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## Key indicators

Single-crystal X-ray study
$T=295 \mathrm{~K}$
Mean $\sigma(\mathrm{C}-\mathrm{C})=0.003 \AA$
Disorder in main residue
$R$ factor $=0.037$
$w R$ factor $=0.117$
Data-to-parameter ratio $=12.4$
For details of how these key indicators were automatically derived from the article, see http://journals.iucr.org/e.
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## Rerefinement of the co-crystal of 4-hydroxypyridine with 4-hydroxypyridinium nitrate

The crystal structure of the title salt adduct, $\mathrm{C}_{5} \mathrm{H}_{6} \mathrm{NO}^{+} \cdot \mathrm{NO}_{3}^{-} \cdot \mathrm{C}_{5} \mathrm{H}_{5} \mathrm{NO}$, has a $\mathrm{C}_{5} \mathrm{H}_{6} \mathrm{NO}^{+} / \mathrm{C}_{5} \mathrm{HNO}$ cation/ molecule assembly (which lies on an inversion center) whose components are linked through their O atoms by a short hydrogen bond $[2.446$ (3) $\AA$ ]. The cation is formally the 4 hydroxypyridinium cation and the neutral component the 4 pyridinone molecule, but the two could not be differentiated as they occupy equivalent positions, with the H atom disordered between them. The nitrogen-bound H atom interacts with the nitrate ion [2.840 (1) $\AA$ ], which lies on another special position of site symmetry 2 .

## Comment

4-Pyridinone, which exists as a $6 / 5$-hydrate (Jones, 2001), is reported to form a 2:1 adduct with nitric acid; the earlier study had placed an H atom midway between two neutral 4 -pyridinone molecules, on a center of inversion (Goodgame et al., 1993). In the present study, the structure was refined without imposing the symmetry constraint on this atom; the structure is consequently interpreted as the title co-crystal (I) (Fig. 1).

(I)

The position of the H atom along the linear internuclear axis, when investigated by a $P M 3$ potential calculation carried out with HYPERCHEM (Hypercube, 2000), coincides with the minimum energy when it is off center, at about $1 \AA$ from the single-bonded O atom. The calculations assumed a structure where one ring has a hydroxy group and the other has a double-bonded carbonyl O atom. The published structure (Goodgame et al., 1993) with symmetrical $\mathrm{O} \cdots \mathrm{H}$ bonds is probably chemically less appropriate.

## Experimental

The title co-crystal was the unexpected compound isolated from the reaction of cadmium nitrate tetrahydrate ( $3.08 \mathrm{~g}, 10 \mathrm{mmol}$ ) and 4 hydroxypyridine $(1.4 \mathrm{~g}, 15 \mathrm{mmol})$ in ethanol. The solvent was allowed to evaporate over several days to furnish colorless prismatic crystals of the organic compound. Calculated for $\mathrm{C}_{10} \mathrm{H}_{10} \mathrm{~N}_{3} \mathrm{O}_{5}$ : C $47.62, \mathrm{H}$ 4.00 , N $16.66 \%$; found: C 47.58, H 3.94, N $16.5 \%$. The reaction of the

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heterocycle with cobalt nitrate had yielded the water-coordinated bisadduct (Lu et al., 2004).
Crystal data
$\mathrm{C}_{5} \mathrm{H}_{6} \mathrm{NO}^{+} \cdot \mathrm{NO}_{3}{ }^{-} \cdot \mathrm{C}_{5} \mathrm{H}_{5} \mathrm{NO}$
$M_{r}=253.22$
Monoclinic, $\mathrm{C} 2 / \mathrm{c}$
$a=17.993(4) \AA$
$b=6.528(1) \AA$
$c=12.501(3) \AA$
$\beta=127.74(3)^{\circ}$
$V=1161.2(4) \AA^{3}$
$Z=4$

$$
\begin{aligned}
& D_{x}=1.448 \mathrm{Mg} \mathrm{~m}^{-3} \\
& \text { Mo } K \alpha \text { radiation } \\
& \text { Cell parameters from } 5340 \\
& \text { reflections } \\
& \theta=3.3-27.5^{\circ} \\
& \mu=0.12 \mathrm{~mm}^{-1} \\
& T=295(2) \mathrm{K} \\
& \text { Prism, colorless } \\
& 0.38 \times 0.24 \times 0.18 \mathrm{~mm}
\end{aligned}
$$

## Data collection

Rigaku R-AXIS RAPID
diffractometer
$\omega$ scans
Absorption correction: multi-scan
(ABSCOR; Higashi, 1995)
$T_{\text {min }}=0.858, T_{\text {max }}=0.979$
5541 measured reflections
1338 independent reflections
1010 reflections with $I>2 \sigma(I)$
$R_{\text {int }}=0.016$
$\theta_{\text {max }}=27.5^{\circ}$
$h=-23 \rightarrow 23$
$k=-7 \rightarrow 8$
$l=-16 \rightarrow 16$

## Refinement

Refinement on $F^{2}$
$R\left[F^{2}>2 \sigma\left(F^{2}\right)\right]=0.037$
$w R\left(F^{2}\right)=0.117$
$S=1.08$
1338 reflections
108 parameters
All H -atom parameters refined

$$
\begin{aligned}
& w=1 /\left[\sigma^{2}\left(F_{o}{ }^{2}\right)+(0.0639 P)^{2}\right. \\
& +0.2278 P] \\
& \text { where } P=\left(F_{o}{ }^{2}+2 F_{c}^{2}\right) / 3 \\
& (\Delta / \sigma)_{\text {max }}=0.001 \\
& \Delta \rho_{\max }=0.21 \mathrm{e} \AA^{-3} \\
& \Delta \rho_{\text {min }}=-0.17 \mathrm{e}^{-3} \\
& \text { Extinction correction: SHELXL97 } \\
& \text { Extinction coefficient: } 0.005 \text { (2) }
\end{aligned}
$$

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

| $\mathrm{O} 1-\mathrm{C} 3$ | $1.291(2)$ | $\mathrm{C} 2-\mathrm{C} 3$ | $1.403(2)$ |
| :--- | :--- | :--- | :--- |
| $\mathrm{N} 1-\mathrm{C} 1$ | $1.339(2)$ | $\mathrm{C} 3-\mathrm{C} 4$ | $1.411(2)$ |
| $\mathrm{N} 1-\mathrm{C} 5$ | $1.339(2)$ | $\mathrm{C} 4-\mathrm{C} 5$ | $1.358(2)$ |
| $\mathrm{C} 1-\mathrm{C} 2$ | $1.354(2)$ |  |  |
| $\mathrm{C} 1-\mathrm{N} 1-\mathrm{C} 5$ | $120.6(1)$ | $\mathrm{O} 1-\mathrm{C} 3-\mathrm{C} 4$ | $123.3(1)$ |
| $\mathrm{N} 1-\mathrm{C} 1-\mathrm{C} 2$ | $121.5(2)$ | $\mathrm{C} 2-\mathrm{C} 3-\mathrm{C} 4$ | $116.8(1)$ |
| $\mathrm{C} 1-\mathrm{C} 2-\mathrm{C} 3$ | $120.0(1)$ | $\mathrm{C} 3-\mathrm{C} 4-\mathrm{C} 5$ | $120.1(1)$ |
| $\mathrm{O} 1-\mathrm{C} 3-\mathrm{C} 2$ | $119.9(1)$ | $\mathrm{N} 1-\mathrm{C} 5-\mathrm{C} 4$ | $121.0(1)$ |

The carbon-bound H atoms were refined with a distance restraint of 0.95 (1) $\AA$; the nitrogen-bound H atom was refined with a restraint of 0.85 (1) $\AA$. The final H atom that would balance the charges was initially located on the $\overline{1}$ site $\left(\frac{1}{4}, \frac{1}{4}, \frac{1}{2}\right)$, the atom being midway between the O atom of the cation and that of the symmetry-related neutral molecule. This H atom was then allowed to refine without this symmetry constraint, but with a distance restraint of 0.85 (1) $\AA$. The


Figure 1
ORTEPII (Johnson, 1976) plot of (I), with ellipsoids drawn at the $50 \%$ probability level. H atoms are drawn as spheres of arbitrary radii. Dashed lines indicate hydrogen bonds.

H atom is thus disordered over two positions, and the cation and neutral molecule are disordered on equivalent sites..

Data collection: RAPID-AUTO (Rigaku, 1998); cell refinement: RAPID-AUTO; data reduction: CrystalStructure (Rigaku/MSC, 2002); program(s) used to solve structure: SHELXS97 (Sheldrick, 1997); program(s) used to refine structure: SHELXL97 (Sheldrick, 1997); molecular graphics: ORTEPII (Johnson, 1976); software used to prepare material for publication: SHELXL97.

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