metal-organic papers

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Key indicators

Single-crystal X-ray study T = 293 KMean $\sigma(C-C) = 0.005 \text{ Å}$ H-atom completeness 51% R factor = 0.040 wR factor = 0.106 Data-to-parameter ratio = 11.4

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

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Dipotassium disodium *trans*-bis(pyrazine-2,3-dicarboxylato)copper(II) dithiocyanate dihydrate

Reaction of pyrazine-2,3-dicarboxylic acid, NaOH, KSCN and $Cu(NO_3)_2 \cdot 3H_2O$ led to the formation of $Na_2K_2[trans-Cu(C_6H_2N_2O_4)_2](SCN)_2 \cdot 2H_2O$, in which the Cu^{II} atom, on an inversion centre, is approximately in a square-planar environment. This is a new coordination geometry for copper in complexes of pyrazine-2,3-dicarboxylate.

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Comment

Takusagawa & Shimada (1973) first determined the structure of pyrazine-2,3-dicarboxlic acid by single-crystal X-ray analysis. Almost at the same time, the first metal-organic compound of pyrazine-2,3-dicarboxylic acid was reported (Richard *et al.*, 1973). More recently, owing to the interest in supramolecular chemistry, pyrazine-2,3-dicarboxylic acid has been intensely investigated (Smith *et al.*, 1995; Neels *et al.*, 1997; Kondo *et al.*, 1999). For further investigation of pyrazine-2,3-dicarboxylic acid, we synthesized the title compound, (I), with a new coordination mode for copper in such complexes. An interesting phenomenon was noted during the synthesis; a blue solid precipitated when pyrazine-2,3-dicarboxylate was added to Cu(NO₃)₂·3H₂O in water, but no product was deposited in the present of SCN⁻. Here we report the synthesis and crystal structure of the title compound.



Compound (I) is composed of K⁺ and Na⁺ cations, [*trans*-Cu(C₆H₂N₂O₄)₂]²⁻ and SCN⁻ anions, and water molecules. As shown in Fig. 1, the Cu^{II} atom has centrosymmetric squareplanar coordination with *trans*-positioned O- and N-donor atoms of two pyrazine-2,3-dicarboxylate ligands. The bond distances Cu1-O3 and Cu1-N3 are 1.971 (3) and 1.970 (3) Å, respectively, almost equal to each other, unlike those found in other Cu^{II} complexes of pyrazine-2,3-dicarboxylate (Smith *et al.*, 1995; Neels *et al.*, 1997; Kondo *et al.*, 1999). As in the structures of C₁₄H₈CuN₂O₈·2H₂O (Yucang, Maochun, Rong & Qian, 2001) and (C₂₈H₂₄CuGd₂. N₄O₂₂)_n·2nH₂O (Yucang, Maochun, Weiping *et al.*, 2001), the C-O bond distance of the carboxylate group involving the O atom coordinated to copper in the present compound [C6-O3 = 1.276 (4) Å] is a little longer than the other [C6-O4 =



Figure 1

The structure of the $[trans-Cu(C_6H_2N_2O_4)_2]^{2-}$ complex anion, shown with 30% probability ellipsoids. H atoms have been omitted for clarity.

1.237 (4) Å]. However, the two C–O bond distances of the uncoordinated carboxylate group are almost equal, as found in the structure of $(C_{14}H_{22}CuN_2Na_2O_{16})_n$ (Sileo *et al.*, 1999).

As shown in Fig. 2, the three-dimensional framework is constructed through electrostatic interactions among [*trans*-Cu(C₆H₂N₂O₄)₂]²⁻, SCN⁻, K⁺ and Na⁺ ions, together with water molecules. O–H···N hydrogen bonds are found between O1W and the uncoordinated N atom of pyrazine-2,3-dicarboxylate with a distance of 2.985 (5) Å. Na is six-coordinated by five O atoms and one N atom, while K is eight-coordinated by seven O atoms and one N atom (Table 1).

Experimental

5 ml of a stirred aqueous solution of pyrazine-2,3-dicarboxylic acid (84 mg, 0.5 mmol) was adjusted to pH *ca* 7 with 1 N NaOH aqueous solution, then KSCN (49 mg, 0.5 mmol) was added. After complete dissolution, $Cu(NO_3)_2$ ·3H₂O (60 mg, 0.25 mmol) was added, giving a clear blue solution, which was filtered. Upon evaporation in air, blue prismatic crystals were obtained in a few days (yield 58%).

Crystal data

Na ₂ K ₂ [Cu(C ₆ H ₂ N ₂ O ₄) ₂]- (SCN) ₂ ·2H ₂ O $M_r = 672.10$ Monoclinic, P2 ₁ /c a = 6.8349 (7) Å b = 6.9692 (7) Å c = 23.184 (2) Å $\beta = 90.797$ (2)° V = 1104.23 (19) Å ³ Z = 2	$D_x = 2.021 \text{ Mg m}^{-3}$ Mo K α radiation Cell parameters from 2150 reflections $\theta = 1.8-25.0^{\circ}$ $\mu = 1.66 \text{ mm}^{-1}$ T = 293 (2) K Prism, blue $0.68 \times 0.22 \times 0.18 \text{ mm}$
Data collection	
Siemens SMART CCD diffractometer ω scans Absorption correction: multi-scan (<i>SADABS</i> ; Sheldrick, 1996) $T_{min} = 0.514, T_{max} = 0.741$ 3306 measured reflections	1929 independent reflections 1663 reflections with $I > 2\sigma(I)$ $R_{int} = 0.025$ $\theta_{max} = 25.0^{\circ}$ $h = -8 \rightarrow 8$ $k = -6 \rightarrow 8$ $l = -27 \rightarrow 9$



Figure 2

The crystal structure of the title compound, viewed along the b axis. H atoms have been omitted and the coordination environments of only one Na ion and one K ion are shown (dotted lines). Dashed lines represent hydrogen bonds.

Refinement

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Refinement on F^2	$w = 1/[\sigma^2(F_o^2) + (0.0473P)^2]$
$R[F^2 > 2\sigma(F^2)] = 0.040$	+ 3.7091P]
$vR(F^2) = 0.106$	where $P = (F_o^2 + 2F_c^2)/3$
S = 1.00	$(\Delta/\sigma)_{\rm max} < 0.001$
929 reflections	$\Delta \rho_{\rm max} = 0.50 \ {\rm e} \ {\rm \AA}^{-3}$
69 parameters	$\Delta \rho_{\rm min} = -0.64 \text{ e } \text{\AA}^{-3}$
I-atom parameters constrained	

Table 1

Selected geometric parameters (Å, °).

Cu1-N3	1.970 (3)	Na1-N1	2.370 (4)
Cu1-O3	1.971 (3)	Na1-O2 ⁱ	2.413 (3)
K1-O4 ⁱ	2.776 (3)	Na1 - O1W	2.445 (4)
K1-O2 ⁱ	2.820 (3)	Na1-O4	2.465 (3)
K1-O4	2.882 (3)	Na1-O1 ^{vi}	2.596 (4)
K1-O1 ⁱⁱ	2.906 (3)	Na1-O1 ⁱ	2.710 (3)
$K1 - O1W^{iii}$	2.912 (4)	C6-O4	1.237 (4)
$K1-O2^{iv}$	2.928 (3)	C6-O3	1.276 (4)
K1-N1 ^v	3.018 (4)	C7-O2	1.248 (5)
K1-O1	3.239 (3)	C7-O1	1.256 (5)
N3-Cu1-O3	82.42 (11)	O2-C7-O1	126.8 (3)
O4-C6-O3	125.0 (3)	O2-C7-C4	118.4 (3)
O4-C6-C5	120.4 (3)	C6-O3-Cu1	114.0 (2)
	1.1 (**) 2	1.1 (***).4	1. 1. (1)

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

H atoms were positioned geometrically and refined as riding on their parent atoms; no H atoms were included for the water molecules.

Data collection: *SMART* (Siemens, 1996); cell refinement: *SMART* and *SAINT* (Siemens, 1994); data reduction: *SAINT* and *SHELXTL* (Siemens, 1994); program(s) used to solve structure: *SHELXTL*; program(s) used to refine structure: *SHELXTL*; molecular graphics: *SHELXTL*; software used to prepare material for publication: *SHELXTL*.

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