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#### **Key indicators**

Single-crystal X-ray study T = 292 KMean  $\sigma(\text{C-C}) = 0.002 \text{ Å}$  R factor = 0.046 wR factor = 0.133 Data-to-parameter ratio = 13.4

For details of how these key indicators were automatically derived from the article, see http://journals.iucr.org/e.

# 1,1'-Methylenebis(4-methyl-1*H*-triazolium) dipicrate

In the crystal structure of the title compound,  $C_7H_{12}N_6^{2+}$ .-2C<sub>6</sub>H<sub>2</sub>N<sub>3</sub>O<sub>7</sub><sup>-</sup>, the asymmetric unit contains two picrate anions and one 1,1'-methylenebis(4-methyl-1*H*-triazolium) dication. The dihedral angle between the triazole rings of the dication is 76.1 (2)°,

### Comment

Ionic liquids have attracted considerable interest as substitutes for volatile organic solvents in synthetic chemistry (Welton, 1999). Organic ionic salts with lower melting points, such as triazolium nitrate or perchlorate, are a new class of energetic materials because of their 'green chemistry' properties (Singh *et al.*, 2006). 1,2,4-Triazolium picrate has been reported as an energetic material (Jin *et al.*, 2006). In this report, we present the crystal structure of the title triazolium picrate, (I).



The asymmetric unit of (I) contains two picrate anions and one 1,1'-methylenebis(4-methyltriazolium) dication (Fig. 1). Within the dication, the dihedral angle between the triazole rings is 76.1 (2)°. Weak  $C-H\cdots N$  hydrogen bonding occurs between dications, and weak  $C-H\cdots O$  hydrogen bonding occurs between anions and dications (Table 1).

### Experimental

Compound (I) was synthesized according to a literature method (Jin *et al.*, 2005). Single crystals were obtained by recrystallization from an acetonitrile solution.

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# organic papers

Crystal data

 $C_7H_{12}N_6^{2+}\cdot 2C_6H_2N_3O_7^{-1}$   $M_r = 636.44$ Triclinic,  $P\overline{1}$  a = 7.0658 (5) Å b = 8.8036 (6) Å c = 20.5831 (14) Å  $\alpha = 96.587$  (1)°  $\beta = 90.122$  (1)°  $\gamma = 94.255$  (1)°

### Data collection

Bruker SMART CCD area-detector diffractometer  $\varphi$  and  $\omega$  scans Absorption correction: none 14366 measured reflections

### Refinement

Refinement on $F^2$	H-atom parameters constrained
$R[F^2 > 2\sigma(F^2)] = 0.046$	$w = 1/[\sigma^2(F_0^2) + (0.0674P)^2]$
$wR(F^2) = 0.133$	where $P = (F_0^2 + 2F_c^2)/3$
S = 1.08	$(\Delta/\sigma)_{\rm max} < 0.001$
5485 reflections	$\Delta \rho_{\rm max} = 0.39 \ {\rm e} \ {\rm \AA}^{-3}$
408 parameters	$\Delta \rho_{\rm min} = -0.25 \text{ e } \text{\AA}^{-3}$

V = 1268.33 (15) Å<sup>3</sup>

 $D_x = 1.666 \text{ Mg m}^{-3}$ 

 $0.30 \times 0.20 \times 0.20$  mm

5485 independent reflections

4435 reflections with  $I > 2\sigma(I)$ 

Mo  $K\alpha$  radiation

 $\mu = 0.14 \text{ mm}^{-1}$ 

T = 292 (2) K

Block, yellow

 $R_{\rm int} = 0.046$ 

 $\theta_{\rm max} = 27.0^{\circ}$ 

Z = 2

### Table 1

Hydrogen-bond geometry (Å, °).

$D - H \cdot \cdot \cdot A$	D-H	$H \cdot \cdot \cdot A$	$D \cdots A$	$D - \mathbf{H} \cdots A$
$C1-H1B\cdots O7^{i}$	0.96	2.31	3.217 (3)	157
$C2-H2\cdots O1^{i}$	0.93	2.42	3.286 (2)	156
C3-H3···O8	0.93	2.17	2.914 (2)	136
C3-H3···O14	0.93	2.35	3.066 (2)	133
$C4-H4A\cdots O8$	0.97	2.41	3.091 (2)	127
$C4-H4A\cdots O13^{ii}$	0.97	2.53	3.211 (2)	127
$C5-H5\cdots O2$	0.93	2.39	2.911 (2)	115
$C5-H5\cdots O8$	0.93	2.48	3.138 (2)	127
C6-H6···O1 <sup>iii</sup>	0.93	2.47	3.350 (2)	159
C6-H6···O2 <sup>iii</sup>	0.93	2.41	3.042 (2)	125
$C7-H7A\cdots N2^{iv}$	0.96	2.62	3.467 (2)	148
C7−H7C···O1	0.96	2.51	3.429 (3)	161

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

Methyl H atoms were placed in calculated positions, with C-H = 0.96 Å, and the torsion angles were refined to fit the electron density, with  $U_{iso}(H) = 1.5U_{eq}(C)$ . Other H atoms were positioned geome-





trically, with C-H = 0.93 (aromatic) and 0.97 Å (methylene), and refined using a riding model with  $U_{iso}(H) = 1.2U_{eq}(C)$ .

Data collection: *SMART* (Bruker, 2001); cell refinement: *SAINT-Plus* (Bruker, 2001); data reduction: *SAINT-Plus*; program(s) used to solve structure: *SHELXS97* (Sheldrick, 1997); program(s) used to refine structure: *SHELXL97* (Sheldrick, 1997); molecular graphics: *SHELXTL* (Bruker, 1997); software used to prepare material for publication: *SHELXTL*.

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