

## Gallic acid monohydrate

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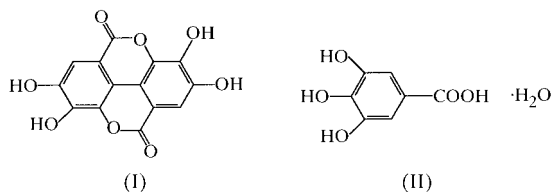
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In the crystal structure of the title compound, 3,4,5-trihydroxybenzoic acid monohydrate,  $C_7H_6O_5 \cdot H_2O$ , the gallic acid molecule has an intramolecular hydrogen bond involving a pair of hydroxyl groups, and it is also linked to a water molecule by a three-centre (bifurcated)  $OW-H \cdots O$  hydrogen bond. The packing of the molecules is stabilized by intermolecular  $O-H \cdots O$  and  $C-H \cdots O$  hydrogen bonds.

### Comment

Tannins are widely distributed in many edible and medicinal plants, and some with strong pharmacological activities have been used as haemostatic and astringent agents. There are two kinds of tannins in nature: one is condensation tannin and the other is hydrolysable tannin. The basic units of hydrolysable tannins are ellagic acid, (I), and gallic acid. The crystal structure of the former was published over three decades ago (Mathieson & Poppleton, 1968). In the present paper, we report the crystal structure of gallic acid monohydrate, (II), which has been shown to exhibit antibacterial and antiviral activities (Jiang & Xiao, 1986).



The molecule of (II) is essentially planar. The mean deviation of the benzene ring is 0.0028 Å and its dihedral angle with the plane of the carboxyl group is 2.9°. The bond distances are all normal.

Within the asymmetric unit, there is an intramolecular hydrogen bond between O1 and O2. The water molecule is linked to the gallic acid by a three-centre (bifurcated) donor hydrogen bond to two acceptor hydroxyl groups, as represented by  $OW-HOWB \cdots O2$  and  $OW-HOWB \cdots O3$  (Fig. 1). These hydrogen bonds form two five-membered hydrogen-bonded rings. The intermolecular hydrogen bonds  $O3-H \cdots OW$ ,  $O2-H \cdots OW$  and  $OW-HOWA \cdots O2$  link the

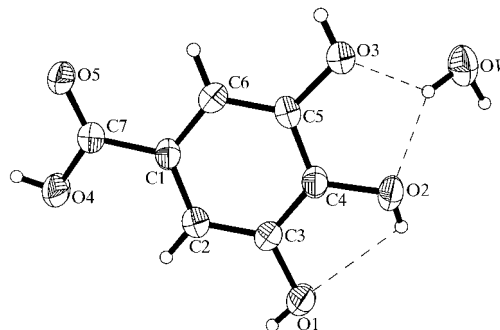


Figure 1

The molecular structure of (II) showing 50% probability displacement ellipsoids and the atom-numbering scheme; the hydrogen bonds in the asymmetric unit are indicated by dashed lines.

asymmetric unit at the junction where the water molecules are spirally distributed, to form a channel running parallel to the  $a$  axis. The intermolecular hydrogen bond  $O4-H \cdots O5$  links the two adjacent channels and results in extended 'way' sheets parallel to the (200) plane. Adjacent sheets are linked by hydrogen bond  $O1-H \cdots O5$  and a weak  $C-H \cdots O$  hydrogen bond between C6 and O1 to form a supramolecular assembly. Details of the hydrogen bonds are listed in Table 2. The crystal packing is shown in Fig. 2 in a projection along the short  $b$  axis.

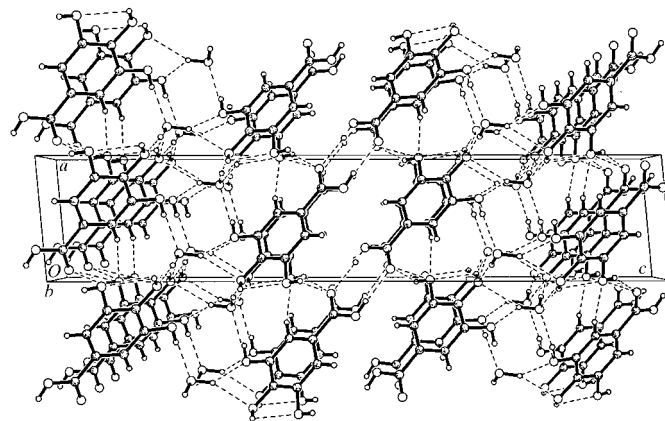


Figure 2

Packing diagram for (II) viewed down the  $b$  axis.

### Experimental

A sample of compound (II) was extracted from the whole plant of *Geum japonicum*. The dried plant material (3 kg) was chopped into small pieces and extracted with 95% ethanol at room temperature. The extracted liquid was evaporated *in vacuo* to yield an ethanol extract (80 g), which was then suspended in distilled water and successively extracted with hexane, ethyl acetate and *n*-butanol. The *n*-butanol extract was subjected to Sephadex LH-20 column chromatography and eluted with EtOH-H<sub>2</sub>O (4:1). The eluted liquid was condensed and allowed to stand at room temperature and yellow prismatic crystals of (II) (50 mg) were obtained.

## Crystal data

C<sub>7</sub>H<sub>6</sub>O<sub>5</sub>·H<sub>2</sub>O  
*M<sub>r</sub>* = 188.13  
 Monoclinic, *P*<sub>2</sub><sub>1</sub>/*c*  
*a* = 5.794 (4) Å  
*b* = 4.719 (5) Å  
*c* = 28.688 (5) Å  
 $\beta$  = 95.08 (3)°  
*V* = 781.4 (3) Å<sup>3</sup>  
*Z* = 4

## Data collection

Rigaku AFC-7R diffractometer  
 $\omega/2\theta$  scans  
 Absorption correction:  $\psi$  scan  
 (Rigaku, 1998)  
*T*<sub>min</sub> = 0.945, *T*<sub>max</sub> = 1.000  
 1514 measured reflections  
 1369 independent reflections  
 907 reflections with *F* > 4 $\sigma$ (*F*<sup>2</sup>)

## Refinement

Refinement on *F*<sup>2</sup>  
*R*(*F*) = 0.042  
*wR*(*F*<sup>2</sup>) = 0.127  
*S* = 1.006  
 1369 reflections  
 122 parameters

*D<sub>x</sub>* = 1.599 Mg m<sup>-3</sup>  
 Mo *K* $\alpha$  radiation  
 Cell parameters from 25  
 reflections  
 $\theta$  = 7.5–12.5°  
 $\mu$  = 0.144 mm<sup>-1</sup>  
*T* = 293 K  
 Prism, yellow  
 0.5 × 0.2 × 0.1 mm

*R*<sub>int</sub> = 0.032  
 $\theta$ <sub>max</sub> = 25°  
*h* = -1 → 6  
*k* = -1 → 5  
*l* = -34 → 33  
 3 standard reflections  
 every 197 reflections  
 intensity decay: none

H atoms treated by a mixture of  
 independent and constrained  
 refinement  
 $w = 1/[\sigma^2(F_o^2) + (0.0746P)^2]$   
 where  $P = (F_o^2 + 2F_c^2)/3$   
 $(\Delta/\sigma)_{\max} = 0.016$   
 $\Delta\rho_{\max} = 0.17 \text{ e } \text{Å}^{-3}$   
 $\Delta\rho_{\min} = -0.25 \text{ e } \text{Å}^{-3}$

Table 1

Selected geometric parameters (Å, °).

O1—C3	1.371 (3)	C1—C6	1.390 (3)
O2—C4	1.375 (3)	C1—C7	1.474 (3)
O3—C5	1.371 (3)	C2—C3	1.385 (3)
O4—C7	1.315 (3)	C3—C4	1.380 (3)
O5—C7	1.229 (3)	C4—C5	1.383 (3)
C1—C2	1.390 (3)	C5—C6	1.381 (3)
C2—C1—C6	120.9 (2)	C3—C4—C5	120.2 (1)
C2—C1—C7	120.9 (2)	O3—C5—C6	123.7 (2)
C6—C1—C7	118.2 (2)	O3—C5—C4	115.9 (1)
C3—C2—C1	119.0 (2)	C6—C5—C4	120.3 (2)
O1—C3—C4	114.6 (2)	C5—C6—C1	119.2 (2)
O1—C3—C2	125.0 (2)	O5—C7—O4	121.1 (2)
C4—C3—C2	120.3 (2)	O5—C7—C1	123.9 (2)
O2—C4—C3	122.6 (2)	O4—C7—C1	115.0 (2)
O2—C4—C5	117.1 (1)		

Table 2

Hydrogen-bonding 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>
OW—HOWA...O2 <sup>i</sup>	1.01	1.90	2.906 (4)	174
OW—HOWB...O2	0.97	2.18	3.058 (4)	151
OW—HOWB...O3	0.97	2.19	2.944 (4)	134
O1—HO1...O5 <sup>ii</sup>	0.82	2.06	2.796 (4)	150
O2—HO2...O1	0.82	2.33	2.693 (4)	108
O2—HO2...OW <sup>i</sup>	0.82	1.94	2.708 (4)	157
O3—HO3...OW <sup>iii</sup>	0.82	1.92	2.699 (4)	160
O4—HO4...O5 <sup>iv</sup>	0.82	1.85	2.663 (4)	175
C6—H6...O1 <sup>v</sup>	0.93	2.44	3.350 (5)	166

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

H(—C) atoms were generated geometrically and treated with a riding model. H(—O) were located from difference maps and refined with isotropic displacement parameters.

Data collection: *Rigaku/AFC Diffractometer Control Software* (Rigaku, 1998); cell refinement: *Rigaku/AFC Diffractometer Control Software*; data reduction: *TEXSAN* (Molecular Structure Corporation, 1995); program(s) used to solve structure: *SHELXS97* (Sheldrick, 1990); program(s) used to refine structure: *SHELXL97* (Sheldrick, 1997); molecular graphics: *XP* (Siemens, 1998).

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Supplementary data for this paper are available from the IUCr electronic archives (Reference: DE1123). Services for accessing these data are described at the back of the journal.

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