

## Propane-1,2-diammonium chromate(VI)

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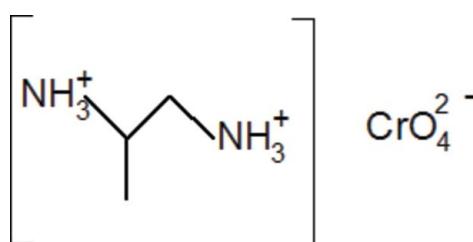
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Key indicators: single-crystal X-ray study;  $T = 150\text{ K}$ ; mean  $\sigma(\text{C}-\text{C}) = 0.003\text{ \AA}$ ;  $R$  factor = 0.026;  $wR$  factor = 0.071; data-to-parameter ratio = 15.1.

In the title molecular salt,  $(\text{C}_3\text{H}_{12}\text{N}_2)[\text{CrO}_4]$ , each chromate anion accepts six  $\text{N}-\text{H}\cdots\text{O}$  and  $\text{C}-\text{H}\cdots\text{O}$  hydrogen bonds from nearby propane-1,2-diammonium cations. Three of the four O atoms of the chromate anion accept these bonds; the remaining Cr–O bond length is notably shorter than the others. In the crystal, the anions and cations stack in layers lying parallel to (100): the hydrogen-bonding pattern leads to a three-dimensional network.

## Related literature

For background to organic chromates, see: Chebbi & Driss (2002, 2004); Srinivasan *et al.* (2003). For the crystal structures of simple salts of the propane-1,2-diammonium cation, see: Pospieszna-Markiewicz *et al.* (2011); Gerrard & Weller (2002); Lee & Harrison (2003); Todd & Harrison (2005). For a discussion on hydrogen bonding, see: Brown (1976); Blessing (1986).



## Experimental

## Crystal data

$(\text{C}_3\text{H}_{12}\text{N}_2)[\text{CrO}_4]$   
 $M_r = 192.15$   
Monoclinic,  $P2_1/c$   
 $a = 5.6462(2)\text{ \AA}$   
 $b = 15.8373(5)\text{ \AA}$   
 $c = 8.4442(3)\text{ \AA}$   
 $\beta = 106.779(1)\text{ }^\circ$   
 $V = 722.94(4)\text{ \AA}^3$   
 $Z = 4$   
Mo  $K\alpha$  radiation

$\mu = 1.54\text{ mm}^{-1}$   
 $T = 150\text{ K}$

$0.55 \times 0.44 \times 0.31\text{ mm}$

## Data collection

Bruker APEXII diffractometer  
Absorption correction: multi-scan  
(SADABS; Sheldrick, 2002)  
 $T_{\min} = 0.477$ ,  $T_{\max} = 0.620$

6302 measured reflections  
1659 independent reflections  
1554 reflections with  $I > 2\sigma(I)$   
 $R_{\text{int}} = 0.032$

## Refinement

$R[F^2 > 2\sigma(F^2)] = 0.026$   
 $wR(F^2) = 0.071$   
 $S = 1.14$   
1659 reflections  
110 parameters

H atoms treated by a mixture of independent and constrained refinement  
 $\Delta\rho_{\max} = 0.34\text{ e \AA}^{-3}$   
 $\Delta\rho_{\min} = -0.52\text{ e \AA}^{-3}$

**Table 1**  
Selected bond lengths ( $\text{\AA}$ ).

Cr–O3	1.6182 (13)	Cr–O1	1.6711 (13)
Cr–O2	1.6378 (13)	Cr–O4	1.6879 (13)

**Table 2**  
Hydrogen-bond geometry ( $\text{\AA}$ ,  $^\circ$ ).

$D-\text{H}\cdots A$	$D-\text{H}$	$\text{H}\cdots A$	$D\cdots A$	$D-\text{H}\cdots A$
N1–H1A $\cdots$ O4 <sup>i</sup>	0.85 (3)	1.94 (3)	2.769 (2)	166 (2)
N1–H1B $\cdots$ O4 <sup>ii</sup>	0.87 (3)	1.97 (3)	2.816 (2)	164 (2)
N1–H1C $\cdots$ O2 <sup>iii</sup>	0.85 (3)	1.97 (3)	2.818 (2)	171 (2)
N2–H2A $\cdots$ O1 <sup>iv</sup>	0.82 (3)	1.96 (3)	2.779 (2)	178 (2)
N2–H2B $\cdots$ O1 <sup>v</sup>	0.85 (2)	1.91 (3)	2.748 (2)	173 (2)
N2–H2C $\cdots$ O4	0.85 (3)	1.95 (3)	2.795 (2)	171 (2)
C1–H1 $\cdots$ O2 <sup>iv</sup>	0.99	2.34	3.313 (2)	167
C3–H3B $\cdots$ O2 <sup>iii</sup>	0.98	2.37	3.301 (2)	158

Symmetry codes: (i)  $x, -y + \frac{1}{2}, z + \frac{1}{2}$ ; (ii)  $-x + 1, y + \frac{1}{2}, -z + \frac{3}{2}$ ; (iii)  $-x + 2, y + \frac{1}{2}, -z + \frac{3}{2}$ ; (iv)  $x - 1, -y + \frac{1}{2}, z - \frac{1}{2}$ ; (v)  $x - 1, y, z$ .

Data collection: *APEX2* (Bruker, 2006); cell refinement: *SAINT* (Bruker, 2006); data reduction: *SAINT*; program(s) used to solve structure: *SIR97* (Altomare *et al.*, 1999); program(s) used to refine structure: *SHELXL97* (Sheldrick, 2008); molecular graphics: *ORTEP-3 for Windows* (Farrugia, 2012) and *DIAMOND* (Brandenburg & Putz, 2005); software used to prepare material for publication: *WinGX* (Farrugia, 2012) and *CRYSCAL* (T. Roisnel, local program).

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Supporting information for this paper is available from the IUCr electronic archives (Reference: HB7193).

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# supporting information

*Acta Cryst.* (2014). E70, m84–m85 [doi:10.1107/S1600536814002463]

## Propane-1,2-diammonium chromate(VI)

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### S1. Comment

In this work, we report the preparation and the structural investigation of a new organic chromate,  $C_3H_{12}N_2\cdot CrO_4$  (I).

The asymmetric unit of (I) consists of one chromate anion and one propane-1,2-ammonium dication (Figure 1). The structure of the compound consists of discrete chromate ions stacked in layers parallel to the (100) plane, separated by organic cations (Figure 2). The structural cohesion is established by a three-dimensional network of  $N-H\cdots O$  and  $C-H\cdots O$  hydrogen bonds. Geometrical characteristics of the chromate anion are slightly different (Table 1). The distance  $Cr-O3$  is notably the shortest (1.6182 (13) Å) because  $O3$  is not applied in any hydrogen bond (Table 2) at the same time as  $Cr-O4$  distance is the longest (1.6879 (13) Å) because  $O4$  is applied in three hydrogen bonds. These geometrical features have also been noticed in other crystal structures (Chebbi & Driss, 2002; 2004; Srinivasan, *et al.*, 2003).

The 1,2-propanediammonium cation is characterized by  $N-C-C-N$  and  $N-C-C-C$  torsion angles of 164.88 (14) and -74.50 (19)°, respectively. Each organic entity is bounded to six different chromate anions through eight  $N-H\cdots O$  and  $C-H\cdots O$  hydrogen bonds forming a three dimensional network. Examination of the 1,2-propanediammonium cation shows that the bond distances and angles show no significant difference from those obtained in other simple salts involving the same organic groups (Pospieszna-Markiewicz, *et al.*, 2011; Gerrard, *et al.*, 2002; Lee, *et al.*, 2003; Todd, *et al.*, 2005).

The established weak H-bonds (Brown, 1976; Blessing, 1986) of types  $N-H\cdots O$  and  $C-H\cdots O$  involve oxygen atoms of the chromate anions as acceptors, and the protonated nitrogen atoms and carbon atoms of 1,2-diammoniumpropane as donors.

### S2. Experimental

$CrO_3$  (0.10 g, 1 mmol) and 1,2-diaminopropane (0.13 ml, 1 mmol) were dissolved in distilled water (20 ml). The resulting solution was stirred for 30 min. and then evaporated slowly at room temperature. Yellow prisms of the title compound were obtained from the solution after one week.

### S3. Refinement

The hydrogen atoms bonded to N1 and N2 were located from a difference map and were allowed to refine. The rest of the H atoms were treated as riding, with  $C-H = 0.99$  Å (methylene) or 0.98 Å (methyl) or 1.00 Å (methine), with  $U_{iso}(H) = 1.2U_{eq}(\text{parent C atoms})$  and  $1.5U_{eq}(\text{parent N or C-methyl atoms})$ .

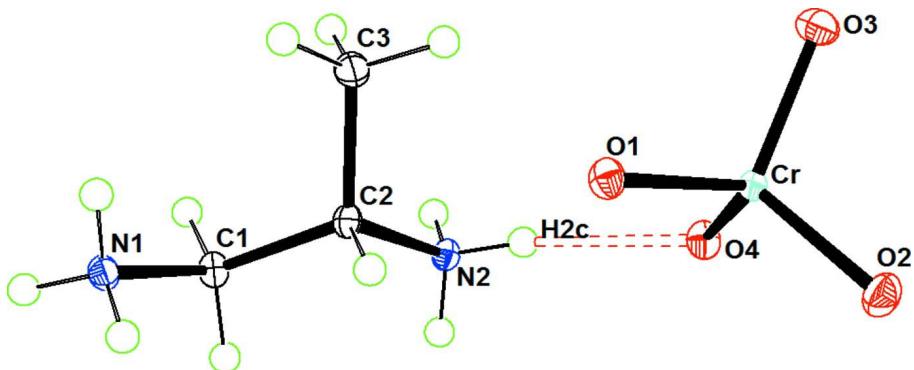
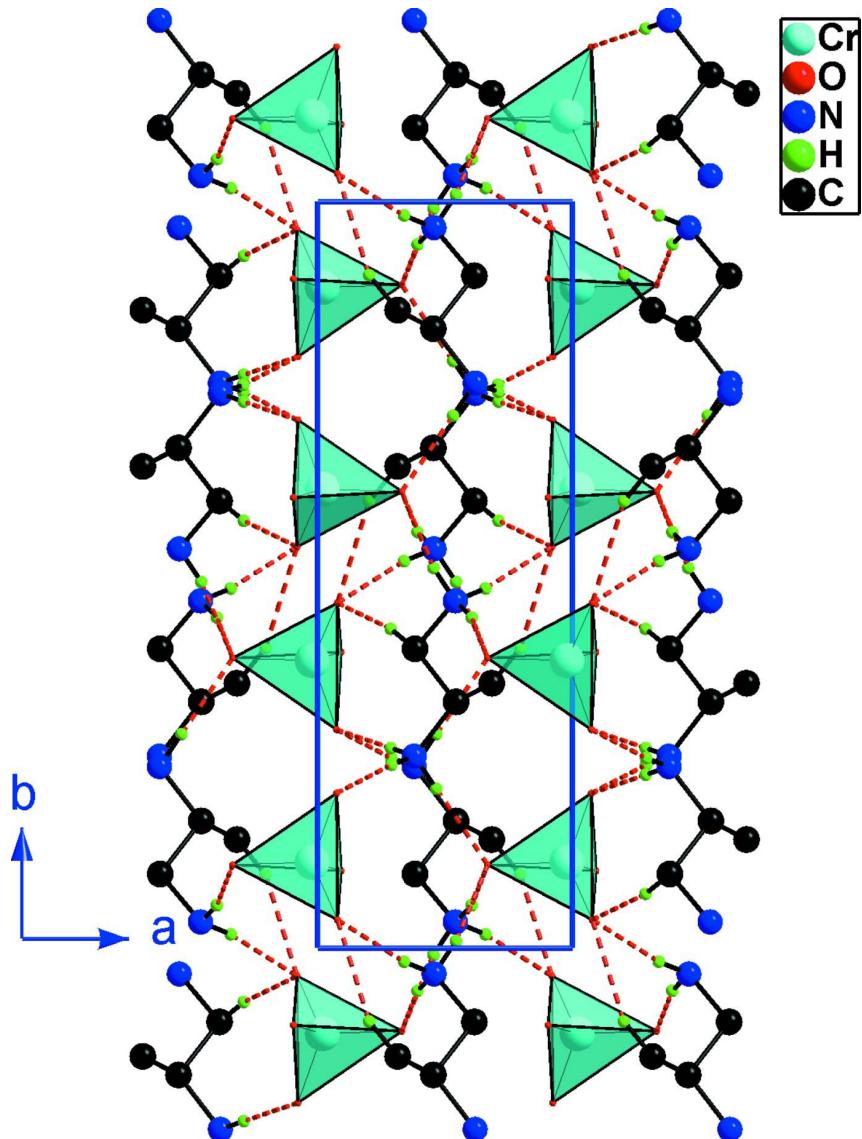


Figure 1

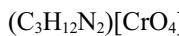
An *ORTEP* view of (I) with displacement ellipsoids drawn at the 30% probability level. H atoms are represented as small spheres of arbitrary radii. Hydrogen bonds are shown as dotted lines.

**Figure 2**

Projection of (I) along the *c* axis. The H-atoms not involved in H-bonding are omitted.

### Propane-1,2-diammonium chromate(VI)

#### *Crystal data*



*M<sub>r</sub>* = 192.15

Monoclinic, *P*2<sub>1</sub>/*c*

Hall symbol: -P 2ybc

*a* = 5.6462 (2) Å

*b* = 15.8373 (5) Å

*c* = 8.4442 (3) Å

β = 106.779 (1)°

*V* = 722.94 (4) Å<sup>3</sup>

*Z* = 4

*F*(000) = 400

*D<sub>x</sub>* = 1.765 Mg m<sup>-3</sup>

Mo *K*α radiation, λ = 0.71073 Å

Cell parameters from 3695 reflections

θ = 2.8–27.5°

μ = 1.54 mm<sup>-1</sup>

*T* = 150 K

Prism, yellow

0.55 × 0.44 × 0.31 mm

*Data collection*

Bruker APEXII  
diffractometer  
Radiation source: fine-focus sealed tube  
Graphite monochromator  
CCD rotation images, thin slices scans  
Absorption correction: multi-scan  
(*SADABS*; Sheldrick, 2002)  
 $T_{\min} = 0.477$ ,  $T_{\max} = 0.620$

6302 measured reflections  
1659 independent reflections  
1554 reflections with  $I > 2\sigma(I)$   
 $R_{\text{int}} = 0.032$   
 $\theta_{\max} = 27.5^\circ$ ,  $\theta_{\min} = 3.6^\circ$   
 $h = -7 \rightarrow 6$   
 $k = -19 \rightarrow 20$   
 $l = -9 \rightarrow 10$

*Refinement*

Refinement on  $F^2$   
Least-squares matrix: full  
 $R[F^2 > 2\sigma(F^2)] = 0.026$   
 $wR(F^2) = 0.071$   
 $S = 1.14$   
1659 reflections  
110 parameters  
0 restraints  
Primary atom site location: structure-invariant  
direct methods

Secondary atom site location: difference Fourier  
map  
Hydrogen site location: inferred from  
neighbouring sites  
H atoms treated by a mixture of independent  
and constrained refinement  
 $w = 1/[\sigma^2(F_o^2) + (0.0303P)^2 + 0.5214P]$   
where  $P = (F_o^2 + 2F_c^2)/3$   
 $(\Delta/\sigma)_{\max} = 0.001$   
 $\Delta\rho_{\max} = 0.34 \text{ e } \text{\AA}^{-3}$   
 $\Delta\rho_{\min} = -0.52 \text{ e } \text{\AA}^{-3}$

*Special details*

**Geometry.** All e.s.d.'s (except the e.s.d. in the dihedral angle between two l.s. planes) are estimated using the full covariance matrix. The cell e.s.d.'s are taken into account individually in the estimation of e.s.d.'s in distances, angles and torsion angles; correlations between e.s.d.'s in cell parameters are only used when they are defined by crystal symmetry. An approximate (isotropic) treatment of cell e.s.d.'s is used for estimating e.s.d.'s involving l.s. planes.

**Refinement.** Refinement of  $F^2$  against ALL reflections. The weighted  $R$ -factor  $wR$  and goodness of fit  $S$  are based on  $F^2$ , conventional  $R$ -factors  $R$  are based on  $F$ , with  $F$  set to zero for negative  $F^2$ . The threshold expression of  $F^2 > \sigma(F^2)$  is used only for calculating  $R$ -factors(gt) etc. and is not relevant to the choice of reflections for refinement.  $R$ -factors based on  $F^2$  are statistically about twice as large as those based on  $F$ , and  $R$ -factors based on ALL data will be even larger.

*Fractional atomic coordinates and isotropic or equivalent isotropic displacement parameters ( $\text{\AA}^2$ )*

	<i>x</i>	<i>y</i>	<i>z</i>	$U_{\text{iso}}^*/U_{\text{eq}}$
Cr	0.97996 (5)	0.114804 (17)	0.67966 (3)	0.00751 (11)
O1	1.0758 (2)	0.20744 (8)	0.77039 (15)	0.0120 (3)
O2	1.0791 (3)	0.03771 (9)	0.81142 (16)	0.0154 (3)
O3	1.0937 (3)	0.10356 (8)	0.52561 (16)	0.0146 (3)
O4	0.6684 (2)	0.11150 (8)	0.60744 (16)	0.0124 (3)
N1	0.5418 (3)	0.46519 (10)	0.7997 (2)	0.0112 (3)
H1A	0.606 (4)	0.4428 (16)	0.894 (3)	0.017*
H1B	0.456 (4)	0.5094 (16)	0.810 (3)	0.017*
H1C	0.660 (5)	0.4816 (15)	0.763 (3)	0.017*
N2	0.3804 (3)	0.25695 (10)	0.5858 (2)	0.0102 (3)
H2A	0.289 (5)	0.2680 (15)	0.494 (3)	0.015*
H2B	0.291 (4)	0.2454 (15)	0.648 (3)	0.015*
H2C	0.467 (4)	0.2134 (16)	0.581 (3)	0.015*
C1	0.3836 (3)	0.40086 (11)	0.6906 (2)	0.0107 (3)
H1	0.2885	0.4275	0.5855	0.013*

H2	0.2647	0.3774	0.7451	0.013*
C2	0.5446 (3)	0.32984 (11)	0.6549 (2)	0.0097 (3)
H3	0.6664	0.3121	0.7612	0.012*
C3	0.6845 (4)	0.35472 (12)	0.5330 (2)	0.0141 (4)
H3A	0.7805	0.3063	0.5133	0.021*
H3B	0.7967	0.4016	0.5786	0.021*
H3C	0.5667	0.3722	0.4284	0.021*

*Atomic displacement parameters ( $\text{\AA}^2$ )*

	$U^{11}$	$U^{22}$	$U^{33}$	$U^{12}$	$U^{13}$	$U^{23}$
Cr	0.00627 (16)	0.00701 (16)	0.00877 (16)	-0.00041 (10)	0.00140 (11)	-0.00009 (9)
O1	0.0121 (6)	0.0105 (6)	0.0129 (6)	-0.0018 (5)	0.0027 (5)	-0.0024 (5)
O2	0.0167 (7)	0.0130 (6)	0.0152 (6)	0.0031 (5)	0.0028 (5)	0.0037 (5)
O3	0.0151 (7)	0.0158 (6)	0.0145 (6)	-0.0032 (5)	0.0069 (5)	-0.0032 (5)
O4	0.0085 (6)	0.0122 (6)	0.0155 (6)	-0.0004 (5)	0.0021 (5)	-0.0011 (5)
N1	0.0127 (8)	0.0099 (7)	0.0110 (7)	0.0005 (6)	0.0035 (6)	-0.0014 (6)
N2	0.0105 (7)	0.0084 (7)	0.0116 (7)	-0.0006 (6)	0.0029 (6)	-0.0004 (6)
C1	0.0086 (8)	0.0102 (8)	0.0123 (8)	0.0002 (7)	0.0016 (7)	-0.0014 (6)
C2	0.0085 (8)	0.0085 (8)	0.0112 (8)	-0.0006 (6)	0.0014 (6)	-0.0011 (6)
C3	0.0135 (9)	0.0125 (8)	0.0185 (9)	-0.0021 (7)	0.0080 (7)	-0.0019 (7)

*Geometric parameters ( $\text{\AA}$ ,  $\text{^\circ}$ )*

Cr—O3	1.6182 (13)	N2—H2B	0.85 (2)
Cr—O2	1.6378 (13)	N2—H2C	0.85 (3)
Cr—O1	1.6711 (13)	C1—C2	1.530 (2)
Cr—O4	1.6879 (13)	C1—H1	0.9900
N1—C1	1.486 (2)	C1—H2	0.9900
N1—H1A	0.85 (3)	C2—C3	1.520 (2)
N1—H1B	0.87 (3)	C2—H3	1.0000
N1—H1C	0.85 (3)	C3—H3A	0.9800
N2—C2	1.490 (2)	C3—H3B	0.9800
N2—H2A	0.82 (3)	C3—H3C	0.9800
O3—Cr—O2	109.08 (7)	N1—C1—C2	109.95 (14)
O3—Cr—O1	108.31 (6)	N1—C1—H1	109.7
O2—Cr—O1	109.94 (7)	C2—C1—H1	109.7
O3—Cr—O4	108.67 (7)	N1—C1—H2	109.7
O2—Cr—O4	109.78 (7)	C2—C1—H2	109.7
O1—Cr—O4	111.01 (6)	H1—C1—H2	108.2
C1—N1—H1A	108.1 (16)	N2—C2—C3	108.76 (14)
C1—N1—H1B	111.1 (16)	N2—C2—C1	108.00 (14)
H1A—N1—H1B	110 (2)	C3—C2—C1	113.43 (15)
C1—N1—H1C	112.0 (16)	N2—C2—H3	108.9
H1A—N1—H1C	107 (2)	C3—C2—H3	108.9
H1B—N1—H1C	108 (2)	C1—C2—H3	108.9
C2—N2—H2A	111.0 (16)	C2—C3—H3A	109.5

C2—N2—H2B	110.0 (16)	C2—C3—H3B	109.5
H2A—N2—H2B	108 (2)	H3A—C3—H3B	109.5
C2—N2—H2C	110.2 (16)	C2—C3—H3C	109.5
H2A—N2—H2C	110 (2)	H3A—C3—H3C	109.5
H2B—N2—H2C	108 (2)	H3B—C3—H3C	109.5
N1—C1—C2—N2	164.88 (14)	N1—C1—C2—C3	−74.50 (19)

*Hydrogen-bond geometry (Å, °)*

D—H···A	D—H	H···A	D···A	D—H···A
N1—H1A···O4 <sup>i</sup>	0.85 (3)	1.94 (3)	2.769 (2)	166 (2)
N1—H1B···O4 <sup>ii</sup>	0.87 (3)	1.97 (3)	2.816 (2)	164 (2)
N1—H1C···O2 <sup>iii</sup>	0.85 (3)	1.97 (3)	2.818 (2)	171 (2)
N2—H2A···O1 <sup>iv</sup>	0.82 (3)	1.96 (3)	2.779 (2)	178 (2)
N2—H2B···O1 <sup>v</sup>	0.85 (2)	1.91 (3)	2.748 (2)	173 (2)
N2—H2C···O4	0.85 (3)	1.95 (3)	2.795 (2)	171 (2)
C1—H1···O2 <sup>iv</sup>	0.99	2.34	3.313 (2)	167
C3—H3B···O2 <sup>iii</sup>	0.98	2.37	3.301 (2)	158

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