# metal-organic compounds

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# Mixed cobalt(III) complexes with aromatic amino acids and diamines. III.<sup>1</sup> Absolute structure of  $\Lambda$ -trans(O)-(1,2-diaminoethane- $\kappa^2$ N,N)bis(S-tyrosinato)cobalt(III) chloride tetrahydrate

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The title compound,  $[Co(C_9H_{10}NO_3)_2(C_2H_8N_2)]Cl·4H_2O$ , is one of six possible diastereomers of the (1,2-diaminoethane) bis(S-tyrosinato)cobalt(III) complex. The cobalt(III) ion has an octahedral coordination, with three five-membered chelate rings which have deformed coordination angles and coordinated O atoms in trans positions. In comparison with the previously reported crystal structure of the  $\Delta$ -C<sub>1</sub>-cis(O) diastereomer [Miodragović et al. (2001). Enantiomer, 6, 299-308], the compound presented in this paper has more planar five-membered aminocarboxylate rings. Complex cations, chloride anions and water molecules of crystallization are linked together by a network of hydrogen bonds. The chloride anions lie approximately between two Co atoms and form hydrogen bonds with all coordinated  $NH<sub>2</sub>$  groups. In the crystal structure, there is a weak intermolecular  $\pi \cdots \pi$ interaction between the phenyl rings.

### Comment

The crystal structure of the title compound, (I), represents further work in our investigation of the structures and stereochemistries of cobalt(III) complexes with aromatic amino acids (Miodragovic et al., 2001). Complexes of cobalt(III) with aromatic amino acids are interesting as simple model systems for the investigation of non-covalent interactions in which aromatic amino acids are involved (Jitsukawa et al., 1997; Kumita et al., 1998, 2001), because it is known that their side chains, through non-covalent interactions, play an important role in molecular recognition in vivo (Fontecave & Pierre, 1996; Ma & Dougherty, 1997; Pomponi et al., 1998;

Hofstädter et al., 1999). Mixed bis(S-aminocarboxylato)(1,2diaminoethane)cobalt(III) complexes can occur in the form of three geometrical isomers, and each of them has an optical counterpart.



In a previous paper (Miodragović et al., 2001), we described the synthesis and characterization of five out of six theoretically possible diastereomers of the (1,2-diaminoethane)bis- (S-tyrosinato)cobalt(III) complex and presented the X-ray crystal structure of the  $\Delta$ -C<sub>1</sub>-cis(O) diastereomer, (II). By a subsequent recrystallization of the diastereomers, high quality single crystals of the  $\Lambda$ -trans(O) diastereomer, (I), were obtained, the crystal structure of which is presented in this paper (Fig. 1).

The greatest differences between complexes (I) and (II) are in the different coordinations of the O and N atoms of the S-tyrosinate ligands,  $viz$ . complex (II) has a  $cis(O), cis(N)$ configuration, while complex  $(I)$  has a trans $(O)$ ,trans $(N)$ configuration, leading to different  $Co-O$  bond lengths. Hence, in complex (I),  $Co-O1 = 1.886$  (2) Å and  $Co-O4 =$ 1.892 (2)  $\AA$ , which are shorter than the distances in (II) [Co– O1 = 1.912 (6)  $\AA$  and Co $-$ O4 = 1.909 (5)  $\AA$ ]. There are also differences in the  $Co-N(S$ -tyrosinate) bond lengths, *i.e.* in (II), they are different  $[Co-N1 = 1.939 (7)$  Å and  $Co-N1 =$ 1.965 (6)  $\AA$ , while in the title complex, (I), they are almost the same  $[Co-N1 = 1.959 (2)$  Å and  $Co-N2 = 1.960 (2)$  Å].

In addition to the differences in coordination, the complexes also exhibit differences in the conformation of the amino acid chelate rings; namely, although these chelate rings exhibit the same conformation (envelope), in (I), the rings are less puckered. The puckering parameters for the  $Co-O1$ 





A view of the molecular structure of (I) with the atom labelling. <sup>1</sup> For part II, see Miodragovic *et al.* (2001). Displacement ellipsoids are shown at the  $50\%$  probability level.

C1–C2–N1 (q2) and Co–O4–C10–C11–N2 (q2') chelate rings for complex (I) are  $q2 = 0.267(2)$  Å and  $q2' =$ 0.230 (3)  $\AA$ , while for complex (II), the puckering parameters are  $q2 = 0.365$  (8) Å and  $q2' = 0.323$  (7) Å (Cremer & Pople, 1975). Such a difference in the planarity of the amino acid chelate rings leads to somewhat smaller  $H-N1-C2-H$  and  $H-N2-C11-H$  torsion angles in complex (I), *i.e.* the H atoms bonded to chelate-ring N and C atoms are closer to being eclipsed in complex (I).

The puckering parameters of the diamine five-membered chelate ring  $(Co-N3-C19-C20-N4)$  are  $q2 = 0.446$  (3) A and  $\varphi = -99.0 \,(3)^{\circ}$ . There are no remarkable differences between the conformations of these chelate rings in the two complexes. In both cases, the chelate ring adopts a conformation which is intermediate between half-chair (with a local pseudo-twofold axis along Co and the midpoint of the C19 $-$ C20 bond) and envelope (with a local pseudo-mirror along C19 and the midpoint of the  $Co-N4$  bond).

Due to the formation of three five-membered chelate rings, the coordinate angles deviate markedly from the ideal values of 90 and  $180^{\circ}$  (Table 1). The crystal structure is stabilized by a large number of hydrogen bonds (Table 2). In the crystal lattice of (I), the chloride anions are located approximately between two neighbouring Co atoms  $[Co \cdots Cl = 4.120 (2)$  Å and  $Cl \cdots Co^{i} = 4.132$  (2) Å; symmetry code: (i)  $x - 1$ , y, z], forming hydrogen bonds with the  $NH<sub>2</sub>$  groups coordinated to the Co atoms. However, in the crystal lattice of complex (II), the complex cations also form intermolecular hydrogen bonds, but these are directly through the  $-C(O)O$  and  $-NH<sub>2</sub>$  groups from the coordination sphere of the Co atom.

In the crystal lattice of complex (I), there is a weak intermolecular  $\pi \cdot \cdot \pi$  interaction between the phenyl rings of the amino acid ligands. The phenyl rings are oriented in such a way that the perpendicular distance from the C4–C9 ring to the C13<sup>i</sup>-C18<sup>i</sup> ring is 3.88 Å, with the closest distance being  $CS \cdot C17^i$  of 3.672 (6) Å. The distance between the ring centroids is 4.142 (3)  $\AA$ . An unhomogeneous arrangement of  $\pi$ -electron density through the phenyl ring is probably responsible for the mutual orientation of the phenyl rings



#### Figure 2

Two projections of the tyrosinate phenyl-ring fragments, showing the  $\pi \cdot \pi$  interactions between the rings. Displacement ellipsoids are shown at the 30% probability level.

(Fig. 2) and the parallel-displaced (PD) type of  $\pi \cdot \cdot \pi$  interactions (Kamishima et al., 2001).

It was subsequently established (i.e. it had not been published earlier) that in (II), the phenyl rings of tyrosine are involved in intermolecular  $C-H \cdots \pi$  interactions with the following geometrical parameters: (i) the distance between an H atom bonded to atom C14 and the centre of the C4-C9 aromatic ring is  $2.67 \text{ Å}$ ; (ii) the distance between an H atom bonded to atom C14 and the plane of the C4–C9 aromatic ring is 2.66  $\AA$ ; (iii) the angle between the line connecting an H atom and the centre of the ring,  $M$ , and the normal to the C4 $-$ C9 plane is 4.6°; (iv) the angle C14 $-H \cdots M$  is 172°.

From the results presented in this paper, it can be concluded that the phenyl rings of amino acids which are present in the same complex, but which have different geometrical configurations, can form two kinds of interactions, namely  $C-H \cdot \cdot \pi$ and  $\pi \cdot \cdot \pi$  interactions. It can also be concluded that in complexes (I) and (II), the complex cations form intermolecular hydrogen bonds of different types. In complex (II), the hydrogen bonds are formed through coordinated  $-C(O)O$ and  $-NH<sub>2</sub>$  groups, while in complex (I), a chloride anion is located approximately between two complex cations and forms multiple hydrogen bonds with the coordinated  $NH<sub>2</sub>$ groups.

## Experimental

The title diastereomer was obtained by the action of the S-tyrosinate ligand on trans-dichlorobis(1,2-diaminoethane)cobalt(III) chloride according to Miodragović et al. (2001). Red crystals of  $(I)$  were recrystallized from water in a 62% yield.

#### Crystal data



# Table 1

Selected geometric parameters  $(\AA, \degree)$ .

$Co-O1$	1.886(2)	$O2 - C1$	1.228(4)
$Co-O4$	1.892(2)	$N1 - C2$	1.487(4)
$Co-N1$	1.959(2)	$C1 - C2$	1.526(4)
$Co-N2$	1.960(2)	$O4 - C10$	1.281(4)
$Co-N3$	1.950(2)	$O5 - C10$	1.235(4)
$Co-N4$	1.957(2)	$N2 - C11$	1.499(4)
$O1 - C1$	1.290(4)	$C10 - C11$	1.527(4)
$O1 - Co - O4$	175.55(10)	$O4 - Co - N4$	92.41 (10)
$O1 - Co - N1$	84.79 (9)	$N1 - Co - N2$	93.81 (11)
$O1 - Co - N2$	91.77(10)	$N1 - Co - N3$	90.49(11)
$O1 - Co - N3$	92.66(11)	$N1 - Co - N4$	173.59 (11)
$O1 - Co - N4$	90.77(11)	$N2 - Co - N3$	174.08 (14)
$O4 - Co - N1$	92.29(10)	$N2 - Co - N4$	90.94 (11)
$O4 - Co - N2$	85.07 (10)	$N3-Co-N4$	85.09 (11)
$O4 - Co - N3$	90.71 (10)		

### Table 2





Symmetry codes: (i)  $-x, \frac{1}{2} + y, 1 - z$ ; (ii)  $1 + x, y, z$ ; (iii)  $1 - x, y - \frac{1}{2}, 1 - z$ ; (iv)  $-x, \frac{1}{2} + y, 2 - z$ ; (vi)  $x - 1, y, z$ ; (vii)  $1 - x, y - \frac{1}{2}, 2 - z$ .

#### Data collection



#### Refinement



All H atoms of the complex cation were placed at calculated positions (C–H = 0.93–0.98 Å and N–H = 0.90 Å) using a riding model and isotropic displacement parameters were set equal to 1.2 times the equivalent isotropic displacement parameter of the parent atoms. Water and hydroxyl H-atom positions were determined by the HYDROGEN program (Nardelli, 1999) and were refined using a riding model with a fixed O–H bond length of 0.85 Å; isotropic displacement parameters were set equal to 1.5 times the equivalent isotropic displacement parameter of the parent atoms. It should be noted that all calculated data for water and hydroxyl H atoms are based on an approximate model. A Gaussian-type absorption correction based on the crystal morphology was applied (Spek, 1990, 1998).

Data collection: CAD-4 Software (Enraf-Nonius, 1989); cell refinement: CAD-4 Software; data reduction: CAD-4 EXPRESS (Enraf±Nonius, 1994); program(s) used to solve structure:  $SHELX$ S97 (Sheldrick, 1997); program(s) used to refine structure: SHELXL97 (Sheldrick, 1997); molecular graphics: ORTEPIII (Burnett & Johnson, 1996); software used to prepare material for publication: SHELXL97, PARST (Nardelli, 1995) and WinGX (Farrugia, 1999).

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Supplementary data for this paper are available from the IUCr electronic archives (Reference: NA1562). Services for accessing these data are described at the back of the journal.

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