

Acta Crystallographica Section E **Structure Reports** Online

ISSN 1600-5368

Methyl 2-(tert-butoxycarbonylamino)-1,3-thiazole-5-carboxylate

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Received 5 August 2010; accepted 11 August 2010

Key indicators: single-crystal X-ray study; T = 293 K; mean σ (C–C) = 0.003 Å; R factor = 0.047; wR factor = 0.142; data-to-parameter ratio = 15.2.

The title compound, C₁₀H₁₄N₂O₄S, was synthesized by the reaction of methyl 2-aminothiazole-5-carboxylate and di-tertbutyl carbonate. In this structure, the thiazole ring is planar (mean deviation = 0.0011 Å). Two weak intramolecular C- $H \cdots O$ hydrogen bonds are formed between two of the methyl groups and one carbonyl O atom, resulting in the formation of two twisted six-membered rings. Intermolecular N-H···N hydrogen bonds link the molecules to form centrosymmetric dimeric units, and the hydrogen-bond scheme is completed by intermolecular C-H···O contacts.

Related literature

For information on the use of the title compound in the synthesis of dasatinib [systematic name: N-(2-chloro-6-methylphenyl)-2-({6-[4-(2-hydroxyethyl)piperazin-1-yl]-2-methylpyrimidin-4-ylamino)-5-thiazolecarboxamide], see: Lombardo et al. (2004). For information on the effectiveness of dasatinib in imatinib-resistant Bcr-Abl kinase domain mutants, see: Shah et al. (2004).



2338 independent reflections 1975 reflections with $I > 2\sigma(I)$

3 standard reflections every 200

 $R_{\rm int} = 0.014$

reflections intensity decay: 1%

Experimental

Crystal data

$C_{10}H_{14}N_2O_4S$	$\gamma = 79.08 \ (3)^{\circ}$
$M_r = 258.29$	V = 642.1 (2) Å ³
Triclinic, P1	Z = 2
a = 7.0700 (14) Å	Mo $K\alpha$ radiation
b = 9.2580 (19) Å	$\mu = 0.26 \text{ mm}^{-1}$
c = 10.708 (2) Å	T = 293 K
$\alpha = 70.10 \ (3)^{\circ}$	$0.30 \times 0.10 \times 0.10$ mm
$\beta = 79.67 \ (3)^{\circ}$	

Data collection

Enrat–Nonius CAD-4
diffractometer
Absorption correction: ψ scan
(North et al., 1968)
$T_{\min} = 0.927, T_{\max} = 0.975$
2543 measured reflections
(North <i>et al.</i> , 1968) $T_{\rm min} = 0.927, T_{\rm max} = 0.975$ 2543 measured reflections

Refinement

 $R[F^2 > 2\sigma(F^2)] = 0.047$ 154 parameters $wR(F^2) = 0.142$ H-atom parameters constrained S = 1.01 $\Delta \rho_{\rm max} = 0.22 \text{ e} \text{ Å}^ \Delta \rho_{\rm min} = -0.36 \text{ e } \text{\AA}^{-3}$ 2338 reflections

Table 1

Hydrogen-bond geometry (Å, °).

$D - H \cdots A$	D-H	$H \cdot \cdot \cdot A$	$D \cdots A$	$D - \mathbf{H} \cdot \cdot \cdot A$
C8-H8A···O3	0.96	2.47	3.030 (4)	117
C10-H10A···O3	0.96	2.45	3.010 (4)	117
$N2-H2A\cdots N1^{i}$	0.86	2.02	2.879 (3)	174
C9−H9 <i>C</i> ···O2 ⁱⁱ	0.96	2.60	3.440 (4)	146
$C10-H10C\cdots O2^{ii}$	0.96	2.57	3.436 (4)	150

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

Data collection: CAD-4 EXPRESS (Enraf-Nonius, 1989); cell refinement: CAD-4 EXPRESS: data reduction: XCAD4 (Harms & Wocadlo, 1995); program(s) used to solve structure: SHELXS97 (Sheldrick, 2008); program(s) used to refine structure: SHELXL97 (Sheldrick, 2008); molecular graphics: SHELXTL (Sheldrick, 2008); software used to prepare material for publication: SHELXL97.

The authors thank Professor Hua-qin Wang (Nanjing University) for carrying out the X-ray crystallographic analysis.

Supplementary data and figures for this paper are available from the IUCr electronic archives (Reference: BH2305).

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Acta Cryst. (2010). E66, o2343 [https://doi.org/10.1107/S1600536810032277]

Methyl 2-(tert-butoxycarbonylamino)-1,3-thiazole-5-carboxylate

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S1. Comment

Methyl 2-((*tert*-butoxycarbonyl)amino)thiazole-5-carboxylate is an important intermediate compound in research on synthesizing Dasatinib (Lombardo *et al.*, 2004). Dasatinib is a high affinity dual Src/Abl and c-Kit inhibitor that has been recently approved for all categories of imatinib-refractory chronic myelogenous leukemia (CML) and acute lymphoblastic leukemia (ALL). Dasatinib is also effective in many imatinib resistant Bcr–Abl kinase domain mutants (Shah *et al.*, 2004).

We report here the crystal structure of the title compound, (I). The molecular structure of (I) is shown in Fig. 1. Ring *A* (S/C5/N1/C4/C3) is a planar five-membered ring with a r.m.s. deviation of 0.0011 Å. In this plane, atoms S, C5, N1, C4 and C3 deviate from the mean plane by less than 0.002 Å. The intramolecular C—H…O hydrogen bonds (Table 1) result in the formation of two twisty six-membered rings *B* (O3/C6/O4/C7/C8/H8A) and *C* (O3/C6/O4/C7/C10/H10A). In the crystal structure, intermolecular N—H…N hydrogen bonds (Table 1) link the molecules to form dimeric units (Fig. 2), stabilizing the crystal structure. The hydrogen bonds scheme is completed by intermolecular C—H…O contacts.

S2. Experimental

Methyl 2-aminothiazole-5-carboxylate (10 mmol), di-*tert*-butyl carbonate (12 mmol) and 4-dimethylamino pyridine (0.66 mmol) were added in THF (30 ml), stirred and refluxed under a nitrogen atmosphere for 10 h. The reaction mixture was left to cool to room temperature, precipitated, filtered, and the filter cake was crystallized from ethanol to give pure compound (I). Crystals of (I) suitable for X-ray diffraction were obtained by slow evaporation of an ethanol solution.

S3. Refinement

All H atoms were positioned geometrically, with C—H = 0.96 and 0.93 Å for methyl and aromatic H atoms, respectively, and N—H = 0.86 Å. All H atoms were constrained to ride on their parent atoms, with $U_{iso}(H) = xU_{eq}(\text{carrier atom})$, where x = 1.5 for methyl H atoms and x = 1.2 for all other H atoms.



Figure 1

A view of the molecular structure of (I). Displacement ellipsoids are drawn at the 50% probability level. Dashed lines indicate intramolecular C—H \cdots O hydrogen bonds.



Figure 2

A packing diagram for (I). Dashed lines indicate C—H…N and C—H…O hydrogen bonds.

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Crystal	data
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$C_{10}H_{14}N_2O_4S$	$\alpha = 70.10 \ (3)^{\circ}$
$M_r = 258.29$	$\beta = 79.67 \ (3)^{\circ}$
Triclinic, $P\overline{1}$	$\gamma = 79.08 \ (3)^{\circ}$
Hall symbol: -P 1	$V = 642.1 (2) \text{ Å}^3$
a = 7.0700 (14) Å	Z = 2
b = 9.2580 (19) Å	F(000) = 272
c = 10.708 (2) Å	$D_{\rm x} = 1.336 {\rm ~Mg} {\rm ~m}^{-3}$

Mo *Ka* radiation, $\lambda = 0.71073$ Å Cell parameters from 25 reflections $\theta = 10-13^{\circ}$ $\mu = 0.26 \text{ mm}^{-1}$

Data collection

Enraf–Nonius CAD-4 diffractometer Radiation source: fine-focus sealed tube Graphite monochromator $\omega/2\theta$ scans Absorption correction: ψ scan (North *et al.*, 1968) $T_{\min} = 0.927, T_{\max} = 0.975$ 2543 measured reflections

Refinement

Refinement on F^2 Least-squares matrix: full $R[F^2 > 2\sigma(F^2)] = 0.047$ $wR(F^2) = 0.142$ S = 1.012338 reflections 154 parameters 0 restraints 0 constraints Primary atom site location: structure-invariant direct methods T = 293 K Block, colourless $0.30 \times 0.10 \times 0.10$ mm

2338 independent reflections 1975 reflections with $I > 2\sigma(I)$ $R_{int} = 0.014$ $\theta_{max} = 25.3^{\circ}, \ \theta_{min} = 2.0^{\circ}$ $h = 0 \rightarrow 8$ $k = -10 \rightarrow 11$ $l = -12 \rightarrow 12$ 3 standard reflections every 200 reflections intensity decay: 1%

Secondary atom site location: difference Fourier map Hydrogen site location: inferred from neighbouring sites H-atom parameters constrained $w = 1/[\sigma^2(F_o^2) + (0.1P)^2 + 0.077P]$ where $P = (F_o^2 + 2F_c^2)/3$ $(\Delta/\sigma)_{max} < 0.001$ $\Delta\rho_{max} = 0.22$ e Å⁻³ $\Delta\rho_{min} = -0.36$ e Å⁻³

Fractional atomic coordinates and isotropic or equivalent isotropic displacement parameters $(Å^2)$

	x	У	Ζ	$U_{ m iso}$ */ $U_{ m eq}$	
S	0.31065 (8)	0.77835 (6)	0.78035 (6)	0.0496 (2)	
N1	0.1959 (3)	1.0086 (2)	0.58090 (19)	0.0477 (5)	
01	0.6194 (3)	0.8268 (2)	0.89511 (19)	0.0668 (5)	
C1	0.7656 (4)	0.8233 (4)	0.9752 (3)	0.0845 (10)	
H1B	0.7815	0.7239	1.0430	0.127*	
H1C	0.7258	0.9029	1.0168	0.127*	
H1D	0.8866	0.8406	0.9190	0.127*	
N2	0.0259 (3)	0.7987 (2)	0.63197 (18)	0.0476 (5)	
H2A	-0.0454	0.8511	0.5698	0.057*	
O2	0.6587 (4)	1.0667 (2)	0.7674 (3)	0.1004 (8)	
C2	0.5780 (3)	0.9562 (3)	0.7965 (3)	0.0538 (6)	
O3	0.0857 (3)	0.56934 (19)	0.79484 (18)	0.0637 (5)	
C3	0.4211 (3)	0.9438 (2)	0.7294 (2)	0.0466 (5)	
O4	-0.1471 (2)	0.61245 (16)	0.65983 (15)	0.0486 (4)	
C4	0.3422 (3)	1.0503 (2)	0.6244 (2)	0.0504 (6)	
H4A	0.3845	1.1467	0.5835	0.061*	
C5	0.1664 (3)	0.8678 (2)	0.6549 (2)	0.0413 (5)	
C6	-0.0051 (3)	0.6494 (2)	0.7045 (2)	0.0462 (5)	
C7	-0.2028 (3)	0.4528 (2)	0.7162 (2)	0.0471 (5)	
C8	-0.2827 (4)	0.4215 (3)	0.8624 (2)	0.0627 (7)	

H8A	-0.1806	0.4165	0.9126	0.094*	
H8B	-0.3338	0.3246	0.8954	0.094*	
H8C	-0.3842	0.5035	0.8718	0.094*	
C9	-0.3586 (4)	0.4627 (3)	0.6324 (3)	0.0622 (7)	
H9A	-0.3035	0.4830	0.5404	0.093*	
H9B	-0.4604	0.5451	0.6403	0.093*	
H9C	-0.4107	0.3663	0.6633	0.093*	
C10	-0.0294 (4)	0.3355 (3)	0.6947 (3)	0.0594 (6)	
H10A	0.0673	0.3313	0.7489	0.089*	
H10B	0.0241	0.3657	0.6021	0.089*	
H10C	-0.0699	0.2351	0.7197	0.089*	

Atomic displacement parameters $(Å^2)$

	U^{11}	U^{22}	U^{33}	U^{12}	U^{13}	U^{23}
S	0.0506 (4)	0.0442 (3)	0.0525 (4)	-0.0135 (2)	-0.0243 (3)	-0.0006 (2)
N1	0.0485 (10)	0.0378 (9)	0.0545 (10)	-0.0097 (8)	-0.0187 (8)	-0.0035 (8)
O1	0.0607 (11)	0.0730 (12)	0.0673 (11)	-0.0191 (9)	-0.0341 (9)	-0.0046 (9)
C1	0.0636 (17)	0.120 (3)	0.081 (2)	-0.0089 (17)	-0.0399 (16)	-0.0318 (19)
N2	0.0527 (11)	0.0383 (9)	0.0512 (10)	-0.0130 (8)	-0.0265 (9)	0.0002 (8)
O2	0.0982 (16)	0.0674 (13)	0.144 (2)	-0.0372 (12)	-0.0668 (16)	-0.0038 (13)
C2	0.0439 (12)	0.0543 (14)	0.0684 (15)	-0.0107 (10)	-0.0157 (11)	-0.0196 (12)
O3	0.0749 (11)	0.0483 (9)	0.0655 (10)	-0.0212 (8)	-0.0409 (9)	0.0079 (8)
C3	0.0406 (11)	0.0447 (12)	0.0558 (13)	-0.0098 (9)	-0.0144 (10)	-0.0114 (10)
O4	0.0564 (9)	0.0403 (8)	0.0488 (9)	-0.0159 (7)	-0.0232 (7)	-0.0005 (6)
C4	0.0496 (12)	0.0383 (11)	0.0628 (14)	-0.0130 (9)	-0.0154 (11)	-0.0074 (10)
C5	0.0424 (11)	0.0377 (10)	0.0426 (11)	-0.0066 (8)	-0.0124 (9)	-0.0069 (8)
C6	0.0519 (12)	0.0406 (11)	0.0466 (11)	-0.0122 (9)	-0.0198 (10)	-0.0046 (9)
C7	0.0482 (12)	0.0409 (11)	0.0506 (12)	-0.0162 (9)	-0.0134 (10)	-0.0035 (9)
C8	0.0643 (15)	0.0627 (15)	0.0529 (14)	-0.0171 (12)	-0.0054 (12)	-0.0043 (12)
C9	0.0623 (15)	0.0575 (14)	0.0714 (16)	-0.0204 (12)	-0.0261 (13)	-0.0107 (12)
C10	0.0605 (14)	0.0496 (13)	0.0689 (16)	-0.0116 (11)	-0.0141 (12)	-0.0146 (11)

Geometric parameters (Å, °)

SC5	1.716 (2)	O4—C6	1.329 (3)
S—C3	1.728 (2)	O4—C7	1.494 (2)
N1—C5	1.309 (3)	C4—H4A	0.9300
N1C4	1.372 (3)	С7—С9	1.512 (3)
O1—C2	1.324 (3)	C7—C8	1.513 (3)
01—C1	1.446 (3)	C7—C10	1.517 (3)
C1—H1B	0.9600	C8—H8A	0.9600
C1—H1C	0.9600	C8—H8B	0.9600
C1—H1D	0.9600	C8—H8C	0.9600
N2—C6	1.373 (3)	С9—Н9А	0.9600
N2—C5	1.375 (3)	C9—H9B	0.9600
N2—H2A	0.8600	С9—Н9С	0.9600
O2—C2	1.187 (3)	C10—H10A	0.9600

$C_{2} - C_{3}$	1 466 (3)	C10—H10B	0 9600
03-C6	1.205 (3)	C10—H10C	0.9600
C3—C4	1 345 (3)		
C5—S—C3	88.29 (10)	O3—C6—N2	122.9 (2)
C5—N1—C4	109.33 (19)	O4—C6—N2	109.62 (17)
C2	117.7 (2)	O4—C7—C9	102.10 (17)
O1—C1—H1B	109.5	O4—C7—C8	109.55 (19)
01—C1—H1C	109.5	C9—C7—C8	111.5 (2)
H1B—C1—H1C	109.5	O4—C7—C10	109.70 (18)
O1—C1—H1D	109.5	C9—C7—C10	111.0 (2)
H1B—C1—H1D	109.5	C8—C7—C10	112.4 (2)
H1C—C1—H1D	109.5	С7—С8—Н8А	109.5
C6—N2—C5	122.54 (18)	С7—С8—Н8В	109.5
C6—N2—H2A	118.7	H8A—C8—H8B	109.5
C5—N2—H2A	118.7	С7—С8—Н8С	109.5
O2—C2—O1	123.7 (2)	H8A—C8—H8C	109.5
O2—C2—C3	125.2 (2)	H8B—C8—H8C	109.5
O1—C2—C3	111.0 (2)	С7—С9—Н9А	109.5
C4—C3—C2	128.1 (2)	С7—С9—Н9В	109.5
C4—C3—S	110.13 (16)	H9A—C9—H9B	109.5
C2—C3—S	121.78 (17)	С7—С9—Н9С	109.5
C6—O4—C7	120.74 (16)	H9A—C9—H9C	109.5
C3—C4—N1	116.2 (2)	H9B—C9—H9C	109.5
C3—C4—H4A	121.9	C7—C10—H10A	109.5
N1—C4—H4A	121.9	C7—C10—H10B	109.5
N1—C5—N2	120.82 (19)	H10A—C10—H10B	109.5
N1—C5—S	116.08 (16)	C7-C10-H10C	109.5
N2—C5—S	123.10 (15)	H10A—C10—H10C	109.5
O3—C6—O4	127.5 (2)	H10B-C10-H10C	109.5
C1—O1—C2—O2	2.8 (4)	C4—N1—C5—S	0.3 (3)
C1C3C3	-177.2 (2)	C6—N2—C5—N1	177.8 (2)
O2—C2—C3—C4	1.0 (5)	C6—N2—C5—S	-2.8 (3)
O1—C2—C3—C4	-178.9 (2)	C3—S—C5—N1	-0.13 (18)
O2—C2—C3—S	-179.3 (2)	C3—S—C5—N2	-179.6 (2)
O1—C2—C3—S	0.7 (3)	C7—O4—C6—O3	-4.2 (4)
C5—S—C3—C4	-0.09 (18)	C7—O4—C6—N2	176.44 (18)
C5—S—C3—C2	-179.8 (2)	C5—N2—C6—O3	1.5 (4)
C2—C3—C4—N1	-180.0 (2)	C5—N2—C6—O4	-179.13 (19)
S-C3-C4-N1	0.3 (3)	C6—O4—C7—C9	-177.3 (2)
C5—N1—C4—C3	-0.4 (3)	C6—O4—C7—C8	64.4 (3)
C4—N1—C5—N2	179.8 (2)	C6—O4—C7—C10	-59.4 (3)

Hydrogen-bond geometry (Å, °)

D—H···A	<i>D</i> —Н	H···A	D···A	<i>D</i> —H··· <i>A</i>
С8—Н8А…О3	0.96	2.47	3.030 (4)	117

C10—H10A····O3	0.96	2.45	3.010 (4)	117	
N2—H2A···N1 ⁱ	0.86	2.02	2.879 (3)	174	
С9—Н9С…О2 ^{іі}	0.96	2.60	3.440 (4)	146	
C10—H10 <i>C</i> ···O2 ⁱⁱ	0.96	2.57	3.436 (4)	150	

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