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Key indicators

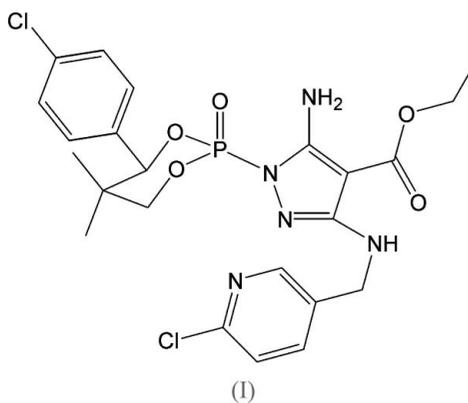
Single-crystal X-ray study
 $T = 292\text{ K}$
Mean $\sigma(\text{C}-\text{C}) = 0.004\text{ \AA}$
 R factor = 0.050
 wR factor = 0.140
Data-to-parameter ratio = 13.6For details of how these key indicators were automatically derived from the article, see <http://journals.iucr.org/e>.Ethyl 5-amino-1-(4-chlorophenyl-5,5-dimethyl-2-oxo-1,3,2-dioxaphosphinan-2-yl)-3-(6-chloro-3-pyridylmethylamino)-1*H*-pyrazole-4-carboxylate

In the title compound, $\text{C}_{23}\text{H}_{26}\text{Cl}_2\text{N}_5\text{O}_5\text{P}$, the P atom adopts a distorted tetrahedral coordination. The terminal ethyl carboxylate group is almost coplanar with the plane of the pyrazole ring. Intramolecular $\text{N}-\text{H}\cdots\text{O}$ hydrogen bonds influence the overall conformation of the molecule. $\text{C}-\text{H}\cdots\pi$ interactions contribute to the crystal packing.

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Comment

Neonicotinoide insecticides as nicotinic acetylcholine receptor inhibitors have attracted increasing attention because of their safety, low toxicity, wide range of activities and high potency (Shiokawa *et al.*, 1986). It has been found that most biologically active nicotinic compounds contain the 3-aminomethylpyridine group (Yamamoto *et al.*, 1994). Pyrazole and phosphorus heterocyclic compounds appear to be very important due to their wide range of biological activities (Tomcufcik *et al.*, 1985; Hirashima *et al.*, 1986).



We report here the crystal structure of the title compound, (I) (Fig. 1), which was synthesized by attachment of a 1,3,2-dioxaphosphinane and a 3-aminomethylpyridine group to pyrazole. The distorted tetrahedral configuration of the P atom can be attributed to the presence of the dioxaphosphinane ring whose sterical and electronic properties influence the coordination. The values of the double $\text{P1}=\text{O3}$ and single $\text{P1}-\text{N1}$ bonds and the angles $\text{O}-\text{P}-\text{O}'$ and $\text{O}-\text{P}-\text{N}$ illustrate the irregularities (Table 1). The dioxaphosphinane ring adopts a distorted chair conformation, with the parameters $Q = 0.5186(28)\text{ \AA}$, $\theta = 151.45(26)^\circ$ and $\varphi = 11.2(6)^\circ$ (Cremer & Pople, 1975). Atoms C11, O1, C7 and O2 of the dioxaphosphinane ring are not coplanar; the average deviation is $0.0245(3)\text{ \AA}$. The $\text{P}-\text{O}$ single and double bonds are shorter (Table 1) than the analogous chemical bonds observed previously [$1.586(2)$, $1.572(2)$ and $1.468(2)\text{ \AA}$; Liu *et al.*,

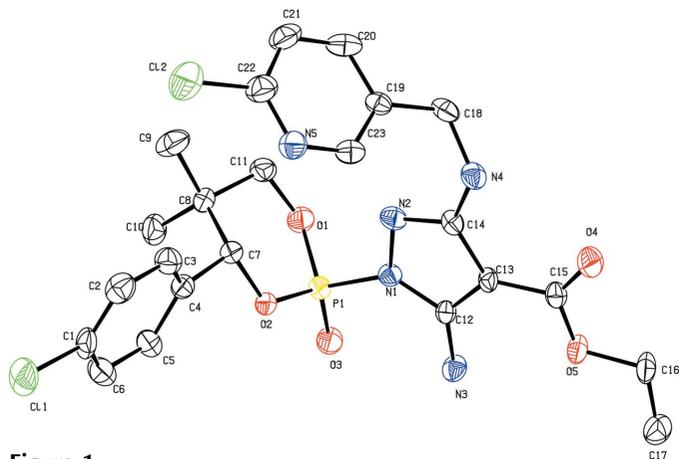


Figure 1
A view of the molecule of (I), showing the atom-numbering scheme and 50% probability displacement ellipsoids. H atoms have been omitted for clarity.

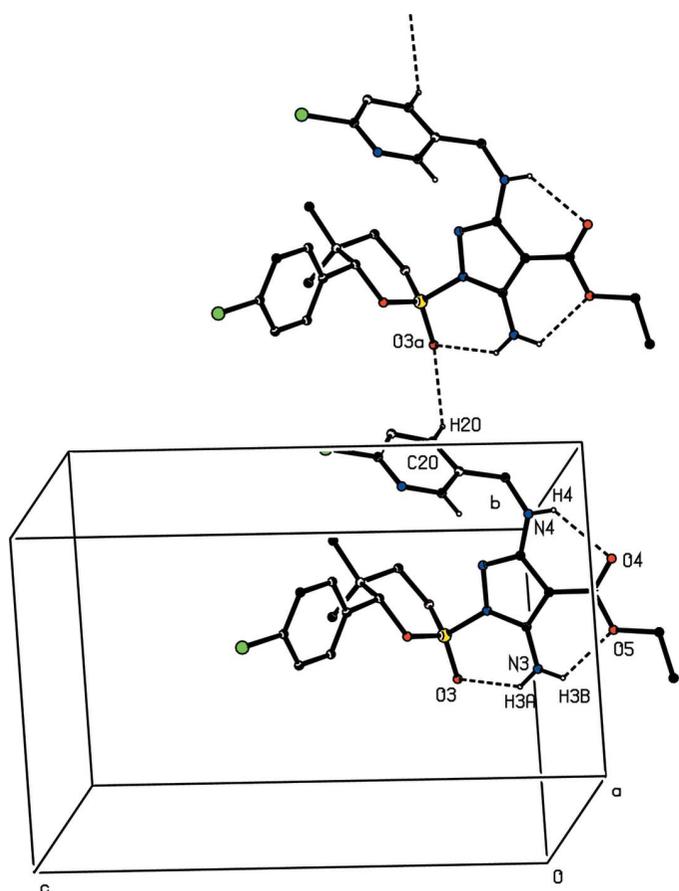


Figure 2
Part of the crystal packing of (I), showing a chain running along *b* formed by C—H...O interactions (dashed lines). The suffix *a* indicates the symmetry position *x*, *y* + 1, *z*.

2005]; on the other hand, the P1—N1 bond [1.677 (2) Å] is longer than the literature value [1.605 (3) Å; Liu *et al.*, 2005]. The terminal ethyl carboxylate group is almost coplanar with the plane of the pyrazole ring (Table 1). The C16—C17 bond distance [1.481 (4) Å] can be attributed to a disorder of the terminal ethyl group.

Intramolecular N—H...O hydrogen bonds form six-membered rings (Table 2) and contribute to the stability of the overall conformation. The intermolecular C20—H...O3 interaction joins molecules into a chain along the *b* axis (Table 2 and Fig. 2). Methyl atom C17 and methylene atom C18 are involved in C—H... π interactions: C17...Cg1 = 3.669 (3) Å, H17A...Cg1 = 2.81 Å and C17—H17A...Cg1 = 148.97°; C18...Cg2 = 3.521 (3) Å, H18A...Cg2 = 2.63 Å and C18—H18A...Cg2 = 153.61° [Cg1 is the centroid of the pyrazole ring of the molecule at (1 - *x*, 1 - *y*, -*z*) and Cg2 is the centroid of the pyrazole ring of the symmetry-related molecule at (-*x*, 2 - *y*, -*z*)]. The interaction of methyl and methylene groups with the aromatic π systems, described by Desiraju (2002), contributes to the crystal packing of (I).

Experimental

A solution of 5-amino-3-(6-chloro-3-pyridylmethylamino)-4-(ethoxycarbonyl)pyrazole (4 mmol) in anhydrous CH₃CN (20 ml) and NaOH powder (5 mmol) was added to a three-necked flask. After vigorously stirring for 5 min, a solution of 2-chloro-4-chlorophenyl-5,5-dimethyl-1,3,2-dioxaphosphinane 2-oxide (4.5 mmol) was added dropwise while cooling in an ice-bath. After the addition was complete, the mixture was stirred at room temperature until the reaction was complete (monitored by thin-layer chromatography). The work-up involved stripping of the solvent followed by addition of water and extraction of the product mixture into chloroform; after phase separation, drying over Na₂SO₄, filtration and evaporation, the crude product was purified by flash column chromatography on silica gel using petroleum ether/ethyl acetate (2:1 *v/v*) as eluent, giving a white solid (yield: 35%, m.p. 453 K). A colourless crystal grown from absolute ethanol was selected for X-ray structure analysis.

Crystal data

C₂₃H₂₆Cl₂N₅O₅P
M_r = 554.36
 Triclinic, *P* $\bar{1}$
a = 8.1968 (11) Å
b = 10.2180 (14) Å
c = 16.581 (2) Å
 α = 86.447 (2)°
 β = 85.668 (3)°
 γ = 70.083 (2)°
V = 1301.0 (3) Å³

Z = 2
D_x = 1.415 Mg m⁻³
 Mo *K* α radiation
 Cell parameters from 2290 reflections
 θ = 2.7–27.3°
 μ = 0.36 mm⁻¹
T = 292 (2) K
 Prism, colourless
 0.40 × 0.30 × 0.20 mm

Data collection

Bruker SMART CCD area-detector diffractometer
 φ and ω scans
 Absorption correction: multi-scan (SADABS; Bruker, 2000)
T_{min} = 0.871, *T_{max}* = 0.932
 6926 measured reflections

4540 independent reflections
 3469 reflections with *I* > 2 σ (*I*)
R_{int} = 0.058
 θ_{max} = 25.0°
h = -9 → 9
k = -12 → 11
l = -18 → 19

Refinement

Refinement on *F*²
R [*F*² > 2 σ (*F*²)] = 0.050
wR(*F*²) = 0.140
S = 1.05
 4540 reflections
 333 parameters
 H-atom parameters constrained

$w = 1/[\sigma^2(F_o^2) + (0.0711P)^2]$
 where $P = (F_o^2 + 2F_c^2)/3$
 $(\Delta/\sigma)_{max} = 0.001$
 $\Delta\rho_{max} = 0.44 \text{ e \AA}^{-3}$
 $\Delta\rho_{min} = -0.45 \text{ e \AA}^{-3}$
 Extinction correction: SHELXL97
 Extinction coefficient: 0.044 (3)

Table 1

Selected geometric parameters (Å, °).

C16—C17	1.481 (4)	O2—P1	1.5584 (17)
N1—P1	1.677 (2)	O3—P1	1.4529 (18)
O1—P1	1.5516 (19)		
O3—P1—O1	114.92 (10)	O3—P1—N1	108.80 (10)
O3—P1—O2	113.32 (10)	O1—P1—N1	105.96 (10)
O1—P1—O2	106.83 (9)	O2—P1—N1	106.44 (10)
C12—C13—C15—O4	176.3 (3)	C17—C16—O5—C15	179.0 (2)
C14—C13—C15—O5	−174.9 (2)		

Table 2

Hydrogen-bond geometry (Å, °).

<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>
C23—H23...N4	0.93	2.55	2.904 (4)	103
C20—H20...O3 ⁱ	0.93	2.58	3.253 (3)	129
N3—H3A...O3	0.86	2.17	2.839 (3)	134
N3—H3B...O5	0.86	2.29	2.832 (3)	121
N3—H3B...O3 ⁱⁱ	0.86	2.63	3.108 (3)	116
N4—H4...O4	0.80 (1)	2.34 (3)	2.938 (3)	132 (3)

Symmetry codes: (i) *x*, *y* + 1, *z*; (ii) *−x*, *−y* + 1, *−z*.

H atoms were treated as riding, with C—H distances in the range 0.93–0.98 Å and N—H distances of 0.86 Å, and $U_{\text{iso}}(\text{H}) = 1.2$ or $1.5U_{\text{eq}}(\text{parent atom})$.

Data collection: *SMART* (Bruker, 2000); cell refinement: *SAINT* (Bruker, 2000); data reduction: *SAINT*; program(s) used to solve structure: *SHELXS97* (Sheldrick, 1997); program(s) used to refine structure: *SHELXL97* (Sheldrick, 1997); molecular graphics: *SHELXTL* (Bruker, 1997); software used to prepare material for publication: *SHELXTL*.

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