

4,7-Phenanthroline

Andrew D. Bond,* Ning Shan
and William Jones

Department of Chemistry, University of
Cambridge, Lensfield Road, Cambridge
CB2 1EW, England

Correspondence e-mail: adb29@cam.ac.uk

Key indicators

Single-crystal X-ray study
T = 180 K
Mean $\sigma(\text{C}-\text{C}) = 0.007 \text{ \AA}$
R factor = 0.063
wR factor = 0.161
Data-to-parameter ratio = 8.7

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

The structure of 4,7-phenanthroline, $\text{C}_{12}\text{H}_8\text{N}_2$, has been determined at 180 K. The molecular unit possesses pseudo- C_{2v} point symmetry but does not possess crystallographic mirror symmetry. The molecules form stacks approximately along the *b* direction, with molecules in adjacent stacks forming an interplane angle of *ca* 54° .

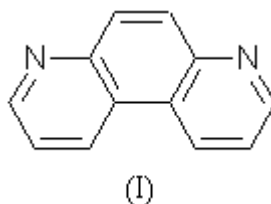
Received 15 December 2000

Accepted 9 January 2001

Online 30 January 2001

Comment

As part of a continuing study of cocrystal formation between organic acids and N-containing organic bases, we have determined the structure of 4,7-phenanthroline, (I), at 180 K. The molecular unit possesses pseudo- C_{2v} point symmetry, but does not exhibit crystallographic mirror symmetry. Similar observations have been made for the isomeric 1,10-phenanthroline (Nishigaki *et al.*, 1978). In the crystal structure, 4,7-phenanthroline forms planar stacks approximately along the *b* direction with molecules in adjacent stacks forming an interplane angle of *ca* 54° (Fig. 2); this contrasts with the observation of two approximately perpendicular layers in 1,10-phenanthroline. There is no conclusive evidence for directional C—H...N contacts in 4,7-phenanthroline, with the shortest H...N contacts, $\text{H1}\cdots\text{N7}^i = 2.72$, $\text{H8}\cdots\text{N4}^{ii} = 3.00$ and $\text{H10}\cdots\text{N7}^{iii} = 3.00 \text{ \AA}$, exhibiting C—H...N angles of 153.6 , 128.6 and 119.5° , respectively [symmetry codes: (i) $\frac{3}{2} - x, -1 + y, \frac{1}{2} + z$; (ii) $\frac{1}{2} + x, 1 - y, z$; (iii) $\frac{3}{2} - x, y, \frac{1}{2} + z$].



Experimental

4,7-Phenanthroline was obtained from Aldrich and recrystallized from ethanol.

Crystal data

$\text{C}_{12}\text{H}_8\text{N}_2$
 $M_r = 180.20$
Orthorhombic, $Pca2_1$
 $a = 19.141 (4) \text{ \AA}$
 $b = 3.8417 (4) \text{ \AA}$
 $c = 11.564 (2) \text{ \AA}$
 $V = 850.4 (3) \text{ \AA}^3$
 $Z = 4$
 $D_x = 1.408 \text{ Mg m}^{-3}$

Mo $K\alpha$ radiation
Cell parameters from 6723
reflections
 $\theta = 1.0\text{--}25.0^\circ$
 $\mu = 0.09 \text{ mm}^{-1}$
 $T = 180 (2) \text{ K}$
Plate, colourless
 $0.30 \times 0.09 \times 0.05 \text{ mm}$

Data collection

Nonius KappaCCD diffractometer
Thin-slice ω and φ scans
Absorption correction: multi-scan
(*SORTAV*; Blessing, 1995)
 $T_{\min} = 0.975$, $T_{\max} = 0.996$
2186 measured reflections
1099 independent reflections

812 reflections with $I > 2\sigma(I)$
 $R_{\text{int}} = 0.080$
 $\theta_{\text{max}} = 25.1^\circ$
 $h = -22 \rightarrow 17$
 $k = -4 \rightarrow 3$
 $l = -13 \rightarrow 10$

Refinement

Refinement on F^2
 $R[F^2 > 2\sigma(F^2)] = 0.063$
 $wR(F^2) = 0.161$
 $S = 1.09$
1099 reflections
127 parameters
H-atom parameters constrained

$w = 1/[\sigma^2(F_o^2) + (0.078P)^2]$
where $P = (F_o^2 + 2F_c^2)/3$
 $(\Delta/\sigma)_{\text{max}} = 0.002$
 $\Delta\rho_{\text{max}} = 0.30 \text{ e \AA}^{-3}$
 $\Delta\rho_{\text{min}} = -0.27 \text{ e \AA}^{-3}$
Absolute structure: Flack (1983)
Flack parameter = $-3 (8)$

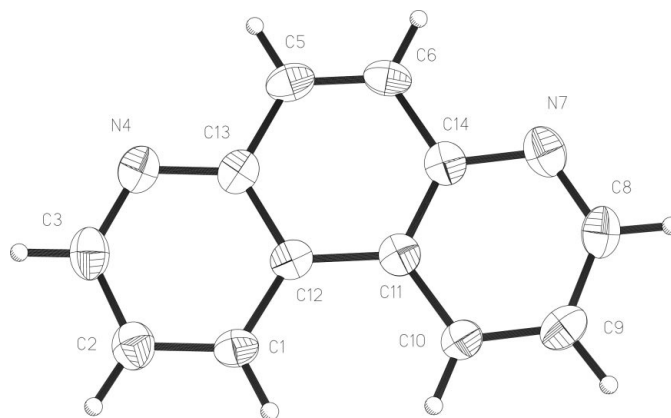


Figure 1
The molecular unit of the title compound showing displacement ellipsoids at the 50% probability level.

The absolute structure was not determined. Friedel opposites merged prior to merging of data in *Pca*₂₁. H atoms were placed geometrically and allowed to ride during subsequent refinement with an isotropic displacement parameter fixed at 1.2 times U_{iso} for the C atom to which they are attached.

Data collection: *COLLECT* (Nonius, 1998); cell refinement: *HKL SCALEPACK* (Otwinowski & Minor, 1997); data reduction: *HKL DENZO* and *SCALEPACK* (Otwinowski & Minor, 1997); program(s) used to solve structure: *SHELXS97* (Sheldrick, 1997); program(s) used to refine structure: *SHELXL97* (Sheldrick, 1997); molecular graphics: *XP* (Sheldrick, 1993); software used to prepare material for publication: *SHELXL97*.

We thank the EPSRC for financial assistance with purchase of the CCD diffractometers, and the Cambridge Overseas Trust and British Council for funding (NS).

References

Blessing, R. H. (1995). *Acta Cryst.* **A51**, 33–38.
Flack, H. D. (1983). *Acta Cryst.* **A39**, 876–881.
Otwinowski Z. & Minor W. (1997). *Methods Enzymol.* **276** 307–316.
Nishigaki, S., Yoshioka, H. & Nakatsu, K. (1978). *Acta Cryst.* **B34**, 875–879.
Nonius (1998). *COLLECT*. Nonius BV, Delft, The Netherlands.
Sheldrick, G. M. (1993). *XP*. University of Göttingen, Germany.
Sheldrick, G. M. (1997). *SHELXL97* and *SHELXS97*. University of Göttingen, Germany.

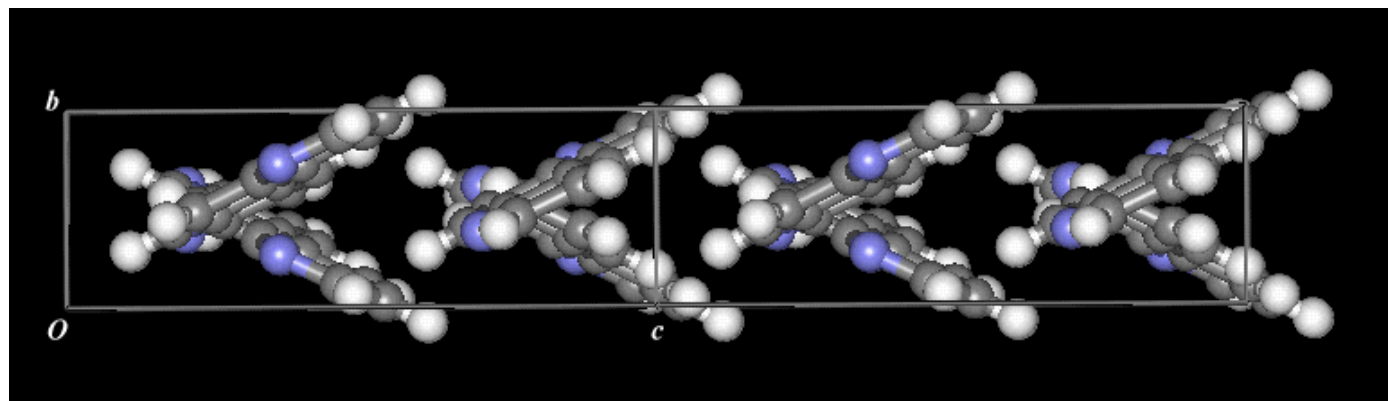


Figure 2
Projection onto (100) showing molecular stacks tilted with respect to each other with molecules in adjacent stacks forming an interplane angle of ca 54°.