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Key indicators

Single-crystal X-ray study

T = 292 K

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

R factor = 0.055

wR factor = 0.174

Data-to-parameter ratio = 17.8

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

2-Benzylmalononitrile

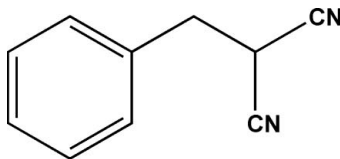
The reaction of 5-methyl-2-phenyl-2*H*-1,2,4-triazol-3(4*H*)-one, benzaldehyde and malononitrile in ethanol in the presence of triethylamine produced the title compound, $\text{C}_{10}\text{H}_8\text{N}_2$, rather than the anticipated polysubstituted 2,6-dicyanoaniline. The plane defined by the phenyl *ipso*-C atom, the methylene C atom and the central C atom of the malononitrile group is roughly orthogonal to the benzene ring plane [dihedral angle = $81.6(2)^\circ$]; one of the nitrile groups of the malononitrile almost lies in this plane [corresponding $\text{C}(\text{Ph})-\text{CH}_2-\text{C}-\text{CN}$ torsion angle = $176.06(17)^\circ$].

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Comment

Cui *et al.* (2005) synthesized polysubstituted 2,6-dicyanoanilines *via* a three-component reaction of aldehydes, ketones and malononitrile in the presence of triethylamine. We attempted to conduct this reaction using 5-methyl-2-phenyl-2*H*-1,2,4-triazol-3(4*H*)-one as a ketone component. However, we were unable to find any of the expected polysubstituted 2,6-dicyanoanilines among the products of this reaction. Instead, we isolated the title compound, 2-benzylmalononitrile, (I).



(I)

The plane of the benzene ring C1–C6 (plane *A*) is roughly orthogonal to the plane defined by atoms C6, C7 and C8 (plane *B*); the dihedral angle *A/B* is $81.6(2)^\circ$. One of the nitrile groups (C9 \equiv N1) lies almost exactly in plane *B*, the C6–C7–C8–C9 torsion angle being $176.06(17)^\circ$.

Experimental

5-Methyl-2-phenyl-2*H*-1,2,4-triazol-3(4*H*)-one (0.175 g, 1 mmol) and malononitrile (0.183 g, 2.5 mmol) were added with stirring to a solution of benzaldehyde (0.106 g, 1 mmol) in EtOH (8 ml) in the presence of triethylamine (0.151 g, 1.5 mmol). The reaction mixture was then refluxed for 10 h and monitored by thin-layer chromatography. Upon completion of the reaction, the solvent was evaporated *in vacuo* and the residue was purified by chromatography on silica using a petroleum ether/diethyl ether (10:1) mixture as eluent to give 2-benzylmalononitrile [yield 15%; m.p. 359–361 K, literature m.p. 362–363 K (Diez-Barra *et al.*, 1991)]. Crystals suitable for X-ray diffraction analysis were grown from ethanol.

Crystal data

$C_{10}H_8N_2$
 $M_r = 156.18$
 Monoclinic, $P2_1/c$
 $a = 5.8391 (13) \text{ \AA}$
 $b = 15.145 (4) \text{ \AA}$
 $c = 9.807 (2) \text{ \AA}$
 $\beta = 99.270 (5)^\circ$
 $V = 856.0 (3) \text{ \AA}^3$

$Z = 4$
 $D_x = 1.212 \text{ Mg m}^{-3}$
 Mo $K\alpha$ radiation
 $\mu = 0.07 \text{ mm}^{-1}$
 $T = 292 (2) \text{ K}$
 Block, colourless
 $0.30 \times 0.20 \times 0.20 \text{ mm}$

Data collection

Bruker SMART CCD area-detector diffractometer
 φ and ω scans
 Absorption correction: multi-scan (SADABS; Sheldrick, 1996)
 $T_{\min} = 0.978, T_{\max} = 0.985$

6068 measured reflections
 1942 independent reflections
 1061 reflections with $I > 2\sigma(I)$
 $R_{\text{int}} = 0.099$
 $\theta_{\text{max}} = 27.5^\circ$

Refinement

Refinement on F^2
 $R[F^2 > 2\sigma(F^2)] = 0.055$
 $wR(F^2) = 0.174$
 $S = 0.94$
 1942 reflections
 109 parameters

H-atom parameters constrained
 $w = 1/[\sigma^2(F_o^2) + (0.0839P)^2]$
 where $P = (F_o^2 + 2F_c^2)/3$
 $(\Delta/\sigma)_{\text{max}} < 0.001$
 $\Delta\rho_{\text{max}} = 0.21 \text{ e \AA}^{-3}$
 $\Delta\rho_{\text{min}} = -0.16 \text{ e \AA}^{-3}$

All H atoms were initially located in a difference Fourier map, then placed in idealized positions (C—H = 0.93 Å for aromatic and 0.97 Å for aliphatic H atoms) and included in the refinement using a riding-model approximation, with $U_{\text{iso}}(\text{H}) = 1.2U_{\text{eq}}(\text{C})$.

Data collection: SMART (Bruker, 1997); cell refinement: SAINT (Bruker, 1999); data reduction: SAINT; program(s) used to solve structure: SHELXS97 (Sheldrick, 1997); program(s) used to refine structure: SHELXL97 (Sheldrick, 1997); molecular graphics: SHELXTL (Bruker, 2001); software used to prepare material for publication: SHELXTL.

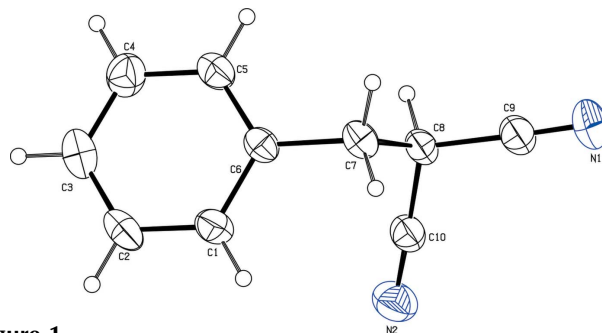


Figure 1
 The molecular structure of (I), showing the atom-labelling scheme and displacement ellipsoids drawn at the 50% probability level.

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