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Key indicators

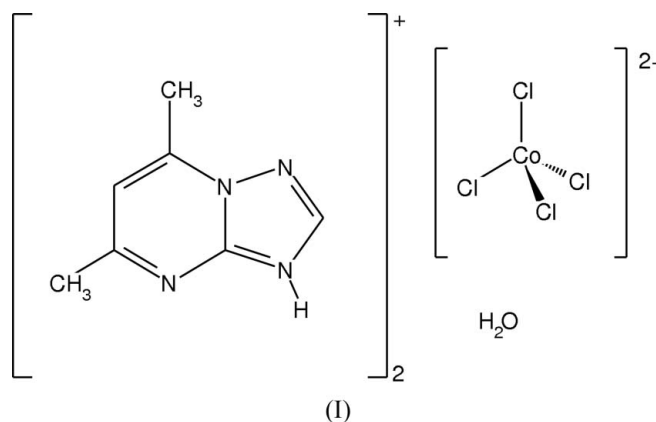
Single-crystal X-ray study
 $T = 298$ K
Mean $\sigma(\text{C}-\text{C}) = 0.003$ Å
 R factor = 0.031
 wR factor = 0.080
Data-to-parameter ratio = 18.5For details of how these key indicators were automatically derived from the article, see <http://journals.iucr.org/e>.5,7-Dimethyl-1,2,4-triazolo[1,5-*a*]pyrimidinium tetrachlorocobaltate(II) monohydrate

The asymmetric unit of the title compound, $(\text{C}_7\text{H}_8\text{N}_4)_2[\text{CoCl}_4]\cdot\text{H}_2\text{O}$, contains one $[\text{CoCl}_4]^{2-}$ anion, two 5,7-dimethyl-1,2,4-triazolo[1,5-*a*]pyrimidinium cations monoprotonated at the N atom at position 3 and one uncoordinated water molecule. Hydrogen bonds define infinite chains along the *a* axis, which link together all residues.

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Comment

1,2,4-Triazolo-[1.5-*a*]pyrimidine derivatives are well known N-donor heterocyclic ligands and have been extensively studied by us, as well as by other research groups (Salas *et al.*, 1999). Some of these ligands are known to be unstable in an acidic medium, where they decompose giving 3-amino-1,2,4-triazole; this decomposition seems to be promoted by the presence of transition metal ions (Sánchez *et al.*, 2004). Other 1,2,4-triazolo[1.5-*a*]pyrimidine derivatives, however, are fairly stable, even at very low pH. The 5,7-dimethyl derivative (dmtp), for example, is easily stabilized as a cationic species protonated at N3, the charge of this cation being balanced by anionic metal complexes such as $[\text{CdBr}_4]^{2-}$ (Salas *et al.*, 1996), $[\text{SnCl}_6]^{2-}$ (Salas *et al.*, 1998) or $[\text{PtCl}_6]^{2-}$ (Szlyk *et al.*, 2002). The structure, which is reported in this paper, adds one more salt to this series, where the charge of the cation is balanced by the tetrachlorocobaltate(II) anion.



The crystal structure of the title compound, (I), is made up of discrete $[\text{CoCl}_4]^{2-}$ anions, 5,7-dimethyl-1,2,4-triazolo-[1,5-*a*]pyrimidinium (Hdmtp^+) cations and uncoordinated water molecules. All residues occupy general positions, the asymmetric unit containing one anion, two crystallographically independent but chemically identical cations and one water molecule. A view of the asymmetric unit is shown in Fig. 1.

The anion displays a considerable distortion from the ideal T_d symmetry (see distances and angles in Table 1), a normal

behaviour for $[\text{CoCl}_4]^{2-}$, which seems to be a very flexible species, and according to the statistical analysis by Haddad *et al.* (2003) exhibits substantial variability in Cl—Co—Cl bond angles. The average Co—Cl bond distance in the title compound, 2.2711 Å, is quite similar to the overall average value of 2.2725 Å, as reported by Haddad *et al.* (2003).

The site of protonation of the organic residue at N3 is clear from the ΔF maps and from the strong hydrogen bonds formed with the water molecule or one of the Cl atoms of the anion as acceptors (see Table 2). The main difference in the geometry of protonated dmp compared with that of the neutral species is the widening of the endocyclic bond angle at N3, which is 106.8 (2) and 106.4 (2)° in the two independent cations of the title compound, whereas the corresponding angle in free dmp is 102.2 (3)° (Salas *et al.*, 1994). Such widening of the bond angle at the protonated N atom is quite common and was also observed in the above-mentioned ionic structures containing dmpH^+ .

It is interesting to mention the existence of a rather short contact between the Cl atoms of the anion and the bridgehead N atoms of the cations, N8A and N8B. These contacts [Cl1...N8B = 3.287 (2) Å and Cl2...N8A = 3.235 (1) Å] are most probably of electrostatic nature, the Cl1...N8B and Cl2...N8A vectors being approximately orthogonal to the plane of the heterocycle; the normal to this plane forms angles of 2.89 (3) and 4.86 (2)° with Cl1...N8B and Cl2...N8A, respectively. In the structure of the title compound, there are also several hydrogen bonds, involving the H atoms at N3A and N3B (see above) as well the H atoms of the water molecule (Table 2). These hydrogen bonds link together all residues, defining infinite chains running along the *a* axis.

Experimental

An aqueous solution of cobalt(II) chloride hexahydrate (1 mmol, 10 ml) and a solution of dmp in 1:1 HCl (2 mmol, 20 ml) were mixed. The mixture was allowed to evaporate at room temperature and, after 24 h, blue prismatic crystals of the title compound were obtained. Analysis found: C 32.48, H 4.23, N 21.75%; calculated: C 32.52, H 3.90, N 21.67%.

Crystal data

$2\text{C}_7\text{H}_9\text{N}_4^+\cdot\text{Cl}_4\text{Co}^{2-}\cdot\text{H}_2\text{O}$	$Z = 2$
$M_r = 517.11$	$D_x = 1.600 \text{ Mg m}^{-3}$
Triclinic, $P\bar{1}$	Mo $K\alpha$ radiation
$a = 7.6369 (4) \text{ \AA}$	Cell parameters from 5772 reflections
$b = 8.5190 (4) \text{ \AA}$	$\theta = 2.2\text{--}27.2^\circ$
$c = 18.2777 (9) \text{ \AA}$	$\mu = 1.32 \text{ mm}^{-1}$
$\alpha = 84.952 (1)^\circ$	$T = 298 (2) \text{ K}$
$\beta = 84.505 (1)^\circ$	Prism, blue
$\gamma = 65.275 (1)^\circ$	$0.46 \times 0.29 \times 0.23 \text{ mm}$
$V = 1073.64 (9) \text{ \AA}^3$	

Data collection

Bruker SMART APEX CCD diffractometer	4866 independent reflections
φ and ω scans	4217 reflections with $I > 2\sigma(I)$
Absorption correction: multi-scan (SADABS; Bruker, 1999)	$R_{\text{int}} = 0.019$
$T_{\text{min}} = 0.668$, $T_{\text{max}} = 0.751$	$\theta_{\text{max}} = 28.3^\circ$
12624 measured reflections	$h = -9 \rightarrow 10$
	$k = -11 \rightarrow 11$
	$l = -24 \rightarrow 23$

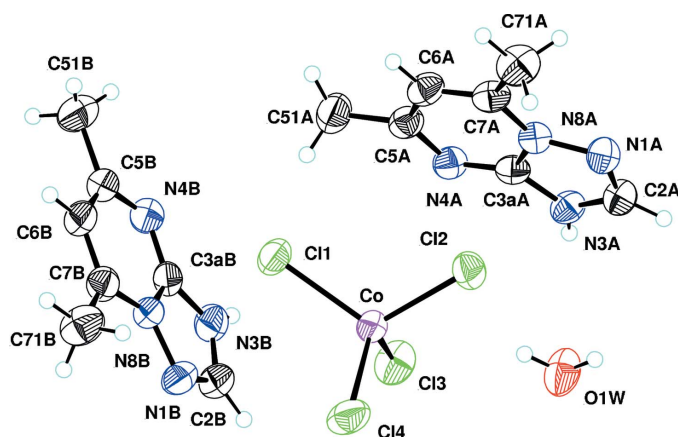


Figure 1

The asymmetric unit of the title compound, with the non-H atoms represented by 50% probability displacement ellipsoids; H atoms are shown as circles of arbitrary radius.

Refinement

Refinement on F^2	$w = 1/[\sigma^2(F_o^2) + (0.04P)^2 + 0.25P]$
$R[F^2 > 2\sigma(F^2)] = 0.031$	where $P = (F_o^2 + 2F_c^2)/3$
$wR(F^2) = 0.080$	$(\Delta/\sigma)_{\text{max}} = 0.002$
$S = 1.04$	$\Delta\rho_{\text{max}} = 0.45 \text{ e \AA}^{-3}$
4866 reflections	$\Delta\rho_{\text{min}} = -0.36 \text{ e \AA}^{-3}$
263 parameters	
H atoms treated by a mixture of independent and constrained refinement	

Table 1

Selected geometric parameters (Å, °).

Co—Cl1	2.2898 (5)	Co—Cl3	2.2583 (6)
Co—Cl2	2.2697 (5)	Co—Cl4	2.2666 (6)
Cl3—Co—Cl4	112.31 (3)	Cl3—Co—Cl1	112.49 (2)
Cl3—Co—Cl2	107.13 (2)	Cl4—Co—Cl1	104.59 (2)
Cl4—Co—Cl2	108.47 (3)	Cl2—Co—Cl1	111.86 (2)

Table 2

Hydrogen-bond geometry (Å, °).

$D\text{—}H\cdots A$	$D\text{—}H$	$H\cdots A$	$D\cdots A$	$D\text{—}H\cdots A$
N3A—H3A...O1W ⁱ	0.86	1.83	2.681 (2)	171
N3B—H3B...Cl1	0.86	2.31	3.1325 (18)	160
O1W—H11W...N1A ⁱⁱ	0.84 (1)	2.19 (1)	3.026 (2)	169 (2)
O1W—H12W...Cl2	0.84 (1)	2.59 (2)	3.3122 (15)	144 (2)

Symmetry codes: (i) $-x, -y + 1, -z$; (ii) $-x + 1, -y + 1, -z$.

H atoms of the organic cations were idealized (C—H 0.93 and 0.96 Å, and N—H = 0.86 Å). Those of the water molecule were located in a ΔF map and refined with restrained O—H [0.86 (1) Å] and H...H distances [1.36 (1) Å]. Isotropic displacement parameters of all H atoms were set to be 1.2 times U_{eq} of their parent atoms.

Data collection: SMART (Bruker, 1999); cell refinement: SAINT (Bruker, 1999); data reduction: SAINT; program(s) used to solve structure: SHELXS97 (Sheldrick, 1997); program(s) used to refine

structure: *SHELXL97* (Sheldrick, 1997); molecular graphics: *Xtal_GX* (Hall & du Boulay, 1997); software used to prepare material for publication: *SHELXL97*.

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