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N-(Quinolin-8-yl)quinoline-2-carboxamide

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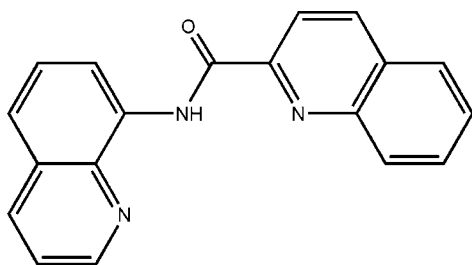
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Key indicators: single-crystal X-ray study; $T = 173$ K; mean $\sigma(\text{C}-\text{C}) = 0.003$ Å; R factor = 0.035; wR factor = 0.082; data-to-parameter ratio = 7.5.

In the title compound, $\text{C}_{19}\text{H}_{13}\text{N}_3\text{O}$, the dihedral angle between the two quinoline systems is $11.54(3)^\circ$. The molecular conformation is stabilized by intramolecular $\text{N}-\text{H}\cdots\text{N}$ and $\text{C}-\text{H}\cdots\text{O}$ hydrogen bonds, with $\text{N}-\text{H}\cdots\text{N}$ being bifurcated towards the two N atoms of the two quinoline rings. In the crystal, there are weak intermolecular $\pi-\pi$ interactions present involving the quinoline rings [centroid-centroid distance $3.7351(14)$ Å].

Related literature

For the synthesis of the title compound and related structures, see: Kim *et al.* (2009). For applications of the title compound and background to the synthesis, see: Wang *et al.* (2011).



Experimental

Crystal data

 $\text{C}_{19}\text{H}_{13}\text{N}_3\text{O}$ $M_r = 299.32$

Orthorhombic, $P2_12_12_1$
 $a = 6.3651(13)$ Å
 $b = 11.475(2)$ Å
 $c = 19.861(4)$ Å
 $V = 1450.6(5)$ Å³

$Z = 4$
Mo $K\alpha$ radiation
 $\mu = 0.09$ mm⁻¹
 $T = 173$ K
 $0.25 \times 0.15 \times 0.15$ mm

Data collection

Rigaku Saturn 724 CCD diffractometer
Absorption correction: multi-scan (ABSCOR; Higashi, 1995)
 $T_{\min} = 0.978$, $T_{\max} = 0.987$

6769 measured reflections
1553 independent reflections
1442 reflections with $I > 2\sigma(I)$
 $R_{\text{int}} = 0.036$

Refinement

$R[F^2 > 2\sigma(F^2)] = 0.035$
 $wR(F^2) = 0.082$
 $S = 1.06$
1553 reflections

208 parameters
H-atom parameters constrained
 $\Delta\rho_{\max} = 0.10$ e Å⁻³
 $\Delta\rho_{\min} = -0.13$ e Å⁻³

Table 1

Hydrogen-bond geometry (Å, °).

$D-H\cdots A$	$D-H$	$H\cdots A$	$D\cdots A$	$D-H\cdots A$
$\text{N2}-\text{H2A}\cdots\text{N1}$	0.88	2.27	2.693 (2)	109
$\text{N2}-\text{H2A}\cdots\text{N3}$	0.88	2.27	2.684 (2)	109
$\text{C12}-\text{H12}\cdots\text{O1}$	0.95	2.25	2.867 (2)	122

Data collection: *CrystalClear* (Rigaku, 2008); cell refinement: *CrystalClear*; data reduction: *CrystalClear*; program(s) used to solve structure: *SHELXS97* (Sheldrick, 2008); program(s) used to refine structure: *SHELXL97* (Sheldrick, 2008); molecular graphics: *DIAMOND* (Brandenburg, 2006); 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: ZL2476).

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supporting information

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***N*-(Quinolin-8-yl)quinoline-2-carboxamide**

Yanfeng Li, Hongbo Zhou and Xiaoping Shen

S1. Comment

The title compound (Hqcq) can act as a tridentate ligand, and has been incorporated into the cyanometalate building block $[\text{Fe}(\text{qcq})(\text{CN})_3]^-$ {qcq = 8-(2-quinolinecarboxamido)quinoline anion}, in which the Fe^{III} ion is coordinated by three carbon atoms of cyanide groups and three N-donors from the qcq ligand in a *mer*-arrangement (Kim *et al.*, 2009).

Through replacement of the cyanide ligands the $\text{Fe}(\text{qcq})$ fragment can coordinate to transition metal ions to form various polynuclear and one-dimensional structures with fascinating magnetic properties such as single molecular magnets and single-chain magnets (Kim *et al.*, 2009; Wang *et al.*, 2011). Herein, the crystal structure of the tridentate ligand of Hqcq is presented.

The molecular structure of the title compound is shown in Fig. 1. The quinoline rings are essentially planar, with a maximum deviation of 0.046 (1) Å for atom C8 in the (N1/C1-C9) ring and 0.016 (1) Å for atom C14 in the (N3/C11-C19) ring. The dihedral angle between the two quinoline rings is 11.54 (3)°. The amide (N2/C10/O1) plane forms dihedral angles of 14.1 (1)° and 4.2 (1)° with the quinoline rings of (N1/C1-C9) and (N3/C11-C19), respectively. The bond lengths of the title molecule are slightly different from those reported for $[\text{Fe}(\text{qcq})(\text{CN})_3]^-$ (Kim *et al.*, 2009), probably owing to the coordination effect to the tridentate ligand. There are intramolecular hydrogen-bonding interactions between the amido N atom and the N atoms of the quinoline rings, and between the O atom of amide group and the C atom of the quinoline ring. The amido N atom (N2) forms bifurcated hydrogen bonds towards the two N atoms (N1, N3) of the two quinoline rings (Table 1).

In the crystal structure, no significant intermolecular hydrogen bonds are observed. The crystal structure features intermolecular π - π interactions between different types of quinoline rings with a distance of *ca.* 3.735 Å between the centroids of the respective rings (Fig. 2), and the adjacent rings tilted against each other.

S2. Experimental

The compound of 8-(2-quinolinecarboxamido)quinoline (Hqcq) was prepared according to a literature method (Kim *et al.*, 2009). Then, 0.3 mmol of Hqcq was added to MeCN (20 mL) with stirring. The resulting solution was filtered and the filtrate was left for slow evaporation in the dark at room temperature. Yellow block-shaped crystals of the title compound suitable for single-crystal X-ray diffraction were obtained after two weeks. Melting point = 429.6–430.5 K. IR (KBr, cm^{-1}): 3314(s), 3044(m), 1678(vs), 1523(vs), 1488(s), 1427(s), 1325(s), 1126(m), 913(s), 834(s), 764(vs), 611(m), 588(m).

S3. Refinement

All non-H atoms were refined with anisotropic thermal parameters. The C- and N-bound H atoms were calculated in idealized positions and included in the refinement in a riding mode (C-H = 0.95 Å, N-H = 0.88 Å) with U_{iso} for H assigned as 1.2 times U_{eq} of the attached atoms. In the absence of atoms heavier than Si and with Mo $K\alpha$ radiation used

Friedel-pair reflections have been merged (using a MERG 3 command) during the refinement. Assignment of the absolute structure is arbitrary.

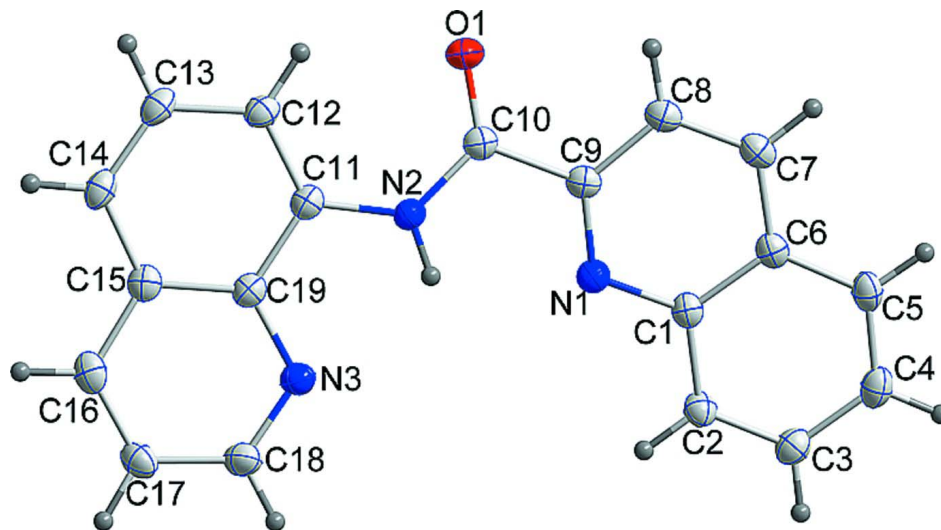


Figure 1

ORTEP diagram of the title compound with displacement ellipsoids drawn at the 30% probability level.

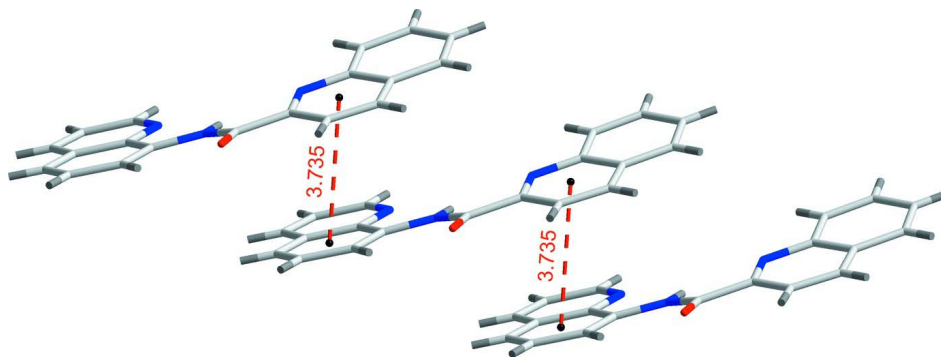


Figure 2

The crystal packing diagram of the title compound showing the intermolecular π - π interactions. The distances shown are between the centroids of the respective rings, and the symmetry operator codes for generating adjacent aromatic rings are $-1+x, y, z$ and $1+x, y, z$.

N-(Quinolin-8-yl)quinoline-2-carboxamide

Crystal data

$C_{19}H_{13}N_3O$

$M_r = 299.32$

Orthorhombic, $P2_12_12_1$

Hall symbol: P2ac2ab

$a = 6.3651(13) \text{ \AA}$

$b = 11.475(2) \text{ \AA}$

$c = 19.861(4) \text{ \AA}$

$V = 1450.6(5) \text{ \AA}^3$

$Z = 4$

$F(000) = 624$

$D_x = 1.371 \text{ Mg m}^{-3}$

Mo $K\alpha$ radiation, $\lambda = 0.71073 \text{ \AA}$

Cell parameters from 6054 reflections

$\theta = 3.4\text{--}29.0^\circ$

$\mu = 0.09 \text{ mm}^{-1}$

$T = 173 \text{ K}$

Block, yellow

$0.25 \times 0.15 \times 0.15 \text{ mm}$

Data collection

Rigaku Saturn 724 CCD
diffractometer
Radiation source: Rotating Anode
Graphite monochromator
 φ and ω scans
Absorption correction: multi-scan
(*ABSCOR*; Higashi, 1995)
 $T_{\min} = 0.978$, $T_{\max} = 0.987$

6769 measured reflections
1553 independent reflections
1442 reflections with $I > 2\sigma(I)$
 $R_{\text{int}} = 0.036$
 $\theta_{\max} = 25.3^\circ$, $\theta_{\min} = 3.4^\circ$
 $h = -7 \rightarrow 6$
 $k = -13 \rightarrow 13$
 $l = -23 \rightarrow 18$

Refinement

Refinement on F^2
Least-squares matrix: full
 $R[F^2 > 2\sigma(F^2)] = 0.035$
 $wR(F^2) = 0.082$
 $S = 1.06$
1553 reflections
208 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
 $w = 1/[\sigma^2(F_o^2) + (0.0494P)^2]$
where $P = (F_o^2 + 2F_c^2)/3$
 $(\Delta/\sigma)_{\max} < 0.001$
 $\Delta\rho_{\max} = 0.10 \text{ e } \text{\AA}^{-3}$
 $\Delta\rho_{\min} = -0.13 \text{ e } \text{\AA}^{-3}$

Special details

Geometry. All esds (except the esd in the dihedral angle between two l.s. planes) are estimated using the full covariance matrix. The cell esds are taken into account individually in the estimation of esds in distances, angles and torsion angles; correlations between esds in cell parameters are only used when they are defined by crystal symmetry. An approximate (isotropic) treatment of cell esds is used for estimating esds 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 $F^2 > 2\sigma(F^2)$ 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>	$U_{\text{iso}}^*/U_{\text{eq}}$
O1	-0.0149 (2)	0.41328 (13)	0.73413 (7)	0.0458 (4)
N1	-0.1752 (2)	0.46907 (14)	0.56729 (8)	0.0328 (4)
C1	-0.3327 (3)	0.44137 (16)	0.52317 (10)	0.0309 (4)
N2	0.1160 (2)	0.54135 (14)	0.65635 (8)	0.0336 (4)
H2A	0.0898	0.5707	0.6163	0.040*
C2	-0.3099 (3)	0.47410 (18)	0.45477 (10)	0.0367 (5)
H2	-0.1874	0.5146	0.4407	0.044*
N3	0.3570 (3)	0.69181 (14)	0.58750 (8)	0.0385 (4)
C3	-0.4630 (3)	0.44780 (19)	0.40896 (10)	0.0405 (5)
H3	-0.4453	0.4691	0.3631	0.049*
C4	-0.6456 (3)	0.38961 (18)	0.42926 (11)	0.0416 (5)
H4	-0.7514	0.3724	0.3970	0.050*
C5	-0.6737 (3)	0.35718 (18)	0.49498 (10)	0.0390 (5)
H5	-0.7993	0.3187	0.5082	0.047*
C6	-0.5164 (3)	0.38082 (16)	0.54318 (10)	0.0324 (5)
C7	-0.5302 (3)	0.34377 (17)	0.61049 (10)	0.0364 (5)

H7	-0.6505	0.3024	0.6257	0.044*
C8	-0.3693 (3)	0.36763 (17)	0.65394 (10)	0.0358 (5)
H8	-0.3739	0.3413	0.6993	0.043*
C9	-0.1960 (3)	0.43222 (16)	0.62995 (10)	0.0317 (5)
C10	-0.0232 (3)	0.46113 (17)	0.67895 (10)	0.0329 (4)
C11	0.2965 (3)	0.58357 (17)	0.68860 (10)	0.0322 (5)
C12	0.3571 (3)	0.55062 (19)	0.75238 (9)	0.0362 (5)
H12	0.2716	0.4994	0.7781	0.043*
C13	0.5465 (3)	0.59343 (19)	0.77909 (10)	0.0409 (6)
H13	0.5880	0.5697	0.8229	0.049*
C14	0.6721 (3)	0.66767 (18)	0.74419 (10)	0.0388 (5)
H14	0.7999	0.6945	0.7635	0.047*
C15	0.6127 (3)	0.70472 (17)	0.67950 (10)	0.0341 (5)
C16	0.7326 (3)	0.78261 (18)	0.64020 (11)	0.0404 (5)
H16	0.8604	0.8134	0.6572	0.049*
C17	0.6643 (4)	0.81326 (18)	0.57795 (11)	0.0423 (5)
H17	0.7423	0.8667	0.5513	0.051*
C18	0.4767 (4)	0.76476 (18)	0.55348 (10)	0.0436 (6)
H18	0.4330	0.7860	0.5094	0.052*
C19	0.4227 (3)	0.66226 (17)	0.65098 (10)	0.0315 (4)

Atomic displacement parameters (Å²)

	U^{11}	U^{22}	U^{33}	U^{12}	U^{13}	U^{23}
O1	0.0481 (9)	0.0504 (9)	0.0389 (9)	-0.0030 (8)	-0.0031 (8)	0.0133 (7)
N1	0.0336 (9)	0.0318 (9)	0.0328 (9)	0.0016 (8)	0.0003 (8)	-0.0013 (7)
C1	0.0303 (10)	0.0286 (10)	0.0337 (10)	0.0012 (9)	0.0019 (9)	-0.0059 (8)
N2	0.0334 (9)	0.0372 (9)	0.0301 (8)	-0.0003 (8)	-0.0032 (8)	0.0028 (7)
C2	0.0339 (11)	0.0405 (12)	0.0356 (11)	-0.0021 (11)	0.0030 (9)	-0.0031 (9)
N3	0.0466 (10)	0.0354 (9)	0.0335 (9)	-0.0013 (9)	-0.0026 (9)	0.0011 (7)
C3	0.0426 (12)	0.0425 (12)	0.0366 (11)	0.0039 (11)	-0.0025 (10)	-0.0047 (10)
C4	0.0383 (12)	0.0374 (12)	0.0492 (14)	0.0031 (11)	-0.0089 (11)	-0.0080 (10)
C5	0.0342 (11)	0.0317 (11)	0.0512 (14)	-0.0025 (11)	-0.0032 (10)	-0.0044 (10)
C6	0.0317 (11)	0.0257 (10)	0.0397 (11)	0.0023 (9)	0.0021 (9)	-0.0040 (8)
C7	0.0357 (11)	0.0287 (10)	0.0448 (12)	0.0019 (10)	0.0075 (10)	-0.0004 (9)
C8	0.0393 (11)	0.0314 (10)	0.0367 (11)	0.0020 (10)	0.0046 (10)	0.0003 (9)
C9	0.0343 (11)	0.0276 (10)	0.0332 (11)	0.0046 (9)	0.0032 (9)	-0.0015 (8)
C10	0.0328 (11)	0.0327 (10)	0.0332 (10)	0.0056 (10)	0.0031 (9)	-0.0010 (9)
C11	0.0324 (11)	0.0337 (10)	0.0307 (11)	0.0043 (10)	-0.0002 (9)	-0.0053 (8)
C12	0.0385 (11)	0.0417 (12)	0.0283 (10)	0.0031 (11)	0.0018 (10)	-0.0015 (9)
C13	0.0418 (12)	0.0510 (14)	0.0300 (11)	0.0074 (12)	-0.0052 (10)	-0.0054 (9)
C14	0.0352 (11)	0.0462 (13)	0.0350 (12)	0.0071 (11)	-0.0055 (10)	-0.0104 (10)
C15	0.0360 (11)	0.0332 (10)	0.0330 (10)	0.0030 (10)	0.0024 (9)	-0.0092 (9)
C16	0.0402 (12)	0.0358 (11)	0.0454 (13)	-0.0030 (11)	0.0036 (10)	-0.0139 (10)
C17	0.0496 (14)	0.0350 (11)	0.0423 (13)	-0.0087 (11)	0.0051 (11)	-0.0044 (9)
C18	0.0563 (14)	0.0372 (12)	0.0374 (11)	-0.0050 (12)	-0.0009 (11)	0.0031 (9)
C19	0.0336 (10)	0.0313 (10)	0.0295 (10)	0.0053 (9)	0.0008 (9)	-0.0052 (8)

Geometric parameters (Å, °)

O1—C10	1.227 (2)	C7—H7	0.9500
N1—C9	1.321 (2)	C8—C9	1.412 (3)
N1—C1	1.369 (2)	C8—H8	0.9500
C1—C2	1.417 (3)	C9—C10	1.505 (3)
C1—C6	1.417 (3)	C11—C12	1.377 (3)
N2—C10	1.354 (3)	C11—C19	1.421 (3)
N2—C11	1.402 (2)	C12—C13	1.406 (3)
N2—H2A	0.8800	C12—H12	0.9500
C2—C3	1.367 (3)	C13—C14	1.358 (3)
C2—H2	0.9500	C13—H13	0.9500
N3—C18	1.318 (3)	C14—C15	1.405 (3)
N3—C19	1.371 (2)	C14—H14	0.9500
C3—C4	1.400 (3)	C15—C16	1.411 (3)
C3—H3	0.9500	C15—C19	1.421 (3)
C4—C5	1.369 (3)	C16—C17	1.357 (3)
C4—H4	0.9500	C16—H16	0.9500
C5—C6	1.412 (3)	C17—C18	1.405 (3)
C5—H5	0.9500	C17—H17	0.9500
C6—C7	1.405 (3)	C18—H18	0.9500
C7—C8	1.367 (3)		
C9—N1—C1	117.09 (17)	C8—C9—C10	117.92 (17)
N1—C1—C2	118.50 (18)	O1—C10—N2	124.89 (19)
N1—C1—C6	122.59 (17)	O1—C10—C9	120.68 (19)
C2—C1—C6	118.90 (18)	N2—C10—C9	114.42 (16)
C10—N2—C11	128.30 (17)	C12—C11—N2	123.71 (19)
C10—N2—H2A	115.8	C12—C11—C19	120.01 (19)
C11—N2—H2A	115.8	N2—C11—C19	116.26 (17)
C3—C2—C1	120.5 (2)	C11—C12—C13	119.4 (2)
C3—C2—H2	119.8	C11—C12—H12	120.3
C1—C2—H2	119.8	C13—C12—H12	120.3
C18—N3—C19	116.89 (19)	C14—C13—C12	122.1 (2)
C2—C3—C4	120.37 (19)	C14—C13—H13	118.9
C2—C3—H3	119.8	C12—C13—H13	118.9
C4—C3—H3	119.8	C13—C14—C15	119.8 (2)
C5—C4—C3	120.8 (2)	C13—C14—H14	120.1
C5—C4—H4	119.6	C15—C14—H14	120.1
C3—C4—H4	119.6	C14—C15—C16	123.5 (2)
C4—C5—C6	120.1 (2)	C14—C15—C19	119.3 (2)
C4—C5—H5	119.9	C16—C15—C19	117.19 (18)
C6—C5—H5	119.9	C17—C16—C15	119.7 (2)
C7—C6—C5	122.86 (19)	C17—C16—H16	120.2
C7—C6—C1	117.81 (18)	C15—C16—H16	120.2
C5—C6—C1	119.30 (18)	C16—C17—C18	119.0 (2)
C8—C7—C6	119.55 (19)	C16—C17—H17	120.5
C8—C7—H7	120.2	C18—C17—H17	120.5

C6—C7—H7	120.2	N3—C18—C17	124.4 (2)
C7—C8—C9	118.52 (19)	N3—C18—H18	117.8
C7—C8—H8	120.7	C17—C18—H18	117.8
C9—C8—H8	120.7	N3—C19—C11	117.95 (18)
N1—C9—C8	124.35 (19)	N3—C19—C15	122.76 (19)
N1—C9—C10	117.73 (18)	C11—C19—C15	119.28 (18)

Hydrogen-bond geometry (Å, °)

<i>D—H...A</i>	<i>D—H</i>	<i>H...A</i>	<i>D...A</i>	<i>D—H...A</i>
N2—H2A...N1	0.88	2.27	2.693 (2)	109
N2—H2A...N3	0.88	2.27	2.684 (2)	109
C12—H12...O1	0.95	2.25	2.867 (2)	122