

Synthesis and crystal structure of *fac*-[3-bromo-6-(1*H*-pyrazol-1-yl- κ N²)pyridazine- κ N¹]tricarbonylchloridorhenium dimethylformamide monosolvate

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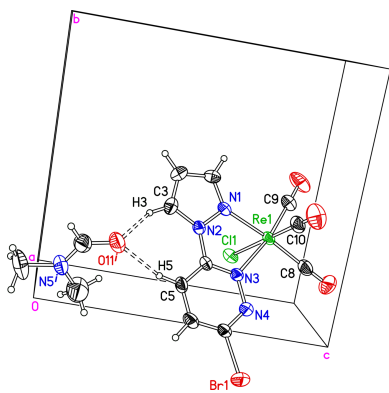
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The crystal structure of the title rhenium(I) complex [ReCl(C₇H₅BrN₄)(CO)₃]-C₃H₇NO or *fac*-[(pypyrBr- κ^2 N,N)Re(CO)₃Cl]-DMF [pypyrBr = 3-bromo-6-(1*H*-pyrazol-1-yl)pyridazine] is reported. The compound was synthesized by reacting Re(CO)₅Cl with pypyrBr [3-bromo-6-(1*H*-pyrazol-1-yl)pyridazine] in toluene under reflux, yielding orange crystals upon purification and crystallization. Structural analysis reveals a distorted octahedral coordination environment around the Re^I center, comprising three facial carbonyl ligands, one chloride, and the bidentate pypyrBr ligand. The ligand exhibits near planarity with minimal torsional deviation, and the N—Re—N bite angle is 73.50 (12)°. The complex crystallizes as a 1:1 dimethylformamide solvate, consolidated by hydrogen bonding between DMF and the ligand. A Cambridge Structural Database search confirmed the novelty of this structure, as no prior reports exist for pypyrBr or its metal complexes. Spectroscopic characterization (¹H NMR, ¹³C NMR, IR) and elemental analysis support the proposed structure. This work expands the family of rhenium(I) tricarbonyl complexes with pyrazolyl-pyridazine ligands, relevant for photophysical and coordination chemistry applications.

1. Chemical context

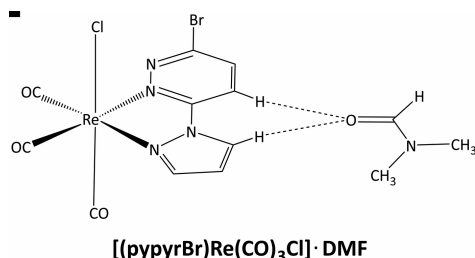
Pyrazolyl-pyridazine derivatives are versatile multidentate and chelating ligands, which are appealing candidates for the synthesis of metal complexes for diverse applications. From the synthetic point of view, their synthesis and derivatization are relatively easy. They are also planar, with limited conformational flexibility, which diminishes non-radiative deactivation paths in photophysical applications (Pizarro *et al.*, 2018). Examples of structurally determined 6-1*H*-pyrazolyl-3-halopyridazine are limited to 3-chloro-6-(1*H*-pyrazol-1-yl)pyridazine (Ather *et al.*, 2010*a*), 3-chloro-6-(3,5-dimethyl-1*H*-pyrazol-1-yl)pyridazine (Ather *et al.*, 2010*c*), ethyl 5-amino-1-(6-chloropyridazin-3-yl)-1*H*-pyrazole-4-carboxylate methyl (Ather *et al.*, 2010*b*), 5-(2-((*t*-butoxycarbonyl)amino)ethyl)-1-(6-chloropyridazin-3-yl)-1*H*-pyrazole-4-carboxylate (Kralj *et al.*, 2009) and 3-chloro-6-(4-chloro-3,5-dimethyl-1*H*-pyrazol-1-yl)pyridazine but to the best of our knowledge no structural determination for 3-bromo-6-(1*H*-pyrazol-1-yl)pyridazine (pypyrBr).

Examples of 6-1*H*-pyrazolyl-3-halopyridazine coordinated to transition metals have been described previously: chloro-[3-chloro-6-(1*H*-pyrazol-1-yl)pyridazine]-[1,2,3,4,5,6- η^6 -1-methyl-4-(propan-2-yl)benzene]ruthenium(II) tetrafluoroborate (Mambanda *et al.*, 2022), chloro-[3-chloro-6-(pyrazol-1-yl)-



pyridazine-*N,N'*]-(η^6 -*p*-cymene)ruthenium(II) hexafluorophosphate (Gupta *et al.*, 2009), diaquabis[3-chloro-6-(pyrazol-1-yl)pyridazine]copper(II) dinitrate (Blake *et al.*, 1998*b*) and (μ_3 -oxo)[μ_2 -3-chloro-6-(pyrazol-1-yl)pyridazine]pentakis(μ_2 -acetato)bis(pyridine)triruthenium(II) hexafluorophosphate (Dai *et al.*, 2009). Examples of rhenium(I) tricarbonyl complexes are limited to [(3-bromo-6-(1*H*-pyrazol-1-yl)pyridazine)Re(CO)₃Br] (Saldías *et al.*, 2019).

In the present paper, we report the synthesis and structure of [(**pyyrBr**)Re(CO)₃Cl] [**pyyrBr**: 3-bromo-6-(1*H*-pyrazol-1-yl)pyridazine].



2. Structural commentary

Fig. 1 shows a displacement ellipsoid plot of [(**pyyrBr**)Re(CO)₃Cl]. The molecule has a rhenium(I) centre, which displays a non-regular octahedron as coordination environment. The coordination number of six is completed by three carbonyl groups in a facial arrangement, a chloro ligand and a bidentate and chelating molecule of **pyyrBr**. The **pyyrBr** ligand displays a highly planar and *C* conformation as required for coordination, which is reflected in the pyrazolyl-pyridazine torsion angles [N1–N2–C4–C5, 176.8 (3)°, C3–N2–C4–C5, 2.1 (6)°, C3–N2–C4–N3, –177.1 (4)° and N1–N2–C4–N3, –2.4 (5)°] and the rather low value of the biting N1–Re1–N3 angle [73.50 (12)°]. The dihedral

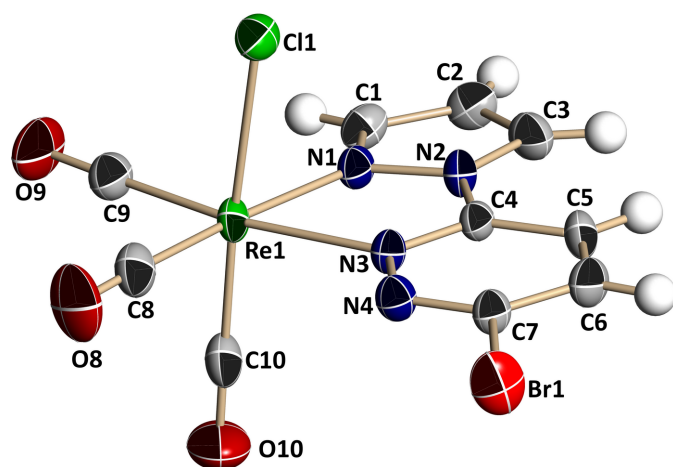


Figure 1
Molecular view of the complex [(**pyyrBr**)Re(CO)₃Cl] showing the partial numbering scheme. Atoms are shown as displacement ellipsoids at the 50% level of probability, except hydrogen, which are shown as arbitrary radii spheres. Solvating dimethylformamide molecule omitted for the sake of clarity.

Table 1
Hydrogen-bond geometry (Å, °).

<i>D</i> –H··· <i>A</i>	<i>D</i> –H	H··· <i>A</i>	<i>D</i> ··· <i>A</i>	<i>D</i> –H··· <i>A</i>
C3–H3···O11 ⁱ	0.93	2.41	3.251 (6)	150
C5–H5···O11 ⁱ	0.93	2.33	3.206 (5)	158
C2–H2···Cl1 ⁱⁱ	0.93	2.79	3.556 (4)	140
C6–H6···Cl1 ⁱⁱⁱ	0.93	2.75	3.614 (4)	154

Symmetry codes: (i) $x - 1, y, z$; (ii) $-x + 1, -y + 1, -z + 1$; (iii) $-x, -y, -z + 1$.

angle between these two planar pyrazolyl and pyridazine groups is 4.0 (2)°.

3. Supramolecular features

As previously commented, the rhenium(I)tricarbonyl molecule crystallizes as a 1:1 dimethylformamide solvate, with no evidence of partial occupancy. Fig. 2 shows how the carbonyl oxygen atom of the dimethylformamide molecule accepts two hydrogen bonds to pyrazolyl, C3–H3, and pyridazine, C5–H5, with $D\cdots A = 3.251$ (6) and 3.206 (5) Å respectively (Table 1).

4. Database survey

A search on the Cambridge Structural Database (v 6.00 updated to August 2025; Groom *et al.*, 2016) found no structural report for 3-bromo-6-(1*H*-pyrazol-1-yl)pyridazine or its transition-metal complexes.

The chloro analogue was reported as DUSZIB (Ather *et al.*, 2010*a*). The 3,5-dimethylpyrazolyl analogue was reported as KUYZIO (Ather *et al.*, 2010*c*). Transition-metal complexes of

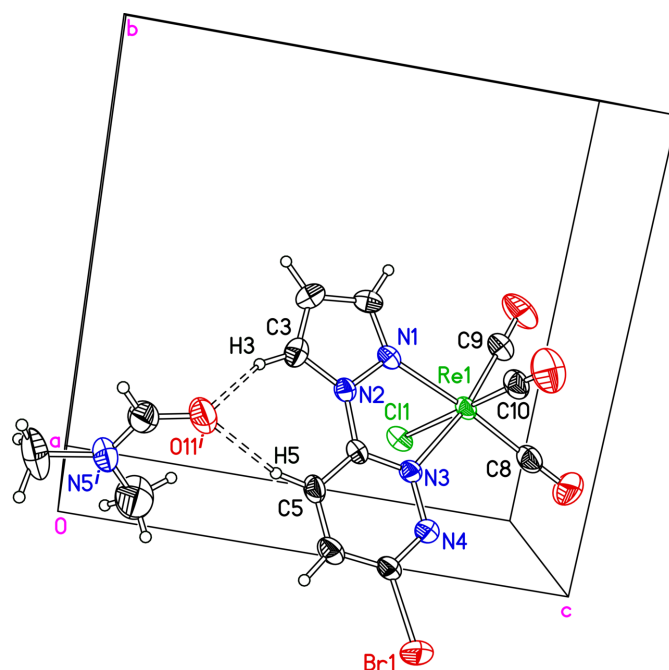
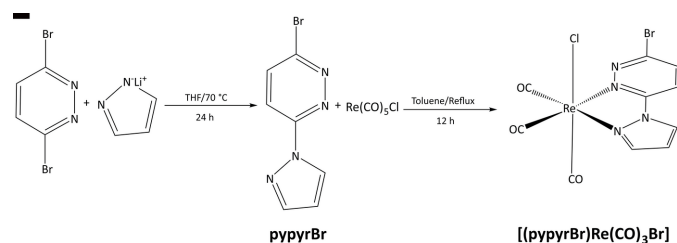


Figure 2
Molecular view of the hydrogen bond between [(**pyyrBr**)Re(CO)₃Cl] and solvating dimethylformamide.

these two chloro analogues are found for copper(II) [COLPOJ and COLPUP (Li *et al.*, 2008), PUCYAN, PUCYER, PUCYIV, PUCYOB and PUCYUH (Blake *et al.*, 1998b), QEPROS and QEPRUY (Blake *et al.*, 1998a)], cobalt(II) (DOXPEM and DOXPIQ; An *et al.*, 2009), nickel(II) (DOXPOW; Li *et al.*, 2008), ruthenium(II/III) [FEWGOH, FEWGUN and FEWHAU (Mambanda *et al.*, 2022), IHEZEB and IHEZIF (Gupta *et al.*, 2009), WUCVAS (Dai *et al.*, 2009)] and rhodium(III) (MUSFOW; Gupta *et al.*, 2010).

5. Synthesis and crystallization

pppyrBr. A solution of 2.83 g (0.0416 mol) of pyrazole in tetrahydrofuran (THF) was prepared under an inert atmosphere. To this solution, 0.2893 g (0.0416 mol) of lithium metal was added in a 1:1 molar ratio. The mixture was stirred at 343 K for approximately 3 h until the lithium was completely dissolved. The excess of unreacted lithium metal was then removed, and a solution of 9.917 g (0.0416 mol) of 3,6-dibromopyridazine in THF was added dropwise. The reaction mixture was stirred at 343 K for 24 h, as shown in the reaction scheme. The resulting solid was washed with cold diethyl ether to yield the pure product **pppyrBr** (4.4238 g), corresponding to an approximate yield of 47%. Elemental analysis: calculated (%): C, 37.36; H, 2.24; N, 24.90. Found (%): C, 37.60; H, 2.43; N, 24.30. ^1H NMR (400 MHz, CDCl_3 , δ ppm): 8.10 (*d*, $J = 9.2$ Hz, 1H, 5) and 7.75 (*d*, $J = 9.2$ Hz, 1H, 6) were assigned to the pyridazine protons, while the signals at 8.71 (*d*, $J = 6.2$ Hz, 1H, 3), 7.81 (*d*, $J = 5.8$ Hz, 1H, 1), and 6.55 (*dd*, $J = 2.7, 1.7$ Hz, 1H, 2) correspond to the pyrazolyl fragment. ^{13}C NMR (101 MHz, CDCl_3 , δ ppm): 154.00 and 145.13 ppm were assigned to pyridazine C7 and C4, bearing the bromo and pyrazolyl groups, respectively. The signals at 143.51 and 133.74 ppm correspond to the nitrogen-bearing ring carbons C5 and C6. The resonances at 127.56, 119.74, and 109.36 ppm were attributed to the pyrazolyl carbons C3, C1, and C2, respectively. IR: $\nu_{\text{C-H}}(\text{arom})$: 3060, 3121, 3140, $\nu_{\text{C=N/C=C}}(\text{rings})$: 1570, 1520, 1455 and 1386 $\nu_{\text{C-Br}}$: 606.



[pppyrBr]Re(CO)₃Cl. The compound was synthesized by the reaction between pentacarbonylchlororhenium(I) and the ligand 3-bromo-6-(1*H*-pyrazol-1-yl)pyridazine (**pppyrBr**) in a 1:1 molar ratio, as shown in the reaction scheme. A solution of $\text{Re}(\text{CO})_5\text{Cl}$ (500 mg, 1.38 mmol) in toluene was prepared in a round-bottom flask equipped for reflux. To this solution, **pppyrBr** (312.4 mg, 1.38 mmol), dissolved in toluene, was added dropwise under an inert atmosphere. The reaction

mixture was refluxed for 12 h. After completion, the solvent was removed under reduced pressure. The crude product was purified by column chromatography using a mixture of dichloromethane/ethyl acetate (2:1) as the mobile phase, obtaining 350 mg of the product (approx. 48% yield). The pure product was crystallized from a THF/DMF (10:1) mixture, affording orange **[(pppyrBr)Re(CO)₃Cl]·C₃H₇NO** crystals. Elemental analysis: calculated (%): C, 25.86; H, 2.00; N, 11.60. Found (%): C, 25.88; H, 2.04; N, 11.29%. The ^1H NMR spectrum in $\text{DMSO-}d_6$ displays the characteristic signals for the aromatic protons. The resonances for the pyridazine ring protons were assigned to the signals at δ 9.24 and 8.58 ppm. Regarding the pyrazole moiety, the protons H1 and H3 appear as doublets at δ 8.75 and 8.65 ppm, respectively, while the central proton H2 is observed at δ 7.04 ppm as a doublet of doublets. ^1H NMR (400 MHz, $\text{DMSO-}d_6$) δ 8.75 (*d*, $J = 9.4$ Hz, 1H, H-pyridazine 5), 9.24 (*d*, $J = 3.1$ Hz, 1H, H-pyrazole 3), 8.58 (*d*, $J = 2.1$ Hz, 1H, H-pyrazole 1), 8.65 (*d*, $J = 9.3$ Hz, 1H, H-pyridazine 6), 7.04 (*dd*, $J = 3.1, 2.1$ Hz, 1H, H-pyrazole 2). The ^{13}C NMR spectrum in $\text{DMSO-}d_6$ confirms the structure of the complex, displaying characteristic signals for the carbonyl ligands in a facial arrangement in the downfield region: a doublet at δ 196.68 ($J = 33.5$ Hz) corresponding to two coupled carbonyls, and a singlet at δ 189.57 ppm assigned to the in-plane carbonyl. Regarding the aromatic moiety, the pyridazine carbons bonded to the bromine atom (C7) and the pyrazole ring (C4) were assigned to δ 152.48 and 145.09 ppm, respectively, while the methine carbons of the same ring (C6/C5) appear at δ 147.36 and 137.29 ppm. Finally, the pyrazole resonances at δ 133.46 and 122.11 ppm were attributed to carbons C3 and C1, respectively, with the central carbon (C2) observed at δ 112.42 ppm. ^{13}C NMR (101 MHz, $\text{DMSO-}d_6$) δ 196.68 (*d*, $J = 33.5$ Hz, CO 8,9), 189.57 (*s*, CO 10), 152.48 (*s*, C-pyridazine 7), 147.36 (*s*, C-pyridazine 6), 145.09 (*s*, C-pyridazine 4), 137.29 (*s*, C-pyridazine 5), 133.46 (*s*, C-pyrazole 3), 122.11 (*s*, C-pyrazole 1), 112.42 (*s*, C-pyrazole 2). IR (cm^{-1}): $\nu_{\text{C-H}}(\text{arom})$ 3090, 3030; $\nu_{\text{C=N}}$: 1670, 1580; $\nu_{\text{C=O}}$: 2025, 1940, 1905.

6. Refinement

Crystal data, data collection and structure refinement details are summarized in Table 2. Hydrogen atoms were placed in calculated positions (C—H: 0.93 Å for aromatic and C—H: 0.96 Å for aliphatic) and refined as riding [$U_{\text{iso}}(\text{H}) = 1.2 U_{\text{eq}}(\text{C})$ for aromatic and $U_{\text{iso}}(\text{H}) = 1.5 U_{\text{eq}}(\text{C})$ for aliphatic].

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Table 2
Experimental details.

Crystal data	
Chemical formula	[ReCl(C ₇ H ₅ BrN ₄)(CO) ₃]-C ₃ H ₇ NO
<i>M_r</i>	603.84
Crystal system, space group	Triclinic, <i>P</i> $\bar{1}$
Temperature (K)	296
<i>a</i> , <i>b</i> , <i>c</i> (Å)	7.2454 (17), 11.136 (3), 11.426 (3)
α , β , γ (°)	90.941 (6), 93.728 (6), 102.141 (6)
<i>V</i> (Å ³)	898.9 (4)
<i>Z</i>	2
Radiation type	Mo <i>K</i> α
μ (mm ⁻¹)	9.16
Crystal size (mm)	0.16 × 0.07 × 0.05
Data collection	
Diffractometer	Bruker CCD area detector
Absorption correction	Multi-scan (<i>SADABS</i> ; Krause <i>et al.</i> , 2015)
<i>T_{min}</i> , <i>T_{max}</i>	0.199, 0.494
No. of measured, independent and observed [<i>I</i> > 2 σ (<i>I</i>)] reflections	7009, 3512, 3277
<i>R_{int}</i>	0.027
(<i>sin</i> θ / λ) _{max} (Å ⁻¹)	0.617
Refinement	
<i>R</i> [<i>F</i> ² > 2 σ (<i>F</i> ²)], <i>wR</i> (<i>F</i> ²), <i>S</i>	0.023, 0.059, 1.04
No. of reflections	3512
No. of parameters	228
H-atom treatment	H-atom parameters constrained
$\Delta\rho_{max}$, $\Delta\rho_{min}$ (e Å ⁻³)	1.14, -1.12

Computer programs: *APEX2* and *SAINT* (Bruker 2012), *SHELXL2014/7* (Sheldrick, 2015), *SHELXTL* (Sheldrick, 2008) and *pubCIF* (Westrip, 2010).

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Computing details

fac-[3-Bromo-6-(1*H*-pyrazol-1-yl- κ N²)pyridazine- κ N¹]tricarbonylchloridorhenium dimethylformamide monosolvate

Crystal data

[ReCl(C₇H₅BrN₄)(CO)₃] \cdot C₃H₇NO

$M_r = 603.84$

Triclinic, $P\bar{1}$

$a = 7.2454$ (17) Å

$b = 11.136$ (3) Å

$c = 11.426$ (3) Å

$\alpha = 90.941$ (6)°

$\beta = 93.728$ (6)°

$\gamma = 102.141$ (6)°

$V = 898.9$ (4) Å³

$Z = 2$

$F(000) = 568$

$D_x = 2.231$ Mg m⁻³

Mo $K\alpha$ radiation, $\lambda = 0.71073$ Å

Cell parameters from 9860 reflections

$\theta = 2.6$ – 29.2 °

$\mu = 9.16$ mm⁻¹

$T = 296$ K

Prism, orange

$0.16 \times 0.07 \times 0.05$ mm

Data collection

Bruker CCD area detector
diffractometer

Radiation source: sealed tube
phi and ω scans

Absorption correction: multi-scan
(SADABS; Krause *et al.*, 2015)

$T_{\min} = 0.199$, $T_{\max} = 0.494$

7009 measured reflections

3512 independent reflections

3277 reflections with $I > 2\sigma(I)$

$R_{\text{int}} = 0.027$

$\theta_{\max} = 26.0$ °, $\theta_{\min} = 1.8$ °

$h = -8 \rightarrow 8$

$k = -13 \rightarrow 13$

$l = -14 \rightarrow 14$

Refinement

Refinement on F^2

Least-squares matrix: full

$R[F^2 > 2\sigma(F^2)] = 0.023$

$wR(F^2) = 0.059$

$S = 1.04$

3512 reflections

228 parameters

0 restraints

Primary atom site location: structure-invariant
direct methods

Secondary atom site location: difference Fourier
map

Hydrogen site location: inferred from
neighbouring sites

H-atom parameters constrained

$w = 1/[\sigma^2(F_o^2) + (0.0299P)^2 + 0.4577P]$

where $P = (F_o^2 + 2F_c^2)/3$

$(\Delta/\sigma)_{\max} = 0.002$

$\Delta\rho_{\max} = 1.14$ e Å⁻³

$\Delta\rho_{\min} = -1.12$ e Å⁻³

Special details

Geometry. All esds (except the esd in the dihedral angle between two l.s. planes) are estimated using the full covariance matrix. The cell esds are taken into account individually in the estimation of esds in distances, angles and torsion angles; correlations between esds in cell parameters are only used when they are defined by crystal symmetry. An approximate (isotropic) treatment of cell esds is used for estimating esds involving l.s. planes.

Least-squares planes (x,y,z in crystal coordinates) and deviations from them (* indicates atom used to define plane)

- 6.3159 (0.0076) x + 6.4497 (0.0188) y + 3.6505 (0.0219) z = 3.4009 (0.0136)

* -0.0024 (0.0023) N1 * 0.0037 (0.0026) C1 * -0.0034 (0.0026) C2 * 0.0020 (0.0025) C3 * 0.0002 (0.0023) N2

Rms deviation of fitted atoms = 0.0027

- 6.0873 (0.0064) x + 7.0610 (0.0142) y + 3.5857 (0.0170) z = 3.6036 (0.0096)

Angle to previous plane (with approximate esd) = The diher. 4.036 (0.230)

* 0.0055 (0.0024) N3 * -0.0044 (0.0026) C4 * 0.0004 (0.0028) C5 * 0.0023 (0.0029) C6 * -0.0012 (0.0028) C7 *

-0.0025 (0.0025) N4

Rms deviation of fitted atoms = 0.0032

Fractional atomic coordinates and isotropic or equivalent isotropic displacement parameters (\AA^2)

	x	y	z	$U_{\text{iso}}^*/U_{\text{eq}}$
Re1	0.24889 (2)	0.33339 (2)	0.76143 (2)	0.02623 (7)
Br1	-0.32866 (7)	-0.11457 (4)	0.69106 (5)	0.05216 (14)
Cl1	0.46364 (14)	0.21798 (9)	0.66737 (10)	0.0349 (2)
N1	0.2276 (4)	0.4143 (3)	0.5929 (3)	0.0277 (7)
N2	0.1065 (4)	0.3432 (3)	0.5096 (3)	0.0280 (7)
N3	0.0312 (4)	0.2057 (3)	0.6545 (3)	0.0260 (7)
N4	-0.0658 (5)	0.0996 (3)	0.6965 (3)	0.0308 (7)
N5	0.7500 (5)	0.1716 (4)	0.0701 (3)	0.0429 (9)
O8	0.2466 (6)	0.1981 (4)	0.9928 (3)	0.0656 (11)
O9	0.5758 (5)	0.5317 (4)	0.8704 (4)	0.0637 (11)
O10	-0.0322 (5)	0.4673 (4)	0.8641 (4)	0.0652 (11)
O11	0.7352 (5)	0.2848 (3)	0.2338 (3)	0.0569 (9)
C1	0.3002 (6)	0.5165 (4)	0.5393 (4)	0.0333 (9)
H1	0.3870	0.5826	0.5750	0.040*
C2	0.2290 (6)	0.5119 (4)	0.4225 (4)	0.0369 (10)
H2	0.2597	0.5720	0.3673	0.044*
C3	0.1068 (6)	0.4025 (4)	0.4057 (4)	0.0368 (10)
H3	0.0360	0.3730	0.3365	0.044*
C4	-0.0017 (5)	0.2319 (3)	0.5441 (3)	0.0253 (8)
C5	-0.1346 (6)	0.1578 (4)	0.4659 (4)	0.0320 (9)
H5	-0.1550	0.1797	0.3887	0.038*
C6	-0.2336 (6)	0.0512 (4)	0.5084 (4)	0.0360 (9)
H6	-0.3248	-0.0033	0.4616	0.043*
C7	-0.1907 (5)	0.0282 (3)	0.6255 (4)	0.0294 (8)
C8	0.2497 (6)	0.2478 (4)	0.9053 (4)	0.0394 (10)
C9	0.4528 (6)	0.4573 (4)	0.8307 (4)	0.0374 (10)
C10	0.0748 (6)	0.4186 (4)	0.8254 (4)	0.0385 (10)
C11	0.7113 (7)	0.2651 (4)	0.1273 (5)	0.0474 (12)
H11	0.6612	0.3217	0.0833	0.057*
C12	0.8330 (11)	0.0826 (6)	0.1358 (6)	0.083 (2)
H12A	0.9135	0.1237	0.2008	0.125*

H12B	0.9061	0.0442	0.0854	0.125*
H12C	0.7340	0.0211	0.1645	0.125*
C13	0.7163 (9)	0.1547 (7)	-0.0550 (5)	0.0749 (19)
H13A	0.6529	0.2163	-0.0853	0.112*
H13B	0.6386	0.0747	-0.0734	0.112*
H13C	0.8348	0.1616	-0.0901	0.112*

Atomic displacement parameters (Å²)

	U^{11}	U^{22}	U^{33}	U^{12}	U^{13}	U^{23}
Re1	0.02852 (10)	0.02494 (10)	0.02296 (10)	0.00326 (6)	-0.00562 (6)	-0.00529 (6)
Br1	0.0578 (3)	0.0378 (3)	0.0503 (3)	-0.0140 (2)	0.0036 (2)	0.0034 (2)
Cl1	0.0346 (5)	0.0301 (5)	0.0399 (6)	0.0083 (4)	-0.0015 (4)	-0.0084 (4)
N1	0.0267 (16)	0.0262 (17)	0.0283 (19)	0.0026 (13)	-0.0024 (14)	-0.0057 (14)
N2	0.0315 (17)	0.0273 (16)	0.0226 (17)	0.0029 (13)	-0.0052 (13)	-0.0017 (13)
N3	0.0263 (16)	0.0243 (16)	0.0249 (18)	0.0014 (12)	-0.0034 (13)	-0.0016 (13)
N4	0.0336 (18)	0.0281 (17)	0.0285 (18)	0.0021 (14)	0.0005 (14)	0.0000 (14)
N5	0.048 (2)	0.049 (2)	0.031 (2)	0.0122 (18)	-0.0049 (17)	-0.0085 (18)
O8	0.093 (3)	0.062 (2)	0.036 (2)	0.006 (2)	-0.0076 (19)	0.0165 (18)
O9	0.043 (2)	0.060 (2)	0.078 (3)	-0.0043 (17)	-0.0128 (18)	-0.034 (2)
O10	0.055 (2)	0.085 (3)	0.065 (3)	0.037 (2)	0.0074 (19)	-0.013 (2)
O11	0.081 (3)	0.051 (2)	0.036 (2)	0.0139 (18)	-0.0114 (18)	-0.0101 (16)
C1	0.028 (2)	0.025 (2)	0.045 (3)	0.0029 (16)	-0.0007 (18)	-0.0006 (18)
C2	0.036 (2)	0.035 (2)	0.040 (3)	0.0069 (17)	0.0049 (19)	0.0120 (19)
C3	0.041 (2)	0.043 (2)	0.028 (2)	0.0114 (19)	0.0022 (18)	0.0054 (19)
C4	0.0251 (18)	0.0260 (18)	0.024 (2)	0.0055 (14)	-0.0019 (15)	-0.0056 (15)
C5	0.036 (2)	0.036 (2)	0.022 (2)	0.0058 (17)	-0.0064 (16)	-0.0049 (17)
C6	0.032 (2)	0.035 (2)	0.037 (3)	-0.0005 (17)	-0.0049 (18)	-0.0094 (18)
C7	0.030 (2)	0.0260 (19)	0.030 (2)	0.0014 (15)	-0.0018 (16)	-0.0052 (16)
C8	0.043 (2)	0.037 (2)	0.035 (3)	0.0051 (18)	-0.0069 (19)	-0.008 (2)
C9	0.035 (2)	0.037 (2)	0.039 (3)	0.0074 (18)	-0.0020 (19)	-0.0091 (19)
C10	0.042 (2)	0.040 (2)	0.030 (2)	0.0060 (19)	-0.0092 (19)	-0.0072 (19)
C11	0.057 (3)	0.043 (3)	0.042 (3)	0.012 (2)	-0.007 (2)	0.001 (2)
C12	0.125 (6)	0.071 (4)	0.070 (5)	0.056 (4)	0.012 (4)	0.000 (3)
C13	0.063 (4)	0.116 (6)	0.042 (3)	0.014 (4)	-0.001 (3)	-0.033 (3)

Geometric parameters (Å, °)

Re1—C10	1.904 (5)	O10—C10	1.143 (6)
Re1—C8	1.914 (5)	O11—C11	1.227 (6)
Re1—C9	1.915 (4)	C1—C2	1.396 (6)
Re1—N1	2.150 (3)	C2—C3	1.350 (6)
Re1—N3	2.180 (3)	C4—C5	1.392 (5)
Re1—Cl1	2.4982 (11)	C5—C6	1.366 (6)
Br1—C7	1.886 (4)	C6—C7	1.394 (6)
N1—C1	1.327 (5)	C1—H1	0.9300
N1—N2	1.370 (5)	C2—H2	0.9300
N2—C3	1.368 (5)	C3—H3	0.9300

N2—C4	1.397 (5)	C5—H5	0.9300
N3—C4	1.318 (5)	C6—H6	0.9300
N3—N4	1.354 (4)	C11—H11	0.9300
N4—C7	1.301 (5)	C12—H12A	0.9600
N5—C11	1.309 (6)	C12—H12B	0.9600
N5—C13	1.437 (7)	C12—H12C	0.9600
N5—C12	1.458 (7)	C13—H13A	0.9600
O8—C8	1.149 (6)	C13—H13B	0.9600
O9—C9	1.145 (5)	C13—H13C	0.9600
C10—Re1—C8	87.5 (2)	C6—C5—C4	116.6 (4)
C10—Re1—C9	89.04 (19)	C5—C6—C7	116.5 (4)
C8—Re1—C9	88.32 (19)	N4—C7—C6	125.5 (4)
C10—Re1—N1	93.10 (16)	N4—C7—Br1	115.9 (3)
C8—Re1—N1	174.39 (15)	C6—C7—Br1	118.6 (3)
C9—Re1—N1	97.27 (16)	O8—C8—Re1	178.1 (5)
C10—Re1—N3	94.22 (16)	O9—C9—Re1	178.9 (4)
C8—Re1—N3	100.89 (16)	O10—C10—Re1	178.5 (4)
C9—Re1—N3	170.35 (15)	O11—C11—N5	125.9 (5)
N1—Re1—N3	73.50 (12)	N1—C1—H1	124.5
C10—Re1—C11	176.65 (13)	C2—C1—H1	124.5
C8—Re1—C11	94.41 (14)	C3—C2—H2	126.9
C9—Re1—C11	93.74 (13)	C1—C2—H2	126.9
N1—Re1—C11	84.70 (9)	C2—C3—N2	107.2 (4)
N3—Re1—C11	82.74 (9)	C2—C3—H3	126.4
C1—N1—N2	105.2 (3)	N2—C3—H3	126.4
C1—N1—Re1	139.4 (3)	C6—C5—H5	121.7
N2—N1—Re1	115.4 (2)	C4—C5—H5	121.7
C3—N2—N1	110.4 (3)	C5—C6—H6	121.7
C3—N2—C4	131.5 (4)	C7—C6—H6	121.7
N1—N2—C4	117.9 (3)	O11—C11—H11	117.0
C4—N3—N4	119.4 (3)	N5—C11—H11	117.0
C4—N3—Re1	118.1 (2)	N5—C12—H12A	109.5
N4—N3—Re1	122.4 (2)	N5—C12—H12B	109.5
C7—N4—N3	117.9 (3)	H12A—C12—H12B	109.5
C11—N5—C13	122.5 (5)	N5—C12—H12C	109.5
C11—N5—C12	118.6 (4)	H12A—C12—H12C	109.5
C13—N5—C12	118.9 (5)	H12B—C12—H12C	109.5
N1—C1—C2	111.1 (4)	N5—C13—H13A	109.5
C3—C2—C1	106.1 (4)	N5—C13—H13B	109.5
C2—C3—N2	107.2 (4)	H13A—C13—H13B	109.5
N3—C4—C5	124.1 (3)	N5—C13—H13C	109.5
N3—C4—N2	114.8 (3)	H13A—C13—H13C	109.5
C5—C4—N2	121.1 (3)	H13B—C13—H13C	109.5
C1—N1—N2—C3	0.3 (4)	Re1—N3—C4—N2	-1.8 (4)
Re1—N1—N2—C3	-178.9 (3)	C3—N2—C4—N3	-177.1 (4)
C1—N1—N2—C4	-175.5 (3)	N1—N2—C4—N3	-2.4 (5)

Re1—N1—N2—C4	5.4 (4)	C3—N2—C4—C5	2.1 (6)
C4—N3—N4—C7	-0.9 (5)	N1—N2—C4—C5	176.8 (3)
Re1—N3—N4—C7	-178.8 (3)	N3—C4—C5—C6	-0.7 (6)
N2—N1—C1—C2	-0.6 (5)	N2—C4—C5—C6	-179.7 (4)
Re1—N1—C1—C2	178.2 (3)	C4—C5—C6—C7	0.0 (6)
N1—C1—C2—C3	0.7 (5)	N3—N4—C7—C6	0.3 (6)
C1—C2—C3—N2	-0.5 (5)	N3—N4—C7—Br1	-177.3 (3)
N1—N2—C3—C2	0.2 (5)	C5—C6—C7—N4	0.2 (7)
C4—N2—C3—C2	175.2 (4)	C5—C6—C7—Br1	177.8 (3)
N4—N3—C4—C5	1.2 (6)	C13—N5—C11—O11	179.5 (5)
Re1—N3—C4—C5	179.1 (3)	C12—N5—C11—O11	-1.5 (8)
N4—N3—C4—N2	-179.7 (3)		

Hydrogen-bond geometry (Å, °)

<i>D</i> —H \cdots <i>A</i>	<i>D</i> —H	H \cdots <i>A</i>	<i>D</i> \cdots <i>A</i>	<i>D</i> —H \cdots <i>A</i>
C3—H3 \cdots O11 ⁱ	0.93	2.41	3.251 (6)	150
C5—H5 \cdots O11 ⁱ	0.93	2.33	3.206 (5)	158
C2—H2 \cdots Cl1 ⁱⁱ	0.93	2.79	3.556 (4)	140
C6—H6 \cdots Cl1 ⁱⁱⁱ	0.93	2.75	3.614 (4)	154

Symmetry codes: (i) $x-1, y, z$; (ii) $-x+1, -y+1, -z+1$; (iii) $-x, -y, -z+1$.