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## Structure Reports

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# Monoclinic polymorph of 3,7-dimethyl-1-(5-oxohexyl)-3,7-dihydro-1H-purine-2,6-dione

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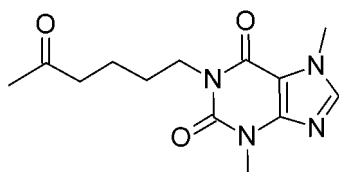
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 Key indicators: single-crystal X-ray study;  $T = 190$  K; mean  $\sigma(\text{C}-\text{C}) = 0.004$  Å;  $R$  factor = 0.066;  $wR$  factor = 0.163; data-to-parameter ratio = 16.7.

The structure of the title compound, pentoxifylline,  $\text{C}_{13}\text{H}_{18}\text{N}_4\text{O}_3$ , has been previously characterized as a triclinic polymorph [Pavelčík *et al.* (1989). *Acta Cryst.* **C45**, 836–837]. We have discovered the monoclinic form. There are no strong hydrogen bonds in the crystal structure, rather, moderate C—H $\cdots$ O hydrogen bonds are present, which serve to stabilize the three-dimensional architecture.

## Related literature

For general background to pentoxifylline, see Dettelbach & Aviado (1985). For the structure and the nature of the hydrogen bonding in the triclinic polymorph, see: Pavelčík *et al.* (1989); Gilli (2002).



## Experimental

### Crystal data

$\text{C}_{13}\text{H}_{18}\text{N}_4\text{O}_3$   
 $M_r = 278.31$   
 Monoclinic,  $P2_1/c$   
 $a = 9.743$  (6) Å

$b = 17.410$  (8) Å  
 $c = 7.956$  (3) Å  
 $\beta = 90.89$  (2)°  
 $V = 1349.4$  (12) Å<sup>3</sup>

$Z = 4$   
 Mo  $K\alpha$  radiation  
 $\mu = 0.10$  mm<sup>-1</sup>

$T = 190$  K  
 $0.40 \times 0.30 \times 0.05$  mm

### Data collection

Nonius KappaCCD diffractometer  
 5073 measured reflections  
 3103 independent reflections

1817 reflections with  $I > 2\sigma(I)$   
 $R_{\text{int}} = 0.053$

### Refinement

$R[F^2 > 2\sigma(F^2)] = 0.066$   
 $wR(F^2) = 0.163$   
 $S = 1.01$   
 3065 reflections

184 parameters  
 H-atom parameters constrained  
 $\Delta\rho_{\text{max}} = 0.24$  e Å<sup>-3</sup>  
 $\Delta\rho_{\text{min}} = -0.20$  e Å<sup>-3</sup>

**Table 1**

Hydrogen-bond geometry (Å, °).

$D-H\cdots A$	$D-H$	$H\cdots A$	$D\cdots A$	$D-H\cdots A$
$\text{C2}-\text{H2}\cdots\text{O18}^i$	0.93	2.39	3.206 (4)	147
$\text{C15}-\text{H15B}\cdots\text{O19}^{ii}$	0.96	2.60	3.439 (4)	147
$\text{C16}-\text{H16A}\cdots\text{O18}^i$	0.96	2.55	3.395 (4)	148

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

Data collection: *KappaCCD Server Software* (Nonius, 1997); cell refinement: *HKL SCALEPACK* (Otwinovski & Minor, 1997); data reduction: *HKL DENZO* (Otwinovski & Minor, 1997) and *SCALEPACK*; program(s) used to solve structure: *SHELXS97* (Sheldrick, 2008); program(s) used to refine structure: *SHELXL97* (Sheldrick, 2008); molecular graphics: *ORTEP-3* (Farrugia, 1997); software used to prepare material for publication: *SHELXL97*.

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Supplementary data and figures for this paper are available from the IUCr electronic archives (Reference: TK2793).

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## supporting information

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## Monoclinic polymorph of 3,7-dimethyl-1-(5-oxohexyl)-3,7-dihydro-1*H*-purine-2,6-dione

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### S1. Comment

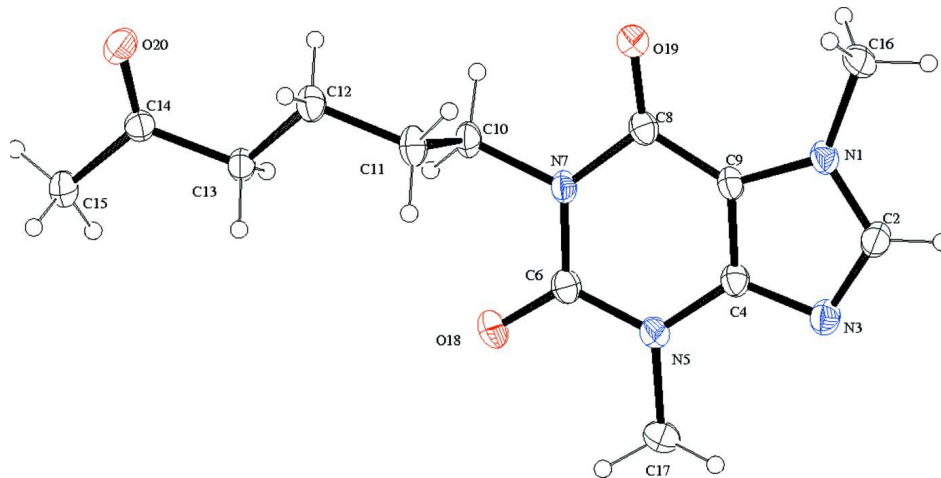
The background to pentoxifylline has been summarized (Dettelbach & Aviado, 1985). The polymorph structure **II** differs from the known form **I** Pavelčík *et al.*, 1989; Gilli, 2002) in their molecular conformation and the packing of the molecules in the lattice. The bond lengths of forms **I** and **II** are close to their standard values, but torsion angle N7—C10—C11—C12 (75.6 (3)° for **I** and 175.8 (2)° for **II**) and some bond angles significantly differ: N7—C10—C11 (114.2 (2)° for **I** and 111.9 (2)° for **II**), C10—C11—C12 (115.4 (2)° for **I** and 115.4 (2)° for **II**), C11—C12—C13 (111.0 (2)° for **I** and 114.1 (2)° for **II**). In the crystal structure of **II**, the molecules are connected by means of C—H···O hydrogen bonds, Table 1.

### S2. Experimental

Title compound was obtained by recrystallization of Trental tablets, produced by Sanofi-Aventis Deutschland GmbH. Crystals were grown by slow evaporation from dichloromethane at temperature range 308–315 K.

### S3. Refinement

All hydrogen atoms were positioned geometrically with C—H distances ranging from 0.93 to 0.97 Å and refined as riding on their parent atoms with  $U_{\text{iso}}(\text{H}) = 1.5U_{\text{eq}}(\text{C})$  for methyl groups and  $U_{\text{iso}}(\text{H}) = 1.2U_{\text{eq}}(\text{C})$  for others.



**Figure 1**

The molecular structure of the title compound, **II**, showing 50% probability ellipsoids and hydrogen atoms are shown as small spheres of arbitrary radii.

**3,7-Dimethyl-1-(5-oxohexyl)-3,7-dihydro-1H-purine-2,6-dione***Crystal data*C<sub>13</sub>H<sub>18</sub>N<sub>4</sub>O<sub>3</sub> $M_r = 278.31$ Monoclinic,  $P2_1/c$  $a = 9.743$  (6) Å $b = 17.410$  (8) Å $c = 7.956$  (3) Å $\beta = 90.89$  (2)° $V = 1349.4$  (12) Å<sup>3</sup> $Z = 4$  $F(000) = 592$  $D_x = 1.370$  Mg m<sup>-3</sup>

Melting point: 365 K

Mo  $K\alpha$  radiation,  $\lambda = 0.71073$  Å

Cell parameters from 6273 reflections

 $\theta = 1.0$ – $27.5^\circ$  $\mu = 0.10$  mm<sup>-1</sup> $T = 190$  K

Plate, colourless

 $0.4 \times 0.3 \times 0.05$  mm*Data collection*

Nonius KappaCCD

diffractometer

Radiation source: fine-focus sealed tube

Graphite monochromator

CCD scans

5073 measured reflections

3065 independent reflections

1817 reflections with  $I > 2\sigma(I)$  $R_{\text{int}} = 0.053$  $\theta_{\text{max}} = 27.5^\circ$ ,  $\theta_{\text{min}} = 2.4^\circ$  $h = -12 \rightarrow 12$  $k = -22 \rightarrow 19$  $l = -10 \rightarrow 10$ *Refinement*Refinement on  $F^2$ 

Least-squares matrix: full

 $R[F^2 > 2\sigma(F^2)] = 0.066$  $wR(F^2) = 0.150$  $S = 1.02$ 

3065 reflections

184 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-atom parameters constrained

 $w = 1/[\sigma^2(F_o^2) + (0.0525P)^2 + 0.5672P]$ where  $P = (F_o^2 + 2F_c^2)/3$  $(\Delta/\sigma)_{\text{max}} = 0.004$  $\Delta\rho_{\text{max}} = 0.24$  e Å<sup>-3</sup> $\Delta\rho_{\text{min}} = -0.20$  e Å<sup>-3</sup>*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 (Å<sup>2</sup>)*

	$x$	$y$	$z$	$U_{\text{iso}}^*/U_{\text{eq}}$
O19	-0.79244 (18)	0.03718 (9)	-0.4841 (2)	0.0407 (5)
O18	-0.54702 (19)	-0.15692 (9)	-0.2345 (2)	0.0431 (5)
N7	-0.6688 (2)	-0.05902 (10)	-0.3550 (2)	0.0314 (5)
O20	-1.1444 (2)	-0.31616 (11)	-0.4130 (3)	0.0550 (6)
N1	-0.5641 (2)	0.14522 (11)	-0.3470 (2)	0.0337 (5)

N5	-0.4583 (2)	-0.03670 (11)	-0.2089 (2)	0.0350 (5)
C9	-0.5831 (2)	0.06649 (13)	-0.3380 (3)	0.0304 (5)
N3	-0.3878 (2)	0.09676 (12)	-0.1963 (3)	0.0380 (5)
C4	-0.4742 (2)	0.04006 (13)	-0.2463 (3)	0.0319 (6)
C10	-0.7740 (3)	-0.11505 (13)	-0.4085 (3)	0.0351 (6)
H10A	-0.7303	-0.1639	-0.4317	0.042*
H10B	-0.8180	-0.0973	-0.5116	0.042*
C6	-0.5568 (3)	-0.08839 (14)	-0.2631 (3)	0.0331 (6)
C13	-0.9288 (3)	-0.26965 (13)	-0.3170 (3)	0.0323 (6)
H13A	-0.8532	-0.2731	-0.3946	0.039*
H13B	-0.8920	-0.2790	-0.2049	0.039*
C14	-1.0310 (3)	-0.33177 (14)	-0.3593 (3)	0.0326 (6)
C12	-0.9861 (2)	-0.18878 (13)	-0.3236 (3)	0.0361 (6)
H12A	-1.0633	-0.1856	-0.2484	0.043*
H12B	-1.0203	-0.1787	-0.4366	0.043*
C15	-0.9876 (3)	-0.41325 (14)	-0.3319 (3)	0.0412 (6)
H15A	-1.0331	-0.4457	-0.4128	0.062*
H15B	-1.0117	-0.4290	-0.2206	0.062*
H15C	-0.8900	-0.4173	-0.3446	0.062*
C17	-0.3361 (3)	-0.06489 (15)	-0.1159 (3)	0.0427 (6)
H17A	-0.2675	-0.0805	-0.1942	0.064*
H17B	-0.3612	-0.1079	-0.0474	0.064*
H17C	-0.3002	-0.0246	-0.0456	0.064*
C8	-0.6905 (3)	0.01842 (13)	-0.4012 (3)	0.0324 (6)
C11	-0.8821 (3)	-0.12644 (14)	-0.2754 (3)	0.0388 (6)
H11A	-0.9303	-0.0784	-0.2583	0.047*
H11B	-0.8373	-0.1403	-0.1700	0.047*
C16	-0.6522 (3)	0.20142 (14)	-0.4321 (3)	0.0420 (6)
H16A	-0.6097	0.2511	-0.4268	0.063*
H16B	-0.7394	0.2034	-0.3778	0.063*
H16C	-0.6653	0.1867	-0.5475	0.063*
C2	-0.4478 (3)	0.15899 (15)	-0.2608 (3)	0.0386 (6)
H2	-0.4118	0.2081	-0.2469	0.046*

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

	$U^{11}$	$U^{22}$	$U^{33}$	$U^{12}$	$U^{13}$	$U^{23}$
O19	0.0381 (11)	0.0348 (10)	0.0490 (10)	-0.0019 (8)	-0.0061 (8)	0.0061 (8)
O18	0.0530 (12)	0.0232 (9)	0.0530 (11)	0.0008 (8)	-0.0011 (8)	0.0021 (8)
N7	0.0359 (12)	0.0224 (10)	0.0358 (11)	-0.0046 (9)	0.0003 (8)	-0.0003 (8)
O20	0.0438 (13)	0.0385 (11)	0.0818 (14)	-0.0051 (9)	-0.0226 (10)	-0.0014 (10)
N1	0.0327 (12)	0.0231 (10)	0.0452 (12)	0.0017 (9)	0.0017 (9)	0.0010 (8)
N5	0.0334 (12)	0.0288 (11)	0.0428 (12)	0.0007 (9)	-0.0018 (9)	0.0033 (9)
C9	0.0322 (13)	0.0217 (11)	0.0374 (13)	-0.0010 (10)	0.0039 (10)	0.0004 (10)
N3	0.0362 (12)	0.0304 (11)	0.0473 (12)	-0.0030 (10)	-0.0003 (9)	-0.0037 (9)
C4	0.0313 (14)	0.0261 (12)	0.0384 (13)	0.0001 (11)	0.0032 (10)	-0.0009 (10)
C10	0.0425 (15)	0.0245 (12)	0.0383 (13)	-0.0080 (11)	0.0011 (11)	-0.0020 (10)
C6	0.0370 (14)	0.0279 (13)	0.0346 (13)	0.0003 (11)	0.0044 (10)	-0.0021 (10)

C13	0.0356 (14)	0.0267 (12)	0.0345 (12)	-0.0052 (10)	-0.0001 (10)	0.0001 (10)
C14	0.0366 (15)	0.0322 (13)	0.0290 (12)	-0.0029 (11)	-0.0008 (10)	-0.0002 (10)
C12	0.0335 (14)	0.0284 (13)	0.0465 (14)	-0.0009 (11)	0.0060 (11)	-0.0015 (11)
C15	0.0443 (16)	0.0282 (13)	0.0510 (15)	-0.0050 (12)	-0.0026 (12)	0.0002 (11)
C17	0.0404 (16)	0.0372 (14)	0.0503 (16)	0.0079 (12)	-0.0051 (12)	0.0051 (12)
C8	0.0332 (14)	0.0294 (13)	0.0347 (13)	0.0002 (11)	0.0061 (11)	0.0011 (10)
C11	0.0453 (16)	0.0253 (12)	0.0459 (15)	-0.0048 (11)	0.0095 (11)	-0.0065 (11)
C16	0.0392 (15)	0.0289 (14)	0.0578 (16)	0.0077 (11)	-0.0040 (12)	0.0066 (11)
C2	0.0322 (15)	0.0328 (14)	0.0509 (15)	-0.0039 (11)	-0.0013 (11)	-0.0039 (12)

*Geometric parameters (Å, °)*

O19—C8	1.228 (3)	C13—C12	1.515 (3)
O18—C6	1.218 (3)	C13—H13A	0.9700
N7—C6	1.401 (3)	C13—H13B	0.9700
N7—C8	1.413 (3)	C14—C15	1.495 (3)
N7—C10	1.473 (3)	C12—C11	1.530 (3)
O20—C14	1.209 (3)	C12—H12A	0.9700
N1—C2	1.337 (3)	C12—H12B	0.9700
N1—C9	1.385 (3)	C15—H15A	0.9600
N1—C16	1.461 (3)	C15—H15B	0.9600
N5—C4	1.377 (3)	C15—H15C	0.9600
N5—C6	1.380 (3)	C17—H17A	0.9600
N5—C17	1.476 (3)	C17—H17B	0.9600
C9—C4	1.359 (3)	C17—H17C	0.9600
C9—C8	1.426 (3)	C11—H11A	0.9700
N3—C2	1.330 (3)	C11—H11B	0.9700
N3—C4	1.353 (3)	C16—H16A	0.9600
C10—C11	1.518 (3)	C16—H16B	0.9600
C10—H10A	0.9700	C16—H16C	0.9600
C10—H10B	0.9700	C2—H2	0.9300
C13—C14	1.505 (3)		
C6—N7—C8	126.6 (2)	C13—C12—H12A	108.7
C6—N7—C10	116.24 (19)	C11—C12—H12A	108.7
C8—N7—C10	117.1 (2)	C13—C12—H12B	108.7
C2—N1—C9	105.3 (2)	C11—C12—H12B	108.7
C2—N1—C16	127.2 (2)	H12A—C12—H12B	107.6
C9—N1—C16	127.5 (2)	C14—C15—H15A	109.5
C4—N5—C6	119.3 (2)	C14—C15—H15B	109.5
C4—N5—C17	121.2 (2)	H15A—C15—H15B	109.5
C6—N5—C17	119.4 (2)	C14—C15—H15C	109.5
C4—C9—N1	105.0 (2)	H15A—C15—H15C	109.5
C4—C9—C8	123.6 (2)	H15B—C15—H15C	109.5
N1—C9—C8	131.4 (2)	N5—C17—H17A	109.5
C2—N3—C4	102.3 (2)	N5—C17—H17B	109.5
N3—C4—C9	112.8 (2)	H17A—C17—H17B	109.5
N3—C4—N5	125.2 (2)	N5—C17—H17C	109.5

C9—C4—N5	121.9 (2)	H17A—C17—H17C	109.5
N7—C10—C11	111.85 (19)	H17B—C17—H17C	109.5
N7—C10—H10A	109.2	O19—C8—N7	120.7 (2)
C11—C10—H10A	109.2	O19—C8—C9	128.1 (2)
N7—C10—H10B	109.2	N7—C8—C9	111.2 (2)
C11—C10—H10B	109.2	C10—C11—C12	112.4 (2)
H10A—C10—H10B	107.9	C10—C11—H11A	109.1
O18—C6—N5	121.9 (2)	C12—C11—H11A	109.1
O18—C6—N7	120.9 (2)	C10—C11—H11B	109.1
N5—C6—N7	117.2 (2)	C12—C11—H11B	109.1
C14—C13—C12	114.7 (2)	H11A—C11—H11B	107.8
C14—C13—H13A	108.6	N1—C16—H16A	109.5
C12—C13—H13A	108.6	N1—C16—H16B	109.5
C14—C13—H13B	108.6	H16A—C16—H16B	109.5
C12—C13—H13B	108.6	N1—C16—H16C	109.5
H13A—C13—H13B	107.6	H16A—C16—H16C	109.5
O20—C14—C15	121.3 (2)	H16B—C16—H16C	109.5
O20—C14—C13	121.0 (2)	N3—C2—N1	114.6 (2)
C15—C14—C13	117.6 (2)	N3—C2—H2	122.7
C13—C12—C11	114.1 (2)	N1—C2—H2	122.7
C2—N1—C9—C4	-0.3 (2)	C8—N7—C6—O18	-177.9 (2)
C16—N1—C9—C4	179.9 (2)	C10—N7—C6—O18	2.0 (3)
C2—N1—C9—C8	177.9 (2)	C8—N7—C6—N5	1.3 (3)
C16—N1—C9—C8	-2.0 (4)	C10—N7—C6—N5	-178.80 (19)
C2—N3—C4—C9	0.1 (3)	C12—C13—C14—O20	8.2 (3)
C2—N3—C4—N5	-179.1 (2)	C12—C13—C14—C15	-171.5 (2)
N1—C9—C4—N3	0.1 (3)	C14—C13—C12—C11	178.31 (19)
C8—C9—C4—N3	-178.2 (2)	C6—N7—C8—O19	179.1 (2)
N1—C9—C4—N5	179.3 (2)	C10—N7—C8—O19	-0.8 (3)
C8—C9—C4—N5	1.0 (4)	C6—N7—C8—C9	-1.5 (3)
C6—N5—C4—N3	177.8 (2)	C10—N7—C8—C9	178.54 (19)
C17—N5—C4—N3	-3.8 (4)	C4—C9—C8—O19	179.7 (2)
C6—N5—C4—C9	-1.3 (3)	N1—C9—C8—O19	1.8 (4)
C17—N5—C4—C9	177.1 (2)	C4—C9—C8—N7	0.4 (3)
C6—N7—C10—C11	87.7 (2)	N1—C9—C8—N7	-177.5 (2)
C8—N7—C10—C11	-92.3 (2)	N7—C10—C11—C12	-175.8 (2)
C4—N5—C6—O18	179.4 (2)	C13—C12—C11—C10	71.4 (3)
C17—N5—C6—O18	1.0 (3)	C4—N3—C2—N1	-0.3 (3)
C4—N5—C6—N7	0.2 (3)	C9—N1—C2—N3	0.3 (3)
C17—N5—C6—N7	-178.2 (2)	C16—N1—C2—N3	-179.8 (2)

Hydrogen-bond geometry ( $\text{\AA}$ ,  $^\circ$ )

$D-H\cdots A$	$D-H$	$H\cdots A$	$D\cdots A$	$D-H\cdots A$
C2—H2 $\cdots$ O18 <sup>i</sup>	0.93	2.39	3.206 (4)	147

C15—H15B···O19 <sup>ii</sup>	0.96	2.60	3.439 (4)	147
C16—H16A···O18 <sup>i</sup>	0.96	2.55	3.395 (4)	148

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Symmetry codes: (i)  $-x-1, y+1/2, -z-1/2$ ; (ii)  $-x-2, y-1/2, -z-1/2$ .