

Quinoline-2-carbonitrile

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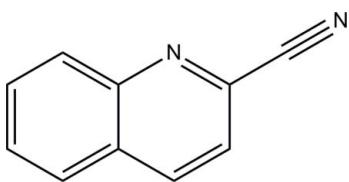
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Key indicators: single-crystal X-ray study; $T = 100\text{ K}$; mean $\sigma(\text{C}-\text{C}) = 0.004\text{ \AA}$; R factor = 0.056; wR factor = 0.141; data-to-parameter ratio = 9.7.

In the title compound, $\text{C}_{10}\text{H}_6\text{N}_2$, the molecule is almost planar, with an r.m.s. deviation of 0.014 \AA . The dihedral angle between the aromatic rings is $1.28(16)^\circ$. In the crystal, molecules are stacked along the a axis by way of weak aromatic π - π stacking interactions between the benzene and pyridine rings of adjacent molecules [centroid–centroid separation = $3.7943(19)\text{ \AA}$].

Related literature

For the biological activity and syntheses of quinoline derivatives, see: Sasaki *et al.* (1998); Reux *et al.* (2009). For related structures, see: Fun *et al.* (2010); Loh *et al.* (2009, 2010). For the stability of the temperature controller used in the data collection, see: Cosier & Glazer (1986). For bond-length data, see: Allen *et al.* (1987).



Experimental

Crystal data

$\text{C}_{10}\text{H}_6\text{N}_2$

$M_r = 154.17$

Orthorhombic, $P2_12_12_1$
 $a = 3.8497(2)\text{ \AA}$
 $b = 9.9559(4)\text{ \AA}$
 $c = 19.9639(13)\text{ \AA}$
 $V = 765.16(7)\text{ \AA}^3$

$Z = 4$
Mo $K\alpha$ radiation
 $\mu = 0.08\text{ mm}^{-1}$
 $T = 100\text{ K}$
 $0.36 \times 0.18 \times 0.03\text{ mm}$

Data collection

Bruker SMART APEXII CCD diffractometer
Absorption correction: multi-scan (*SADABS*; Bruker, 2009)
 $T_{\min} = 0.971$, $T_{\max} = 0.997$

4086 measured reflections
1056 independent reflections
838 reflections with $I > 2\sigma(I)$
 $R_{\text{int}} = 0.059$

Refinement

$R[F^2 > 2\sigma(F^2)] = 0.056$
 $wR(F^2) = 0.141$
 $S = 1.09$
1056 reflections

109 parameters
H-atom parameters constrained
 $\Delta\rho_{\max} = 0.31\text{ e \AA}^{-3}$
 $\Delta\rho_{\min} = -0.24\text{ e \AA}^{-3}$

Data collection: *APEX2* (Bruker, 2009); cell refinement: *SAINT* (Bruker, 2009); data reduction: *SAINT*; program(s) used to solve structure: *SHELXTL* (Sheldrick, 2008); program(s) used to refine structure: *SHELXTL*; molecular graphics: *SHELXTL*; software used to prepare material for publication: *SHELXTL* and *PLATON* (Spek, 2009).

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Supplementary data and figures for this paper are available from the IUCr electronic archives (Reference: HB5597).

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supplementary materials

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Quinoline-2-carbonitrile

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Comment

Heterocyclic molecules containing cyano group are useful as drug intermediates. Syntheses of the quinoline derivatives were discussed earlier (Sasaki *et al.*, 1998; Reux *et al.*, 2009). Recently, we have synthesized a number of quinoline compounds to investigate the hydrogen bonding patterns in these compounds (Loh *et al.*, 2009; 2010). Herein we report the crystal structure of quinoline-2-carbonitrile.

In the title compound, Fig. 1, the molecule is almost planar with an r.m.s. deviation of 0.014 Å. The dihedral angle between the benzene (C3–C8) and pyridine (C1–C3/N1/C8/C9) rings is 1.28 (16)°. The bond lengths (Allen *et al.*, 1987) and angles in the title compound are within normal ranges and comparable to the related structure of quinoxaline-2-carbonitrile (Fun *et al.*, 2010).

In the crystal packing, Fig. 2, the molecules are stacked along the a axis by way of weak aromatic π – π stacking interactions between the centroid of the benzene ($Cg1$) and the centroid of the pyridine ($Cg2$) rings of adjacent molecules [$Cg1\cdots Cg2$ separation = 3.7943 (19) Å]. There is no significant hydrogen bond observed in this compound.

Experimental

A hot methanol solution (20 ml) of quinoline-2-carbonitrile (39 mg, Aldrich) was warmed over a magnetic stirrer hotplate for a few minutes. The resulting solution was allowed to cool slowly to room temperature. Colourless plates of (I) appeared after a few days.

Refinement

All H atoms were positioned geometrically with the bond length of C–H being 0.93 Å and were refined using a riding model, with $U_{iso}(\text{H}) = 1.2 U_{eq}(\text{C})$. In the absence of significant anomalous dispersion, 616 Friedel pairs were merged for the final refinement.

Figures

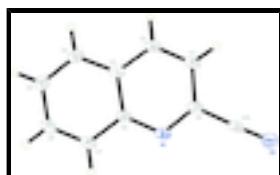


Fig. 1. The molecular structure of (I) showing 50% probability displacement ellipsoids for non-H atoms.

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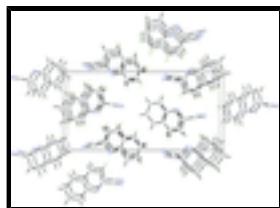


Fig. 2. The crystal structure of (I), viewed along the a axis.

Quinoline-2-carbonitrile

Crystal data

| | |
|---|---|
| C ₁₀ H ₆ N ₂ | $F(000) = 320$ |
| $M_r = 154.17$ | $D_x = 1.338 \text{ Mg m}^{-3}$ |
| Orthorhombic, $P2_12_12_1$ | Mo $K\alpha$ radiation, $\lambda = 0.71073 \text{ \AA}$ |
| Hall symbol: P 2ac 2ab | Cell parameters from 930 reflections |
| $a = 3.8497 (2) \text{ \AA}$ | $\theta = 3.7\text{--}27.4^\circ$ |
| $b = 9.9559 (4) \text{ \AA}$ | $\mu = 0.08 \text{ mm}^{-1}$ |
| $c = 19.9639 (13) \text{ \AA}$ | $T = 100 \text{ K}$ |
| $V = 765.16 (7) \text{ \AA}^3$ | Plate, colourless |
| $Z = 4$ | $0.36 \times 0.18 \times 0.03 \text{ mm}$ |

Data collection

| | |
|---|---|
| Bruker SMART APEXII CCD diffractometer | 1056 independent reflections |
| Radiation source: fine-focus sealed tube graphite | 838 reflections with $I > 2\sigma(I)$ |
| φ and ω scans | $R_{\text{int}} = 0.059$ |
| Absorption correction: multi-scan (<i>SADABS</i> ; Bruker, 2009) | $\theta_{\text{max}} = 27.5^\circ, \theta_{\text{min}} = 3.7^\circ$ |
| $T_{\text{min}} = 0.971, T_{\text{max}} = 0.997$ | $h = -4 \rightarrow 4$ |
| 4086 measured reflections | $k = -12 \rightarrow 12$ |
| | $l = -25 \rightarrow 21$ |

Refinement

| | |
|---------------------------------|---|
| Refinement on F^2 | Primary atom site location: structure-invariant direct methods |
| Least-squares matrix: full | Secondary atom site location: difference Fourier map |
| $R[F^2 > 2\sigma(F^2)] = 0.056$ | Hydrogen site location: inferred from neighbouring sites |
| $wR(F^2) = 0.141$ | H-atom parameters constrained |
| $S = 1.09$ | $w = 1/[\sigma^2(F_o^2) + (0.0827P)^2]$ where $P = (F_o^2 + 2F_c^2)/3$ |
| 1056 reflections | $(\Delta/\sigma)_{\text{max}} < 0.001$ |
| 109 parameters | $\Delta\rho_{\text{max}} = 0.31 \text{ e \AA}^{-3}$ |
| 0 restraints | $\Delta\rho_{\text{min}} = -0.24 \text{ e \AA}^{-3}$ |

Special details

Experimental. The crystal was placed in the cold stream of an Oxford Cryosystems Cobra open-flow nitrogen cryostat (Cosier & Glazer, 1986) operating at 100.0 (1) K.

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 $F^2 > \sigma(F^2)$ 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)

| | x | y | z | $U_{\text{iso}}^*/U_{\text{eq}}$ |
|-----|------------|------------|--------------|----------------------------------|
| N1 | 0.7386 (8) | 0.5083 (2) | 0.20498 (14) | 0.0185 (6) |
| N2 | 0.3930 (9) | 0.4374 (2) | 0.35822 (16) | 0.0296 (7) |
| C1 | 0.7983 (9) | 0.2683 (3) | 0.22614 (16) | 0.0193 (7) |
| H1A | 0.7552 | 0.1964 | 0.2547 | 0.023* |
| C2 | 0.9617 (9) | 0.2502 (3) | 0.16647 (16) | 0.0192 (7) |
| H2A | 1.0368 | 0.1649 | 0.1541 | 0.023* |
| C3 | 1.0178 (9) | 0.3604 (3) | 0.12314 (16) | 0.0174 (7) |
| C4 | 1.1815 (8) | 0.3487 (3) | 0.06064 (16) | 0.0201 (7) |
| H4A | 1.2662 | 0.2658 | 0.0468 | 0.024* |
| C5 | 1.2177 (9) | 0.4589 (3) | 0.01965 (18) | 0.0224 (7) |
| H5A | 1.3204 | 0.4498 | -0.0223 | 0.027* |
| C6 | 1.0989 (8) | 0.5855 (3) | 0.04129 (17) | 0.0239 (8) |
| H6A | 1.1257 | 0.6595 | 0.0133 | 0.029* |
| C7 | 0.9466 (9) | 0.6018 (3) | 0.10188 (17) | 0.0213 (7) |
| H7A | 0.8736 | 0.6866 | 0.1154 | 0.026* |
| C8 | 0.8981 (8) | 0.4894 (3) | 0.14492 (17) | 0.0162 (7) |
| C9 | 0.6962 (8) | 0.4002 (3) | 0.24322 (16) | 0.0169 (7) |
| C10 | 0.5263 (9) | 0.4229 (3) | 0.30714 (17) | 0.0196 (7) |

Atomic displacement parameters (\AA^2)

| | U^{11} | U^{22} | U^{33} | U^{12} | U^{13} | U^{23} |
|----|-------------|-------------|-------------|--------------|--------------|--------------|
| N1 | 0.0149 (13) | 0.0120 (10) | 0.0284 (15) | -0.0014 (10) | -0.0013 (13) | -0.0012 (10) |
| N2 | 0.0336 (17) | 0.0199 (13) | 0.0353 (17) | 0.0000 (12) | 0.0055 (16) | -0.0013 (13) |
| C1 | 0.0146 (16) | 0.0128 (12) | 0.0306 (18) | 0.0016 (12) | -0.0030 (15) | 0.0009 (12) |
| C2 | 0.0155 (16) | 0.0113 (12) | 0.0307 (18) | 0.0009 (12) | -0.0020 (15) | -0.0027 (12) |
| C3 | 0.0130 (15) | 0.0150 (13) | 0.0243 (17) | 0.0006 (11) | -0.0063 (14) | -0.0028 (12) |
| C4 | 0.0134 (16) | 0.0184 (14) | 0.0286 (19) | 0.0016 (12) | -0.0010 (15) | -0.0060 (13) |
| C5 | 0.0164 (16) | 0.0251 (15) | 0.0256 (17) | -0.0004 (14) | -0.0003 (15) | -0.0006 (13) |
| C6 | 0.0218 (17) | 0.0167 (13) | 0.0333 (19) | -0.0009 (13) | -0.0003 (16) | 0.0053 (14) |

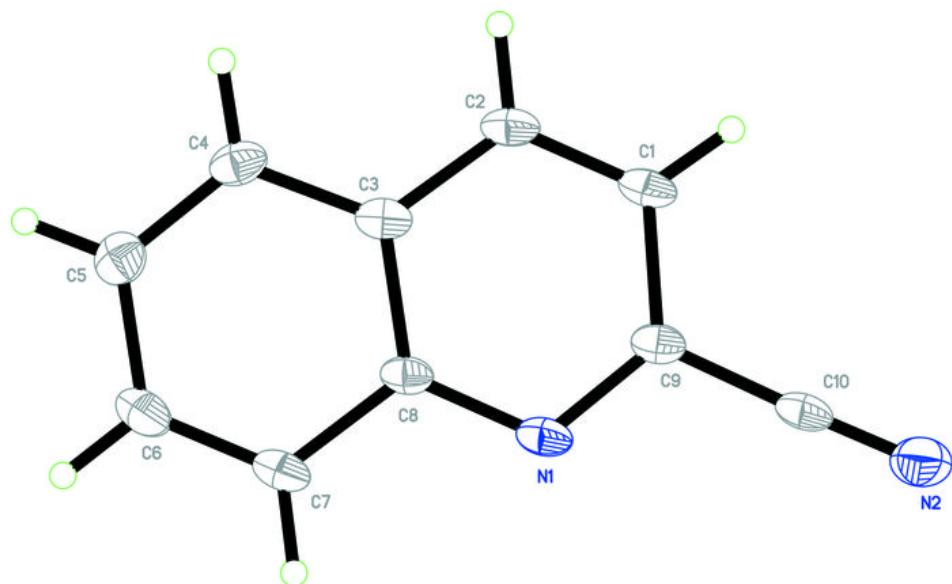
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|-----|-------------|-------------|-------------|--------------|--------------|--------------|
| C7 | 0.0202 (17) | 0.0128 (13) | 0.0311 (19) | -0.0003 (13) | -0.0039 (16) | 0.0021 (12) |
| C8 | 0.0124 (15) | 0.0111 (12) | 0.0249 (16) | -0.0016 (11) | -0.0027 (14) | -0.0033 (12) |
| C9 | 0.0130 (14) | 0.0130 (12) | 0.0247 (17) | -0.0019 (12) | -0.0035 (14) | -0.0018 (11) |
| C10 | 0.0186 (17) | 0.0098 (12) | 0.0306 (19) | -0.0019 (12) | -0.0015 (15) | 0.0008 (12) |

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

| | | | |
|-------------|------------|--------------|------------|
| N1—C9 | 1.329 (4) | C4—C5 | 1.376 (4) |
| N1—C8 | 1.360 (4) | C4—H4A | 0.9300 |
| N2—C10 | 1.151 (4) | C5—C6 | 1.409 (4) |
| C1—C2 | 1.359 (4) | C5—H5A | 0.9300 |
| C1—C9 | 1.413 (4) | C6—C7 | 1.354 (4) |
| C1—H1A | 0.9300 | C6—H6A | 0.9300 |
| C2—C3 | 1.414 (4) | C7—C8 | 1.424 (4) |
| C2—H2A | 0.9300 | C7—H7A | 0.9300 |
| C3—C4 | 1.402 (4) | C9—C10 | 1.452 (4) |
| C3—C8 | 1.432 (4) | | |
| C9—N1—C8 | 116.7 (2) | C6—C5—H5A | 120.1 |
| C2—C1—C9 | 117.6 (3) | C7—C6—C5 | 121.5 (3) |
| C2—C1—H1A | 121.2 | C7—C6—H6A | 119.3 |
| C9—C1—H1A | 121.2 | C5—C6—H6A | 119.3 |
| C1—C2—C3 | 120.3 (3) | C6—C7—C8 | 120.1 (3) |
| C1—C2—H2A | 119.9 | C6—C7—H7A | 120.0 |
| C3—C2—H2A | 119.9 | C8—C7—H7A | 120.0 |
| C4—C3—C2 | 123.3 (2) | N1—C8—C7 | 118.8 (2) |
| C4—C3—C8 | 119.3 (3) | N1—C8—C3 | 122.5 (3) |
| C2—C3—C8 | 117.4 (3) | C7—C8—C3 | 118.7 (3) |
| C5—C4—C3 | 120.6 (3) | N1—C9—C1 | 125.4 (3) |
| C5—C4—H4A | 119.7 | N1—C9—C10 | 115.7 (2) |
| C3—C4—H4A | 119.7 | C1—C9—C10 | 118.8 (3) |
| C4—C5—C6 | 119.9 (3) | N2—C10—C9 | 178.2 (3) |
| C4—C5—H5A | 120.1 | | |
| C9—C1—C2—C3 | -1.4 (4) | C6—C7—C8—N1 | -178.8 (3) |
| C1—C2—C3—C4 | -179.4 (3) | C6—C7—C8—C3 | 0.9 (5) |
| C1—C2—C3—C8 | 0.3 (4) | C4—C3—C8—N1 | -179.7 (3) |
| C2—C3—C4—C5 | 177.7 (3) | C2—C3—C8—N1 | 0.6 (4) |
| C8—C3—C4—C5 | -2.0 (4) | C4—C3—C8—C7 | 0.6 (4) |
| C3—C4—C5—C6 | 1.9 (5) | C2—C3—C8—C7 | -179.1 (3) |
| C4—C5—C6—C7 | -0.4 (5) | C8—N1—C9—C1 | -1.0 (5) |
| C5—C6—C7—C8 | -1.0 (5) | C8—N1—C9—C10 | 179.8 (3) |
| C9—N1—C8—C7 | 179.4 (3) | C2—C1—C9—N1 | 1.8 (5) |
| C9—N1—C8—C3 | -0.3 (4) | C2—C1—C9—C10 | -179.0 (3) |

Fig. 1



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Fig. 2

