

Tetraethylammonium tetra- μ -benzamidato- κ^4 N:O; κ^4 O:N-dichlorodirhodate(II,III)(Rh—Rh) dichloromethane disolvate

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Key indicators

Single-crystal X-ray study
T = 173 K
Mean $\sigma(C-C)$ = 0.007 Å
Disorder in solvent or counterion
R factor = 0.049
wR factor = 0.101
Data-to-parameter ratio = 18.6

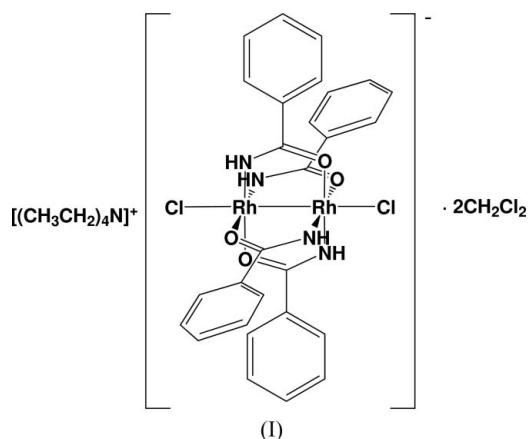
For details of how these key indicators were automatically derived from the article, see <http://journals.iucr.org/e>.

The title compound, $(C_8H_{20}N)[Rh_2(C_7H_6NO)_4Cl_2] \cdot 2CH_2Cl_2$, has the acetamide-bridged-Rh₂ anion on an inversion centre. The tetraethylammonium cation is disordered on another inversion centre. The Rh—Rh bond distance is 2.4265 (13) Å; the metal atom shows octahedral coordination. The chloro ligands both occupy axial sites.

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Comment

Paddlewheel dinuclear complexes are good modules for constructing assembled structures since they have linearly arranged axial sites. Halide ions are the most common linker ligands for cationic paddlewheel dinuclear complexes in the construction of chain structures, such as $[Ru_2(O_2CR)_4(\mu-Cl)]$ (Aquino, 1998, 2004). For dirhodium complexes, we have reported chain structures constructed from the acetamide-bridged cationic complex and halide ions, $[Rh_2(acam)_4(\mu-X)] \cdot nH_2O$ (Hacam = acetamide; *X* = Cl, Br, I; *n* = 0, 2, 3, 7; Yang *et al.*, 2000, 2001). Discrete paddlewheel dirhodium complexes with chloro ligands may be important modules for the construction of assembled structures with metal cations, but so far examples have only been reported for the oxidation state Rh₂⁴⁺ (Miskowski *et al.*, 1984, 1987; Zhou *et al.*, 1991; Galdecka *et al.*, 1999). In this paper, we report the title dirhodium(II,III) complex with chloro ligands at both axial sites, $[(C_2H_5)_4N][Rh_2(bzam)_4Cl_2] \cdot 2CH_2Cl_2$, (I) (Hbzam = benzamide).



The structure of (I) is shown in Fig. 1. The dirhodium complex lies on an inversion centre at $(\frac{1}{2}, \frac{1}{2}, \frac{1}{2})$. The tetraethylammonium ion lies on another inversion centre at $(0, \frac{1}{2}, 0)$. The Rh—Rh distance of 2.4265 (13) Å in (I) is similar to that in $[Rh_2(bzam)_4(py)_2]$ [2.437 (1) Å; Chakravarty *et al.*, 1985]. This small difference in the Rh—Rh distance between the

Rh_2^{4+} and Rh_2^{5+} complexes is consistent with the δ^* odd-electron orbital of amidate-bridged dirhodium complexes (Kawamura *et al.*, 1989, 1998). The Rh–Rh distance is also similar to those of the chloro-bridged chain structures of the Rh_2^{5+} complexes $[\text{Rh}_2(\text{acam})_4\text{Cl}]$ [2.428 (1) Å; Yang *et al.*, 2001] and $[\text{Rh}_2(\text{acam})_4\text{Cl}] \cdot 7\text{H}_2\text{O}$ [2.4172 (7) Å; Yang *et al.*, 2000].

The Rh–Cl distance of 2.5168 (16) Å is shorter than those observed in the chain complexes $[\text{Rh}_2(\text{acam})_4\text{Cl}]$ [2.581 (1) Å] and $[\text{Rh}_2(\text{acam})_4\text{Cl}] \cdot 7\text{H}_2\text{O}$ [2.564 (1) Å]. It is also shorter than that in the Rh_2^{4+} complexes $[\text{C}(\text{NH}_2)_3]_2[\text{Rh}_2(\text{O}_2\text{CCH}_3)_4\text{Cl}_2]$ [2.5853 (6) Å; Miskowski *et al.*, 1987], $\text{Li}_2[\text{Rh}_2(\text{O}_2\text{CCH}_3)_4\text{Cl}_2] \cdot 8\text{H}_2\text{O}$ [2.601 (1) Å; Miskowski *et al.*, 1984] and $\text{Na}_2[\text{Rh}_2(\text{O}_2\text{CCH}_3)_4\text{Cl}_2] \cdot 4\text{H}_2\text{O}$ [2.564 (1) Å; Galdecka *et al.*, 1999].

Experimental

$[\text{Rh}_2(\text{bzam})_4(\text{py})_2]$ (py = pyridine) was prepared using the literature method of Chakravarty *et al.* (1985). To a CH_2Cl_2 solution of $[\text{Rh}_2(\text{bzam})_4(\text{py})_2]$ (0.090 g, 0.11 mmol), AgPF_6 (0.234 g, 0.93 mmol) in toluene (10 ml) was added. The resulting black precipitate was removed by filtration and the remaining solution was evaporated to dryness to afford the crude product, $[\text{Rh}_2(\text{bzam})_4(\text{py})_2]\text{PF}_6$. This crude product was washed with H_2O and toluene and dissolved in CH_2Cl_2 . $[(\text{C}_2\text{H}_5)_4\text{N}] \text{Cl}$ in CH_2Cl_2 was slowly diffused into a CH_2Cl_2 solution of $[\text{Rh}_2(\text{bzam})_4(\text{py})_2]\text{PF}_6$ to obtain crystals of (I).

Crystal data

$(\text{C}_8\text{H}_{20}\text{N})[\text{Rh}_2(\text{C}_7\text{H}_6\text{NO})_4\text{Cl}_2] \cdot 2\text{CH}_2\text{Cl}_2$	$\gamma = 83.479$ (18)°
$M_r = 1057.33$	$V = 1107.1$ (10) Å ³
Triclinic, $P\bar{1}$	$Z = 1$
$a = 9.649$ (5) Å	$D_x = 1.586$ Mg m ⁻³
$b = 10.506$ (6) Å	Mo $K\alpha$ radiation
$c = 12.202$ (6) Å	$\mu = 1.15$ mm ⁻¹
$\alpha = 76.571$ (16)°	$T = 173$ (2) K
$\beta = 66.996$ (13)°	Prism, brown
	0.10 × 0.10 × 0.10 mm

Data collection

Rigaku/MSC Mercury CCD area-detector diffractometer
 ω scans
Absorption correction: none
9060 measured reflections

5028 independent reflections
4109 reflections with $I > 2\sigma(I)$
 $R_{\text{int}} = 0.032$
 $\theta_{\text{max}} = 27.5$ °

Refinement

Refinement on F^2
 $R[F^2 > 2\sigma(F^2)] = 0.049$
 $wR(F^2) = 0.101$
 $S = 1.07$
5028 reflections
270 parameters
H-atom parameters constrained

$$w = 1/[\sigma^2(F_o^2) + (0.0319P)^2 + 2.8887P]$$

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

$$(\Delta/\sigma)_{\text{max}} = 0.001$$

$$\Delta\rho_{\text{max}} = 0.72 \text{ e } \text{\AA}^{-3}$$

$$\Delta\rho_{\text{min}} = -0.80 \text{ e } \text{\AA}^{-3}$$

Table 1
Selected bond lengths (Å).

$\text{Rh1} - \text{Rh1}^i$	2.4266 (13)	$\text{Rh1} - \text{O2}^i$	2.043 (3)
$\text{Rh1} - \text{Cl1}$	2.5167 (16)	$\text{Rh1} - \text{N1}$	1.975 (3)
$\text{Rh1} - \text{O1}^i$	2.038 (3)	$\text{Rh1} - \text{N2}$	1.970 (3)

Symmetry code: (i) $-x + 1, -y + 1, -z + 1$.

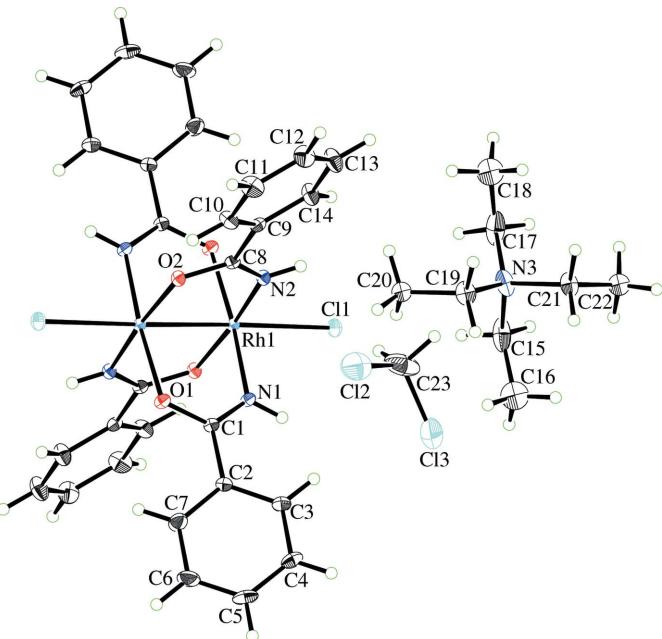


Figure 1

The N–C and C–C distances in the tetraethylammonium ion were restrained (SADI, 0.005) and methyl C atoms were refined with isotropic displacement parameters. All H atoms were placed in idealized positions and treated as riding atoms, with C–H distances in the range 0.95–0.99 Å and with $U_{\text{iso}}(\text{H}) = 1.2U_{\text{eq}}(\text{C})$ or $1.5U_{\text{eq}}(\text{C})$, and with N–H = 0.88 Å and $U_{\text{iso}}(\text{H}) = 1.2U_{\text{eq}}(\text{N})$. The occupancy factors of all C atoms (C15–C22) in the disordered tetraethylammonium ion were set at 0.5.

Data collection: *CrystalClear* (Molecular Structure Corporation & Rigaku, 2001); cell refinement: *CrystalClear*; data reduction: *TEXSAN* (Rigaku/MSC, 2004); program(s) used to solve structure: *SIR97* (Altomare *et al.*, 1999); program(s) used to refine structure: *SHELXL97* (Sheldrick, 1997); molecular graphics: *ORTEPII* (Johnson, 1976); software used to prepare material for publication: *SHELXL97* and *TEXSAN*.

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