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## Structure Reports

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## Key indicators

Single-crystal X-ray study
$T=297 \mathrm{~K}$
Mean $\sigma(\mathrm{C}-\mathrm{C})=0.004 \AA$
$R$ factor $=0.049$
$w R$ factor $=0.137$
Data-to-parameter ratio $=15.3$
For details of how these key indicators were automatically derived from the article, see http://journals.iucr.org/e.

[^0]
## Trianilinophosphine oxide

The title compound, $\mathrm{C}_{18} \mathrm{H}_{18} \mathrm{~N}_{3} \mathrm{OP}$, was synthesized by the condensation of phosphoryl chloride with an excess of aniline. The P atom is in a distorted tetrahedral environment. The molecules are stabilized by a weak intramolecular $\mathrm{C}-\mathrm{H} \cdots \mathrm{O}$ hydrogen bond and $\mathrm{C}-\mathrm{H} \cdots \pi$ interactions. The molecules are stacked along the $b$ axis.

## Comment

Variation in $\mathrm{P}-\mathrm{N}$ bond lengths has been observed in a survey of $p$-substituted phosphoric trianilides (Cameron et al., 1976). We therefore decided to investigate the $\mathrm{P}-\mathrm{N}$ bond lengths in the simple phosphoryl trianilide, (I), which was synthesized by the condensation of phosphoryl chloride and excess aniline.

(I)

All bond lengths in (I) show normal values (Allen et al., 1987). In this simple phosphoryl trianilide system there is little difference in the $\mathrm{P}-\mathrm{N}$ bond lengths (Table 1) compared with the reported $\mathrm{P}-\mathrm{N}$ bond lengths for $p$-tolylphosphoryl triamide $[$ i.e. 1.620 (7), 1.635 (7) and 1.669 (9) $\AA$ ] and other similar compounds (Cameron et al., 1976). This may be due to complete and equal overlap of the non-bonding $p_{z}$ orbitals of all three $\mathrm{N} s p^{3}$ atoms with an unoccupied $d$ orbital on the P atom. This is further supported by the shorter phosphoryl $\mathrm{P}=\mathrm{O}$ bond length in (I) compared with those in the $N, N^{\prime}, N^{\prime \prime}-$ tri-p-tolylphosphoryl triamide system [1.495 (8) $\AA$; Cameron et al., 1976]. The three $\mathrm{N}-\mathrm{P}-\mathrm{N}$ bond angles (Table 1) indicate that the P atom is in a distorted tetrahedral environment in order to minimize the steric interactions between the three aromatic rings.

The intramolecular $\mathrm{C} 14-\mathrm{H} 14 A \cdots \mathrm{O} 1$ hydrogen bond generates an $S(6)$ graph-set motif (Bernstein et al., 1995) and
$\qquad$


Figure 1
The molecular structure of (I), showing $50 \%$ probability displacement ellipsoids and the atom-numbering scheme. A dashed line indicates the intramolecular hydrogen bond.


Figure 2
The crystal packing of (I), viewed down the $b$ axis.
weak intermolecular $\mathrm{C}-\mathrm{H} \cdots \pi$ interactions stabilize the structure (Table 2). The molecules are stacked along the $b$ axis in parallel layers (Fig. 2).

## Experimental

Phosphoryl chloride $(1.5 \mathrm{ml}, 0.98 \mathrm{mmol})$ was added dropwise to an excess of aniline ( 8 ml ) at 273 K . After stirring for 20 min , the solid mass was filtered off, washed with dilute hydrochloric acid and crystallized from methanol to afford (I) (m.p. $546 \mathrm{~K}, 74 \%$ yield).

## Crystal data

$\mathrm{C}_{18} \mathrm{H}_{18} \mathrm{~N}_{3} \mathrm{OP}$
$M_{r}=323.32$
Orthorhombic, $P b c a$
$a=9.9173(8) \AA$
$b=9.2463(7) \AA$
$c=35.405(3) \AA$
$V=3246.6(4) \AA^{3}$

## Data collection

| Bruker SMART CCD area-detector | 16288 measured reflections |
| :--- | :--- |
| $\quad$ diffractometer | 3183 independent reflections |
| $\omega$ scans | 2735 reflections with $I>2 \sigma(I)$ |
| Absorption correction: multi-scan | $R_{\text {int }}=0.021$ |
| $\quad(S A D A B S ;$ Sheldrick, 1996) | $\theta_{\max }=26.0^{\circ}$ |
| $\quad T_{\min }=0.931, T_{\max }=0.959$ |  |

## Refinement

Refinement on $F^{2}$

$$
\begin{aligned}
& w=1 /\left[\sigma^{2}\left(F_{\mathrm{o}}^{2}\right)+(0.0685 P)^{2}\right. \\
& \quad+1.0693 P] \\
& \text { where } P=\left(F_{\mathrm{o}}^{2}+2 F_{\mathrm{c}}^{2}\right) / 3 \\
& (\Delta / \sigma)_{\max }=0.001 \\
& \Delta \rho_{\max }=0.18 \mathrm{e}^{-3} \\
& \Delta \rho_{\min }=-0.39 \mathrm{e} \mathrm{~A}^{-3}
\end{aligned}
$$

Table 1
Selected geometric parameters ( $\AA^{\circ},{ }^{\circ}$ ).

| P1-O1 | $1.4672(14)$ | $\mathrm{P} 1-\mathrm{N} 3$ | $1.6436(18)$ |
| :--- | :--- | :--- | :--- |
| $\mathrm{P} 1-\mathrm{N} 1$ | $1.6368(16)$ | $\mathrm{P} 1-\mathrm{N} 2$ | $1.6447(17)$ |
|  |  |  |  |
| $\mathrm{N} 1-\mathrm{P} 1-\mathrm{N} 3$ | $105.06(9)$ | $\mathrm{C} 1-\mathrm{N} 1-\mathrm{P} 1$ | $127.92(13)$ |
| $\mathrm{N} 1-\mathrm{P} 1-\mathrm{N} 2$ | $105.16(9)$ | $\mathrm{C} 7-\mathrm{N} 2-\mathrm{P} 1$ | $128.31(14)$ |
| $\mathrm{N} 3-\mathrm{P} 1-\mathrm{N} 2$ | $103.58(9)$ | $\mathrm{C} 13-\mathrm{N} 3-\mathrm{P} 1$ | $127.46(14)$ |

Table 2
Hydrogen-bond geometry ( $\AA$, ${ }^{\circ}$ ).
$C g 1$ and $C g 2$ are the centroids rings $\mathrm{C} 1-\mathrm{C} 6$ and $\mathrm{C} 7-\mathrm{C} 12$, respectively.

| $D-\mathrm{H} \cdots A$ | $D-\mathrm{H}$ | $\mathrm{H} \cdots A$ | $D \cdots A$ | $D-\mathrm{H} \cdots A$ |
| :---: | :---: | :---: | :---: | :---: |
| C14-H14A $\cdots$ O1 | 0.93 | 2.57 | 3.229 (3) | 128 |
| $\mathrm{C} 3-\mathrm{H} 3 B \cdots \mathrm{Cg} 2^{\text {i }}$ | 0.93 | 3.05 | 3.3714 | 132 |
| $\mathrm{C} 6-\mathrm{H} 6 A \cdots \mathrm{Cg} 1^{\text {ii }}$ | 0.93 | 3.17 | 3.3423 | 157 |
| $\mathrm{C} 11-\mathrm{H} 11 A \cdots C 22^{\text {iii }}$ | 0.93 | 2.97 | 3.3258 | 117 |

Symmetry codes: (i) $-x+\frac{3}{2}, y-\frac{3}{2}, z$; (ii) $-x+\frac{3}{2}, y-\frac{1}{2}, z$; (iii) $-x+\frac{5}{2}, y-\frac{1}{2}, z$.

H atoms were placed in calculated positions, with $\mathrm{N}-\mathrm{H}$ distances of $0.86 \AA$ and $\mathrm{C}-\mathrm{H}$ distances of $0.93 \AA$, and with $U_{\text {iso }}(\mathrm{H})=$ $1.2 U_{\text {eq }}(\mathrm{C}, \mathrm{N})$.

Data collection: SMART (Siemens, 1996); cell refinement: SAINT (Siemens, 1996); data reduction: SAINT; program(s) used to solve structure: SHELXTL (Sheldrick, 1997); program(s) used to refine structure: SHELXTL; molecular graphics: SHELXTL; software used to prepare material for publication: SHELXTL, PARST (Nardelli, 1995) and PLATON (Spek, 2003).

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