

1-(4-Methylbenzylideneamino)-pyridinium iodide

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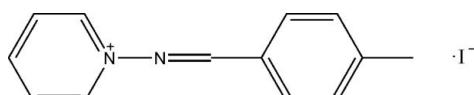
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Key indicators: single-crystal X-ray study; $T = 291\text{ K}$; mean $\sigma(\text{C}-\text{C}) = 0.017\text{ \AA}$; R factor = 0.049; wR factor = 0.125; data-to-parameter ratio = 9.6.

The title compound, $\text{C}_{13}\text{H}_{13}\text{N}_2^+\text{I}^-$, is a derivative of 1-aminopyridinium iodide. The pyridine and benzene rings are oriented at a dihedral angle of $45.78(3)^\circ$. In the crystal structure, weak intermolecular $\text{C}-\text{H}\cdots\text{I}$ hydrogen bonds link the molecules.

Related literature

For bond-length data, see: Allen *et al.* (1987).



Experimental

Crystal data

$\text{C}_{13}\text{H}_{13}\text{N}_2^+\text{I}^-$	$V = 1335.6(5)\text{ \AA}^3$
$M_r = 324.15$	$Z = 4$
Orthorhombic, $P2_12_12_1$	Mo $K\alpha$ radiation
$a = 7.1690(14)\text{ \AA}$	$\mu = 2.37\text{ mm}^{-1}$
$b = 12.399(3)\text{ \AA}$	$T = 291(2)\text{ K}$
$c = 15.026(3)\text{ \AA}$	$0.30 \times 0.10 \times 0.10\text{ mm}$

Data collection

Enraf–Nonius CAD-4 diffractometer	1408 independent reflections
Absorption correction: ψ scan (North <i>et al.</i> , 1968)	1015 reflections with $I > 2\sigma(I)$
$T_{\min} = 0.536$, $T_{\max} = 0.797$	3 standard reflections
1408 measured reflections	frequency: 120 min
	intensity decay: none

Refinement

$R[F^2 > 2\sigma(F^2)] = 0.048$	$\Delta\rho_{\max} = 1.10\text{ e \AA}^{-3}$
$wR(F^2) = 0.125$	$\Delta\rho_{\min} = -0.54\text{ e \AA}^{-3}$
$S = 1.08$	Absolute structure: Flack (1983), with no Friedel pairs
1408 reflections	Flack parameter: 0.05 (10)
146 parameters	H-atom parameters constrained

Table 1

Hydrogen-bond geometry (\AA , $^\circ$).

$D-\text{H}\cdots A$	$D-\text{H}$	$\text{H}\cdots A$	$D\cdots A$	$D-\text{H}\cdots A$
C9—H9 \cdots I ⁱ	0.93	3.06	3.795 (12)	138
C12—H12 \cdots I ⁱⁱ	0.93	3.03	3.929 (13)	162
Symmetry codes: (i) $-x + 1, y - \frac{1}{2}, -z + \frac{3}{2}$; (ii) $x - \frac{1}{2}, -y + \frac{3}{2}, -z + 1$.				

Data collection: *CAD-4 Software* (Enraf–Nonius, 1989); cell refinement: *CAD-4 Software*; data reduction: *XCAD4* (Harms & Wocadlo, 1995); program(s) used to solve structure: *SHELXS97* (Sheldrick, 2008); program(s) used to refine structure: *SHELXL97* (Sheldrick, 2008); molecular graphics: *ORTEP-3 for Windows* (Farrugia, 1997); software used to prepare material for publication: *SHELXTL* (Sheldrick, 2008).

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Supplementary data and figures for this paper are available from the IUCr electronic archives (Reference: HK2492).

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1-(4-Methylbenzylideneamino)pyridinium iodide

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S1. Comment

Some derivatives of 1-aminopyridinium iodide is important chemical materials. We report herein the crystal structure of the title compound.

In the molecule of the title compound (Fig. 1), the bond lengths (Allen *et al.*, 1987) and angles are within normal ranges. Rings A (C2-C7) and B (N1/C9-C13) are, of course, planar and they are oriented at a dihedral angle of A/B = 45.78 (3)°.

In the crystal structure, weak intermolecular C-H···I hydrogen bonds (Table 1) link the molecules (Fig. 2), in which they may be effective in the stabilization of the structure.

S2. Experimental

For the preparation of the title compound, 1-aminopyridinium iodide (22.2 g, 0.10 mol) was dissolved in ethanol (20 ml). 4-Methylbenzaldehyde (32.4 g, 0.1 mol) was added with stirring, and then the mixture was heated at reflux for 5 h. Upon cooling to room temperature, a precipitate formed, which was collected by filtration and washed with cold ethanol (2 x 10 ml) to obtain a yellow solid (yield; 38 g, 70%). Crystals suitable for X-ray analysis were obtained by slow evaporation of an ethanol solution.

S3. Refinement

H atoms were positioned geometrically, with C-H = 0.93 and 0.96 Å for aromatic and methyl H, respectively, and constrained to ride on their parent atoms with $U_{\text{iso}}(\text{H}) = xU_{\text{eq}}(\text{C})$, where $x = 1.5$ for methyl H and $x = 1.2$ for aromatic H atoms.

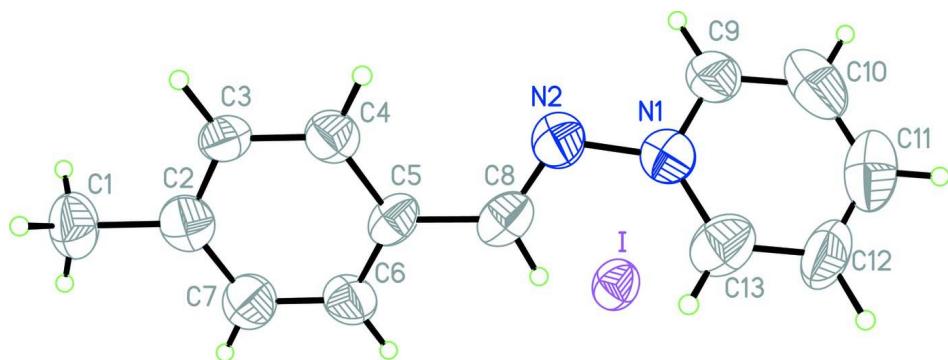
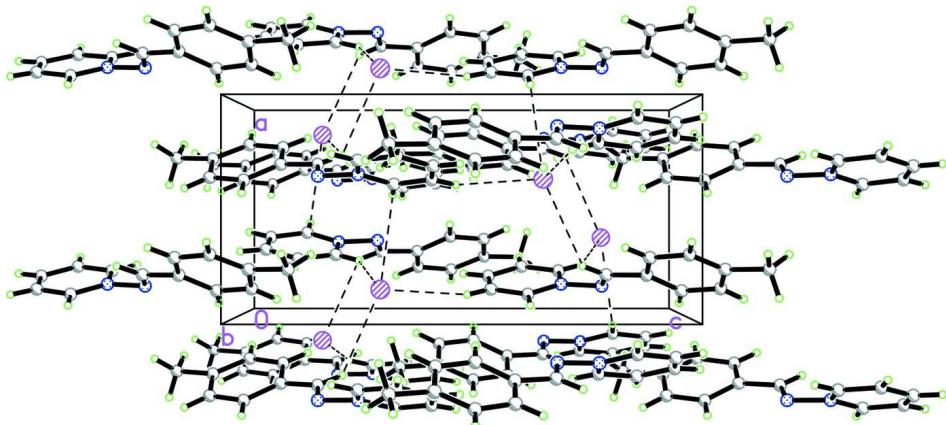


Figure 1

The molecular structure of the title molecule, with the atom-numbering scheme. Displacement ellipsoids are drawn at the 50% probability level

**Figure 2**

A packing diagram of the title molecule. Hydrogen bonds are shown as dashed lines.

1-(4-Methylbenzylideneamino)pyridinium iodide

Crystal data

$C_{13}H_{13}N_2^+ \cdot I^-$
 $M_r = 324.15$
Orthorhombic, $P2_12_12_1$
Hall symbol: P 2ac 2ab
 $a = 7.1690 (14)$ Å
 $b = 12.399 (3)$ Å
 $c = 15.026 (3)$ Å
 $V = 1335.6 (5)$ Å³
 $Z = 4$

$F(000) = 632$
 $D_x = 1.612$ Mg m⁻³
Mo $K\alpha$ radiation, $\lambda = 0.71073$ Å
Cell parameters from 25 reflections
 $\theta = 2.1\text{--}25.3^\circ$
 $\mu = 2.37$ mm⁻¹
 $T = 291$ K
Block, yellow
 $0.30 \times 0.10 \times 0.10$ mm

Data collection

Enraf–Nonius CAD-4
diffractometer

Radiation source: fine-focus sealed tube

Graphite monochromator
 $\omega/2\theta$ scans

Absorption correction: ψ scan
(North *et al.*, 1968)

$T_{\min} = 0.536$, $T_{\max} = 0.797$

1408 measured reflections

1408 independent reflections
1015 reflections with $I > 2\sigma(I)$

$R_{\text{int}} = 0.000$
 $\theta_{\max} = 25.3^\circ$, $\theta_{\min} = 2.1^\circ$
 $h = 0 \rightarrow 8$
 $k = 0 \rightarrow 14$
 $l = 0 \rightarrow 18$

3 standard reflections every 120 min
intensity decay: none

Refinement

Refinement on F^2

Least-squares matrix: full

$R[F^2 > 2\sigma(F^2)] = 0.048$

$wR(F^2) = 0.125$

$S = 1.08$

1408 reflections

146 parameters

0 restraints

Primary atom site location: structure-invariant
direct methods

Secondary atom site location: difference Fourier
map

Hydrogen site location: inferred from
neighbouring sites

H-atom parameters constrained

$w = 1/[\sigma^2(F_o^2) + (0.0599P)^2 + 0.581P]$
where $P = (F_o^2 + 2F_c^2)/3$

$(\Delta/\sigma)_{\max} < 0.001$

$\Delta\rho_{\max} = 1.10$ e Å⁻³

$\Delta\rho_{\min} = -0.54$ e Å⁻³

Absolute structure: Flack (1983), with no
Friedel pairs

Absolute structure parameter: 0.05 (10)

Special details

Geometry. All e.s.d.'s (except the e.s.d. in the dihedral angle between two l.s. planes) are estimated using the full covariance matrix. The cell e.s.d.'s are taken into account individually in the estimation of e.s.d.'s in distances, angles and torsion angles; correlations between e.s.d.'s in cell parameters are only used when they are defined by crystal symmetry. An approximate (isotropic) treatment of cell e.s.d.'s is used for estimating e.s.d.'s involving l.s. planes.

Refinement. Refinement of F^2 against ALL reflections. The weighted R-factor wR and goodness of fit S are based on F^2 , conventional R-factors R are based on F, with F set to zero for negative F^2 . The threshold expression of $F^2 > 2\text{sigma}(F^2)$ is used only for calculating R-factors(gt) etc. and is not relevant to the choice of reflections for refinement. R-factors based on F^2 are statistically about twice as large as those based on F, and R-factors based on ALL data will be even larger.

Fractional atomic coordinates and isotropic or equivalent isotropic displacement parameters (\AA^2)

	<i>x</i>	<i>y</i>	<i>z</i>	$U_{\text{iso}}^*/U_{\text{eq}}$
I	0.63765 (9)	0.75120 (8)	0.67606 (4)	0.0632 (3)
N1	0.1554 (13)	0.6174 (7)	0.7189 (6)	0.049 (2)
N2	0.1485 (13)	0.6612 (8)	0.8065 (6)	0.051 (2)
C1	0.241 (2)	1.0008 (11)	1.1391 (8)	0.068 (4)
H1A	0.3685	1.0169	1.1541	0.102*
H1B	0.1854	0.9596	1.1862	0.102*
H1C	0.1735	1.0669	1.1313	0.102*
C2	0.2358 (15)	0.9358 (10)	1.0529 (7)	0.053 (3)
C3	0.1465 (17)	0.8349 (9)	1.0519 (7)	0.055 (3)
H3	0.0964	0.8069	1.1041	0.065*
C4	0.1329 (14)	0.7771 (8)	0.9731 (7)	0.048 (3)
H4	0.0746	0.7101	0.9727	0.058*
C5	0.2057 (14)	0.8187 (9)	0.8955 (7)	0.044 (3)
C6	0.2865 (17)	0.9199 (8)	0.8960 (8)	0.055 (3)
H6	0.3312	0.9490	0.8431	0.066*
C7	0.3023 (16)	0.9792 (10)	0.9746 (8)	0.059 (3)
H7	0.3570	1.0472	0.9742	0.071*
C8	0.2037 (13)	0.7578 (10)	0.8115 (6)	0.051 (3)
H8	0.2447	0.7918	0.7599	0.062*
C9	0.2109 (17)	0.5139 (9)	0.7163 (9)	0.061 (3)
H9	0.2361	0.4769	0.7688	0.073*
C10	0.230 (2)	0.4624 (11)	0.6341 (12)	0.081 (5)
H10	0.2715	0.3914	0.6305	0.097*
C11	0.1874 (17)	0.5193 (13)	0.5595 (9)	0.072 (4)
H11	0.1953	0.4859	0.5042	0.087*
C12	0.1332 (19)	0.6237 (12)	0.5642 (8)	0.070 (4)
H12	0.1104	0.6627	0.5125	0.084*
C13	0.1126 (18)	0.6708 (11)	0.6443 (8)	0.065 (3)
H13	0.0681	0.7411	0.6478	0.078*

Atomic displacement parameters (\AA^2)

	U^{11}	U^{22}	U^{33}	U^{12}	U^{13}	U^{23}
C1	0.078 (10)	0.073 (8)	0.054 (7)	0.006 (8)	-0.006 (8)	-0.019 (7)
C2	0.046 (7)	0.058 (8)	0.055 (7)	0.013 (6)	-0.012 (6)	0.001 (6)
C3	0.068 (8)	0.046 (6)	0.050 (7)	0.009 (7)	0.016 (6)	0.011 (5)

C4	0.052 (6)	0.044 (7)	0.048 (5)	0.002 (5)	-0.007 (5)	0.001 (5)
C5	0.033 (5)	0.060 (7)	0.038 (6)	0.007 (5)	0.006 (5)	0.006 (5)
C6	0.059 (7)	0.043 (6)	0.062 (7)	-0.007 (6)	0.020 (6)	0.001 (6)
C7	0.054 (7)	0.064 (8)	0.059 (7)	0.007 (6)	0.002 (7)	0.000 (7)
C8	0.039 (4)	0.073 (8)	0.042 (5)	0.013 (8)	0.001 (4)	0.011 (8)
C9	0.068 (8)	0.055 (8)	0.061 (7)	-0.011 (7)	-0.013 (7)	0.007 (6)
C10	0.070 (10)	0.064 (9)	0.108 (12)	-0.016 (9)	0.011 (10)	-0.026 (10)
C11	0.046 (8)	0.117 (13)	0.053 (8)	-0.017 (9)	0.015 (6)	-0.025 (9)
C12	0.072 (9)	0.094 (11)	0.044 (7)	-0.016 (9)	-0.018 (7)	0.001 (7)
C13	0.064 (8)	0.075 (8)	0.054 (7)	0.010 (8)	0.007 (7)	0.009 (7)
I	0.0631 (5)	0.0721 (5)	0.0544 (4)	0.0063 (8)	0.0064 (4)	-0.0020 (6)
N1	0.044 (5)	0.050 (5)	0.053 (5)	0.000 (5)	0.006 (5)	0.001 (5)
N2	0.049 (5)	0.054 (5)	0.049 (5)	-0.012 (5)	0.021 (5)	-0.005 (4)

Geometric parameters (\AA , $^{\circ}$)

C1—C2	1.526 (15)	C8—N2	1.264 (13)
C1—H1A	0.9600	C8—H8	0.9300
C1—H1B	0.9600	C9—N1	1.344 (13)
C1—H1C	0.9600	C9—C10	1.398 (17)
C2—C7	1.379 (15)	C9—H9	0.9300
C2—C3	1.405 (16)	C10—C11	1.360 (19)
C3—C4	1.387 (15)	C10—H10	0.9300
C3—H3	0.9300	C11—C12	1.353 (17)
C4—C5	1.379 (13)	C11—H11	0.9300
C4—H4	0.9300	C12—C13	1.345 (16)
C5—C6	1.382 (14)	C12—H12	0.9300
C5—C8	1.471 (13)	C13—N1	1.338 (14)
C6—C7	1.396 (15)	C13—H13	0.9300
C6—H6	0.9300	N1—N2	1.425 (11)
C7—H7	0.9300		
C2—C1—H1A	109.5	C6—C7—H7	120.4
C2—C1—H1B	109.5	N2—C8—C5	122.7 (9)
H1A—C1—H1B	109.5	N2—C8—H8	118.6
C2—C1—H1C	109.5	C5—C8—H8	118.6
H1A—C1—H1C	109.5	N1—C9—C10	119.4 (12)
H1B—C1—H1C	109.5	N1—C9—H9	120.3
C7—C2—C3	119.7 (11)	C10—C9—H9	120.3
C7—C2—C1	120.6 (12)	C11—C10—C9	118.0 (12)
C3—C2—C1	119.4 (12)	C11—C10—H10	121.0
C4—C3—C2	120.1 (10)	C9—C10—H10	121.0
C4—C3—H3	120.0	C12—C11—C10	121.2 (13)
C2—C3—H3	120.0	C12—C11—H11	119.4
C5—C4—C3	120.2 (10)	C10—C11—H11	119.4
C5—C4—H4	119.9	C13—C12—C11	119.6 (13)
C3—C4—H4	119.9	C13—C12—H12	120.2
C4—C5—C6	119.6 (10)	C11—C12—H12	120.2

C4—C5—C8	122.0 (10)	N1—C13—C12	120.6 (12)
C6—C5—C8	118.4 (10)	N1—C13—H13	119.7
C5—C6—C7	121.1 (11)	C12—C13—H13	119.7
C5—C6—H6	119.4	C13—N1—C9	121.1 (11)
C7—C6—H6	119.4	C13—N1—N2	125.3 (9)
C2—C7—C6	119.2 (11)	C9—N1—N2	113.6 (10)
C2—C7—H7	120.4	C8—N2—N1	113.9 (9)
C7—C2—C3—C4	2.9 (16)	N1—C9—C10—C11	-2 (2)
C1—C2—C3—C4	177.1 (11)	C9—C10—C11—C12	2 (2)
C2—C3—C4—C5	-0.5 (17)	C10—C11—C12—C13	-3 (2)
C3—C4—C5—C6	-2.2 (16)	C11—C12—C13—N1	4 (2)
C3—C4—C5—C8	176.6 (10)	C12—C13—N1—C9	-3.4 (19)
C4—C5—C6—C7	2.5 (17)	C12—C13—N1—N2	176.5 (11)
C8—C5—C6—C7	-176.3 (10)	C10—C9—N1—C13	2.3 (18)
C3—C2—C7—C6	-2.6 (17)	C10—C9—N1—N2	-177.6 (11)
C1—C2—C7—C6	-176.7 (12)	C5—C8—N2—N1	179.2 (9)
C5—C6—C7—C2	0.0 (18)	C13—N1—N2—C8	-38.8 (15)
C4—C5—C8—N2	-5.4 (16)	C9—N1—N2—C8	141.1 (10)
C6—C5—C8—N2	173.3 (10)		

Hydrogen-bond geometry (Å, °)

D—H···A	D—H	H···A	D···A	D—H···A
C9—H9···I ⁱ	0.93	3.06	3.795 (12)	138
C12—H12···I ⁱⁱ	0.93	3.03	3.929 (13)	162

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