

Supplementary

Table S1: The values of the Ramsey terms that define all the metal-ligand spin spin coupling of the complexes from two different systems treated with DGDZVP(Ru)|6-31+G(d,p) and 3-21G respectively

DGDZVP(Ru) 6-31+G(d,p)						3-21G					
Complex 1						Complex 1					
	FC	SD	PSO	DSO	J (HZ)		FC	SD	PSO	DSO	J (HZ)
Ru-C9	-	-	-	-	-	Ru-C9	-	-	-	-	-
	7.06E+00		1.71E+00	-3.20E-	8.44E+00		2.15E+00		1.74E+00	-3.21E-	3.74E+00
	0	3.62E-001	0	002	0		0	1.81E-001	0	002	0
Ru-C10	-	-	-	-	-	Ru-C10	-	-	-	-	-
	7.83E+00	-3.13E-	1.67E+00	-3.37E-	9.85E+00		2.32E+00	-2.63E-	1.65E+00	-3.31E-	4.26E+00
	0	001	0	002	0		0	001	0	002	0
Ru-C13	-	-	-	-	-	Ru-C13	-	-	-	-	-
	7.44E+00	-2.17E-	1.53E+00	-3.37E-	9.22E+00		2.14E+00	-1.73E-	1.60E+00	-3.32E-	3.95E+00
	0	001	0	002	0		0	001	0	002	0
Ru-C16	-	-	-	-	-	Ru-C16	-	-	-	-	-
	9.25E+00		1.38E+00	-3.31E-	1.04E+00		2.76E+00		1.31E+00	-3.30E-	4.01E+00
	0	2.82E-001	0	002	1		0	9.19E-002	0	002	0
Ru-C18	-	-	-	-	-	Ru-C18	-	-	-	-	-
	8.82E+00		1.74E+00	-3.29E-	1.03E+00		2.54E+00		1.69E+00	-3.26E-	4.19E+00
	0	2.44E-001	0	002	1		0	7.33E-002	0	002	0
Ru-C19	-	-	-	-	-	Ru-C19	-	-	-	-	-
	8.16E+00	-3.04E-	1.04E+00	-3.34E-	9.53E+00		2.55E+00	-2.52E-	1.12E+00	-3.31E-	3.96E+00
	0	001	0	002	0		0	001	0	002	0
Ru-Cl22	-	-	-	-	-	Ru-Cl22	-	-	-	-	-
	1.57E+00	2.33E+00	2.32E+00	-9.87E-	2.71E+00		1.11E+00	2.49E+00	2.20E+00	-9.88E-	2.56E+00
	0	0	1	003	1		0	0	1	003	1
Ru-N12	-	-	-	-	-	Ru-N12	-	-	-	-	-
	1.24E+00	-1.01E-	-7.08E-	-1.46E-	1.32E+00		8.92E+00	-1.07E-	-6.04E-	-1.42E-	9.65E+00
	1	001	001	002	1		0	001	001	002	0
Ru-N17	-	-	-	-	-	Ru-N17	-	-	-	-	-
	1.22E+00	-1.03E-	-6.93E-	-1.46E-	1.30E+00		8.86E+00	-1.04E-	-5.18E-	-1.42E-	9.49E+00
	1	001	001	002	1		0	001	001	002	0
Complex 2						Complex 2					
Ru-C7	-	-	-	-	-	Ru-C7	-	-	-	-	-
	7.31E+00	-3.58E-	1.31E+00	-3.33E-	9.01E+00		2.42E+00	-3.80E-	1.32E+00	-3.31E-	4.16E+00

	0	001	0	002	0		0	001	0	002	0
	-		-		-		-		-		-
Ru-C8	8.36E+00		1.26E+00	-3.21E-	9.21E+00	Ru-C8	2.76E+00		1.26E+00	-3.19E-	3.74E+00
	0	4.45E-001	0	002	0		0	3.22E-001	0	002	0
	-		-		-		-		-		-
Ru-C13	6.18E+00		1.71E+00	-3.08E-	7.47E+00	Ru-C13	2.13E+00		1.73E+00	-3.07E-	3.55E+00
	0	4.40E-001	0	002	0		0	3.39E-001	0	002	0
	-		-		-		-		-		-
Ru-C14	7.30E+00	-4.17E-	-8.50E-	-3.30E-	8.61E+00	Ru-C14	2.11E+00	-4.33E-	-9.09E-	-3.29E-	3.48E+00
	0	001	001	002	0		0	001	001	002	0
	-		-		-		-		-		-
Ru-C18	7.30E+00	-3.58E-	1.31E+00	-3.33E-	9.01E+00	Ru-C18	2.42E+00	-3.80E-	1.33E+00	-3.30E-	4.17E+00
	0	001	0	002	0		0	001	0	002	0
	-		-		-		-		-		-
Ru-C19	8.34E+00		1.26E+00	-3.21E-	9.18E+00	Ru-C19	2.75E+00		1.26E+00	-3.19E-	3.72E+00
	0	4.46E-001	0	002	0		0	3.23E-001	0	002	0
	-	-	-		-		-	-	-		-
Ru-C110	1.17E+00	2.36E+00	2.40E+00	-9.72E-	2.76E+00	Ru-C110	-9.17E-	2.48E+00	2.26E+00	-9.71E-	2.60E+00
	0	0	1	003	1		001	0	1	003	1
	-		-		-		-		-		-
Ru-N11	1.28E+00	-8.72E-	-2.53E-	-1.41E-	1.29E+00	Ru-N11	8.59E+00	-1.01E-		-1.37E-	8.71E+00
	1	002	002	002	1		0	001	1.99E-003	002	0
	-		-		-		-		-		-
Ru-N16	1.28E+00	-8.64E-	-9.98E-	-1.41E-	1.29E+00	Ru-N16	8.59E+00	-1.01E-		-1.37E-	8.69E+00
	1	002	003	002	1		0	001	1.20E-002	002	0
	Complex 3						Complex 3				
	-		-		-		-		-		-
Ru-C13	8.75E+00	-1.31E-	1.30E+00	-3.59E-	1.01E+00	Ru-C13	2.83E+00	-6.54E-	1.31E+00	-3.59E-	4.25E+00
	0	002	0	002	1		0	002	0	002	0
	-		-		-		-		-		-
Ru-C14	7.71E+00	-7.27E-	1.84E+00	-3.59E-	9.66E+00	Ru-C14	2.42E+00	-1.44E-	1.79E+00	-3.58E-	4.39E+00
	0	002	0	002	0		0	001	0	002	0
	-		-		-		-		-		-
Ru-C18	8.63E+00	-5.35E-	1.29E+00	-3.58E-	1.00E+00	Ru-C18	2.79E+00	-1.41E-	1.33E+00	-3.58E-	4.29E+00
	0	002	0	002	1		0	001	0	002	0
	-		-		-		-		-		-
Ru-C21	7.02E+00	-2.37E-	1.63E+00	-3.47E-	8.68E+00	Ru-C21	2.02E+00	-5.09E-	1.68E+00	-3.45E-	3.79E+00
	0	003	0	002	0		0	002	0	002	0
	-		-		-		-		-		-
Ru-C23	7.68E+00	-2.66E-	1.86E+00	-3.57E-	9.60E+00	Ru-C23	2.40E+00	-7.16E-	1.88E+00	-3.55E-	4.38E+00
	0	002	0	002	0		0	002	0	002	0

	-	-	-	-	-		-	-	-	-	-
	7.09E+00	-4.84E-	1.62E+00	-3.48E-	8.80E+00		2.13E+00	-1.25E-	1.70E+00	-3.47E-	3.98E+00
Ru-C25	0	002	0	002	0	Ru-C25	0	001	0	002	0
	-	-	-	-	-		-	-	-	-	-
	2.06E+00	2.04E+00	2.29E+00	-1.11E-	2.70E+00		1.45E+00	2.17E+00	2.19E+00	-1.11E-	2.55E+00
Ru-C17	0	0	1	002	1	Ru-C17	0	0	1	002	1
	-	-	-	-	-		-	-	-	-	-
	1.15E+00	-5.83E-		-1.49E-	1.13E+00		8.07E+00	-6.77E-		-1.45E-	7.83E+00
Ru-N10	1	002	3.27E-001	002	1	Ru-N10	0	002	3.26E-001	002	0
	-	-	-	-	-		-	-	-	-	-
	1.15E+00	-6.05E-		-1.49E-	1.13E+00		8.01E+00	-7.72E-		-1.46E-	7.80E+00
Ru-N16	1	002	3.14E-001	002	1	Ru-N16	0	002	2.97E-001	002	0
	Complex 4						Complex 4				
	-	-	-	-	-		-	-	-	-	-
	7.36E+00	-2.69E-	1.17E+00	-3.17E-	8.56E+00		2.45E+00	-6.12E-	1.30E+00	-3.17E-	3.84E+00
Ru-C2	0	003	0	002	0	Ru-C2	0	002	0	002	0
	-	-	-	-	-		-	-	-	-	-
	6.27E+00	-1.82E-	1.65E+00	-3.24E-	8.13E+00		1.53E+00	-2.32E-	1.69E+00	-3.24E-	3.48E+00
Ru-C3	0	001	0	002	0	Ru-C3	0	001	0	002	0
	-	-	-	-	-		-	-	-	-	-
	7.37E+00		1.16E+00	-3.17E-	8.56E+00		2.46E+00	-5.27E-	1.30E+00	-3.17E-	3.84E+00
Ru-C4	0	6.30E-003	0	002	0	Ru-C4	0	002	0	002	0
	-	-	-	-	-		-	-	-	-	-
	7.96E+00	-7.38E-	2.60E+00	-3.23E-	1.07E+00		2.91E+00	-1.26E-	2.50E+00	-3.22E-	5.57E+00
Ru-C5	0	002	0	002	1	Ru-C5	0	001	0	002	0
	-	-	-	-	-		-	-	-	-	-
	6.26E+00	-1.90E-	1.64E+00	-3.24E-	8.13E+00		1.53E+00	-2.40E-	1.68E+00	-3.24E-	3.48E+00
Ru-C6	0	001	0	002	0	Ru-C6	0	001	0	002	0
	-	-	-	-	-		-	-	-	-	-
	7.95E+00	-6.55E-	2.60E+00	-3.23E-	1.06E+00		2.90E+00	-1.19E-	2.51E+00	-3.22E-	5.55E+00
Ru-C7	0	002	0	002	1	Ru-C7	0	001	0	002	0
	-	-	-	-	-		-	-	-	-	-
	1.28E+00		-7.79E-	-1.33E-	1.35E+00		8.84E+00		-6.00E-	-1.30E-	9.43E+00
Ru-N10	1	3.65E-002	001	002	1	Ru-N10	0	2.47E-002	001	002	0
	-	-	-	-	-		-	-	-	-	-
	1.28E+00		-7.79E-	-1.33E-	1.35E+00		8.84E+00		-5.99E-	-1.30E-	9.43E+00
Ru-N13	1	3.67E-002	001	002	1	Ru-N13	0	2.50E-002	001	002	0
	-	-	-	-	-		-	-	-	-	-
	1.07E+00	-1.64E-	-6.73E-	-1.47E-	1.16E+00		6.86E+00	-1.59E-	-4.56E-	-1.45E-	7.49E+00
Ru-N15	1	001	001	002	1	Ru-N15	0	001	001	002	0
	Complex 5						Complex 5				

	-	-	-	-	-		-	-	-	-	-
	7.94E+00	-8.67E-	2.14E+00	-3.26E-	1.01E+00		2.81E+00	-6.52E-	2.07E+00	-3.25E-	4.98E+00
Ru-C18	0	003	0	002	1	Ru-C18	0	002	0	002	0
	-	-	-	-	-		-	-	-	-	-
	6.08E+00	-1.52E-	1.42E+00	-3.27E-	7.69E+00		1.43E+00	-1.99E-	1.45E+00	-3.26E-	3.11E+00
Ru-C20	0	001	0	002	0	Ru-C20	0	001	0	002	0
	-	-	-	-	-		-	-	-	-	-
	7.95E+00	-1.74E-	2.14E+00	-3.26E-	1.01E+00		2.82E+00	-7.17E-	2.07E+00	-3.25E-	4.99E+00
Ru-C22	0	002	0	002	1	Ru-C22	0	002	0	002	0
	-	-	-	-	-		-	-	-	-	-
	6.77E+00	-8.44E-	-3.18E-	7.57E+00			2.28E+00	-9.64E-	-3.17E-	3.26E+00	
Ru-C23	0	7.09E-002	001	002	0	Ru-C23	0	1.27E-002	001	002	0
	-	-	-	-	-		-	-	-	-	-
	6.10E+00	-1.46E-	1.43E+00	-3.27E-	7.71E+00		1.43E+00	-1.96E-	1.45E+00	-3.26E-	3.11E+00
Ru-C26	0	001	0	002	0	Ru-C26	0	001	0	002	0
	-	-	-	-	-		-	-	-	-	-
	6.75E+00	-8.46E-	-3.18E-	7.57E+00			2.27E+00	-9.66E-	-3.17E-	3.26E+00	
Ru-C27	0	6.16E-002	001	002	0	Ru-C27	0	6.47E-003	001	002	0
	-	-	-	-	-		-	-	-	-	-
	1.27E+00	-2.15E-	-1.33E-	1.29E+00			8.69E+00	-9.55E-	-1.30E-	8.80E+00	
Ru-N14	1	1.90E-002	001	002	1	Ru-N14	0	5.82E-003	002	002	0
	-	-	-	-	-		-	-	-	-	-
	1.24E+00	-1.60E-	-3.08E-	-1.50E-	1.29E+00		7.90E+00	-1.68E-	-1.85E-	-1.48E-	8.26E+00
Ru-N15	1	001	001	002	1	Ru-N15	0	001	001	002	0
	-	-	-	-	-		-	-	-	-	-
	1.27E+00	-2.34E-	-1.33E-	1.30E+00			8.71E+00	-1.08E-	-1.30E-	8.82E+00	
Ru-N21	1	1.81E-002	001	002	1	Ru-N21	0	5.23E-003	001	002	0

Table S2: The values of the Ramsey terms that define all the metal-ligand spin spin coupling of the complexes from two different systems treated with ECP(Ru)|6-31+G(d,p) and ECP(Ru,Cl)|6-31G* respectively

Ramsey Terms	ECP(6- 31+G(d,p)	ECP(6- 31G*)	Ramsey Terms
Complex 1		Complex 1	

	FC (HZ) for Ru	SD for Ru	PSO for Ru	DSO for Ru	J (HZ) for Ru		FC (HZ) for Ru	SD for Ru	PSO for Ru	DSO for Ru	J (HZ) for Ru
Ru-C9	-3.67E-006	4.55E-002	-2.43E-001	-3.18E-002	-2.29E-001	Ru-C9	-1.79E-006	3.65E-002	-2.55E-001	-3.00E-002	-2.48E-001
Ru-C10	-6.04E-006	-2.15E-002	-2.38E-001	-3.35E-002	-2.93E-001	Ru-C10	0.00E+000	-1.41E-002	-2.43E-001	-3.06E-002	-2.88E-001
Ru-C13	-6.00E-006	-7.58E-003	-2.17E-001	-3.35E-002	-2.58E-001	Ru-C13	-2.90E-006	8.14E-004	-2.34E-001	-3.09E-002	-2.64E-001
Ru-C16	-3.66E-006	4.04E-002	-2.09E-001	-3.29E-002	-2.01E-001	Ru-C16	0.00E+000	3.01E-002	-2.00E-001	-2.98E-002	-2.00E-001
Ru-C18	-3.00E-006	3.33E-002	-2.52E-001	-3.27E-002	-2.52E-001	Ru-C18	1.33E-006	2.31E-002	-2.51E-001	-2.99E-002	-2.58E-001
Ru-C19	-6.44E-006	-1.89E-002	-1.57E-001	-3.32E-002	-2.09E-001	Ru-C19	0.00E+000	-9.33E-003	-1.69E-001	-2.98E-002	-2.08E-001
Ru-Cl22	0.00E+000	-1.71E-001	3.17E+000	-9.93E-003	3.35E+000	Ru-Cl22	0.00E+000	-1.17E-002	-2.17E-001	-9.88E-003	-2.38E-001
Ru-N12	-3.55E-006	-3.63E-003	-9.07E-002	-1.46E-002	-1.09E-001	Ru-N12	1.35E-006	-4.11E-003	-1.09E-001	-1.37E-002	-1.27E-001
Ru-N17	-3.49E-006	-3.73E-003	-8.93E-002	-1.46E-002	-1.08E-001	Ru-N17	1.18E-006	-3.82E-003	-9.36E-002	-1.37E-002	-1.11E-001
Complex 2						Complex 2					
Ru-C7	-8.10E-006	-2.42E-002	-1.86E-001	-3.31E-002	-2.43E-001	Ru-C7	-5.02E-006	-2.49E-002	-1.95E-001	-3.06E-002	-2.51E-001
Ru-C8	-7.05E-006	5.42E-002	-1.89E-001	-3.18E-002	-1.67E-001	Ru-C8	0.00E+000	5.35E-002	-1.94E-001	-2.89E-002	-1.69E-001
Ru-C13	-6.10E-006	5.10E-002	-2.39E-001	-3.06E-002	-2.19E-001	Ru-C13	-5.93E-006	5.09E-002	-2.56E-001	-2.86E-002	-2.34E-001
Ru-C14	-4.52E-006	-2.90E-002	-1.27E-001	-3.28E-002	-1.89E-001	Ru-C14	-1.35E-006	-2.96E-002	-1.36E-001	-2.95E-002	-1.96E-001
Ru-C18	-8.10E-006	-2.42E-002	-1.87E-001	-3.31E-002	-2.44E-001	Ru-C18	-5.03E-006	-2.49E-002	-1.96E-001	-3.06E-002	-2.51E-001
Ru-C19	-7.04E-006	5.42E-002	-1.88E-001	-3.18E-002	-1.66E-001	Ru-C19	0.00E+000	5.36E-002	-1.93E-001	-2.89E-002	-1.69E-001
Ru-Cl10	0.00E+000	-1.69E-001	3.28E+000	-9.78E-003	3.46E+000	Ru-Cl10	0.00E+000	-1.10E-002	-2.25E-001	-9.74E-003	-2.46E-001
Ru-N11	-3.33E-006	-3.49E-003	8.27E-003	-1.41E-002	-9.33E-001	Ru-N11	0.00E+000	-3.67E-003	4.69E-004	-1.30E-002	-1.62E-001

	006	003		002	003		0	003		002	002
Ru-N16	-3.28E-006	-3.57E-003	1.05E-002	-1.41E-002	-7.21E-003	Ru-N16	0.00E+000	-3.73E-003	2.52E-003	-1.30E-002	-1.42E-002
	Complex 3						Complex 3				
Ru-C13	-6.29E-006	8.47E-003	-1.86E-001	-3.57E-002	-2.13E-001	Ru-C13	0.00E+000	-1.87E-001	-3.27E-002	-2.10E-001	
Ru-C14	-7.82E-006	2.83E-003	-2.47E-001	-3.56E-002	-2.79E-001	Ru-C14	-1.39E-006	-2.42E-001	-3.31E-002	-2.74E-001	
Ru-C18	-6.23E-006	4.71E-003	-1.84E-001	-3.56E-002	-2.15E-001	Ru-C18	1.32E-006	1.50E-003	-1.89E-001	-3.26E-002	-2.20E-001
Ru-C21	-7.37E-006	1.06E-002	-2.25E-001	-3.45E-002	-2.49E-001	Ru-C21	-2.65E-006	-2.40E-001	-3.24E-002	-2.61E-001	
Ru-C23	-7.69E-006	7.47E-003	-2.49E-001	-3.55E-002	-2.77E-001	Ru-C23	-1.47E-006	-2.54E-001	-3.29E-002	-2.78E-001	
Ru-C25	-7.51E-006	6.02E-003	-2.24E-001	-3.46E-002	-2.53E-001	Ru-C25	-2.80E-006	-2.41E-001	-3.25E-002	-2.71E-001	
Ru-Cl7	0.00E+000	-1.24E-001	3.06E+000	-1.11E-002	3.20E+000	Ru-Cl7	0.00E+000	-7.99E-003	-2.11E-001	-1.11E-002	-2.30E-001
Ru-N10	-3.25E-006	-2.18E-003	6.56E-002	-1.49E-002	4.85E-002	Ru-N10	-1.10E-006	-1.94E-003	-1.39E-002	4.50E-002	
Ru-N16	-3.25E-006	-2.24E-003	6.39E-002	-1.49E-002	4.67E-002	Ru-N16	-1.02E-006	-2.53E-003	-1.39E-002	4.11E-002	
	Complex 4						Complex 4				
Ru-C2	-3.54E-006	8.88E-003	-1.74E-001	-3.14E-002	-1.97E-001	Ru-C2	5.22E-006	8.93E-003	-1.80E-001	-3.13E-002	-2.03E-001
Ru-C3	-8.33E-006	-8.29E-003	-2.29E-001	-3.22E-002	-2.70E-001	Ru-C3	-2.18E-006	-8.75E-003	-2.34E-001	-3.21E-002	-2.74E-001
Ru-C4	-3.54E-006	9.73E-003	-1.74E-001	-3.14E-002	-1.96E-001	Ru-C4	5.25E-006	9.81E-003	-1.80E-001	-3.13E-002	-2.02E-001
Ru-C5	-7.34E-006	1.34E-004	-3.43E-001	-3.22E-002	-3.75E-001	Ru-C5	0.00E+000	-5.09E-004	-3.47E-001	-3.21E-002	-3.80E-001
Ru-C6	-8.31E-006	-9.04E-003	-2.28E-001	-3.22E-002	-2.69E-001	Ru-C6	-2.17E-006	-9.54E-003	-2.32E-001	-3.21E-002	-2.74E-001
Ru-C7	-7.33E-006	9.98E-004	-3.44E-001	-3.22E-002	-3.75E-001	Ru-C7	0.00E+000	-3.48E-001	-3.21E-002	-3.79E-001	
Ru-N10	-3.37E-006	1.11E-002	-1.07E-001	-1.34E-002	-1.09E-002	Ru-N10	0.00E+000	1.10E-002	-1.10E-001	-1.33E-002	-1.13E-002

	006		001	002	001	0		001	002	001
Ru-N13	-3.37E-006	1.11E-002	-1.07E-001	-1.34E-002	-1.09E-001	0.00E+00	0	-1.10E-001	-1.33E-002	-1.13E-001
Ru-N15	-2.89E-006	-1.15E-002	-6.88E-002	-1.47E-002	-9.51E-002	0.00E+00	0	-1.17E-002	-7.21E-002	-1.47E-002
	Complex 5					Complex 5				
Ru-C18	-6.68E-006	7.27E-003	-2.86E-001	-3.25E-002	-3.11E-001	0.00E+00	0	-2.91E-001	-3.24E-002	-3.17E-001
Ru-C20	-7.65E-006	-5.87E-003	-2.02E-001	-3.24E-002	-2.40E-001	0.00E+00	0	-6.18E-003	-2.07E-001	-3.23E-002
Ru-C22	-6.70E-006	6.32E-003	-2.86E-001	-3.25E-002	-3.12E-001	0.00E+00	0	-2.91E-001	-3.24E-002	-3.17E-001
Ru-C23	-2.90E-006	1.57E-002	-1.33E-001	-3.14E-002	-1.49E-001	4.56E-006	1.59E-002	-1.40E-001	-3.13E-002	-1.55E-001
Ru-C26	-7.62E-006	-5.40E-003	-2.02E-001	-3.24E-002	-2.40E-001	0.00E+00	0	-5.95E-003	-2.07E-001	-3.23E-002
Ru-C27	-2.93E-006	1.49E-002	-1.33E-001	-3.14E-002	-1.50E-001	4.62E-006	1.54E-002	-1.40E-001	-3.13E-002	-1.56E-001
Ru-N14	-3.40E-006	1.07E-002	-1.97E-002	-1.33E-002	-2.23E-002	-1.06E-006	1.08E-002	-2.29E-002	-1.33E-002	-2.54E-002
Ru-N15	-3.65E-006	-1.37E-002	-1.20E-002	-1.51E-002	-4.08E-002	0.00E+00	0	-1.41E-002	-1.94E-002	-1.50E-002
Ru-N21	-3.39E-006	1.07E-002	-2.23E-002	-1.33E-002	-2.50E-002	-1.08E-006	1.08E-002	-2.54E-002	-1.33E-002	-2.79E-002

Table S3: The atomic NMR magnetazability and shielding for the selected atoms in the complexes using combined basis set DGDZVP(Ru)|6-31+G(d,p)

Complex 1															
q(A)	Vol(A),		DI(A,A'		$\chi_{\text{intra_Is}}$		$\chi_{\text{bond_Is}}$		$\chi_{\text{intra_Is}}$		$\sigma_{\text{Iso}}(A,$		$\sigma_{\text{Iso}}(A',$		Isotrop
	0.001	L(A)	K(A)	LI(A))/2	o	o	o(A)	Mu(A)	A)	A)	A)	$\sigma_{\text{Iso}}(A)$	ic	opy
Ru1	9.50E-01	1.01E+02	6.36E-04	4.44E+03	4.05E+01	2.54E+00	2.24E+01	2.27E+01	-3.35E-01	1.20E+00	3.51E+03	2.52E+01	3.49E+03	3.49E+03	6.16E+03

		-	-	-											
C9	-6.26E-02	7.38E+01	-5.40E-05	3.78E+01	3.97E+00	2.09E+00	3.09E+00	4.98E+00	8.07E+00	9.26E-02	8.50E+01	2.29E+01	1.08E+02	1.08E+02	1.28E+02
		-	-	-											
C10	-4.68E-02	7.14E+01	-3.51E-04	3.78E+01	3.96E+00	2.09E+00	2.65E+00	9.19E+00	1.18E+01	6.08E-02	7.30E+01	2.35E+01	9.65E+01	9.65E+01	1.35E+02
		-	-	-											
C13	-4.96E-02	7.16E+01	7.50E-05	3.78E+01	3.96E+00	2.09E+00	2.68E+00	9.21E+00	1.19E+01	1.02E-01	7.55E+01	2.33E+01	9.88E+01	9.88E+01	1.31E+02
		-	-	-											
C16	-5.13E-02	7.37E+01	-1.58E-04	3.78E+01	3.97E+00	2.08E+00	3.09E+00	4.90E+00	7.99E+00	5.52E-02	8.44E+01	2.30E+01	1.07E+02	1.07E+02	1.32E+02
		-	-	-											
C18	-5.82E-02	7.48E+01	-1.32E-04	3.78E+01	3.98E+00	2.08E+00	3.22E+00	4.95E+00	8.17E+00	6.78E-02	8.75E+01	2.35E+01	1.11E+02	1.11E+02	1.27E+02
		-	-	-											
C19	-2.85E-02	6.99E+01	6.90E-05	3.78E+01	3.95E+00	2.08E+00	2.34E+00	9.02E+00	1.14E+01	1.02E-01	6.40E+01	2.25E+01	8.65E+01	8.65E+01	1.51E+02
		-	-	-											
Cl22	-5.27E-01	2.32E+02	8.30E-05	4.59E+02	1.69E+01	6.46E+01	2.15E+01	3.69E+00	2.52E+01	1.29E+00	1.06E+03	4.24E+00	1.06E+03	1.06E+03	5.20E+02
		-	-	-											
N12	-7.59E-01	7.81E+01	2.16E-04	5.48E+01	5.90E+00	1.86E+00	3.51E+00	6.63E+00	1.01E+01	6.58E-01	3.46E+00	1.70E+01	1.35E+01	1.35E+01	1.92E+02
		-	-	-											
N17	-7.59E-01	7.80E+01	2.41E-04	5.48E+01	5.90E+00	1.86E+00	3.50E+00	6.64E+00	1.01E+01	6.17E-01	3.94E+00	1.69E+01	1.29E+01	1.29E+01	1.93E+02
		-	-	-											
O2	1.17E+00	1.30E+02	8.20E-05	7.54E+01	8.23E+00	9.47E-01	6.12E+00	3.48E+00	9.60E+00	1.03E+00	9.64E+01	2.41E+00	9.40E+01	9.40E+01	5.67E+02
		-	-	-											
H24	1.23E-01	4.05E+01	2.40E-05	5.68E-01	3.45E-01	5.32E-01	-9.18E-01	1.36E+00	2.28E+00	4.80E-01	1.77E+01	5.68E+00	2.33E+01	2.33E+01	6.65E+00
		-	-	-											
H29	1.23E-01	4.06E+01	2.60E-05	5.68E-01	3.45E-01	5.32E-01	-9.20E-01	1.36E+00	2.29E+00	4.92E-01	1.77E+01	5.67E+00	2.33E+01	2.33E+01	6.63E+00
Complex 2															
		-	-	-											
Ru1	9.59E-01	1.04E+02	2.47E-04	4.44E+03	4.05E+01	2.57E+00	2.00E+01	2.28E+01	2.79E+00	1.18E+00	3.12E+03	2.45E+01	3.09E+03	3.09E+03	5.61E+03
		-	-	-											
C7	-5.32E-01	7.01E+01	-2.27E-01	3.78E+01	3.96E+00	2.09E+00	2.53E+00	9.07E+00	1.16E+01	6.14E-01	6.95E+01	2.28E+01	9.23E+01	9.23E+01	1.38E+02

	02	01	04	01	00	00	00	00	01	02	01	01	01	01	02
							-	-	-						
C8	-6.29E-02	7.46E+01	9.20E-05	3.78E+01	3.98E+00	2.08E+00	3.26E+00	4.86E+00	8.12E+00	8.23E-02	8.88E+01	2.26E+01	1.11E+02	1.11E+02	1.30E+02
							-	-	-						
C13	-5.29E-02	7.64E+01	-6.00E-06	3.78E+01	3.97E+00	2.08E+00	3.20E+00	4.90E+00	8.10E+00	7.84E-02	8.70E+01	2.27E+01	1.10E+02	1.10E+02	1.30E+02
							-	-	-						
C14	-2.67E-02	6.96E+01	-5.71E-04	3.78E+01	3.95E+00	2.08E+00	2.13E+00	8.85E+00	1.10E+01	9.18E-02	5.83E+01	2.19E+01	8.02E+01	8.02E+01	1.57E+02
							-	-	-						
C18	-5.42E-02	7.02E+01	-8.17E-04	3.78E+01	3.96E+00	2.09E+00	2.53E+00	9.07E+00	1.16E+01	9.29E-02	6.96E+01	2.29E+01	9.24E+01	9.24E+01	1.37E+02
							-	-	-						
C19	-6.30E-02	7.46E+01	9.00E-05	3.78E+01	3.98E+00	2.08E+00	3.26E+00	4.86E+00	8.12E+00	7.40E-02	8.89E+01	2.26E+01	1.11E+02	1.11E+02	1.30E+02
							-	-	-						
Cl10	-5.21E-01	2.37E+02	8.60E-05	4.59E+02	1.69E+01	6.23E+01	2.20E+01	3.76E+00	2.57E+01	1.26E+00	1.10E+03	4.72E+00	1.10E+03	1.10E+03	4.20E+02
							-	-	-						
N11	1.22E+00	8.21E+01	8.40E-05	5.51E+01	6.36E+00	1.86E+00	3.55E+00	7.30E+00	1.09E+01	5.42E-01	3.05E+01	1.29E+01	1.76E+01	1.76E+01	3.87E+02
							-	-	-						
N16	1.22E+00	8.21E+01	1.68E-04	5.51E+01	6.36E+00	1.86E+00	3.54E+00	7.32E+00	1.09E+01	5.10E-01	3.11E+01	1.29E+01	1.82E+01	1.82E+01	3.89E+02
							-	-	-						
H34	8.99E-02	4.09E+01	8.10E-05	5.87E-01	3.64E-01	5.46E-01	-9.14E-01	1.27E+00	2.19E+00	6.96E-02	1.82E+01	4.25E+00	2.25E+01	2.25E+01	1.07E+01
							-	-	-						
H38	9.65E-02	4.10E+01	5.80E-05	5.81E-01	3.59E-01	5.45E-01	-9.15E-01	1.28E+00	2.19E+00	5.34E-02	1.81E+01	4.22E+00	2.23E+01	2.23E+01	1.01E+01
Complex 3															
							-	-	-						
Ru1	9.55E-01	9.99E+01	3.57E-04	4.44E+03	4.05E+01	2.52E+00	2.45E+01	2.56E+01	1.05E+00	8.40E-01	3.76E+03	2.39E+01	3.73E+03	3.73E+03	6.00E+03
							-	-	-						
C13	-4.01E-02	7.06E+01	-1.19E-04	3.78E+01	3.96E+00	2.08E+00	2.72E+00	6.38E+00	9.11E+00	8.08E-02	7.46E+01	2.26E+01	9.72E+01	9.72E+01	1.39E+02
							-	-	-						
C14	-4.85E-02	6.98E+01	-1.15E-04	3.79E+01	3.97E+00	2.08E+00	2.69E+00	6.81E+00	9.50E+00	4.00E-01	7.44E+01	2.31E+01	9.75E+01	9.75E+01	1.39E+02

		-	-	-											
C18	-3.97E-02	7.07E+01	-9.40E-05	3.78E+01	3.96E+00	2.08E+00	2.69E+00	6.38E+00	9.06E+00	8.56E-02	7.35E+01	2.26E+01	9.61E+01	9.61E+01	1.40E+02
C21	-5.97E-02	7.36E+01	-6.60E-05	3.78E+01	3.97E+00	2.09E+00	2.89E+00	6.48E+00	9.37E+00	1.32E-01	7.88E+01	2.31E+01	1.02E+02	1.02E+02	1.34E+02
C23	-5.00E-02	7.02E+01	-1.81E-04	3.79E+01	3.97E+00	2.08E+00	2.73E+00	6.81E+00	9.53E+00	4.12E-01	7.53E+01	2.31E+01	9.84E+01	9.84E+01	1.39E+02
C25	-5.93E-02	7.33E+01	-2.62E-04	3.78E+01	3.97E+00	2.09E+00	2.87E+00	6.46E+00	9.33E+00	1.40E-01	7.81E+01	2.32E+01	1.01E+02	1.01E+02	1.34E+02
Cl7	-5.33E-01	2.36E+02	1.08E-04	4.59E+02	1.69E+01	6.31E+01	2.18E+01	3.57E+00	2.54E+01	1.30E+00	1.08E+03	4.16E+00	1.09E+03	1.09E+03	5.79E+02
N10	1.20E+00	8.30E+01	2.90E-04	5.50E+01	6.36E+00	1.84E+00	3.65E+00	7.31E+00	1.10E+01	9.47E-01	2.31E+01	1.21E+01	1.11E+01	1.11E+01	3.83E+02
N16	1.20E+00	8.30E+01	8.80E-05	5.50E+01	6.36E+00	1.84E+00	3.65E+00	7.31E+00	1.10E+01	9.12E-01	2.24E+01	1.21E+01	1.03E+01	1.03E+01	3.81E+02
O2	1.19E+00	1.28E+02	1.16E-04	7.54E+01	8.22E+00	9.69E+01	6.05E+00	2.15E+00	8.20E+00	9.26E-01	8.06E+01	2.34E+00	7.83E+01	7.83E+01	5.28E+02
O28	1.19E+00	1.28E+02	1.27E-04	7.54E+01	8.23E+00	9.67E+01	6.07E+00	2.15E+00	8.22E+00	9.10E-01	8.12E+01	2.31E+00	7.89E+01	7.89E+01	5.26E+02
H31	1.12E-01	3.92E+01	-2.90E-05	5.80E-01	3.48E-01	5.39E-01	-9.26E-01	1.24E+00	2.16E+00	3.59E-01	1.82E+01	6.49E+00	2.47E+01	2.47E+01	6.46E+00
H36	1.17E-01	3.87E+01	2.90E-05	5.79E-01	3.45E-01	5.39E-01	-9.12E-01	1.23E+00	2.15E+00	3.62E-01	1.81E+01	6.49E+00	2.46E+01	2.46E+01	6.83E+00
Complex 4															
Ru1	9.92E-01	1.10E+02	-9.65E-04	4.43E+03	4.06E+01	2.46E+00	2.65E+01	2.59E+01	5.96E-01	3.99E-01	4.14E+03	2.40E+01	4.12E+03	4.12E+03	6.61E+03
C2	-4.55E-02	7.33E+01	1.09E-04	3.78E+01	3.97E+00	2.08E+00	2.80E+00	6.45E+00	9.25E+00	6.59E-02	7.44E+01	2.23E+01	9.67E+01	9.67E+01	1.37E+02
C3	-4.44E-01	7.07E+01	-1.18E-01	3.78E+01	3.96E+00	2.08E+00	2.70E+00	6.37E+00	9.07E+00	6.64E-01	7.52E+01	2.36E+01	9.87E+01	9.87E+01	1.40E+02

		-	-	-											
	-4.90E-	7.29E+	-1.03E-	3.78E+	3.97E+	2.08E+	2.71E+	3.64E+	6.35E+	1.07E-	7.11E+	2.16E+	9.26E+	9.26E+	1.43E+
C27	02	01	04	01	00	00	00	00	00	01	01	01	01	01	02
	-														
	1.23E+	8.58E+	2.21E-	5.50E+	6.40E+	1.83E+	3.88E+	7.35E+	1.12E+	4.62E-	1.31E+	1.19E+	1.24E+	1.24E+	3.37E+
N14	00	01	04	01	00	00	00	00	01	01	01	01	00	00	02
	-														
	1.23E+	7.90E+	4.56E-	5.51E+	6.35E+	1.88E+	3.71E+	7.16E+	1.09E+	2.32E-	1.01E+	1.22E+	2.12E+	2.12E+	3.05E+
N15	00	01	04	01	00	00	00	00	01	01	01	01	00	00	02
	-														
	1.23E+	8.58E+	2.75E-	5.50E+	6.40E+	1.83E+	3.89E+	7.37E+	1.13E+	3.70E-	1.23E+	1.19E+	-4.32E-	-4.31E-	3.35E+
N21	00	01	04	01	00	00	00	00	01	01	01	01	01	01	02

q(A) is total charge on atom A, Vol(A) is the volume bounded by an isosurface of the electron density distribution (0.001) and by interatomic surfaces of atom A; K(A) is electronic kinetic energy of atom A (Hamiltonian Form), Loc(A) is number of electrons localized in atom A, K_Scaled(A) is approximation to virial-based total energy of atom A, DI(A,A') is number of electron delocalization from atom A, $\chi_{\text{intra_iso}}$ is intraatomic magnetizability contribution of atom A (cgs-ppm); $\chi_{\text{bond_iso}}$ is bonding magnetizability contribution of atom A; χ_{iso} is total magnetizability contribution of atom A (i.e. magnetizability isotropic); Mu(A) is total dipole moment contribution of atom A; $\sigma_{\text{iso}}(\text{A},\text{A})$ is the intraatomic shielding tensor; $\sigma_{\text{iso}}(\text{A}',\text{A})$ is total induced magnetic field on nucleus a by other nuclei A'; $\sigma_{\text{iso}}(\text{A})$ is total magnetic shielding tensor of atom A (i.e. nmr shielding isotropic).

Table S4: The atomic NMR magnetazability and shielding for the selected atoms in the complexes using combined basis sets 3-21G

Complex 1															
	Vol(A),														
	q(A)	0.001	L(A)	K(A)	LI(A)	DI(A,A')/2	$\chi_{\text{intra_iso}}$ o(A)	$\chi_{\text{bond_iso}}$ o(A)	χ_{iso} (A)	Mu(A)	$\sigma_{\text{iso}}(\text{A},\text{A})$	$\sigma_{\text{iso}}(\text{A}',\text{A})$ $\sigma_{\text{iso}}(\text{A})$	Isotrop ic	Anisotr opy	
								-			-		-		
	8.63E-	9.73E+	5.52E-	4.40E+	4.07E+	2.46E+	2.47E+	2.50E+	-2.65E-	1.19E+	3.10E+	2.41E+	3.07E+	3.07E+	6.03E+
Ru1	01	01	04	03	01	00	01	01	01	00	03	01	03	03	03
								-							
	-1.14E-	7.66E+	-7.00E-	3.76E+	4.04E+	2.08E+	3.64E+	6.30E+	9.95E+	1.35E-	1.04E+	2.27E+	1.26E+	1.26E+	1.19E+
C9	01	01	05	01	00	00	00	00	00	01	02	01	02	02	02
								-							
	-1.04E-	7.49E+	-1.97E-	3.76E+	4.03E+	2.07E+	3.35E+	6.23E+	9.58E+	9.47E-	9.56E+	2.30E+	1.19E+	1.19E+	1.24E+
C10	01	01	04	01	00	00	00	00	00	02	01	01	02	02	02
								-							
	-1.09E-	7.52E+	-1.11E-	3.76E+	4.03E+	2.08E+	3.37E+	6.24E+	9.61E+	1.16E-	9.77E+	2.28E+	1.20E+	1.20E+	1.21E+
C13	01	01	04	01	00	00	00	00	00	01	01	01	02	02	02

		-	-	-											
C16	-9.09E-02	7.57E+01	-5.60E-05	3.76E+01	4.02E+00	2.07E+00	3.52E+00	6.13E+00	9.66E+00	9.32E-02	1.00E+02	2.24E+01	1.23E+02	1.23E+02	1.26E+02
C18	-1.04E-01	7.71E+01	-4.80E-05	3.76E+01	4.04E+00	2.07E+00	3.71E+00	6.23E+00	9.94E+00	1.14E-01	1.04E+02	2.30E+01	1.27E+02	1.27E+02	1.21E+02
C19	-8.40E-02	7.33E+01	-1.10E-04	3.76E+01	4.02E+00	2.07E+00	3.05E+00	6.06E+00	9.11E+00	9.99E-02	8.86E+01	2.21E+01	1.11E+02	1.11E+02	1.35E+02
Cl22	-4.89E-01	2.29E+02	6.20E-05	4.57E+02	1.69E+01	6.10E+01	2.15E+01	3.19E+00	2.47E+01	1.33E+00	9.98E+02	3.76E+00	1.00E+03	1.00E+03	4.24E+02
N12	-6.13E-01	7.39E+01	4.00E-05	5.43E+01	5.75E+00	1.86E+00	3.74E+00	6.36E+00	1.01E+01	6.53E-01	2.36E+01	1.74E+01	4.10E+01	4.10E+01	1.58E+02
N17	-6.03E-01	7.39E+01	1.41E-04	5.43E+01	5.74E+00	1.86E+00	3.68E+00	6.52E+00	1.02E+01	8.83E-01	1.93E+01	1.74E+01	3.66E+01	3.66E+01	1.68E+02
O2	-8.53E-01	1.04E+02	8.40E-05	7.47E+01	7.78E+00	1.07E+00	5.00E+00	2.28E+00	7.27E+00	9.06E-01	9.86E+01	4.65E+00	9.40E+01	9.40E+01	5.71E+02
H24	2.00E-01	3.41E+01	3.70E-05	5.27E+01	2.92E-01	5.07E+01	-8.31E-01	-9.37E-01	1.77E+00	4.44E-01	1.80E+01	5.96E+00	2.40E+01	2.40E+01	6.47E+00
H29	1.55E-01	3.93E+01	4.10E-05	5.43E+01	3.30E-01	5.14E+01	-9.62E-01	-4.06E-01	1.37E+00	3.57E-02	1.90E+01	6.38E+00	2.54E+01	2.54E+01	5.79E+00
Complex 2															
Ru1	8.62E-01	9.91E+01	4.02E-04	4.40E+03	4.06E+01	2.49E+00	2.21E+01	2.48E+01	2.66E+00	1.17E+00	2.71E+03	2.35E+01	2.69E+03	2.69E+03	5.42E+03
C7	-1.16E-01	7.31E+01	-8.10E-05	3.76E+01	4.04E+00	2.08E+00	3.14E+00	6.11E+00	9.25E+00	8.08E-02	9.07E+01	2.25E+01	1.13E+02	1.13E+02	1.27E+02
C8	-1.05E-01	7.72E+01	8.60E-05	3.76E+01	4.04E+00	2.07E+00	3.83E+00	6.13E+00	9.96E+00	1.24E-01	1.07E+02	2.23E+01	1.29E+02	1.29E+02	1.23E+02
C13	-9.59E-02	7.87E+01	1.57E-04	3.76E+01	4.03E+00	2.07E+00	3.78E+00	6.21E+00	9.99E+00	1.31E-01	1.05E+02	2.24E+01	1.28E+02	1.28E+02	1.24E+02
C14	-8.33E-01	7.24E+01	7.10E-05	3.76E+01	4.02E+00	2.07E+00	2.78E+00	5.96E+00	8.74E+00	8.95E-01	8.18E+01	2.17E+01	1.03E+02	1.03E+02	1.41E+02

	02	01	05	01	00	00	00	00	00	02	01	01	02	02	02
							-	-	-						
C18	-1.16E-01	7.32E+01	-8.80E-05	3.76E+01	4.04E+00	2.08E+00	3.15E+00	6.13E+00	9.27E+00	1.08E-01	9.08E+01	2.25E+01	1.13E+02	1.13E+02	1.27E+02
							-	-	-						
C19	-1.05E-01	7.73E+01	5.90E-05	3.76E+01	4.04E+00	2.07E+00	3.83E+00	6.15E+00	9.97E+00	1.36E-01	1.07E+02	2.23E+01	1.29E+02	1.29E+02	1.23E+02
							-	-	-						
Cl10	-4.78E-01	2.34E+02	9.30E-05	4.57E+02	1.69E+01	5.87E-01	2.19E+01	3.41E+00	2.53E+01	1.29E+00	1.03E+03	4.26E+00	1.03E+03	1.03E+03	3.66E+02
							-	-	-						
N11	-9.49E-01	7.74E+01	5.00E-05	5.45E+01	6.07E+00	1.88E+00	3.67E+00	6.91E+00	1.06E+01	6.40E-01	1.00E+00	1.39E+01	1.49E+01	1.49E+01	3.56E+02
							-	-	-						
N16	-9.49E-01	7.75E+01	6.00E-05	5.45E+01	6.07E+00	1.88E+00	3.67E+00	6.90E+00	1.06E+01	6.04E-01	4.94E-01	1.39E+01	1.44E+01	1.44E+01	3.58E+02
							-	-	-						
H34	1.49E-01	3.72E+01	3.70E-05	5.50E-01	3.27E-01	5.25E-01	-9.03E-01	-4.11E-01	1.31E+00	3.97E-02	1.89E+01	4.77E+00	2.37E+01	2.37E+01	1.00E+01
							-	-	-						
H38	1.51E-01	3.74E+01	8.50E-05	5.47E-01	3.24E-01	5.25E-01	-9.09E-01	-4.18E-01	1.33E+00	4.12E-02	1.88E+01	4.70E+00	2.35E+01	2.35E+01	9.30E+00
Complex 3															
							-	-	-						
Ru1	8.73E-01	9.61E+01	-6.31E-04	4.40E+03	4.07E+01	2.45E+00	2.73E+01	2.50E+01	2.30E+00	8.55E-01	3.41E+03	2.29E+01	3.38E+03	3.38E+03	5.84E+03
							-	-	-						
C13	-7.91E-02	7.32E+01	-2.80E-05	3.76E+01	4.01E+00	2.07E+00	3.29E+00	6.18E+00	9.47E+00	8.53E-02	9.47E+01	2.23E+01	1.17E+02	1.17E+02	1.29E+02
							-	-	-						
C14	-9.33E-02	7.21E+01	-4.90E-05	3.76E+01	4.02E+00	2.07E+00	3.27E+00	6.65E+00	9.92E+00	4.11E-01	9.41E+01	2.29E+01	1.17E+02	1.17E+02	1.27E+02
							-	-	-						
C18	-8.72E-02	7.36E+01	8.60E-05	3.76E+01	4.02E+00	2.07E+00	3.26E+00	6.21E+00	9.47E+00	9.86E-02	9.37E+01	2.23E+01	1.16E+02	1.16E+02	1.29E+02
							-	-	-						
C21	-1.18E-01	7.64E+01	-3.70E-05	3.76E+01	4.04E+00	2.08E+00	3.50E+00	6.31E+00	9.81E+00	1.79E-01	9.88E+01	2.29E+01	1.22E+02	1.22E+02	1.25E+02
							-	-	-						
C23	-9.43E-02	7.27E+01	-8.40E-05	3.76E+01	4.03E+00	2.06E+00	3.35E+00	6.66E+00	1.00E+01	4.09E-01	9.62E+01	2.29E+01	1.19E+02	1.19E+02	1.29E+02

		-	-	-											
C25	-1.17E-01	7.61E+01	1.72E-04	3.76E+01	4.04E+00	2.08E+00	3.46E+00	6.30E+00	9.76E+00	1.83E-01	9.81E+01	2.30E+01	1.21E+02	1.21E+02	1.25E+02
		-	-	-											
CI7	-4.97E-01	2.32E+02	7.30E-05	4.57E+02	1.69E+01	5.90E+01	2.18E+01	3.20E+00	2.50E+01	1.35E+00	1.01E+03	3.77E+00	1.02E+03	1.02E+03	4.99E+02
		-	-	-											
N10	-9.26E-01	7.82E+01	2.83E-04	5.44E+01	6.06E+00	1.86E+00	3.72E+00	7.07E+00	1.08E+01	9.41E-01	2.92E+00	1.31E+01	1.60E+01	1.60E+01	3.60E+02
		-	-	-											
N16	-9.30E-01	7.78E+01	2.81E-04	5.45E+01	6.06E+00	1.87E+00	3.73E+00	6.99E+00	1.07E+01	9.13E-01	4.52E+00	1.31E+01	1.76E+01	1.76E+01	3.55E+02
		-	-	-											
O2	-8.96E-01	1.04E+02	1.68E-04	7.47E+01	7.80E+00	1.09E+00	5.04E+00	2.27E+00	7.31E+00	9.29E-01	7.73E+01	4.31E+00	7.30E+01	7.30E+01	5.56E+02
		-	-	-											
O28	-8.94E-01	1.05E+02	1.36E-04	7.47E+01	7.81E+00	1.09E+00	5.06E+00	2.21E+00	7.27E+00	8.86E-01	8.26E+01	4.19E+00	7.84E+01	7.84E+01	5.53E+02
		-	-	-											
H31	1.61E-01	3.65E+01	3.80E-05	5.47E-01	3.19E-01	5.20E-01	-9.69E-01	1.06E+00	2.03E+00	3.16E-01	1.93E+01	7.78E+00	2.71E+01	2.71E+01	5.20E+00
		-	-	-											
H36	1.77E-01	3.48E+01	1.80E-05	5.41E-01	3.05E-01	5.17E-01	-9.20E-01	1.05E+00	1.98E+00	3.08E-01	1.89E+01	7.65E+00	2.66E+01	2.66E+01	6.36E+00
Complex 4															
		-	-	-											
Ru1	9.46E-01	1.02E+02	-2.96E-04	4.40E+03	4.07E+01	2.38E+00	2.97E+01	2.51E+01	4.51E+00	4.45E-01	3.77E+03	2.31E+01	3.75E+03	3.75E+03	6.50E+03
		-	-	-											
C2	-9.53E-02	7.64E+01	9.40E-05	3.76E+01	4.03E+00	2.06E+00	3.39E+00	6.18E+00	9.57E+00	9.19E-02	9.45E+01	2.19E+01	1.16E+02	1.16E+02	1.28E+02
		-	-	-											
C3	-9.30E-02	7.35E+01	-9.30E-05	3.76E+01	4.03E+00	2.06E+00	3.29E+00	6.17E+00	9.46E+00	1.09E-01	9.52E+01	2.33E+01	1.18E+02	1.18E+02	1.32E+02
		-	-	-											
C4	-9.54E-02	7.65E+01	-5.80E-05	3.76E+01	4.03E+00	2.06E+00	3.40E+00	6.18E+00	9.58E+00	1.18E-01	9.46E+01	2.19E+01	1.17E+02	1.17E+02	1.28E+02
		-	-	-											
C5	-8.74E-02	7.61E+01	-7.70E-05	3.76E+01	4.03E+00	2.06E+00	3.50E+00	6.23E+00	9.73E+00	1.29E-01	1.01E+02	2.36E+01	1.25E+02	1.25E+02	1.26E+02
		-	-	-											
C6	-9.30E-01	7.35E+01	-5.60E-05	3.76E+01	4.03E+00	2.06E+00	3.28E+00	6.16E+00	9.45E+00	1.27E-01	9.49E+01	2.33E+01	1.18E+02	1.18E+02	1.32E+02

	02	01	05	01	00	00	00	00	00	01	01	01	02	02	02
							-	-	-						
C7	-8.74E-02	7.61E+01	-8.00E-05	3.76E+01	4.03E+00	2.06E+00	3.51E+00	6.23E+00	9.74E+00	1.46E-01	1.01E+02	2.36E+01	1.25E+02	1.25E+02	1.26E+02
							-	-	-						
N10	-6.07E-01	7.84E+01	1.22E-04	5.42E+01	5.78E+00	1.82E+00	3.84E+00	6.59E+00	1.04E+01	6.51E-01	1.74E+01	1.59E+01	3.33E+01	3.33E+01	1.33E+02
							-	-	-						
N13	-6.07E-01	7.84E+01	-4.29E-04	5.42E+01	5.78E+00	1.82E+00	3.84E+00	6.59E+00	1.04E+01	6.36E-01	1.74E+01	1.59E+01	3.33E+01	3.33E+01	1.33E+02
							-	-	-						
N15	-9.65E-01	7.53E+01	1.70E-04	5.46E+01	6.10E+00	1.86E+00	4.55E+00	6.58E+00	1.11E+01	3.83E-01	6.20E+01	1.36E+01	7.56E+01	7.56E+01	2.10E+02
Complex 5															
							-	-	-						
Ru1	9.26E-01	1.01E+02	9.21E-04	4.40E+03	4.06E+01	2.43E+00	2.82E+01	2.44E+01	3.84E+00	4.94E-01	3.57E+03	2.23E+01	3.55E+03	3.55E+03	5.25E+03
							-	-	-						
C18	-8.75E-02	7.60E+01	-2.80E-04	3.76E+01	4.03E+00	2.06E+00	3.44E+00	6.11E+00	9.55E+00	7.76E-02	9.92E+01	2.27E+01	1.22E+02	1.22E+02	1.29E+02
							-	-	-						
C20	-9.77E-02	7.36E+01	-1.45E-04	3.76E+01	4.03E+00	2.07E+00	3.26E+00	6.04E+00	9.29E+00	9.89E-02	9.45E+01	2.26E+01	1.17E+02	1.17E+02	1.32E+02
							-	-	-						
C22	-8.70E-02	7.59E+01	7.70E-05	3.76E+01	4.03E+00	2.06E+00	3.43E+00	6.10E+00	9.53E+00	9.16E-02	9.91E+01	2.27E+01	1.22E+02	1.22E+02	1.29E+02
							-	-	-						
C23	-1.01E-01	7.60E+01	-4.40E-05	3.76E+01	4.03E+00	2.07E+00	3.31E+00	6.03E+00	9.34E+00	1.18E-01	9.15E+01	2.12E+01	1.13E+02	1.13E+02	1.33E+02
							-	-	-						
C26	-9.72E-02	7.36E+01	-2.40E-05	3.76E+01	4.03E+00	2.07E+00	3.26E+00	6.03E+00	9.29E+00	1.14E-01	9.46E+01	2.26E+01	1.17E+02	1.17E+02	1.32E+02
							-	-	-						
C27	-1.01E-01	7.59E+01	-1.10E-04	3.76E+01	4.03E+00	2.07E+00	3.30E+00	6.03E+00	9.33E+00	1.38E-01	9.13E+01	2.12E+01	1.13E+02	1.13E+02	1.33E+02
							-	-	-						
N14	-9.58E-01	8.06E+01	1.32E-04	5.44E+01	6.10E+00	1.85E+00	3.97E+00	7.06E+00	1.10E+01	5.43E-01	1.27E+01	1.30E+01	2.57E+01	2.57E+01	3.17E+02
							-	-	-						
N15	-9.52E-01	7.36E+01	3.41E-04	5.46E+01	6.05E+00	1.90E+00	3.79E+00	6.70E+00	1.05E+01	2.78E-01	1.69E+01	1.34E+01	3.04E+01	3.04E+01	2.83E+02

							-	-	-						
N21	-9.59E-	8.06E+	8.90E-	5.44E+	6.11E+	1.85E+	3.97E+	7.07E+	1.10E+	4.49E-	1.32E+	1.30E+	2.62E+	2.62E+	3.16E+
	01	01	05	01	00	00	00	00	01	01	01	01	01	01	02

Table S5: The atomic NMR magnetizability and shielding for the selected atoms in the complexes using combined basis sets ECP(Ru)|6-31+G(d,p)

Complex 1															
q(A)	Vol(A),				DI(A,A')		$\chi_{\text{intra_Iso}}$	$\chi_{\text{bond_Is}}$	$\sigma_{\text{Iso}}(A,$		$\sigma_{\text{Iso}}(A',$	Isotropi		Anisotr	
	0.001	L(A)	K(A)	Li(A)	/2	(A)	(A)	$\chi_{\text{Iso}}(A)$	Mu(A)	A)	A)	$\sigma_{\text{Iso}}(A)$	c	opy	
							-	-							-
	9.40E-	1.03E+	2.84E-	3.27E+	1.24E+	2.70E+	1.42E+	4.48E+	3.06E+	1.07E+	1.52E+	2.53E+	1.55E+	4.41E+	6.61E+0
	01	02	04	01	01	00	01	01	01	00	03	01	03	02	2
							-	-							
	-7.56E-	7.39E+	2.30E-	3.78E+	3.98E+	2.09E+	3.04E+	6.24E+	9.29E+	1.03E-	8.71E+	1.29E+	1.00E+	1.10E+	1.21E+0
	02	01	05	01	00	00	00	00	00	01	01	01	02	02	2
							-	-							
	-5.98E-	7.16E+	-2.78E-	3.78E+	3.97E+	2.09E+	2.69E+	6.57E+	9.26E+	5.74E-	7.77E+	7.36E+	7.04E+	1.01E+	1.28E+0
	02	01	04	01	00	00	00	00	00	02	01	00	01	02	2
							-	-							
	-6.39E-	7.18E+	-2.07E-	3.78E+	3.97E+	2.09E+	2.75E+	8.16E+	1.09E+	1.06E-	8.19E+	6.66E+	8.85E+	1.05E+	1.23E+0
	02	01	04	01	00	00	00	00	01	01	01	00	01	02	2
							-	-							
	-6.54E-	7.37E+	1.51E-	3.78E+	3.98E+	2.09E+	3.07E+	2.09E+	5.16E+	8.08E-	8.69E+	3.55E+	5.14E+	1.09E+	1.26E+0
	02	01	04	01	00	00	00	00	00	02	01	01	01	02	2
							-	-							
	-7.15E-	7.47E+	2.20E-	3.78E+	3.98E+	2.09E+	3.17E+	3.53E+	6.70E+	9.33E-	8.93E+	2.09E+	6.84E+	1.12E+	1.21E+0
	02	01	05	01	00	00	00	00	00	02	01	01	01	02	2
							-	-							
	-4.23E-	7.01E+	-2.32E-	3.78E+	3.96E+	2.09E+	2.42E+	7.31E-	1.69E+	8.09E-	7.01E+	4.33E+	2.68E+	9.18E+	1.43E+0
	02	01	04	01	00	00	00	01	00	02	01	01	01	01	2
							-	-							
	-4.92E-	2.30E+	6.40E-	4.59E+	1.68E+	6.81E-	2.03E+	3.56E+	2.38E+	1.16E+	9.04E+	3.28E+	9.01E+	9.08E+	5.31E+0
	01	02	05	02	01	01	01	00	01	00	02	00	02	02	2
							-	-							
	-7.26E-	7.72E+	2.69E-	5.47E+	5.84E+	1.89E+	3.36E+	3.66E+	3.99E+	6.76E-	2.99E+	6.84E+	7.13E+	1.95E+	1.73E+0
	01	01	04	01	00	00	00	01	01	01	00	01	01	01	2
							-	-							
	-7.26E-	7.72E+	2.02E-	5.47E+	5.84E+	1.89E+	3.36E+	3.86E+	4.19E+	6.36E-	2.77E+	6.84E+	7.12E+	1.93E+	1.74E+0

01	01	04	01	00	00	00	01	01	01	00	01	01	01	2
-						-	-	-		-		-	-	
1.13E+0	1.29E+02	7.20E-05	7.54E+01	8.15E+00	9.82E-01	5.90E+00	4.31E+00	1.02E+01	1.00E+00	8.96E+01	1.29E+01	7.67E+01	8.70E+01	5.43E+02
1.28E-01	4.04E+01	1.60E-05	5.64E-01	3.40E-01	5.32E-01	-9.11E-01	2.81E+00	3.72E+00	4.61E-01	1.75E+01	1.67E+01	3.42E+01	2.32E+01	6.57E+00
1.27E-01	4.04E+01	2.30E-05	5.64E-01	3.41E-01	5.32E-01	-9.13E-01	3.04E+00	3.95E+00	4.72E-01	1.75E+01	1.67E+01	3.43E+01	2.33E+01	6.51E+00
Complex 2														
9.57E-01	1.05E+02	4.83E-04	3.27E+01	1.23E+01	2.73E+00	1.23E+01	6.54E+01	5.31E+01	1.06E+00	1.53E+03	2.44E+01	1.56E+03	4.29E+02	5.46E+02
-6.65E-02	7.04E+01	4.00E-05	3.78E+01	3.97E+00	2.10E+00	2.58E+00	1.72E+00	4.30E+00	7.96E-02	7.53E+01	3.11E+01	4.43E+01	9.75E+01	1.31E+02
-7.69E-02	7.49E+01	-1.02E-04	3.78E+01	3.99E+00	2.09E+00	3.19E+00	4.54E+00	7.73E+00	9.09E-02	8.97E+01	5.68E+01	3.29E+01	1.11E+02	1.26E+02
-6.59E-02	7.67E+01	-2.96E-04	3.78E+01	3.98E+00	2.08E+00	3.15E+00	6.68E+00	9.83E+00	8.29E-02	8.70E+01	1.67E+01	7.03E+01	1.09E+02	1.26E+02
-4.00E-02	6.99E+01	-5.43E-04	3.78E+01	3.96E+00	2.08E+00	2.20E+00	4.69E+00	2.49E+00	9.34E-02	6.43E+01	7.19E+01	7.61E+00	8.54E+01	1.50E+02
-6.67E-02	7.04E+01	1.40E-05	3.78E+01	3.97E+00	2.10E+00	2.58E+00	3.40E-01	2.24E+00	1.05E-01	7.54E+01	3.10E+01	4.45E+01	9.75E+01	1.30E+02
-7.67E-02	7.48E+01	2.55E-04	3.78E+01	3.99E+00	2.09E+00	3.19E+00	3.50E+00	6.69E+00	8.19E-02	8.98E+01	5.67E+01	3.31E+01	1.11E+02	1.26E+02
-4.84E-01	2.34E+02	6.80E-05	4.59E+02	1.68E+01	6.57E-01	2.08E+01	4.12E+00	2.49E+01	1.12E+00	9.50E+02	1.89E+01	9.69E+02	9.55E+02	4.05E+02
1.16E+00	8.10E+01	6.60E-05	5.50E+01	6.27E+00	1.89E+00	3.34E+00	5.67E+01	6.00E+01	5.24E-01	2.40E+01	9.76E+01	7.36E+01	1.13E+01	3.45E+02
1.16E+00	8.09E+01	2.46E-04	5.50E+01	6.27E+00	1.89E+00	3.34E+00	5.92E+01	6.25E+01	4.90E-01	2.44E+01	9.81E+01	7.37E+01	1.17E+01	3.47E+02

Complex 1															
9.67E-02	4.09E+01	9.50E-05	5.81E-01	3.58E-01	5.45E-01	-9.04E-01	1.89E+00	2.80E+00	6.89E-02	1.80E+01	2.20E+01	4.00E+01	2.23E+01	1.05E+01	
Complex 2															
1.03E-01	4.10E+01	7.70E-05	5.76E-01	3.53E-01	5.45E-01	-9.10E-01	1.87E+00	2.78E+00	5.26E-02	1.79E+01	2.19E+01	3.98E+01	2.22E+01	9.71E+00	
Complex 3															
9.59E-01	1.01E+02	2.52E-04	3.27E+01	1.23E+01	2.71E+00	1.57E+01	2.51E+01	9.33E+00	7.69E-01	1.45E+03	2.38E+01	1.47E+03	5.15E+02	5.85E+02	
Complex 4															
-5.37E-02	7.08E+01	-6.10E-05	3.78E+01	3.97E+00	2.09E+00	2.74E+00	1.78E+00	4.52E+00	7.69E-02	7.80E+01	3.43E+01	4.37E+01	9.77E+01	1.36E+02	
Complex 5															
-6.12E-02	6.99E+01	-1.90E-04	3.78E+01	3.97E+00	2.09E+00	2.66E+00	1.16E+00	3.81E+00	4.01E-01	7.74E+01	1.80E+01	5.94E+01	9.83E+01	1.35E+02	
Complex 6															
-5.33E-02	7.09E+01	-1.48E-04	3.78E+01	3.97E+00	2.09E+00	2.71E+00	-8.01E-01	3.52E+00	8.25E-02	7.71E+01	3.46E+01	4.25E+01	9.68E+01	1.37E+02	
Complex 7															
-7.24E-02	7.38E+01	-1.20E-04	3.78E+01	3.98E+00	2.09E+00	2.90E+00	3.34E+00	6.24E+00	1.34E-01	8.08E+01	2.52E+00	7.83E+01	1.02E+02	1.32E+02	
Complex 8															
-6.28E-02	7.02E+01	-2.92E-04	3.78E+01	3.98E+00	2.09E+00	2.69E+00	-3.87E-01	3.08E+00	4.12E-01	7.80E+01	1.87E+01	5.94E+01	9.88E+01	1.36E+02	
Complex 9															
-7.20E-02	7.35E+01	-2.53E-04	3.78E+01	3.98E+00	2.09E+00	2.88E+00	2.91E+00	5.79E+00	1.42E-01	8.04E+01	2.87E+00	7.76E+01	1.02E+02	1.32E+02	
Complex 10															
-4.94E-01	2.33E+02	8.90E-05	4.59E+02	1.68E+01	6.69E-01	2.04E+01	3.83E+00	2.42E+01	1.16E+00	9.18E+02	1.43E+01	9.32E+02	9.23E+02	6.30E+02	
Complex 11															
1.14E+00	8.19E+01	2.23E-04	5.49E+01	6.27E+00	1.87E+00	3.46E+00	3.07E+01	3.42E+01	9.04E-01	1.84E+01	6.25E+01	4.41E+01	4.93E+00	3.41E+02	
Complex 12															
1.14E+00	8.18E+01	5.60E-05	5.49E+01	6.27E+00	1.87E+00	3.46E+00	3.25E+01	3.60E+01	8.69E-01	1.79E+01	6.25E+01	4.46E+01	4.43E+00	3.41E+02	
Complex 13															
1.15E+00	1.27E+02	1.11E-04	7.54E+01	8.14E+00	1.01E+00	5.80E+00	1.14E+01	1.72E+01	8.93E-01	7.45E+01	-1.02E-01	7.46E+01	7.20E+01	5.01E+02	
Complex 14															
1.15E+00	1.27E+02	1.17E-04	7.54E+01	8.14E+01	1.01E+01	5.82E+01	1.19E+01	1.77E+01	8.78E-01	7.53E+01	-5.23E-01	7.54E+01	7.28E+01	4.99E+02	

0	02	04	01	00	00	00	01	01	01	01	02	01	01	2
1.18E-01	3.91E+01	2.20E-05	5.75E-01	3.43E-01	5.39E-01	-9.17E-01	5.56E+00	6.48E+00	3.52E-01	1.80E+01	9.34E+00	8.68E+00	2.41E+01	6.77E+00
1.22E-01	3.86E+01	3.20E-05	5.74E-01	3.40E-01	5.39E-01	-9.03E-01	5.80E+00	6.70E+00	3.54E-01	1.79E+01	9.83E+00	8.07E+00	2.40E+01	7.09E+00
Complex 4														
9.82E-01	1.12E+02	-8.61E-04	3.27E+01	1.24E+01	2.63E+00	1.72E+01	3.72E+01	2.00E+01	3.23E-01	1.38E+03	2.40E+01	1.41E+03	5.81E+02	7.13E+02
-5.75E-02	7.36E+01	-2.60E-05	3.78E+01	3.98E+00	2.08E+00	2.80E+00	3.30E+00	6.10E+00	7.55E-02	7.77E+01	2.22E+01	5.55E+01	9.86E+01	1.32E+02
-5.54E-02	7.08E+01	-2.19E-04	3.78E+01	3.97E+00	2.08E+00	2.71E+00	3.84E+00	6.55E+00	7.39E-02	7.94E+01	7.69E+00	7.17E+01	1.02E+02	1.32E+02
-5.77E-02	7.36E+01	-3.00E-05	3.78E+01	3.98E+00	2.08E+00	2.81E+00	2.41E+00	5.22E+00	9.46E-02	7.79E+01	2.22E+01	5.57E+01	9.87E+01	1.32E+02
-4.58E-02	7.27E+01	-9.20E-05	3.78E+01	3.97E+00	2.08E+00	2.85E+00	5.17E+00	8.02E+00	9.35E-02	8.34E+01	6.44E+00	8.98E+01	1.07E+02	1.31E+02
-5.55E-02	7.07E+01	-3.53E-04	3.78E+01	3.97E+00	2.08E+00	2.71E+00	3.48E+00	6.19E+00	8.67E-02	7.92E+01	7.80E+00	7.14E+01	1.02E+02	1.32E+02
-4.61E-02	7.27E+01	-7.90E-05	3.78E+01	3.97E+00	2.08E+00	2.85E+00	4.95E+00	7.80E+00	1.01E-01	8.35E+01	6.38E+00	8.99E+01	1.07E+02	1.31E+02
-7.38E-01	8.21E+01	2.73E-04	5.47E+01	5.89E+00	1.85E+00	3.51E+00	1.44E+01	1.79E+01	5.90E-01	1.80E+00	3.50E+01	3.68E+01	1.66E+01	1.21E+02
-7.38E-01	8.21E+01	2.80E-04	5.47E+01	5.89E+00	1.85E+00	3.51E+00	1.46E+01	1.81E+01	5.75E-01	1.80E+00	3.50E+01	3.68E+01	1.66E+01	1.21E+02
1.18E+00	7.93E+01	4.18E-04	5.51E+01	6.30E+00	1.87E+00	4.22E+00	2.86E+01	3.28E+01	3.23E-01	3.98E+01	6.59E+01	1.06E+02	5.34E+01	1.95E+02
Complex 5														
9.80E-01	1.09E+02	1.36E-03	3.27E+01	1.23E+01	2.69E+00	1.65E+01			5.13E-01	1.36E+03	2.32E+01	1.38E+03		4.66E+02
							5.14E+01	3.49E+01					6.08E+01	

															01		01		02	
-4.88E-	7.28E+	-3.90E-	3.78E+	3.97E+	2.08E+	2.82E+	5.26E+	8.08E+	6.77E-	8.26E+	1.05E+	8.15E+	1.04E+	1.33E+0	-	-	-	-	-	-
02	01	04	01	00	00	00	00	00	02	01	00	01	02	2						
-5.86E-	7.09E+	-3.20E-	3.78E+	3.97E+	2.08E+	2.72E+	1.28E+	4.00E+	1.08E-	7.85E+	2.39E+	5.46E+	9.99E+	1.34E+0	-	-	-	-	-	-
02	01	05	01	00	00	00	00	00	01	01	01	01	01	2						
-4.81E-	7.26E+	4.30E-	3.78E+	3.97E+	2.08E+	2.81E+	5.08E+	7.89E+	6.63E-	8.24E+	1.02E+	8.14E+	1.04E+	1.33E+0	-	-	-	-	-	-
02	01	05	01	00	00	00	00	00	02	01	00	01	02	2						
-6.19E-	7.32E+	-2.73E-	3.78E+	3.98E+	2.08E+	2.73E+	2.89E+	5.62E+	1.00E-	7.44E+	4.68E+	2.76E+	9.43E+	1.38E+0	-	-	-	-	-	-
02	01	04	01	00	00	00	00	00	01	01	01	01	01	2						
-5.90E-	7.11E+	-3.44E-	3.78E+	3.97E+	2.08E+	2.72E+	1.14E+	3.86E+	1.17E-	7.88E+	2.38E+	5.49E+	1.00E+	1.34E+0	-	-	-	-	-	-
02	01	04	01	00	00	00	00	00	01	01	01	01	02	2						
-6.16E-	7.31E+	-2.82E-	3.78E+	3.98E+	2.08E+	2.73E+	2.89E+	5.62E+	9.87E-	7.42E+	4.68E+	2.74E+	9.40E+	1.38E+0	-	-	-	-	-	-
02	01	04	01	00	00	00	00	00	02	01	01	01	01	2						
1.17E+0	8.46E+	2.33E-	5.49E+	6.31E+	1.86E+	3.60E+	2.02E+	2.38E+	4.48E-	6.06E+	4.15E+	3.55E+	6.43E+	2.96E+0	-	-	-	-	-	-
0	01	04	01	00	00	00	01	01	01	00	01	01	00	2						
1.17E+0	7.77E+	4.56E-	5.51E+	6.26E+	1.91E+	3.45E+	4.33E+	4.68E+	2.20E-	9.09E+	1.00E+	9.11E+	4.13E+	2.62E+0	-	-	-	-	-	-
0	01	04	01	00	00	00	01	01	01	00	02	01	00	2						
1.17E+0	8.46E+	2.77E-	5.49E+	6.31E+	1.86E+	3.61E+	2.10E+	2.46E+	3.55E-	5.54E+	4.13E+	3.57E+	6.90E+	2.94E+0	-	-	-	-	-	-
0	01	04	01	00	00	00	01	01	01	00	01	01	00	2						