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## Structure Reports

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## Key indicators

Single-crystal X-ray study
$T=90 \mathrm{~K}$
Mean $\sigma(\mathrm{C}-\mathrm{C})=0.003 \AA$
$R$ factor $=0.047$
$w R$ factor $=0.131$
Data-to-parameter ratio $=18.7$

For details of how these key indicators were automatically derived from the article, see http://journals.iucr.org/e.
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## 2-Amino-5-butyl-4-methyl-1,3-thiazol-3-ium nitrate

The title compound, $\mathrm{C}_{8} \mathrm{H}_{15} \mathrm{~N}_{3} \mathrm{O}_{3} \mathrm{~S}$, shows bond lengths and angles that are typical and are in accordance with expected values. The structure comprises a substituted thiazolium ring that is connected to a nitrate ion via $\mathrm{N}-\mathrm{H} \cdots \mathrm{O}$ hydrogenbonding interactions.

## Comment

The cation of the title compound, (I), is nearly planar. The torsion angles which deviate most significantly from 0 or $180^{\circ}$ are $\mathrm{S} 1-\mathrm{C} 5-\mathrm{C} 8-\mathrm{C} 9 \quad\left[5.5(3)^{\circ}\right]$ and $\mathrm{C} 4-\mathrm{C} 5-\mathrm{C} 8-\mathrm{C} 9$ [ $\left.-175.5(2)^{\circ}\right]$. This corresponds to the twist of the butyl group with respect to the thiazole ring. The bond lengths and angles are in accordance with the expected values (Fitzsimons \& Gallagher, 1999; Lynch \& McClenaghan, 2001). Even the C2$\mathrm{S} 1-\mathrm{C} 5$ angle of $90.9(1)^{\circ}$ is typical for substituted thiamines. A survey of the Cambridge Structural Database (Allen, 2002) for 1,3,4-thiazole rings gave a mean value of $90.8^{\circ}$.

(I)

Strong interactions are observed between the S atom and the nitrate ions. These interactions have been observed in other structures and it has been suggested that they have a mechanistic importance in C2-substituted thiamines (Yang et al., 1987). This relatively strong short contact [S1 . O O15 = 3.15 (2) $\AA$ ] contributes to the stabilization of the molecular packing along the $a$ axis. The packing is stabilized by ionic interactions between the nitrate anions, the organic cations, and $\mathrm{N}-\mathrm{H} \cdots \mathrm{O}$ hydrogen bonds. Two of the cations are joined head-to-head by the hydrogen bonds between them and the nitrate ions. These pairs of molecules form the characteristic chains extended along the $a$ axis.

## Experimental

To 50 ml of dry ether containing 0.5 g of $\mathrm{AlCl}_{3}$ was added 2-heptanone ( $10 \mathrm{~g}, 87 \mathrm{mmol}$ ). The solution was cooled and 4.5 ml $(0.087 \mathrm{~mol})$ of bromine was added dropwise. HBr and ether were evaporated under a stream of dry nitrogen at 313 K . To the 6.7 g ( 88 mmol ) of brown residue thiourea in 65 ml of water was added. The solution was warmed in a boiling water bath for 3 h . To the solution was then added 2 g of charcoal and $3.6 \mathrm{~g}(45 \mathrm{mmol})$ of NaOH . The solution was extracted with diethyl ether, diluted with petroleum ether and dried with $\mathrm{MgSO}_{4}$. Crystals suitable for X-ray

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Figure 1
The molecular structure of the title compound. Displacement ellipsoids are drawn at the $50 \%$ probability level.


Figure 2
The crystal packing of the title compound. Displacement ellipsoids are drawn at the $50 \%$ probability level. [Symmetry codes: $(A) x, y, z ;(B)$ $1-x, y, z ;(C) 1-x,-y,-z ;(D)-x,-y,-z$.]
diffraction studies were grown by slow evaporation from $n$-heptane. $9.2 \mathrm{~g}(62.2 \%)$ of 2-amino-5-butyl-4-methylthiazole was obtained as light yellow flakes melting at $352-353 \mathrm{~K}$. The title compound was prepared by dissolving $1.7 \mathrm{~g}(0.01 \mathrm{~mol})$ of 2-amino-5-butyl-4methylthiazole in 10 ml of methanol. Then, $0.76 \mathrm{ml}(0.011 \mathrm{~mol})$ of a $65 \%$ water solution of $\mathrm{HNO}_{3}$ was added. $1.91 \mathrm{~g}(82 \%)$ of the product was filtered off and dried in a vacuum (m.p. 365-367 K). The products were recrystallized from methanol.

## Crystal data

$$
\begin{aligned}
& \mathrm{C}_{8} \mathrm{H}_{15} \mathrm{~N}_{3} \mathrm{O}_{3} \mathrm{~S} \\
& M_{r}=233.29 \\
& \text { Triclinic, } P \overline{1} \\
& a=7.486(2) \AA \\
& b=8.918(2) \AA \\
& c=9.240(2) \AA \\
& \alpha=76.26(3)^{\circ} \\
& \beta=87.15(3)^{\circ} \\
& \gamma=75.08(3)^{\circ} \\
& V=579.0(2) \AA^{\circ} \\
& Z=2
\end{aligned}
$$

## Data collection

| Oxford Diffraction Xcalibur | $R_{\text {int }}=0.056$ |
| :--- | :--- |
| $\quad$ diffractometer | $\theta_{\max }=28.0^{\circ}$ |
| $\omega$ scans | $h=-10 \rightarrow 8$ |
| 4395 measured reflections | $k=-12 \rightarrow 11$ |
| 2712 independent reflections | $l=-12 \rightarrow 11$ |

2712 independent reflections 2016 reflections with $I>2 \sigma(I)$

## Refinement

Refinement on $F^{2}$
$R\left[F^{2}>2 \sigma\left(F^{2}\right)\right]=0.047$
$w R\left(F^{2}\right)=0.131$
H atoms treated by a mixture of independent and constrained
$S=0.98$ refinement

2712 reflections
145 parameters
$w=1 /\left[\sigma^{2}\left(F_{o}{ }^{2}\right)+(0.0762 P)^{2}\right]$
where $P=\left(F_{o}{ }^{2}+2 F_{c}^{2}\right) / 3$
$(\Delta / \sigma)_{\max }<0.001$
$\Delta \rho_{\text {max }}=0.40 \mathrm{e}^{\mathrm{A}}{ }^{-3}$
$\Delta \rho_{\text {min }}=-0.43 \mathrm{e}^{-3}$

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

| $\mathrm{C} 2-\mathrm{S} 1-\mathrm{C} 5$ | $90.9(1)$ | $\mathrm{C} 8-\mathrm{C} 5-\mathrm{S} 1$ | $121.7(1)$ |
| :--- | ---: | :--- | ---: |
| $\mathrm{C} 4-\mathrm{C} 5-\mathrm{C} 8$ | $127.3(2)$ |  |  |
| $\mathrm{C} 4-\mathrm{C} 5-\mathrm{C} 8-\mathrm{C} 9$ | $-175.5(2)$ | $\mathrm{S} 1-\mathrm{C} 5-\mathrm{C} 8-\mathrm{C} 9$ | $5.5(3)$ |

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

| $D-\mathrm{H} \cdots A$ | $D-\mathrm{H}$ | $\mathrm{H} \cdots A$ | $D \cdots A$ | $D-\mathrm{H} \cdots A$ |
| :--- | :--- | :--- | :--- | :--- |
| N6-H6A $\cdots$ O13 | $0.81(3)$ | $2.10(3)$ | $2.901(3)$ | $171(2)$ |
| N3-H3 $\cdots$ O14 | $0.92(3)$ | $2.36(3)$ | $3.090(3)$ | $135(2)$ |
| N3-H3 $A \cdots$ O15 | $0.92(3)$ | $1.97(3)$ | $2.849(3)$ | $157(2)$ |
| N6-H6B $\quad$ O14 | $0.90(3)$ | $2.25(3)$ | $3.045(3)$ | $146(2)$ |
| N6-H6B $\cdots$ O14 |  |  |  |  |

Symmetry codes: (i) $x-1, y, z$; (ii) $1-x,-y,-z$.
H atoms bonded to C atoms were refined freely using a riding model. The coordinates of the H atoms bonded to N atoms were refined.

Data collection: CrysAlisCCD (Oxford Diffraction, 2002); cell refinement: CrysAlisCCD; data reduction: CrysAlisRED (Oxford Diffraction, 2002); program(s) used to solve structure: SHELXS97 (Sheldrick, 1997); program(s) used to refine structure: SHELXL97 (Sheldrick, 1997); molecular graphics: SHELXTL (Sheldrick, 1990); software used to prepare material for publication: SHELXL97.

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