## Acta Crystallographica Section E

## Structure Reports

Online
ISSN 1600-5368

## 5-(Pyridinium-4-yl)-1 H-1,2,3,4-tetrazol-1-ide

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Received 17 November 2010; accepted 1 December 2010

Key indicators: single-crystal X-ray study; $T=298 \mathrm{~K}$; mean $\sigma(\mathrm{C}-\mathrm{C})=0.004 \AA$; $R$ factor $=0.040 ; w R$ factor $=0.096$; data-to-parameter ratio $=7.2$.

In the title zwitterionic molecule, $\mathrm{C}_{6} \mathrm{H}_{5} \mathrm{~N}_{5}$, the tetrazole and pyridine rings are nearly coplanar, making a dihedral angle of 2.08 (1) ${ }^{\circ}$. In the crystal, molecules are connected by classical $\mathrm{N}-\mathrm{H} \cdots \mathrm{N}$ and weak $\mathrm{C}-\mathrm{H} \cdots \mathrm{N}$ hydrogen bonds.

## Related literature

For applications of tetrazole derivatives, see: Zhao et al. (2008); Fu et al. (2008, 2009). For the crystal structures and properties of related compounds, see: Fu et al. (2007, 2009); Fu \& Xiong (2008).


## Experimental

## Crystal data

$\mathrm{C}_{6} \mathrm{H}_{5} \mathrm{~N}_{5}$
$M_{r}=147.15$
Monoclinic, Cc
$a=7.0508$ (14) $\AA$
$b=7.4007$ (15) A
$c=11.926$ (2) $\AA$
$\beta=96.56$ (3) ${ }^{\circ}$

Data collection
Rigaku Mercury2 diffractometer
3122 measured reflections
719 independent reflections

## Refinement

$R\left[F^{2}>2 \sigma\left(F^{2}\right)\right]=0.040 \quad 2$ restraints
$w R\left(F^{2}\right)=0.096$
$S=1.13$
719 reflections
100 parameters

H -atom parameters constrained
$\Delta \rho_{\text {max }}=0.23 \mathrm{e}^{-3}$
633 reflections with $I>2 \sigma(I)$ $R_{\text {int }}=0.039$
$\Delta \rho_{\text {min }}=-0.21 \mathrm{e}^{-3}$

Table 1
Hydrogen-bond geometry ( $\AA \AA^{\circ}$ ).

| $D-\mathrm{H} \cdots A$ | $D-\mathrm{H}$ | $\mathrm{H} \cdots A$ | $D \cdots A$ | $D-\mathrm{H} \cdots A$ |
| :--- | :--- | :--- | :--- | :--- |
| $\mathrm{~N} 1-\mathrm{H} 1 A \cdots \mathrm{~N}^{\mathrm{i}}$ | 0.86 | 1.89 | $2.745(4)$ | 176 |
| $\mathrm{~N} 1-\mathrm{H} 1 A \cdots \mathrm{~N}^{\mathrm{i}}$ | 0.86 | 2.52 | $3.306(4)$ | 152 |
| $\mathrm{C} 1-\mathrm{H} 1 \cdots \mathrm{~N} 5^{\mathrm{ii}}$ | 0.93 | 2.46 | $3.308(4)$ | 152 |
| $\mathrm{C} 5-\mathrm{H} 5 \cdots \mathrm{~N} 4^{\text {iii }}$ | 0.93 | 2.38 | $3.168(4)$ | 142 |
| Symmetry codes: (i) $x+\frac{1}{2},-y+\frac{1}{2}, z+\frac{1}{2} ;$ |  |  |  | (ii) $x, y+1, z ;$ (iii) $x+\frac{1}{2},-y-\frac{1}{2}, z+\frac{1}{2}$. |

Data collection: CrystalClear (Rigaku, 2005); cell refinement: CrystalClear; data reduction: CrystalClear; program(s) used to solve structure: SHELXTL (Sheldrick, 2008); program(s) used to refine structure: SHELXTL; molecular graphics: SHELXTL; software used to prepare material for publication: SHELXTL.

This work was supported by a start-up grant from Southeast University, China.

Supplementary data and figures for this paper are available from the IUCr electronic archives (Reference: XU5093).

## References

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## supporting information

Acta Cryst. (2011). E67, o43 [https://doi.org/10.1107/S1600536810050257]
5-(Pyridinium-4-yl)-1H-1,2,3,4-tetrazol-1-ide

## Qian Xu and Jie Xu

## S1. Comment

Tetrazole compounds attracted more attention as phase transition dielectric materials for its application in microelectronics, memory storage. With the purpose of obtaining phase transition crystals of tetrazole compound, a series of new materials have been elaborated with this organic molecule (Zhao et al., 2008; Fu et al., 2008; Fu et al., 2007; Fu \& Xiong 2008). We report here the crystal structure of the title compound, 5-(pyridinium-4-yl)tetrazol-1-ide.

The dielectric constant of title compound as a function of temperature indicates that the permittivity is basically temperature-independent, suggesting that this compound should be not a real ferroelectrics or there may be no distinct phase transition occurred within the measured temperature range. Similarly, below the melting point (413K) of the compound, the dielectric constant as a function of temperature also goes smoothly, and there is no dielectric anomaly observed (dielectric constant equaling to 6.1 to 7.9 ).
In the title compound (Fig.1), the pyridine N atom is protonated, thus indicating a positive charge in the pyridine N atom. And the tetrazole ring was showing a negative charge to make the charge balance. The tetrazole and pyridine rings are twisted from each other by a dihedral angle of $2.08(1)^{\circ}$. The geometric parameters of the tetrazole rings are comparable to those in related molecules (Fu et al., 2009).
In the crystal structure the molecules are connected by classic $\mathrm{N}-\mathrm{H} \cdots \mathrm{N}$ and weak $\mathrm{C}-\mathrm{H} \cdots \mathrm{N}$ hydrogen bonds (Table 1).

## S2. Experimental

5-(Pyridinium-4-yl)tetrazol-1-ide was obtained commercially, and the single crystals were obtained from an ethanol solution.

## S3. Refinement

H atoms attached to N atoms were located in a difference Fourier map, and refined in riding mode with $\mathrm{N}-\mathrm{H}=0.86 \AA$ and $\mathrm{U}_{\text {iso }}(\mathrm{H})=1.2 \mathrm{U}_{e q}(\mathrm{~N})$. Other H atoms were fixed geometrically and treated as riding with $\mathrm{C}-\mathrm{H}=0.93 \AA$ and $\mathrm{U}_{\text {iso }}(\mathrm{H})=$ $1.2 \mathrm{U}_{e q}(\mathrm{C})$. As no significant anomalous scattering, Friedel pairs were merged.


Figure 1
A view of the title compound with the atomic numbering scheme. Displacement ellipsoids were drawn at the $30 \%$ probability level.

## 5-(Pyridinium-4-yl)-1H-1,2,3,4-tetrazol-1-ide

## Crystal data

$\mathrm{C}_{6} \mathrm{H}_{5} \mathrm{~N}_{5}$
$M_{r}=147.15$
Monoclinic, Cc
Hall symbol: C -2yc
$a=7.0508$ (14) $\AA$
$b=7.4007(15) \AA$
$c=11.926(2) \AA$
$\beta=96.56(3)^{\circ}$
$V=618.2(2) \AA^{3}$
$Z=4$

## Data collection

Rigaku Mercury2
diffractometer
Radiation source: fine-focus sealed tube
Graphite monochromator
Detector resolution: 13.6612 pixels $\mathrm{mm}^{-1}$
CCD profile fitting scans
3122 measured reflections

## Refinement

Refinement on $F^{2}$
Least-squares matrix: full
$R\left[F^{2}>2 \sigma\left(F^{2}\right)\right]=0.040$
$w R\left(F^{2}\right)=0.096$
$S=1.13$
719 reflections
100 parameters
2 restraints
$F(000)=304$
$D_{\mathrm{x}}=1.581 \mathrm{Mg} \mathrm{m}^{-3}$
Mo $K \alpha$ radiation, $\lambda=0.71073 \AA$
Cell parameters from 1425 reflections
$\theta=3.4-24.5^{\circ}$
$\mu=0.11 \mathrm{~mm}^{-1}$
$T=298 \mathrm{~K}$
Block, colorless
$0.30 \times 0.20 \times 0.15 \mathrm{~mm}$

719 independent reflections
633 reflections with $I>2 \sigma(I)$
$R_{\text {int }}=0.039$
$\theta_{\text {max }}=27.5^{\circ}, \theta_{\text {min }}=3.4^{\circ}$
$h=-9 \rightarrow 9$
$k=-9 \rightarrow 9$
$l=-15 \rightarrow 15$

Primary atom site location: structure-invariant direct methods
Secondary atom site location: difference Fourier map
Hydrogen site location: inferred from neighbouring sites
H -atom parameters constrained

$$
\begin{aligned}
& w=1 /\left[\sigma^{2}\left(F_{\mathrm{o}}^{2}\right)+(0.056 P)^{2}\right] \\
& \text { where } P=\left(F_{\mathrm{o}}^{2}+2 F_{\mathrm{c}}^{2}\right) / 3 \\
& (\Delta / \sigma)_{\max }<0.001
\end{aligned}
$$

$$
\begin{aligned}
& \Delta \rho_{\max }=0.23 \mathrm{e} \AA^{-3} \\
& \Delta \rho_{\min }=-0.21 \mathrm{e} \AA^{-3}
\end{aligned}
$$

## Special details

Geometry. All esds (except the esd in the dihedral angle between two 1.s. planes) are estimated using the full covariance matrix. The cell esds are taken into account individually in the estimation of esds in distances, angles and torsion angles; correlations between esds in cell parameters are only used when they are defined by crystal symmetry. An approximate (isotropic) treatment of cell esds is used for estimating esds involving l.s. planes.
Refinement. Refinement of $\mathrm{F}^{2}$ against ALL reflections. The weighted R -factor wR and goodness of fit S are based on $\mathrm{F}^{2}$, conventional $R$-factors $R$ are based on $F$, with $F$ set to zero for negative $F^{2}$. The threshold expression of $F^{2}>2 \operatorname{sigma}\left(\mathrm{~F}^{2}\right)$ is used only for calculating R-factors (gt) etc. and is not relevant to the choice of reflections for refinement. R-factors based on $\mathrm{F}^{2}$ are statistically about twice as large as those based on F , and R - factors based on ALL data will be even larger.

Fractional atomic coordinates and isotropic or equivalent isotropic displacement parameters ( $\AA^{2}$ )

|  | $x$ | $y$ | $z$ | $U_{\text {iso }} * / U_{\text {eq }}$ |
| :--- | :--- | :--- | :--- | :--- |
| N1 | $1.0961(4)$ | $0.3035(4)$ | $0.4442(2)$ | $0.0382(6)$ |
| H1A | 1.1462 | 0.3791 | 0.4937 | $0.046^{*}$ |
| N2 | $0.7525(4)$ | $-0.0341(4)$ | $0.1099(2)$ | $0.0375(6)$ |
| N3 | $0.7094(4)$ | $-0.1968(3)$ | $0.0653(2)$ | $0.0443(7)$ |
| N4 | $0.7884(4)$ | $-0.3224(3)$ | $0.1321(2)$ | $0.0444(8)$ |
| N5 | $0.8864(3)$ | $-0.2457(4)$ | $0.2218(2)$ | $0.0395(7)$ |
| C1 | $0.9963(5)$ | $0.3659(4)$ | $0.3504(3)$ | $0.0397(9)$ |
| H1 | 0.9805 | 0.4896 | 0.3394 | $0.048^{*}$ |
| C2 | $0.9185(4)$ | $0.2487(4)$ | $0.2718(3)$ | $0.0368(7)$ |
| H2 | 0.8496 | 0.2920 | 0.2062 | $0.044^{*}$ |
| C3 | $0.9406(4)$ | $0.0634(4)$ | $0.2879(2)$ | $0.0290(6)$ |
| C4 | $1.0442(4)$ | $0.0050(4)$ | $0.3861(2)$ | $0.0372(8)$ |
| H4 | 1.0622 | -0.1179 | 0.3996 | $0.045^{*}$ |
| C5 | $1.1202(4)$ | $0.1280(4)$ | $0.4632(3)$ | $0.0391(7)$ |
| H5 | 1.1896 | 0.0888 | 0.5298 | $0.047^{*}$ |
| C6 | $0.8605(4)$ | $-0.0704(4)$ | $0.2067(2)$ | $0.0305(7)$ |
|  |  |  |  |  |

Atomic displacement parameters ( $\hat{A}^{2}$ )

|  | $U^{11}$ | $U^{22}$ | $U^{33}$ | $U^{12}$ | $U^{13}$ | $U^{23}$ |
| :--- | :--- | :--- | :--- | :--- | :--- | :--- |
| N1 | $0.0486(14)$ | $0.0342(16)$ | $0.0299(14)$ | $-0.0035(13)$ | $-0.0037(10)$ | $-0.0059(13)$ |
| N2 | $0.0465(15)$ | $0.0288(12)$ | $0.0353(13)$ | $0.0003(11)$ | $-0.0040(10)$ | $0.0010(11)$ |
| N3 | $0.0595(17)$ | $0.0348(13)$ | $0.0359(13)$ | $-0.0060(15)$ | $-0.0060(11)$ | $-0.0082(14)$ |
| N4 | $0.062(2)$ | $0.0293(14)$ | $0.0404(19)$ | $-0.0004(13)$ | $-0.0010(15)$ | $-0.0045(12)$ |
| N5 | $0.0533(16)$ | $0.0251(13)$ | $0.0377(16)$ | $0.0014(11)$ | $-0.0051(13)$ | $-0.0026(11)$ |
| C1 | $0.0455(17)$ | $0.029(2)$ | $0.0431(16)$ | $0.0041(15)$ | $0.0005(13)$ | $0.0020(15)$ |
| C2 | $0.0422(17)$ | $0.0311(16)$ | $0.0346(16)$ | $0.0031(13)$ | $-0.0059(12)$ | $0.0052(13)$ |
| C3 | $0.0355(14)$ | $0.0240(15)$ | $0.0264(13)$ | $0.0001(12)$ | $-0.0007(11)$ | $-0.0009(11)$ |
| C4 | $0.0514(19)$ | $0.0257(18)$ | $0.0319(15)$ | $0.0008(12)$ | $-0.0060(12)$ | $-0.0004(13)$ |
| C5 | $0.0512(18)$ | $0.0331(16)$ | $0.0304(14)$ | $0.0035(14)$ | $-0.0069(13)$ | $-0.0014(13)$ |
| C6 | $0.0369(16)$ | $0.0263(15)$ | $0.0272(14)$ | $0.0002(12)$ | $-0.0016(11)$ | $0.0014(13)$ |

Geometric parameters (A, ${ }^{\circ}$ )

| N1-C5 | 1.325 (4) | C1-H1 | 0.9300 |
| :---: | :---: | :---: | :---: |
| N1-C1 | 1.334 (5) | C2-C3 | 1.391 (4) |
| N1-H1A | 0.8600 | C2-H2 | 0.9300 |
| N2-C6 | 1.335 (4) | C3-C4 | 1.377 (4) |
| N2-N3 | 1.337 (4) | C3-C6 | 1.453 (4) |
| N3-N4 | 1.306 (4) | C4-C5 | 1.359 (4) |
| N4-N5 | 1.333 (4) | C4-H4 | 0.9300 |
| N5-C6 | 1.320 (4) | C5-H5 | 0.9300 |
| C1-C2 | 1.348 (5) |  |  |
| C5-N1-C1 | 121.8 (3) | C4-C3-C2 | 117.8 (3) |
| C5-N1-H1A | 119.1 | C4-C3-C6 | 118.8 (2) |
| C1-N1-H1A | 119.1 | C2-C3-C6 | 123.4 (2) |
| C6-N2-N3 | 104.1 (3) | C5-C4-C3 | 119.6 (3) |
| N4-N3-N2 | 109.6 (3) | C5-C4-H4 | 120.2 |
| N3-N4-N5 | 109.4 (2) | C3-C4-H4 | 120.2 |
| C6-N5-N4 | 104.9 (2) | N1-C5-C4 | 120.5 (3) |
| N1-C1-C2 | 119.6 (3) | N1-C5-H5 | 119.7 |
| N1-C1-H1 | 120.2 | C4-C5-H5 | 119.7 |
| $\mathrm{C} 2-\mathrm{C} 1-\mathrm{H} 1$ | 120.2 | N5-C6-N2 | 111.8 (3) |
| $\mathrm{C} 1-\mathrm{C} 2-\mathrm{C} 3$ | 120.5 (3) | N5-C6-C3 | 122.8 (2) |
| $\mathrm{C} 1-\mathrm{C} 2-\mathrm{H} 2$ | 119.7 | N2-C6-C3 | 125.4 (3) |
| C3-C2-H2 | 119.7 |  |  |

Hydrogen-bond geometry ( $A,{ }^{\circ}$ )

| $D — \mathrm{H} \cdots A$ | $D-\mathrm{H}$ | $\mathrm{H} \cdots A$ | $D \cdots A$ | $D-\mathrm{H} \cdots A$ |
| :--- | :--- | :--- | :--- | :--- |
| $\mathrm{~N} 1 — \mathrm{H} 1 A \cdots \mathrm{~N} 2^{\mathrm{i}}$ | 0.86 | 1.89 | $2.745(4)$ | 176 |
| $\mathrm{~N} 1 — \mathrm{H} 1 A \cdots \mathrm{~N} 3^{\mathrm{i}}$ | 0.86 | 2.52 | $3.306(4)$ | 152 |
| $\mathrm{C} 1 — \mathrm{H} 1 \cdots \mathrm{~N} 5^{\mathrm{ii}}$ | 0.93 | 2.46 | $3.308(4)$ | 152 |
| $\mathrm{C} 5 — \mathrm{H} 5 \cdots \mathrm{~N} 4^{\text {iii }}$ | 0.93 | 2.38 | $3.168(4)$ | 142 |

Symmetry codes: (i) $x+1 / 2,-y+1 / 2, z+1 / 2$; (ii) $x, y+1, z$; (iii) $x+1 / 2,-y-1 / 2, z+1 / 2$.

