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Key indicators

Single-crystal X-ray study
 T = 100 K
 Mean $\sigma(\text{C}-\text{C}) = 0.002 \text{ \AA}$
 R factor = 0.044
 wR factor = 0.112
 Data-to-parameter ratio = 17.1

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

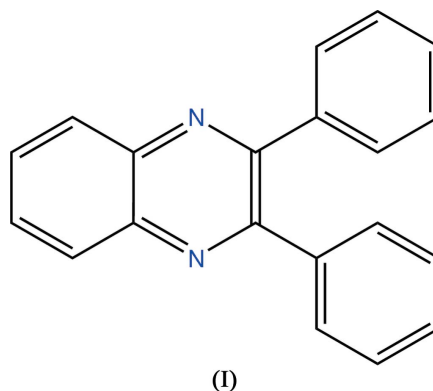
2,3-Diphenylquinoxaline

In the title compound, $\text{C}_{20}\text{H}_{14}\text{N}_2$, the presence of adjacent phenyl rings causes an out-of-plane twist in the quinoxaline system of approximately 12° .

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Comment

Compounds containing a quinoxaline group are of considerable interest due to the potential use of their metal complexes as molecular devices and DNA probes (Holmlin & Barton, 1995; Balzani, Campagna *et al.*, 1998; Balzani, Gomez-Loez & Stoddart, 1998; Du & Zhao, 2003).



In (I), the quinoxaline unit makes a dihedral angle of $37.1 (1)^\circ$ with the C11–C16 phenyl ring and $53.3 (1)^\circ$ with the C17–C22 phenyl ring. The presence of the two adjacent phenyl rings causes an out-of-plane twist in the quinoxaline system [$\text{C}17-\text{C}2-\text{C}3-\text{C}11 = 12.0 (2)^\circ$].

Experimental

Benzil (0.50 mg, 2.38 mmol), 1,2-diaminobenzene (0.26 mg, 2.40 mmol), molecular sieves (1 g) and manganese oxide (0.10 mg) as a catalyst were heated in a glass vial using domestic microwave oven. After 1 min the vial was cooled to room temperature and the solution was filtered. The product was recrystallized from ethanol to give (I) (yield 0.570 mg, 76.49%).

Crystal data

$\text{C}_{20}\text{H}_{14}\text{N}_2$	$Z = 4$
$M_r = 282.33$	$D_x = 1.268 \text{ Mg m}^{-3}$
Monoclinic, $P2_1/n$	Mo $K\alpha$ radiation
$a = 6.0306 (2) \text{ \AA}$	$\mu = 0.08 \text{ mm}^{-1}$
$b = 10.9269 (5) \text{ \AA}$	$T = 100 (2) \text{ K}$
$c = 22.5309 (8) \text{ \AA}$	Prism, light yellow
$\beta = 95.005 (3)^\circ$	$0.5 \times 0.45 \times 0.3 \text{ mm}$
$V = 1479.03 (10) \text{ \AA}^3$	

Data collection

Oxford Diffraction CrysAlisCCD
diffractometer
 ω - 2θ scans
Absorption correction: multi-scan
(SADABS; Sheldrick, 2003)
 $T_{\min} = 0.93$, $T_{\max} = 0.98$

16180 measured reflections
3413 independent reflections
2773 reflections with $I > 2\sigma(I)$
 $R_{\text{int}} = 0.027$
 $\theta_{\text{max}} = 27.6^\circ$

Refinement

Refinement on F^2
 $R[F^2 > 2\sigma(F^2)] = 0.044$
 $wR(F^2) = 0.112$
 $S = 1.13$
3413 reflections
200 parameters
H-atom parameters constrained

$w = 1/[\sigma^2(F_o^2) + (0.0404P)^2 + 0.715P]$
where $P = (F_o^2 + 2F_c^2)/3$
 $(\Delta/\sigma)_{\text{max}} < 0.001$
 $\Delta\rho_{\text{max}} = 0.30 \text{ e } \text{Å}^{-3}$
 $\Delta\rho_{\text{min}} = -0.17 \text{ e } \text{Å}^{-3}$
Extinction correction: SHELXL97
Extinction coefficient: 0.0092 (16)

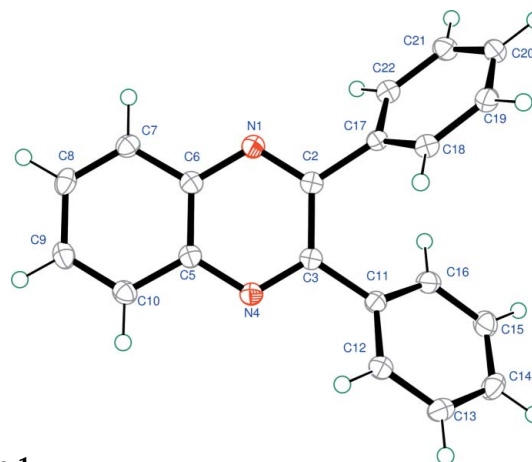


Figure 1

The molecular structure, with displacement ellipsoids drawn at the 50% probability level.

Table 1

Selected geometric parameters (Å, °).

N1—C2	1.3228 (18)	C3—N4	1.3213 (19)
N1—C6	1.3678 (19)	N4—C5	1.3683 (19)
C3—C2—C17	123.34 (13)	C2—C3—C11	122.99 (12)

All H atoms were positioned geometrically ($C-H = 0.95 \text{ Å}$) refined as riding on their parent atoms, with $U_{\text{iso}}(\text{H}) = 1.2U_{\text{eq}}(\text{parent})$.

Data collection: *CrysAlis CCD* (Oxford Diffraction, 2003); cell refinement: *CrysAlis RED* (Oxford Diffraction, 2003); data reduction: *CrysAlis RED*; program(s) used to solve structure: *SHELXS97* (Sheldrick, 1997); program(s) used to refine structure: *SHELXL97* (Sheldrick, 1997); molecular graphics: *ORTEP-3 for Windows* (Farrugia, 1997); software used to prepare material for publication: *PARST95* (Nardelli, 1995) and *PLATON* (Spek, 2003).

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