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Pyridine-3-carboxamidinium chloride

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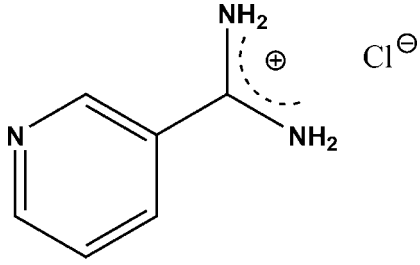
Key indicators: single-crystal X-ray study; T = 293 K; mean σ(C–C) = 0.002 Å; R factor = 0.029; wR factor = 0.071; data-to-parameter ratio = 17.7.

The title compound, $C_6H_8N_3^+\cdot Cl^-$, crystallizes with two formula units in the asymmetric unit. The cations are non-planar with the $-C(NH_2)_2$ groups twisted relative to the ring planes by 36.7 (3) and 37.8 (3)°. The cations are linked into chains through N—H···N hydrogen bonds. N—H···Cl hydrogen bonds link the chains into a three-dimensional network.

Related literature

For structures of pyridine-carboxamidinium chlorides, see: Fan *et al.* (2009); Chen *et al.* (2010).



Experimental

Crystal data

$C_6H_8N_3^+\cdot Cl^-$
M_r = 157.60
Orthorhombic, $Pna2_1$
a = 10.9485 (7) Å
b = 33.1581 (14) Å
c = 4.1488 (5) Å
V = 1506.1 (2) Å^3
Z = 8
Mo $K\alpha$ radiation
μ = 0.43 mm^{-1}
T = 293 K
0.40 × 0.35 × 0.17 mm

Data collection

Rigaku R-AXIS RAPID diffractometer
Absorption correction: multi-scan (*ABSCOR*; Higashi, 1995)
T_{min} = 0.845, T_{max} = 0.930
14201 measured reflections
3213 independent reflections
2953 reflections with $I > 2\sigma(I)$
R_{int} = 0.033

Refinement

$R[F^2 > 2\sigma(F^2)]$ = 0.029
$wR(F^2)$ = 0.071
S = 1.00
3213 reflections
182 parameters
1 restraint
H-atom parameters constrained
$\Delta\rho_{max}$ = 0.18 e Å^{-3}
$\Delta\rho_{min}$ = −0.18 e Å^{-3}
Absolute structure: Flack (1983), 1246 Friedel pairs
Flack parameter: 0.0019 (5)

Table 1
Hydrogen-bond geometry (Å, °).

D—H···A	D—H	H···A	D···A	D—H···A
N3—H3B···N4	0.86	2.07	2.880 (2)	157
N5—H5B···Cl2	0.86	2.29	3.1403 (15)	170
N6—H6B···Cl1	0.86	2.36	3.1562 (16)	155
N2—H2A···N1^{i}	0.86	2.22	2.990 (2)	149
N2—H2B···Cl2^{ii}	0.86	2.31	3.1452 (13)	165
N3—H3A···Cl2^{iii}	0.86	2.27	3.1040 (16)	164
N5—H5A···Cl1^{iv}	0.86	2.46	3.2373 (16)	150
N6—H6A···Cl1^{iv}	0.86	2.41	3.2013 (16)	152

Symmetry codes: (i) $x+\frac{1}{2}, -y+\frac{1}{2}, z+1$; (ii) $x-\frac{1}{2}, -y+\frac{1}{2}, z+1$; (iii) $x, y, z+1$; (iv) $-x+2, -y+1, z-\frac{1}{2}$.

Data collection: *PROCESS-AUTO* (Rigaku, 2006); cell refinement: *PROCESS-AUTO*; data reduction: *CrystalStructure* (Rigaku/MSC, 2007); program(s) used to solve structure: *SHELXS97* (Sheldrick, 2008); program(s) used to refine structure: *SHELXL97* (Sheldrick, 2008); molecular graphics: *ORTEP-3 for Windows* (Farrugia, 1997); software used to prepare material for publication: *WinGX* (Farrugia, 1999).

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Supplementary data and figures for this paper are available from the IUCr electronic archives (Reference: GK2326).

References

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supporting information

Acta Cryst. (2011). E**67**, o781 [doi:10.1107/S1600536811002704]

Pyridine-3-carboxamidinium chloride

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S1. Comment

During the last decade, much interest has been focused on the synthesis of pyridine carboximidamidate derivatives because of their very potent antibacterial activities, interesting biological properties and applications in coordination chemistry.

The title compound is an organic salt which crystallizes with two formula units in the asymmetric unit (Fig. 1). The molecules are nonplanar with the $C(NH_2)_2$ groups twisted out of the ring planes with the twist angles of 36.7 (3) ° and 37.8 (3) °. The bond lengths and angles in title compound are in the normal ranges comparable with those in the related structure (Fan *et al.*, 2009).

In the crystal structure, pyridine-3-carboximidamidate cations are linked by N—H···N hydrogen bonds to form one-dimensional supramolecular molecular chain (Fig. 2). Cl1 ion bridges two pyridine carboximidamidate cations through N—H···Cl hydrogen bonding and Cl2 ion triple-bridges three cations through N—H···Cl hydrogen bonding. Thus, one-dimensional chains of cations are linked by N—H···Cl interactions into a three-dimensional network (Fig. 3).

S2. Experimental

To a solution of sodium methoxide (5.15 mmol) in methnol (50 ml) was added 3-cyanopyridine (5.2 g, 4.99 mmol). The mixture was stirred at room temperature for 2 h. Then ammonium chloride (2.9 g, 5.42 mmol) was slowly added to the resulting solution and the mixture was stirred at room temperature for 48 h. The resulting suspension was filtered and the solvent was removed from the filtrate under reduced pressure. Purification by washing with diethylether gave 3-amidino-pyridine chloride (5.74 g, 94%) as an off-white solid. Block-shaped crystals suitable for X-ray diffraction were obtained by recrystallization from from ethanol.

S3. Refinement

H atoms were placed in calculated positions and treated as riding on their parent atoms (C—H = 0.93 Å, N-H = 0.86 Å) and U_{iso}(H) = 1.2U_{eq}(C,N).

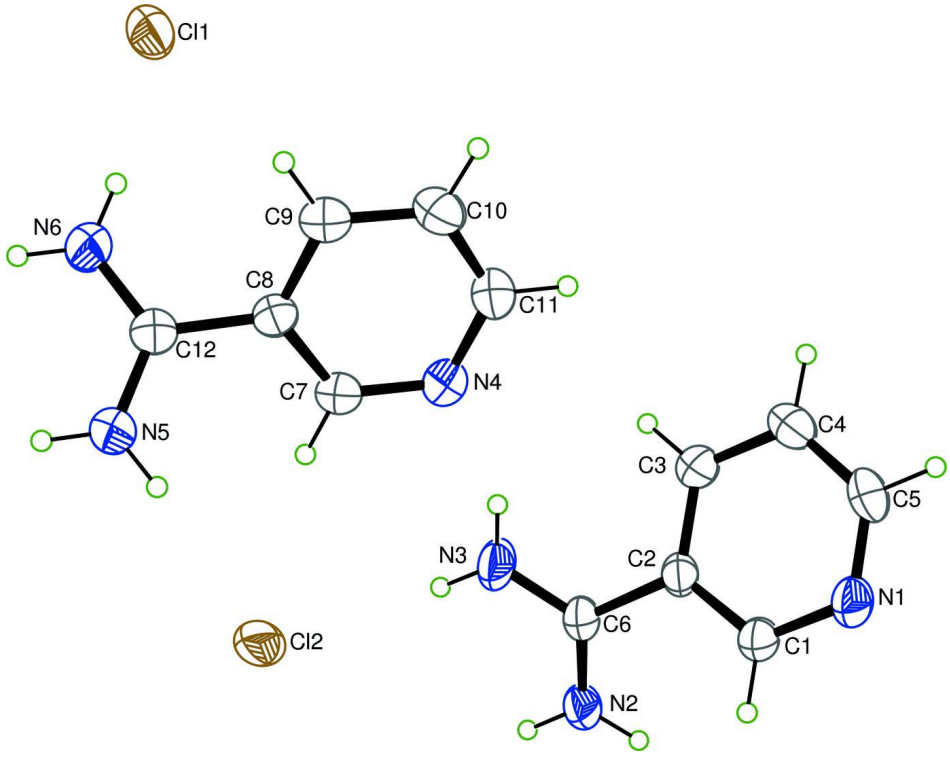

Figure 1

Structure of the title compound showing 30% probability displacement ellipsoids.

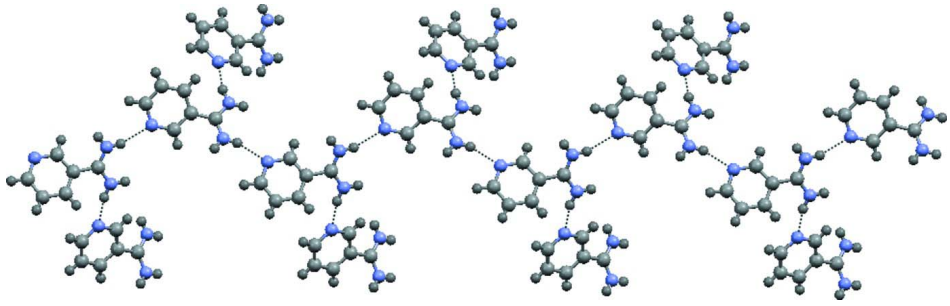

Figure 2

View of one-dimensional supramolecular chain in the title compound.

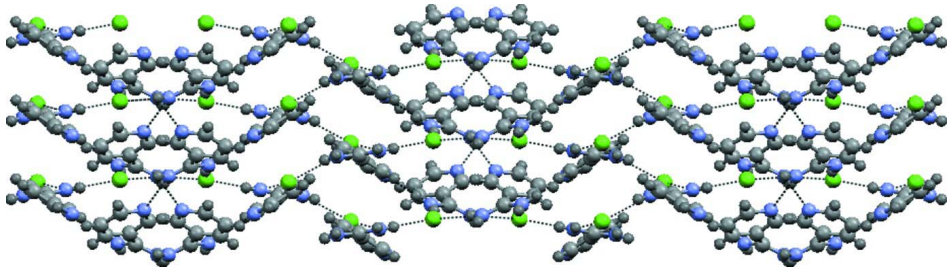

Figure 3

The crystal packing of the title compound shown down the *x* axis. Hydrogen bonds are shown with dashed lines.

Pyridine-3-carboxamidinium chloride

Crystal data

$C_6H_8N_3^+\cdot Cl^-$
M_r = 157.60
Orthorhombic, $Pna2_1$
Hall symbol: P 2c -2n
a = 10.9485 (7) Å
b = 33.1581 (14) Å
c = 4.1488 (5) Å
V = 1506.1 (2) Å^3
Z = 8
$F(000)$ = 656
D_x = 1.390 Mg m^{-3}
Mo $K\alpha$ radiation, λ = 0.71073 Å
Cell parameters from 3213 reflections
θ = 3.1–27.5°
μ = 0.43 mm^{-1}
T = 293 K
Block, colorless
0.40 × 0.35 × 0.17 mm

Data collection

Rigaku R-AXIS RAPID diffractometer
Radiation source: fine-focus sealed tube
Graphite monochromator
Detector resolution: 10.0 pixels mm^{-1}
ω scans
Absorption correction: multi-scan (*ABSCOR*; Higashi, 1995)
T_{min} = 0.845, T_{max} = 0.930
14201 measured reflections
3213 independent reflections
2953 reflections with $I > 2\sigma(I)$
R_{int} = 0.033
θ_{max} = 27.5°, θ_{min} = 3.1°
h = −14→14
k = −42→42
l = −4→5

Refinement

Refinement on F^2
Least-squares matrix: full
$R[F^2 > 2\sigma(F^2)]$ = 0.029
$wR(F^2)$ = 0.071
S = 1.00
3213 reflections
182 parameters
1 restraint
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
$w = 1/[\sigma^2(F_o^2) + (0.0334P)^2 + 0.3352P]$ where $P = (F_o^2 + 2F_c^2)/3$
$(\Delta/\sigma)_{max}$ = 0.002
$\Delta\rho_{max}$ = 0.18 e Å^{-3}
$\Delta\rho_{min}$ = −0.18 e Å^{-3}
Extinction correction: *SHELXL97* (Sheldrick, 2008), $Fc^*=kFc[1+0.001xFc^2\lambda^3/\sin(2\theta)]^{-1/4}$
Extinction coefficient: 0.061 (2)
Absolute structure: Flack (1983), 1246 Friedel pairs
Absolute structure parameter: 0.0019 (5)

Special details

Geometry. All e.s.d.'s (except the e.s.d. in the dihedral angle between two l.s. planes) are estimated using the full covariance matrix. The cell e.s.d.'s are taken into account individually in the estimation of e.s.d.'s in distances, angles and torsion angles; correlations between e.s.d.'s in cell parameters are only used when they are defined by crystal symmetry. An approximate (isotropic) treatment of cell e.s.d.'s is used for estimating e.s.d.'s involving l.s. planes.
Refinement. Refinement of F^2 against ALL reflections. The weighted R-factor wR and goodness of fit S are based on F^2, conventional R-factors R are based on F, with F set to zero for negative F^2. The threshold expression of $F^2 > \sigma(F^2)$ is used only for calculating R-factors(gt) *etc*. and is not relevant to the choice of reflections for refinement. R-factors based on 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 (Å^2)

	x	y	z	U_{iso}*/U_{eq}
Cl1	0.80408 (4)	0.549620 (13)	0.75756 (15)	0.05193 (15)

Cl2	0.82982 (4)	0.318730 (12)	0.22533 (15)	0.04986 (15)
N2	0.49341 (12)	0.25905 (4)	1.1812 (4)	0.0384 (3)
H2A	0.5635	0.2541	1.2661	0.046*
H2B	0.4372	0.2409	1.1835	0.046*
N1	0.17495 (13)	0.28055 (4)	0.6167 (4)	0.0392 (4)
N4	0.56768 (13)	0.39520 (4)	0.6649 (4)	0.0427 (4)
C3	0.30154 (15)	0.34172 (5)	0.9320 (5)	0.0359 (4)
H3	0.3444	0.3622	1.0348	0.043*
C2	0.35090 (13)	0.30307 (5)	0.9096 (4)	0.0294 (3)
N3	0.55529 (14)	0.32232 (5)	1.0408 (6)	0.0540 (5)
H3A	0.6261	0.3181	1.1239	0.065*
H3B	0.5390	0.3451	0.9521	0.065*
C1	0.28379 (13)	0.27351 (4)	0.7498 (5)	0.0321 (3)
H1	0.3164	0.2477	0.7352	0.038*
N5	0.93657 (14)	0.40511 (4)	0.3554 (5)	0.0515 (5)
H5A	1.0143	0.4075	0.3329	0.062*
H5B	0.9030	0.3817	0.3431	0.062*
N6	0.91881 (14)	0.47268 (4)	0.4274 (5)	0.0511 (5)
H6A	0.9965	0.4754	0.4053	0.061*
H6B	0.8739	0.4935	0.4622	0.061*
C7	0.68741 (16)	0.40022 (5)	0.6164 (5)	0.0376 (4)
H7	0.7409	0.3809	0.6971	0.045*
C12	0.86919 (16)	0.43713 (5)	0.4083 (5)	0.0366 (4)
C6	0.47248 (14)	0.29402 (5)	1.0493 (5)	0.0328 (3)
C5	0.12821 (16)	0.31783 (5)	0.6479 (5)	0.0448 (5)
H5	0.0512	0.3228	0.5624	0.054*
C8	0.73540 (15)	0.43301 (4)	0.4505 (4)	0.0332 (4)
C4	0.18844 (16)	0.34905 (5)	0.7998 (5)	0.0440 (5)
H4	0.1533	0.3745	0.8126	0.053*
C9	0.65596 (16)	0.46198 (5)	0.3279 (5)	0.0400 (4)
H9	0.6853	0.4846	0.2198	0.048*
C11	0.49241 (17)	0.42297 (5)	0.5413 (5)	0.0432 (4)
H11	0.4089	0.4196	0.5718	0.052*
C10	0.53172 (17)	0.45609 (5)	0.3720 (5)	0.0433 (4)
H10	0.4757	0.4744	0.2879	0.052*

Atomic displacement parameters (Å²)

	U^{11}	U^{22}	U^{33}	U^{12}	U^{13}	U^{23}
Cl1	0.0348 (2)	0.0464 (2)	0.0746 (4)	0.00362 (16)	0.0023 (3)	−0.0112 (3)
Cl2	0.0345 (2)	0.03388 (19)	0.0812 (4)	0.00596 (15)	−0.0204 (2)	−0.0130 (2)
N2	0.0260 (6)	0.0361 (7)	0.0531 (10)	−0.0005 (5)	−0.0083 (6)	0.0086 (7)
N1	0.0318 (7)	0.0404 (7)	0.0454 (9)	−0.0045 (6)	−0.0117 (6)	0.0005 (7)
N4	0.0342 (7)	0.0342 (6)	0.0597 (11)	−0.0032 (5)	0.0011 (7)	0.0042 (7)
C3	0.0339 (8)	0.0309 (8)	0.0428 (10)	−0.0015 (6)	−0.0054 (7)	−0.0025 (8)
C2	0.0230 (7)	0.0320 (7)	0.0332 (8)	−0.0020 (6)	−0.0022 (6)	0.0043 (7)
N3	0.0291 (7)	0.0421 (8)	0.0907 (14)	−0.0092 (6)	−0.0213 (8)	0.0190 (9)
C1	0.0292 (7)	0.0307 (7)	0.0363 (9)	−0.0015 (5)	−0.0041 (8)	0.0015 (8)

N5	0.0326 (7)	0.0328 (7)	0.0890 (15)	0.0001 (6)	0.0051 (8)	−0.0079 (8)
N6	0.0359 (8)	0.0336 (7)	0.0837 (14)	−0.0037 (6)	0.0103 (8)	−0.0099 (9)
C7	0.0348 (8)	0.0285 (7)	0.0495 (11)	0.0022 (6)	−0.0024 (7)	0.0008 (8)
C12	0.0347 (8)	0.0318 (7)	0.0433 (10)	0.0002 (6)	0.0015 (7)	−0.0025 (8)
C6	0.0239 (7)	0.0348 (8)	0.0396 (9)	−0.0021 (6)	−0.0046 (7)	0.0004 (7)
C5	0.0281 (8)	0.0510 (10)	0.0552 (13)	0.0046 (7)	−0.0141 (8)	0.0008 (9)
C8	0.0324 (8)	0.0276 (7)	0.0397 (10)	−0.0003 (6)	−0.0010 (7)	−0.0046 (7)
C4	0.0364 (8)	0.0391 (8)	0.0565 (13)	0.0098 (7)	−0.0067 (8)	−0.0020 (9)
C9	0.0413 (9)	0.0313 (7)	0.0473 (12)	0.0021 (7)	−0.0011 (8)	0.0036 (7)
C11	0.0306 (8)	0.0410 (9)	0.0580 (12)	−0.0010 (7)	−0.0038 (8)	−0.0026 (9)
C10	0.0400 (9)	0.0383 (9)	0.0516 (11)	0.0076 (7)	−0.0066 (8)	0.0031 (8)

Geometric parameters (Å, º)

N2—C6	1.303 (2)	N5—H5A	0.8600
N2—H2A	0.8600	N5—H5B	0.8600
N2—H2B	0.8600	N6—C12	1.300 (2)
N1—C1	1.334 (2)	N6—H6A	0.8600
N1—C5	1.344 (2)	N6—H6B	0.8600
N4—C7	1.337 (2)	C7—C8	1.390 (2)
N4—C11	1.338 (2)	C7—H7	0.9300
C3—C4	1.376 (2)	C12—C8	1.482 (2)
C3—C2	1.394 (2)	C5—C4	1.380 (3)
C3—H3	0.9300	C5—H5	0.9300
C2—C1	1.393 (2)	C8—C9	1.392 (2)
C2—C6	1.483 (2)	C4—H4	0.9300
N3—C6	1.305 (2)	C9—C10	1.386 (3)
N3—H3A	0.8600	C9—H9	0.9300
N3—H3B	0.8600	C11—C10	1.373 (3)
C1—H1	0.9300	C11—H11	0.9300
N5—C12	1.311 (2)	C10—H10	0.9300
C6—N2—H2A	120.0	C8—C7—H7	118.6
C6—N2—H2B	120.0	N6—C12—N5	120.60 (16)
H2A—N2—H2B	120.0	N6—C12—C8	119.31 (15)
C1—N1—C5	117.47 (14)	N5—C12—C8	120.08 (14)
C7—N4—C11	117.45 (16)	N2—C6—N3	121.94 (15)
C4—C3—C2	118.99 (16)	N2—C6—C2	120.16 (14)
C4—C3—H3	120.5	N3—C6—C2	117.88 (15)
C2—C3—H3	120.5	N1—C5—C4	123.50 (16)
C1—C2—C3	118.31 (14)	N1—C5—H5	118.2
C1—C2—C6	121.15 (14)	C4—C5—H5	118.2
C3—C2—C6	120.54 (14)	C7—C8—C9	118.99 (16)
C6—N3—H3A	120.0	C7—C8—C12	120.30 (15)
C6—N3—H3B	120.0	C9—C8—C12	120.71 (15)
H3A—N3—H3B	120.0	C3—C4—C5	118.66 (15)
N1—C1—C2	123.03 (14)	C3—C4—H4	120.7
N1—C1—H1	118.5	C5—C4—H4	120.7

C2—C1—H1	118.5	C10—C9—C8	117.87 (16)
C12—N5—H5A	120.0	C10—C9—H9	121.1
C12—N5—H5B	120.0	C8—C9—H9	121.1
H5A—N5—H5B	120.0	N4—C11—C10	123.61 (17)
C12—N6—H6A	120.0	N4—C11—H11	118.2
C12—N6—H6B	120.0	C10—C11—H11	118.2
H6A—N6—H6B	120.0	C11—C10—C9	119.20 (17)
N4—C7—C8	122.86 (16)	C11—C10—H10	120.4
N4—C7—H7	118.6	C9—C10—H10	120.4
C4—C3—C2—C1	0.8 (3)	N4—C7—C8—C12	179.21 (18)
C4—C3—C2—C6	−179.98 (18)	N6—C12—C8—C7	−141.6 (2)
C5—N1—C1—C2	−1.7 (3)	N5—C12—C8—C7	37.8 (3)
C3—C2—C1—N1	0.2 (3)	N6—C12—C8—C9	37.8 (3)
C6—C2—C1—N1	−178.99 (18)	N5—C12—C8—C9	−142.8 (2)
C11—N4—C7—C8	1.2 (3)	C2—C3—C4—C5	−0.4 (3)
C1—C2—C6—N2	−36.7 (3)	N1—C5—C4—C3	−1.2 (3)
C3—C2—C6—N2	144.10 (19)	C7—C8—C9—C10	−1.4 (3)
C1—C2—C6—N3	144.72 (19)	C12—C8—C9—C10	179.20 (17)
C3—C2—C6—N3	−34.5 (3)	C7—N4—C11—C10	−0.6 (3)
C1—N1—C5—C4	2.1 (3)	N4—C11—C10—C9	−1.0 (3)
N4—C7—C8—C9	−0.2 (3)	C8—C9—C10—C11	1.9 (3)

Hydrogen-bond geometry (Å, º)

D—H···*A*	*D*—H	H···*A*	*D*···*A*	*D*—H···*A*
N3—H3*B*···N4	0.86	2.07	2.880 (2)	157
N5—H5*B*···Cl2	0.86	2.29	3.1403 (15)	170
N6—H6*B*···Cl1	0.86	2.36	3.1562 (16)	155
N2—H2*A*···N1^{i}	0.86	2.22	2.990 (2)	149
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N3—H3*A*···Cl2^{iii}	0.86	2.27	3.1040 (16)	164
N5—H5*A*···Cl1^{iv}	0.86	2.46	3.2373 (16)	150
N6—H6*A*···Cl1^{iv}	0.86	2.41	3.2013 (16)	152

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