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## Structure Reports

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## Key indicators

Single-crystal X-ray study
$T=295 \mathrm{~K}$
Mean $\sigma(\mathrm{C}-\mathrm{C})=0.007 \AA$
$R$ factor $=0.049$
$w R$ factor $=0.098$
Data-to-parameter ratio $=14.9$
For details of how these key indicators were automatically derived from the article, see http://journals.iucr.org/e.
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# 4-(4-Bromo-5-methylthiophen-2-yl)pyridine 

The title compound, $\mathrm{C}_{10} \mathrm{H}_{8} \mathrm{BrNS}$, is used as a precursor to diarylethene derivatives. The dihedral angle between the thiophene and pyridine rings is $4.9(1)^{\circ}$, and there is evidence for conjugation throughout the molecule. The structure is stabilized by $\pi-\pi$ stacking interactions down the $c$ axis.

## Comment

Photochromic diarylethenes are among the most promising materials for optical memories and other optoelectronic devices (Irie, 2000). The title compound, (I), can be used to produce 1,2-bis[2-methyl-5-(4-pyridyl)-3-thienyl]perfluorocyclopentene and other photochromic diarylethene derivatives (Nakashima et al., 1996; Alvaro \& Lehn, 1999; Sasai et al., 2000; Matsuda et al., 2004). The dihedral angle between the pyridine and thiophene ring planes is $4.9(1)^{\circ}\left[5.2(1)^{\circ}\right.$ if the thiophene ring is extended to include the Br and methyl C atoms], suggesting a considerable degree of conjugation throughout the molecule.

(I)

Compound (I) can be compared to the structure of the related compound 1,2-bis[5-(4-pyridyl)-3-thienyl]perfluorocyclopentene [(A); Matsuda et al., 2001]. This system contains two discrete 4-pyridylthienyl groups linked through a cyclopentene ring and crystallizes with two molecules in the asymmetric unit. The C1-C6 bond in (I) $[1.456$ (6) $\AA$ ] is shorter than the equivalent bonds in $(A)$ [1.463 (4), 1.469 (4), 1.470 (4) and 1.478 (4) $\AA$ ]. Furthermore, the dihedral angles between the pyridine and thiophene rings in $(A)$ range from 3.8 (1) to $27.8(1)^{\circ}$, suggesting that steric interactions within and between the molecules of $(A)$ may be of greater importance than for (I). The structure is stabilized by columnar $\pi-\pi$ stacking interactions down the $c$ axis. The distance between molecular planes in the columns is 3.426 (5) $\AA$, with adjacent molecules stacked in an obverse fashion (Fig. 2).

## Experimental

The title compound was prepared according to the procedure of Gilat et al. (1993). Crystals were obtained by evaporation of a solution in chloroform. ${ }^{1} \mathrm{H}$ NMR $\left(\mathrm{CDCl}_{3}\right): \delta 8.59(d, 2 \mathrm{H}), 7.41(d, 1 \mathrm{H}), 7.39(d$, $1 \mathrm{H}), 7.32(s, 1 \mathrm{H}), 2.45(s, 3 \mathrm{H})$.

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Figure 1
View of (I), showing the atom-numbering scheme. Displacement ellipsoids are drawn at the $35 \%$ probability level. H atoms are represented by circles of arbitrary radius.

## Crystal data

## $\mathrm{C}_{10} \mathrm{H}_{8} \mathrm{BrNS}$

$M_{r}=254.14$
Orthorhombic, Pbca
$a=12.726$ (4) $\AA$
$b=11.629$ (4) $\AA$
$c=13.705$ (5) $\AA$
$V=2028.3(11) \AA^{3}$
$Z=8$
$D_{x}=1.665 \mathrm{Mg} \mathrm{m}^{-3}$

## Data collection

Bruker $P 4$ diffractometer $\omega$ scans
Absorption correction: multi-scan (North et al., 1968)
$T_{\text {min }}=0.217, T_{\text {max }}=0.283$
2321 measured reflections
1776 independent reflections 1028 reflections with $I>2 \sigma(I)$

## Mo $K \alpha$ radiation

Cell parameters from 47 reflections
$\theta=5.1-12.5^{\circ}$
$\mu=4.21 \mathrm{~mm}^{-1}$
$T=295$ (2) K
Prism, yellow
$0.4 \times 0.4 \times 0.3 \mathrm{~mm}$

## Refinement

Refinement on $F^{2}$
$R\left[F^{2}>2 \sigma\left(F^{2}\right)\right]=0.049$
$w R\left(F^{2}\right)=0.098$
$S=1.04$
1776 reflections
119 parameters
H -atom parameters constrained


Figure 2
A view, down the $c$ axis, of the molecular packing of (I). H atoms have been omitted.

All H atoms were refined using a riding model, with $\mathrm{C}-\mathrm{H}=0.93 \AA$ and $U_{\text {iso }}(\mathrm{H})=1.2 U_{\text {eq }}(\mathrm{C})$ for aromatic, and $\mathrm{C}-\mathrm{H}=0.96 \AA$ and $U_{\text {iso }}(\mathrm{H})=1.5 U_{\text {eq }}(\mathrm{C})$ for methyl C atoms.

Data collection: XSCANS (Bruker, 1997); cell refinement: $X S C A N S$; data reduction: XSCANS; program(s) used to solve structure: SHELXTL (Bruker, 1997); program(s) used to refine structure: SHELXTL; molecular graphics: SHELXTL; software used to prepare material for publication: SHELXTL.

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