

(R)-N-(1-Phenylethyl)quinoline-2-carboxamide

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Key indicators

Single-crystal X-ray study

$T = 294$ K

Mean $\sigma(\text{C}-\text{C}) = 0.003$ Å

R factor = 0.038

wR factor = 0.090

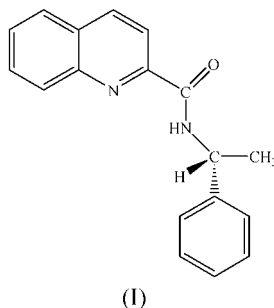
Data-to-parameter ratio = 12.5

For details of how these key indicators were automatically derived from the article, see <http://journals.iucr.org/e>.

The quinolyl and phenyl rings in the title compound, $\text{C}_{18}\text{H}_{16}\text{N}_2\text{O}$, form a dihedral angle of $89.07(5)^\circ$. The amide group is rotated out of the quinoline ring plane, with a dihedral angle of $21.19(1)^\circ$.

Comment

Recently, we have reported the structure of the tridentate ligand *N,N'*-di-2-naphthylpyridine-2,6-dicarboxamide (Qi *et al.*, 2001). Because of the steric effect, the bulky naphthyl ring will hinder coordination of the ligand to a metal ion, so a new chiral bidentate ligand, (I), containing the quinoline-2-carboxamide moiety was synthesized. It is expected that the N atom of the quinoline ring and the amide N atom will coordinate to a metal ion and form a complex with a five-membered ring structure. The quinolyl and phenyl rings in (I) (Fig. 1) are not coplanar, and form a dihedral angle of $89.07(5)^\circ$. The amide group is rotated out of the quinoline ring plane, forming a dihedral angle of $21.19(1)^\circ$. The title compound could have practical applications as a new chiral ligand (Noyori, 1989).

**Experimental**

The title compound was synthesized from 2-quinolinecarboxylic acid and (*D*)- α -methylbenzylamine according to the general procedure of Johnson *et al.* (1960). Therefore, the absolute configuration of the chiral centre was known in advance as *R*. The crystal used for the data collection was obtained by slow evaporation from a saturated DMF-water (10:1) solution at room temperature.

Crystal data

$\text{C}_{18}\text{H}_{16}\text{N}_2\text{O}$
 $M_r = 276.33$
 Monoclinic, $P2_1$
 $a = 5.5955(11)$ Å
 $b = 10.684(2)$ Å
 $c = 12.224(2)$ Å
 $\beta = 98.687(4)^\circ$
 $V = 722.4(2)$ Å³
 $Z = 2$

$D_x = 1.270$ Mg m⁻³
 Mo $K\alpha$ radiation
 Cell parameters from 2871 reflections
 $\theta = 1-27.5^\circ$
 $\mu = 0.08$ mm⁻¹
 $T = 294(2)$ K
 Plate, colourless
 $0.28 \times 0.20 \times 0.10$ mm

Data collection

Bruker CCD area-detector
diffractometer
 φ and ω scans
Absorption correction: empirical
(*SADABS*; Sheldrick, 1996)
 $T_{\min} = 0.978$, $T_{\max} = 0.992$
4900 measured reflections

2371 independent reflections
1768 reflections with $I > 2\sigma(I)$
 $R_{\text{int}} = 0.023$
 $\theta_{\text{max}} = 27.5^\circ$
 $h = -7 \rightarrow 7$
 $k = -13 \rightarrow 9$
 $l = -15 \rightarrow 15$

Refinement

Refinement on F^2
 $R[F^2 > 2\sigma(F^2)] = 0.038$
 $wR(F^2) = 0.090$
 $S = 0.96$
2371 reflections
190 parameters

H-atom parameters constrained
 $w = 1/[\sigma^2(F_o^2) + (0.1P)^2]$
where $P = (F_o^2 + 2F_c^2)/3$
 $(\Delta/\sigma)_{\text{max}} < 0.001$
 $\Delta\rho_{\text{max}} = 0.11 \text{ e } \text{\AA}^{-3}$
 $\Delta\rho_{\text{min}} = -0.12 \text{ e } \text{\AA}^{-3}$

Table 1

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

<i>D</i> —H... <i>A</i>	<i>D</i> —H	H... <i>A</i>	<i>D</i> ... <i>A</i>	<i>D</i> —H... <i>A</i>
N1—H1A...N2	0.86	2.41	2.754 (2)	104

The C-bound H atoms were placed at geometrically calculated positions and included in the final refinement using the riding-model approximation.

Data collection: *SMART* (Siemens, 1995); cell refinement: *SMART*; data reduction: *SHELXTL-NT* (Siemens, 1995); program(s) used to solve structure: *SHELXS97* (Sheldrick, 1990); program(s) used to refine structure: *SHELXL97* (Sheldrick, 1997); molecular graphics: *SHELXTL-NT*; software used to prepare material for publication: *SHELXTL-NT*.

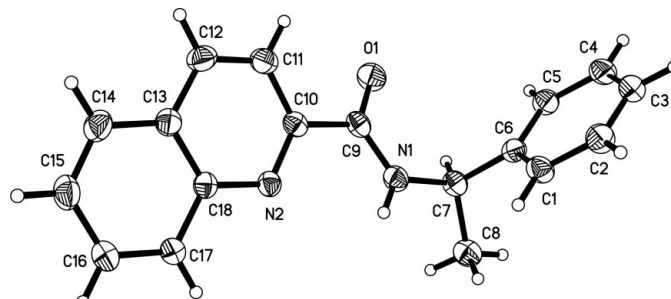


Figure 1
The molecular structure of (I) showing ellipsoids plotted at the 30% probability level (Siemens, 1995).

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References

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