

Crystal structure of tetrakis(μ -*n*-butyrato- κ^2 O:O')bis[chloridorhenium(III)](Re—Re)

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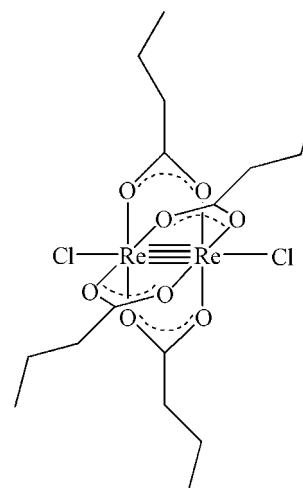
With an inversion center at the mid-point of the two Re^{III} atoms, the title compound, [Re₂Cl₂{O₂C(CH₂)₂CH₃}]₄, exhibits a paddle-wheel or lantern-type structure with four *n*-butyrate groups bridging two Re^{III} atoms in a *syn-syn* fashion. The axial chloride ligands together with the Re—Re quadruple bond [2.2330 (3) Å] complete an essentially octahedral geometry around each Re^{III} atom. There is little distortion, with an Re—Re—Cl bond angle of 176.18 (3)° and typical *cis*-O—Re—O bond angles ranging from 89.39 (11) to 90.68 (11)°. There are two molecules in the unit cell, and no significant intermolecular interactions were noticed between molecules in the crystal.

Keywords: crystal structure; dirhenium core; butyrate bridging ligand.

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1. Related literature

For the synthesis and structure of five related structures, see: Taha & Wilkinson (1963); Calvo *et al.* (1969); Collins *et al.* (1979); Lydon *et al.* (2003).



2. Experimental

2.1. Crystal data

[Re ₂ (C ₄ H ₇ O ₂) ₄ Cl ₂]	<i>V</i> = 1173.0 (1) Å ³
<i>M_r</i> = 791.68	<i>Z</i> = 2
Monoclinic, <i>P</i> 2 ₁ / <i>n</i>	Mo <i>K</i> α radiation
<i>a</i> = 6.7292 (4) Å	<i>μ</i> = 10.57 mm ⁻¹
<i>b</i> = 12.0367 (8) Å	<i>T</i> = 180 K
<i>c</i> = 14.6737 (9) Å	0.30 × 0.06 × 0.06 mm
<i>β</i> = 99.262 (1)°	

2.2. Data collection

Bruker APEXII CCD diffractometer	4625 measured reflections
Absorption correction: multi-scan (SADABS; Bruker, 2010)	2277 independent reflections
<i>T</i> _{min} = 0.144, <i>T</i> _{max} = 0.570	2077 reflections with <i>I</i> > 2σ(<i>I</i>)
	<i>R</i> _{int} = 0.016

2.3. Refinement

<i>R</i> [<i>F</i> ² > 2σ(<i>F</i> ²)] = 0.019	129 parameters
<i>wR</i> (<i>F</i> ²) = 0.049	H-atom parameters constrained
<i>S</i> = 1.06	Δ <i>ρ</i> _{max} = 0.90 e Å ⁻³
2277 reflections	Δ <i>ρ</i> _{min} = -1.32 e Å ⁻³

Table 1

Selected bond lengths (Å).

Re1—O2 ⁱ	2.008 (3)	Re1—O1	2.026 (3)
Re1—O3	2.019 (2)	Re1—Cl1	2.5135 (9)
Re1—O4 ⁱ	2.025 (2)		

Symmetry code: (i) $-x + 2, -y + 1, -z + 2$.

Data collection: APEX2 (Bruker, 2010); cell refinement: SAINT (Bruker, 2010); data reduction: SAINT; program(s) used to solve structure: SHELXS97 (Sheldrick, 2008); program(s) used to refine structure: SHELXL97 (Sheldrick, 2008); molecular graphics: SHELXTL (Sheldrick, 2008); software used to prepare material for publication: SHELXTL.

Acknowledgements

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Supporting information for this paper is available from the IUCr electronic archives (Reference: WM5055).

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supporting information

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Crystal structure of tetrakis(μ -*n*-butyrato- κ^2 O:O')bis[chloridorhenium(III)](Re—Re)

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S1. Synthesis and crystallization

This compound was prepared following a method analogous to the procedure used by Taha & Wilkinson (1963). Rhenium trichloride (1.25 g, 4.27 mmol) was reacted with 13 ml of butyric acid and 0.5 ml of butyric anhydride at 423 K, with stirring, under a steady stream of nitrogen for five days. The dark solution was removed from heat and allowed to cool overnight. The brown product was collected *via* suction filtration and washed with a 50 ml portion of petroleum ether. The product was dried *in vacuo* for 24 hours and then re-crystallized from dichloromethane. Yield = 0.481 g, (56.9 %).

S2. Refinement

All H atoms were placed in geometrically calculated positions, with C—H = 0.99 Å (CH₂), and 0.98 Å (CH₃), and refined as riding atoms, with $U_{\text{iso}}(\text{H}) = 1.5U_{\text{eq}}(\text{C})$ (CH₃) or $1.2U_{\text{eq}}(\text{C})$ (CH₂).

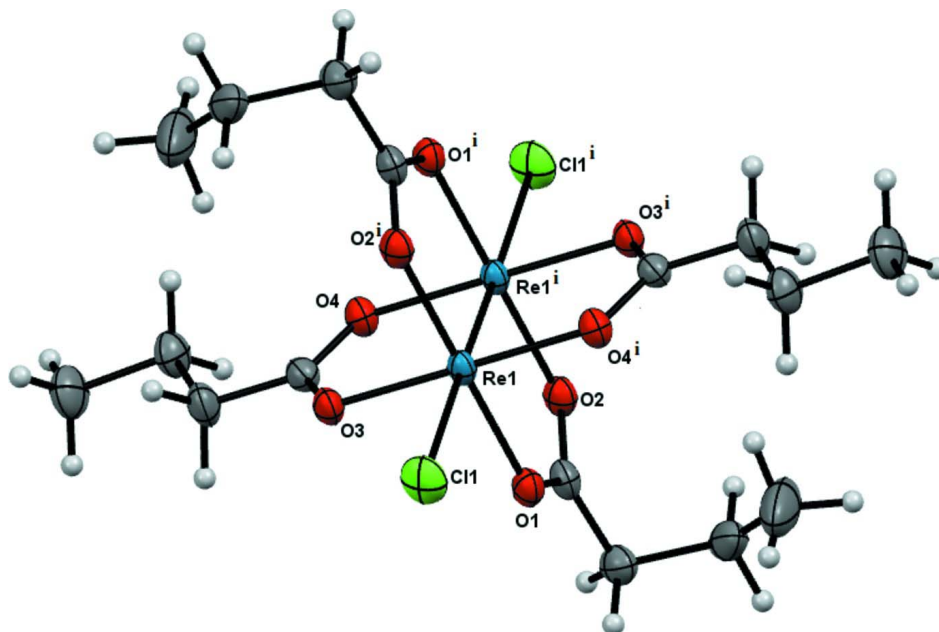
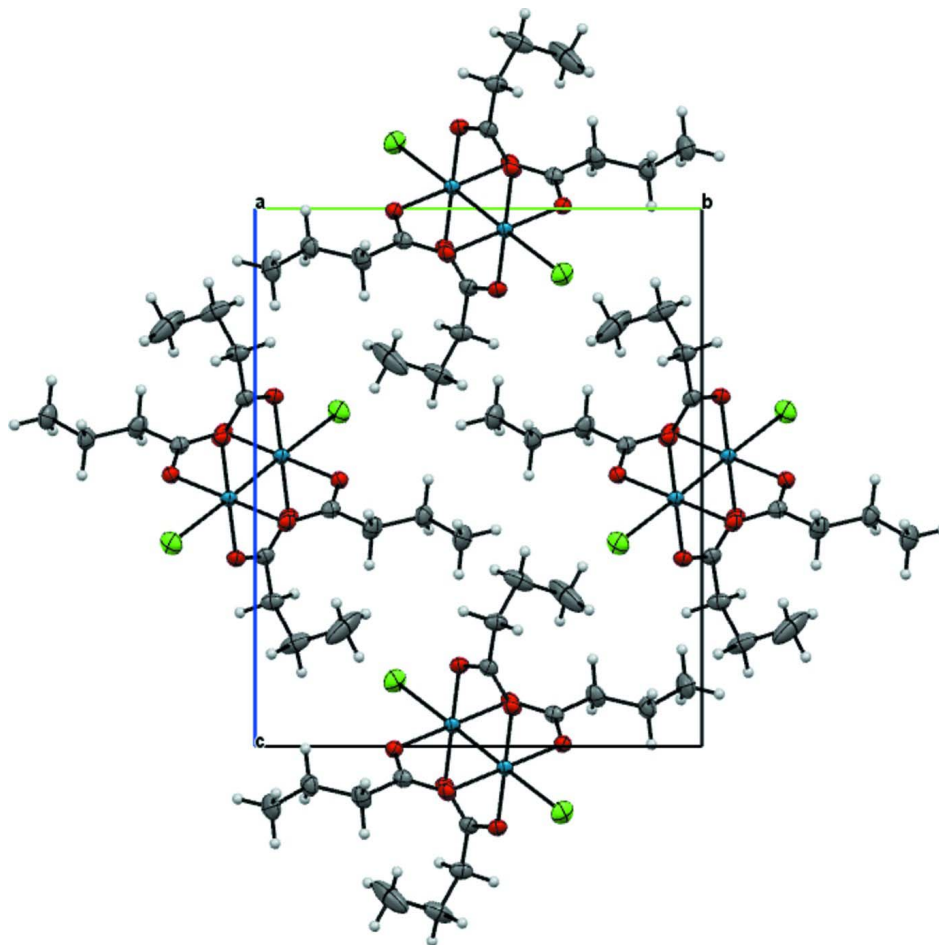


Figure 1

The molecular structure of the title compound with displacement ellipsoids drawn at the 50% probability level. Hydrogen atoms are drawn as small spheres of arbitrary radius. [Symmetry code i) $-x + 2, -y + 1, -z + 2$.]

**Figure 2**

The packing diagram for the title compound viewed along [100].

Tetrakis(μ -*n*-butyrato- κ^2 O:O')bis[chloridorhenium(III)](Re—Re)

Crystal data

[Re₂(C₄H₇O₂)₄Cl₂]

$M_r = 791.68$

Monoclinic, $P2_1/n$

Hall symbol: $-P\ 2_1n$

$a = 6.7292$ (4) Å

$b = 12.0367$ (8) Å

$c = 14.6737$ (9) Å

$\beta = 99.262$ (1)°

$V = 1173.0$ (1) Å³

$Z = 2$

$F(000) = 744$

$D_x = 2.241$ Mg m⁻³

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

Cell parameters from 3188 reflections

$\theta = 3.7$ – 27.1 °

$\mu = 10.57$ mm⁻¹

$T = 180$ K

Needle, orange

$0.30 \times 0.06 \times 0.06$ mm

Data collection

Bruker APEXII CCD
diffractometer

Radiation source: fine-focus sealed tube

Graphite monochromator

φ and ω scans

Absorption correction: multi-scan
(*SADABS*; Bruker, 2010)

$T_{\min} = 0.144$, $T_{\max} = 0.570$

4625 measured reflections

2277 independent reflections

2077 reflections with $I > 2\sigma(I)$

$R_{\text{int}} = 0.016$
 $\theta_{\text{max}} = 26.0^\circ$, $\theta_{\text{min}} = 3.7^\circ$
 $h = -8 \rightarrow 7$

$k = -12 \rightarrow 14$
 $l = -18 \rightarrow 18$

Refinement

Refinement on F^2
 Least-squares matrix: full
 $R[F^2 > 2\sigma(F^2)] = 0.019$
 $wR(F^2) = 0.049$
 $S = 1.06$
 2277 reflections
 129 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
 $w = 1/[\sigma^2(F_o^2) + (0.0278P)^2 + 0.2053P]$
 where $P = (F_o^2 + 2F_c^2)/3$
 $(\Delta/\sigma)_{\text{max}} = 0.005$
 $\Delta\rho_{\text{max}} = 0.90 \text{ e } \text{\AA}^{-3}$
 $\Delta\rho_{\text{min}} = -1.32 \text{ e } \text{\AA}^{-3}$

Special details

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

Refinement. Refinement of F^2 against ALL reflections. The weighted R -factor wR and goodness of fit S are based on F^2 , conventional R -factors R are based on F , with F set to zero for negative F^2 . The threshold expression of $F^2 > \sigma(F^2)$ is used only for calculating R -factors(gt) *etc.* and is not relevant to the choice of reflections for refinement. R -factors based on F^2 are statistically about twice as large as those based on F , and R -factors based on ALL data will be even larger.

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

	<i>x</i>	<i>y</i>	<i>z</i>	$U_{\text{iso}}^*/U_{\text{eq}}$
Re1	0.91824 (2)	0.559012 (11)	1.039538 (10)	0.02050 (7)
Cl1	0.71261 (17)	0.68750 (8)	1.12219 (8)	0.0397 (3)
O1	0.7020 (4)	0.57529 (19)	0.92707 (18)	0.0254 (6)
O2	0.8665 (4)	0.45750 (19)	0.84940 (19)	0.0256 (6)
O3	0.7697 (4)	0.42938 (19)	1.08438 (19)	0.0243 (6)
O4	0.9312 (4)	0.31123 (19)	1.00486 (18)	0.0241 (6)
C1	0.7167 (6)	0.5223 (3)	0.8532 (3)	0.0253 (8)
C2	0.5652 (6)	0.5386 (3)	0.7691 (3)	0.0308 (9)
H2A	0.4565	0.5877	0.7838	0.037*
H2B	0.5045	0.4661	0.7486	0.037*
C3	0.6593 (7)	0.5899 (4)	0.6916 (3)	0.0451 (12)
H3A	0.7561	0.5366	0.6719	0.054*
H3B	0.5526	0.6040	0.6380	0.054*
C4	0.7670 (9)	0.6974 (5)	0.7202 (5)	0.083 (2)
H4A	0.8092	0.7325	0.6662	0.125*
H4B	0.8857	0.6821	0.7666	0.125*
H4C	0.6761	0.7474	0.7464	0.125*
C5	0.8023 (6)	0.3306 (3)	1.0595 (3)	0.0247 (8)
C6	0.6899 (6)	0.2376 (3)	1.0943 (3)	0.0338 (10)
H6A	0.5453	0.2464	1.0688	0.041*
H6B	0.7037	0.2445	1.1623	0.041*
C7	0.7539 (7)	0.1221 (3)	1.0718 (3)	0.0333 (9)

H7A	0.8978	0.1111	1.0975	0.040*
H7B	0.7373	0.1126	1.0040	0.040*
C8	0.6260 (8)	0.0359 (3)	1.1127 (3)	0.0426 (11)
H8A	0.6674	-0.0389	1.0974	0.064*
H8B	0.4836	0.0470	1.0871	0.064*
H8C	0.6449	0.0445	1.1800	0.064*

Atomic displacement parameters (Å²)

	U^{11}	U^{22}	U^{33}	U^{12}	U^{13}	U^{23}
Re1	0.02322 (11)	0.02022 (10)	0.02042 (10)	0.00156 (6)	0.01068 (7)	0.00100 (5)
Cl1	0.0449 (6)	0.0377 (5)	0.0419 (6)	0.0105 (5)	0.0232 (5)	-0.0039 (5)
O1	0.0283 (15)	0.0278 (13)	0.0217 (14)	0.0015 (11)	0.0092 (12)	0.0030 (10)
O2	0.0294 (16)	0.0275 (13)	0.0222 (14)	0.0013 (11)	0.0108 (12)	-0.0003 (10)
O3	0.0256 (15)	0.0252 (13)	0.0243 (14)	-0.0007 (10)	0.0105 (12)	0.0029 (10)
O4	0.0267 (14)	0.0235 (13)	0.0238 (13)	-0.0025 (11)	0.0086 (11)	-0.0006 (11)
C1	0.027 (2)	0.0276 (18)	0.024 (2)	-0.0031 (16)	0.0112 (17)	0.0024 (15)
C2	0.029 (2)	0.041 (2)	0.024 (2)	0.0023 (18)	0.0081 (17)	0.0013 (17)
C3	0.034 (3)	0.071 (3)	0.033 (2)	0.015 (2)	0.012 (2)	0.021 (2)
C4	0.063 (4)	0.095 (5)	0.084 (5)	-0.027 (4)	-0.010 (3)	0.060 (4)
C5	0.0217 (19)	0.0284 (19)	0.0246 (19)	-0.0020 (16)	0.0058 (15)	0.0014 (15)
C6	0.035 (2)	0.031 (2)	0.039 (2)	-0.0015 (18)	0.0177 (19)	0.0082 (17)
C7	0.038 (2)	0.025 (2)	0.038 (2)	-0.0064 (17)	0.011 (2)	0.0019 (17)
C8	0.048 (3)	0.031 (2)	0.049 (3)	-0.012 (2)	0.012 (2)	0.0063 (19)

Geometric parameters (Å, °)

Re1—O2 ⁱ	2.008 (3)	C3—C4	1.509 (7)
Re1—O3	2.019 (2)	C3—H3A	0.9900
Re1—O4 ⁱ	2.025 (2)	C3—H3B	0.9900
Re1—O1	2.026 (3)	C4—H4A	0.9800
Re1—Re1 ⁱ	2.2330 (3)	C4—H4B	0.9800
Re1—Cl1	2.5135 (9)	C4—H4C	0.9800
O1—C1	1.275 (5)	C5—C6	1.487 (5)
O2—C1	1.283 (4)	C6—C7	1.508 (5)
O2—Re1 ⁱ	2.008 (3)	C6—H6A	0.9900
O3—C5	1.273 (4)	C6—H6B	0.9900
O4—C5	1.294 (4)	C7—C8	1.531 (5)
O4—Re1 ⁱ	2.025 (2)	C7—H7A	0.9900
C1—C2	1.482 (6)	C7—H7B	0.9900
C2—C3	1.520 (6)	C8—H8A	0.9800
C2—H2A	0.9900	C8—H8B	0.9800
C2—H2B	0.9900	C8—H8C	0.9800
O2 ⁱ —Re1—O3	89.39 (11)	C4—C3—H3B	109.2
O2 ⁱ —Re1—O4 ⁱ	90.26 (10)	C2—C3—H3B	109.2
O3—Re1—O4 ⁱ	179.64 (11)	H3A—C3—H3B	107.9
O2 ⁱ —Re1—O1	179.70 (10)	C3—C4—H4A	109.5

O3—Re1—O1	90.68 (11)	C3—C4—H4B	109.5
O4 ⁱ —Re1—O1	89.67 (10)	H4A—C4—H4B	109.5
O2 ⁱ —Re1—Re1 ⁱ	90.42 (7)	C3—C4—H4C	109.5
O3—Re1—Re1 ⁱ	89.36 (7)	H4A—C4—H4C	109.5
O4 ⁱ —Re1—Re1 ⁱ	90.55 (7)	H4B—C4—H4C	109.5
O1—Re1—Re1 ⁱ	89.29 (7)	O3—C5—O4	120.7 (3)
O2 ⁱ —Re1—Cl1	92.90 (8)	O3—C5—C6	118.9 (3)
O3—Re1—Cl1	88.76 (8)	O4—C5—C6	120.4 (3)
O4 ⁱ —Re1—Cl1	91.34 (7)	C5—C6—C7	116.1 (3)
O1—Re1—Cl1	87.40 (8)	C5—C6—H6A	108.3
Re1 ⁱ —Re1—Cl1	176.18 (3)	C7—C6—H6A	108.3
C1—O1—Re1	120.0 (2)	C5—C6—H6B	108.3
C1—O2—Re1 ⁱ	119.6 (2)	C7—C6—H6B	108.3
C5—O3—Re1	120.7 (2)	H6A—C6—H6B	107.4
C5—O4—Re1 ⁱ	118.7 (2)	C6—C7—C8	109.9 (4)
O1—C1—O2	120.7 (4)	C6—C7—H7A	109.7
O1—C1—C2	120.3 (3)	C8—C7—H7A	109.7
O2—C1—C2	119.0 (3)	C6—C7—H7B	109.7
C1—C2—C3	111.3 (4)	C8—C7—H7B	109.7
C1—C2—H2A	109.4	H7A—C7—H7B	108.2
C3—C2—H2A	109.4	C7—C8—H8A	109.5
C1—C2—H2B	109.4	C7—C8—H8B	109.5
C3—C2—H2B	109.4	H8A—C8—H8B	109.5
H2A—C2—H2B	108.0	C7—C8—H8C	109.5
C4—C3—C2	112.3 (4)	H8A—C8—H8C	109.5
C4—C3—H3A	109.2	H8B—C8—H8C	109.5
C2—C3—H3A	109.2		
O2 ⁱ —Re1—O1—C1	13 (19)	Re1 ⁱ —O2—C1—O1	-0.9 (5)
O3—Re1—O1—C1	-89.7 (3)	Re1 ⁱ —O2—C1—C2	176.6 (2)
O4 ⁱ —Re1—O1—C1	90.2 (3)	O1—C1—C2—C3	115.7 (4)
Re1 ⁱ —Re1—O1—C1	-0.3 (3)	O2—C1—C2—C3	-61.7 (5)
Cl1—Re1—O1—C1	-178.4 (3)	C1—C2—C3—C4	-55.2 (5)
O2 ⁱ —Re1—O3—C5	-90.7 (3)	Re1—O3—C5—O4	-0.1 (5)
O4 ⁱ —Re1—O3—C5	-77 (17)	Re1—O3—C5—C6	179.9 (3)
O1—Re1—O3—C5	89.0 (3)	Re1 ⁱ —O4—C5—O3	0.5 (5)
Re1 ⁱ —Re1—O3—C5	-0.3 (3)	Re1 ⁱ —O4—C5—C6	-179.5 (3)
Cl1—Re1—O3—C5	176.4 (3)	O3—C5—C6—C7	-172.7 (4)
Re1—O1—C1—O2	0.8 (5)	O4—C5—C6—C7	7.2 (6)
Re1—O1—C1—C2	-176.6 (2)	C5—C6—C7—C8	179.5 (4)

Symmetry code: (i) $-x+2, -y+1, -z+2$.