## Structure Reports

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## Key indicators

Single-crystal X-ray study
$T=93 \mathrm{~K}$
Mean $\sigma(\mathrm{C}-\mathrm{C})=0.007 \AA$
$R$ factor $=0.074$
$w R$ factor $=0.147$
Data-to-parameter ratio $=10.3$

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

## $N, N^{\prime}$-Di-4-pyridylperylene-3,4:9,10-bis(dicarboximide)

The title compound, $\mathrm{C}_{34} \mathrm{H}_{16} \mathrm{~N}_{4} \mathrm{O}_{4}$, is a perylene-imide pigment utilized for $\mathrm{H}_{2}$ gas sensors. The molecule has $C_{i}$ symmetry. The angle between each of the pyridyl rings and the peryleneimide skeleton is 74.5 (2) ${ }^{\circ}$. The molecules are stacked along the $c$ axis with a tilt angle of about $31.5^{\circ}$ between adjacent molecules.

## Comment

Perylene compounds are industrially important pigments, covering a variety of shades from red via maroon to black (Herbst \& Hunger, 1993). The title compound, (I), is a parapyridyl derivative, abbreviated to PPP.

(I)

We have carried out a series of investigations on $\mathrm{H}_{2}$ gas sensors utilizing a high proton affinity of organic pigments that have pyridyl rings connected directly to the choromophore (Takahashi \& Mizuguchi, 2005). The N atom of the pyridyl ring works as a strong proton acceptor and plays a decisive role for signal detection. PPP exhibits a resistivity change of about three orders of magnitude even for $0.05 \% \mathrm{H}_{2}$. Other than the present para derivative, there are also ortho and meta-derivatives and their sensitivity is slightly different, depending the site of the N atom. In this connection, structure analyses of these derivatives have been carried out. The structures of the ortho and meta derivatives are reported in the preceding two papers (Mizuguchi et al., 2005a,b). This paper reports the structure of the para derivative.

The molecule of PPP (Fig. 1) is characterized by $C_{i}$ symmetry. The angle between each of the pyridyl rings and the perylene-imide skeleton is $74.5(2)^{\circ}$. The perylene-imide skeleton is planar (r.m.s. deviation $=0.032 \AA$ ). The molecules are stacked along the $c$ axis with a tilt angle of $31.5^{\circ}$ between adjacent molecules, as shown in Fig. 2.
$\qquad$

## Experimental

PPP was synthesized by reaction of perylenetetracarboxylic dianhydride with 1,2-di-4-pyridyldiamine in dimethylnaphthalene at 490 K for 3 h according to the method of Herbst \& Hunger (1993). The product was then purified three times by sublimation at 760 K , using a two-zone furnace (Mizuguchi, 1981). Single crystals of PPP were grown from a solution in 1-chloronaphthalene After 72 h , a number of single crystals were obtained in the form of needles.

## Crystal data

| $\mathrm{C}_{34} \mathrm{H}_{16} \mathrm{~N}_{4} \mathrm{O}_{4}$ | $\mathrm{Cu} \mathrm{K} \alpha$ radiation |
| :--- | :--- |
| $M_{r}=544.51$ | Cell parameters from 8331 |
| Orthorhombic, Pcccn | $\quad$ reflections |
| $a=21.232(2) \AA$ | $\theta=3.5-68.2^{\circ}$ |
| $b=15.890(2) \AA$ | $\mu=0.85 \mathrm{~mm}^{-1}$ |
| $c=6.9311(8) \AA$ | $T=93.2 \mathrm{~K}$ |
| $V=2338.4(5) \AA^{3}$ | Needle, red |
| $Z=4$ | $0.25 \times 0.05 \times 0.05 \mathrm{~mm}$ |
| $D_{x}=1.547 \mathrm{Mg} \mathrm{m}^{-3}$ |  |

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## Data collection

Rigaku R-AXIS RAPID Imaging Plate diffractometer
$\omega$ scans
Absorption correction: multi-scan (ABSCOR; Higashi, 1995)
$T_{\text {min }}=0.554, T_{\text {max }}=0.958$
18572 measured reflections

> 1965 independent reflections
> 839 reflections with $F^{2}>2 \sigma\left(F^{2}\right)$
> $R_{\text {int }}=0.077$
> $\theta_{\max }=68.2^{\circ}$
> $h=-24 \rightarrow 23$
> $k=-18 \rightarrow 18$
> $l=-7 \rightarrow 6$

## Refinement

Refinement on $F^{2}$
H -atom parameters constrained
$R\left[F^{2}>2 \sigma\left(F^{2}\right)\right]=0.074$
$w R\left(F^{2}\right)=0.148$
$w=1 /\left[\sigma^{2}\left(F_{o}^{2}\right)\right]$
$(\Delta / \sigma)_{\text {max }}=0.001$
$\Delta \rho_{\max }=0.36$ e $\AA^{-3}$
$\Delta \rho_{\min }=-0.33 \mathrm{e}^{-3}$
1965 reflections

190 parameters

Table 1
Selected geometric parameters ( $\AA{ }^{\circ}{ }^{\circ}$ ).

| O1-C1 | 1.211 (5) | C5-C6 | 1.429 (6) |
| :---: | :---: | :---: | :---: |
| O2-C11 | 1.231 (5) | C5-C7 ${ }^{\text {i }}$ | 1.466 (6) |
| N1-C1 | 1.415 (6) | C6-C7 | 1.424 (5) |
| N1-C11 | 1.402 (5) | C6-C12 | 1.426 (6) |
| C1-C2 | 1.459 (6) | C7-C8 | 1.401 (6) |
| C2-C3 | 1.386 (6) | C8-C9 | 1.397 (6) |
| C2-C12 | 1.415 (6) | C9-C10 | 1.373 (6) |
| C3-C4 | 1.380 (6) | C10-C11 | 1.476 (6) |
| C4-C5 | 1.399 (6) | C10-C12 | 1.408 (6) |
| C1-N1-C11 | 124.4 (4) | C5 ${ }^{\text {i }}$ - $77-\mathrm{C} 6$ | 119.8 (4) |
| $\mathrm{O} 1-\mathrm{C} 1-\mathrm{N} 1$ | 119.4 (4) | $\mathrm{C} 5{ }^{\mathrm{i}}-\mathrm{C} 7-\mathrm{C} 8$ | 121.9 (4) |
| $\mathrm{O} 1-\mathrm{C} 1-\mathrm{C} 2$ | 123.5 (5) | C6-C7-C8 | 118.3 (4) |
| $\mathrm{N} 1-\mathrm{C} 1-\mathrm{C} 2$ | 117.1 (4) | C7-C8-C9 | 121.9 (4) |
| C1-C2-C3 | 119.0 (4) | C8-C9-C10 | 120.5 (4) |
| C1-C2-C12 | 120.8 (4) | C9-C10-C11 | 119.3 (4) |
| C3-C2-C12 | 120.2 (4) | C9-C10-C12 | 119.7 (4) |
| C2-C3-C4 | 120.1 (4) | C11-C10-C12 | 121.0 (4) |
| C3-C4-C5 | 122.7 (4) | $\mathrm{O} 2-\mathrm{C} 11-\mathrm{N} 1$ | 120.2 (4) |
| C4-C5-C6 | 117.7 (4) | $\mathrm{O} 2-\mathrm{C} 11-\mathrm{C} 10$ | 123.2 (4) |
| $\mathrm{C} 4-\mathrm{C} 5-\mathrm{C} 7^{\text {i }}$ | 123.2 (4) | N1-C11-C10 | 116.6 (4) |
| C6-C5-C7 ${ }^{\text {i }}$ | 119.0 (4) | C2-C12-C6 | 119.4 (4) |
| C5-C6-C7 | 121.2 (4) | C2-C12-C10 | 120.0 (4) |
| C5-C6-C12 | 119.8 (4) | C6-C12-C10 | 120.6 (4) |
| C7-C6-C12 | 119.0 (4) |  |  |

Symmetry code: (i) $\frac{3}{2}-x, \frac{3}{2}-y, z$.

All H atoms were positioned geometrically $[\mathrm{C}-\mathrm{H}=0.95 \AA$ and $\left.U_{\text {iso }}=1.2 U_{\text {eq }}(\mathrm{C})\right]$ and refined using a riding model.


Figure 1
A view of the molecular conformation of (I), showing 50\% displacement ellipsoids for the non-H atoms. Unlabelled atoms are related to labelled atoms by $\frac{3}{2}-x, \frac{3}{2}-y, z$.


The packing arrangement of PPP.

Data collection: PROCESS-AUTO (Rigaku, 1998); cell refinement: PROCESS-AUTO; data reduction: TEXSAN (Molecular Structure Corporation, 2001); program(s) used to solve structure: SHELXS86 (Sheldrick, 1985); program(s) used to refine structure: TEXSAN; molecular graphics: ORTEPIII (Burnett \& Johnson, 1996); software used to prepare material for publication: TEXSAN.

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