TEXSAN (Molecular Structure Corporation, 1989). Program(s) used to solve structure: TEXSAN. Program(s) used to refine structure: TEXSAN. Software used to prepare material for publication: TEXSAN.

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(Benzenethiolato-S)(4-tert-butylpyridine-N)bis(dimethylglyoximato-N,N')cobalt(III) and (4-tert-Butylpyridine-N)bis(dimethylglyoximato-N,N')(4-methoxybenzenethiolato-S)cobalt(III)

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Abstract

The Co—S distances in the title compounds, [Co- $(C_4H_7N_2O_2)_2(C_9H_{13}N)(C_6H_5S)$] and [Co($C_4H_7N_2O_2)_2-(C_9H_{13}N)(C_7H_7OS)$], are 2.2800 (8) and 2.2885 (15) Å,

respectively. The electron-donating substituent on the para position of the arylthiolate lengthens the Co—S bond.

Comment

A Co—S coordination is one of the possible models of coenzyme B_{12} involvement in biological methyl transfer from a tetrahydrofolate coenzyme to homosystein, which is mediated by coenzyme B_{12} (Matthews, 1984; Taylor, 1982). Model studies using cobaloxime, $Co(DH)_2$, as a coenzyme B_{12} model showed that a variety of reaction schemes are envisaged for the reaction of the cobalt complex and thiol (Schrauzer & Windgassen, 1967; Jacobsen *et al.*, 1993; Polson *et al.*, 1997). Although the crystal structures of cobaloxime derivatives, $Co(DH)_2L^1L^2$, have been widely investigated and reviewed (Pahor *et al.*, 1985; Randaccio *et al.*, 1989), only a few compounds involving sulfur ligands have been reported (L^1 and/or $L^2 = SR$; Polson *et al.*, 1997, and references therein).

As a model reaction of biological methyl transfer from methyltetrahydrofolic acid to coenzyme M, we studied the methyl transfer from the N5-methyltetrahydropteridinium ion to arylthiocobaloxime. The reaction simulated the methyl transfer from the ammonium ion to the arylthio group to give methyl aryl sulfide, and the relative reactivity of the arylthiocobaloximes decreased in the order *p*-methoxyphenylthio-, phenylthio-, *p*-chlorophenylthio- and *p*-cyanophenylthiocobaloxime (Tada *et al.*, 1998).

If the alkyl transfer starts by the initial homolysis of the Co—S bond, the reactivity must follow the bond strength and hence the bond distance of the Co—S bond. The crystal structures of [Co(DH)₂(4-'BuPy)(PhS)], (1), and [Co(DH)₂(4-'BuPy)(4-MeO-PhS)], (2), were investigated on this premis.

The geometries around the cobaloxime moiety [Co(DH)₂] are normal in both crystals. The deviation of cobalt from the best plane of the four equatorial N atoms (N1–N4) is 0.050(1) and 0.059(2) Å in (1) and (2), respectively, while the twisting of the two DH

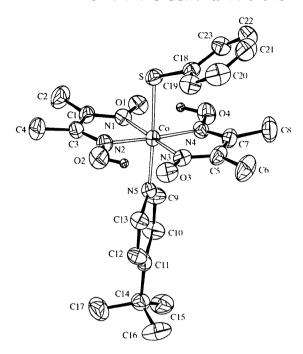


Fig. 1. The molecular structure of (1) shown with 50% probability displacement ellipsoids (*ORTEPII*; Johnson, 1976). Only the hydroxyl H atoms are shown.

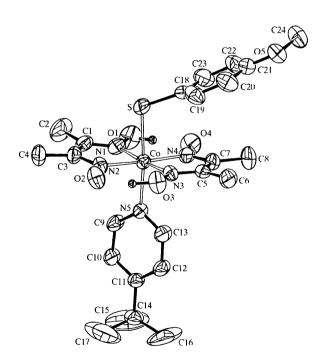


Fig. 2. The molecular structure of (2) shown with 50% probability displacement ellipsoids (*ORTEPII*; Johnson, 1976). Only the hydroxyl H atoms are shown.

planes in (1) and (2) is 9.5 (2) and 9.1 (3)°, respectively. The distances between cobalt and the axial ligands are 2.2800 (8) (Co—S) and 2.000 (2) Å (Co—N5) in (1), and 2.2885 (15) (Co—S) and 2.045 (4) Å (Co—N5) in (2). These values show clearly that the electron-donating substituent lengthens the Co—S bond. Such a trend has previously been observed in the cases of axial carbon ligands (Pahor *et al.*, 1985). The relation between the Co—S bond length and the reactivity of [Co(DH)₂(4-'BuPy)(4-X-PhS)] is satisfactorily explained by assuming that the reaction starts by the initial homolysis of the Co—S bond.

Experimental

A mixture of CoCl₂.6H₂O (6 mmol), dimethylglyoxime (DH₂, 12 mmol), and NaOH (12.8 mmol) was stirred under argon in 10 ml of degassed methanol (10 ml) for 10 min, 4-tert-Butylpyridine (6 mmol) was added and the mixture was stirred for 30 min. A solution of diaryl disulfide (5.94 mmol) in 5 ml of benzene was added and the whole was stirred at room temperature for 5 h. After filtration through a celite pad and solvent evaporation, the residue was extracted with dichloromethane $(3 \times 30 \text{ ml})$ and the extract dried over anhydrous Na₂SO₄. After removal of the solvent, the residue was purified by chromatography (Florisil, dichloromethane). Crystallization was carried out by diffusional mixing of hexane into a dichloromethane solution of the cobaloxime. Compound (1): m.p. 477-478 K; found: C 51.45, H 6.20, N 13.08%; calculated for C₂₃H₃₂CoN₅O₄S: C 51.78, H 6.05, N 13.13%. Compound (2): m.p. 482-484 K; found: C 51.22, H 6.21, N 12.39%; calculated for C₂₄H₃₄CoN₅O₅S: C 51.15, H 6.08, N 12.43%.

Compound (1)

Crystal data

$[Co(C_4H_7N_2O_2)_2(C_9H_{13}N)-$	Mo $K\alpha$ radiation
(C_6H_5S)	$\lambda = 0.71073 \text{ Å}$
$M_r = 533.53$	Cell parameters from 25
Monoclinic	reflections
$P2_1/a$	$\theta = 11.56 - 13.85^{\circ}$
a = 23.630 (6) Å	$\mu = 0.797 \text{ mm}^{-1}$
b = 10.9850 (10) Å	T = 293 (2) K
c = 9.779(3) Å	Rod
$\beta = 91.060 (10)^{\circ}$	$0.20 \times 0.20 \times 0.10 \text{ mm}$
$V = 2538.0 (10) \text{ Å}^3$	Dark green
Z = 4	
$D_x = 1.396 \text{ Mg m}^{-3}$	
D_m not measured	

Data collection

c
,

Refinement

Refinement on F^2	$w = 1/[\sigma^2(F_o^2) + (0.0433P)^2$
$R[F^2 > 2\sigma(F^2)] = 0.034$	+ 1.0976P]
$wR(F^2) = 0.090$	where $P = (F_o^2 + 2F_c^2)/3$
S = 1.076	$(\Delta/\sigma)_{\text{max}} = -0.025$
4469 reflections	$\Delta \rho_{\text{max}} = 0.280 \text{ e Å}^{-3}$
340 parameters	$\Delta \rho_{\min} = -0.315 \text{ e Å}^{-3}$
H atoms treated by a	Extinction correction: none
mixture of independent	Scattering factors from
and constrained refinement	International Tables for
	Crystallography (Vol. C)

Table 1. Selected geometric parameters (\mathring{A} , \circ) for (1)

Co—N3 Co—N4 Co—N2 Co—N1	1.888 (2) 1.890 (2) 1.892 (2) 1.893 (2)	Co—N5 Co—S S—C18	2.000 (2) 2.2800 (8) 1.761 (3)
N3—Co—N4	81.33 (9)	N2—Co—N5	91.73 (8)
N3—Co—N2	98.59 (9)	N1—Co—N5	90.62 (8)
N4—Co—N2	175.96 (9)	N3—Co—S	93.95 (7)
N3—Co—N1	178.00 (9)	N4—Co—S	88.31 (6)
N4—Co—N1	98.92 (9)	N2—Co—S	87.66 (6)
N2—Co—N1	81.01 (9)	N1—Co—S	84.08 (7)
N3—Co—N5	91.35 (8)	N5CoS	174.70 (6)
N4—Co—N5	92.31 (8)	C18SCo	111.30 (8)

Compound (2)

Crystal data

$[C_0(C_4H_7N_2O_2)_2(C_9H_{13}N)$ -	Mo $K\alpha$ radiation
(C_7H_7OS)	$\lambda = 0.71073 \text{ Å}$
$M_r = 563.55$	Cell parameters from 25
Orthorhombic	reflections
$P2_12_12_1$	$\theta = 11.64 - 13.90^{\circ}$
a = 12.8980 (10) Å	$\mu = 0.750 \text{ mm}^{-1}$
b = 18.482 (2) Å	T = 293 (2) K
c = 11.4120 (10) Å	Rod
$V = 2720.4 (4) \text{ Å}^3$	$0.20 \times 0.20 \times 0.15$ mm
Z = 4	Dark green
$D_x = 1.376 \text{ Mg m}^{-3}$	-
D_m not measured	

Data collection

Enraf-Nonius CAD-4	$\theta_{\text{max}} = 24.98^{\circ}$
diffractometer	$h = 0 \rightarrow 15$
ω –2 θ scans	$k = 0 \longrightarrow 21$
Absorption correction: none	$l = 0 \rightarrow 13$
2705 measured reflections	3 standard reflections
2705 independent reflections	frequency: 120 min
2311 reflections with	intensity decay: -1.1%
$I > 2\sigma(I)$	

Refinement

Refinement on F^2	$w = 1/[\sigma^2(F_o^2) + (0.0498P)^2]$
$R[F^2 > 2\sigma(F^2)] = 0.035$	+ 3.1260 <i>P</i>]
$wR(F^2) = 0.090$	where $P = (F_o^2 + 2F_c^2)/3$
S = 1.028	$(\Delta/\sigma)_{\text{max}} = 0.045$
2704 reflections	$\Delta \rho_{\text{max}} = 0.301 \text{ e Å}^{-3}$
363 parameters	$\Delta \rho_{\min} = -0.205 \text{ e Å}^{-3}$
H atoms treated by a	Extinction correction: none
mixture of independent	Scattering factors from
and constrained refinement	International Tables for
	Crystallography (Vol. C)

Table 2. Selected geometric parameters (Å, °) for (2)

Co-N1	1.880(4)	Co-N5	2.045 (4)
Co-N4	1.888 (4)	Co—S	2.2885 (15)
Co-N2	1.891(4)	S—C18	1.764 (5)
Co-N3	1.896 (4)		
NI-CoN4	98.4(2)	N2—Co—N5	89.86 (15)
N1—Co—N2	81.1(2)	N3—Co—N5	92.2 (2)
N4CoN2	177.2(2)	N1—Co—S	87.16 (13)
N1Co-N3	175.6(2)	N4CoS	92.42 (12)
N4CoN3	80.9(2)	N2CoS	84.78 (12)
N2—Co—N3	99.3(2)	N3—Co—S	88.47 (12)
N1—Co—N5	92.2(2)	N5—Co—S	174.64 (11)
N4—Co—N5	92.9(2)	C18—S—Co	110.2 (2)

The positions of H atoms, except those of the methyl groups, were refined, while methyl H atoms were fixed geometrically (C—H = 0.96 Å). The isotropic displacement parameters of the methyl and hydroxyl H atoms were set at 1.5 times those of the attached non-H atoms; others were set at 1.2 times. Although the *tert*-butyl group in [Co(DH)₂(4-'BuPy)(4-MeO-PhS)], (2), is highly disordered, only one conformer was included in calculations because trials on several disorder models were not successful.

For both compounds, data collection: *CAD-4 Software* (Enraf–Nonius, 1989); cell refinement: *CAD-4 Software*; data reduction: *MolEN* (Fair, 1990). Program(s) used to solve structures: *SHELXS86* (Sheldrick, 1990) for (1); *MolEN* for (2). For both compounds, program(s) used to refine structures: *SHELXL*93 (Sheldrick, 1993); molecular graphics: *ORTEPII* (Johnson, 1976).

Supplementary data for this paper are available from the IUCr electronic archives (Reference: OA1029). Services for accessing these data are described at the back of the journal.

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