Acta Crystallographica Section E
Structure Reports
Online
ISSN 1600-5368

Rodolfo Moreno-Fuquen, ${ }^{\text {a* }}$ Mercé Font i Carot, ${ }^{\text {b }}$ Miquel Garriga, ${ }^{\text {b }}$ Felix Cano, ${ }^{\text {c }}$ Martin Martinez-Ripoll, ${ }^{\text {c }}$ Jaime<br>Valderrama-Naranjo ${ }^{\text {d }}$ and Luis Manuel Serratto ${ }^{\text {e }}$<br>a Departamento de Química, Facultad de Ciencias, Universidad del Valle, Apartado 25360, Santiago de Cali, Colombia, ${ }^{\text {b }}$ CSIC, Institut de Ciencia de Materials de Barcelona, Universidad Autónoma de Barcelona, Bellaterra, Spain, ${ }^{\text {CInstituto de Química, Física Rocasolano, }}$ CSIC, Departamento de Cristalografía, Serrano 119, E-28006 Madrid, Spain, ${ }^{\text {d Departamento de }}$ Física, Facultad de Ciencias, Universidad del Valle, Apartado 25360, Santiago de Cali, Colombia, and ${ }^{\mathbf{e}}$ Departamento de Procesos Químicos, Facultad de Ingenieria, Universidad del Valle, Apartado 25360, Santiago de Cali, Colombia

Correspondence e-mail: rodolfu@telesat.com.co

## Key indicators

Single-crystal X-ray study
$T=293 \mathrm{~K}$
Mean $\sigma(\mathrm{C}-\mathrm{C})=0.002 \AA$
$R$ factor $=0.044$
$w R$ factor $=0.146$
Data-to-parameter ratio $=21.2$
For details of how these key indicators were automatically derived from the article, see http://journals.iucr.org/e.
(C) 2003 International Union of Crystallography Printed in Great Britain - all rights reserved

# The 2:1 complex of 4 -aminobenzoic acid and 4,4'-bipyridyl $N, N^{\prime}$-dioxide 

The co-crystal $\mathrm{C}_{10} \mathrm{H}_{8} \mathrm{~N}_{2} \mathrm{O}_{2} \cdot 2 \mathrm{C}_{7} \mathrm{H}_{7} \mathrm{NO}_{2}$ owes its formation to an intermolecular hydrogen bond between the $\mathrm{O}-\mathrm{H}$ and $\mathrm{N}-\mathrm{O}$ groups; the $\mathrm{O} \cdots \mathrm{O}$ distance is $2.5796(17) \AA$. The dihedral angle between the planes of the pyridyl $N$-oxide and aminobenzoic rings of the complex is $10.2(1)^{\circ}$. An additional intermolecular $\mathrm{C}-\mathrm{H} \cdots \mathrm{O}$ hydrogen bond is formed, giving rise to an eight-membered ring within the asymmetric unit. The structure is centrosymmetric about the mid-point of the bond joining the two pyridine rings. The crystal structure exhibits overlap between the aromatic rings of the molecules in the [110] direction.

## Comment

The title compound, (I), was investigated as part of a study on $D-\mathrm{H} \cdots A$ hydrogen bonding in systems containing 4,4'bipyridyl $N, N^{\prime}$-dioxide and hydrogen-bond donors (Thaimattam et al., 1998). Four molecular structures similar to (I) were found in the Cambridge Structural Database (Allen, 2002), namely $4,4^{\prime}$-bipyridine $N, N^{\prime}$-dioxide urea clathrate, $4,4^{\prime}$ bipyridine $N, N^{\prime}$-dioxide thiourea clathrate and $4,4^{\prime}$-bipyridine $N, N^{\prime}$-dioxide dihydrate clathrate (Thaimattam et al., 1998), and 4-aminobenzoic acid (Lai \& Marsh, 1967); they were used as reference standards in analysing the structural characteristics of complex (I).

PABA


DIPNO


PABA
(I)

The complex is held together by a strong intermolecular hydrogen bond (Emsley, 1984) between the $\mathrm{O}-\mathrm{H}$ group of the 4 -aminobenzoic acid (PABA) molecule and the $\mathrm{N}-\mathrm{O}$ group of the $4,4^{\prime}$-bipyridyl $N, N^{\prime}$-dioxide (DIPNO) molecule. In the crystal structure, two acid components, related by a centre of symmetry, are linked to a centrosymmetric 4, $4^{\prime}$ bipyridyl $N, N^{\prime}$-dioxide, not only by short $\mathrm{O}-\mathrm{H} \cdots \mathrm{O}$ bonds but also by longer $\mathrm{C} 5-\mathrm{H} 5 \cdots \mathrm{O} 3$ hydrogen bonds, giving rise to an eight-membered ring within the asymmetric unit. This results in an approximately linear chain of the three molecules. The formation of these hydrogen bonds is probably related to the rotation of the carboxylic acid group around the C9-C12 bond [the dihedral angle between this group and the $\mathrm{C} 6-\mathrm{C} 11$

Received 18 February 2003
Accepted 10 March 2003
Online 21 March 2003


Figure 1
An ORTEP-3 (Farrugia, 1997) plot of the title complex, with the atomic labelling scheme. Displacement ellipsoids are plotted at the $50 \%$ probability level. H atoms are shown as spheres of arbitrary radius.


Figure 2
A ZORTEP (Zsolnai, 1995) diagram of (I), showing the overlap between the molecules along the [ 110 ] direction.
ring being $\left.7.5(1)^{\circ}\right]$ and to the approximate planarity of the whole centrosymmetric unit. Hydrogen-bond $\mathrm{O}-\mathrm{H} \cdots \mathrm{O}, \mathrm{N}-$ $\mathrm{H} \cdots \mathrm{O}$ and $\mathrm{C}-\mathrm{H} \cdots \mathrm{O}$ interactions are presented in Table 2. A perspective view of the DIPNO-PABA (1:2) adduct, showing the atomic numbering scheme, is given in Fig. 1. The dihedral angle formed by the planes which contain the rings of the DIPNO and PABA molecules is $10.2(1)^{\circ}$. The values of bond lengths and other internal parameters of DIPNO are very close to values reported in the literature (Thaimattam et al., 1998). The central $\mathrm{C} 3-\mathrm{C} 3^{\mathrm{i}}$ bond length $[1.475$ (3) $\AA$; symmetry code: (i) $-x,-y,-z]$ is shorter than a normal $\mathrm{C}-\mathrm{C}$ single-bond distance, showing that this $\mathrm{C}-\mathrm{C}$ bond probably has a little $\pi$-bond character. The presence of a strong O $\mathrm{H} \cdots \mathrm{O}$ and other relatively weak $\mathrm{C}-\mathrm{H} \cdots \mathrm{O}$ intermolecular hydrogen bonds induces some structural changes in the donor complex. Indeed, the $\mathrm{C} 9-\mathrm{C} 12, \mathrm{C} 12-\mathrm{O} 2$ and $\mathrm{C} 12-\mathrm{O} 3$ bond lengths change from 1.471 (2), 1.320 (2) and 1.218 (2) $\AA$ in (I) to 1.455 (4), 1.295 (3) and 1.248 (5) $\AA$ in the uncomplexed PABA molecule. There are no unusual short intermolecular contacts in the crystal structure. The molecules of PABA and $\operatorname{DIPNO}\left(\frac{1}{2}-x,-y, z-\frac{1}{2}\right)$ overlap along the [ $\overline{1} 10$ ] direction at a distance of 3.454 (2) $\AA$ (Fig. 2). Additionally, hydrogen-bond interactions between atom N 2 with atoms $\mathrm{O} 3{ }^{\mathrm{ii}}$ and $\mathrm{O} 1^{\mathrm{iii}}$ of neighboring molecules [symmetry codes: (ii) $x, \frac{1}{2}-y, \frac{1}{2}+z$; (iii) $\left.\frac{1}{2}+x, \frac{1}{2}-y, 1-z\right]$ are observed (Fig. 3).


Figure 3
A ZORTEP (Zsolnai, 1995) view of (I), showing additional hydrogen bonds, indicated by dashed lines.

## Experimental

Yellow single crystals of DIPNO-PABA (1:2) [m.p. 453 (1) K] suitable for X-ray analysis were obtained by slow evaporation from an equimolar solution of DIPNO and PABA in methanol. Initial reagents were purchased from Aldrich and were used without additional purification.

## Crystal data

$\mathrm{C}_{10} \mathrm{H}_{8} \mathrm{~N}_{2} \mathrm{O}_{2} \cdot 2 \mathrm{C}_{7} \mathrm{H}_{7} \mathrm{NO}_{2}$
$M_{r}=462.46$
Orthorhombic, Pbca
$a=11.188$ (1) $\AA$
$b=12.138$ (2) $\AA$
$c=16.354$ (2) $\AA$
$V=2220.9(5) \AA^{3}$
$Z=4$
$D_{x}=1.383 \mathrm{Mg} \mathrm{m}^{-3}$

## Data collection

Enraf-Nonius CAD-4

## diffractometer

$\omega / 2 \theta$ scans
Absorption correction: none
6757 measured reflections 3350 independent reflections
2057 reflections with $I>2 \sigma(I)$
$R_{\text {int }}=0.026$

## Refinement

Refinement on $F^{2}$
$R\left[F^{2}>2 \sigma\left(F^{2}\right)\right]=0.044$
$w R\left(F^{2}\right)=0.146$
$S=1.02$
3350 reflections
158 parameters
H atoms treated by a mixture of independent and constrained refinement

Mo $K \alpha$ radiation
Cell parameters from 25 reflections
$\theta=9-18^{\circ}$
$\mu=0.10 \mathrm{~mm}^{-1}$
$T=293$ (2) K
Regular block, yellow
$0.22 \times 0.18 \times 0.12 \mathrm{~mm}$

$$
\begin{aligned}
& \theta_{\max }=30.4^{\circ} \\
& h=0 \rightarrow 15 \\
& k=0 \rightarrow 17 \\
& l=-23 \rightarrow 23 \\
& 3 \text { standard reflections } \\
& \quad \text { frequency: } 120 \mathrm{~min} \\
& \quad \text { intensity decay: } 1.9 \%
\end{aligned}
$$

$$
\begin{gathered}
w=1 /\left[\sigma^{2}\left(F_{o}{ }^{2}\right)+(0.0639 P)^{2}\right. \\
+0.5535 P] \\
\text { where } P=\left(F_{o}{ }^{2}+2 F_{c}^{2}\right) / 3 \\
(\Delta / \sigma)_{\max }<0.001 \\
\Delta \rho_{\max }=0.22 \mathrm{e} \AA^{-3} \\
\Delta \rho_{\min }=-0.20 \mathrm{e}^{-3}
\end{gathered}
$$

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

| $\mathrm{O} 1-\mathrm{N} 1$ | $1.3243(16)$ | $\mathrm{C} 3-\mathrm{C} 3^{\mathrm{i}}$ | $1.475(3)$ |
| :--- | :--- | :--- | :--- |
|  |  |  |  |
| $\mathrm{O} 1-\mathrm{N} 1-\mathrm{C} 1$ | $119.23(13)$ | $\mathrm{O} 3-\mathrm{C} 12-\mathrm{O} 2$ | $122.43(15)$ |
| $\mathrm{O} 1-\mathrm{N} 1-\mathrm{C} 5$ | $120.95(13)$ | $\mathrm{O} 3-\mathrm{C} 12-\mathrm{C} 9$ | $123.00(15)$ |
| $\mathrm{N} 2-\mathrm{C} 6-\mathrm{C} 7$ | $120.26(16)$ | $\mathrm{O} 2-\mathrm{C} 12-\mathrm{C} 9$ | $114.53(13)$ |
| $\mathrm{C} 8-\mathrm{C} 9-\mathrm{C} 12-\mathrm{O} 2$ | $174.67(14)$ |  |  |

Symmetry code: (i) $-x,-y,-z$.

Table 2
Hydrogen-bonding geometry $\left(\AA \AA^{\circ}\right)$.

| $D-\mathrm{H} \cdots A$ | $D-\mathrm{H}$ | $\mathrm{H} \cdots A$ | $D \cdots A$ | $D-\mathrm{H} \cdots A$ |
| :--- | :--- | :--- | :--- | :--- |
| O2-HO2 $\cdots$ O1 | $0.90(2)$ | $1.73(2)$ | $2.580(2)$ | $157(2)$ |
| N2-HN21 $\cdots$ O3 $^{\text {ii }}$ | 0.86 | 2.23 | $2.975(2)$ | 146 |
| N2-HN22 $\cdots 1^{\text {iii }}$ | 0.86 | 2.30 | $3.095(2)$ | 154 |
| C5-H5 $\cdots$ O3 | 0.93 | 2.39 | $3.273(2)$ | 158 |

Symmetry codes: (ii) $x, \frac{1}{2}-y, \frac{1}{2}+z$; (iii) $\frac{1}{2}+x, \frac{1}{2}-y, 1-z$.
All H atoms, with the exception of the carboxyl atom HO 2 , were placed at geometrically idealized positions and were allowed to ride on their parent atoms; $\mathrm{N}-\mathrm{H}=0.86, \mathrm{C}-\mathrm{H}=0.93 \AA$ and $U_{\text {iso }}(\mathrm{H})=$
$1.2 U_{\text {eq }}$ of the carrier atom. Hydroxyl atom HO 2 was located from a Fourier difference map and its coordinates were refined freely.

Data collection: CAD-4 EXPRESS (Enraf-Nonius, 1994); cell refinement: CAD-4 EXPRESS; data reduction: XCAD4 (Harms \& Wocadlo, 1995); program(s) used to solve structure: SHELXS86 (Sheldrick, 1990); program(s) used to refine structure: SHELXL97 (Sheldrick, 1997); molecular graphics: ORTEP-3 (Farrugia, 1997) and ZORTEP (Zsolnai, 1995); software used to prepare material for publication: SHELXL97.

One of the authors (RMF) acknowledges Universidad del Valle, Colombia, for partial support of this work.

## References

Allen, F. H. (2002). Acta Cryst. B58, 380-388.
Emsley, J. (1984). Complex Chemistry, Structure and Bonding. Vol. 57, pp. 147191. Berlin: Springer-Verlag.

Enraf-Nonius (1994). CAD-4 EXPRESS. Version 5.1/1.2. Enraf-Nonius, Delft, The Netherlands.
Farrugia, L. J. (1997). J. Appl. Cryst. 30, 565.
Harms, K. \& Wocadlo, S. (1995). XCAD4. University of Marburg, Germany.
Lai, T. F. \& Marsh, R. E. (1967). Acta Cryst. 22, 885-893.
Sheldrick, G. M. (1990). Acta Cryst. A46, 467-473.
Sheldrick, G. M. (1997). SHELXL97. University of Göttingen, Germany.
Thaimattam, R., Reddy, D. S., Xue, F., Mak, T. W. C., Nangia, A. \& Desiraju, G. R. (1998). J. Chem. Soc. Perkin Trans. 2, pp. 1783-1789.

Zsolnai, L. (1995). ZORTEP. University of Heidelberg, Germany.

