

Methyl 2-(4,6-dichloro-1,3,5-triazin-2-yl-amino)acetate

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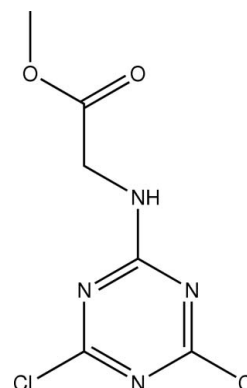
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Key indicators: single-crystal X-ray study; $T = 150$ K; mean $\sigma(\text{C}-\text{C}) = 0.004$ Å; R factor = 0.059; wR factor = 0.165; data-to-parameter ratio = 19.3.

The title compound, $\text{C}_6\text{H}_6\text{Cl}_2\text{N}_4\text{O}_2$, was prepared by the nucleophilic substitution of 2,4,6-trichloro-1,3,5-triazine by glycine methyl ester hydrochloride, and was isolated from the reaction by using flash chromatography. The crystal structure at 150 K reveals the presence two crystallographically independent molecules in the asymmetric unit which differ in the orientation of the pendant methoxycarbonyl group. Each molecular unit is engaged in strong and highly directional $\text{N}-\text{H}\cdots\text{N}$ hydrogen-bonding interactions with a symmetry-related molecule, forming supramolecular dimers which act as the synthons in the crystal packing.

Related literature

For background to nucleophilic reactions based on 1,3,5-triazine derivatives, see: Blotny (2006); Giacomelli *et al.* (2004). For coordination polymers based 1,3,5-triazine derivatives, see: Wang, Xing *et al.* (2007); Wang, Bai, Xing *et al.* (2007); Wang, Bai, Li *et al.* (2007). For general background studies on crystal-engineering approaches from our research group, see: Vilela *et al.* (2009); Shi *et al.* (2008); Paz & Klinowski (2003, 2007); Paz *et al.* (2002, 2005). For a description of the graph-set notation for hydrogen-bonded aggregates, see: Bernstein *et al.* (1995). For a description of the Cambridge Structural Database and the Mercury software package, see: Allen (2002); Macrae *et al.* (2008).



Experimental

Crystal data

$\text{C}_6\text{H}_6\text{Cl}_2\text{N}_4\text{O}_2$
 $M_r = 237.05$
 Triclinic, $P\bar{1}$
 $a = 7.3543$ (4) Å
 $b = 9.7523$ (5) Å
 $c = 13.4133$ (7) Å
 $\alpha = 97.714$ (3)°
 $\beta = 92.714$ (3)°
 $\gamma = 90.225$ (3)°
 $V = 952.19$ (9) Å³
 $Z = 4$
 Mo $K\alpha$ radiation
 $\mu = 0.66$ mm⁻¹
 $T = 150$ K
 $0.18 \times 0.16 \times 0.04$ mm

Data collection

Bruker X8 Kappa CCD APEXII diffractometer
 Absorption correction: multi-scan (SADABS; Sheldrick, 1997)
 $T_{\min} = 0.890$, $T_{\max} = 0.974$
 23605 measured reflections
 5043 independent reflections
 3753 reflections with $I > 2\sigma(I)$
 $R_{\text{int}} = 0.047$

Refinement

$R[F^2 > 2\sigma(F^2)] = 0.059$
 $wR(F^2) = 0.165$
 $S = 1.04$
 5043 reflections
 261 parameters
 2 restraints
 H atoms treated by a mixture of independent and constrained refinement
 $\Delta\rho_{\max} = 1.78$ e Å⁻³
 $\Delta\rho_{\min} = -0.42$ e Å⁻³

Table 1

Hydrogen-bond geometry (Å, °).

$D-\text{H}\cdots A$	$D-\text{H}$	$\text{H}\cdots A$	$D\cdots A$	$D-\text{H}\cdots A$
$\text{N4}-\text{H4}\cdots\text{N2}^i$	0.945 (10)	2.092 (12)	3.028 (3)	171 (3)
$\text{N8}-\text{H8}\cdots\text{N6}^ii$	0.943 (10)	2.083 (11)	3.022 (3)	173 (3)

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

Data collection: APEX2 (Bruker, 2006); cell refinement: SAINT-Plus (Bruker, 2005); data reduction: SAINT-Plus; program(s) used to solve structure: SHELXTL (Sheldrick, 2008); program(s) used to refine structure: SHELXTL; molecular graphics: DIAMOND (Brandenburg, 2009); software used to prepare material for publication: SHELXTL.

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Supplementary data and figures for this paper are available from the IUCr electronic archives (Reference: TK2507).

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supporting information

Acta Cryst. (2009). E65, o1985–o1986 [doi:10.1107/S1600536809028670]

Methyl 2-(4,6-dichloro-1,3,5-triazin-2-ylamino)acetate

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S1. Comment

Worldwide research on 1,3,5-triazine derivatives has increased quite considerably in recent years driven by the versatility of this molecule which allows the nucleophilic substitution of the chloride atoms by various functional groups such as carboxylic acids, amines, amides, chlorides, nitriles, among others (Blotny, 2006; Giacomelli *et al.*, 2004). These reactions allow the engineering of novel derivative compounds which exhibit markedly different properties from their precursors. Hence, the isolated products can be ultimately employed in various areas such as in pharmaceutical sciences, in the textile industry, and in analytical chemistry. Following our interest in crystal engineering (Vilela *et al.*, 2009; Shi *et al.*, 2008; Paz & Klinowski, 2003, 2007; Paz *et al.*, 2002, 2005), we started using 2,4,6-trichloro-1,3,5-triazine as a molecular canvas for the preparation of novel multipodal organic ligands. A search in the literature and in the Cambridge Structural Database (CSD, Version of November 2008 with three updates; Allen, 2002) shows that the group of Bai (Wang, Xing *et al.*, 2007; Wang, Bai, Xing *et al.*, 2007; Wang, Bai, Li *et al.*, 2007) reported the only known examples of transition metal coordination polymers containing *N,N',N''*-1,3,5-triazine-2,4,6-triyltriglycine. We intend to further develop their concept by preparing mono-, di- and tri-substituted derivatives with several amino acid pendant groups. By using glycine methyl ester hydrochloride (Vilela *et al.*, 2009) we isolated the pure title compound (*i.e.*, the monosubstituted derivative, I).

At 150 K compound (I) contains two identical molecular units in the asymmetric unit (Fig. 1). The bond lengths and angles observed for the two molecules are statistically identical. The pendant methoxycarbonyl group exhibits considerable conformational flexibility due to the possibility of rotation around the —CH₂— moiety. Indeed, while the rings and the —NH— moiety of the two crystallographically independent molecular units are almost co-planar, the pendant group is rotated by *ca* 180° (Fig. 2), with this feature arising with the objective to minimize steric repulsion in the crystal structure (see below).

The co-planarity of the —NH— bond with the ring of each molecular unit seems to be promoted by the existence of two strong ($d_{D\cdots A}$ being *ca* 3.02 Å) and highly directional [$\angle(DHA)$ angles above 170° - see Table 1] N—H \cdots N hydrogen bonding interactions that form a $R_2^2(8)$ graph set motif (Bernstein *et al.*, 1995). This arrangement leads to the existence of supramolecular dimers (one for each molecular unit) in the crystal structure, with Fig. 3 depicting one of these. The close packing in the solid-state is based on the spatial interdigitation of the two dimers to effectively occupy the available space, hence the two conformations for the pendant groups which ultimately help promoting a more effective packing (Fig. 4).

S2. Experimental

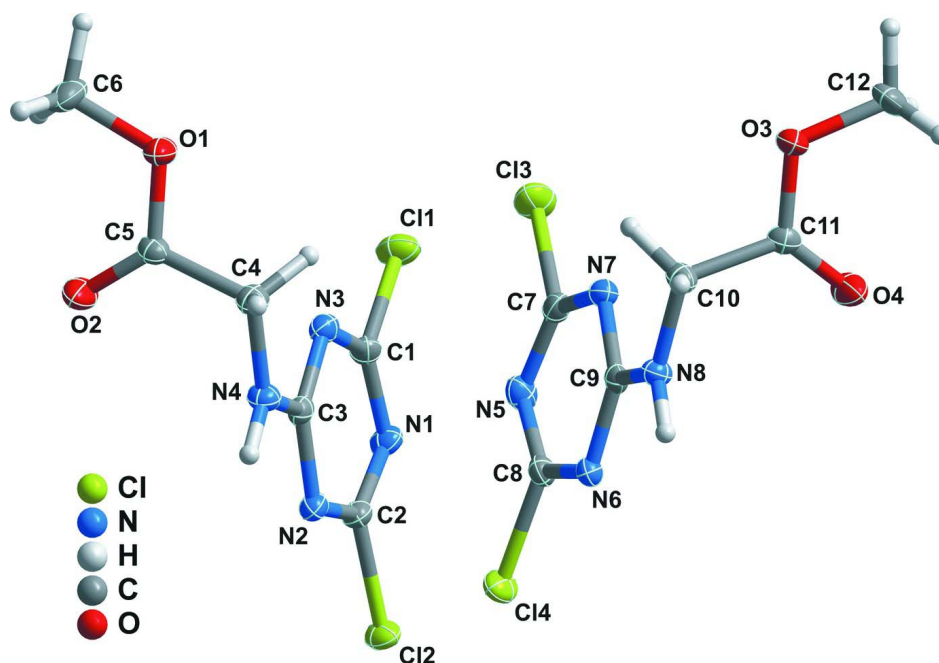
Glycine methyl ester hydrochloride (193 mg, 2.169 mmol; Sigma-Aldrich, 99%) and potassium carbonate (200 mg, 1.447 mmol; Sigma-Aldrich, >99.0%) were added at 273 K to a solution of 2,4,6-trichloro-1,3,5-triazine (100 mg, 0.542 mmol; Sigma-Aldrich, >98,0%) in dried toluene (*ca* 5 ml). The reaction mixture was kept under magnetic stirring and slowly heated to reflux under an anhydrous atmosphere. The reaction was controlled by TLC and stopped after 24 h. The reaction mixture was separated by flash column chromatography using as eluent a gradient of methanol in dichloromethane. The first isolated fraction was identified as (I) (7% yield). Single crystals were isolated from recrystallization of the crude product from a solution in dichloromethane: methanol (*ca* 1: 1). All employed solvents were of analytical grade and purchased from commercial sources.

$^1\text{H NMR}$ (300.13 MHz, CDCl_3) δ : 3.83 (*s*, 3H, OCH_3), 4.27 (*d*, 2H, $J = 2.7$ Hz, CH_2), 6.35 (*br s*, 1H, NH). $^{13}\text{C NMR}$ (75.47 MHz, CDCl_3) δ : 42.8 (CH_2), 52.8 (OCH_3), 165.8 (CNH), 168.9 (CCl), 170.2 (CCl), 171.1 (CO_2Me). **MS (TOF MS ES+)** m/z : 237.0 ($M+H$) $^+$. Selected FT—IR data (ATR, in cm^{-1}): $\nu(\text{N—H}) = 3264m$; $\nu_{\text{asym}}(\text{—CH}_3) = 2961m$; $\nu(\text{C=O}) = 1751vs$; $\nu_{\text{in-plane}}(\text{ring}) = 1549s$ and $1524s$ (doublet); $\delta(\text{—CH}_3) = 1417m$; $\nu(\text{C}_{\text{aromatic}}\text{—N}) = 1322m$; $\nu_{\text{asym}}(\text{C—O—C}) = 1205s$; $\nu_{\text{sym}}(\text{C—O—C}) = 1134s$; $\gamma(\text{ring}) = 841s$.

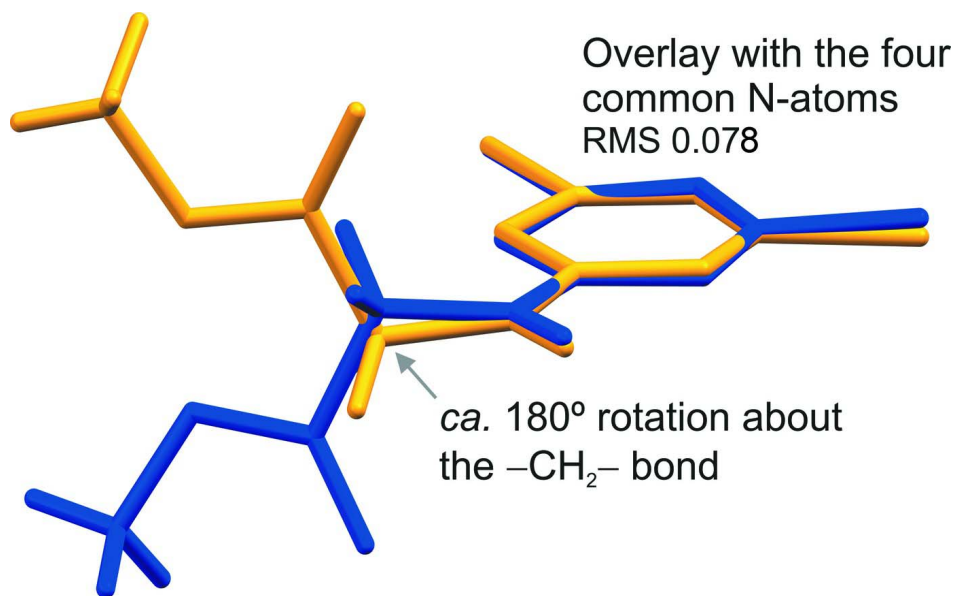
S3. Refinement

Hydrogen atoms bound to carbon were located at their idealized positions and were included in the model in the riding model approximation with $\text{C—H} = 0.99$ Å (for the $\text{—CH}_2\text{—}$ groups) or 0.98 Å (for the —CH_3 moieties). The isotropic thermal displacement parameters for these atoms were fixed at 1.2 (methylene) or 1.5 (methyl) times U_{eq} of the carbon atom to which they are attached. The N—H atoms were located from difference Fourier maps and included in the structure with the N—H distance restrained to $0.95(1)$ Å and with U_{iso} fixed at 1.5 times U_{eq} of the N atom.

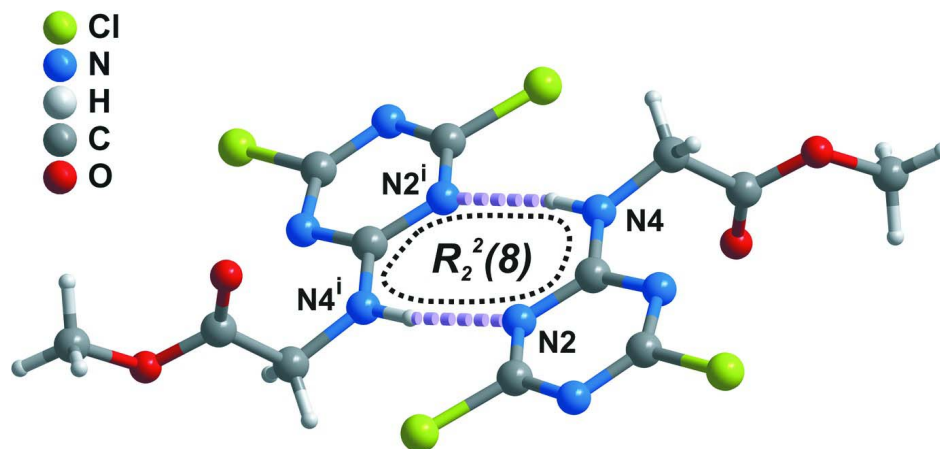
The structure contains a large residual electron density of $1.78 \text{ e}\cdot\text{\AA}^{-3}$ located at 1.36 Å of H4A. Attempts to include this peak as a disordered C atom did not lead to sensible structural refinements.

**Figure 1**

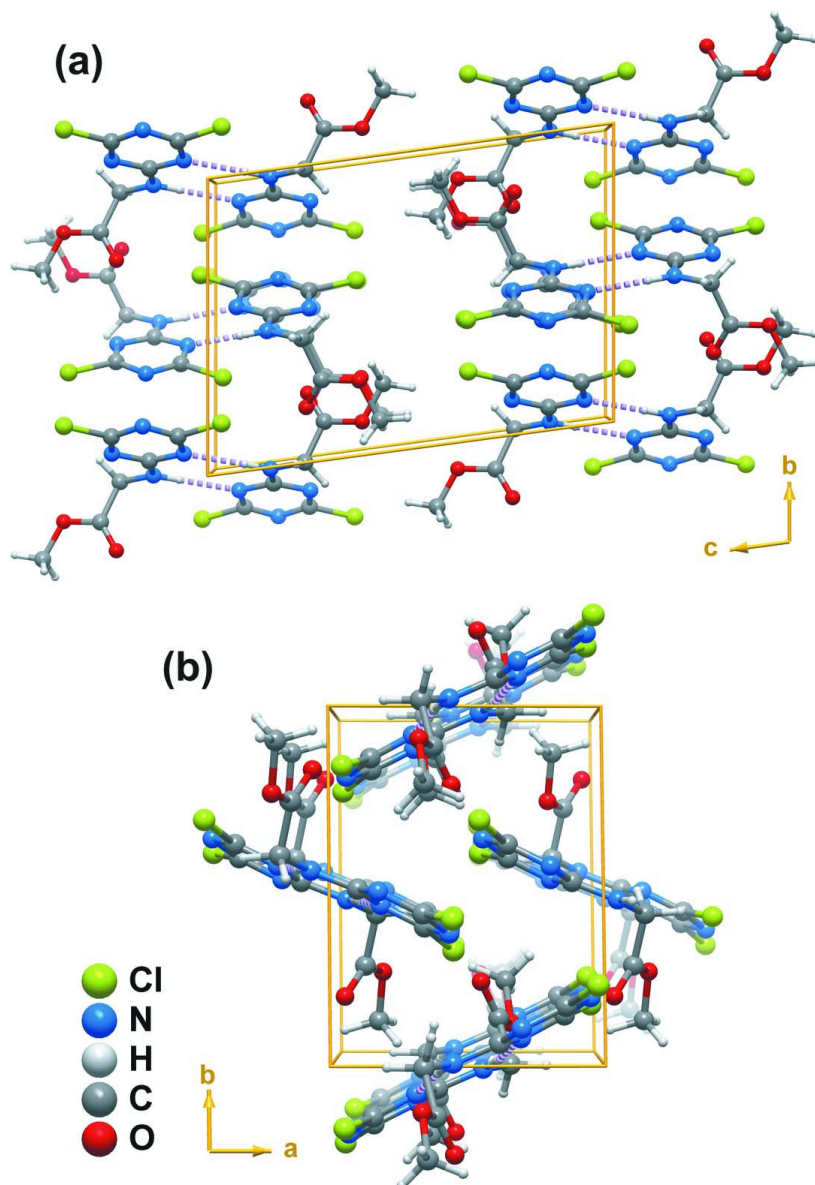
Molecular structures of the two independent molecules in (I). Non-hydrogen atoms are represented as thermal displacement ellipsoids drawn at the 50% probability level and hydrogen atoms as small spheres with arbitrary radii. The atomic labeling is provided for all non-hydrogen atoms.

**Figure 2**

Structure overlay of the two crystallographically independent molecular units comprising the asymmetric unit in (I): while the -NH group remains almost co-planar with the aromatic ring, the two methoxycarbonyl groups are mutually rotated by *ca* 180° around the $\text{-CH}_2\text{-}$ bond.

**Figure 3**

Strong N—H \cdots N hydrogen bonding interactions connecting adjacent molecular units *via* a $R_2^2(8)$ synthon. For details on the hydrogen bonding geometry see Table 1.

**Figure 4**

Crystal packing of (I) viewed in perspective along the (a) [100] and (b) [001] directions of the unit cell. N—H \cdots N hydrogen bonds are represented as violet dashed lines.

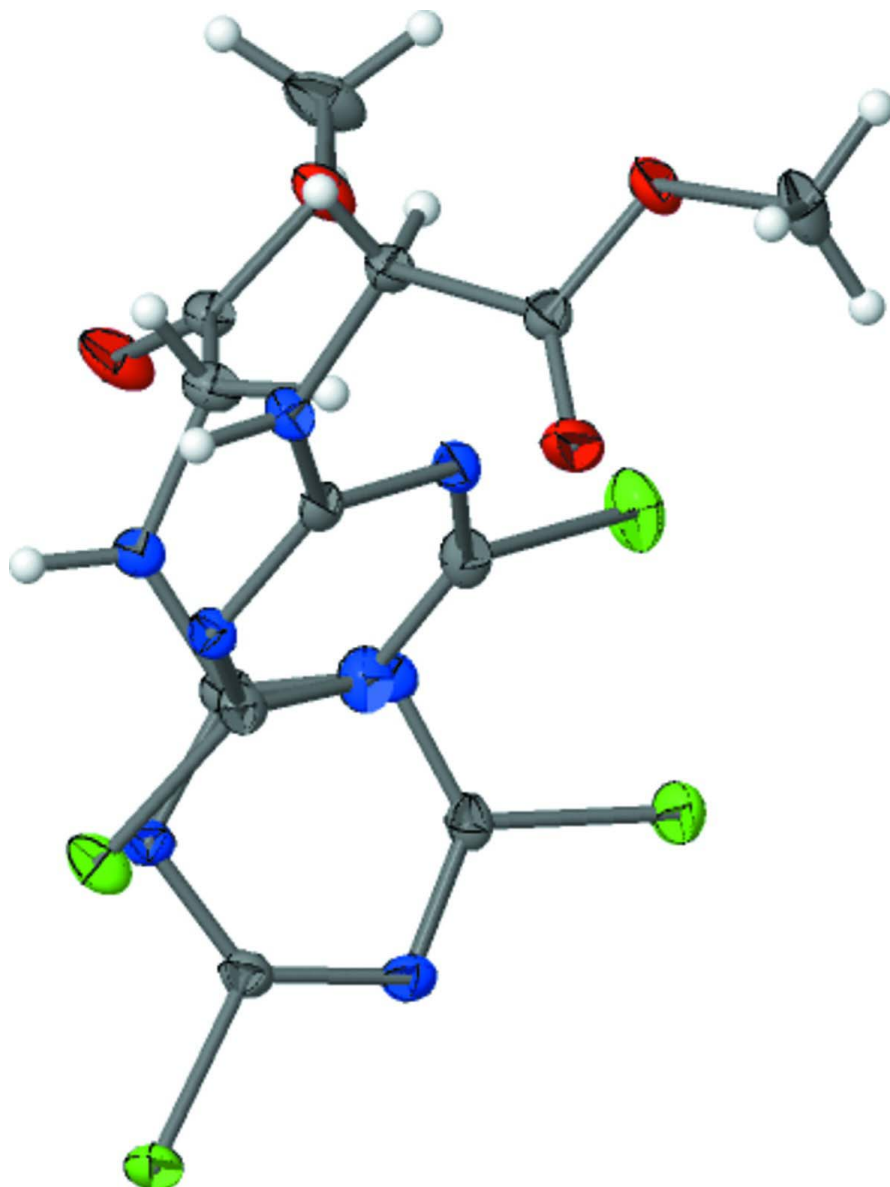
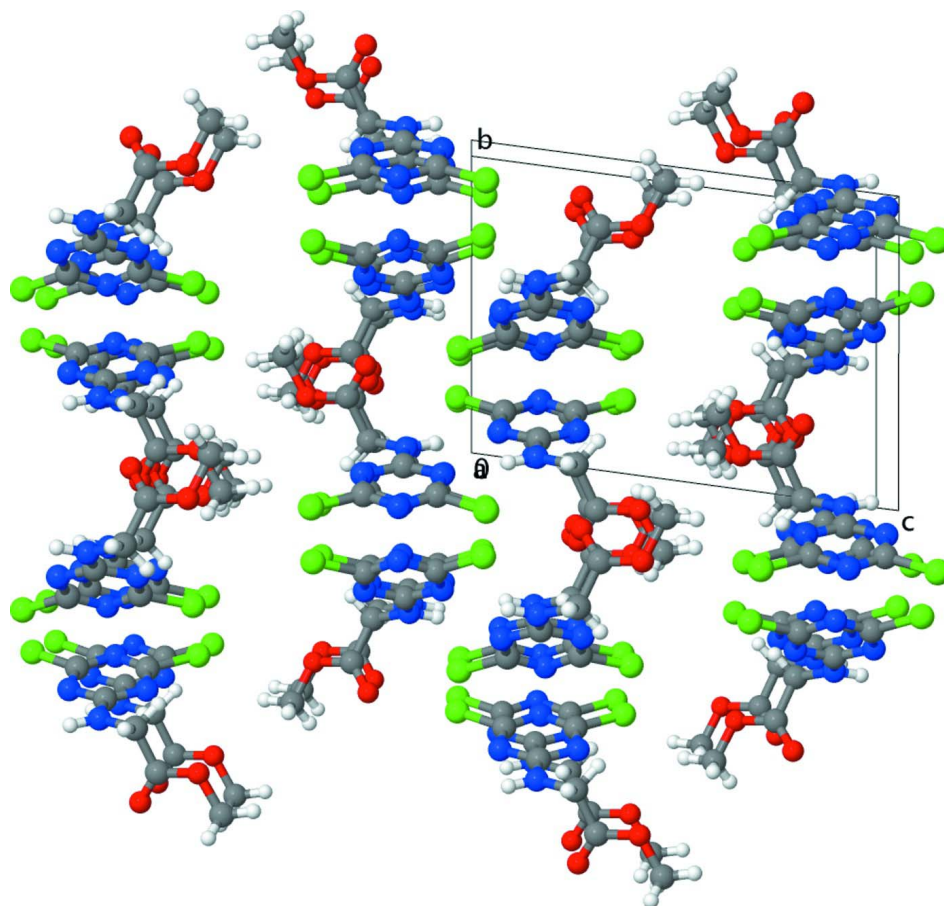


Figure 5

Asymmetric unit of the title compound depicting the two crystallographically independent molecular units. Non-hydrogen atoms are represented as thermal ellipsoids drawn at the 50% probability level.

**Figure 6**

Crystal packing of the title compound viewed in perspective along the [100] direction of the unit cell.

Methyl 2-(4,6-dichloro-1,3,5-triazin-2-ylamino)acetate

Crystal data

$C_6H_6Cl_2N_4O_2$

$M_r = 237.05$

Triclinic, $P\bar{1}$

Hall symbol: -P 1

$a = 7.3543$ (4) Å

$b = 9.7523$ (5) Å

$c = 13.4133$ (7) Å

$\alpha = 97.714$ (3)°

$\beta = 92.714$ (3)°

$\gamma = 90.225$ (3)°

$V = 952.19$ (9) Å³

$Z = 4$

$F(000) = 480$

$D_x = 1.654$ Mg m⁻³

Mo $K\alpha$ radiation, $\lambda = 0.71073$ Å

Cell parameters from 6830 reflections

$\theta = 2.8$ – 28.9 °

$\mu = 0.66$ mm⁻¹

$T = 150$ K

Plate, colourless

$0.18 \times 0.16 \times 0.04$ mm

Data collection

Bruker X8 Kappa CCD APEXII
diffractometer

Radiation source: fine-focus sealed tube

Graphite monochromator

ω and φ scans

Absorption correction: multi-scan
(*SADABS*; Sheldrick, 1997)

$T_{\min} = 0.890$, $T_{\max} = 0.974$

23605 measured reflections

5043 independent reflections

3753 reflections with $I > 2\sigma(I)$

$R_{\text{int}} = 0.047$
 $\theta_{\text{max}} = 29.1^\circ$, $\theta_{\text{min}} = 3.7^\circ$
 $h = -10 \rightarrow 10$

$k = -13 \rightarrow 13$
 $l = -18 \rightarrow 18$

Refinement

Refinement on F^2
 Least-squares matrix: full
 $R[F^2 > 2\sigma(F^2)] = 0.059$
 $wR(F^2) = 0.165$
 $S = 1.04$
 5043 reflections
 261 parameters
 2 restraints
 Primary atom site location: structure-invariant
 direct methods

Secondary atom site location: difference Fourier
 map
 Hydrogen site location: mixed
 H atoms treated by a mixture of independent
 and constrained refinement
 $w = 1/[\sigma^2(F_o^2) + (0.0886P)^2 + 1.283P]$
 where $P = (F_o^2 + 2F_c^2)/3$
 $(\Delta/\sigma)_{\text{max}} = 0.001$
 $\Delta\rho_{\text{max}} = 1.78 \text{ e } \text{\AA}^{-3}$
 $\Delta\rho_{\text{min}} = -0.42 \text{ e } \text{\AA}^{-3}$

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 (\AA^2)

	<i>x</i>	<i>y</i>	<i>z</i>	$U_{\text{iso}}^*/U_{\text{eq}}$
C11	0.03910 (11)	0.76811 (9)	0.64289 (6)	0.02802 (19)
C12	0.05165 (10)	0.83098 (7)	1.03313 (5)	0.02279 (18)
C13	0.55099 (12)	0.59177 (9)	0.63658 (6)	0.0312 (2)
C14	0.54953 (10)	0.68082 (7)	1.02721 (6)	0.02374 (18)
N1	0.0584 (3)	0.8055 (2)	0.83787 (18)	0.0195 (5)
N2	0.3083 (3)	0.9206 (2)	0.93391 (18)	0.0172 (5)
N3	0.3084 (3)	0.8875 (2)	0.75349 (18)	0.0177 (5)
N4	0.5544 (3)	0.9858 (2)	0.85124 (18)	0.0182 (5)
H4	0.610 (5)	1.014 (4)	0.9155 (14)	0.027*
N5	0.5630 (3)	0.6316 (2)	0.83184 (19)	0.0214 (5)
N6	0.8095 (3)	0.5542 (2)	0.93194 (17)	0.0160 (5)
N7	0.8146 (3)	0.5166 (2)	0.75176 (18)	0.0178 (5)
N8	1.0585 (3)	0.4599 (2)	0.85293 (18)	0.0174 (5)
H8	1.109 (5)	0.456 (4)	0.9184 (13)	0.026*
C1	0.1492 (4)	0.8273 (3)	0.7573 (2)	0.0179 (5)
C2	0.1510 (4)	0.8563 (3)	0.9232 (2)	0.0173 (5)
C3	0.3874 (4)	0.9307 (3)	0.8456 (2)	0.0163 (5)
C4	0.6573 (4)	0.9941 (3)	0.7636 (2)	0.0194 (6)
H4A	0.6299	0.9116	0.7136	0.023*
H4B	0.7888	0.9934	0.7830	0.023*
C5	0.6151 (4)	1.1235 (3)	0.7156 (2)	0.0175 (5)
C6	0.6440 (5)	1.2261 (4)	0.5677 (3)	0.0321 (8)

H6A	0.5217	1.2643	0.5759	0.048*
H6B	0.6629	1.1990	0.4959	0.048*
H6C	0.7349	1.2963	0.5957	0.048*
C7	0.6559 (4)	0.5773 (3)	0.7536 (2)	0.0194 (6)
C8	0.6527 (4)	0.6144 (3)	0.9185 (2)	0.0172 (5)
C9	0.8913 (4)	0.5102 (3)	0.8450 (2)	0.0156 (5)
C10	1.1643 (4)	0.4189 (3)	0.7663 (2)	0.0181 (5)
H10A	1.2954	0.4260	0.7871	0.022*
H10B	1.1405	0.4833	0.7162	0.022*
C11	1.1208 (4)	0.2724 (3)	0.7175 (2)	0.0175 (5)
C12	1.1656 (5)	0.1125 (3)	0.5745 (2)	0.0296 (7)
H12A	1.2338	0.0502	0.6137	0.044*
H12B	1.2142	0.1072	0.5073	0.044*
H12C	1.0368	0.0849	0.5682	0.044*
O1	0.6625 (3)	1.1059 (2)	0.62027 (16)	0.0241 (5)
O2	0.5510 (3)	1.2274 (2)	0.75767 (17)	0.0289 (5)
O3	1.1833 (3)	0.2526 (2)	0.62497 (15)	0.0222 (4)
O4	1.0454 (3)	0.1858 (2)	0.75626 (17)	0.0298 (5)

Atomic displacement parameters (Å²)

	U^{11}	U^{22}	U^{33}	U^{12}	U^{13}	U^{23}
C11	0.0294 (4)	0.0362 (4)	0.0181 (4)	-0.0088 (3)	-0.0034 (3)	0.0043 (3)
C12	0.0263 (4)	0.0243 (3)	0.0186 (3)	-0.0070 (3)	0.0063 (3)	0.0042 (3)
C13	0.0374 (5)	0.0337 (4)	0.0217 (4)	0.0070 (3)	-0.0087 (3)	0.0045 (3)
C14	0.0263 (4)	0.0231 (3)	0.0221 (4)	0.0093 (3)	0.0070 (3)	0.0016 (3)
N1	0.0204 (12)	0.0204 (11)	0.0188 (12)	-0.0048 (9)	0.0021 (10)	0.0058 (9)
N2	0.0194 (12)	0.0155 (10)	0.0173 (11)	-0.0006 (9)	0.0032 (9)	0.0041 (9)
N3	0.0194 (12)	0.0180 (11)	0.0165 (11)	0.0009 (9)	0.0011 (9)	0.0045 (9)
N4	0.0176 (12)	0.0208 (11)	0.0163 (11)	-0.0014 (9)	0.0026 (9)	0.0017 (9)
N5	0.0217 (13)	0.0195 (11)	0.0223 (13)	0.0046 (10)	-0.0028 (10)	0.0012 (9)
N6	0.0179 (11)	0.0140 (10)	0.0161 (11)	0.0015 (8)	0.0017 (9)	0.0015 (8)
N7	0.0220 (12)	0.0167 (11)	0.0146 (11)	-0.0015 (9)	0.0013 (9)	0.0017 (8)
N8	0.0190 (12)	0.0186 (11)	0.0144 (11)	0.0020 (9)	0.0017 (9)	0.0010 (9)
C1	0.0212 (14)	0.0182 (12)	0.0143 (13)	0.0001 (10)	-0.0026 (10)	0.0030 (10)
C2	0.0198 (13)	0.0168 (12)	0.0161 (13)	-0.0009 (10)	0.0039 (10)	0.0040 (10)
C3	0.0190 (13)	0.0138 (11)	0.0170 (13)	0.0031 (10)	0.0021 (10)	0.0048 (10)
C4	0.0185 (13)	0.0215 (13)	0.0190 (14)	0.0020 (10)	0.0046 (11)	0.0040 (11)
C5	0.0160 (13)	0.0193 (12)	0.0170 (13)	-0.0012 (10)	0.0035 (10)	0.0007 (10)
C6	0.043 (2)	0.0308 (16)	0.0269 (17)	0.0052 (14)	0.0134 (15)	0.0146 (14)
C7	0.0263 (15)	0.0153 (12)	0.0169 (13)	-0.0008 (11)	-0.0031 (11)	0.0043 (10)
C8	0.0185 (13)	0.0137 (12)	0.0191 (13)	0.0016 (10)	0.0033 (10)	0.0002 (10)
C9	0.0185 (13)	0.0107 (11)	0.0171 (13)	-0.0024 (9)	0.0017 (10)	0.0003 (9)
C10	0.0176 (13)	0.0186 (12)	0.0177 (13)	-0.0016 (10)	0.0043 (10)	-0.0006 (10)
C11	0.0154 (13)	0.0209 (13)	0.0160 (13)	0.0006 (10)	0.0001 (10)	0.0015 (10)
C12	0.0411 (19)	0.0255 (15)	0.0194 (15)	0.0042 (13)	0.0031 (13)	-0.0075 (12)
O1	0.0330 (12)	0.0227 (10)	0.0178 (10)	0.0049 (9)	0.0089 (9)	0.0046 (8)
O2	0.0384 (13)	0.0257 (11)	0.0242 (12)	0.0106 (10)	0.0135 (10)	0.0047 (9)

O3	0.0316 (12)	0.0195 (10)	0.0152 (10)	0.0002 (8)	0.0057 (8)	-0.0006 (8)
O4	0.0389 (13)	0.0254 (11)	0.0252 (12)	-0.0109 (10)	0.0121 (10)	0.0002 (9)

Geometric parameters (Å, °)

C11—C1	1.728 (3)	N8—C10	1.442 (4)
C12—C2	1.723 (3)	N8—H8	0.943 (10)
C13—C7	1.739 (3)	C4—C5	1.519 (4)
C14—C8	1.725 (3)	C4—H4A	0.9900
N1—C1	1.337 (4)	C4—H4B	0.9900
N1—C2	1.337 (4)	C5—O2	1.200 (4)
N2—C2	1.306 (4)	C5—O1	1.331 (3)
N2—C3	1.359 (4)	C6—O1	1.451 (4)
N3—C1	1.314 (4)	C6—H6A	0.9800
N3—C3	1.354 (4)	C6—H6B	0.9800
N4—C3	1.333 (4)	C6—H6C	0.9800
N4—C4	1.439 (4)	C10—C11	1.516 (4)
N4—H4	0.945 (10)	C10—H10A	0.9900
N5—C7	1.330 (4)	C10—H10B	0.9900
N5—C8	1.340 (4)	C11—O4	1.197 (4)
N6—C8	1.312 (4)	C11—O3	1.334 (3)
N6—C9	1.357 (4)	C12—O3	1.444 (4)
N7—C7	1.311 (4)	C12—H12A	0.9800
N7—C9	1.357 (4)	C12—H12B	0.9800
N8—C9	1.330 (4)	C12—H12C	0.9800
C1—N1—C2	111.0 (2)	O1—C6—H6B	109.5
C2—N2—C3	114.0 (2)	H6A—C6—H6B	109.5
C1—N3—C3	113.1 (2)	O1—C6—H6C	109.5
C3—N4—C4	122.6 (2)	H6A—C6—H6C	109.5
C3—N4—H4	119 (2)	H6B—C6—H6C	109.5
C4—N4—H4	119 (2)	N7—C7—N5	129.7 (3)
C7—N5—C8	110.5 (2)	N7—C7—C13	115.6 (2)
C8—N6—C9	113.7 (2)	N5—C7—C13	114.6 (2)
C7—N7—C9	113.1 (2)	N6—C8—N5	128.6 (3)
C9—N8—C10	122.3 (2)	N6—C8—C14	115.4 (2)
C9—N8—H8	117 (2)	N5—C8—C14	116.0 (2)
C10—N8—H8	120 (2)	N8—C9—N6	117.2 (2)
N3—C1—N1	129.1 (3)	N8—C9—N7	118.7 (2)
N3—C1—C11	116.2 (2)	N6—C9—N7	124.1 (3)
N1—C1—C11	114.7 (2)	N8—C10—C11	112.5 (2)
N2—C2—N1	128.4 (3)	N8—C10—H10A	109.1
N2—C2—C12	115.8 (2)	C11—C10—H10A	109.1
N1—C2—C12	115.8 (2)	N8—C10—H10B	109.1
N4—C3—N3	118.5 (3)	C11—C10—H10B	109.1
N4—C3—N2	117.2 (3)	H10A—C10—H10B	107.8
N3—C3—N2	124.3 (3)	O4—C11—O3	124.6 (3)
N4—C4—C5	112.4 (2)	O4—C11—C10	125.6 (3)

N4—C4—H4A	109.1	O3—C11—C10	109.8 (2)
C5—C4—H4A	109.1	O3—C12—H12A	109.5
N4—C4—H4B	109.1	O3—C12—H12B	109.5
C5—C4—H4B	109.1	H12A—C12—H12B	109.5
H4A—C4—H4B	107.9	O3—C12—H12C	109.5
O2—C5—O1	124.8 (3)	H12A—C12—H12C	109.5
O2—C5—C4	125.2 (3)	H12B—C12—H12C	109.5
O1—C5—C4	109.9 (2)	C5—O1—C6	115.8 (2)
O1—C6—H6A	109.5	C11—O3—C12	114.9 (2)
C3—N3—C1—N1	0.7 (4)	C8—N5—C7—N7	-1.6 (4)
C3—N3—C1—Cl1	-179.70 (19)	C8—N5—C7—Cl3	179.3 (2)
C2—N1—C1—N3	0.7 (4)	C9—N6—C8—N5	3.2 (4)
C2—N1—C1—Cl1	-178.9 (2)	C9—N6—C8—Cl4	-176.48 (19)
C3—N2—C2—N1	-2.7 (4)	C7—N5—C8—N6	-0.1 (4)
C3—N2—C2—Cl2	176.07 (19)	C7—N5—C8—Cl4	179.6 (2)
C1—N1—C2—N2	0.5 (4)	C10—N8—C9—N6	-175.6 (2)
C1—N1—C2—Cl2	-178.3 (2)	C10—N8—C9—N7	3.4 (4)
C4—N4—C3—N3	-3.4 (4)	C8—N6—C9—N8	173.8 (2)
C4—N4—C3—N2	175.9 (2)	C8—N6—C9—N7	-5.1 (4)
C1—N3—C3—N4	175.9 (2)	C7—N7—C9—N8	-175.2 (2)
C1—N3—C3—N2	-3.4 (4)	C7—N7—C9—N6	3.7 (4)
C2—N2—C3—N4	-175.1 (2)	C9—N8—C10—C11	-84.5 (3)
C2—N2—C3—N3	4.3 (4)	N8—C10—C11—O4	-18.0 (4)
C3—N4—C4—C5	86.3 (3)	N8—C10—C11—O3	163.6 (2)
N4—C4—C5—O2	21.8 (4)	O2—C5—O1—C6	4.0 (4)
N4—C4—C5—O1	-159.3 (2)	C4—C5—O1—C6	-174.9 (3)
C9—N7—C7—N5	-0.1 (4)	O4—C11—O3—C12	-4.2 (4)
C9—N7—C7—Cl3	178.98 (19)	C10—C11—O3—C12	174.2 (2)

Hydrogen-bond geometry (Å, °)

<i>D</i> —H... <i>A</i>	<i>D</i> —H	H... <i>A</i>	<i>D</i> ... <i>A</i>	<i>D</i> —H... <i>A</i>
N4—H4...N2 ⁱ	0.95 (1)	2.09 (1)	3.028 (3)	171 (3)
N8—H8...N6 ⁱⁱ	0.94 (1)	2.08 (1)	3.022 (3)	173 (3)

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