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Hua-Yue Wu^{a*} and Zi-Kai Pan^a^aDepartment of Chemistry and Material Science, Wenzhou Normal College, Wenzhou 325027, People's Republic of China, and ^bCollege of Pharmaceutical Sciences, Zhejiang University of Technology, Hangzhou 310014, People's Republic of China

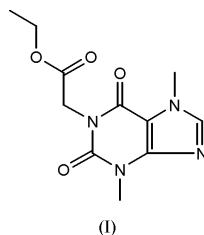
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Key indicatorsSingle-crystal X-ray study
T = 298 K
Mean $\sigma(C-C)$ = 0.003 Å
R factor = 0.044
wR factor = 0.109
Data-to-parameter ratio = 12.7For details of how these key indicators were automatically derived from the article, see <http://journals.iucr.org/e>.**Ethyl (3,7-dimethyl-2,6-dioxo-2,3,6,7-tetrahydro-purin-1-yl)acetate**

In the title compound, C₁₁H₁₄N₄O₄, the ethyl acetate portion of the organic molecule is nearly planar and makes a dihedral angle of 74.55 (6)° with the plane of the theobromine heterocycle. The crystal structure is stabilized by inter- and intramolecular C—H···O hydrogen bonds and intermolecular C—H···N hydrogen bonds.

Comment

Theobromine is an important class of alkaloid which can cause significant physiological effects, such as strong diuresis, cardiac stimulation as well as arterial dilation and is a useful product in the food and pharmaceutical industries (Mohamed *et al.*, 2002). This has stimulated our interest in its derivatives and the synthesis of ethyl theobromineacetate, (I), is another example of microwave-assisted synthesis (Kingston & Haswell, 1997; Loupy, 2002) without using any solvent (Loupy *et al.*, 1998).



In (I), there is a theobromine heterocycle and an ethyl acetate portion (Fig. 1). The theobromine ring atoms C5, C6, C7, C8, N1, N2, N3, N4 and C11 are essentially planar, with an average deviation from planarity of 0.0156 Å. This is similar to that found in the parent theobromine (Ford *et al.*, 1998). Atoms C1, C2, O1, O2, C3 and C4 of the ethyl acetate portion are also coplanar, with an average deviation of 0.0137 Å; the

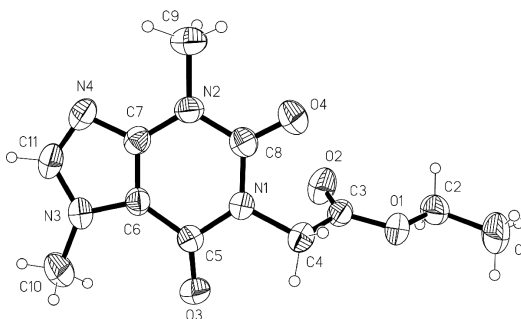


Figure 1
The molecule structure of (I), with the atom numbering. Displacement ellipsoids are drawn at the 50% probability level. H atoms are shown as spheres of arbitrary radius.

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dihedral angle between these two planes is 74.55 (6)°. Moreover, in the crystal structure there are two intramolecular and three intermolecular C—H···O hydrogen-bond interactions, as well as an intermolecular C—H···N hydrogen-bond interaction, with distances in the range 2.741 (2)–3.516 (3) Å (Table 1), resulting in the formation of a three-dimensional network structure (Fig. 2).

Experimental

The title compound was prepared by the hydrocarbylation of theobromine (1.8 g, 10 mmol) and ethyl bromoacetate (1.2 ml, 11 mmol) in the presence of tetrabutylammonium bromide catalyst (0.3 g). The reaction was carried out in a 630 W microwave oven for 1 min. Single crystals were obtained by using ethanol as the solvent for recrystallization (m.p. 451–452 K). ¹H NMR (300 MHz, CDCl₃, p.p.m.): 7.53 (s, 1H, —N=CH), 4.76 (s, 2H, —NCH₂), 4.22 (q, 2H, *J* = 7.14 Hz, OCH₂CH₃), 3.98 (s, 3H, —NCH₃), 3.59 (s, 3H, —NCH₃), 1.29 (t, 3H, *J* = 7.14 Hz, OCH₂CH₃); IR (KBr, cm⁻¹): 2925, 1657, 1547, 1484, 1236, 963, 749, 612; MS: *m/z*, 266 (*M*⁺). Analysis calculated for C₁₁H₁₄N₄O₄: C 49.62, H 5.30, N 21.04%; found: C 49.57, H 5.33, N 21.11%.

Crystal data

C ₁₁ H ₁₄ N ₄ O ₄	<i>D</i> _x = 1.432 Mg m ⁻³
<i>M</i> _r = 266.26	Mo <i>K</i> α radiation
Monoclinic, <i>P</i> 2 ₁ / <i>c</i>	Cell parameters from 2072 reflections
<i>a</i> = 8.0564 (3) Å	<i>θ</i> = 2.6–25.0°
<i>b</i> = 20.5089 (9) Å	<i>μ</i> = 0.11 mm ⁻¹
<i>c</i> = 7.8112 (3) Å	<i>T</i> = 298 (2) K
<i>β</i> = 106.833 (2)°	Block, colorless
<i>V</i> = 1235.33 (9) Å ³	0.25 × 0.23 × 0.18 mm
<i>Z</i> = 4	

Data collection

Bruker SMART APEX area-detector diffractometer	2214 independent reflections
<i>φ</i> and <i>ω</i> scans	1851 reflections with <i>I</i> > 2σ(<i>I</i>)
Absorption correction: multi-scan (SADABS; Bruker, 2002)	<i>R</i> _{int} = 0.022
<i>T</i> _{min} = 0.972, <i>T</i> _{max} = 0.973	<i>θ</i> _{max} = 25.2°
6424 measured reflections	<i>h</i> = -8 → 9
	<i>k</i> = -24 → 14
	<i>l</i> = -9 → 9

Refinement

Refinement on <i>F</i> ²	$w = 1/[\sigma^2(F_o^2) + (0.0479P)^2 + 0.3391P]$
$R[F^2 > 2\sigma(F^2)] = 0.044$	where $P = (F_o^2 + 2F_c^2)/3$
$wR(F^2) = 0.109$	(Δ/σ) _{max} < 0.001
<i>S</i> = 1.05	$\Delta\rho_{max} = 0.18 \text{ e } \text{Å}^{-3}$
2214 reflections	$\Delta\rho_{min} = -0.20 \text{ e } \text{Å}^{-3}$
175 parameters	
H-atom parameters constrained	

Table 1

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>
C10—H10A···O4 ⁱ	0.96	2.54	3.502 (2)	175
C9—H9A···O2 ⁱⁱ	0.96	2.53	3.404 (2)	152
C9—H9A···O4	0.96	2.43	2.766 (2)	100
C4—H4A···O3 ⁱⁱⁱ	0.97	2.41	3.346 (2)	161
C4—H4A···O3	0.97	2.30	2.741 (2)	107
C2—H2A···N4 ^{iv}	0.97	2.61	3.516 (3)	156

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

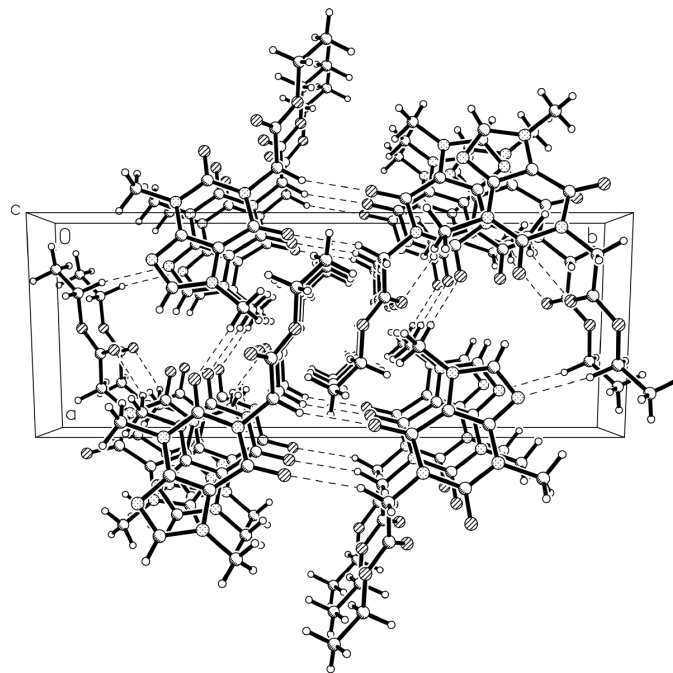


Figure 2

The three-dimensional network of (I), viewed down the *c* axis. Dashed lines indicate hydrogen bonds.

All H atoms were positioned geometrically and allowed to ride on their parent atoms, with *Csp*²—H = 0.93 Å and *U*_{iso} = 1.2*U*_{eq}(C), and *Csp*³—H = 0.96 Å and *U*_{iso} = 1.5*U*_{eq}(C).

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: XP (Bruker, 2002); software used to prepare material for publication: SHELXL97.

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