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Crystal structure of 5-(adamantan-1-yl)-3-[(4-trifluoromethylanilino)methyl]-2,3-dihydro-1,3,4-oxadiazole-2-thione, $C_{20}H_{22}F_3N_3OS$

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Abstract

$C_{20}H_{22}F_3N_3OS$, triclinic, $P1$ (no. 1), $a = 6.9678(8)$ Å, $b = 10.7614(14)$ Å, $c = 13.0503(14)$ Å, $\alpha = 76.870(3)^\circ$, $\beta = 88.004(4)^\circ$, $\gamma = 87.275(4)^\circ$, $V = 951.60(19)$ Å³, $Z = 2$, $R_{gt}(F) = 0.0629$, $wR_{ref}(F^2) = 0.1626$, $T = 100$ K.

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Table 1 contains crystallographic data and Table 2 contains the list of the atoms including atomic coordinates and displacement parameters.

Source of material

Formaldehyde solution (37%, 1.5 mL) and 4-trifluoromethylaniline (1.61 g, 0.01 mol) were added to a solution of 5-(adamantan-1-yl)-1,3,4-oxadiazole-2(3H)-thione (2.36 g, 0.01 mol) in ethanol (15 mL), and the mixture was stirred at 293 K for 2 h and allowed to stand overnight. The precipitated crude product was filtered, washed with water, dried and crystallised from aqueous ethanol to yield 3.36 g (82%) of the title compound as colourless prisms. M.pt: 447–449 K (uncorrected). **Anal. Calc.** for $C_{20}H_{22}F_3N_3OS$: C, 58.67; H, 5.42; N, 10.26; S, 7.83%. Found: C, 58.63; H, 5.45;

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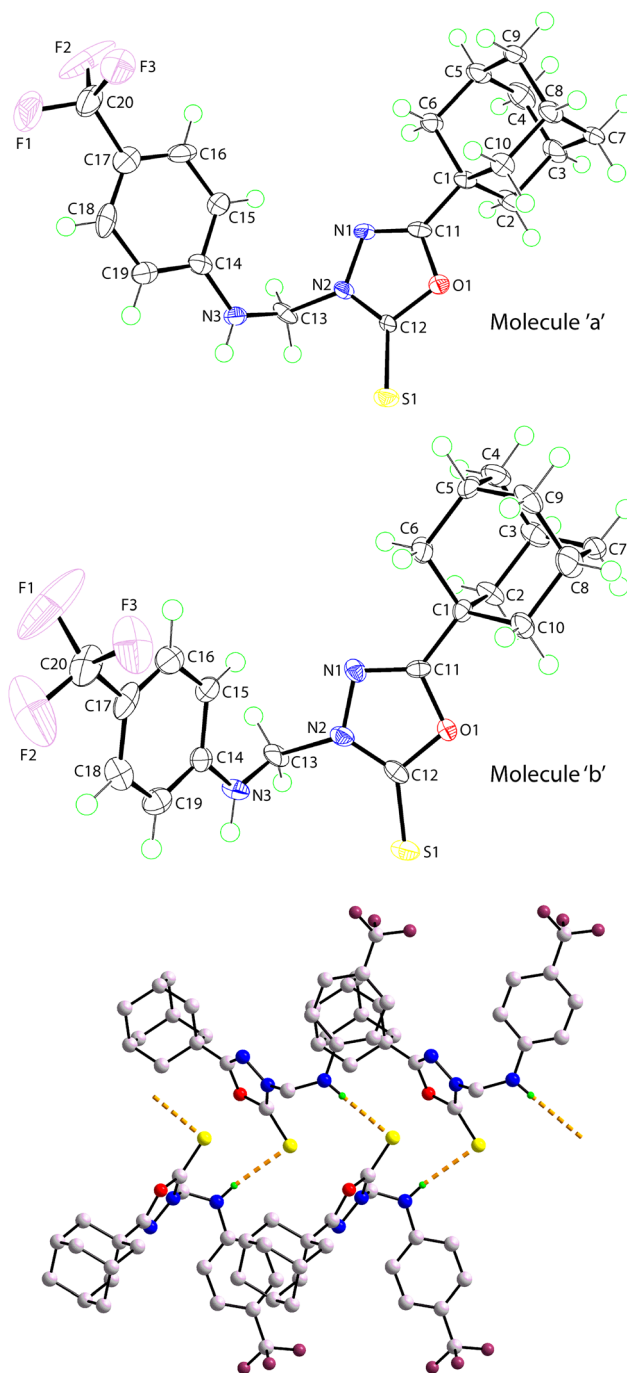


Table 1: Data collection and handling.

Crystal:	Colourless prism
Size:	0.54 × 0.23 × 0.11 mm
Wavelength:	Mo K α radiation (0.71073 Å)
μ :	0.22 mm ⁻¹
Diffractometer, scan mode:	Bruker APEX-II D8 venture, φ and ω
$\theta_{\max}$, completeness:	25.0°, >99%
$N(hkl)_{\text{measured}}$, $N(hkl)_{\text{unique}}$, R_{int} :	21,069, 6660, 0.041
Criterion for I_{obs} , $N(hkl)_{\text{gt}}$:	$I_{\text{obs}} > 2\sigma(I_{\text{obs}})$, 6248
$N(\text{param})_{\text{refined}}$:	511
Programs:	Bruker [1], SHELX [2, 3], WinGX/ORTEP [4]

Table 2: Fractional atomic coordinates and isotropic or equivalent isotropic displacement parameters (Å²).

Atom	<i>x</i>	<i>y</i>	<i>z</i>	<i>U</i> _{iso} */ <i>U</i> _{eq}
S1A	0.3734 (2)	−0.06030 (14)	0.44740 (12)	0.0166 (4)
O1A	0.1263 (6)	0.0610 (4)	0.3004 (3)	0.0131 (9)
N1A	0.0897 (7)	0.2555 (5)	0.3300 (4)	0.0136 (11)
N2A	0.2170 (7)	0.1807 (5)	0.4005 (4)	0.0133 (10)
N3A	0.4741 (7)	0.2654 (5)	0.4833 (4)	0.0153 (11)
H3AB	0.572 (7)	0.215 (6)	0.510 (5)	0.018*
F1A	0.8885 (10)	0.7712 (6)	0.2557 (4)	0.0644 (19)
F2A	0.5965 (10)	0.8309 (5)	0.2232 (5)	0.0644 (18)
F3A	0.7488 (7)	0.7206 (4)	0.1308 (3)	0.0379 (11)
C1A	−0.1006 (9)	0.2087 (6)	0.1860 (4)	0.0131 (13)
C2A	−0.2851 (9)	0.1361 (7)	0.2270 (5)	0.0188 (14)
H2AA	−0.254193	0.043201	0.247696	0.023*
H2AB	−0.338981	0.164466	0.289477	0.023*
C3A	−0.4324 (9)	0.1639 (7)	0.1390 (5)	0.0211 (14)
H3AA	−0.551853	0.117547	0.164934	0.025*
C4A	−0.4816 (9)	0.3078 (7)	0.1086 (5)	0.0235 (15)
H4AA	−0.578978	0.325877	0.053066	0.028*
H4AB	−0.535929	0.337093	0.170541	0.028*
C5A	−0.2999 (10)	0.3787 (6)	0.0684 (5)	0.0193 (14)
H5AA	−0.332776	0.472425	0.048326	0.023*
C6A	−0.1501 (9)	0.3533 (6)	0.1556 (5)	0.0157 (13)
H6AA	−0.202663	0.382915	0.217727	0.019*
H6AB	−0.032805	0.400366	0.129951	0.019*
C7A	−0.3492 (10)	0.1177 (6)	0.0425 (5)	0.0182 (14)
H7AA	−0.316444	0.024952	0.062525	0.022*
H7AB	−0.445849	0.132579	−0.013265	0.022*
C8A	−0.1686 (10)	0.1909 (6)	0.0014 (5)	0.0179 (13)
H8AA	−0.114654	0.161871	−0.061632	0.022*
C9A	−0.2162 (9)	0.3350 (6)	−0.0287 (5)	0.0170 (13)
H9AA	−0.310615	0.354184	−0.085687	0.020*
H9AB	−0.098380	0.381364	−0.054519	0.020*
C10A	−0.0181 (9)	0.1631 (6)	0.0893 (5)	0.0153 (13)
H10A	0.014370	0.070330	0.109004	0.018*
H10B	0.100871	0.208062	0.063410	0.018*
C11A	0.0406 (8)	0.1820 (6)	0.2727 (4)	0.0116 (12)
C12A	0.2407 (8)	0.0624 (6)	0.3843 (4)	0.0103 (12)
C13A	0.2859 (9)	0.2254 (6)	0.4916 (5)	0.0158 (13)
H13A	0.199478	0.297127	0.502702	0.019*
H13B	0.274240	0.155203	0.555075	0.019*

Table 2: (continued)

Atom	<i>x</i>	<i>y</i>	<i>z</i>	<i>U</i> _{iso} */ <i>U</i> _{eq}
C14A	0.5328 (9)	0.3822 (6)	0.4224 (5)	0.0143 (13)
C15A	0.4044 (10)	0.4654 (6)	0.3557 (5)	0.0194 (14)
H15A	0.275917	0.442040	0.350387	0.023*
C16A	0.4670 (11)	0.5816 (6)	0.2979 (5)	0.0220 (15)
H16A	0.379578	0.639995	0.255293	0.026*
C17A	0.6570 (11)	0.6128 (7)	0.3021 (5)	0.0223 (15)
C18A	0.7850 (10)	0.5317 (7)	0.3681 (5)	0.0216 (15)
H18A	0.914460	0.553968	0.371951	0.026*
C19A	0.7184 (9)	0.4165 (6)	0.4287 (5)	0.0172 (13)
H19A	0.803501	0.360982	0.475187	0.021*
C20A	0.7246 (12)	0.7329 (7)	0.2305 (6)	0.0302 (18)
S1B	0.8594 (2)	1.06071 (14)	0.55707 (12)	0.0181 (4)
O1B	0.6111 (6)	0.9361 (4)	0.7018 (3)	0.0121 (9)
N1B	0.6062 (7)	0.7430 (5)	0.6693 (4)	0.0138 (11)
N2B	0.7321 (7)	0.8202 (5)	0.6002 (4)	0.0144 (11)
N3B	1.0073 (8)	0.7352 (5)	0.5130 (4)	0.0166 (11)
H3BB	1.098 (8)	0.787 (6)	0.484 (5)	0.020*
F1B	1.1988 (13)	0.1684 (5)	0.7620 (5)	0.081 (2)
F2B	1.4793 (10)	0.2431 (6)	0.7529 (4)	0.072 (2)
F3B	1.2775 (7)	0.2737 (5)	0.8712 (3)	0.0413 (12)
C1B	0.3907 (9)	0.7883 (6)	0.8138 (5)	0.0116 (12)
C2B	0.1965 (9)	0.8522 (7)	0.7710 (5)	0.0171 (13)
H2BA	0.209521	0.945584	0.746741	0.021*
H2BB	0.159649	0.818357	0.710316	0.021*
C3B	0.0406 (9)	0.8245 (7)	0.8585 (5)	0.0197 (14)
H3BA	−0.084207	0.866512	0.831068	0.024*
C4B	0.0191 (9)	0.6804 (7)	0.8921 (5)	0.0227 (15)
H4BA	−0.084525	0.661078	0.946554	0.027*
H4BB	−0.016148	0.646910	0.830956	0.027*
C5B	0.2072 (10)	0.6167 (6)	0.9358 (5)	0.0182 (14)
H5BA	0.191746	0.522441	0.958370	0.022*
C6B	0.3673 (10)	0.6434 (6)	0.8500 (5)	0.0187 (14)
H6BA	0.489875	0.601295	0.878336	0.022*
H6BB	0.333932	0.608230	0.789419	0.022*
C7B	0.0991 (10)	0.8778 (6)	0.9527 (5)	0.0199 (14)
H7BA	0.114163	0.971073	0.929898	0.024*
H7BB	−0.002375	0.862112	1.008562	0.024*
C8B	0.2873 (10)	0.8127 (7)	0.9953 (5)	0.0242 (15)
H8BA	0.323632	0.846582	1.056910	0.029*
C9B	0.2664 (10)	0.6671 (6)	1.0307 (5)	0.0198 (14)
H9BA	0.389877	0.625215	1.057680	0.024*
H9BB	0.167595	0.647761	1.087787	0.024*
C10B	0.4459 (9)	0.8404 (6)	0.9099 (5)	0.0168 (13)
H10C	0.569004	0.799209	0.938070	0.020*
H10D	0.462561	0.933660	0.887926	0.020*
C11B	0.5385 (8)	0.8157 (6)	0.7282 (4)	0.0116 (12)
C12B	0.7359 (8)	0.9358 (6)	0.6192 (5)	0.0136 (13)
C13B	0.8156 (9)	0.7771 (6)	0.5067 (5)	0.0168 (13)
H13C	0.738634	0.706928	0.494871	0.020*
H13D	0.801939	0.848788	0.444338	0.020*
C14B	1.0726 (9)	0.6173 (6)	0.5754 (5)	0.0157 (13)
C15B	0.9477 (10)	0.5316 (6)	0.6376 (5)	0.0183 (13)
H15B	0.813251	0.550759	0.638783	0.022*
C16B	1.0248 (10)	0.4166 (7)	0.6984 (5)	0.0216 (15)
H16B	0.940911	0.356650	0.739494	0.026*
C17B	1.2184 (11)	0.3892 (6)	0.6996 (5)	0.0214 (15)

Table 2: (continued)

Atom	<i>x</i>	<i>y</i>	<i>z</i>	U_{iso}^*/U_{eq}
C18B	1.3421 (9)	0.4726 (6)	0.6362 (5)	0.0183 (14)
H18B	1.476321	0.452406	0.635334	0.022*
C19B	1.2684 (10)	0.5859 (6)	0.5737 (5)	0.0188 (14)
H19B	1.352845	0.642595	0.529361	0.023*
C20B	1.2923 (12)	0.2697 (7)	0.7700 (5)	0.0284 (17)

N, 10.24; S, 7.82%. **¹H-NMR** (CDCl₃, 500.13 MHz): δ 1.74–1.80 (m, 6H, Adamantane-H), 1.97–1.99 (m, 6H, Adamantane-H), 2.12 (s, 3H, Adamantane-H), 5.46 (s, 1H, NH), 5.78 (d, 2H, CH₂, *J* = 6.8 Hz), 6.96 (d, 2H, Ar-H, *J* = 8.5 Hz), 7.43 (d, 2H, Ar-H, *J* = 8.5 Hz). **¹³C-NMR** (CDCl₃, 125.76 MHz): δ 27.41, 34.33, 36.04, 39.04 (adamantane-C), 59.0 (CH₂), 123.85 (CF₃), 114.50, 119.88, 128.09, 147.12 (Ar-C), 165.45 (oxadiazole-C), 177.43 (C=S). **ESI-MS** *m/z*: 410.3 [M + H]⁺.

Experimental details

The C-bound H atoms were geometrically placed (C–H = 0.95–1.00 Å) and refined as riding with $U_{iso}(H) = 1.2U_{eq}(C)$. The N-bound H atoms were located from a difference Fourier map and refined with N–H = 0.88 + –0.01 Å, and with $U_{iso}(H)$ set to $1.2U_{eq}(N)$.

Comment

The highly lipophilic adamantane cage is frequently found in biologically-active compounds possessing various pharmacological activities [5–7]. Amantadine was discovered early as an efficient drug for the control of Influenza A infection [8, 9] and further approved for the treatment of Parkinson's disease [10]. Further studies based on adamantane derivatives resulted in the evolution of more active anti-viral drugs such as tromantadine [11] and rimantadine [12]. Adamantane-based analogues were also reported to display significant inhibitory effects against human immunodeficiency virus (HIV) [13–15]. The adamantane ethylenediamine analogue, SQ109, and the related dipiperidine derivative, SQ609, were further approved as efficient therapies against drug-susceptible and drug-resistant *Mycobacterium tuberculosis* strains [16, 17]. The adamantane carboxamide derivative Opaganib (ABC294640) is a newly approved anti-tumour drug for treating patients suffering from advanced solid tumours [18–20]. Furthermore, various adamantane derivatives were recognised as efficient anti-bacterial and anti-fungal candidates [21–24]. On the other

hand, the 1,3,4-oxadiazole nucleus was reported to constitute the core of several chemotherapeutic agents possessing anti-bacterial [25, 26], anti-fungal [27], anti-cancer [28, 29] and anti-viral [30] activities. Besides, 1,3,4-oxadiazole derivatives are presently utilised as safe herbicides for crop protection [31]. In an earlier study, we described the synthesis, anti-HIV, and anti-bacterial activities of adamantane-1,3,4-oxadiazole hybrid N-Mannich bases [15]. The previously reported compounds showed strong inhibitory activity against Gram-positive bacteria and limited activity against Gram-negative bacteria. The anti-bacterial structure-activity relationship of these compounds proved that the spectrum of the anti-bacterial efficacy is fundamentally contingent on the nature of the substituents of the aryl moiety. Further optimisation studies based on the previously prepared compounds resulted in development of the title compound (I), which exhibited strong broad-spectrum anti-bacterial activity.

The molecular structures of the independent molecules, i.e. molecule a and molecule b, comprising the crystallographic asymmetric-unit of (I) are shown in the figure (70% probability ellipsoids). The independent molecules resemble each other very closely as seen in the r.m.s. bond and angle fits of 0.0128 Å and 0.745°, respectively [32]. Crucially, there is no pseudo centre of inversion between the molecules, they being directly superimposable.

Each molecule comprises a central, planar 1,3,4-oxadiazole core connected to a thione-S [1.649(6) & 1.665(6) Å for molecules a and b, respectively], an N-bound (4-trifluoromethylanilino)methyl group and the remaining carbon atom carries the adamantan-1-yl group. Within the five-membered ring, the C11–N1 bond of 1.270(8) Å is significantly shorter than the C12–N2 bond of 1.339(8) Å, and the C11–O1 [1.383(7) Å] and C12–O1 [1.379(7) Å] bond lengths are experimentally equivalent, results suggesting limited delocalisation of π -electron density in the ring; the equivalent bond lengths for molecule b are 1.280(8), 1.324(8), 1.379(7) and 1.361(7) Å, respectively. The dihedral angle between the five- and six-membered rings is 47.1(3)°; the equivalent angle for molecule b is 51.3(3)°. Globally, the two central ring-substituents tend to be oriented to the same side of the molecule in a direction away from the thione-S atom.

There are four closely related structures in the literature, namely the phenyl compound, i.e. the parent compound [33], the 4-fluorophenyl [34], the 4-chlorophenyl [35] and the 2-trifluoromethyl [36] derivatives. Each of the previously reported structure resemble that noted above for (I).

The most prominent feature of the molecular packing in the crystal of (I) is a twisted chain aligned along the *a*-axis. The chains comprise alternating independent molecules

and feature amine-N–H...S(thione) hydrogen bonds [N3a–H3ab...S1bⁱ: H3ab...S1bⁱ = 2.54(6) Å, N3a–H3ab...S1bⁱ = 3.412(5) Å with angle at H3ab = 171(6)° and N3b–H3bb...S1aⁱⁱ: H3bb...S1aⁱⁱ = 2.55(6) Å, N3b–H3bb...S1aⁱⁱ = 3.407(6) Å with angle at H3bb = 166(5)° for symmetry operations (i): $x, -1 + y, z$ and (ii): $1 + x, 1 + y, z$]. A portion of the supramolecular chain is shown in the lower view of the Figure, where the hydrogen bonds are shown as dashed lines and non-participating hydrogen atoms are omitted.

As indicated above, the molecular structures are very similar as is the primary mode of association between the molecules. This is confirmed by the calculation of the Hirshfeld surfaces and the full and decomposed two-dimensional fingerprint plots. Crystal Explorer 17 [37] was employed for this purpose following established procedures [38]. The differences between the Hirshfeld surfaces for each independent molecule differ by no more than 1%. The most significant contribution are from H...H contacts being 39.9 and 39.2% for molecules a and b, respectively. The next most important contributions are of the type F...H/H...F, i.e. 24.5 and 24.0%, respectively. There are also significant contributions to the Hirshfeld surface by C...H/H...C [13.6 and 13.9%], S...H/H...S [9.0 and 9.8%], N...H/H...N [4.1 and 4.3%] and O...H/H...O [3.4 and 3.5%]. The only other surface contacts greater than 1.0% are S...F/F...S [1.5 and 0.9%], S...N/N...S [1.2 and 1.2%] and S...C/C...S [1.0 and 1.0%].

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