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Sodium 1-(4-chlorophenyl)-5-methyl-1*H*-1,2,3-triazole-4-carboxylate

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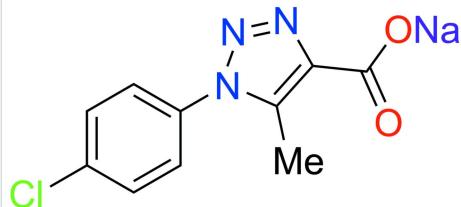
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In the title molecular salt, $\text{Na}^+\cdot\text{C}_{10}\text{H}_7\text{ClN}_3\text{O}_2^-$, the dihedral angles between the planes of adjacent chlorophenyl, methyltriazole and carboxylate groups of the anion are 50.2 (1) and 9.0 (3) $^\circ$, respectively. The shortest distance between sodium cations is 4.0595 (9) Å. The sodium cation is coordinated by two N atoms and three O atoms, generating layers of ions lying parallel to the *bc* plane.

3D view



Chemical scheme



Structure description

1,2,3-Triazole-4-carboxylic acids are important precursors for various biologically active compounds that act as xanthine oxidase inhibitors (Zhang *et al.*, 2017; Ojha *et al.*, 2017) and antibacterial (Maji & Haldar, 2017) and antitubercular (Kamal *et al.*, 2013) agents. As part of our studies in this area, we now describe the synthesis and structure of the title salt.

The asymmetric unit comprises a Na^+ cation and a $\text{C}_{10}\text{H}_7\text{ClN}_3\text{O}_2^-$ anion (Fig. 1). The anion consists of chlorophenyl, methyltriazolyl and carboxylate groups, and the twist angles between the planes through adjacent groups are 50.2 (1) and 9.0 (3) $^\circ$, respectively. The shortest distance between sodium cations is 4.0595 (9) Å and each cation is coordinated by two N atoms, with an average $\text{Na} - \text{N}$ distance of 2.5654 (6) Å, and by three O atoms, with an average $\text{Na} \cdots \text{O}$ distance of 2.37 (4) Å. The resulting coordination polyhedron is a very distorted trigonal bipyramidal with one N atom and one O atom in the axial sites. The extended structure generates layers of ions parallel to the *bc* plane (Fig. 2). A short C–H \cdots N contact is also observed (Table 1).

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

$D-\text{H}\cdots A$	$D-\text{H}$	$\text{H}\cdots A$	$D\cdots A$	$D-\text{H}\cdots A$
C4—H4A···N2 ⁱ	0.96	2.52	3.432 (3)	159

Symmetry code: (i) $x, y - 1, z$.

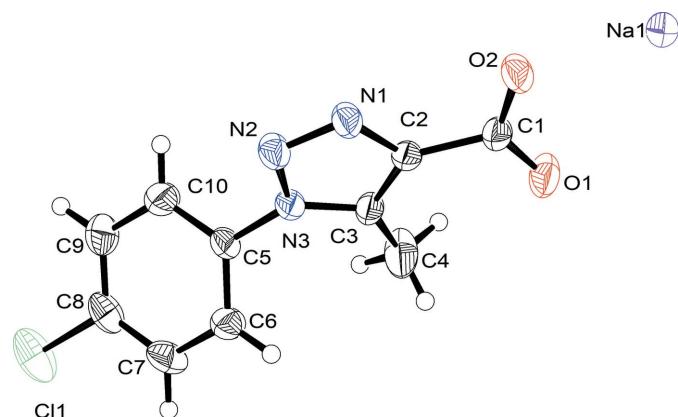


Figure 1

The asymmetric unit of the title salt, showing 50% probability displacement ellipsoids.

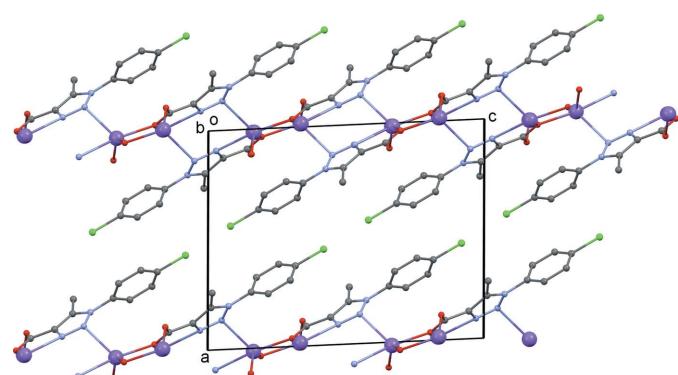


Figure 2

The crystal packing of the title salt, viewed down the b axis.

Synthesis and crystallization

The title compound was synthesized from the reaction of 1-(4-chlorophenyl)-5-methyl-1*H*-1,2,3-triazole-4-carboxylic acid and sodium hydroxide (10%) in ethanol under reflux for 4 h. The crude product obtained after work-up was recrystallized from dimethylformamide solution to give colourless blocks.

Refinement

Crystal data, data collection and structure refinement details are summarized in Table 2.

Table 2
Experimental details.

Crystal data	$\text{Na}^+\cdot\text{C}_{10}\text{H}_7\text{ClN}_3\text{O}_2^-$
Chemical formula	$\text{Na}^+\cdot\text{C}_{10}\text{H}_7\text{ClN}_3\text{O}_2^-$
M_r	259.63
Crystal system, space group	Monoclinic, $P2_1/c$
Temperature (K)	293
a, b, c (\AA)	11.6462 (8), 6.3754 (4), 14.7003 (9)
β ($^\circ$)	92.711 (7)
V (\AA^3)	1090.26 (12)
Z	4
Radiation type	Mo $K\alpha$
μ (mm^{-1})	0.38
Crystal size (mm)	0.42 \times 0.16 \times 0.07
Data collection	Rigaku OD SuperNova Dual source diffractometer with an Atlas detector
Diffractometer	Gaussian (<i>CrysAlis PRO</i> ; Rigaku OD, 2015)
Absorption correction	8333, 2665, 1875
T_{\min}, T_{\max}	0.467, 1.000
No. of measured, independent and observed [$I > 2\sigma(I)$] reflections	8333, 2665, 1875
R_{int}	0.030
$(\sin \theta/\lambda)_{\max}$ (\AA^{-1})	0.696
Refinement	Rigaku OD SuperNova Dual source diffractometer with an Atlas detector
$R[F^2 > 2\sigma(F^2)], wR(F^2), S$	0.045, 0.115, 1.05
No. of reflections	2665
No. of parameters	155
H-atom treatment	H-atom parameters constrained
$\Delta\rho_{\max}, \Delta\rho_{\min}$ ($e \text{\AA}^{-3}$)	0.25, -0.40

Computer programs: *CrysAlis PRO* (Rigaku OD, 2015), *SHELXS97* (Sheldrick, 2008), *SHELXL2018* (Sheldrick, 2015), *ORTEP-3* for Windows and *WinGX* (Farrugia, 2012) and *CHEMDRAW Ultra* (CambridgeSoft, 2001).

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References

- CambridgeSoft (2001). *CHEMDRAW Ultra*. CambridgeSoft Corporation, Cambridge, Massachusetts, USA.
- Farrugia, L. J. (2012). *J. Appl. Cryst.* **45**, 849–854.
- Kamal, A., Hussaini, S. M. A., Faazil, S., Poornachandra, Y., Narendra Reddy, G., Kumar, C. G., Rajput, V. S., Rani, C., Sharma, R., Khan, I. A. & Jagadeesh Babu, N. (2013). *Bioorg. Med. Chem. Lett.* **23**, 6842–6846.
- Maji, K. & Haldar, D. R. (2017). *R. Soc. Open Sci.* **4**, 170684.
- Ojha, R., Singh, J., Ojha, A., Singh, H., Sharma, S. & Nepali, K. (2017). *Expert Opin. Ther. Pat.* **27**, 311–345.
- Rigaku OD (2015). *CrysAlis PRO*. Rigaku Oxford Diffraction, Yarnton, England.
- Sheldrick, G. M. (2008). *Acta Cryst. A* **64**, 112–122.
- Sheldrick, G. M. (2015). *Acta Cryst. C* **71**, 3–8.
- Zhang, T. J., Wu, Q. X., Li, S. Y., Wang, L., Sun, Q., Zhang, Y., Meng, F. H. & Gao, H. (2017). *Bioorg. Med. Chem. Lett.* **27**, 3812–3816

full crystallographic data

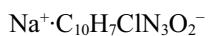
IUCrData (2018). **3**, x181127 [https://doi.org/10.1107/S2414314618011276]

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Sodium 1-(4-chlorophenyl)-5-methyl-1*H*-1,2,3-triazole-4-carboxylate

Crystal data



$M_r = 259.63$

Monoclinic, $P2_1/c$

$a = 11.6462 (8) \text{ \AA}$

$b = 6.3754 (4) \text{ \AA}$

$c = 14.7003 (9) \text{ \AA}$

$\beta = 92.711 (7)^\circ$

$V = 1090.26 (12) \text{ \AA}^3$

$Z = 4$

$F(000) = 528$

$D_x = 1.582 \text{ Mg m}^{-3}$

Mo $K\alpha$ radiation, $\lambda = 0.71073 \text{ \AA}$

Cell parameters from 2649 reflections

$\theta = 3.5\text{--}29.2^\circ$

$\mu = 0.38 \text{ mm}^{-1}$

$T = 293 \text{ K}$

Block, colourless

$0.42 \times 0.16 \times 0.07 \text{ mm}$

Data collection

Rigaku OD SuperNova Dual source diffractometer with an Atlas detector

ω scans

Absorption correction: gaussian

(CrysAlis PRO; Rigaku OD, 2015)

$T_{\min} = 0.467$, $T_{\max} = 1.000$

8333 measured reflections

2665 independent reflections

1875 reflections with $I > 2\sigma(I)$

$R_{\text{int}} = 0.030$

$\theta_{\max} = 29.7^\circ$, $\theta_{\min} = 3.2^\circ$

$h = -14 \rightarrow 15$

$k = -8 \rightarrow 8$

$l = -20 \rightarrow 20$

Refinement

Refinement on F^2

Least-squares matrix: full

$R[F^2 > 2\sigma(F^2)] = 0.045$

$wR(F^2) = 0.115$

$S = 1.05$

2665 reflections

155 parameters

0 restraints

Primary atom site location: structure-invariant direct methods

Hydrogen site location: inferred from neighbouring sites

H-atom parameters constrained

$w = 1/[\sigma^2(F_o^2) + (0.0446P)^2 + 0.2829P]$
where $P = (F_o^2 + 2F_c^2)/3$

$(\Delta/\sigma)_{\max} < 0.001$

$\Delta\rho_{\max} = 0.25 \text{ e \AA}^{-3}$

$\Delta\rho_{\min} = -0.40 \text{ e \AA}^{-3}$

Special details

Geometry. All esds (except the esd in the dihedral angle between two l.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.

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

	<i>x</i>	<i>y</i>	<i>z</i>	$U_{\text{iso}}^*/U_{\text{eq}}$
C1	0.08988 (16)	0.5137 (3)	0.64261 (13)	0.0306 (4)
C2	0.12619 (15)	0.6087 (3)	0.55520 (12)	0.0277 (4)
C3	0.19398 (17)	0.5260 (3)	0.48997 (13)	0.0304 (4)
C4	0.2601 (2)	0.3285 (3)	0.48670 (17)	0.0597 (8)
H4A	0.209664	0.215967	0.467891	0.090*
H4B	0.294116	0.298547	0.546045	0.090*
H4C	0.319562	0.342836	0.444051	0.090*
C5	0.25655 (16)	0.6837 (3)	0.34347 (13)	0.0302 (4)
C6	0.24624 (18)	0.5191 (3)	0.28293 (14)	0.0378 (5)
H6	0.201366	0.403343	0.296042	0.045*
C7	0.30294 (19)	0.5273 (4)	0.20265 (14)	0.0448 (6)
H7	0.297119	0.416527	0.161513	0.054*
C8	0.36800 (18)	0.7002 (4)	0.18405 (14)	0.0421 (5)
C9	0.3785 (2)	0.8660 (4)	0.24377 (15)	0.0481 (6)
H9	0.422407	0.982653	0.230039	0.058*
C10	0.32262 (19)	0.8562 (3)	0.32453 (15)	0.0429 (5)
H10	0.329600	0.965933	0.366091	0.052*
N1	0.09199 (14)	0.8041 (2)	0.52889 (11)	0.0337 (4)
N2	0.13405 (15)	0.8474 (2)	0.45008 (11)	0.0355 (4)
N3	0.19683 (13)	0.6783 (2)	0.42611 (10)	0.0292 (4)
O1	0.11449 (14)	0.3261 (2)	0.65583 (10)	0.0469 (4)
O2	0.03699 (12)	0.6316 (2)	0.69418 (9)	0.0406 (4)
Cl1	0.43802 (6)	0.70889 (13)	0.08201 (4)	0.0687 (3)
Na1	-0.01461 (7)	0.49886 (11)	0.83416 (5)	0.0360 (2)

Atomic displacement parameters (\AA^2)

	U^{11}	U^{22}	U^{33}	U^{12}	U^{13}	U^{23}
C1	0.0339 (11)	0.0277 (10)	0.0310 (10)	-0.0038 (8)	0.0085 (9)	0.0010 (8)
C2	0.0327 (10)	0.0239 (9)	0.0269 (9)	-0.0009 (7)	0.0067 (8)	-0.0022 (7)
C3	0.0376 (11)	0.0241 (9)	0.0303 (10)	-0.0014 (7)	0.0100 (9)	0.0016 (8)
C4	0.0875 (19)	0.0378 (12)	0.0569 (15)	0.0222 (12)	0.0378 (14)	0.0114 (11)
C5	0.0345 (10)	0.0332 (9)	0.0234 (9)	0.0004 (8)	0.0067 (8)	-0.0006 (8)
C6	0.0419 (12)	0.0384 (11)	0.0336 (11)	-0.0062 (9)	0.0081 (9)	-0.0057 (9)
C7	0.0498 (14)	0.0542 (13)	0.0310 (11)	0.0016 (10)	0.0075 (10)	-0.0134 (10)
C8	0.0330 (11)	0.0646 (14)	0.0293 (11)	0.0056 (10)	0.0081 (9)	0.0040 (10)
C9	0.0502 (14)	0.0527 (13)	0.0425 (13)	-0.0119 (11)	0.0142 (11)	0.0075 (11)
C10	0.0533 (14)	0.0382 (11)	0.0381 (12)	-0.0091 (10)	0.0119 (10)	-0.0050 (9)
N1	0.0462 (10)	0.0287 (8)	0.0271 (8)	0.0056 (7)	0.0107 (7)	0.0023 (7)
N2	0.0487 (10)	0.0299 (8)	0.0287 (9)	0.0090 (7)	0.0103 (8)	0.0025 (7)
N3	0.0363 (9)	0.0265 (8)	0.0253 (8)	0.0013 (6)	0.0079 (7)	-0.0016 (6)
O1	0.0652 (10)	0.0282 (7)	0.0494 (9)	0.0057 (7)	0.0242 (8)	0.0101 (7)
O2	0.0584 (9)	0.0334 (7)	0.0318 (7)	0.0055 (7)	0.0205 (7)	0.0019 (6)
Cl1	0.0569 (4)	0.1123 (6)	0.0390 (3)	0.0133 (4)	0.0240 (3)	0.0101 (3)
Na1	0.0500 (5)	0.0288 (4)	0.0300 (4)	-0.0048 (3)	0.0083 (4)	0.0008 (3)

Geometric parameters (\AA , ^\circ)

Na1—N1 ⁱ	2.5649 (16)	C4—H4C	0.9600
Na1—N2 ⁱⁱ	2.5658 (19)	C5—C6	1.377 (3)
Na1—O1 ⁱⁱⁱ	2.3964 (16)	C5—C10	1.378 (3)
Na1—O2	2.3299 (14)	C5—N3	1.429 (2)
Na1—O2 ⁱ	2.3902 (15)	C6—C7	1.380 (3)
Na1—Na1 ⁱ	4.0595 (9)	C6—H6	0.9300
Na1—Na1 ⁱⁱⁱ	4.0595 (9)	C7—C8	1.373 (3)
C1—O1	1.243 (2)	C7—H7	0.9300
C1—O2	1.250 (2)	C8—C9	1.376 (3)
C1—C2	1.499 (2)	C8—Cl1	1.742 (2)
C2—N1	1.359 (2)	C9—C10	1.382 (3)
C2—C3	1.375 (2)	C9—H9	0.9300
C3—N3	1.352 (2)	C10—H10	0.9300
C3—C4	1.478 (3)	N1—N2	1.308 (2)
C4—H4A	0.9600	N2—N3	1.358 (2)
C4—H4B	0.9600		
O1—C1—O2	126.89 (17)	N1—N2—N3	106.93 (14)
O1—C1—C2	116.76 (16)	N1—N2—Na1 ^{iv}	113.55 (12)
O2—C1—C2	116.34 (16)	N3—N2—Na1 ^{iv}	119.04 (12)
O1—C1—Na1	85.80 (11)	C3—N3—N2	111.00 (14)
O2—C1—Na1	41.15 (9)	C3—N3—C5	129.55 (15)
C2—C1—Na1	157.39 (12)	N2—N3—C5	119.42 (14)
N1—C2—C3	108.84 (15)	C1—O1—Na1 ⁱ	137.58 (14)
N1—C2—C1	121.58 (15)	C1—O2—Na1	118.17 (12)
C3—C2—C1	129.59 (17)	C1—O2—Na1 ⁱⁱⁱ	122.42 (11)
N3—C3—C2	104.06 (15)	Na1—O2—Na1 ⁱⁱⁱ	118.64 (6)
N3—C3—C4	124.00 (16)	O2—Na1—O2 ⁱ	103.44 (4)
C2—C3—C4	131.76 (18)	O2—Na1—O1 ⁱⁱⁱ	83.35 (5)
C3—C4—H4A	109.5	O2 ⁱ —Na1—O1 ⁱⁱⁱ	144.50 (7)
C3—C4—H4B	109.5	O2—Na1—N1 ⁱ	169.72 (6)
H4A—C4—H4B	109.5	O2 ⁱ —Na1—N1 ⁱ	67.86 (5)
C3—C4—H4C	109.5	O1 ⁱⁱⁱ —Na1—N1 ⁱ	100.58 (5)
H4A—C4—H4C	109.5	O2—Na1—N2 ⁱⁱ	104.69 (6)
H4B—C4—H4C	109.5	O2 ⁱ —Na1—N2 ⁱⁱ	123.69 (6)
C6—C5—C10	120.64 (17)	O1 ⁱⁱⁱ —Na1—N2 ⁱⁱ	86.61 (6)
C6—C5—N3	120.01 (16)	N1 ⁱ —Na1—N2 ⁱⁱ	85.11 (5)
C10—C5—N3	119.33 (16)	O2—Na1—C1	20.68 (5)
C5—C6—C7	119.56 (18)	O2 ⁱ —Na1—C1	85.20 (5)
C5—C6—H6	120.2	O1 ⁱⁱⁱ —Na1—C1	103.90 (5)
C7—C6—H6	120.2	N1 ⁱ —Na1—C1	152.77 (5)
C8—C7—C6	119.43 (19)	N2 ⁱⁱ —Na1—C1	107.94 (5)
C8—C7—H7	120.3	O2—Na1—Na1 ⁱ	73.40 (4)
C6—C7—H7	120.3	O2 ⁱ —Na1—Na1 ⁱ	30.25 (3)
C7—C8—C9	121.55 (18)	O1 ⁱⁱⁱ —Na1—Na1 ⁱ	140.89 (6)
C7—C8—Cl1	118.84 (17)	N1 ⁱ —Na1—Na1 ⁱ	98.11 (4)

C9—C8—Cl1	119.61 (17)	N2 ⁱⁱ —Na1—Na1 ⁱ	129.16 (5)
C8—C9—C10	118.82 (19)	C1—Na1—Na1 ⁱ	55.00 (4)
C8—C9—H9	120.6	O2—Na1—Na1 ⁱⁱⁱ	31.12 (4)
C10—C9—H9	120.6	O2 ⁱ —Na1—Na1 ⁱⁱⁱ	132.42 (5)
C5—C10—C9	119.99 (19)	O1 ⁱⁱⁱ —Na1—Na1 ⁱⁱⁱ	53.76 (3)
C5—C10—H10	120.0	N1 ⁱ —Na1—Na1 ⁱⁱⁱ	154.32 (4)
C9—C10—H10	120.0	N2 ⁱⁱ —Na1—Na1 ⁱⁱⁱ	91.80 (4)
N2—N1—C2	109.17 (14)	C1—Na1—Na1 ⁱⁱⁱ	51.60 (4)
N2—N1—Na1 ⁱⁱⁱ	138.87 (11)	Na1 ⁱ —Na1—Na1 ⁱⁱⁱ	103.49 (3)
C2—N1—Na1 ⁱⁱⁱ	109.24 (11)		

Symmetry codes: (i) $-x, y-1/2, -z+3/2$; (ii) $x, -y+3/2, z+1/2$; (iii) $-x, y+1/2, -z+3/2$; (iv) $x, -y+3/2, z-1/2$.

Hydrogen-bond geometry (\AA , °)

$D\text{—H}\cdots A$	$D\text{—H}$	$H\cdots A$	$D\cdots A$	$D\text{—H}\cdots A$
C4—H4A ^v —N2 ^v	0.96	2.52	3.432 (3)	159

Symmetry code: (v) $x, y-1, z$.