

***In vivo* and *in vitro* studies of  $[M(\eta^6\text{-pseudoerlotinib})_2]^+$  sandwich complexes (M = Re,  $^{99m}\text{Tc}$ )**

Federica Battistin<sup>a\*</sup>, Célia Fernandes<sup>b</sup>, Paula. D. Raposinho<sup>b</sup>, Olivier Blacque<sup>a</sup>, António Paulo<sup>b\*</sup>, Roger Alberto<sup>a</sup>

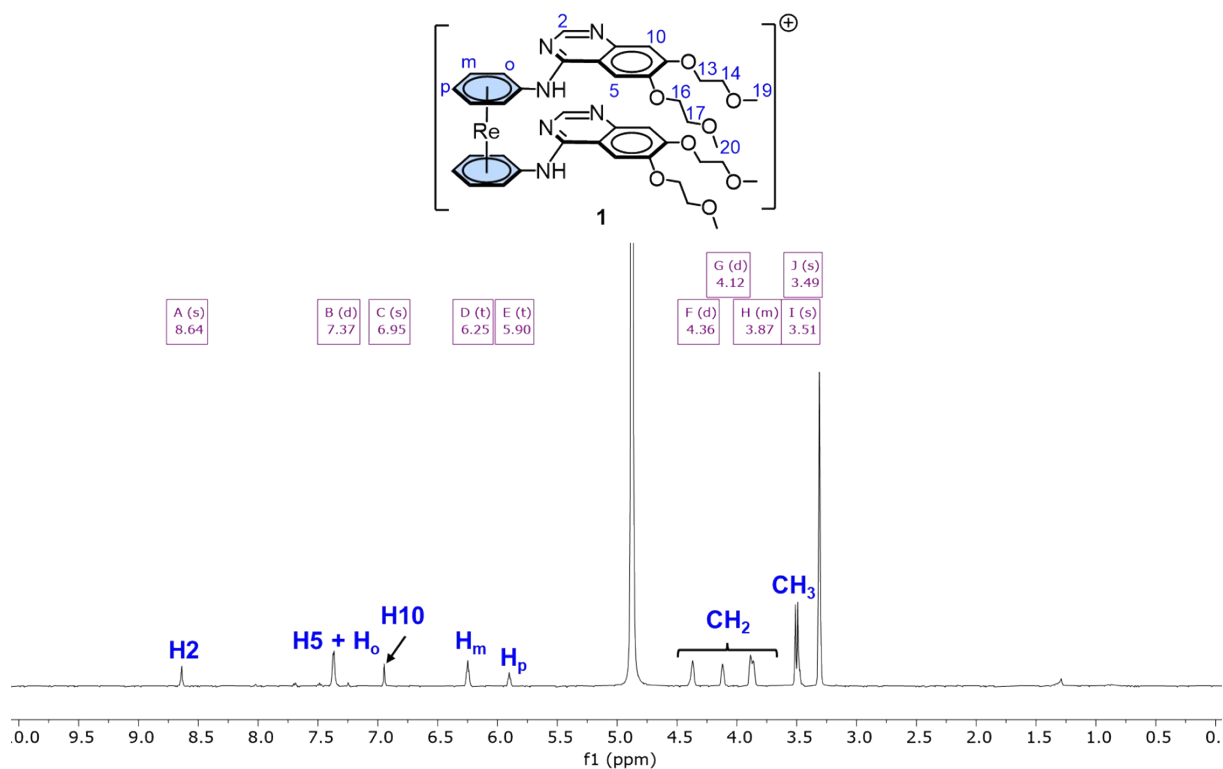
<sup>a</sup> Department of Chemistry, University of Zurich, Winterthurerstr. 190, 8057 Zurich, Switzerland. federica.battistin@chem.uzh.ch

<sup>b</sup> Centro de Ciências e Tecnologias Nucleares, Instituto Superior Técnico, Universidade de Lisboa, Campus Tecnológico e Nuclear, Bobadela, Portugal. apaulo@ctn.tecnico.ulisboa.pt

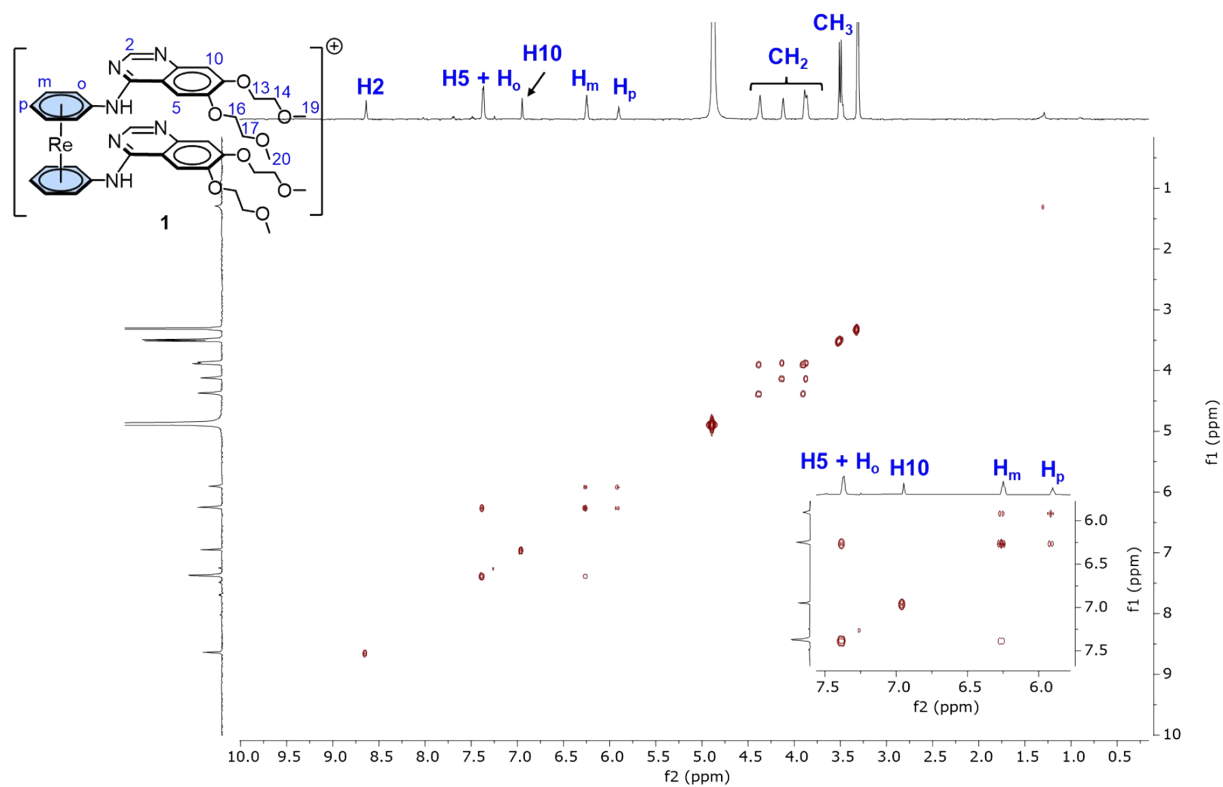
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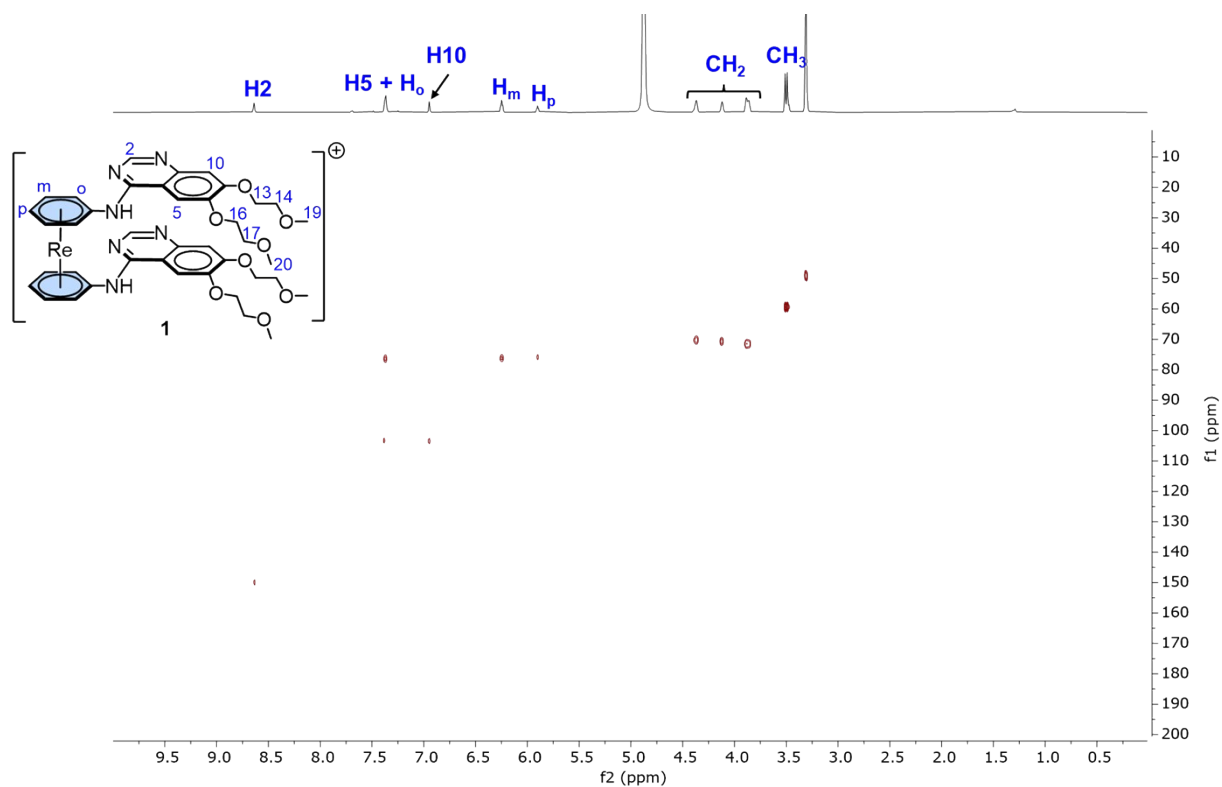
## NMR spectra



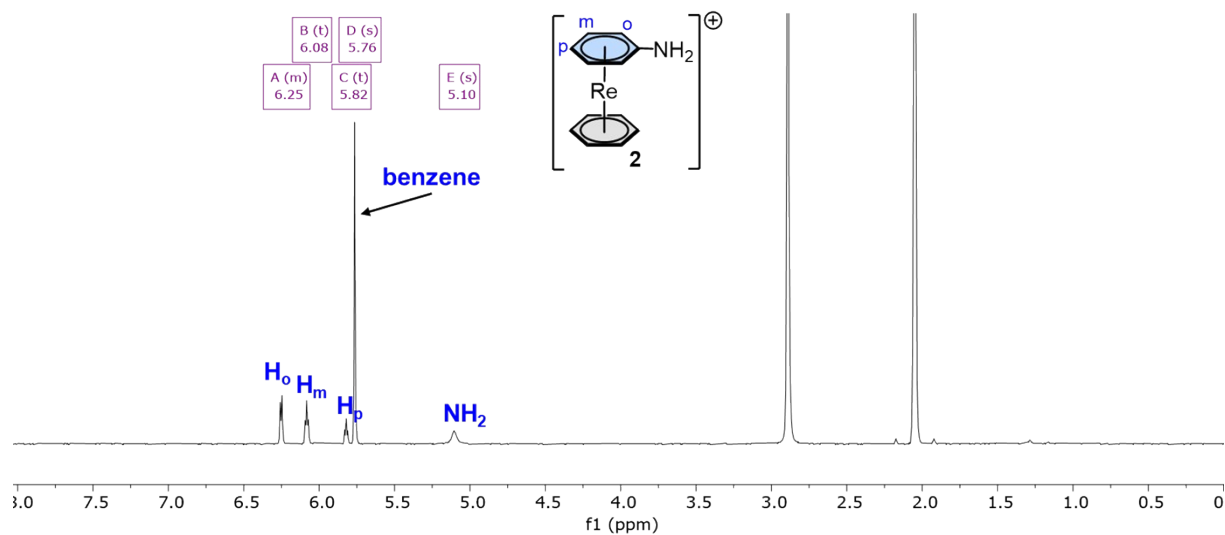
**Figure S1.**  $^1\text{H}$  NMR spectrum in  $\text{CD}_3\text{OD}$  of  $[\text{Re}(\eta^6\text{-pseudoerlotinib})_2]^+$  (1).



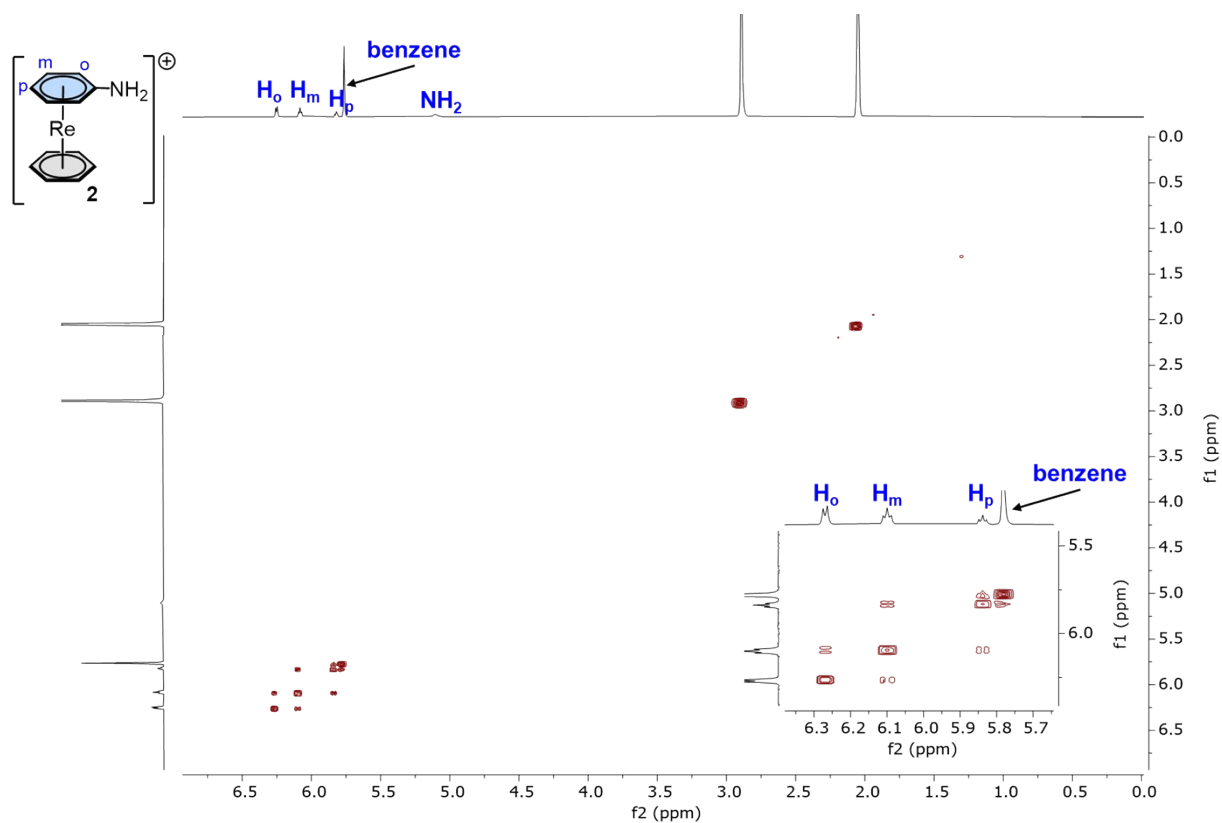
**Figure S2.**  $^1\text{H}$ - $^1\text{H}$  COSY spectrum in  $\text{CD}_3\text{OD}$  of  $[\text{Re}(\eta^6\text{-pseudoerlotinib})_2]^+$  (1).



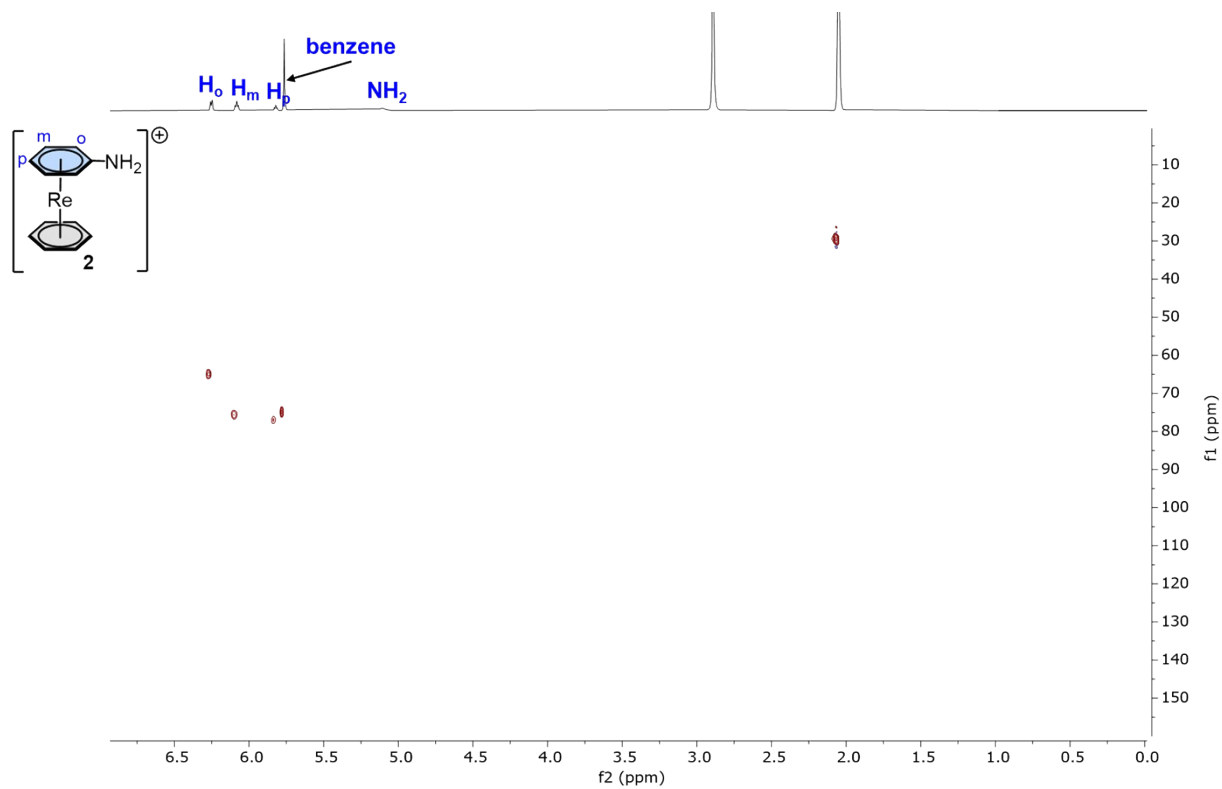
**Figure S3.**  $^1\text{H}$ - $^{13}\text{C}$  HSQC spectrum in  $\text{CD}_3\text{OD}$  of  $[\text{Re}(\eta^6\text{-pseudoerlotinib})_2]^+$  (**1**).



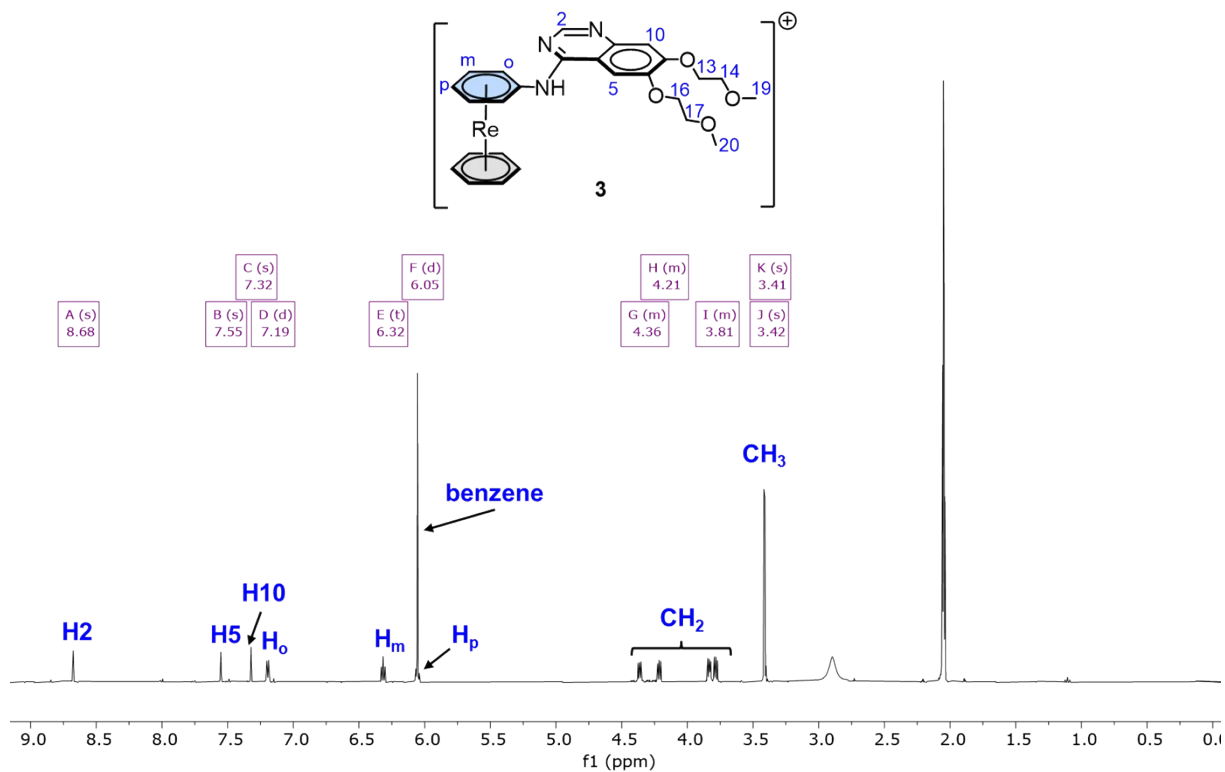
**Figure S4.**  $^1\text{H}$  NMR spectrum in  $\text{acetone-}d_6$  of  $[\text{Re}(\eta^6\text{-bz})(\eta^6\text{-aniline})]^+$  (**2**).



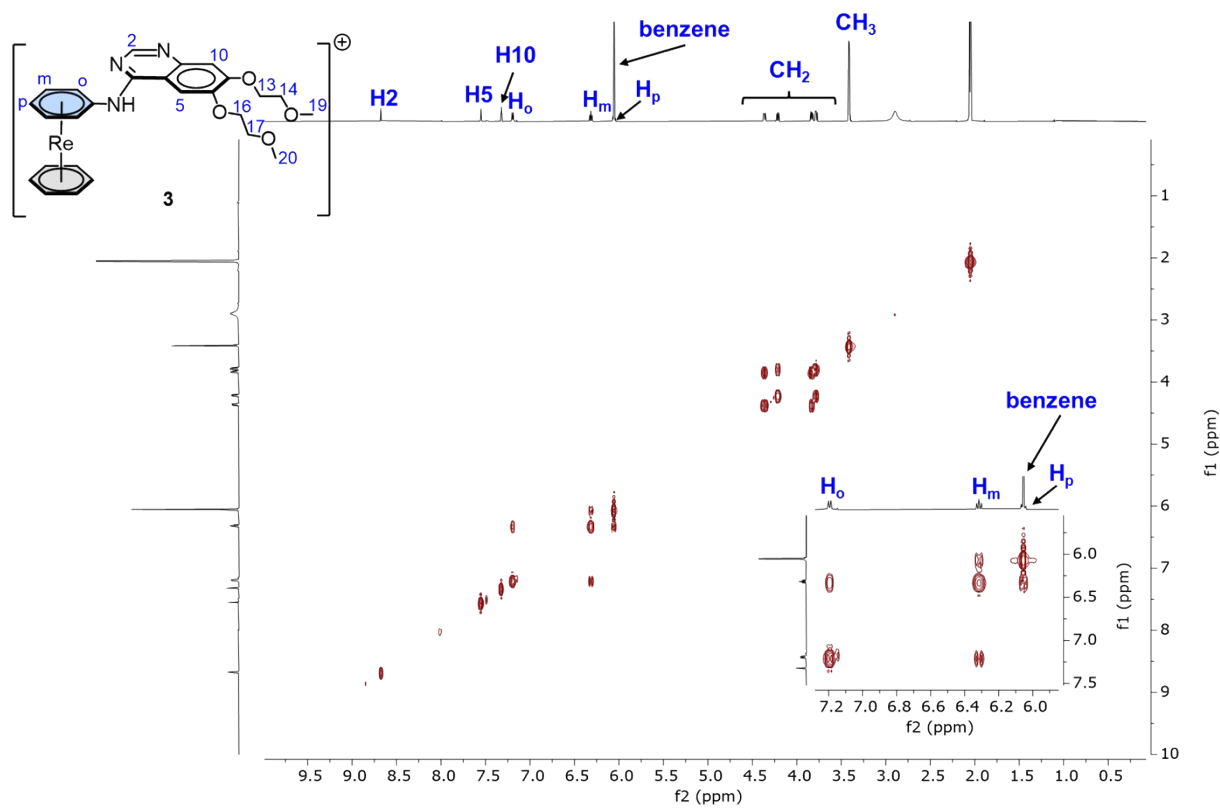
**Figure S5.**  $^1\text{H}$ - $^1\text{H}$  COSY spectrum in acetone- $d_6$  of  $[\text{Re}(\eta^6\text{-bz})(\eta^6\text{-aniline})]^+$  (**2**).



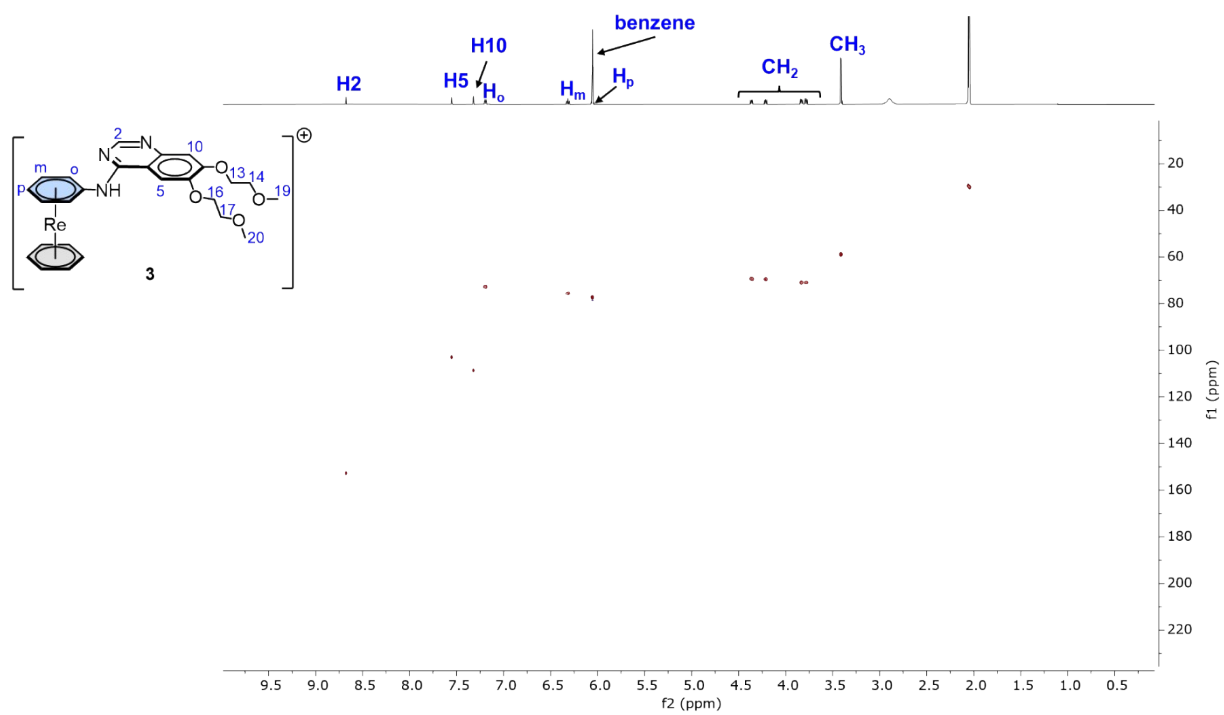
**Figure S6.**  $^1\text{H}$ - $^{13}\text{C}$  HSQC spectrum in acetone- $d_6$  of  $[\text{Re}(\eta^6\text{-bz})(\eta^6\text{-aniline})]^+$  (**2**).



**Figure S7.**  $^1\text{H}$  NMR spectrum in acetone- $d_6$  of  $[\text{Re}(\eta^6\text{-bz})(\eta^6\text{-pseudoerlotinib})]^+$  (**3**).

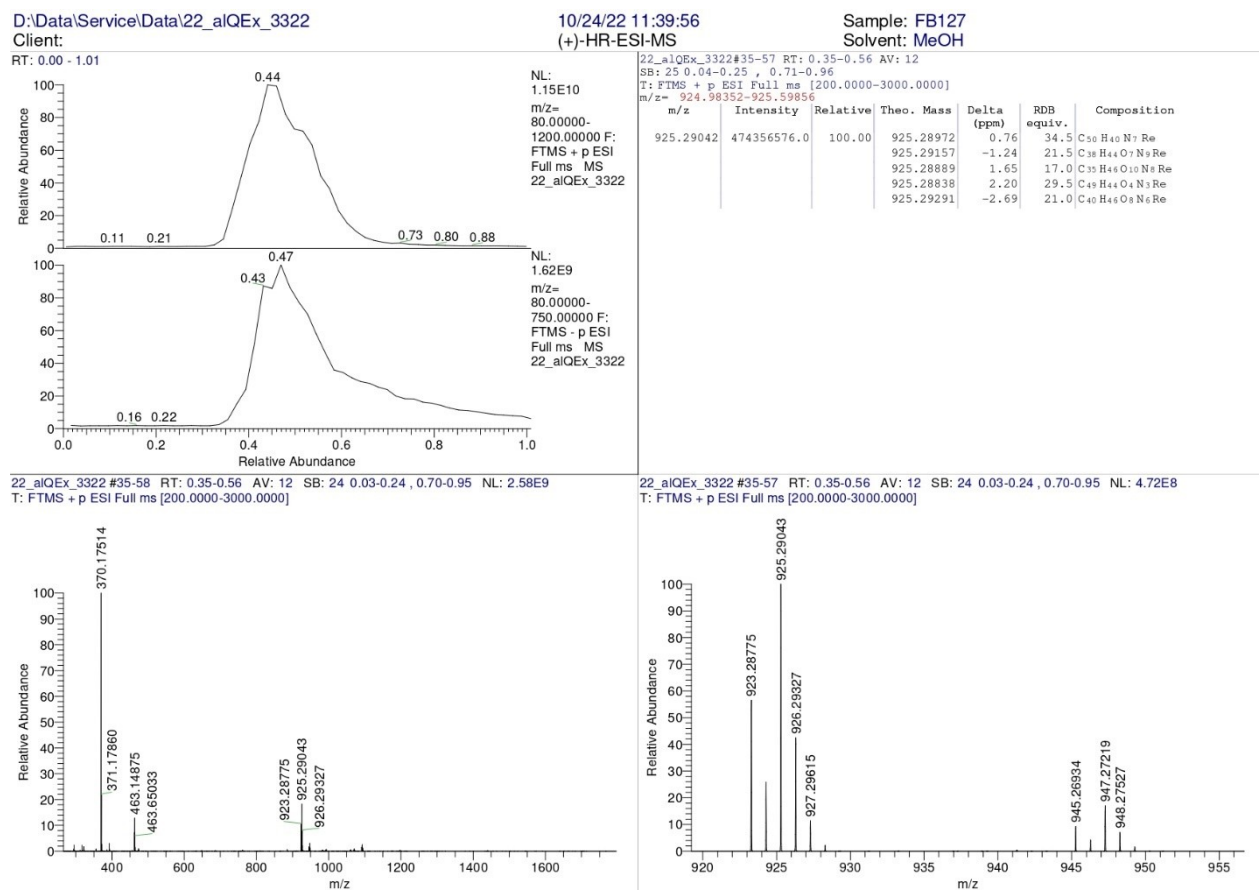


**Figure S8.**  $^1\text{H}$ - $^1\text{H}$  COSY spectrum in acetone- $d_6$  of  $[\text{Re}(\eta^6\text{-bz})(\eta^6\text{-pseudoerlotinib})]^+$  (**3**).



**Figure S9.**  $^1\text{H}$ - $^{13}\text{C}$  HSQC spectrum in acetone- $d_6$  of  $[\text{Re}(\eta^6\text{-bz})(\eta^6\text{-pseudoerlotinib})]^+$  (**3**).

# HR-ESI spectra



**Figure S10.** HR-ESI-MS spectrum of  $[\text{Re}(\eta^6\text{-pseudoerlotinib})_2]^+$  (**1**).

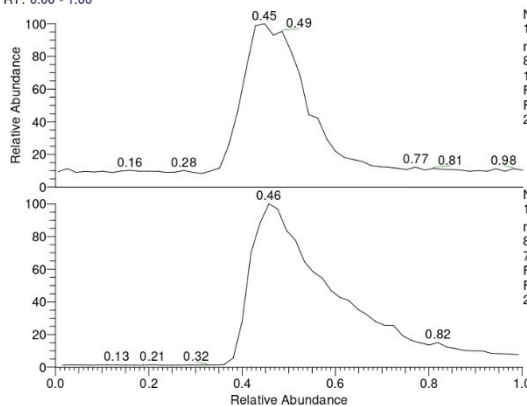


D:\Data\Service\Data\22\_aIQEx\_3299  
Client:

10/24/22 10:53:53  
(+)-HR-ESI-MS

Sample: FB123  
Solvent: MeOH

RT: 0.00 - 1.00

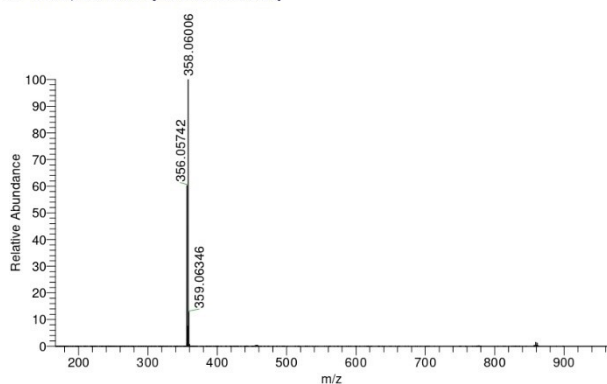


NL:  
1.23E10  
m/z=  
80.00000-  
1200.00000 F:  
FTMS + p ESI  
Full ms MS  
22\_aIQEx\_3299

22\_aIQEx\_3299#37-59 RT: 0.35-0.56 AV: 12  
SB: 25 0.03-0.24 , 0.70-0.94  
T: FTMS + p ESI Full ms [100.0000-1500.0000]  
m/z= 357.97477-358.16251

m/z	Intensity	Relative	Theo. Mass	Delta (ppm)	RDB equiv.	Composition
358.06006	3566684416.0	100.00	358.06000	0.17	7.0	C <sub>12</sub> H <sub>13</sub> NRe
			358.06053	-1.32	2.5	C <sub>5</sub> H <sub>12</sub> O <sub>2</sub> N <sub>4</sub> BRe
			358.05919	2.43	3.0	C <sub>3</sub> H <sub>10</sub> ON <sup>+</sup> BRe

22\_aIQEx\_3299 #37-59 RT: 0.35-0.56 AV: 12 SB: 26 0.03-0.24 , 0.70-0.94 NL: 3.54E9  
T: FTMS + p ESI Full ms [100.0000-1500.0000]



22\_aIQEx\_3299 #37-60 RT: 0.35-0.56 AV: 12 SB: 25 0.03-0.24 , 0.70-0.94 NL: 3.54E9  
T: FTMS + p ESI Full ms [100.0000-1500.0000]

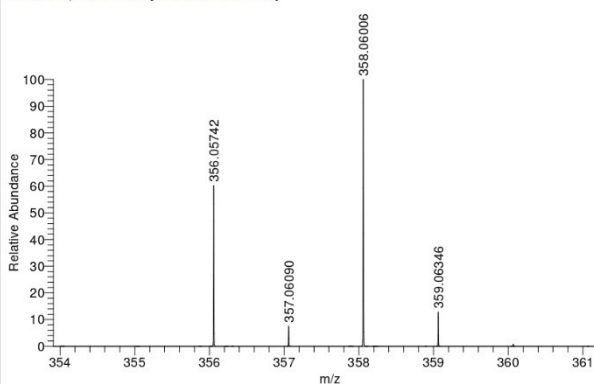


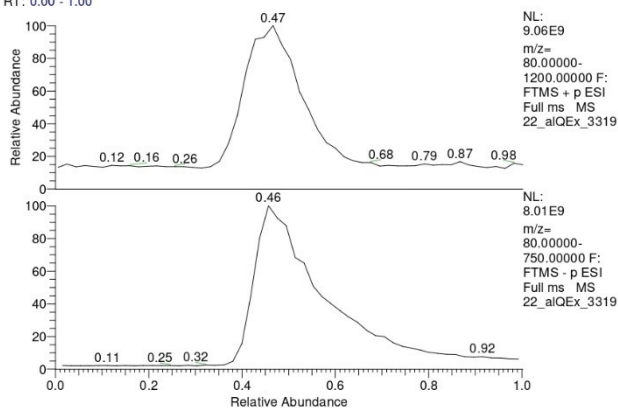
Figure S11. HR-ESI-MS spectrum of  $[\text{Re}(\eta^6\text{-bz})(\eta^6\text{-aniline})]^+$  (**2**).

D:\Data\Service\Data\22\_aIQEx\_3319  
Client:

10/24/22 11:33:52  
(+)-HR-ESI-MS

Sample: FB124  
Solvent: MeOH

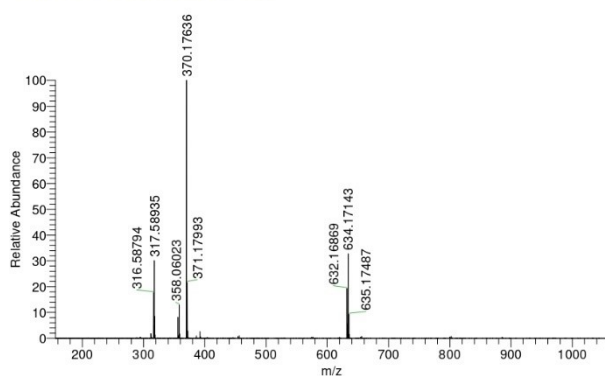
RT: 0.00 - 1.00



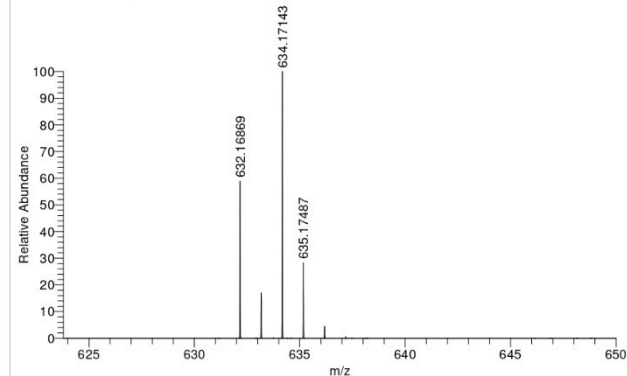
22\_aIQEx\_3319#37-61 RT: 0.35-0.58 AV: 13  
SB: 25 0.03-0.24 , 0.70-0.94  
T: FTMS + p ESI Full ms [100.0000-1500.0000]  
m/z= 633.92064-634.46847

m/z	Intensity	Relative	Theo. Mass	Delta (ppm)	RDB equiv.	Composition
634.17143	490399584.0	100.00	634.17101	0.66	14.0	C <sub>26</sub> H <sub>29</sub> O <sub>4</sub> N <sub>3</sub> Re
			634.17235	-1.45	19.0	C <sub>27</sub> H <sub>23</sub> N <sub>7</sub> Re
			634.17235	-1.46	13.5	C <sub>28</sub> H <sub>31</sub> O <sub>3</sub> Re
			634.16967	2.78	14.5	C <sub>24</sub> H <sub>27</sub> O <sub>3</sub> N <sub>6</sub> Re

22\_aIQEx\_3319 #37-61 RT: 0.35-0.58 AV: 13 SB: 24 0.03-0.24 , 0.70-0.95 NL: 1.49E9  
T: FTMS + p ESI Full ms [100.0000-1500.0000]

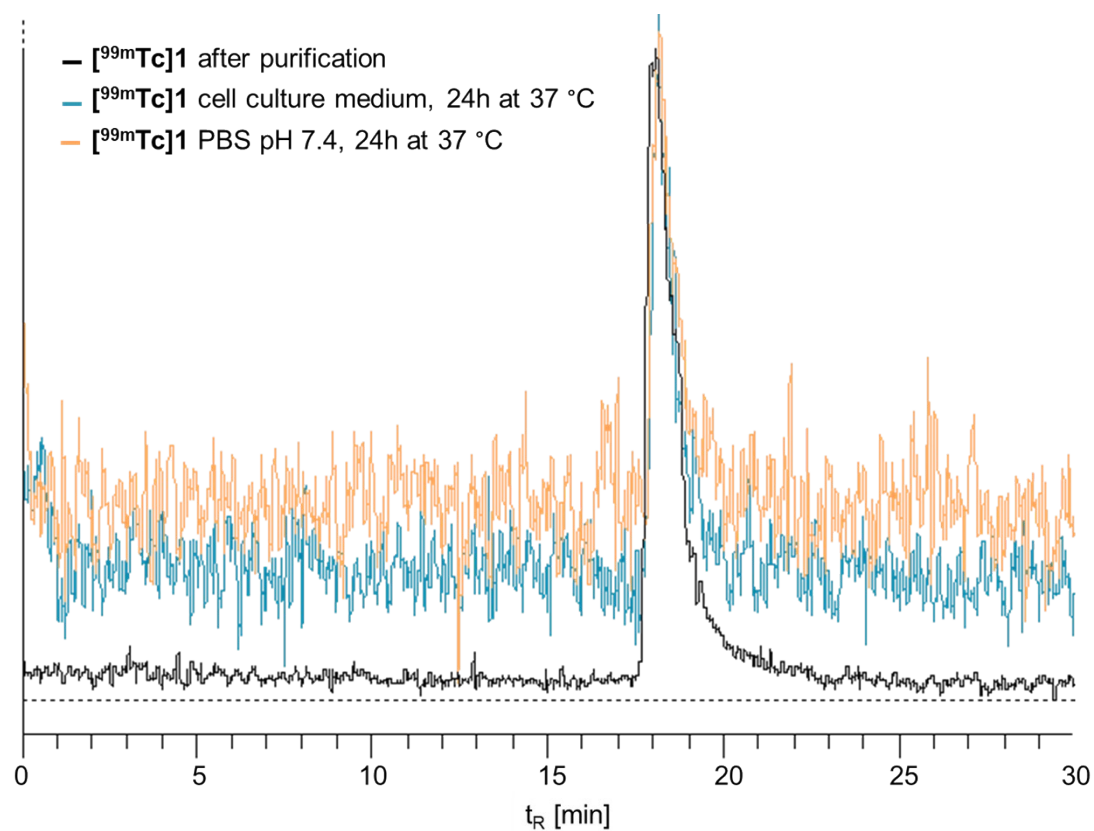


22\_aIQEx\_3319 #37-61 RT: 0.35-0.58 AV: 13 SB: 24 0.03-0.24 , 0.70-0.95 NL: 4.89E8  
T: FTMS + p ESI Full ms [100.0000-1500.0000]



**Figure S12.** HR-ESI-MS spectrum of  $[\text{Re}(\eta^6\text{-bz})(\eta^6\text{-pseudoerlotinib})]^+$  (**3**).

*in vitro* stability of [<sup>99m</sup>Tc]1



**Figure S13.**  $\gamma$ -traces of [<sup>99m</sup>Tc]1 after purification (black line), after incubation in cell culture medium for 24h at 37 °C (blue line), after incubation in PBS at pH 7.4 for 24h at 37 °C (blue line).

## X-ray crystallography

Single-crystal X-ray diffraction data of **1**, **2**, **3** were collected at 160(1) K on a Rigaku OD XtaLAB Synergy, Dualflex, Pilatus 200K diffractometer using a single wavelength X-ray source (Cu K $\alpha$  radiation:  $\lambda = 1.54184 \text{ \AA}$ ) using the copper X-ray radiation ( $\lambda = 1.54184 \text{ \AA}$ ) from a dual wavelength X-ray source and an Oxford Instruments Cryojet XL cooler. The selected suitable single crystal was mounted using polybutene oil on a flexible loop fixed on a goniometer head and immediately transferred to the diffractometer. Pre-experiment, data collection, data reduction and analytical absorption correction<sup>1</sup> were performed with the program suite CrysAlisPro.<sup>2</sup> Using Olex2,<sup>3</sup> the structure was solved with the SHELXT<sup>4</sup> small molecule structure solution program and refined with the SHELXL2018/3 program package<sup>5</sup> by full-matrix least-squares minimization on F<sup>2</sup>. PLATON<sup>6</sup> was used to check the result of the X-ray analysis. CCDC 2293185-2293187 contain the supplementary crystallographic data for this paper. These data are provided free of charge by The Cambridge Crystallographic Data Centre via [www.ccdc.cam.ac.uk/structures](http://www.ccdc.cam.ac.uk/structures).

**Table S1.** Crystal data and data collection of complexes [1](PF<sub>6</sub>)Cl<sub>2</sub>, [2](PF<sub>6</sub>), and [3](TFA).

	[Re( $\eta^6$ -pseudoerlotinib) <sub>2</sub> ] (PF <sub>6</sub> )Cl <sub>2</sub> ·2(H <sub>2</sub> O) ([1]PF <sub>6</sub> Cl <sub>2</sub> ·2(H <sub>2</sub> O))	[Re( $\eta^6$ -bz)( $\eta^6$ -aniline)]PF <sub>6</sub> ([2]PF <sub>6</sub> )	4[Re( $\eta^6$ -bz)( $\eta^6$ -pseudoerlotinib)] 4(PF <sub>6</sub> )·2.25(H <sub>2</sub> O)·0.667(CH <sub>2</sub> Cl <sub>2</sub> ) (4[3](PF <sub>6</sub> )·2.25(H <sub>2</sub> O)·0.667(CH <sub>2</sub> Cl <sub>2</sub> ))
Empirical formula	C <sub>40</sub> H <sub>52</sub> Cl <sub>2</sub> F <sub>6</sub> N <sub>6</sub> O <sub>10</sub> PRE	C <sub>12</sub> H <sub>13</sub> F <sub>6</sub> NPRE	C <sub>104.67</sub> H <sub>120.43</sub> Cl <sub>1.33</sub> F <sub>24</sub> N <sub>12</sub> O <sub>17.55</sub> P <sub>4</sub> Re <sub>4</sub>
Formula weight	1178.94	502.40	3199.22
Temperature/K	160(1)	160(1)	160.0(1)
Crystal system	triclinic	orthorhombic	triclinic
Space group	P-1	Immm	P-1
a/Å	11.06150(10)	8.09180(10)	11.04098(7)
b/Å	13.8305(3)	12.2476(2)	14.03029(9)
c/Å	16.0904(3)	7.01550(10)	19.54390(14)
$\alpha$ /°	73.359(2)	90	77.5660(6)
$\beta$ /°	71.7530(10)	90	84.5485(6)
$\gamma$ /°	75.7690(10)	90	83.5136(5)
Volume/Å <sup>3</sup>	2206.68(7)	695.272(17)	2929.77(4)
Z	2	2	1
$\rho_{\text{calc}}/\text{cm}^3$	1.774	2.400	1.813
$\mu/\text{mm}^{-1}$	7.636	18.792	9.607
F(000)	1184.0	472.0	1571.0
Crystal size/mm <sup>3</sup>	0.15 × 0.04 × 0.03	0.18 × 0.12 × 0.09	0.15 × 0.09 × 0.01
Radiation	Cu K $\alpha$ ( $\lambda$ = 1.54184)	Cu K $\alpha$ ( $\lambda$ = 1.54184)	Cu K $\alpha$ ( $\lambda$ = 1.54184)
2 $\theta$ range for data collection/°	5.94 to 148.998	13.114 to 148.64	7.15 to 149.004
Index ranges	-12 ≤ h ≤ 13, -17 ≤ k ≤ 17, -20 ≤ l ≤ 20	-10 ≤ h ≤ 9, -15 ≤ k ≤ 15, -8 ≤ l ≤ 8	-13 ≤ h ≤ 13, -17 ≤ k ≤ 14, -23 ≤ l ≤ 24
Reflections collected	46323	3744	61434
Independent reflections	8981 [R <sub>int</sub> = 0.0251, R <sub>sigma</sub> = 0.0172]	436 [R <sub>int</sub> = 0.0280, R <sub>sigma</sub> = 0.0112]	11918 [R <sub>int</sub> = 0.0237, R <sub>sigma</sub> = 0.0171]
Data/restraints/parameters	8981/0/627	436/108/72	11918/409/832
Goodness-of-fit on F <sup>2</sup>	1.130	1.172	1.050
Final R indexes [I ≥ 2 $\sigma$ (I)]	R <sub>1</sub> = 0.0287, wR <sub>2</sub> = 0.0690	R <sub>1</sub> = 0.0162, wR <sub>2</sub> = 0.0396	R <sub>1</sub> = 0.0293, wR <sub>2</sub> = 0.0755
Final R indexes [all data]	R <sub>1</sub> = 0.0292, wR <sub>2</sub> = 0.0692	R <sub>1</sub> = 0.0162, wR <sub>2</sub> = 0.0396	R <sub>1</sub> = 0.0306, wR <sub>2</sub> = 0.0764
Largest diff. peak/hole / e Å <sup>-3</sup>	1.30/-1.61	0.47/-0.70	1.09/-1.52
CCDC Nr.	2293186	2293185	2293187

## Bibliography:

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- (2) CrysAlisPro (version 1.171.42.57a). Rigaku Oxford Diffraction Ltd. Yarnton, Oxfordshire, England, 2022.
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- (4) Sheldrick, G. M. SHELXT—Integrated Space-Group and Crystal-Structure Determination. *Acta Crystallogr. Sect. A Found. Adv.* **2015**, *71* (1), 3–8.
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- (6) Spek, A. L. Structure Validation in Chemical Crystallography. *Acta Crystallogr. Sect. D Biol. Crystallogr.* **2009**, *65* (2), 148–155.