04—S1—O2—C1	5.6 (3)	C6C5C10C9	-20.4 (5
C10C5C6C7	49.4 (5)	C3-C2-C11-O1	36.5 (5
C5—C6—C7—C8	-55.1 (5)	C1-C2-C11-C12	38.2 (5
C6C7C8C9	29.4 (5)	O3-S1-C19-C20	23.4 (3
C7—C8—C9—C10	1.3 (5)	O4—S1—C19—C24	-22.9 (3
C8C9C10C5	-4.3 (5)		

Table 2. Hydrogen-bonding geometry (Å, °)

D— $H$ ··· $A$	<i>D</i> H	$\mathbf{H} \cdot \cdot \cdot \mathbf{A}$	$D \cdot \cdot \cdot A$	$D = H \cdot \cdot \cdot A$
C12-H12C···O2	0.96	2.28	2.987 (5)	130
C7—H7A···O3′	0.97	2.47	3.238 (6)	136
Summatry and a: (i) 2	v 1	-		

Symmetry code: (i) 2 - x, 1 - y, -z.

Due to the low r/p ratio of 9.7 for observed reflections, all the H atoms were geometrically fixed and allowed to ride on the atoms to which they were attached. Rotating-group refinement was used for the methyl groups. The reflections  $\bar{4}13$ , 400, 523 and 107 were suppressed during refinement as they showed very poor agreement between  $F_o^2$  and  $F_c^2$ . At the end of the refinement, a distinct peak of 0.74 e Å<sup>-3</sup> at 1.02 Å from C6 suggested the possibility of a conformational disorder for the cyclohexene ring. But the refinement based on a disorder model resulted in extreme displacement parameters for the minor component (0.2 occupancy) of C6 and unrealistic bond lengths and angles involving the same. Hence, the disorder model was abandoned.

Data collection: *SMART* (Siemens, 1996). Cell refinement: *SAINT* (Siemens, 1996). Data reduction: *SAINT*. Program(s) used to solve structure: *SHELXTL* (Sheldrick, 1997). Program(s) used to refine structure: *SHELXTL*. Molecular graphics: *SHELXTL*. Software used to prepare material for publication: *SHELXTL*, *PARST* (Nardelli, 1995) and *PLATON* (Spek, 1990).

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# 9-Hydroxy-8-iodo-4-methyl-7,8,9,10-tetrahydrobenzo[*h*]coumarin

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#### Abstract

In the title molecule,  $C_{14}H_{13}IO_3$ , the coumarin ring system is planar and the fused cyclohexane ring adopts a half-chair conformation. The planes of the coumarin and cyclohexane rings form a dihedral angle of 8.5 (1)°. In the crystal, the hydroxyl group and the carbonyl O atom are involved in an O—H···O intermolecular hydrogen bond. The molecules are stacked at a perpendicular distance of 3.588 (8) Å along the *a* axis.

## Comment

Coumarin derivatives are found in many natural products displaying diverse biological properties. They are found to exhibit antifungal and anticoagulant properties (Parrish *et al.*, 1974). The amino and hydroxy coumarin derivatives are widely used in dye lasers (Maeda, 1984). The structural study of the title compound, (I), one of the coumarin derivatives, was performed as part of our study on coumarin derivatives.



As observed in other coumarin derivatives, the C3— C4 and C5—C6 bonds in the coumarin moiety show double-bond character, and steric interactions cause the widening of angles C3—C2—O11 and C4—C10— C5, and the narrowing of angles O1—C2—O11 and O1—C9—C8 from 120° (Chinnakali *et al.*, 1997*a*,*b*; Kumar *et al.*, 1997). The coumarin ring system is planar, with the O11 atom deviating by a maximum

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of -0.056(5) Å. The fused cyclohexane ring adopts a half-chair conformation, with C13 and C14 deviating from the mean plane by -0.372(6) and 0.306(6) Å, respectively; the asymmetry parameter  $\Delta C_2$ (C14—C13) is 0.045(2) (Nardelli, 1983). The mean plane through the fused cyclohexane ring forms a dihedral angle of 8.5(1)° with the coumarin ring plane. In the solid state, the glide-related molecules are linked by an O—H···O hydrogen bond involving the hydroxyl group and the carbonyl O atom (Table 2) to form an infinite one-dimensional chain. The molecules translated a unit along the *a* axis are stacked stepwise at a perpendicular distance of 3.588(8) Å, with no  $\pi$ - $\pi$  interactions.



Fig. 1. The structure of (1), showing 50% probability displacement ellipsoids and the atom-numbering scheme.

#### Experimental

Our ongoing synthetic project required the synthesis of 9-hydroxy-4,8-dimethyl-7,8,9,10-tetrahydro-7,8-benzocoumarin. To achieve our target compound, 4-methyl-7,10-dihydro-7,8-benzocoumarin 8,9-oxide (0.5 g, 2.19 mmol) in tetrahydro-furan (THF) was added to a solution of methyl magnesium iodide (1.2 equivalents, 0.437 g, 2.63 mmol) in THF under a nitrogen atmosphere. After completion of the reaction, the reaction mixture was quenched with saturated ammonium chloride. The resulting organic layer was separated and dried. The residue after chromatographic purification furnished the title compound, (I), instead of our target compound. Single crystals of (I) were grown by slow evaporation of a chloroform/methanol (1:1) solution of the compound.

Crystal	data
~	

$C_{14}H_{13}IO_3$	Mo $K\alpha$ radiation
$M_r = 356.14$	$\lambda = 0.71073 \text{ Å}$

Monoclinic  

$$P2_1/c$$
  
 $a = 5.4931 (1) Å$   
 $b = 22.2521 (3) Å$   
 $c = 11.0350 (2) Å$   
 $\beta = 104.230 (1)^{\circ}$   
 $V = 1307.45 (4) Å^{3}$   
 $Z = 4$   
 $D_x = 1.809 \text{ Mg m}^{-3}$ 

 $D_m$  not measured

· ·

## Data collection

Siemens SMART CCD area-
detector diffractometer
$\omega$ scans
Absorption correction:
multi-scan empirical
(SADABS; Sheldrick,
1997)
$T_{\rm min} = 0.361, T_{\rm max} = 0.415$

## Refinement

Refinement on  $F^2$  (A  $R[F^2 > 2\sigma(F^2)] = 0.053$  (A  $wR(F^2) = 0.151$  (A) S = 1.208 (B) 2824 reflections (S) 165 parameters H atoms: see below  $w = 1/[\sigma^2(F_o^2) + (0.0530P)^2 + 4.5488P]$ where  $P = (F_o^2 + 2F_o^2)/3$ 

Cell parameters from 2021 reflections  $\theta = 2.64-26.99^{\circ}$  $\mu = 2.446 \text{ mm}^{-1}$ T = 293 (2) K Parallelepiped  $0.44 \times 0.38 \times 0.36 \text{ mm}$ Light brown

8143 measured reflections 2824 independent reflections 2463 reflections with  $l > 2\sigma(l)$   $R_{int} = 0.021$   $\theta_{max} = 26.99^{\circ}$   $h = -6 \rightarrow 6$   $k = 0 \rightarrow 28$  $l = 0 \rightarrow 14$ 

 $\begin{array}{l} (\Delta/\sigma)_{\rm max} < 0.001 \\ \Delta\rho_{\rm max} = 1.03 \ {\rm e} \ {\rm \AA}^{-3} \\ \Delta\rho_{\rm min} = -1.08 \ {\rm e} \ {\rm \AA}^{-3} \\ {\rm Extinction \ correction: \ none} \\ {\rm Scattering \ factors \ from} \\ International \ Tables \ for \\ Crystallography \ (Vol. \ C) \end{array}$ 

## Table 1. Selected geometric parameters (Å, °)

11—C13	2.175 (6)	C12C13	1.505 (9)
C3—C4	1.357 (9)	C13—C14	1.527 (8)
C5-C6	1.356 (10)	C14016	1.395 (7)
C7C12	1.528 (8)	C14—C15	1.526 (8)
C8—C15	1.509 (7)		
011—C2—O1	115.8 (6)	01	114.5 (5)
O11C2C3	127.1 (6)	C5C10C4	125.2 (5)
C12-C7-C8-C15	1.3 (8)	C12-C13-C14-C15	64.2 (6)
C8C7C12C13	20.6 (7)	C7-C8-C15-C14	9.7 (7)
C7-C12-C13-C14	-53.9 (6)	C13-C14-C15-C8	-40.7 (6)

#### Table 2. Hydrogen-bonding geometry (Å, °)

$D = H \cdot \cdot \cdot A$	D - H	H. · · A	$D \cdot \cdot \cdot A$	$D = H \cdots A$
016—H16· · · O11'	0.82	1.95	2.761 (7)	168
Symmetry code: (i) $x = 1, \frac{1}{2} - y, \frac{1}{2} + z$ .				

After checking their presence in the difference map, the H atoms were fixed geometrically and allowed to ride on their attached atoms. The reflections 140 and 141 were suppressed during refinement as they showed very poor agreement between  $F_a^2$  and  $F_c^2$ . The highest difference peak, 1.03 e Å<sup>-3</sup>, was found near II at a distance of 0.94 Å.

Data collection: *SMART* (Siemens, 1996). Cell refinement: *SAINT* (Siemens, 1996). Data reduction: *SAINT*. Program(s) used to solve structure: *SHELXTL* (Sheldrick, 1997). Program(s) used to refine structure: *SHELXTL*. Molecular graphics: *SHELXTL*. Software used to prepare material for publication: *SHELXTL* and *PARST* (Nardelli, 1995).

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Supplementary data for this paper are available from the IUCr electronic archives (Reference: NA1400). Services for accessing these data are described at the back of the journal.

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## 7-Methyl-3-(2-methylphenyl)-1,2,4-triazolo-[3,4-*b*][1,3]benzothiazole, (I), and 7-methyl-3-(4-methylphenyl)-1,2,4-triazolo[3,4-*b*]-[1,3]benzothiazole, (II)

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#### Abstract

The two title compounds, both  $C_{16}H_{13}N_3S$ , were synthesized by photolysis of 4-(2-bromo-4-methylphenyl)-5-(2-methylphenyl)-1,2,4-triazole-3-thione and 4-(2-bromo-4-methylphenyl)-5-(4-methylphenyl)-1,2,4-triazole-3-thione, respectively. Both molecules consist of a 7-

© 1999 International Union of Crystallography Printed in Great Britain – all rights reserved methyl-1,2,4-triazolo[3,4-b][1,3]benzothiazole moiety linked to a methyl-substituted phenyl ring. The interplanar angle between the triazole-benzothiazole fused fragment and the phenyl ring is  $66.8(1)^{\circ}$  in (I) and  $35.4(1)^{\circ}$  in (II). The structures of both compounds are stabilized by weak C—H···N-type interactions.

## Comment

The benzothiazole moiety possesses many useful pharmacological properties, such as acting as antitumour (Shi *et al.*, 1996), antifungal (Kü *et al.*, 1997), antimicrobial (Kü *et al.*, 1995), antineoplastic (Alarcón *et al.*, 1998) and anti-inflammatory (Dannhardt *et al.*, 1997) agents. Condensed [1,2,4]triazoles are also medicinally important compounds (Francis & Gelotte, 1988; Kottke *et al.*, 1983). Substituted 1,2,4-triazolo[3,4-*b*]-benzothiazoles have been prepared from the respective triazole-3-thiones by a photochemical method. In order to confirm the identity of the compounds and to study the stereochemistry and the structure–activity relationships, a detailed crystallographic study has been carried out and the results are presented here.

In the following discussion, the values given without brackets are for (I), and those within square brackets are for (II). Figs. 1 and 2 show the *ORTEP* (Johnson, 1965) drawings of the two molecules in the same relative orientation. The structure consists of methylbenzothia-



zole-triazole fused fragments linked to a methylphenyl group. The N-N, C-N and S-C bond distances in the fused fragment system are comparable with those found in related structures (Ruiwu et al., 1997; Palmer & Parsons, 1996). The S1-C7 distance of 1.728(3) [1.736(3) Å], and the C7-N1 and C7-N2 distances of 1.367(3) [1.370(3)Å] and 1.307(4) [1.302(3)Å]. respectively, clearly show the resonance of the thiourea system (Valle et al., 1970). The C7-S1-C1 angle of 89.3(1) [ $89.4(1)^{\circ}$ ] suggests that the S atom uses only p-orbitals to form bonds with C1 and C7. In order to maintain the planar conformation of the thiazole ring. the other bond angles, namely, C7-N1-C6, N1-C6-C1, C6-C1-S1 and S1-C7-N1, are in the range 110.1 (2)–114.4 (2) $^{\circ}$  [109.6 (2)–114.1 (2) $^{\circ}$ ], which is less than the ideal  $120^{\circ}$  for  $sp^2$  orbital hybridizations of C and N atoms and greater than the ideal 108° required for a regular planar pentagon. These results agree with a similar structure found in 2-amino-6fluorobenzothiazole (Jai-Nhuknan et al., 1997). The