

4-(4-Methoxyphenyl)-2-oxo-1,2,5,6-tetrahydrobenzo[*h*]quinoline-3-carbonitrile

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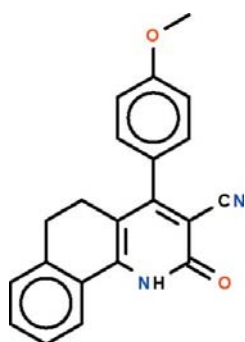
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Key indicators: single-crystal X-ray study; *T* = 100 K; mean $\sigma(\text{C}-\text{C})$ = 0.002 Å; *R* factor = 0.035; *wR* factor = 0.096; data-to-parameter ratio = 14.0.

In the molecule of the title compound, C₂₁H₁₆N₂O₂, the tetrahydrobenzo[*h*]quinoline fused-ring system is buckled owing to the ethylene –CH₂CH₂– fragment, the benzene ring and the pyridine ring being twisted by 19.7 (1)°. The 4-substituted aromatic ring is bent away from the pyridine ring by 50.3 (1)° in order to avoid crowding the cyanide substituent. In the crystal, two molecules are linked by a pair of N–H···O hydrogen bonds to form a centrosymmetric dimer.

Related literature

For background to the anticancer properties of this class of compounds, see: Rostom *et al.* (2011).



Experimental

Crystal data

C₂₁H₁₆N₂O₂
M_r = 328.36
Monoclinic, *P*2₁/*c*
a = 14.2016 (2) Å
b = 14.4725 (2) Å
c = 7.9935 (1) Å
 β = 96.017 (1)°

V = 1633.87 (4) Å³
Z = 4
Cu *K*α radiation
 μ = 0.70 mm⁻¹
T = 100 K
0.30 × 0.25 × 0.20 mm

Data collection

Agilent SuperNova Dual diffractometer with Atlas detector
Absorption correction: multi-scan (*CrysAlis PRO*; Agilent, 2010)
*T*_{min} = 0.818, *T*_{max} = 0.873

6187 measured reflections
3211 independent reflections
3011 reflections with *I* > 2σ(*I*)
*R*_{int} = 0.014

Refinement

R[*F*² > 2σ(*F*²)] = 0.035
wR(*F*²) = 0.096
S = 1.03
3211 reflections
230 parameters

H atoms treated by a mixture of independent and constrained refinement
 $\Delta\rho_{\text{max}}$ = 0.21 e Å⁻³
 $\Delta\rho_{\text{min}}$ = -0.21 e Å⁻³

Table 1

Hydrogen-bond 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>
N1–H1···O1 ⁱ	0.90 (2)	1.94 (2)	2.823 (1)	166 (1)

Symmetry code: (i) $-x + 1, -y + 1, -z + 1$.

Data collection: *CrysAlis PRO* (Agilent, 2010); cell refinement: *CrysAlis PRO*; data reduction: *CrysAlis PRO*; program(s) used to solve structure: *SHELXS97* (Sheldrick, 2008); program(s) used to refine structure: *SHELXL97* (Sheldrick, 2008); molecular graphics: *X-SEED* (Barbour, 2001); software used to prepare material for publication: *publCIF* (Westrip, 2010).

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Supplementary data and figures for this paper are available from the IUCr electronic archives (Reference: XU5292).

References

- Agilent (2010). *CrysAlis PRO*. Agilent Technologies, Yarnton, Oxfordshire, England.
Barbour, L. J. (2001). *J. Supramol. Chem.* **1**, 189–191.
Rostom, S. A. F., Faidallah, H. M. & Al-Saadi, M. S. (2011). *Med. Chem. Res.* **20** (DOI: 10.1007/s00044-010-9469-0).
Sheldrick, G. M. (2008). *Acta Cryst.* **A64**, 112–122.
Westrip, S. P. (2010). *J. Appl. Cryst.* **43**, 920–925.