

## Supporting Information:

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### Tuning of Redox Properties for the Design of Ruthenium Anticancer Drugs: Part 2. Syntheses, Crystal Structures and Electrochemistry of Potentially Antitumor $[\text{Ru}^{\text{III/II}}\text{Cl}_{6-n}(\text{azole})_n]^z$ ( $n = 3,4,6$ ) Complexes

S1–S2: Geometrical Details of complexes **B** and **C** studied by X-ray diffraction methods

S3–S7: Selected Bond Distances and Angles for **B** and **C**

S8: UV-vis Absorption Bands for **B** and **C** in Methanol at 298K

S9–S10: Figures showing ORTEP views of **B1b**, **B7b**, packing details of **B7**, **C2**, **C3** and **C6**

S11: UV-vis Absorption Bands against  $E_{1/2}$  ( $\text{Ru}^{\text{III}}/\text{Ru}^{\text{II}}$ ) for **B** and **C**.

S12: The Electronic Absorption Spectra of **C2** and **C6** in 0.2 M phosphate buffer at pH 7.0.

**Crystal structures.** Complex **B1**. The dihedral angles of mutually *trans* butylimidazole ligands are –30.6 and –9.9° in **B1a** and **B1b**, correspondingly. The butylimidazol ligands coordinated to ruthenium via N10 and N19 in **B1a** are tilted relative to the mean plane through Ru1N10N19Cl11Cl13, the corresponding angles being at 36.7 and 6.3°. In **B1b** the dihedral angles between imidazole rings bonded to ruthenium via N37 and N46, and the mean plane through Ru2N37N46Cl4Cl6 are at 45.2 and 35.6°, respectively. The imidazole rings ligated to ruthenium via N1 and N28 nearly bisect the angles Cl3Ru1N10 and Cl6Ru2N46, the torsion angles  $\Theta_{\text{C2N1Ru1Cl3}}$  and  $\Theta_{\text{C29N28Ru2Cl6}}$  being at –38.60(15) and 33.90(15)°, correspondingly.

Complex **B4**. The bond angles at the ruthenium center have almost idealized values; the largest deviations from 90 and 180° were found for Cl3–Ru–Cl2 at 92.67(3)° and N4–Ru–N10 at 176.94(7)°. The triazole ligands coordinated *via* N4 and N10 are tilted relative to the mean plane through RuN4N10Cl11Cl13, the corresponding angles being at 48.7 and 14.5°, while the triazole ligand ligated *via* N16 nearly bisects the angle Cl1–Ru–N4, the torsion angle  $\Theta_{\text{Cl1RuN16Cl15}}$  being at 37.87(19)°.

Complex **B5**. The 4-methylpyrazole ligands linked to the Ru atom *via* N1 and N7 are nearly coplanar to the mean planes through RuCl11Cl2Cl3N1 and RuCl2N1N7N13, correspondingly. The torsion angles

$\Theta_{\text{N1RuN7C11}}$  and  $\Theta_{\text{Cl3RuN1N2}}$  are at  $-4.25(18)$  and  $2.31(15)^\circ$ , respectively. The thirdazole ring ligated to Ru atom *via* N13 is tilted to the RuCl2N7N1N13 mean plane by  $\sim 26^\circ$ .

Complex **B6**·H<sub>2</sub>O. The triazole ligand ligated *via* N2 bisects the angle Cl1–Ru–N3, the torsion angle  $\Theta_{\text{Cl1RuN2C3}}$  being at  $-45.44(15)^\circ$ . The other two torsion angles  $\Theta_{\text{Cl3RuN7N6}}$  and  $\Theta_{\text{Cl3RuN12C13}}$  indicating the position of triazole ligands coordinated *via* N7 and N12 are at  $48.38(12)$  and  $-48.52(15)^\circ$ , respectively.

Complex **B7**. The pyrazole ligands *trans* to each other are almost coplanar in both cases, the corresponding dihedral angle being at  $1.7$  (**B7a**) and  $3.8^\circ$  (**B7b**). Their position in respect to the Ru1N6N11Cl1Cl3 and Ru2N21N26Cl4Cl6 can be defined by torsion angles  $\Theta_{\text{N12N11Ru1Cl3}}$  at  $-29.1(2)$  and  $\Theta_{\text{N7N6Ru1Cl3}}$  at  $30.14(18)^\circ$  in molecule **B7a** and  $\Theta_{\text{Cl6Ru2N21C25}}$  at  $156.2(2)$  and  $\Theta_{\text{Cl6Ru2N26C30}}$  at  $-154.6(2)^\circ$  in **B7b**. The positions of the pyrazole rings bonded to ruthenium through N1 and N16 in respect to Ru1Cl1Cl2Cl3N1 and Ru2Cl4Cl5Cl6N16 are defined by the torsion angles  $\Theta_{\text{N2N1Ru1Cl3}}$  at  $-18.63(18)^\circ$  and  $\Theta_{\text{N17N16Ru2Cl6}}$  at  $40.75(18)^\circ$ .

Complex **C2**. The orientation of the imidazole rings with respect to the Ru–Cl1 vector is almost staggered, the  $\Theta_{\text{C3N1RuCl1}}$  and  $\Theta_{\text{C6N3RuCl1}}$  torsion angles being at  $45.63(11)$  and  $50.48(11)^\circ$ .

Complex **C3**·CH<sub>3</sub>OH·(C<sub>2</sub>H<sub>5</sub>)<sub>2</sub>O. The orientation of benzimidazole ligands in respect to Ru–Cl1 and Ru–Cl2 vectors is very similar, tending to minimize steric crowding between them and is determined by torsion angles  $\Theta_{\text{C1N1RuCl1}}$  at  $-41.35(13)$ ,  $\Theta_{\text{C8N3RuCl1}}$  at  $-44.66(12)$ ,  $\Theta_{\text{C21N5RuCl2}}$  at  $-45.79(14)$  and  $\Theta_{\text{C28N7RuCl2}}$  at  $-45.41(14)^\circ$ .

Complex **C6a**·2H<sub>2</sub>O. The orientation of the Htrz planes with respect to the Ru–Cl vector is almost staggered, the  $\Theta_{\text{C3N2RuCl}}$  and  $\Theta_{\text{N6N7RuCl}}$  torsion angles being at  $31.8(3)$  and  $30.3(3)^\circ$ , respectively.

Complex **C7**. The *trans*-coordinated pyrazole ligands are parallel to each other. The dihedral angle between them is small and does not exceed  $1^\circ$ . The N–N bonds in *trans*-coordinated ligands are also *trans* to each other. The orientation of pyrazole ligands in respect to Ru–Cl1 and Ru–Cl2 vectors is described by the torsion angles  $\Theta_{\text{C5N1Ru1Cl1}}$  at  $-17.61(16)$ ,  $\Theta_{\text{N7N6Ru1Cl1}}$  at  $24.20(13)$ ,  $\Theta_{\text{N12N11Ru2Cl2}}$  at  $-33.33(14)$  and  $\Theta_{\text{N17–N16–Ru2–Cl2}}$  at  $21.67(14)^\circ$ . Both the coordinated chloro ligands and the chloride counter anions are involved in hydrogen bonding interactions.

**Table S1.** Selected Bond Distances (Å) and Angles (°) in Two Independent Molecules of **B1**

Atom1–Atom2	<b>B1a</b>	Atom1–Atom2	<b>B1b</b>
Ru1–N1	2.0623(15)	Ru2–N28	2.0670(15)
Ru1–N10	2.0768(14)	Ru2–N37	2.0763(15)
Ru1–N19	2.0886(14)	Ru2–N46	2.0744(15)
Ru1–Cl1	2.3602(6)	Ru2–Cl4	2.3597(6)
Ru1–Cl2	2.3696(6)	Ru2–Cl5	2.3800(6)
Ru1–Cl3	2.3603(6)	Ru2–Cl6	2.3522(6)
Atom1-Atom2-Atom3		Atom1-Atom2-Atom3	
N10–Ru1–N19	176.88(6)	N37–Ru2–N46	178.02(6)
N1–Ru1–Cl2	178.55(4)	N28–Ru2–Cl5	177.94(4)
Cl1–Ru1–Cl3	178.343(15)	Cl4–Ru2–Cl6	178.389(16)

**Table S2.** Selected Interatomic Bond Distances (Å) and Angles (°) in **B4**

Atom1–Atom2	Å	Atom1–Atom2	Å
Ru1–N4	2.0691(18)	Ru1–Cl1	2.3627(7)
Ru1–N10	2.0864(18)	Ru1–Cl2	2.3541(8)
Ru1–N16	2.0748(19)	Ru1–Cl3	2.3395(6)
Atom1-Atom2-Atom3	deg	Atom1-Atom2-Atom3	deg
N4–Ru1–N10	176.94(7)	N16–Ru1–Cl2	177.30(5)
Cl1–Ru1–Cl3	177.37(2)		

**Table S3.** Selected Bond Distances (Å) and Angles (°) in **B5**

Atom1–Atom2	Å	Atom1–Atom2	Å
Ru1–N1	2.0793(17)	Ru1–Cl1	2.3468(6)
Ru1–N7	2.0576(16)	Ru1–Cl2	2.3912(6)
Ru1–N13	2.0504(16)	Ru1–Cl3	2.3413(6)
Atom1-Atom2-Atom3	deg	Atom1-Atom2-Atom3	deg
N7–Ru1–N13	178.50(7)	N1–Ru1–Cl2	178.66(5)
Cl1–Ru1–Cl3	178.654(19)		

**Table S4.** Selected Bond Distances (Å) and Angles (°) in **B6·H<sub>2</sub>O**

Atom1–Atom2	Å	Atom1–Atom2	Å
Ru1–N2	2.0764(15)	Ru1–Cl1	2.3434(7)
Ru1–N7	2.0734(16)	Ru1–Cl2	2.3469(6)
Ru1–N12	2.0667(16)	Ru1–Cl3	2.3427(8)
Atom1-Atom2-Atom3	deg	Atom1-Atom2-Atom3	deg
N7–Ru1–N12	178.66(6)	N2–Ru1–Cl2	179.51(4)
Cl1–Ru1–Cl3	178.539(16)		

**Table S5.** Selected Bond Distances (Å) and Angles (°) in Two Independent Molecules in **B7**

Atom1–Atom2	<b>B7a</b>	Atom1–Atom2	<b>B7b</b>
Ru1–N1	2.073(2)	Ru2–N16	2.0630(19)
Ru1–N6	2.0527(19)	Ru2–N21	2.0665(19)
Ru1–N11	2.0600(19)	Ru2–N26	2.0641(19)
Ru1–Cl1	2.3584(8)	Ru2–Cl4	2.3536(8)
Ru1–Cl2	2.3552(9)	Ru2–Cl5	2.3539(8)
Ru1–Cl3	2.3777(8)	Ru2–Cl6	2.3820(8)
Atom1-Atom2-Atom3		Atom1-Atom2-Atom3	
N6–Ru1–N11	177.54(8)	N21–Ru2–N26	179.28(7)
N1–Ru1–Cl2	178.37(5)	N16–Ru2–Cl5	179.06(5)
Cl1–Ru1–Cl3	179.41(2)	Cl4–Ru2–Cl6	177.61(2)

**Table S6.** Selected Bond Distances (Å) and Angles (°) in **C2**

Atom1–Atom2	Å	Atom1–Atom2	Å
Ru1–N1	2.0687(12)	Ru1–N3	2.0682(12)
Ru1–Cl1	2.3379(6)	Ru1–Cl2	2.3458(6)
Atom1-Atom2-Atom3 deg		Atom1-Atom2-Atom3 deg	
N3–Ru1–N1	90.19(5)	N3–Ru1–Cl1	90.43(3)
N1–Ru1–Cl1	90.34(3)	N3–Ru1–Cl2	89.57(3)
N1–Ru1–Cl2	89.66(3)	Cl1–Ru1–Cl2	180.0

**Table S7.** Selected Bond Distances (Å) and Angles (°) in **C3**·CH<sub>3</sub>OH·(C<sub>2</sub>H<sub>5</sub>)<sub>2</sub>O

Atom1–Atom2	Å	Atom1–Atom2	Å
Ru1–N1	2.0862(15)	Ru1–N7	2.0792(14)
Ru1–N3	2.0697(14)	Ru1–Cl1	2.3570(8)
Ru1–N5	2.0732(15)	Ru1–Cl2	2.3232(8)
Atom1-Atom2-Atom3	deg	Atom1-Atom2-Atom3	deg
N1–Ru1–N5	177.14(5)	N3–Ru1–N7	178.09(5)
Cl1–Ru1–Cl2	178.601(15)		

**Table S8.** Selected Bond Distances (Å) and Angles (°) in **C6**·2H<sub>2</sub>O

Atom1–Atom2	Å	Atom1–Atom2	Å
Ru1–N2	2.073(3)	Ru1–N7	2.074(3)
Ru1–Cl1	2.3289(10)		
Atom1-Atom2-Atom3	deg	Atom1-Atom2-Atom3	deg
N2–Ru1–N7	86.84(12)	N2–Ru1–Cl1	89.58(9)
N7–Ru1–Cl1	91.07(9)		

**Table S9.** Selected Bond Distances (Å) and Angles (°) in **C7**

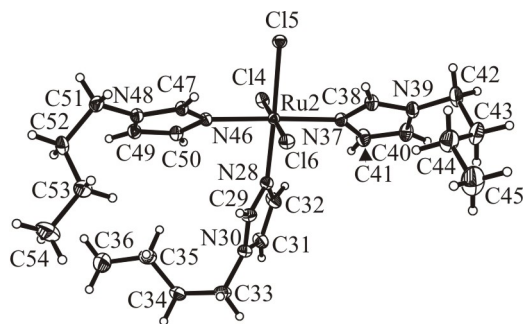
Atom1–Atom2	Å	Atom1–Atom2	Å
Ru1–N1	2.0557(15)	Ru2–N11	2.0624(15)
Ru1–N6	2.0571(15)	Ru2–N16	2.0659(18)
Ru1–Cl1	2.3767(9)	Ru2–Cl2	2.3565(7)
Atom1-Atom2-Atom3	deg	Atom1-Atom2-Atom3	deg
N1–Ru1–N6	89.14(6)	N11–Ru2–N16	93.03(6)
N1–Ru1–Cl1	90.86(5)	N11–Ru2–Cl2	91.09(5)
N6–Ru1–Cl1	89.13(5)	N16–Ru2–Cl2	89.34(5)

**Table S10.** UV-vis Absorption Bands [Wavelength and Molar Absorptivity (in Brackets)] for Complexes **B** and **C** in Methanol at 298K used for the establishment of equations (3) and (4). The metal-centered Ru<sup>III</sup>/Ru<sup>II</sup> redox potential increases with the increase of the absorption wavelengths  $\lambda$  (Fig. S7)

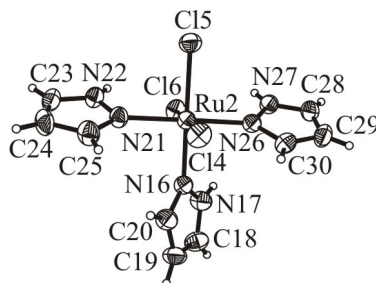
Complex	$\lambda_{\text{max}}$ (nm) / $\varepsilon$ (mM <sup>-1</sup> cm <sup>-1</sup> )	Complex	$\lambda_{\text{max}}$ (nm) / $\varepsilon$ (mM <sup>-1</sup> cm <sup>-1</sup> )
<b>B1</b>	347 (3.29)	<b>C2</b>	346 (2.71)
<b>B3</b>	358 (3.12)	<b>C3</b>	363 (2.67)
<b>B4</b>	371 (3.55)	<b>C6</b>	400 (2.77)
<b>B5</b>	369 (3.08)	<b>C7</b>	401 (3.71) <sup>a</sup>
<b>B6·H<sub>2</sub>O</b>	376 (3.35)	<b>C8</b>	452 (2.69)
<b>B7</b>	376 (3.18)		
<b>B8</b>	381 (3.71)		

<sup>a</sup> overlapped band

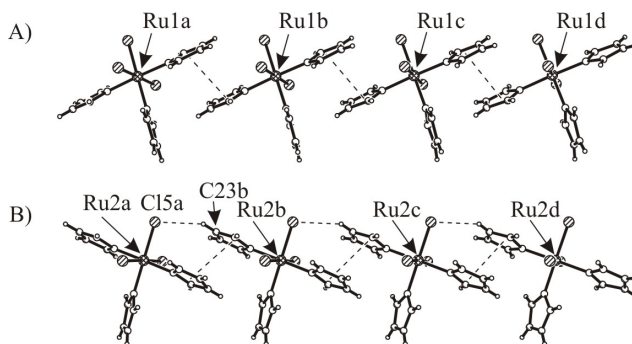




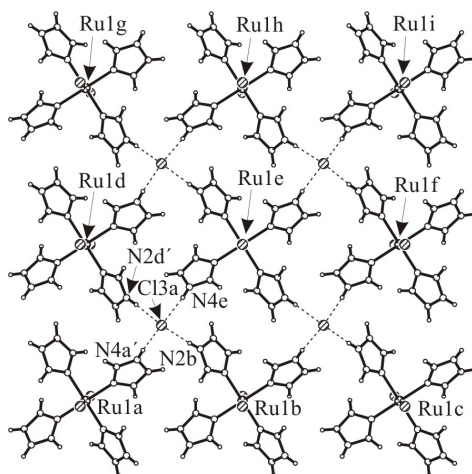
**Fig. S1.** ORTEP view of the second independent molecule of  $[\text{Ru}^{\text{III}}\text{Cl}_3(\text{buim})_3]$  (**B1b**), showing the atom-numbering scheme. Thermal displacement ellipsoids are drawn at the 50% probability level.



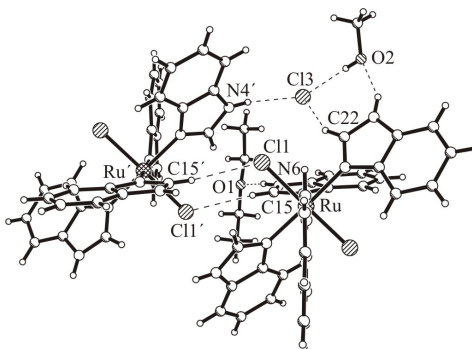
**Fig. S2.** ORTEP view of the second independent molecule of  $[\text{Ru}^{\text{III}}\text{Cl}_3(\text{Hpz})_3]$  (**B7b**), showing the atom-numbering scheme. Thermal displacement ellipsoids are drawn at the 50% probability level.



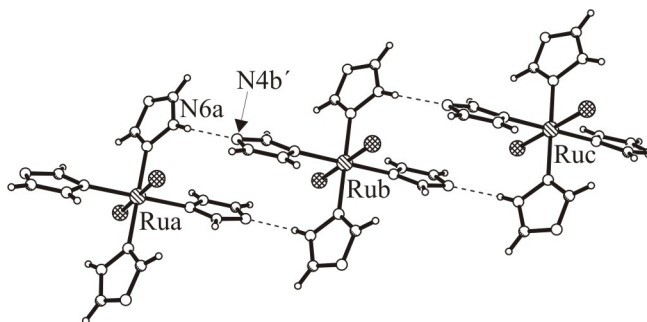
**Fig. S3.** Infinite chains in **B7** formed by the two independent molecules **B7a** (A) and **B7b** (B), which are stabilized by stacking  $\pi$ - $\pi$  interactions between *trans*-coordinated pyrazole ligands of adjacent molecules and *intermolecular* hydrogen bonding.



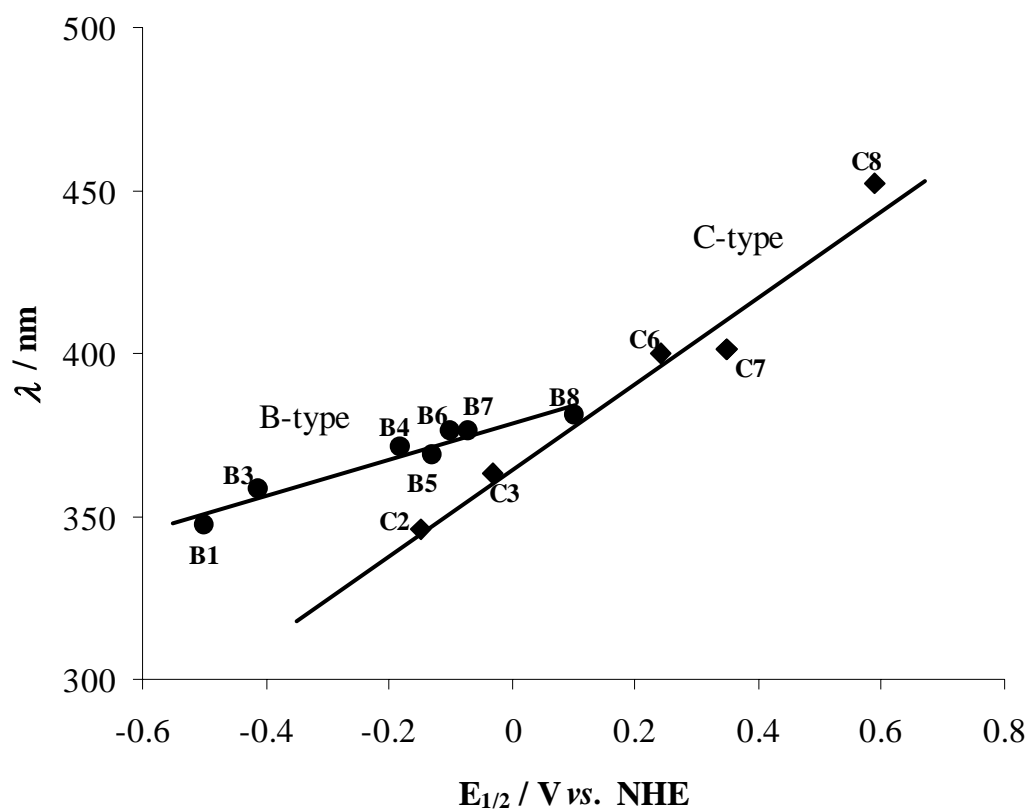
**Fig. S4.** N–H...Cl hydrogen bonding interactions of  $[\text{Ru}^{\text{III}}\text{Cl}_2(\text{Him})_4]\text{Cl}$  (**C2**) parallel to ac cell planes.



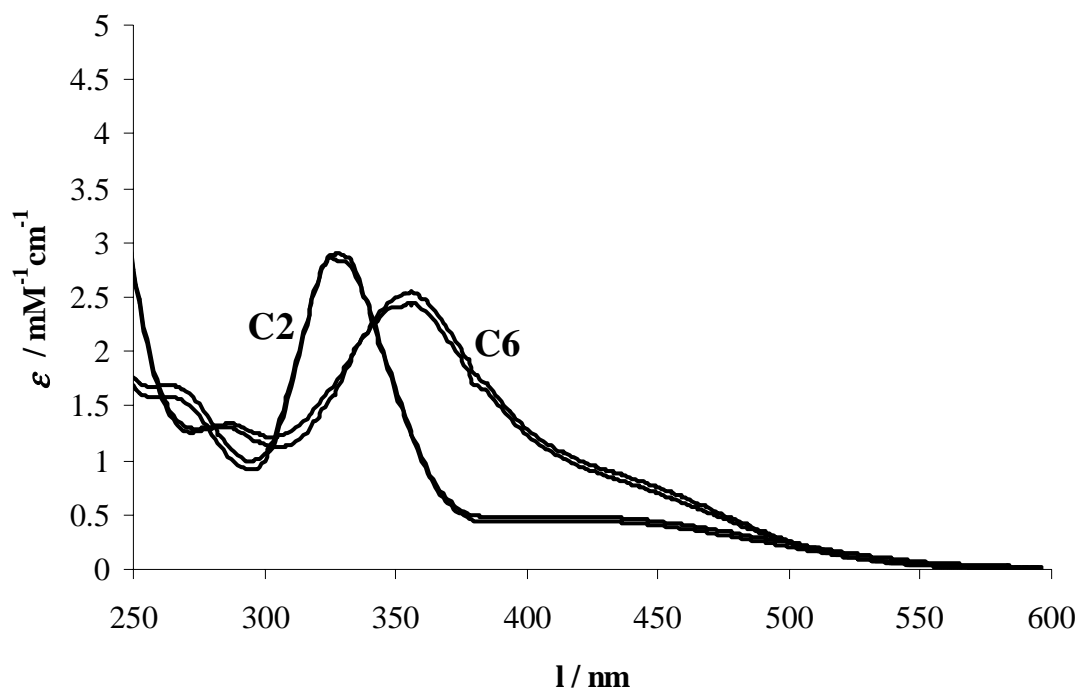
**Fig. S5.** A part of the crystal structure of **C3**·CH<sub>3</sub>OH·(C<sub>2</sub>H<sub>5</sub>)<sub>2</sub>O showing the formation of multiple hydrogen bonds between the complex cations, counter anion and solvent molecules.



**Fig. S6.** Parallel chains in the structure of **C6a**·2H<sub>2</sub>O, which are stabilized by strong *intrachain* hydrogen bonding interactions between  $[\text{Ru}^{\text{III}}\text{Cl}_2(\text{Htrz})_4]^+$ .



**Fig. S7.** Plot of the UV-vis absorption bands against  $E_{1/2}$  ( $\text{Ru}^{\text{III}}/\text{Ru}^{\text{II}}$ ) for complexes **B** (●) and **C** (◆). The electrochemical data is shown in Table 3 and the optical data in Table S10:  $\lambda$  (nm) = 55.9  $E_{1/2}$  (V) + 378.6 ( $r = 0.97$ ) (**B**) and  $\lambda$  (nm) = 136.1  $E_{1/2}$  (V) + 365.2 ( $r = 0.99$ ) (**C**).



**Fig. S8.** The electronic absorption spectra of **C2** [measured directly after dissolution (upper line) and after 24 h (lower line)] and **C6** [measured directly after dissolution (upper line) and after 3 h (lower line)] in 0.2 M phosphate buffer at pH 7.0 at room temperature.