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#### **Key indicators**

Single-crystal X-ray study T = 298 KMean  $\sigma(\text{C}-\text{C}) = 0.002 \text{ Å}$  R factor = 0.052 wR factor = 0.136 Data-to-parameter ratio = 17.4

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

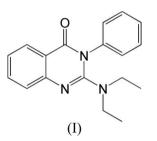
# 2-Diethylamino-3-phenylquinazolin-4(3H)-one

In the title compound,  $C_{18}H_{19}N_3O$ , the quinazoline system is approximately planar. The plane of the substituent phenyl ring forms a dihedral angle of 63.9 (2)° with respect to the mean plane of the quinazolinone ring system.

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## Comment

Quinazolinones are important heterocyclic compounds which exhibit good biological and pharmaceutical activity, including anti-inflammatory, antifungal, anticancer and AMPA-receptor antagonistic properties (Shiba *et al.*, 1997). As part of our work on the preparation of potentially active heterocycles (Ding *et al.*, 2000), we have obtained the title compound, (I).

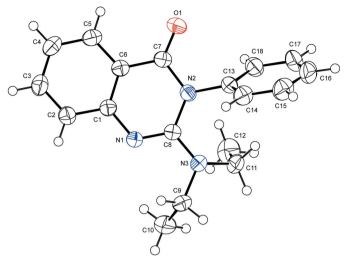


In (I), the quinazoline system is approximately planar (Fig. 1). The plane of the phenyl ring bound to N2 forms a dihedral angle of  $63.9 (2)^{\circ}$  with respect to the mean plane of the quinazolinone ring system, and the N2–C13 bond is twisted out of the quinazolinone plane [C14–C13–N2–C8 torsion angle =  $-60.28 (19)^{\circ}$ ]. The C8–N3, C8–N2, C13–N2 and C7–N2 bond lengths [1.3677 (18), 1.4072 (17), 1.4499 (18) and 1.4139 (19) Å, respectively] indicate some degree of delocalization around the heterocyclic sixmembered ring (Yuan & Ding, 2006).

## **Experimental**

Phenyl isocyanate (0.36 g, 3 mmol) was added to a solution of *N*-(2ethoxycarbonylphenyl)iminophosphorane (1.45 g, 3 mmol) in anhydrous dichloromethane (15 ml) under dry nitrogen at room temperature. The reaction mixture was left unstirred for 8 h at room temperature and the solvent was then removed under reduced pressure and diethyl ether–petroleum ether (1:2  $\nu/\nu$ , 20 ml) was added to precipitate triphenylphosphine oxide. After filtration, a solution of the carbodiimide was added to a solution of diethylamine in anhydrous dichloromethane (5 ml). After stirring the reaction mixture for 8 h, the solvent was removed under reduced pressure and the residue was recrystallized from ethanol to give the title compound (yield 88%; m.p. 388 K). Single crystals suitable for X-ray diffraction were obtained by recrystallization from a hexane–dichloromethane (1:3  $\nu/\nu$ ) solution at room temperature.

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#### Figure 1

The molecular structure of (I), showing displacement ellipsoids drawn at the 50% probability level. H atoms are shown as spheres of arbitrary radius.

#### Crystal data

 $\begin{array}{l} C_{18}H_{19}N_{3}O\\ M_{r}=293.36\\ Orthorhombic, Pbca\\ a=8.4259 (7) ~\text{\AA}\\ b=10.9029 (9) ~\text{\AA}\\ c=33.263 (3) ~\text{\AA}\\ V=3055.7 (4) ~\text{\AA}^{3} \end{array}$ 

#### Data collection

Bruker SMART CCD diffractometer  $\varphi$  and  $\omega$  scans Absorption correction: multi-scan (*SADABS*; Sheldrick, 1996)  $T_{\min} = 0.976, T_{\max} = 0.992$  Z = 8  $D_x = 1.275 \text{ Mg m}^{-3}$ Mo K $\alpha$  radiation  $\mu = 0.08 \text{ mm}^{-1}$ T = 298 (2) K Block, white  $0.30 \times 0.20 \times 0.10 \text{ mm}$ 

17133 measured reflections 3493 independent reflections 2931 reflections with  $I > 2\sigma(I)$  $R_{\text{int}} = 0.029$  $\theta_{\text{max}} = 27.5^{\circ}$  Refinement

Refinement on  $F^2$   $R[F^2 > 2\sigma(F^2)] = 0.052$   $wR(F^2) = 0.136$  S = 1.053493 reflections 201 parameters H-atom parameters constrained 
$$\begin{split} w &= 1/[\sigma^2(F_o^2) + (0.0701P)^2 \\ &+ 0.7969P] \\ \text{where } P &= (F_o^2 + 2F_c^2)/3 \\ (\Delta/\sigma)_{\text{max}} &= 0.001 \\ \Delta\rho_{\text{max}} &= 0.27 \text{ e } \text{ \AA}^{-3} \\ \Delta\rho_{\text{min}} &= -0.22 \text{ e } \text{ \AA}^{-3} \end{split}$$

H atoms were placed at calculated positions, with C–H = 0.93, 0.96 and 0.97 Å for aromatic, methylene and methyl H atoms, respectively. They were refined using a riding model, with  $U_{\rm iso}({\rm H}) = 1.2U_{\rm eq}({\rm C})$ , or  $1.5U_{\rm eq}({\rm C})$  for the methyl groups. The methyl groups were allowed to rotate about their local threefold axes.

Data collection: *SMART* (Bruker, 1997); cell refinement: *SAINT* (Bruker, 1997); data reduction: *SAINT*; program(s) used to solve structure: *SHELXS97* (Sheldrick, 1997); program(s) used to refine structure: *SHELXL97* (Sheldrick, 1997); molecular graphics: *SHELXTL* (Sheldrick, 2001); software used to prepare material for publication: *SHELXTL*.

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