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Key indicators

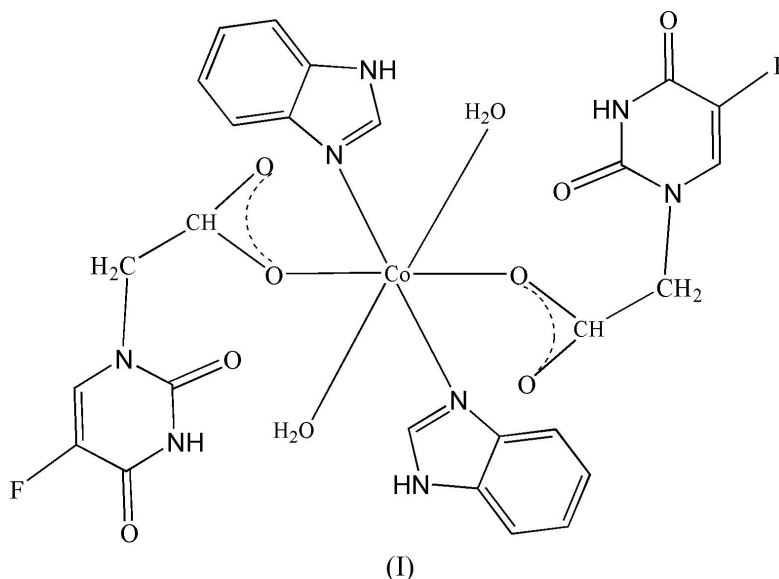
Single-crystal X-ray study
 $T = 298\text{ K}$
Mean $\sigma(\text{C}-\text{C}) = 0.005\text{ \AA}$
 R factor = 0.054
 wR factor = 0.142
Data-to-parameter ratio = 11.3For details of how these key indicators were
automatically derived from the article, see
<http://journals.iucr.org/e>.Bis(aqua)bis(benzimidazole- κN)-
bis(5-fluorouracil-1-acetate- κO)cobalt(II)

In the centrosymmetric molecule of the title compound, $[\text{Co}(\text{C}_6\text{H}_4\text{N}_2\text{O}_4\text{F})_2(\text{C}_7\text{H}_6\text{N}_2)_2(\text{H}_2\text{O})_2]$, the Co^{II} ion is coordinated by two 5-fluorouracil-1-acetate anions *via* carboxylate O atoms, two water molecules and two benzimidazole ligands, forming a six-coordinate octahedral environment. $\text{N}-\text{H}\cdots\text{O}$ and $\text{O}-\text{H}\cdots\text{O}$ hydrogen-bond interactions are observed in the structure, leading to the formation of a three-dimensional network.

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Comment

5-Fluorouracil-1-acetic acid is a derivative of 5-fluorouracil (5FU) from the pyrimidine family of bases. It is a cytotoxic analogue of the natural base thymine, which has proved useful in the chemotherapy of a number of cancers, particularly colorectal cancer (Markova Venelin Enchev & Timtcheva, 2005). Increasing attention has been paid to the anticancer activity of 5FU and its derivatives (Akgerman & Guney, 2000), but only a few of their transition metal complexes have been reported so far (Wang *et al.*, 1993). In an extension of this research, we report here the crystal structure of the title compound, (I).



The mononuclear molecule of (I) consists of a Co^{II} ion, two coordinated water molecules, two 5-fluorouracil-1-acetate anions coordinated through carboxylate O atoms, and two benzimidazole molecules. The Co^{II} cation lies on an inversion centre and the geometry around the Co^{II} ion is octahedral (Fig. 1 and Table 1). The equatorial square plane is formed by atoms O1, O1ⁱ, N3 and N3ⁱ atoms [symmetry code: (i) $-x, -y,$

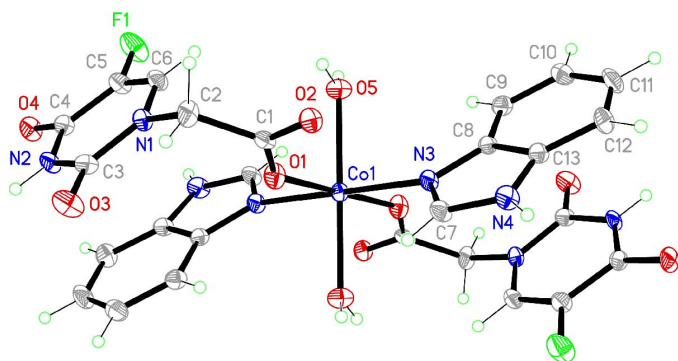


Figure 1

The coordination environment of the Co^{II} ion in (I), showing the atom-numbering scheme. Displacement ellipsoids are drawn at the 50% probability level and H atoms are shown as small spheres of arbitrary radii. Unlabelled atoms are related to labelled atoms by $-x, -y, 1-z$.

$1-z$], and is essentially planar (r.m.s. deviation 0.0011 \AA). The two axial sites are occupied by atoms O5 and O5ⁱ, with a Co—O bond distance of $2.173(3) \text{ \AA}$, longer than the average equatorial Co—O bond distance of $2.081(3) \text{ \AA}$.

Each molecule is linked to four adjacent molecules *via* N—H \cdots O and O—H \cdots O hydrogen-bond interactions, to form a three-dimensional network (Fig. 2 and Table 2).

Experimental

The title compound was synthesized in a hydrothermal process from a mixture of benzimidazole (2 mmol, 0.24 g), $\text{CoCl}_2 \cdot 2\text{H}_2\text{O}$ (1 mmol, 0.16 g), 5-fluorouracil-1-acetic acid (2 mmol, 0.75 g) and water (20 ml) in a 30 ml Teflon-lined stainless steel reactor. The solution was heated to 421 K for 5 d. After the reaction, the system was slowly cooled to room temperature, and red crystals of (I) were collected and washed with distilled water.

Crystal data

$[\text{Co}(\text{C}_6\text{H}_4\text{N}_2\text{O}_4\text{F})_2(\text{C}_7\text{H}_6\text{N}_2)_2 \cdot (\text{H}_2\text{O})_2]$	$Z = 1$
$M_r = 705.46$	$D_x = 1.677 \text{ Mg m}^{-3}$
Triclinic, $P\bar{1}$	Mo $K\alpha$ radiation
$a = 7.368(13) \text{ \AA}$	Cell parameters from 2778 reflections
$b = 8.368(14) \text{ \AA}$	$\theta = 2.6\text{--}24.9^\circ$
$c = 12.07(2) \text{ \AA}$	$\mu = 0.70 \text{ mm}^{-1}$
$\alpha = 88.59(3)^\circ$	$T = 298(2) \text{ K}$
$\beta = 88.38(3)^\circ$	Block, red
$\gamma = 69.92(3)^\circ$	$0.43 \times 0.33 \times 0.23 \text{ mm}$
$V = 699(2) \text{ \AA}^3$	

Data collection

Bruker APEX area-detector diffractometer	2410 independent reflections
φ and ω scans	2064 reflections with $I > 2\sigma(I)$
Absorption correction: multi-scan (SADABS; Bruker, 2002)	$R_{\text{int}} = 0.035$
$T_{\text{min}} = 0.752, T_{\text{max}} = 0.855$	$\theta_{\text{max}} = 25.0^\circ$
4448 measured reflections	$h = -8 \rightarrow 8$
	$k = -9 \rightarrow 9$
	$l = -14 \rightarrow 14$

Refinement

Refinement on F^2	H-atom parameters constrained
$R[F^2 > 2\sigma(F^2)] = 0.054$	$w = 1/[\sigma^2(F_o^2) + (0.0997P)^2]$
$wR(F^2) = 0.142$	where $P = (F_o^2 + 2F_c^2)/3$
$S = 1.06$	$(\Delta/\sigma)_{\text{max}} < 0.001$
2410 reflections	$\Delta\rho_{\text{max}} = 1.26 \text{ e \AA}^{-3}$
214 parameters	$\Delta\rho_{\text{min}} = -0.87 \text{ e \AA}^{-3}$

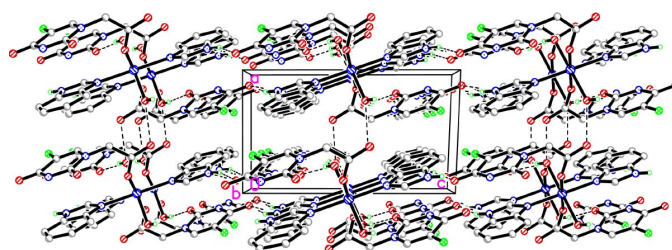


Figure 2

The three-dimensional network of (I), formed by hydrogen-bonding interactions, which are shown as dashed lines. H atoms not involved in these interactions have been omitted for clarity.

Table 1

Selected geometric parameters ($\text{\AA}, ^\circ$).

Co1—O1	2.081(3)	Co1—N3	2.125(4)
Co1—O1 ⁱ	2.081(3)	Co1—O5 ⁱ	2.173(3)
Co1—N3 ⁱ	2.125(4)	Co1—O5	2.173(3)
O1—Co1—O1 ⁱ	180.00(7)	N3 ⁱ —Co1—O5 ⁱ	90.65(12)
O1—Co1—N3 ⁱ	86.88(11)	N3—Co1—O5 ⁱ	89.35(12)
O1 ⁱ —Co1—N3 ⁱ	93.12(11)	O1—Co1—O5	90.38(16)
O1—Co1—N3	93.12(11)	O1 ⁱ —Co1—O5	89.62(16)
O1 ⁱ —Co1—N3	86.88(11)	N3 ⁱ —Co1—O5	89.35(12)
N3 ⁱ —Co1—N3	180.00(4)	N3—Co1—O5	90.65(12)
O1—Co1—O5 ⁱ	89.62(16)	O5 ⁱ —Co1—O5	180.000(1)
O1 ⁱ —Co1—O5 ⁱ	90.38(16)		

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

Table 2

Hydrogen-bond geometry ($\text{\AA}, ^\circ$).

$D\text{---}H\cdots A$	$D\text{---}H$	$H\cdots A$	$D\cdots A$	$D\text{---}H\cdots A$
N4—H4 \cdots O4 ⁱⁱ	0.86	2.12	2.905(5)	152
O5—H5B \cdots O3 ⁱⁱⁱ	0.82	2.07	2.877(4)	169

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

H atoms were positioned geometrically and allowed to ride on their parent atoms at distances of $\text{Csp}^2\text{---}H = 0.93 \text{ \AA}$ with $U_{\text{iso}}(H) = 1.2U_{\text{eq}}(C)$, $\text{Csp}^3\text{---}H = 0.97 \text{ \AA}$ with $U_{\text{iso}}(H) = 1.5U_{\text{eq}}(C)$, $\text{N---}H = 0.86 \text{ \AA}$ with $U_{\text{iso}}(H) = 1.2U_{\text{eq}}(N)$ and $\text{O---}H = 0.82 \text{ \AA}$ with $U_{\text{iso}}(H) = 1.2U_{\text{eq}}(O)$. The highest peak is located 0.94 \AA from atom Co1.

Data collection: SMART (Bruker, 2002); cell refinement: SAINT (Bruker, 2002); data reduction: SAINT; program(s) used to solve structure: SHELXS97 (Sheldrick, 1997); program(s) used to refine structure: SHELXL97 (Sheldrick, 1997); molecular graphics: SHELXTL (Bruker, 2002); software used to prepare material for publication: SHELXL97.

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