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## Key indicators

Single-crystal X-ray study
$T=298 \mathrm{~K}$
Mean $\sigma(\mathrm{C}-\mathrm{C})=0.007 \AA$
$R$ factor $=0.045$
$w R$ factor $=0.108$
Data-to-parameter ratio $=13.3$

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## 4-Bromo-2-(2-pyridylmethyliminomethyl)phenol

Two molecules of the title compound, $\mathrm{C}_{13} \mathrm{H}_{11} \mathrm{BrN}_{2} \mathrm{O}$, are linked by an intermolecular $\mathrm{Br} \cdots \mathrm{N}_{\text {pyridyl }}$ interaction of 3.263 (4) $\AA$ across a center of inversion. The imino $N$ atom is engaged in intramolecular hydrogen bonding with the phenol group $[\mathrm{O} \cdots \mathrm{N}=2.595(5) \AA$ ].

## Comment

Schiff bases derived from the condensation of salicylaldehyde and a primary amine can be further reduced by sodium borohydride to form substituted phenol derivatives having both secondary and tertiary amino groups on the same substituent, e.g. $2-\mathrm{OH}-4-\mathrm{NO}_{2}-3-\mathrm{CH}_{2}-\mathrm{NH}-\mathrm{CH}_{2} \mathrm{CH}_{2}-\mathrm{N}\left(\mathrm{CH}_{3}\right)_{2}$, which, owing to its existence in the zwitterionic form (Hazell et al., 1997), is an excellent Lewis base that can coordinate to organotin Lewis acids. With the less electron-withdrawing Br atom in place of the $-\mathrm{NO}_{2}$ group, the compound probably does not exist in this zwitterionic form. However, the molecule is able to form complexes (Khoo, Yan, Goh \& Ng, 2000; Khoo et al., 2001). A previous study on the Schiff base 7-methoxy-3(salicyldene)aminocoumarin (Khoo, Zhang \& Ng, 2000) has revealed potentially useful lasing activity.

(I)

The bromo-substituted title compound, (I), is a monomeric compound whose hydroxy group is engaged in hydrogen bonding with the imino N atom (Fig. 1). Two molecules are linked by a $\mathrm{Br} \cdots \mathrm{N}_{\text {pyridyl }}$ interaction across an inversion center, forming a dimeric entity (Fig. 2).


Figure 1
ORTEP plot of (I), with displacement ellipsoids drawn at the $50 \%$ probability level. H atoms are drawn as spheres of arbitrary radii.

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Figure 2
ORTEP plot of the $\mathrm{Br} \cdots \mathrm{N}$-linked dimeric unit.

## Experimental

The compound was synthesized by condensing equimolar quantities of 5-bromosalicylaldehyde and 2-(aminomethyl)pyridine in chloroform, duplicating the method used for the synthesis of 2-(3-pyridylmethyliminomethyl)phenol (Cimerman et al., 1994).

## Crystal data

$\mathrm{C}_{13} \mathrm{H}_{11} \mathrm{BrN}_{2} \mathrm{O}$
$M_{r}=291.15$
Triclinic, $P \overline{1}$
$a=4.474$ (1) A
$b=9.529$ (2) $\AA$
$c=14.271$ (2) $\AA$
$\alpha=92.65(1)^{\circ}$
$\beta=93.72(1)^{\circ}$
$\gamma=95.16(2)^{\circ}$
$V=603.8$ (2) $\AA^{3}$

$$
\begin{aligned}
& Z=2 \\
& D_{x}=1.601 \mathrm{Mg} \mathrm{~m}^{-3} \\
& \text { Mo } K \alpha \text { radiation } \\
& \text { Cell parameters from } 25 \\
& \quad \text { reflections } \\
& \theta=4.5-12.7^{\circ} \\
& \mu=3.39 \mathrm{~mm}^{-1} \\
& T=298(2) \mathrm{K} \\
& \text { Block, yellow } \\
& 0.20 \times 0.20 \times 0.10 \mathrm{~mm}
\end{aligned}
$$

## Data collection

Siemens $P 4$ four-circle diffractometer
$\omega-2 \theta$ scans
Absorption correction: $\psi$ scan (North et al., 1968) $T_{\text {min }}=0.531, T_{\text {max }}=0.713$
2912 measured reflections
2069 independent reflections
1416 reflections with $I>2 \sigma(I)$

## Refinement

Refinement on $F^{2}$
$R\left[F^{2}>2 \sigma\left(F^{2}\right)\right]=0.045$
$w R\left(F^{2}\right)=0.108$
$S=1.01$
2069 reflections
155 parameters
H -atom parameters constrained

Table 1
Selected geometric parameters ( $\left({ }^{\circ},{ }^{\circ}\right)$.

| $\mathrm{Br} 1-\mathrm{C} 5$ | $1.901(4)$ | $\mathrm{C} 2-\mathrm{C} 3$ | $1.395(6)$ |
| :--- | :--- | :--- | :--- |
| $\mathrm{O} 1-\mathrm{C} 2$ | $1.343(5)$ | $\mathrm{C} 3-\mathrm{C} 4$ | $1.379(6)$ |
| $\mathrm{N} 1-\mathrm{C} 7$ | $1.269(5)$ | $\mathrm{C} 4-\mathrm{C} 5$ | $1.384(6)$ |
| $\mathrm{N} 1-\mathrm{C} 8$ | $1.458(6)$ | $\mathrm{C} 5-\mathrm{C} 6$ | $1.370(6)$ |
| $\mathrm{N} 2-\mathrm{C} 9$ | $1.330(5)$ | $\mathrm{C} 8-\mathrm{C} 9$ | $1.513(6)$ |
| $\mathrm{N} 2-\mathrm{C} 13$ | $1.319(6)$ | $\mathrm{C} 9-\mathrm{C} 10$ | $1.371(6)$ |
| $\mathrm{C} 1-\mathrm{C} 2$ | $1.396(6)$ | $\mathrm{C} 10-\mathrm{C} 11$ | $1.377(7)$ |
| $\mathrm{C} 1-\mathrm{C} 6$ | $1.385(6)$ | $\mathrm{C} 11-\mathrm{C} 12$ | $1.361(8)$ |
| $\mathrm{C} 1-\mathrm{C} 7$ | $1.461(6)$ | $\mathrm{C} 12-\mathrm{C} 13$ | $1.361(7)$ |
|  |  |  |  |
| $\mathrm{C} 7-\mathrm{N} 1-\mathrm{C} 8$ | $118.3(4)$ | $\mathrm{C} 6-\mathrm{C} 5-\mathrm{Br} 1$ | $120.4(3)$ |
| $\mathrm{C} 9-\mathrm{N} 2-\mathrm{C} 13$ | $116.9(4)$ | $\mathrm{C} 5-\mathrm{C} 6-\mathrm{C} 1$ | $120.7(4)$ |
| $\mathrm{C} 6-\mathrm{C} 1-\mathrm{C} 2$ | $119.1(4)$ | $\mathrm{N} 1-\mathrm{C} 7-\mathrm{C} 1$ | $122.2(4)$ |
| $\mathrm{C} 6-\mathrm{C} 1-\mathrm{C} 7$ | $120.1(4)$ | $\mathrm{N} 1-\mathrm{C} 8-\mathrm{C} 9$ | $109.9(4)$ |
| $\mathrm{C} 2-\mathrm{C} 1-\mathrm{C} 7$ | $120.8(4)$ | $\mathrm{N} 2-\mathrm{C} 9-\mathrm{C} 8$ | $115.0(4)$ |
| $\mathrm{O} 1-\mathrm{C} 2-\mathrm{C} 1$ | $121.8(4)$ | $\mathrm{N} 2-\mathrm{C} 9-\mathrm{C} 10$ | $122.5(4)$ |
| $\mathrm{O} 1-\mathrm{C} 2-\mathrm{C} 3$ | $118.5(4)$ | $\mathrm{C} 10-\mathrm{C} 9-\mathrm{C} 8$ | $122.5(4)$ |
| $\mathrm{C} 1-\mathrm{C} 2-\mathrm{C} 3$ | $119.6(4)$ | $\mathrm{C} 9-\mathrm{C} 10-\mathrm{C} 11$ | $119.0(5)$ |
| $\mathrm{C} 2-\mathrm{C} 3-\mathrm{C} 4$ | $120.5(4)$ | $\mathrm{C} 12-\mathrm{C} 11-\mathrm{C} 10$ | $118.7(5)$ |
| $\mathrm{C} 3-\mathrm{C} 4-\mathrm{C} 5$ | $119.2(4)$ | $\mathrm{C} 11-\mathrm{C} 12-\mathrm{C} 13$ | $118.1(5)$ |
| $\mathrm{C} 4-\mathrm{C} 5-\mathrm{C} 6$ | $120.8(4)$ | $\mathrm{N} 2-\mathrm{C} 13-\mathrm{C} 12$ | $124.7(5)$ |
| $\mathrm{C} 4-\mathrm{C} 5-\mathrm{Br} 1$ | $118.8(3)$ |  |  |

H atoms were positioned geometrically $\left(\mathrm{C}-\mathrm{H}=0.93 \AA\right.$ for $s p^{2}$ hybridized C atoms, $\mathrm{C}-\mathrm{H}=0.97 \AA$ for the $s p^{3}$-hybridized C atom and $\mathrm{O}-\mathrm{H}=0.82 \AA$ for the phenol H atom). Displacement parameters were set to $1.2 U_{\text {eq }}$ of the parent atoms for the aromatic H atoms and $1.5 U_{\text {eq }}$ for the other H atoms.

Data collection: XSCANS (Bruker, 1997); cell refinement: LEAST SQUARES in XSCANS; data reduction: REDUCE in XSCANS; program(s) used to solve structure: SHELXS97 (Sheldrick, 1997); program(s) used to refine structure: SHELXL97 (Sheldrick, 1997); molecular graphics: ORTEPII (Johnson, 1976); software used to prepare material for publication: SHELXL97.

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