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## Key indicators

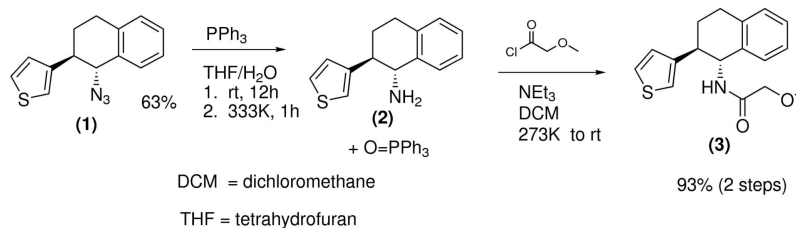
Single-crystal X-ray study  
 $T = 150\text{ K}$   
Mean  $\sigma(\text{C}-\text{C}) = 0.006\text{ \AA}$   
 $R$  factor = 0.059  
 $wR$  factor = 0.158  
Data-to-parameter ratio = 16.8For details of how these key indicators were  
automatically derived from the article, see  
<http://journals.iucr.org/e>.2-Methoxy-*N*-[2-(3-thienyl)-1,2,3,4-tetra-  
hydro-1-naphthyl]acetamide

The asymmetric unit of the title compound,  $\text{C}_{17}\text{H}_{19}\text{NO}_2\text{S}$ , contains two independent molecules with different conformations. Both molecules are linked by intermolecular  $\text{N}-\text{H}\cdots\text{O}$  hydrogen bonds [ $\text{N}\cdots\text{O} = 3.033(4)$  and  $2.969(4)\text{ \AA}$ ] to form two types of one-dimensional chains running in the  $b$  direction.

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## Comment

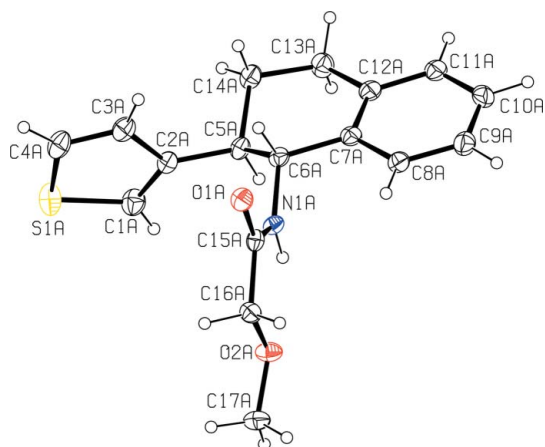
Tetralins (tetrahydronaphthalenes) can be considered as privileged structures, as they are present in a variety of important bioactive compounds, such as the antidepressant sertraline and the anticancer agent podophyllotoxin. One useful method for the synthesis of these scaffolds involves the metal-catalyzed ring-opening reaction of heterobicyclic alkenes (for a review, see Lautens *et al.*, 2003). Earlier, we reported that  $\text{Rh}^{\text{I}}$  catalysts can promote the highly enantioselective ring-opening of oxabicyclic alkenes with boronic acids (Lautens *et al.*, 2002). Recently, we have utilized this method to synthesize several tetralin scaffolds for drug discovery efforts with our industrial collaborators (Dockendorff, 2006). During the course of this project, the title amide, (3), was synthesized from azide (1), and we report its X-ray crystal structure here.



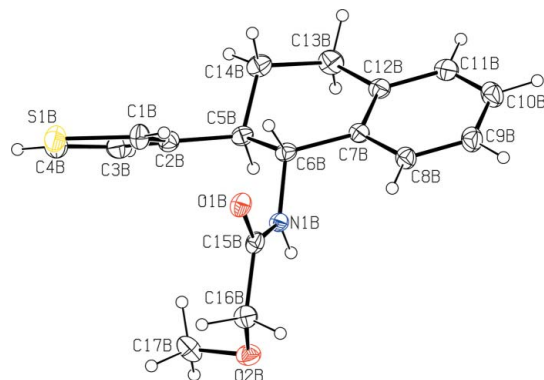
The two independent molecules, *A* and *B*, in the asymmetric unit of (3) are shown in Figs. 1 and 2, respectively. Table 1 lists the pertinent torsion angles, which describe the differences in the conformations between the two molecules. The dihedral angles between the mean planes of the thiophene and benzene rings are  $81.4(2)$  and  $44.9(2)^\circ$  for molecules *A* and *B*, respectively.

The conformational analysis of the cyclohexene ring ( $\text{C}5/\text{C}6/\text{C}7/\text{C}12-\text{C}14$ ) in each molecule (Duax *et al.*, 1976) shows that the conformation is a half-chair, with a local pseudotwofold axis running through the midpoints of the  $\text{C}5\text{A}-\text{C}14\text{A}$  ( $\text{C}5\text{B}-\text{C}14\text{B}$ ) and  $\text{C}7\text{A}-\text{C}12\text{A}$  ( $\text{C}7\text{B}-\text{C}12\text{B}$ ) bonds.

In the crystal structure of (3), two independent one-dimensional  $C_4$  chains (Bernstein *et al.*, 1995) running along the  $b$  direction are formed *via* intermolecular  $\text{N}-\text{H}\cdots\text{O}$  hydrogen bonds (Table 2 and Fig. 2). Chains of both types of



**Figure 1**  
A view of molecule *A* of (3), showing 30% probability displacement ellipsoids (arbitrary spheres for the H atoms).

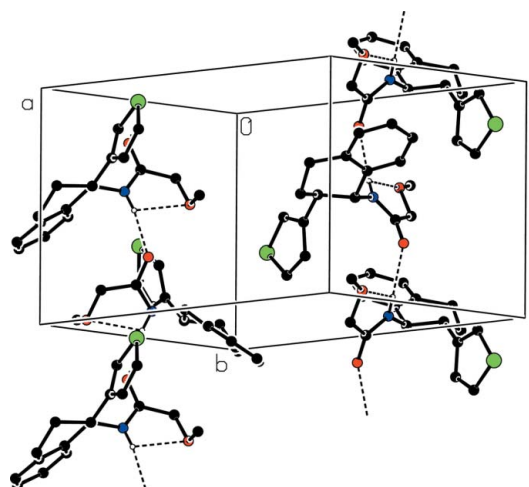


**Figure 2**  
A view of molecule *B* of (3), showing 30% probability displacement ellipsoids (arbitrary spheres for the H atoms).

molecule are, in turn, linked together into a two-dimensional framework *via* weak intermolecular C—H...O hydrogen bonds (Table 2).

## Experimental

Azide (1) (253 mg, 0.991 mmol) and PPh<sub>3</sub> (520 mg, 1.98 mmol) were added to a 50 ml round-bottomed flask with a stirrer bar. The flask was sealed and flushed with nitrogen before tetrahydrofuran (THF; 9 ml) and water (deionized, 0.9 ml) were added. The reaction mixture was stirred for 12 h, after which time analysis by thin-layer chromatography (TLC) [50% dichloromethane (DCM)/hexanes] showed no azide remaining. The reaction mixture was then heated in an oil bath at 333 K for 2 h to hydrolyze all the phosphoimine intermediate. The reaction mixture was then evaporated and dried under high vacuum to remove most of the water. The resulting crude amine, (2), was dissolved in DCM, dried over MgSO<sub>4</sub>, filtered and transferred to a flame-dried 25 ml round-bottomed flask with a stirrer bar and further DCM (5 ml). The flask was sealed and flushed with N<sub>2</sub> and then cooled in an ice bath, before NEt<sub>3</sub> (275 μl, 1.98 mmol) and methoxyacetyl chloride (136 μl, 1.49 mmol) were added dropwise *via* a syringe. The reaction mixture was removed from the ice bath and stirred for 4 h, after which time TLC analysis (5% MeOH/DCM) showed no amine remaining. The reaction mixture was diluted with saturated aqueous NH<sub>4</sub>Cl and DCM in a 60 ml separating funnel, separated and the aqueous layer re-extracted twice with DCM. The



**Figure 3**  
A partial packing plot of (3), showing hydrogen bonds as dashed lines. H atoms not involved in hydrogen bonding have been omitted.

combined organics were dried over MgSO<sub>4</sub>, filtered and evaporated down to a pale-yellow oil. The crude product was purified by flash chromatography (2.5 × 15.2 cm SiO<sub>2</sub>, crude product loaded as solution in DCM, eluted with 0–70% EtOAc/hexanes) (yield 93%). X-ray quality crystals of (3) were obtained by dissolving the compound in a minimum volume of boiling EtOH and then cooling to room temperature in a sealed 1-dram vial, before storing in a freezer at 258 K for 20 h. Spectroscopic analysis: <sup>1</sup>H NMR (300 MHz, CDCl<sub>3</sub>, δ, p.p.m.): 7.31–7.23 (2H, *m*), 7.22–7.16 (2H, *m*), 7.12 (1H, *m*), 7.02 (1H, *d*, *J* = 4.1 Hz), 6.67 (1H, *br d*, *J* = 9.4 Hz), 5.46 (1H, *dd*, *J* = 9.1 Hz), 3.92 (1H, *d*, *J* = 15.2 Hz), 3.78 (1H, *d*, *J* = 15.2 Hz), 3.29 (3H, *s*), 3.13 (1H, *ddd*, *J* = 9.7, 9.7 and 3.2 Hz), 3.03–2.77 (2H, *m*), 2.25–1.99 (2H, *m*).

### Crystal data

C<sub>17</sub>H<sub>19</sub>NO<sub>2</sub>S  
*M<sub>r</sub>* = 301.39  
 Monoclinic, *P*2<sub>1</sub>  
*a* = 10.5344 (3) Å  
*b* = 9.7241 (4) Å  
*c* = 15.1999 (6) Å  
 β = 99.594 (2)°  
*V* = 1535.2 (1) Å<sup>3</sup>  
*Z* = 4

*D<sub>x</sub>* = 1.304 Mg m<sup>-3</sup>  
 Mo Kα radiation  
 Cell parameters from 12544 reflections  
 θ = 2.6–27.5°  
 μ = 0.22 mm<sup>-1</sup>  
*T* = 150 (1) K  
 Needle, colourless  
 0.20 × 0.08 × 0.05 mm

### Data collection

Bruker–Nonius KappaCCD area-detector diffractometer  
 φ scans, and ω scans with κ offsets  
 Absorption correction: multi-scan (SORTAV; Blessing, 1995)  
*T<sub>min</sub>* = 0.619, *T<sub>max</sub>* = 0.993  
 12544 measured reflections

6434 independent reflections  
 4012 reflections with *I* > 2σ(*I*)  
*R<sub>int</sub>* = 0.076  
 θ<sub>max</sub> = 27.5°  
*h* = −13 → 13  
*k* = −12 → 11  
*l* = −18 → 19

### Refinement

Refinement on *F*<sup>2</sup>  
*R* [*F*<sup>2</sup> > 2σ(*F*<sup>2</sup>)] = 0.059  
*wR* (*F*<sup>2</sup>) = 0.158  
*S* = 1.01  
 6434 reflections  
 382 parameters  
 H-atom parameters constrained  
*w* = 1/[σ<sup>2</sup>(*F<sub>o</sub>*<sup>2</sup>) + (0.0723*P*)<sup>2</sup>]  
 where *P* = (*F<sub>o</sub>*<sup>2</sup> + 2*F<sub>c</sub>*<sup>2</sup>)/3

(Δ/σ)<sub>max</sub> = 0.001  
 Δρ<sub>max</sub> = 0.25 e Å<sup>-3</sup>  
 Δρ<sub>min</sub> = −0.32 e Å<sup>-3</sup>  
 Extinction correction: SHELXTL/PC (Sheldrick, 2001)  
 Extinction coefficient: 0.012 (2)  
 Absolute structure: Flack (1983), with 2710 Friedel pairs  
 Flack parameter: −0.07 (9)

**Table 1**Selected torsion angles ( $^{\circ}$ ).

C6A—C5A—C2A—C1A	-130.6 (4)	C15A—C16A—O2A—C17A	-172.1 (3)
C6B—C5B—C2B—C1B	23.4 (5)	C15B—C16B—O2B—C17B	-75.7 (4)

**Table 2**Hydrogen-bond geometry ( $\text{\AA}$ ,  $^{\circ}$ ).

$D-H\cdots A$	$D-H$	$H\cdots A$	$D\cdots A$	$D-H\cdots A$
N1A—H1NA $\cdots$ O2A	0.88	2.19	2.613 (4)	109
N1B—H1NB $\cdots$ O2B	0.88	2.35	2.741 (4)	107
N1A—H1NA $\cdots$ O1A <sup>i</sup>	0.88	2.28	3.033 (4)	144
N1B—H1NB $\cdots$ O1B <sup>ii</sup>	0.88	2.10	2.969 (4)	168
C1B—H1B $\cdots$ O2B <sup>iii</sup>	0.95	2.54	3.386 (5)	148
C3A—H3AA $\cdots$ O2A <sup>iv</sup>	0.95	2.52	3.441 (5)	164
C4B—H4B $\cdots$ O1A <sup>v</sup>	0.95	2.49	3.419 (5)	165
C10A—H10A $\cdots$ O1B <sup>vi</sup>	0.95	2.55	3.276 (5)	134

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

H atoms were placed in calculated positions, with C—H distances in the range 0.95–1.00  $\text{\AA}$  and an N—H distance of 0.88  $\text{\AA}$ . They were included in the refinement in the riding-model approximation, with  $U_{\text{iso}}(\text{H}) = 1.2U_{\text{eq}}(\text{C})$ , or  $1.5U_{\text{eq}}(\text{C})$  for methyl H atoms.

Data collection: *COLLECT* (Nonius, 2002); cell refinement: *DENZO-SMN* (Otwinowski & Minor, 1997); data reduction:

*DENZO-SMN*; program(s) used to solve structure: *SIR92* (Altomare *et al.*, 1994); program(s) used to refine structure: *SHELXTL/PC* (Sheldrick, 2001); molecular graphics: *PLATON* (Spek, 2003); software used to prepare material for publication: *SHELXTL/PC*.

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