A novel heterocyclic sulfonamide: *N*-benzyl-5-[*N*-benzyl-*N*-(*tert*-butyl oxycarbonyl)amino]-*N*-(*tert*-butyl oxycarbonyl)-1,3,4-thiadiazole-2-sulfonamide

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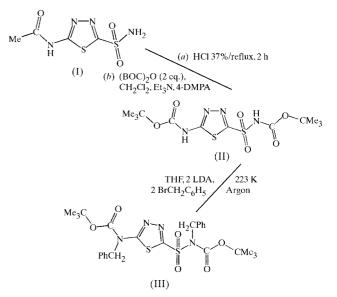
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The title compound, $C_{26}H_{32}N_4O_6S_2$, is a heterocyclic sulfonamide which is a 1,3,4-thiadiazole derivative. Structural data for this compound are compared with those of related compounds.

Comment

Since Davenport (1945) reported that thiophene-2-sulfonamide is a carbonic anhydrase (CA) inhibitor 40 times more active than other sulfonamides, many heterocyclic sulfonamides have been prepared. We have obtained and determined the crystal structures of several 1,3,4thiadiazolesulfonamides following the method proposed by Young et al. (1956) with minor modifications (Alzuet et al., 1991a; Pedregosa et al., 1993). The results obtained by our group with these sulfonamides have rendered data which indicate a notable increase in CA inhibitory properties (Chufán et al., 2000). In addition, Krebs' (1948) investigations on compounds presenting the modified sulfonamido group has led to the conclusion that substitution within the sulfonamido group drastically affects inhibitory activity and increases lipoaffinity (Duffel et al., 1986). Based on these investigations, it was interesting to prepare compounds with the substituted sulfonamido group and to determine their crystal structures. Furthermore, following our programme for systematic studies on unsubstituted and substituted heterocyclic sulfonamides and their metal complexes (Alzuet et al., 1991b; Pedregosa et al., 1995), we will study the role of these bulky groups with regard to CA inhibition and lipoaffinity.

In this work, we obtained the crystal structure of the title compound, (III), and compared it with related compounds, such as acetazolamide (5-acetylamino-1,3,4-thiadiazole-2-sulfonamide; Mathew & Palenik, 1974), 5-amino-1,3,4-thiadiazole-2-sulfonamide (Pedregosa *et al.*, 1993), 5-*tert*-butyloxycarbonylamino-1,3,4-thiadiazole-2-sulfonamide (Pedregosa *et al.*, 1995), 5-amino-1,3,4-thiadiazole-2-thiol (Downie *et al.*, 1972) and 5-*tert*-butyloxycarbonylamino-1,3,4-thiadiazole-2-sulfonyl chloride (Pedregosa *et al.*, 1996). The main features of the new structure, *i.e.* planarity of the 1,3,4-thiadiazole ring, distorted tetrahedral geometry in the sulfonamide group and a coplanar arrangement of the disubstituted 5-amino group and the 1,3,4-thiadiazole ring, are in agreement with the compared compounds.



Experimental

The synthesis of (III) was performed in three steps. In the first step, the acid hydrolysis of acetazolamide (I) to give 5-amino-1,3,4-thiadiazole-2-sulfonamide was performed following the literature procedure of Supuran et al. (1990). In the second step (group protection), the product (0.5 g, 2.77 mmol) of the previous reaction, triethylamine (Et₃N, Aldrich, 1 ml, 7.2 mmol), 4-dimethyl-aminopyridine (4-DMAP, Aldrich, 100 mg, 4.6 mmol) and di-tert-butyldicarbonate [(BOC)₂O, Aldrich, 1.2 g, 5.54 mmol] were added in order and with permanent stirring at room temperature to CH2Cl2 (50 ml, Merck) under an inert argon atmosphere. The reaction was followed by thin-layer chromatography (TLC). After 18 h, the solvent was removed by vacuum extraction. The obtained solid (II) was purified by treatment with 1 N HCl, filtered and washed with distilled water (3-4 times) and hexane (3-4 times). Recrystallization from ethanol/dichloromethane (1:1) rendered crystals that were washed with ethanol/hexane (1:20) giving 0.9 g (86%) of (II) (m.p. 436–438 K). Finally, the third step consisted of deprotonation of (II). Lithium diisopropylamide (LDA) was prepared by addition of diisopropylamine (2 ml, 14.28 mmol) and *n*-butyllithium (1.32 ml, 14.23 mmol) to tetrahydrofuran (50 ml, Supelco) on an acetone/solid CO₂ cold bath (213 K) under an inert argon atmosphere. The mixture was stabilized at 213 K for 30 min. Subsequently, (II) (0.5 g,

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1.32 mmol) was added. After 30 min of stabilization at the same temperature, benzylbromide (0.4 ml, 3.29 mmol) was added. The reaction was followed by TLC. After 24 h, when the bath reached room temperature, the solvent was removed by vacuum extraction. The obtained solid (III), was purified with 1 N HCl and washed with distilled water and hexane. Recrystallization was performed as described in step two and 0.55 g (75% yield) of (III) were isolated (m.p. 389–391 K).

Crystal data

| $C_{26}H_{32}N_4O_6S_2$ |
|--------------------------------|
| $M_r = 560.68$ |
| Monoclinic, $P2_1/n$ |
| a = 15.408 (1) Å |
| b = 10.256 (1) Å |
| c = 19.402 (1) Å |
| $\beta = 107.91 \ (1)^{\circ}$ |
| $V = 2917.4 (4) \text{ Å}^3$ |
| Z = 4 |

Data collection

 $\begin{array}{l} {\rm Enraf-Nonius\ CAD-4\ diffract-}\\ {\rm ometer\ }\\ \omega/2\theta\ {\rm scans\ }\\ 5807\ {\rm measured\ reflections\ }\\ 5114\ {\rm independent\ reflections\ }\\ 3022\ {\rm reflections\ with\ }I>2\sigma(I)\ R_{\rm int\ }=0.0074 \end{array}$

Refinement

Refinement on F^2 $R[F^2 > 2\sigma(F^2)] = 0.049$ $wR(F^2) = 0.156$ S = 1.0575114 reflections 360 parameters Mo $K\alpha$ radiation Cell parameters from 25 reflections $\theta = 9.5-13.6^{\circ}$ $\mu = 0.227 \text{ mm}^{-1}$ T = 293 (2) K Prismatic, colourless $0.30 \times 0.25 \times 0.20 \text{ mm}$

 $D_x = 1.277 \text{ Mg m}^{-3}$

 $\theta_{\max} = 24.99^{\circ}$ $h = 0 \rightarrow 18$ $k = 0 \rightarrow 12$ $l = -23 \rightarrow 21$ 3 standard reflections frequency: 90 min intensity decay: none

| Н | -atom parameters constrained |
|----|-----------------------------------------------------|
| w | $r = 1/[\sigma^2(F_o^2) + (0.0755P)^2]$ |
| | where $P = (F_o^2 + 2F_c^2)/3$ |
| (4 | $\Delta/\sigma)_{\rm max} = 0.001$ |
| Δ | $\rho_{\rm max} = 0.23 \ {\rm e} \ {\rm \AA}^{-3}$ |
| Δ | $A \rho_{\rm min} = -0.27 \text{ e} \text{ Å}^{-3}$ |

The H atoms were placed geometrically and treated with a riding model in which the C-H distance, common to all H atoms bonded to the same C, was free to refine.

Data collection: *CAD-4 Software* (Enraf–Nonius, 1989); cell refinement: *CAD-4 Software*; data reduction: *XRAY*80 (Stewart, 1980); program(s) used to solve structure: *SIR*92 (Altomare *et al.*, 1994); program(s) used to refine structure: *SHELXL*97 (Sheldrick, 1997).

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