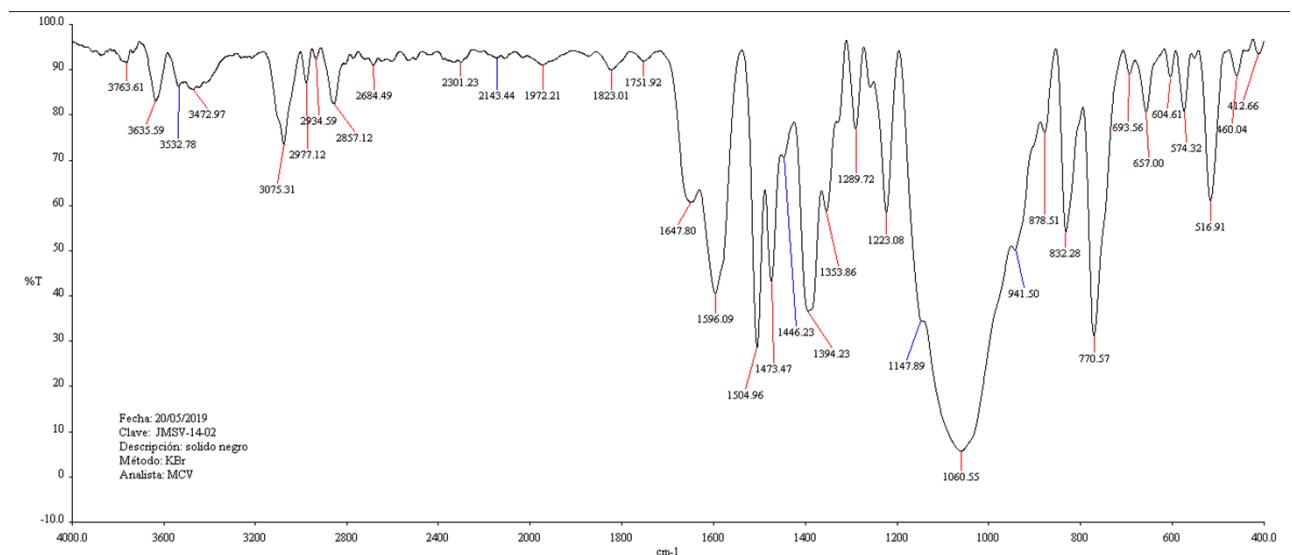


Figure S7. IR spectrum for complex 2.

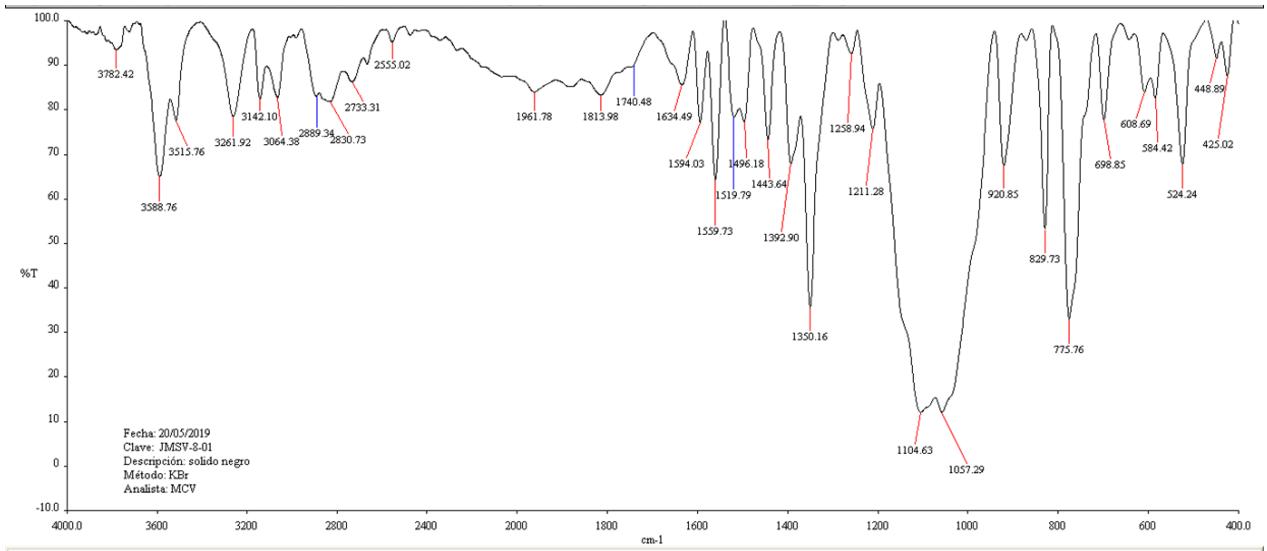


Figure S8. IR spectrum for complex 3.

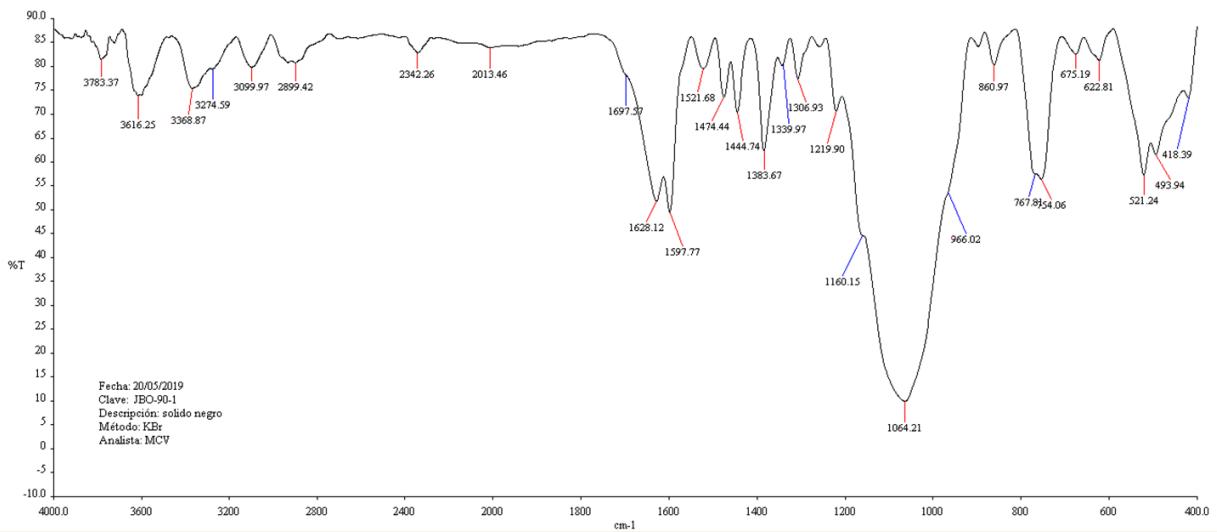


Figure S9. IR spectrum for complex 4.

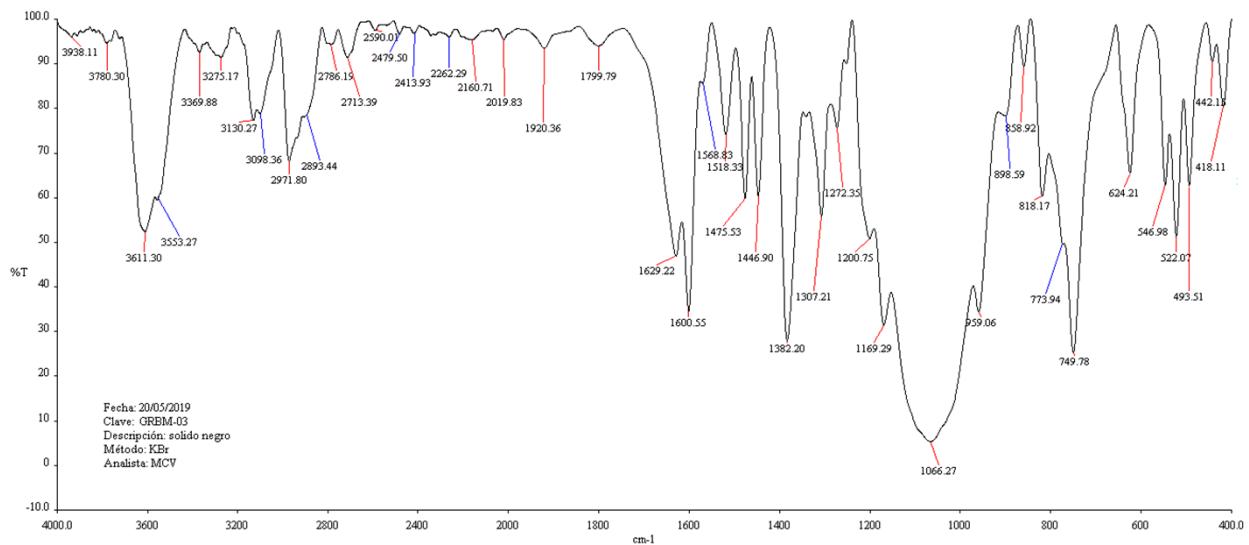


Figure S10. IR spectrum for complex 5.

3. Comments on checkcif alert for complex 2

The following alert level B for complex 2 (JMSV14sq.cif) was not possible to remove. The alert in this file is:

! Alert level B
PLAT230_ALERT_2_B Hirshfeld Test Diff for O19 --C31 . 9.0 s.u.

O19 and C31 are the atoms corresponding to the methoxy group in the α -aminoether, which has some disorder. Attempts to model the disorder were not successful. Thus, this alert B is caused by vibration of the entire methoxy group and not due to wrong atom assignment.^{S1}

- S1. G. Valluru, S. Rahman, P. E. Georghiou, L. N. Dawe, A. N. Alodhayb and L. Y. Beaulieu, *N. J. Chem.*, 2014, **38**, 5868-5872.

Table S1. Crystal data and structure refinement for complexes **2**, **3** and **5**.

Compound	2 ·solvents	3 ·solvents	5 ·(MeOH)(Et ₂ O) _{0.5}
Identification code	JMSV14sq	JMSV01sq	ICQ02
Empirical formula	C ₃₁ H ₂₅ B ₂ CoF ₈ N ₆ O	C ₂₆ H ₂₀ B ₂ F ₈ FeN ₈	C ₆₀ H ₆₆ B ₄ Co ₂ F ₁₆ N ₁₆ O ₅
Formula weight	730.12	673.97	1556.38
Temperature	298(2) K	293(2) K	166(2) K
Wavelength	0.71073 Å	0.71073 Å	0.71073 Å
Crystal system	Monoclinic	Monoclinic	Monoclinic
Space group	P2 ₁ /n	I2/a	C2/c
Unit cell dimensions	a = 12.1427(7) Å, α = 90° b = 15.5443(9) Å, β = 106.570(2)° c = 19.5164(12) Å, γ = 90°	a = 16.375(3) Å, α = 90° b = 12.327(3) Å, β = 102.25(3)° c = 17.399(4), γ = 90°	a = 21.699(9) Å, α = 90° b = 12.540(4) Å, β = 119.63(2)° c = 15.873(8) Å, γ = 90°
Volume	3530.7(4) Å ³	3431.9(13) Å ³	3754(3) Å ³
Z	4	4	2
Density (calculated)	1.374 mg/m ³	1.304 mg/m ³	1.377 mg/m ³
Absorption coefficient	0.562 mm ⁻¹	0.511 mm ⁻¹	0.537 mm ⁻¹
F(000)	1480	1360	1592
Crystal size	0.390 x 0.250 x 0.150 mm ³	0.320 x 0.290 x 0.180 mm ³	0.460 x 0.300 x 0.200 mm ³
Theta range for data collection	2.186 to 25.026°.	2.396 to 26.575°.	2.660 to 28.907°.
Index ranges	-14 ≤ h ≤ 14, -18 ≤ k ≤ 18, -23 ≤ l ≤ 23	-20 ≤ h ≤ 20, -15 ≤ k ≤ 15, -21 ≤ l ≤ 21	-29 ≤ h ≤ 29, -17 ≤ k ≤ 17, -21 ≤ l ≤ 21
Reflections collected	113448	55537	136671
Independent reflections	6231 [R(int) = 0.0997]	3536 [R(int) = 0.1307]	4897 [R(int) = 0.0784]
Completeness to theta	99.8 %	99.8 %	99.6 %
Max. and min. transmission	0.7458 and 0.6496	0.7455 and 0.5768	0.7458 and 0.6754
Data / restraints / parameters	6231 / 306 / 516	3536 / 0 / 204	4897 / 21 / 279
Goodness-of-fit on F ²	0.971	1.037	1.222
Final R indices [I>2sigma(I)]	R1 = 0.0620, wR2 = 0.1709	R1 = 0.0587, wR2 = 0.1570	R1 = 0.0548, wR2 = 0.1578
R indices (all data)	R1 = 0.0840, wR2 = 0.1892	R1 = 0.1046, wR2 = 0.1812	R1 = 0.0701, wR2 = 0.1660
Largest diff. peak and hole	0.945 and -0.488 e.Å ⁻³	0.420 and -0.343 e.Å ⁻³	0.658d -0.541 e.Å ⁻³

4. Supramolecular interactions

Complex 3

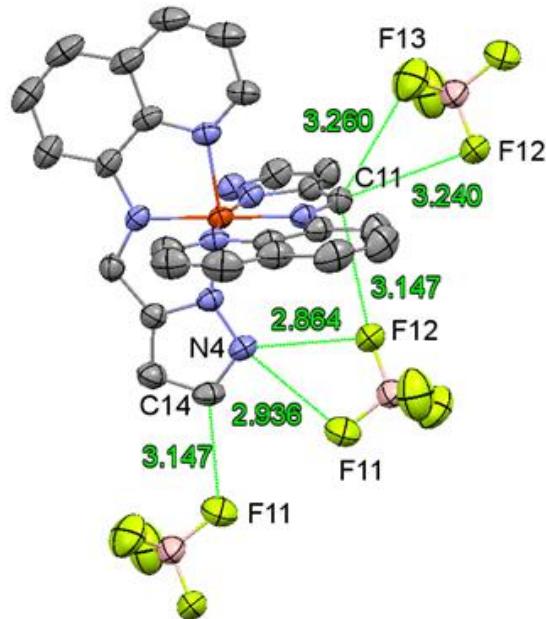


Figure S11. H-bonds of the type $\text{NH}\cdots\text{F-BF}_3$, $\text{N}=\text{C}-\text{H}\cdots\text{F-BF}_3$ and $\text{C}-\text{H}\cdots\text{F-BF}_3$ found in the crystal structure of complex 3.

Table S2. Hydrogen bonds for complex 3 [\AA and $^\circ$].

D-H...A	d(D-H)	d(H...A)	d(D...A)	$\angle(\text{DHA})$
C(2)-H(2)...N(2)#1	0.93	2.65	3.116(5)	111.3
N(4)-H(4A)...F(11)#1	0.86	2.31	2.936(4)	129.5
N(4)-H(4A)...F(12)#1	0.86	2.01	2.864(4)	169.9
C(3)-H(3)...F(14)#2	0.93	2.53	3.434(6)	164.9
C(11)-H(11)...F(12)#3	0.93	2.36	3.240(4)	157.4
C(14)-H(14)...F(11)#4	0.93	2.31	3.148(5)	149.6

Symmetry transformations used to generate equivalent atoms:

#1 -x+1/2,y,-z+1 #2 -x+1,-y+1,-z+1 #3 -x+1/2,-y+1/2,-z+1/2
#4 x-1/2,-y,z

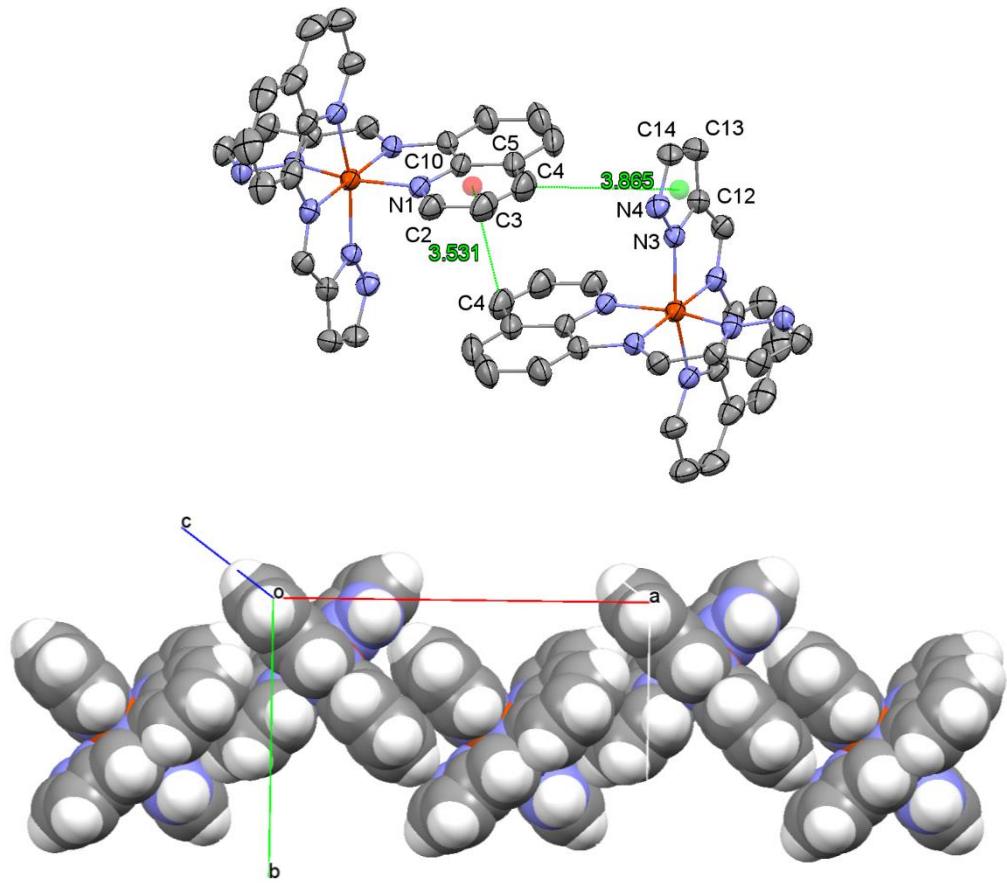


Figure S12. Embracing π - π and C-H- π interactions found in the crystal structure of complex 3 (top). Supramolecular zig-zag polymer running along *a*-axis (bottom).

Complex 2

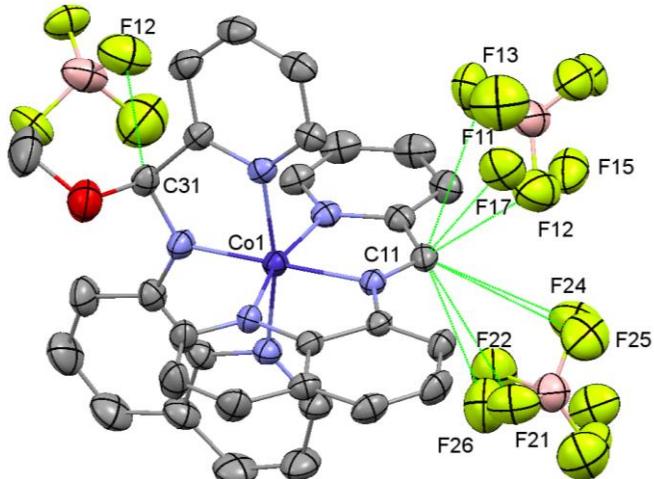


Figure S13. H-bonds interactions (green dotted lines) present in complex 2.

Table S3. Hydrogen bonds for Complex **2** [Å and °].

D-H...A	d(D-H)	d(H...A)	d(D...A)	<(DHA)
C(31)-H(31)...F(12)#3	0.98	2.36	3.279(10)	156.4
C(22)-H(22)...F(26)	0.93	2.45	3.144(14)	131.2
C(11)-H(11)...F(12)	0.93	2.34	2.939(10)	122.1
C(11)-H(11)...F(15)	0.93	2.48	3.142(10)	128.3
C(11)-H(11)...F(25)	0.93	2.56	3.330(13)	140.6

Symmetry transformations used to generate equivalent atoms:

#1 x+1/2,-y+1/2,z-1/2 #2 x+1,y,z #3 -x+1/2,y+1/2,-z+1/2

#4 -x+1,-y+1,-z+1

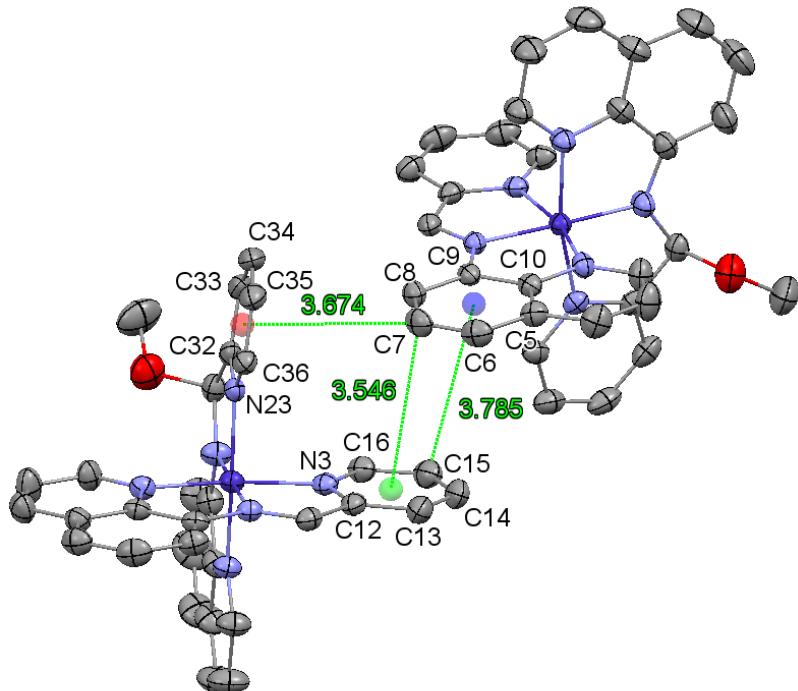


Figure S14. Perspective view of the dimers formed via $\pi\cdots\pi$ and $\text{C}-\text{H}\cdots\pi$ interactions observed in the crystal lattice of complex **2**.

Complex 5

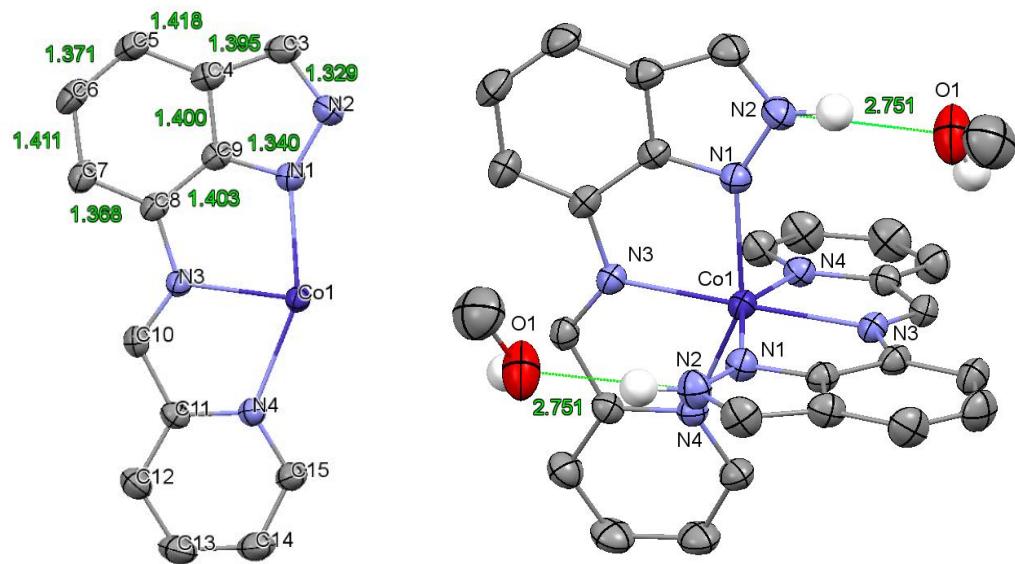


Figure S15. Perspective views of complex 5 showing the C-C bond lengths for the 7-amino-2*H*-indazole unit (left) and the hydrogen bonding of the NH_{indazole} unit to the methanol solvent molecule (right).

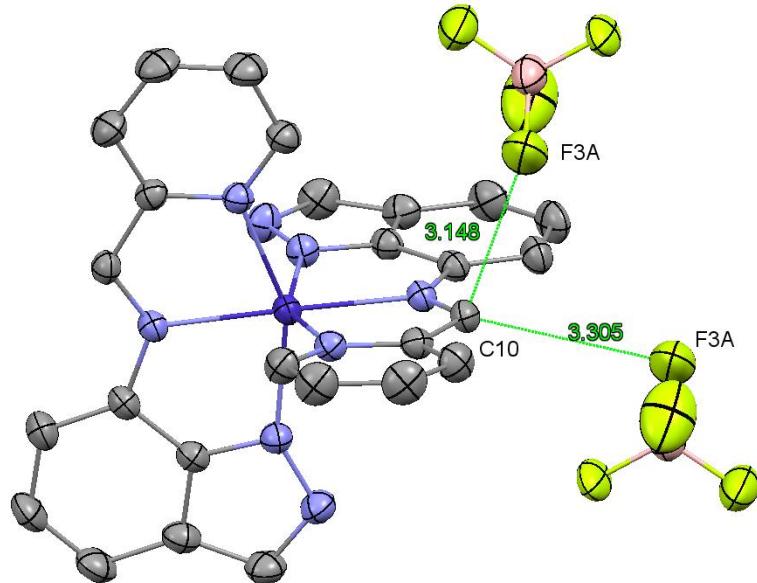


Figure S16. Perspective view of complex 5 showing the H-bond interactions between the imine C-H and the F-BF₃ anions.

Table S4. Hydrogen bonds for complex 5 [Å and °].

D-H...A	d(D-H)	d(H...A)	d(D...A)	<(DHA)
N(2)-H(2)...O(1)	0.88	1.88	2.751(4)	168.3
C(3)-H(3)...F(4A ^a)	0.95	2.53	3.277(9)	136.1
C(15)-H(15)...F(1) ^{#2}	0.95	2.52	3.225(4)	131.4
C(12)-H(12)...F(4A ^a) ^{#3}	0.95	2.57	3.440(10)	151.8
C(10)-H(10)...F(3A ^a) ^{#3}	0.95	2.40	3.305(5)	158.3
C(20)-H(1B)...F(4B ^b)	0.98	2.63	3.58(3)	163.1
C(20)-H(1C)...F(3B ^b) ^{#4}	0.98	2.64	3.152(18)	112.5
O(1)-H(1)...F(2A ^a) ^{#4}	0.953(10)	2.08(8)	2.796(5)	131(8)
O(1)-H(1)...F(3B ^b) ^{#4}	0.953(10)	2.19(8)	2.89(2)	129(8)
O(1)-H(1)...F(4B ^b) ^{#4}	0.953(10)	2.61(8)	3.30(3)	130(8)

Symmetry transformations used to generate equivalent atoms:

#1 -x+1,y,-z+1/2 #2 x,-y+1,z-1/2 #3 x+1/2,y-1/2,z

#4 -x+1/2,y-1/2,-z+1/2

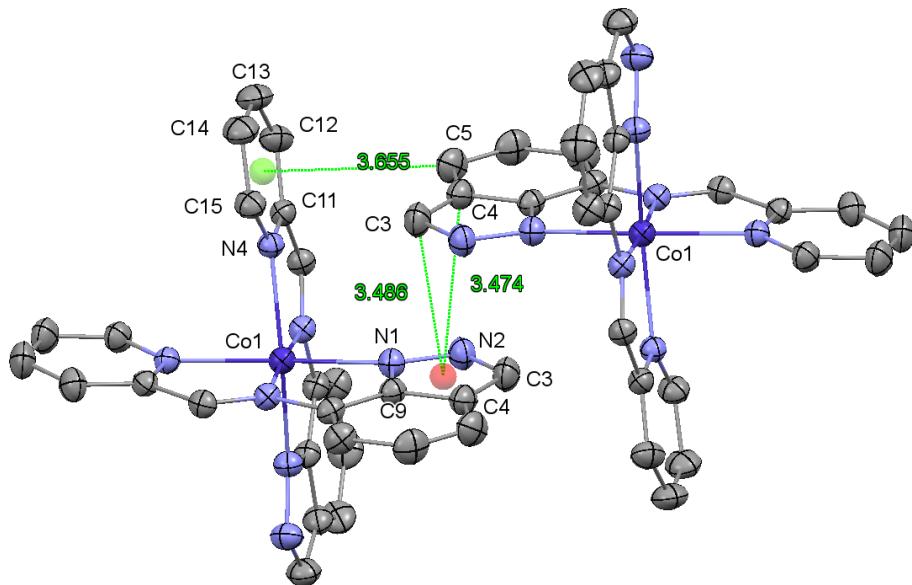


Figure S17. Perspective view of the dimers formed via π - π and C-H... π interactions observed in the crystal lattice of complex 5.