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## Christian B. Nielsen, ${ }^{\text {a* }}$ Michael Pittelkow ${ }^{\text {b }}$ and Henning O. Sørensen ${ }^{\text {c }}$

${ }^{\text {a }}$ The Danish Polymer Center, Ris $\varnothing$ National Laboratory, PO Box 49, DK-4000 Roskilde, Denmark, ${ }^{\text {b }}$ University of Copenhagen, Department of Chemistry, Universitetsparken 5, 2100 Copenhagen, Denmark, and ${ }^{\text {c }}$ Center for Fundamental Research: Metal Structures in Four Dimensions, Risø National Laboratory, PO Box 49, DK-4000 Roskilde, Denmark

Correspondence e-mail:
christian.nielsen@risoe.dk

## Key indicators

Single-crystal X-ray study
$T=122 \mathrm{~K}$
Mean $\sigma(\mathrm{C}-\mathrm{C})=0.002 \AA$
$R$ factor $=0.024$
$w R$ factor $=0.057$
Data-to-parameter ratio $=22.7$
For details of how these key indicators were automatically derived from the article, see http://journals.iucr.org/e.
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## 2-(4-Bromophenyl)-1-methyl-1H-phenanthro-[9,10-d]imidazole

In the title compound, $\mathrm{C}_{22} \mathrm{H}_{15} \mathrm{BrN}_{2}$, the phenanthrene moiety is slightly skewed. The dihedral angle between the phenanthro $[9,10-d]$ imidazole mean plane and the benzene ring is $37.78(6)^{\circ}$. The crystal packing is stabilized mainly by aromatic interactions, though a weak intermolecular C $\mathrm{H} \cdots \mathrm{N}$ hydrogen bond is also observed.

## Comment

The title compound, (I) (Fig. 1), was prepared for use as a building block in the syntheses of oligo-phenylene vinylenes for non-linear optical studies. In (I), the imidazole ring is slightly bent from the skewed phenanthrene ring, with atom C16 located 0.092 (2) $\AA$ out of the least-squares plane of the phenanthro[9,10-d]imidazole system. The dihedral angle between the phenanthro $[9,10-d]$ imidazole mean plane and the benzene ring is $37.78(6)^{\circ}$. This rotation is most probably due to the adjacent $N$-methyl group.

(I)

The crystal packing (Fig. 2) is stabilized mainly by aromatic interactions. A herring-bone pattern is formed by the phenanthrene rings at $y=\frac{1}{4}$ and $\frac{3}{4}$, along with stacking of bromobenzene rings in the planes at $y=0$ and $\frac{1}{2}$. A weak intermolecular $\mathrm{C}-\mathrm{H} \cdots \mathrm{N}$ hydrogen bond (Table 1) is observed in the crystal structure.

## Experimental

The title compound was prepared in accordance with a known procedure (Krebs \& Jørgensen, 2001).

## Crystal data

$$
\begin{aligned}
& \mathrm{C}_{22} \mathrm{H}_{15} \mathrm{BrN} \mathrm{~N}_{2} \\
& M_{r}=387.27 \\
& \text { Orthorhombic, } \text { Pna }_{1} \\
& a=6.0163(4) \AA \\
& b=29.332(3) \AA \\
& c=9.1726() \AA \AA \\
& V=1618.7(2) \AA^{3} \\
& Z=4 \\
& D_{x}=1.589 \mathrm{Mg} \mathrm{~m}^{-3}
\end{aligned}
$$

$$
\text { Mo } K \alpha \text { radiation }
$$

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Figure 1
View of (I), with displacement ellipsoids at the $50 \%$ probability level. H atoms are shown as spheres of arbitrary radius.

## Data collection

Nonius KappaCCD diffractometer
$\omega$ and $\varphi$ scans
Absorption correction: Gaussian integration (Coppens, 1970)
$T_{\text {min }}=0.365, T_{\text {max }}=0.903$
36851 measured reflections
5137 independent reflections

## Refinement

Refinement on $F^{2}$
$R\left[F^{2}>2 \sigma\left(F^{2}\right)\right]=0.024$
$w R\left(F^{2}\right)=0.057$
$S=1.15$
5137 reflections
226 parameters
H -atom parameters constrained

4771 reflections with $I>2 \sigma(I)$
$R_{\text {int }}=0.046$
$\theta_{\text {max }}=31.0^{\circ}$
$h=-8 \rightarrow 8$
$k=-42 \rightarrow 42$
$l=-13 \rightarrow 13$

$$
\begin{aligned}
& w=1 /\left[\sigma^{2}\left(F_{o}{ }^{2}\right)+(0.0299 P)^{2}\right. \\
& +0.9474 P] \\
& \text { where } P=\left(F_{o}{ }^{2}+2 F_{c}{ }^{2}\right) / 3 \\
& (\Delta / \sigma)_{\max }=0.001 \\
& \Delta \rho_{\text {max }}=0.45 \mathrm{e}^{-3} \\
& \Delta \rho_{\min }=-0.34 \mathrm{e}^{-3} \\
& \text { Absolute structure: Flack (1983), } \\
& 2414 \text { Friedel reflections } \\
& \text { Flack parameter }=-0.009(6)
\end{aligned}
$$

Table 1
Hydrogen-bonding geometry $\left(\AA,^{\circ}\right)$.

| $D-\mathrm{H} \cdots A$ | $D-\mathrm{H}$ | $\mathrm{H} \cdots A$ | $D \cdots A$ | $D-\mathrm{H} \cdots A$ |
| :--- | :--- | :--- | :--- | :--- |
| $\mathrm{C} 23-\mathrm{H} 23 \cdots \mathrm{~N} 15^{\mathrm{i}}$ | 0.95 | 2.54 | $3.419(2)$ | 154 |

Symmetry code: (i) $-x,-y, \frac{1}{2}+z$.
All H atoms were located in a difference Fourier map and refined using a riding model, with fixed individual displacement parameters set at 1.2-1.5 times $U_{\text {eq }}$ of the parent atom ( $\mathrm{C}-\mathrm{H}=0.95-0.98 \AA$ ).

Data collection: COLLECT (Nonius, 1999); cell refinement: DIRAX (Duisenberg, 1992); data reduction: EvalCCD (Duisenberg et al., 2003); program(s) used to solve structure: SIR97 (Altomare et al., 1999); program(s) used to refine structure: SHELXL97 (Sheldrick, 1997); molecular graphics: ORTEPIII (Burnett \& Johnson, 1996); software used to prepare material for publication: SHELXL97.


Figure 2
The crystal packing of (I). H atoms have been omitted for clarity.

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