SUPPLEMENTARY INFORMATION

Diruthenium complexes as pH-responsive delivery systems: a quantitative assessment

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• Figure S7. Electronic spectra of **RuNAA** and **Ru'NAA** in DMSO/water solution using an HEPES KOH (4-(2-hydroxyethyl)-1-piperazineethanesulfonic acid) buffer at pH 6.5 over 24 h.

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- Figure S9. Temperature dependence of the molar susceptibility χ_{M} and χ_{M} . T for **Ru'IAA**
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1. SINGLE CRYSTAL X-RAY DIFFRACTION

Empirical formula	
Eormula woight	1772 49
	1223.46
Temperature/K	296.15
Crystal system	triclinic
Space group	P-1
a/Å	11.869(2)
b/Å	15.027(3)
c/Å	16.745(3)
α/°	101.077(4)
β/°	107.283(3)
γ/°	106.402(3)
Volume/Å ³	2608.4(8)
Z	2
ρ _{calc} g/cm ³	1.558
µ/mm⁻¹	0.795
F(000)	1242.0
Crystal size/mm ³	$0.34 \times 0.2 \times 0.07$
Radiation	ΜοΚα (λ = 0.71073)
20 range for data collection/°	2.672 to 50.698
Index ranges	-14 ≤ h ≤ 14, -18 ≤ k ≤ 18, -20 ≤ l ≤ 20
Reflections collected	22965
Independent reflections	9497 [R _{int} = 0.0774, R _{sigma} = 0.1203]
Data/restraints/parameters	9497/0/664
Goodness-of-fit on F ²	0.971
Final R indexes [I>=2σ (I)]	$R_1 = 0.0547, wR_2 = 0.1162$
Final R indexes [all data]	R ₁ = 0.1270, wR ₂ = 0.1498
Largest diff. peak/hole / e Å ⁻³	0.96/-0.61

Table S1. Crystal and structure refinement data for Ru'2,4-D.



Figure S1. Asymmetric unit of Ru'2,4-D. Hydrogen atoms are omitted for clarity.



Figure S2. Intra- and intermolecular interactions found in the structure of of Ru'2,4-D (distances in Å).

			-
Ru1	Ru2	2.3288(8)	
Ru1	Cl1	2.416(2)	
Ru1	01	2.113(5)	
Ru1	N1	2.096(5)	
Ru1	N3	2.060(6)	
Ru1	N5	2.102(5)	
Ru2	02	2.068(5)	
Ru2	N2	2.044(6)	
Ru2	N4	2.005(6)	
Ru2	N6	2.029(5)	
Ru1	Ru2	Cl1	173.69(5)
Ru1	01	N3	176.1(2)
Ru1	N1	N5	175.7(2)
Ru2	02	N4	178.4(2)
Ru2	N2	N6	174.5(2)

Table S2. Selected bond distances (Å) and angles (deg) for Ru'2,4-D.

Empirical formula	$C_{57}H_{54}CIN_6O_8Ru_2$
Formula weight	1188.65
Temperature/K	100.0
Crystal system	triclinic
Space group	P-1
a/Å	10.3478(4)
b/Å	16.5979(7)
c/Å	16.8204(8)
α/°	110.212(2)
β/°	91.855(2)
γ/°	104.522(2)
Volume/Å ³	2601.8(2)
Z	2
ρ _{calc} g/cm ³	1.517
µ/mm⁻¹	5.678
F(000)	1214.0
Crystal size/mm ³	$0.21 \times 0.04 \times 0.02$
Radiation	CuKα (λ = 1.54178)
20 range for data collection/°	5.648 to 136.49
Index ranges	-11 ≤ h ≤ 12, -19 ≤ k ≤ 19, -20 ≤ l ≤ 20
Reflections collected	99612
Independent reflections	9537 [R _{int} = 0.1196, R _{sigma} = 0.0536]
Data/restraints/parameters	9537/0/673
Goodness-of-fit on F ²	1.041
Final R indexes [I>=2σ (I)]	$R_1 = 0.0524$, $wR_2 = 0.1112$
Final R indexes [all data]	$R_1 = 0.0854, wR_2 = 0.1338$
Largest diff. peak/hole / e Å ⁻³	2.55/-1.03

Table S3. Crystal and structure refinement data for Ru'NAA.



Figure S3. Intermolecular interactions found in the structure of Ru'NAA (distances in Å).

Ru1	Ru2	2.3155(6)	
Ru1	Cl1	2.395(1)	
Ru1	01	2.079(4)	
Ru1	N1	2.107(5)	
Ru1	N3	2.083(4)	
Ru1	N5	2.091(5)	
Ru2	02	2.066(4)	
Ru2	N2	2.046(4)	
Ru2	N4	2.024(4)	
Ru2	N6	2.039(5)	
Ru1	Ru2	Cl1	173.92(4)
Ru1	01	N3	177.0(2)
Ru1	N1	N5	175.5(2)
Ru2	02	N4	178.5(2)
Ru2	N2	N6	176.2(2)

Table S4. Selected bond distances (Å) and angles (deg) for Ru'NAA.

2. MASS SPECTROMETRY



Figure S4. ESI⁺ spectra of **Ru'IAA**, **Ru'2,4-D** and **Ru'NAA** (left). Enlargement of the peak corresponding to [M-CI]⁺ base peak (centre) and calculated spectra for the [M-CI]⁺ fragments (right) of **Ru'IAA** (top), **Ru'2,4-D** (middle) and **Ru'NAA** (bottom). Nominal molecular masses and distribution isotopes were calculated with the MASAS¹ software.

¹ F. A. Urbanos. Software MASAS, Universidad Complutense de Madrid: Madrid, 2002.



Figure S8. ESI⁻ analyses of **Ru'2,4-D** (black) and **Ru'2,4-D** after 24 h at pH = 6.5 (pink). Both samples were dissolved in methanol. The second one was slightly acidified with HCl keeping the pH at 6.5. The intensities of both spectra were normalized to their base peak, which is the same in both cases.

3. IR SPECTROSCOPY



Figure S5. IR spectra of $[Ru_2Cl(\mu-DAniF)_3(\mu-O_2CMe)]$ (black), $[Ru_2Cl_2(\mu-DAniF)_3]$ (blue), **Ru'IAA** (pink), **Ru'2,4-D** (blue) and **Ru'NAA** (red). The antisymmetric O-C-O stretching bands are marked with a red asterisk, the symmetric O-C-O stretching bands with a blue asterisk and the N-C-N stretching bands with a black asterisk.

Table S5. Tentative assignment of the most relevant bands (cm⁻¹) in the IR spectra of $[Ru_2Cl(\mu-DPhF)_3(\mu-O_2CMe)]$, $[Ru_2Cl_2(\mu-DPhF)_3]$, $[Ru_2(\mu-DPhF)_3]$, [R

Compound	v N-H	v C _{ar} -H	v _{as} C-H	v _s C-H	v C=C _{ar}	v _{as} O-C-O + v N-C-N	v _s O-C-O	v C-N	Δ (v _{as} - v _s O-C-O)	
[BuaCl(u-DPbE)a(u-OaCMe)]	_	3057	2957	_	1592	1529	1430	1308	99	
		3037	2557		1485	1525	1430	1211	55	
					1607			1292		
[Ru₂Cl(µ-DAniF)₃(µ-O₂CMe)]	-	3005	2957	2835	1578	1540	1438	1212	102	
					1484					
					1591			1312		
[Ru ₂ Cl ₂ (µ-DPhF) ₃]	-	3063	2951	-	1582	1520*	-	1208	-	
					1484			1200		
[Bu ₂ Cl ₂ (u-DAniF) ₂]	_	3042	2011	2834	1602	1530*	-	1294	_	
[1022012([27 1111 75]		5012	2311	2001	1498	1000		1210		
RulAA	3260	3056	2965	-	1595	1530	1408	1317	122	
	5200		2505		1485	1550		1100	1214	
Ru2 4-D	_	3060	2968	-	1592	1547	1426	1317	121	
		5000			1486		1420	1219	121	
BuNAA	_	3056	2954	_	1593	1531	1408	1318	123	
		5050	1486	1486	1486	1400	1220	125		
BullAA	3276	3030	2957	2833	1606	15/10	1/12	1291	138	
		2557	2000	1497	1340	1712	1239	100		
Bu/2 4-D	Ru'2,4-D - 3039 2959 28	2832	1606	1551	1/121	1291	130			
Ru 2,4-D		5039	2939	2052	1497	1991	1721	1210	130	
Bu/NAA		2026	2054	2022	1606	15/1	1407	1292	12/	
NU NAA	- 3036	2554 20	2035	1499	1541	1407	1212	134		

*Corresponds only to v_{as} O-C-O.

4. ELECTRONIC SPECTROSCOPY



Figure S6. Electronic spectra of $[Ru_2Cl(\mu-DAniF)_3(\mu-O_2CMe)]$ (black), Ru'IAA (red), Ru'2,4-D (blue), Ru'NAA (green) and $[Ru_2Cl(\mu-DPhF)_3(\mu-O_2CMe)]$ (purple).

Transition	LMCT π(Cl)→π*(Ru₂)	π(RuO/N,Ru₂)→ π*(Ru₂)	π*(Ru₂)→ σ*(RuO/N)	δ(Ru₂) → π*(Ru₂)
[Ru₂Cl(µ-DPhF)₃(µ-O₂CMe)]	340 sh	515	565 sh	665 sh
RuIAA	335 sh	517	570 sh	640 sh
Ru2,4-D	340 sh	527	570 sh	640 sh
RuNAA	340 sh	525	570 sh	640 sh
[Ru ₂ Cl(μ-DAniF) ₃ (μ-O ₂ CMe)]	380 sh	511	594	690 sh
Ru'IAA	385 sh	498	594	690 sh
Ru'2,4-D	385 sh	478 sh	603	690 sh
Ru'NAA	360 sh	500	590	690 sh

Table S6. Tentative assignment, of the transitions (nm) observed in the electronic spectra of $[Ru_2Cl(\mu-DPhF)_3(\mu-O_2CMe)]$, **Ru1AA**, **Ru2,4-D**, **RuNAA**, $[Ru_2Cl(\mu-DAniF)_3(\mu-O_2CMe)]$, **Ru1AA**, **Ru2,4-D** and **Ru2AA**.²

 ² (a) Lin, C.; Ren, T.; Valente, E. J.; Zubkowski, J. D.; Smith, E. T. *Chem. Lett.* **1997**, *26* (8), 753–754. (b) Barral, M. C.; Herrero, S.; Jiménez-Aparicio, R.; Torres, M. R.; Urbanos, F. A. *Angew. Chem. Int. Ed.* **2005**, *44* (2), 305–307. (c) Chen, W.-Z.; Ren, T. *Organometallics* **2005**, *24* (11), 2660–2669. (d) Bear, J. L.; Han, B.; Huang, S.; Kadish, K. M. *Inorg. Chem.* **1996**, *35* (10), 3012–3021. (e) Castro, M. A.; Roitberg, A. E.; Cukiernik, F. D. *Inorg. Chem.* **2008**, *47* (11), 4682–4690.



Figure S7. Electronic spectra of **RuNAA** (left) and **Ru'NAA** (right) in DMSO/water solution using an HEPES KOH (4-(2-hydroxyethyl)-1-piperazineethanesulfonic acid) buffer at pH 6.5 over 24 h.

5. MAGNETIC MEASUREMENTS



Figure S9. Temperature dependence of the molar susceptibility χ_M (circles) and $\chi_M \cdot T$ (squares) for **Ru'IAA**. Solid lines are the best fit to the data as described in the text.



Figure S10. Temperature dependence of the molar susceptibility χ_M (circles) and $\chi_M \cdot T$ (squares) for **Ru'2,4-D**. Solid lines are the best fit to the data as described in the text.



Figure S11. Temperature dependence of the molar susceptibility χ_M (circles) and $\chi_M \cdot T$ (squares) for **Ru'NAA**. Solid lines are the best fit to the data as described in the text.

Equations employed in the fitting of the magnetic data:

$$\chi_M = \frac{\chi_{\parallel} + 2\chi_{\perp}}{3}$$

Equation S1

$$\chi_{\parallel} = \left(\frac{Ng^2\beta^2}{k_BT}\right) \left[\frac{1+9\exp\left(-2D/k_BT\right)}{4\left(1+\exp\left(-2D/k_BT\right)\right)}\right]$$

Equation S2

$$\chi_{\perp} = \left(\frac{Ng^2\beta^2}{k_BT}\right) \left[\frac{4 + \left(\frac{3k_BT}{D}\right)\left(1 - \exp\left(\frac{-2D}{k_BT}\right)\right)}{4\left(1 + \exp\left(\frac{-2D}{k_BT}\right)\right)}\right]$$

Equation S3

$$\chi_{M}' = \frac{\chi_{M}}{1 - \left(\frac{2zJ}{Ng^{2}\beta^{2}}\right)\chi_{M}}$$

Equation S4

(*N*, *g*, β , and $k_{\rm B}$ have their usual meanings)

6. BIOLOGICAL ASSAYS AND CHEMOMETRICS

2,4-D					
Source	Sum of squares	Df	Mean square	F-ratio	<i>p</i> -value
Main effects (pH)	0.0472109	2	0.0236054	115.81	0.0000
Main effect (time, h)	0.700818	2	0.350409	1710.06	0.0000
Interactions (pH-time)	0.0455153	4	0.0113788	55.82	0.0000
Ru2,4-D					
Source	Sum of squares	Df	Mean square	F-ratio	<i>p</i> -value
Main effects (pH)	0.0528029	2	0.0264014	20.19	0.0000
Main effect (time, h)	0.426837	2	0.213418	163.20	0.0000
Interactions (pH-time)	0.00732122	4	0.0183031	14.00	0.0000
Ru'2,4-D					
Source	Sum of squares	Df	Mean square	F-ratio	<i>p</i> -value
Main effects (pH)	0.0708965	2	0.0354482	202.64	0.0000
Main effect (time, h)	0.74159	2	0.370795	2119.69	0.0000
Interactions (pH-time)	0.0877365	4	0.02119341	125.39	0.0000
NAA					
Source	Sum of squares	Df	Mean square	F-ratio	<i>p</i> -value
Main effects (pH)	0.0659725	2	0.0329862	11.64	0.0009
Main effect (time, h)	0.804324	2	0.402162	141.90	0.0000
Interactions (pH-time)	0.060053	4	0.0150133	5.30	0.0073
RuNAA					
Source	Sum of squares	Df	Mean square	F-ratio	<i>p</i> -value
Main effects (pH)	0.0838757	2	0.041937	233.62	0.0000
Main effect (time, h)	0.355385	2	0.177692	989.84	0.0000
Interactions (pH-time)	0.0923937	4	0.0230984	128.67	0.0000
Ru'NAA	·				
Source	Sum of squares	Df	Mean square	F-ratio	<i>p</i> -value
Main effects (pH)	0.0734904	2	0.0367452	25.392	0.0000
Main effect (time, h)	0.416898	2	0.208449	144.03	0.0000
Interactions (pH-time)	0.0668784	4	0.0167196	11.55	0.0001

 Table S7. Multifactorial ANOVA.³ Influence of pH and time on auxin activity.

³ C. Mongay Fernández, *Quimiometría*, UNIVERSITAT DE VALÈNCIA, 2014.



Figure S12. Multifactor ANOVA plots. Factors evaluated and *p*-values obtained. a) 2,4 D; b) Ru2,4-D; c) Ru'2,4-D; d) NAA; e) RuNAA; f) Ru'NAA.



Figure S13. a) Variation of the auxin activity of NAA as a function of pH and time (the pH time crossdependence is not observed for **RuNAA** and **Ru'NAA**). b) Variation of the auxin activity as a function of pH and time of 2,4-D. c) Variation of the auxin activity as a function of pH and time of **Ru2,4-D**. d) Variation of the auxin activity as a function of pH and time of **Ru'2,4-D**. Intervals have been estimated using the LSD statistic at 95% probability. Auxin activity is measured in 4-MU nmol·min⁻¹·µg⁻¹ protein.