

3-Ammonio-4-hydroxybenzoate monohydrate

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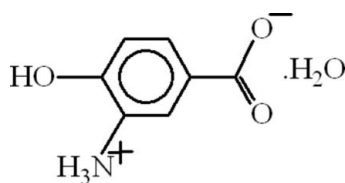
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Key indicators: single-crystal X-ray study; $T = 296$ K; mean $\sigma(\text{C}-\text{C}) = 0.002$ Å; R factor = 0.042; wR factor = 0.118; data-to-parameter ratio = 13.6.

The title compound, $\text{C}_7\text{H}_7\text{NO}_3 \cdot \text{H}_2\text{O}$, which crystallized as a hydrate, was obtained from an extraction of the plant species *Saussurea atkinsonii* of the asteraceae family collected from the hilly area (Ayubia) of Pakistan during the flowering season. The dihedral angle between the benzene ring and the carboxylate group is 25.64 (5)°. In the crystal, the packing is consolidated by $\text{N}-\text{H} \cdots \text{O}$ and $\text{O}-\text{H} \cdots \text{O}$ hydrogen bonds, as well as weak aromatic $\pi-\pi$ stacking [centroid-centroid separation = 3.9365 (9) Å] and $\text{C}=\text{O} \cdots \pi$ interactions.

Related literature

For a related structure, see: Bertasso *et al.* (2001). For reference structural data, see: Allen *et al.* (1987).



Experimental

Crystal data

$\text{C}_7\text{H}_7\text{NO}_3 \cdot \text{H}_2\text{O}$

$M_r = 171.15$

Orthorhombic, $Pbca$

$a = 8.7711$ (3) Å

$b = 12.7193$ (7) Å

$c = 12.9289$ (6) Å

$V = 1442.38$ (11) Å³

$Z = 8$

Mo $K\alpha$ radiation

$\mu = 0.13$ mm⁻¹

$T = 296$ K

$0.26 \times 0.20 \times 0.20$ mm

Data collection

Bruker Kappa APEXII CCD

diffractometer

Absorption correction: multi-scan

(*SADABS*; Bruker, 2005)

$T_{\min} = 0.971$, $T_{\max} = 0.976$

8827 measured reflections

1725 independent reflections

1277 reflections with $I > 2\sigma(I)$

$R_{\text{int}} = 0.033$

Refinement

$R[F^2 > 2\sigma(F^2)] = 0.042$

$wR(F^2) = 0.118$

$S = 1.06$

1725 reflections

127 parameters

H atoms treated by a mixture of independent and constrained refinement

$\Delta\rho_{\text{max}} = 0.31$ e Å⁻³

$\Delta\rho_{\text{min}} = -0.24$ e Å⁻³

Table 1

Hydrogen-bond geometry (Å, °).

$D-H \cdots A$	$D-H$	$H \cdots A$	$D \cdots A$	$D-H \cdots A$
$\text{N1}-\text{H1A} \cdots \text{O1}^{\text{i}}$	0.952 (18)	1.945 (18)	2.8884 (16)	170.4 (14)
$\text{N1}-\text{H1A} \cdots \text{O2}^{\text{i}}$	0.952 (18)	2.335 (18)	2.9008 (18)	117.6 (13)
$\text{N1}-\text{H1B} \cdots \text{O4}^{\text{ii}}$	0.944 (19)	2.001 (19)	2.8957 (19)	157.4 (16)
$\text{N1}-\text{H1C} \cdots \text{O1}^{\text{iii}}$	0.933 (17)	1.860 (17)	2.7846 (18)	170.5 (16)
$\text{O3}-\text{H3} \cdots \text{O4}^{\text{iv}}$	0.904 (18)	1.760 (18)	2.6456 (15)	166.0 (19)
$\text{O4}-\text{H41} \cdots \text{O2}^{\text{iv}}$	0.90 (2)	1.80 (2)	2.6945 (18)	171.1 (17)
$\text{O4}-\text{H42} \cdots \text{O1}^{\text{iii}}$	0.884 (19)	2.03 (2)	2.9027 (18)	168.8 (17)
$\text{C6}-\text{H6} \cdots \text{O3}^{\text{v}}$	0.93	2.55	3.446 (2)	161
$\text{C7}-\text{O2} \cdots \text{CgA}^{\text{vi}}$	1.25 (1)	3.49 (1)	3.9313 (16)	101 (1)

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

Data collection: *APEX2* (Bruker, 2007); cell refinement: *SAINT* (Bruker, 2007); data reduction: *SAINT*; program(s) used to solve structure: *SHELXS97* (Sheldrick, 2008); program(s) used to refine structure: *SHELXL97* (Sheldrick, 2008); molecular graphics: *ORTEP-3 for Windows* (Farrugia, 1997) and *PLATON* (Spek, 2009); software used to prepare material for publication: *WinGX* (Farrugia, 1999) and *PLATON*.

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Supplementary data and figures for this paper are available from the IUCr electronic archives (Reference: HB2969).

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supplementary materials

Acta Cryst. (2009). E65, o1304 [doi:10.1107/S1600536809017462]

3-Ammonio-4-hydroxybenzoate monohydrate

S. Ullah, M. N. Tahir, D. Shahwar, Z.-D. Khan and M. A. Khan

Comment

The medicinal plants are available all over the world. Locally available plant specie *Saussurea atkinsonii* of asteraceae family was collected from hilly area (Ayubia) of Pakistan during the flowering season. The plant was dried inside room for 20–25 days. The title compound (I), (Fig. 1), is an extract of it in chloroform and methanol. The study of its bio-activity is in progress.

The crystal structure of (II) (4-vinylphenyl 3-amino-4-hydroxybenzoate or bagremycin A (Bertasso *et al.*, 2001), has been reported which contains the aromatic ring along with heavy atoms of the substituents of (I). In the title compound, the bond distances and bond angles are within normal ranges (Allen *et al.*, 1987). The benzene ring A (C1–C6) is planar and is oriented at a dihedral angle of 25.64 (5)° with the CO₂ group. The N-atom of ammonium is in plane of the ring A, whereas the O-atom of hydroxy group is at a distance of -0.0580 (21) Å from the same.

There exist intensive intermolecular H-bonding (Table 1), resulting in three-dimensional polymeric network. There also exist CgA...CgAⁱ [symmetry code $i = 1 - x, 1 - y, -z$] interaction at a distance of 3.9365 (9) Å, where CgA is the centroid of aromatic ring. The molecules may also be stabilized due to C=O... π interaction (Table 1).

Experimental

The Specie *Saussurea atkinsonii* of asteraceae family was dried inside room for 20–25 days as a whole and grinded. The extract was obtained using soxhlet apparatus in 50% chloroform and 50% methanol and it was subjected to isolation by performing column chromatography and thin layer chromatography. The extract obtained was recrystallized from methanol and light brown rods of (I) were obtained. The water found in the structure was presumably incorporated from the atomsphere.

Refinement

The coordinates of H-atoms of hydroxy, ammonium moiety and water molecule were refined. The other H-atoms were positioned geometrically, with C-H = 0.93 Å for aromatic type and constrained to ride on their parent atoms, with $U_{\text{iso}}(\text{H}) = 1.2U_{\text{eq}}(\text{C, N, O})$.

Figures

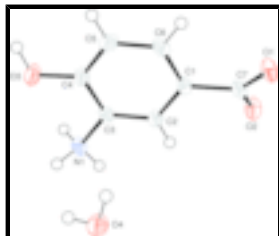


Fig. 1. View of (I) with displacement ellipsoids drawn at the 50% probability level. H-atoms are shown by small circles of arbitrary radius.

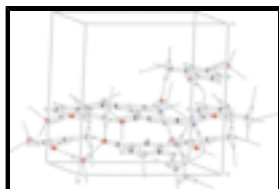


Fig. 2. The partial packing of (I), showing that molecules form three-dimensional polymeric network.

3-Ammonio-4-hydroxybenzoate monohydrate

Crystal data

$C_7H_7NO_3 \cdot H_2O$

$M_r = 171.15$

Orthorhombic, *Pbca*

Hall symbol: -P 2ac 2ab

$a = 8.7711$ (3) Å

$b = 12.7193$ (7) Å

$c = 12.9289$ (6) Å

$V = 1442.38$ (11) Å³

$Z = 8$

$F_{000} = 720$

$D_x = 1.576$ Mg m⁻³

Mo $K\alpha$ radiation

$\lambda = 0.71073$ Å

Cell parameters from 1725 reflections

$\theta = 3.2$ – 28.3°

$\mu = 0.13$ mm⁻¹

$T = 296$ K

Rod, light brown

$0.26 \times 0.20 \times 0.20$ mm

Data collection

Bruker Kappa APEXII CCD diffractometer

Radiation source: fine-focus sealed tube

Monochromator: graphite

Detector resolution: 7.40 pixels mm⁻¹

$T = 296$ K

ω scans

Absorption correction: multi-scan (SADABS; Bruker, 2005)

$T_{\min} = 0.971$, $T_{\max} = 0.976$

8827 measured reflections

1752 independent reflections

1277 reflections with $I > 2\sigma(I)$

$R_{\text{int}} = 0.033$

$\theta_{\text{max}} = 28.3^\circ$

$\theta_{\text{min}} = 3.2^\circ$

$h = -11 \rightarrow 11$

$k = -15 \rightarrow 16$

$l = -11 \rightarrow 17$

Refinement

Refinement on F^2

Secondary atom site location: difference Fourier map

Least-squares matrix: full

$$R[F^2 > 2\sigma(F^2)] = 0.042$$

$$wR(F^2) = 0.118$$

$$S = 1.06$$

1725 reflections

127 parameters

Primary atom site location: structure-invariant direct methods

Hydrogen site location: inferred from neighbouring sites

H atoms treated by a mixture of independent and constrained refinement

$$w = 1/[\sigma^2(F_o^2) + (0.0684P)^2 + 0.0409P]$$

$$\text{where } P = (F_o^2 + 2F_c^2)/3$$

$$(\Delta/\sigma)_{\max} < 0.001$$

$$\Delta\rho_{\max} = 0.31 \text{ e } \text{\AA}^{-3}$$

$$\Delta\rho_{\min} = -0.24 \text{ e } \text{\AA}^{-3}$$

Special details

Geometry. Bond distances, angles *etc.* have been calculated using the rounded fractional coordinates. All su's are estimated from the variances of the (full) variance-covariance matrix. The cell e.s.d.'s are taken into account in the estimation of distances, angles and torsion angles

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 (\AA^2)

	x	y	z	$U_{\text{iso}}^*/U_{\text{eq}}$
O1	0.69598 (12)	0.67023 (9)	0.16442 (9)	0.0284 (4)
O2	0.83764 (11)	0.53206 (10)	0.12269 (9)	0.0302 (4)
O3	0.20823 (11)	0.29862 (10)	0.11448 (10)	0.0319 (4)
N1	0.48231 (14)	0.21872 (11)	0.16388 (11)	0.0232 (4)
C1	0.57236 (15)	0.50443 (12)	0.14203 (11)	0.0205 (5)
C2	0.58869 (15)	0.39732 (13)	0.15731 (11)	0.0203 (4)
C3	0.46511 (14)	0.33137 (12)	0.14912 (11)	0.0193 (4)
C4	0.32078 (15)	0.37051 (12)	0.12444 (12)	0.0212 (4)
C5	0.30300 (15)	0.47766 (13)	0.11230 (12)	0.0241 (5)
C6	0.42742 (15)	0.54444 (13)	0.12153 (12)	0.0232 (4)
C7	0.71208 (15)	0.57284 (13)	0.14299 (11)	0.0216 (5)
O4	0.43081 (12)	0.34489 (11)	0.45169 (10)	0.0308 (4)
H1A	0.587 (2)	0.2008 (14)	0.1721 (12)	0.0278*
H1B	0.4519 (19)	0.1828 (14)	0.1034 (15)	0.0278*
H1C	0.4229 (18)	0.1948 (14)	0.2189 (14)	0.0278*
H2	0.68403	0.36974	0.17325	0.0243*
H3	0.119 (2)	0.3252 (16)	0.0913 (15)	0.0383*
H5	0.20714	0.50524	0.09785	0.0289*
H6	0.41402	0.61656	0.11398	0.0279*
H41	0.3898 (19)	0.4057 (16)	0.4289 (15)	0.0369*
H42	0.381 (2)	0.2927 (16)	0.4217 (15)	0.0369*

supplementary materials

Atomic displacement parameters (\AA^2)

	U^{11}	U^{22}	U^{33}	U^{12}	U^{13}	U^{23}
O1	0.0316 (5)	0.0154 (7)	0.0383 (7)	-0.0051 (4)	0.0053 (4)	-0.0044 (5)
O2	0.0231 (5)	0.0245 (7)	0.0430 (7)	-0.0032 (5)	0.0043 (4)	-0.0063 (5)
O3	0.0193 (5)	0.0217 (7)	0.0548 (8)	-0.0032 (4)	-0.0059 (5)	0.0000 (6)
N1	0.0214 (6)	0.0150 (8)	0.0331 (8)	0.0010 (5)	0.0027 (5)	0.0015 (6)
C1	0.0236 (7)	0.0164 (9)	0.0216 (8)	-0.0029 (5)	0.0013 (5)	-0.0026 (6)
C2	0.0183 (6)	0.0190 (9)	0.0235 (8)	0.0006 (5)	0.0011 (5)	-0.0006 (6)
C3	0.0209 (6)	0.0142 (9)	0.0228 (8)	0.0003 (5)	0.0018 (5)	0.0013 (6)
C4	0.0195 (6)	0.0188 (9)	0.0254 (8)	-0.0024 (5)	-0.0004 (5)	-0.0019 (6)
C5	0.0205 (6)	0.0218 (9)	0.0300 (9)	0.0041 (6)	-0.0024 (6)	0.0003 (7)
C6	0.0290 (7)	0.0137 (8)	0.0270 (8)	0.0014 (6)	0.0000 (5)	0.0005 (7)
C7	0.0265 (7)	0.0169 (9)	0.0214 (8)	-0.0044 (6)	0.0013 (5)	-0.0002 (6)
O4	0.0257 (5)	0.0255 (8)	0.0412 (7)	0.0017 (5)	-0.0044 (4)	0.0023 (6)

Geometric parameters (\AA , $^\circ$)

O1—C7	1.277 (2)	C1—C7	1.503 (2)
O2—C7	1.2453 (17)	C1—C6	1.3948 (19)
O3—C4	1.3518 (18)	C1—C2	1.384 (2)
O3—H3	0.904 (18)	C2—C3	1.375 (2)
O4—H41	0.90 (2)	C3—C4	1.3972 (19)
O4—H42	0.884 (19)	C4—C5	1.381 (2)
N1—C3	1.453 (2)	C5—C6	1.388 (2)
N1—H1C	0.933 (17)	C2—H2	0.9300
N1—H1A	0.952 (18)	C5—H5	0.9300
N1—H1B	0.944 (19)	C6—H6	0.9300
C4—O3—H3	114.3 (13)	O3—C4—C5	125.06 (12)
H41—O4—H42	107.7 (17)	C3—C4—C5	118.69 (13)
C3—N1—H1A	110.5 (11)	O3—C4—C3	116.26 (13)
C3—N1—H1B	109.8 (11)	C4—C5—C6	120.36 (13)
H1A—N1—H1B	104.4 (14)	C1—C6—C5	120.64 (15)
H1A—N1—H1C	112.1 (14)	O1—C7—O2	123.20 (14)
C3—N1—H1C	111.3 (11)	O2—C7—C1	118.57 (14)
H1B—N1—H1C	108.4 (15)	O1—C7—C1	118.24 (12)
C2—C1—C6	118.72 (13)	C3—C2—H2	120.00
C2—C1—C7	118.97 (12)	C1—C2—H2	120.00
C6—C1—C7	122.24 (14)	C4—C5—H5	120.00
C1—C2—C3	120.53 (13)	C6—C5—H5	120.00
N1—C3—C2	120.64 (12)	C1—C6—H6	120.00
C2—C3—C4	120.97 (14)	C5—C6—H6	120.00
N1—C3—C4	118.38 (12)		
C6—C1—C2—C3	2.1 (2)	C1—C2—C3—C4	0.5 (2)
C7—C1—C2—C3	-175.02 (13)	N1—C3—C4—O3	-1.1 (2)
C2—C1—C6—C5	-2.7 (2)	N1—C3—C4—C5	178.75 (14)
C7—C1—C6—C5	174.30 (14)	C2—C3—C4—O3	177.60 (14)

C2—C1—C7—O1	-156.35 (14)	C2—C3—C4—C5	-2.5 (2)
C2—C1—C7—O2	23.8 (2)	O3—C4—C5—C6	-178.25 (15)
C6—C1—C7—O1	26.7 (2)	C3—C4—C5—C6	1.9 (2)
C6—C1—C7—O2	-153.17 (15)	C4—C5—C6—C1	0.7 (2)
C1—C2—C3—N1	179.24 (13)		

Hydrogen-bond geometry (Å, °)

<i>D</i> —H... <i>A</i>	<i>D</i> —H	H... <i>A</i>	<i>D</i> ... <i>A</i>	<i>D</i> —H... <i>A</i>
N1—H1A...O1 ⁱ	0.952 (18)	1.945 (18)	2.8884 (16)	170.4 (14)
N1—H1A...O2 ⁱ	0.952 (18)	2.335 (18)	2.9008 (18)	117.6 (13)
N1—H1B...O4 ⁱⁱ	0.944 (19)	2.001 (19)	2.8957 (19)	157.4 (16)
N1—H1C...O1 ⁱⁱⁱ	0.933 (17)	1.860 (17)	2.7846 (18)	170.5 (16)
O3—H3...O4 ^{iv}	0.904 (18)	1.760 (18)	2.6456 (15)	166.0 (19)
O4—H41...O2 ^{iv}	0.90 (2)	1.80 (2)	2.6945 (18)	171.1 (17)
O4—H42...O1 ⁱⁱⁱ	0.884 (19)	2.03 (2)	2.9027 (18)	168.8 (17)
C6—H6...O3 ^v	0.93	2.55	3.446 (2)	161
C7—O2...CgA ^{vi}	1.2453 (17)	3.4940 (13)	3.9313 (16)	101.24 (9)

Symmetry codes: (i) $-x+3/2, y-1/2, z$; (ii) $x, -y+1/2, z-1/2$; (iii) $-x+1, y-1/2, -z+1/2$; (iv) $x-1/2, y, -z+1/2$; (v) $-x+1/2, y+1/2, z$; (vi) $x+1/2, y, -z+1/2$.

Fig. 1

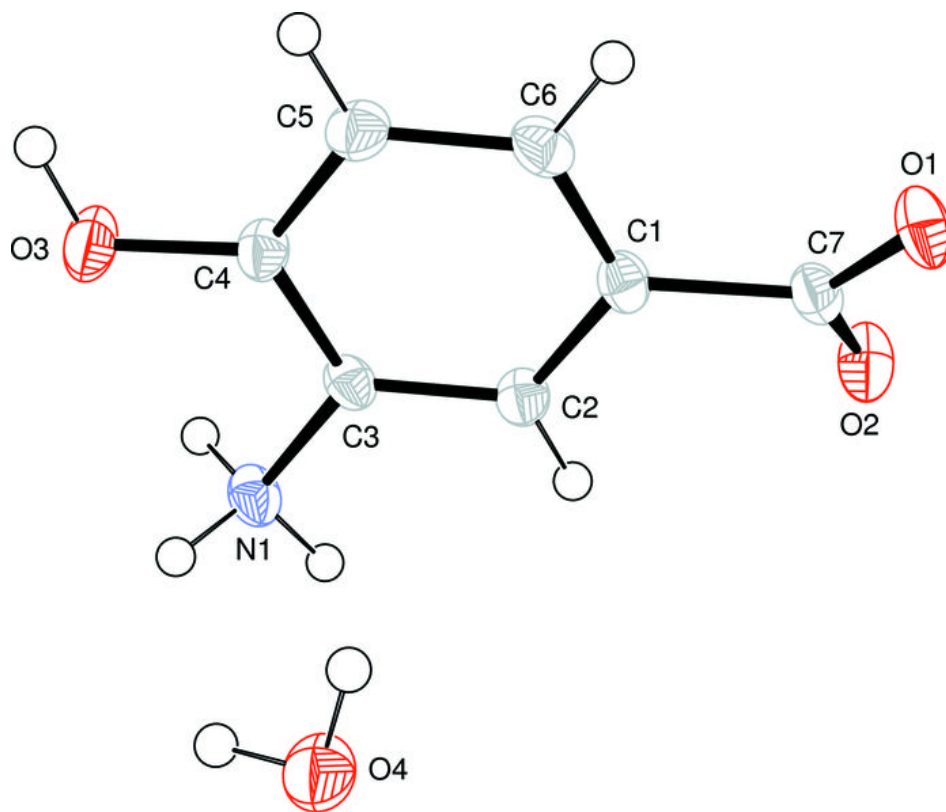


Fig. 2

