# data reports



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# Crystal structure of 2-*tert*-butyl-1,3-thiazolo[4,5-*b*]pyridine

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Received 2 July 2014; accepted 14 July 2014

Edited by G. Smith, Queensland University of Technology, Australia

The title compound,  $C_{10}H_{12}N_2S$ , does not contain any strong hydrogen-bond donors but two long  $C-H\cdots N$  contacts are observed in the crystal structure, with the most linear interaction linking molecules along [010]. The ellipsoids of the *tert*-butyl group indicate large librational motion.

**Keywords:** crystal structure; C—H···N contacts; 1,3-thiazolo[4,5-*b*]pyri-]pyridine.

#### CCDC reference: 1013859

#### 1. Related literature

For the synthesis of substituted thiazolopyridines, see: Smith *et al.* (1994, 1995); El-Hiti (2003); Johnson *et al.* (2006); Rao *et al.* (2009); Sahasrabudhe *et al.* (2009); Lee *et al.* (2010); Chaban *et al.* (2013). For the crystal structure of a related compound, see: Yu *et al.* (2007).



### 2. Experimental

2.1. Crystal data

 $\begin{array}{l} C_{10}H_{12}N_2S\\ M_r = 192.28\\ Orthorhombic, P2_12_12_1\\ a = 9.4606 \ (3) \ {\rm \AA}\\ b = 9.7999 \ (3) \ {\rm \AA}\\ c = 11.1155 \ (4) \ {\rm \AA} \end{array}$ 

 $V = 1030.55 (6) \text{ Å}^{3}$  Z = 4Cu K\alpha radiation  $\mu = 2.42 \text{ mm}^{-1}$  T = 296 K $0.40 \times 0.29 \times 0.14 \text{ mm}$  2.2. Data collection

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Agilent SuperNova (Dual, Cu at
zero, Atlas) diffractometer
Absorption correction: multi-scan
(CrysAlis PRO; Agilent, 2014)
T_{min} = 0.721, T_{max} = 1.000
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2.3. Refinement

$$\begin{split} R[F^2 > 2\sigma(F^2)] &= 0.033 & \Delta\rho_{\min} = -0.22 \text{ e}^{-3} \\ & wR(F^2) &= 0.090 & \text{Absolute structure: Flack $x$ determined using 791 quotients [(I+)-1996 reflections & (I-)]/[(I+)+(I-)] (Parsons $et$ al., 121 parameters constrained \\ H-atom parameters constrained & Absolute structure parameter: \\ & \Delta\rho_{\max} = 0.16 \text{ e}^{-3} & 0.027 (7) \end{split}$$

Table 1Hydrogen-bond geometry (Å, °).

$D - H \cdots A$	D-H	$H \cdot \cdot \cdot A$	$D \cdots A$	$D - \mathbf{H} \cdot \cdot \cdot A$
$\begin{array}{c} C4-H4\cdots N1^{i}\\ C6-H6\cdots N1^{ii} \end{array}$	0.93 0.93	2.81 2.72	3.564 (3) 3.620 (3)	138 164

Symmetry codes: (i)  $-x + \frac{3}{2}, -y + 1, z + \frac{1}{2}$ ; (ii)  $-x + 1, y + \frac{1}{2}, -z + \frac{1}{2}$ .

Data collection: *CrysAlis PRO* (Agilent, 2014); cell refinement: *CrysAlis PRO*; data reduction: *CrysAlis PRO*; program(s) used to solve structure: *SHELXS2013* (Sheldrick, 2008); program(s) used to refine structure: *SHELXL2013* (Sheldrick, 2008); molecular graphics: *CHEMDRAW ultra* (Cambridge Soft, 2001); software used to prepare material for publication: *SHELXTL* (Sheldrick, 2008).

#### Acknowledgements

The authors extend their appreciation to the Cornea Research Chair, Department of Optometry, College of Applied Medical Sciences, King Saud University, for funding this research.

Supporting information for this paper is available from the IUCr electronic archives (Reference: ZS2307).

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3395 measured reflections

 $R_{\rm int} = 0.016$ 

1996 independent reflections

1951 reflections with  $I > 2\sigma(I)$ 

# supporting information

Acta Cryst. (2014). E70, o932 [doi:10.1107/S160053681401633X]

# Crystal structure of 2-tert-butyl-1,3-thiazolo[4,5-b]pyridine

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# S1. Structural commentary

Various methods have been reported for the synthesis of substituted thiazolopyridines (Smith *et al.*, 1994, 1995; El-Hiti, 2003; Johnson *et al.*, 2006; Rao *et al.*, 2009; Sahasrabudhe *et al.*, 2009; Lee *et al.*, 2010; Chaban *et al.*, 2013). In a continuation of our research focused on new synthetic routes towards substituted heterocycles we have synthesized the title compound 2-*tert*-butylthiazolo[4,5-*b*]pyridine in high yield (Smith *et al.*, 1995; El-Hiti, 2003) and now report its X-ray crystal structure. The X-ray structure for a related compound has been reported previously (Yu *et al.*, 2007). In the title compound (Fig. 1), the ellipsoids of the methyl groups of the *tert*-butyl group are large which is consistent with librational motion of the group. Assumption of a disordered model did not show significant improvement in the refinement. The molecule does not contain strong hydrogen bond donors. In the crystal, two long C—H…N contacts are observed, the most linear of which links the molecules in chains along [010] (Fig. 2).

# S2. Synthesis and crystallization

2-*tert*-Butylthiazolo[4,5-*b*]pyridine was obtained in 97% yield from acid hydrolysis of 3-(diisopropylaminothiocarbonylthio)-2-(pivalamido)pyridine under reflux (Smith *et al.*, 1995). The compound may also be synthesized in 66% yield from reaction of 3-(diisopropylaminothiocarbonylthio)-2-aminopyridine with 2,2-dimethylpropionic acid in the presence of phosphorus oxychloride under reflux (El-Hiti, 2003). Crystallization from a mixture of ethyl acetate and diethyl ether (1:3 by volume) gave the title compound as colourless crystals. The NMR and low and high resolution mass spectra for the title compound were consistent with those previously reported (Smith *et al.*, 1995).

# **S3. Refinement**

Crystal data, data collection and structure refinement details are summarized in the crystal data table. The hydrogen atoms were positioned geometrically and refined using a riding model with  $U_{iso}(H) = 1.2$  times  $U_{eq}$  for the atom to which they are bonded in the case of aromatic rings, NH and CH<sub>2</sub> groups and 1.5 times  $U_{eq}$  for the methyl hydrogens. Crystal data, data collection and structure refinement details are summarized in Table 1. The Flack parameter (Parsons *et al.*, 2013) was 0.027 (7) but is not of any structural relevance with this compound.



## Figure 1

A molecule of the title compound showing atom labels and 50% probability displacement ellipsoids for non-H atoms.



## Figure 2

Crystal structure packing with the long linear C—H···N contacts shown as dashed lines.

# 2-tert-Butyl-1,3-thiazolo[4,5-b]pyridine

Crystal data  $C_{10}H_{12}N_2S$   $M_r = 192.28$ Orthorhombic,  $P2_12_12_1$  a = 9.4606 (3) Å b = 9.7999 (3) Å c = 11.1155 (4) Å V = 1030.55 (6) Å<sup>3</sup> Z = 4F(000) = 408

 $D_{\rm x} = 1.239 \text{ Mg m}^{-3}$ Cu K\alpha radiation,  $\lambda = 1.54184 \text{ Å}$ Cell parameters from 1951 reflections  $\theta = 6.0-73.4^{\circ}$  $\mu = 2.42 \text{ mm}^{-1}$ T = 296 KPlate, colourless  $0.40 \times 0.29 \times 0.14 \text{ mm}$  Data collection

Agilent SuperNova (Dual, Cu at zero, Atlas) diffractometer Radiation source: SuperNova (Cu) X-ray Source $\omega$ scans Absorption correction: multi-scan ( <i>CrysAlis PRO</i> ; Agilent, 2014) $T_{\min} = 0.721, T_{\max} = 1.000$ <i>Refinement</i>	3395 measured reflections 1996 independent reflections 1951 reflections with $I > 2\sigma(I)$ $R_{int} = 0.016$ $\theta_{max} = 73.4^{\circ}, \theta_{min} = 6.0^{\circ}$ $h = -7 \rightarrow 11$ $k = -11 \rightarrow 12$ $l = -13 \rightarrow 10$
Refinement on $F^2$	H-atom parameters constrained
Least-squares matrix: full	$w = 1/[\sigma^2(F_o^2) + (0.0513P)^2 + 0.0815P]$
$R[F^2 > 2\sigma(F^2)] = 0.033$	where $P = (F_o^2 + 2F_c^2)/3$
$wR(F^2) = 0.090$	$(\Delta/\sigma)_{max} < 0.001$
S = 1.12	$\Delta\rho_{max} = 0.16 \text{ e} \text{ Å}^{-3}$
1996 reflections	$\Delta\rho_{min} = -0.22 \text{ e} \text{ Å}^{-3}$
121 parameters	Absolute structure: Flack x determined using
0 restraints	791 quotients [(I+)-(I-)]/[(I+)+(I-)] (Parsons <i>et</i>
Hydrogen site location: inferred from	<i>al.</i> , 2013).
neighbouring sites	Absolute structure parameter: 0.027 (7)

# 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.

	x	у	Ζ	$U_{ m iso}$ */ $U_{ m eq}$	
C1	0.8245 (2)	0.2846 (2)	0.37042 (19)	0.0458 (5)	
C2	0.6967 (2)	0.4685 (2)	0.33748 (19)	0.0435 (5)	
C3	0.7716 (2)	0.5154 (3)	0.4378 (2)	0.0483 (5)	
C4	0.7455 (3)	0.6454 (3)	0.4822 (3)	0.0645 (6)	
H4	0.7924	0.6789	0.5496	0.077*	
C5	0.6470 (3)	0.7218 (3)	0.4218 (3)	0.0677 (7)	
H5	0.6260	0.8097	0.4477	0.081*	
C6	0.5791 (3)	0.6682 (3)	0.3225 (3)	0.0656 (7)	
H6	0.5138	0.7234	0.2833	0.079*	
C7	0.8831 (3)	0.1417 (2)	0.3577 (2)	0.0556 (6)	
C8	0.8599 (5)	0.0641 (4)	0.4747 (3)	0.0938 (11)	
H8A	0.9096	0.1090	0.5387	0.141*	
H8B	0.8948	-0.0274	0.4662	0.141*	
H8C	0.7608	0.0617	0.4929	0.141*	
C9	1.0411 (4)	0.1506 (5)	0.3319 (5)	0.1151 (16)	
H9A	1.0560	0.1983	0.2575	0.173*	
H9B	1.0798	0.0603	0.3260	0.173*	
H9C	1.0871	0.1990	0.3961	0.173*	
C10	0.8067 (6)	0.0657 (4)	0.2577 (4)	0.124 (2)	
H10A	0.7076	0.0614	0.2758	0.186*	

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

# supporting information

H10B	0.8440	-0.0251	0.2513	0.186*
H10C	0.8203	0.1129	0.1829	0.186*
N1	0.72864 (19)	0.3364 (2)	0.30173 (16)	0.0453 (4)
N2	0.6004 (2)	0.5424 (2)	0.2787 (2)	0.0576 (5)
<b>S</b> 1	0.88585 (7)	0.39028 (7)	0.48644 (5)	0.0607 (2)

Atomic displacement parameters  $(\mathring{A}^2)$ 

	$U^{11}$	$U^{22}$	$U^{33}$	$U^{12}$	$U^{13}$	U <sup>23</sup>
C1	0.0431 (10)	0.0571 (12)	0.0371 (10)	0.0007 (10)	-0.0013 (8)	0.0016 (9)
C2	0.0423 (10)	0.0483 (11)	0.0398 (10)	-0.0043 (9)	0.0022 (8)	0.0053 (9)
C3	0.0443 (11)	0.0550 (12)	0.0457 (11)	-0.0048 (9)	0.0022 (9)	-0.0035 (9)
C4	0.0664 (15)	0.0611 (14)	0.0661 (15)	-0.0075 (11)	0.0034 (13)	-0.0160 (13)
C5	0.0707 (17)	0.0500 (13)	0.0823 (19)	-0.0008 (12)	0.0129 (15)	-0.0023 (13)
C6	0.0686 (16)	0.0542 (13)	0.0740 (17)	0.0102 (12)	0.0041 (14)	0.0157 (13)
C7	0.0564 (12)	0.0560 (13)	0.0544 (12)	0.0118 (11)	-0.0010 (11)	0.0012 (10)
C8	0.122 (3)	0.078 (2)	0.081 (2)	0.0248 (19)	0.009 (2)	0.0219 (17)
C9	0.074 (2)	0.097 (3)	0.174 (5)	0.0241 (19)	0.044 (3)	-0.007 (3)
C10	0.187 (5)	0.073 (2)	0.112 (3)	0.052 (3)	-0.072 (3)	-0.034 (2)
N1	0.0468 (9)	0.0492 (9)	0.0401 (9)	0.0001 (8)	-0.0045 (8)	0.0005 (8)
N2	0.0588 (11)	0.0570 (11)	0.0570 (12)	0.0070 (10)	-0.0078 (10)	0.0101 (9)
<b>S</b> 1	0.0568 (3)	0.0761 (4)	0.0493 (3)	0.0096 (3)	-0.0161 (3)	-0.0112 (3)

Geometric parameters (Å, °)

C6—H6 C7—C10 C7—C8 C7—C9 C8—H8A C8—H8B	0.9300 1.521 (4) 1.522 (4) 1.524 (4) 0.9600
C7—C10 C7—C8 C7—C9 C8—H8A C8—H8B	1.521 (4) 1.522 (4) 1.524 (4) 0.9600
C7—C8 C7—C9 C8—H8A C8—H8B	1.522 (4) 1.524 (4) 0.9600
C7—C9 C8—H8A C8—H8B	1.524 (4) 0.9600
C8—H8A C8—H8B	0.9600
C8—H8B	0.0600
	0.9000
C8—H8C	0.9600
С9—Н9А	0.9600
C9—H9B	0.9600
С9—Н9С	0.9600
C10—H10A	0.9600
C10—H10B	0.9600
C10—H10C	0.9600
C8C7C9	109.3 (3)
C7 - C8 - H8A	109.5
C7 - C8 - H8B	109.5
H8A - C8 - H8B	109.5
C7—C8—H8C	109.5
H8A—C8—H8C	109.5
H8B-C8-H8C	109.5
С7—С9—Н9А	109.5
С7—С9—Н9В	109.5
	C8—H8B C8—H8C C9—H9A C9—H9B C9—H9C C10—H10A C10—H10B C10—H10C C8—C7—C9 C7—C8—H8A C7—C8—H8B H8A—C8—H8B H8A—C8—H8C H8B—C8—H8C H8B—C8—H8C C7—C9—H9A C7—C9—H9B

116.6 (3)	H9A—C9—H9B	109.5
121.7	С7—С9—Н9С	109.5
121.7	Н9А—С9—Н9С	109.5
120.0 (3)	Н9В—С9—Н9С	109.5
120.0	C7-C10-H10A	109.5
120.0	C7—C10—H10B	109.5
124.8 (3)	H10A—C10—H10B	109.5
117.6	C7—C10—H10C	109.5
117.6	H10A—C10—H10C	109.5
110.3 (2)	H10B—C10—H10C	109.5
109.3 (2)	C1—N1—C2	110.6 (2)
108.1 (3)	C2—N2—C6	115.0 (2)
108.9 (3)	C3—S1—C1	88.96 (11)
110.9 (3)		
	116.6 (3) 121.7 121.7 120.0 (3) 120.0 120.0 124.8 (3) 117.6 117.6 110.3 (2) 109.3 (2) 108.1 (3) 108.9 (3) 110.9 (3)	116.6(3) $H9A-C9-H9B$ $121.7$ $C7-C9-H9C$ $121.7$ $H9A-C9-H9C$ $120.0(3)$ $H9B-C9-H9C$ $120.0$ $C7-C10-H10A$ $120.0$ $C7-C10-H10B$ $124.8(3)$ $H10A-C10-H10B$ $117.6$ $C7-C10-H10C$ $117.6$ $H10A-C10-H10C$ $110.3(2)$ $H10B-C10-H10C$ $109.3(2)$ $C1-N1-C2$ $108.1(3)$ $C2-N2-C6$ $108.9(3)$ $C3-S1-C1$

# Hydrogen-bond geometry (Å, °)

D—H···A	D—H	H···A	D···A	D—H…A
C4— $H4$ ···N1 <sup>i</sup>	0.93	2.81	3.564 (3)	138
C6—H6…N1 <sup>ii</sup>	0.93	2.72	3.620 (3)	164

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