

Ethyl 4-(3-hydroxy-6-methyl-4-oxo-4*H*-pyran-2-ylmethyl)piperazine-1-carboxylateNazan Ocak,<sup>a\*</sup> Şamil Işık<sup>a</sup> and Mutlu Dilsiz Aytemir<sup>b</sup><sup>a</sup>Department of Physics, Faculty of Arts and Sciences, Ondokuz Mayıs University, TR-55139, Kurupelit-Samsun, Turkey, and<sup>b</sup>Department of Pharmaceutical Chemistry, Faculty of Pharmacy, Hacettepe University, TR-06100, Sıhhiye, Ankara, Turkey

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## Key indicators

Single-crystal X-ray study

T = 293 K

Mean  $\sigma(\text{C}-\text{C}) = 0.003 \text{ \AA}$ 

R factor = 0.049

wR factor = 0.162

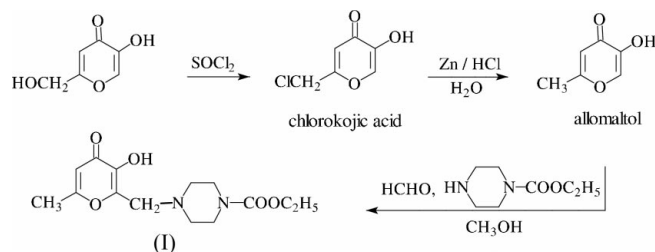
Data-to-parameter ratio = 15.1

For details of how these key indicators were automatically derived from the article, see <http://journals.iucr.org/e>.

The title compound,  $\text{C}_{14}\text{H}_{20}\text{N}_2\text{O}_5$ , was synthesized as a Mannich base and characterized by IR,  $^1\text{H}$  NMR, GC mass spectra and elemental analysis. The piperazine ring displays a chair conformation, and the crystal structure is stabilized by  $\text{O}-\text{H}\cdots\text{O}$  and  $\text{C}-\text{H}\cdots\text{O}$  intra- and intermolecular hydrogen bonds.

## Comment

The title compound, (I), is being studied for its possible biological properties due to the presence in it of the allomaltol group.



The title compound consists of 2-methyl-5-hydroxy-4*H*-pyran-4-one (allomaltol) and a piperazine ring, which is connected on one N side to the methylene bridge at the 2-position of the pyranone ring and on the other N side to the carboxylic acid ethyl ester group.

The bond lengths and angles observed in the allomaltol group are comparable to those found in maltol (3-hydroxy-2-methyl-4*H*-pyran-4-one; Burgess *et al.*, 1996).

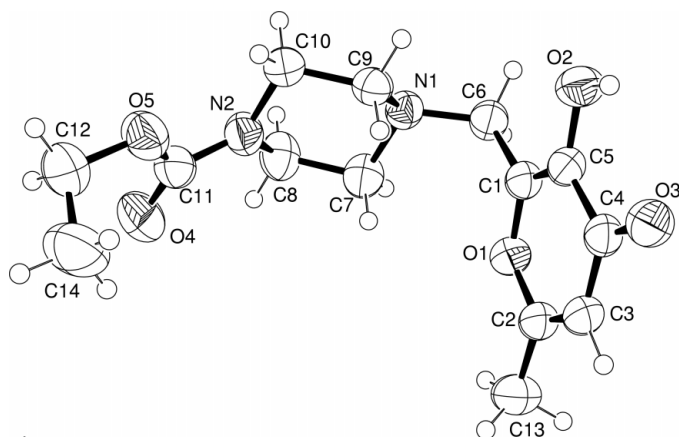
In the piperazine ring, the bond lengths and angles conform to those found previously (Yogavel *et al.*, 2002; Thirumurugan *et al.*, 1998; Koysal *et al.*, 2003). The piperazine ring adopts a chair conformation, with a total puckering amplitude of  $Q_T = 0.566(2) \text{ \AA}$  (Cremer & Pople, 1975). The sums of the bond angles around atoms N1 and N2 are  $337.1$  and  $359.8^\circ$ , respectively, because atom N1 is  $0.411(1) \text{ \AA}$  out of the plane through atoms C6, C7 and C9, and atom N2 is  $0.040(2) \text{ \AA}$  out of the plane through atoms C8, C10 and C11, indicative that atom N1 is  $sp^3$  while atom N2 is  $sp^2\pi$ -conjugated with the carboxy group. This is also shown by the values of the N1—C6 and N2—C11 bond distances. The plane through the C atoms of the piperazine ring makes a dihedral angle of  $77.43(4)^\circ$  with the allomaltol group.

There are one intermolecular ( $\text{O}-\text{H}\cdots\text{O}$ ) and five intramolecular ( $\text{O}-\text{H}\cdots\text{O}$  and  $\text{C}-\text{H}\cdots\text{O}$ ) hydrogen bonds. Atom O2 is involved as a donor in an inter- and intramolecular bifurcated hydrogen bond. The  $\text{C}-\text{H}\cdots\text{O}$  intramolecular interactions, shown in Fig. 2, help to stabilize the structure.

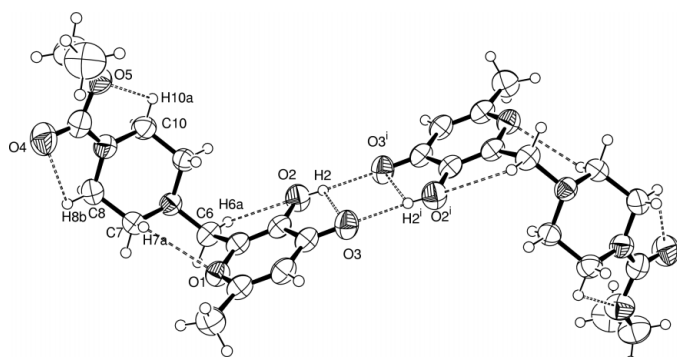
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**Figure 1**  
A view of (I), with the atom-numbering scheme. Displacement ellipsoids of non-H atoms are drawn at the 50% probability level.



**Figure 2**  
The hydrogen-bond network in (I).

## Experimental

All chemicals used in this study were supplied by Merck (Darmstadt, Germany) or Aldrich Chemical Co. (Steinheim, Germany). Compound (I) was prepared by the reaction of ethyl 1-piperazine-carboxylate (0.01 mol) and allomaltol (0.01 mol) in methanol (20 ml) with 37% formalin (1 ml). The mixture was stirred vigorously for 25 min. The resulting precipitate was collected by filtration and washed with cold methanol. Recrystallization from chloroform/petroleum ether (313–333 K) gave a white crystalline solid in 30% yield (m.p. 431–432 K). IR ( $\text{cm}^{-1}$ ): 1700 (C=O, s), 1613 (C=O, s, pyranone), 1459 (C=C, s) and 1221 (C–O, s);  $^1\text{H}$  NMR ( $\text{CDCl}_3$ , 80 MHz, p.p.m.): 1.20 (3H, t,  $-\text{CH}_3$ ), 2.30 (3H, s, 6- $\text{CH}_3$ ), 2.50 (4H, t,  $J = 4$  Hz, piperazine  $-\text{CH}_2-$ ), 3.40 (4H, t,  $J = 4$  Hz, piperazine  $-\text{CH}_2-$ ), 3.85 (2H, s,  $-\text{CH}_2-$ ), 4.10 (2H, q,  $-\text{CH}_2\text{CH}_3$ ), 6.20 (1H, s,  $\text{H}^5$ ); GC (MS) *m/e*: 116, 111, 85, 69, 56 (base peak). Analysis calculated for  $\text{C}_{14}\text{H}_{20}\text{N}_2\text{O}_5$ : C 56.74, H 6.80, N 9.45%; found: C 56.67, H 6.42, N 9.42%.

### Crystal data

$\text{C}_{14}\text{H}_{20}\text{N}_2\text{O}_5$   
 $M_r = 296.32$   
Monoclinic,  $C2/c$   
 $a = 24.069$  (3) Å  
 $b = 6.1796$  (4) Å  
 $c = 19.788$  (2) Å  
 $\beta = 95.513$  (8)°  
 $V = 2929.7$  (5) Å<sup>3</sup>  
 $Z = 8$

$D_x = 1.344$  Mg  $\text{m}^{-3}$   
Mo  $K\alpha$  radiation  
Cell parameters from 15 639 reflections  
 $\theta = 1.7$ – $29.6^\circ$   
 $\mu = 0.10$   $\text{mm}^{-1}$   
 $T = 293$  (2) K  
Prism, colourless  
 $0.60 \times 0.55 \times 0.32$  mm

### Data collection

Stoe IPDS-II diffractometer  
 $\omega$  scans  
Absorption correction: by integration (*X-RED32*; Stoe & Cie, 2002)  
 $T_{\min} = 0.937$ ,  $T_{\max} = 0.970$   
16 025 measured reflections

2888 independent reflections  
2388 reflections with  $I > 2\sigma(I)$   
 $R_{\text{int}} = 0.074$   
 $\theta_{\max} = 26.0^\circ$   
 $h = -29 \rightarrow 29$   
 $k = -7 \rightarrow 7$   
 $l = -24 \rightarrow 24$

### Refinement

Refinement on  $F^2$   
 $R[F^2 > 2\sigma(F^2)] = 0.049$   
 $wR(F^2) = 0.162$   
 $S = 1.10$   
2888 reflections  
191 parameters  
H-atom parameters constrained

$w = 1/[\sigma^2(F_o^2) + (0.0946P)^2 + 0.909P]$   
where  $P = (F_o^2 + 2F_c^2)/3$   
 $(\Delta/\sigma)_{\max} < 0.001$   
 $\Delta\rho_{\max} = 0.37$  e Å<sup>-3</sup>  
 $\Delta\rho_{\min} = -0.33$  e Å<sup>-3</sup>  
Extinction correction: *SHELXL97*  
Extinction coefficient: 0.0072 (13)

**Table 1**

Selected geometric parameters (Å, °).

O1–C2	1.351 (2)	C9–C10	1.511 (2)
O1–C1	1.376 (2)	N2–C11	1.340 (2)
C4–O3	1.235 (2)	N2–C8	1.446 (2)
C4–C3	1.427 (3)	N2–C10	1.459 (2)
C4–C5	1.457 (2)	C3–C2	1.342 (3)
C5–C1	1.338 (2)	O4–C11	1.213 (2)
C5–O2	1.357 (2)	C8–C7	1.519 (3)
C9–N1	1.452 (2)	N1–C6	1.463 (2)
N1–C9–C10	110.40 (14)	N2–C8–C7	109.25 (14)
C11–N2–C8	121.14 (16)	N1–C7–C8	109.62 (15)
C11–N2–C10	125.39 (17)	C9–N1–C7	111.21 (13)
C8–N2–C10	113.23 (15)	C9–N1–C6	112.41 (13)
N2–C10–C9	110.17 (14)	C7–N1–C6	113.51 (14)

**Table 2**

Hydrogen-bonding geometry (Å, °).

$D-H\cdots A$	$D-H$	$H\cdots A$	$D\cdots A$	$D-H\cdots A$
O2–H2 $\cdots$ O3 <sup>i</sup>	0.82	1.97	2.6941 (18)	147
O2–H2 $\cdots$ O3	0.82	2.33	2.7527 (18)	113
C6–H6A $\cdots$ O2	0.97	2.53	2.909 (2)	103
C7–H7A $\cdots$ O1	0.97	2.54	3.060 (2)	114
C8–H8B $\cdots$ O4	0.97	2.39	2.784 (3)	104
C10–H10A $\cdots$ O5	0.97	2.27	2.696 (2)	106

Symmetry code: (i)  $\frac{1}{2} - x, \frac{3}{2} - y, -z$ .

H atoms were included in calculated positions and treated using a riding model [ $\text{C}-\text{H}(\text{aromatic}) = 0.93$  Å and  $\text{C}-\text{H}(\text{CH}_2) = 0.97$  Å, with  $U_{\text{iso}}(\text{H}) = 1.2U_{\text{eq}}(\text{parent C atom})$ ;  $\text{C}-\text{H}(\text{CH}_3) = 0.96$  Å and  $\text{O}-\text{H} = 0.82$  Å, with  $U_{\text{iso}}(\text{H}) = 1.5U_{\text{eq}}(\text{parent C, O atom})$ ].

Data collection: *X-AREA* (Stoe & Cie, 2002); cell refinement: *X-AREA*; data reduction: *X-RED32* (Stoe & Cie, 2002); program(s) used to solve structure: *SHELXS97* (Sheldrick, 1997); program(s) used to refine structure: *SHELXL97* (Sheldrick, 1997); molecular graphics: *ORTEPIII* (Burnett & Johnson, 1996); software used to prepare material for publication: *WinGX* (Farrugia, 1999) and *PARST* (Nardelli, 1995).

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