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## Key indicators

Single-crystal X-ray study
$T=293 \mathrm{~K}$
Mean $\sigma(\mathrm{C}-\mathrm{C})=0.004 \AA$
Disorder in main residue
$R$ factor $=0.063$
$w R$ factor $=0.186$
Data-to-parameter ratio $=12.5$

For details of how these key indicators were automatically derived from the article, see http://journals.iucr.org/e.
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## 6,7-Dimethoxy- N -[3-(trifluoromethyl)phenyl]-quinazolin-4-amine ethanol disolvate

The title compound, $\mathrm{C}_{17} \mathrm{H}_{14} \mathrm{~F}_{3} \mathrm{~N}_{3} \mathrm{O}_{2} \cdot 2 \mathrm{C}_{2} \mathrm{H}_{5} \mathrm{OH}$, was synthesized by the reaction of 4-chloro-6,7-dimethoxyquinazoline and 3-(trifluromethyl)aniline at 580 W in a domestic microwave oven and obtained in $93 \%$ yield. The bond lengths and angles are normal. Intermolecular $\mathrm{N}-\mathrm{H} \cdots \mathrm{O}, \mathrm{O}-\mathrm{H} \cdots \mathrm{N}$ and $\mathrm{O}-\mathrm{H} \cdots \mathrm{O}$ hydrogen bonds and $\pi-\pi$ stacking interactions between the quinazoline rings of neighbouring molecules stabilize the crystal packing.

## Comment

Recently, epidermal growth factor receptor (EGFR) has became one of the significant target proteins in medicinal developments (Arteaga, 2002), since its excess always leads to a variety of vicious tumours and cancers. Among the small molecular inhibitors of EGFR, 4-anilinoquinazoline derivatives are clinically confirmed to be among the most effective compounds (Hou et al., 2002). The results of quantitative structure-activity relationship research indicate that electrondonating groups at the 6,7-positions will improve their degree of EGFR inhibition (Bridges et al., 1996). We report here the synthesis and crystal structure of the title compound, (I).

(I)

In compound (I) (Fig. 1), the bond lengths and angles are normal (Table 1); two ethanol solvent molecules complete the asymmetric unit. The quinazoline system and non-fused benzene ring make a dihedral angle of 43.2 (1) ${ }^{\circ}$. Interestingly, atoms N1, C6, C8 and H3 are almost coplanar, with an r.m.s. deviation of 0.026 (8) $\AA$, in spite of steric hindrance between the quinazoline system and the non-fused benzene ring.

The solvent molecules play an important role in the crystal formation, participating in a number of hydrogen bonds with the quinazolin-4-amine molecules (Table 2). These hydrogen bonds stabilize the crystal packing (Fig. 2), along with $\pi-\pi$

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Figure 1
View of (I), showing the atom-labelling scheme and displacement ellipsoids at the $40 \%$ probability level. The two ethanol solvent molecules and the minor component of the disordered $\mathrm{CF}_{3}$ group have been omitted for clarity. H atoms are represented by circles of arbitrary size.
stacking interactions between the quinazoline rings of neighbouring molecules; the interplanar distance to the symmetryrelated molecule at $(1-x, 2-y, 2-z)$ is $3.356(8) \AA$.

## Experimental

At room temperature, basic alumina ( 2 g ) was added to a solution of 4-chloro-6,7-dimethoxyquinazoline ( 1 mmol ) and 3-(trifluromethyl)aniline ( 1 mmol ) in diethyl ether ( 10 ml ). After removal of the ether, the solid was poured into an open vessel and irradiated for 3 min at 580 W in a domestic microwave oven. After cooling to room temperature, ethanol ( 15 ml ) was added and the filtrate was concentrated on a rotary evaporator. The residue was isolated by silica column chromatography with $\mathrm{CH}_{2} \mathrm{Cl}_{2}-\mathrm{Et}_{2} \mathrm{O}(5: 3 \mathrm{v} / \mathrm{v})$ as the eluent in $93 \%$ yield. Suitable crystals were obtained by evaporation of an ethanol-water ( $1: 1 \mathrm{v} / \mathrm{v}$ ) mixed solution (m.p. 442-443 K). ${ }^{1} \mathrm{H} \operatorname{NMR}\left(\mathrm{CDCl}_{3}\right): \delta 4.01(s, 3 \mathrm{H}), 4.03(s, 3 \mathrm{H}), 7.08(s, 1 \mathrm{H}), 7.27(d, 1 \mathrm{H}$, $J=7.5 \mathrm{~Hz}), 7.38-7.50(m, 2 \mathrm{H}), 7.51(t, 1 \mathrm{H}, J=8.0 \mathrm{~Hz}), 8.70(s, 1 \mathrm{H})$. IR ( $\mathrm{KBr}, \nu \mathrm{cm}^{-1}$ ): 3448, 2925, 2851, 1625, 1584, 1516, 1449, 1332, 1242, $1166,1123,1069,993,793,700$. Analysis calculated for $\mathrm{C}_{21} \mathrm{H}_{26} \mathrm{~F}_{3} \mathrm{~N}_{3} \mathrm{O}_{4}$ : C 57.14, H 5.94, N 9.52\%; found C 57.39, H 5.78, N $9.67 \%$.

## Crystal data

$\mathrm{C}_{17} \mathrm{H}_{14} \mathrm{~F}_{3} \mathrm{~N}_{3} \mathrm{O}_{2} \cdot 2 \mathrm{C}_{2} \mathrm{H}_{6} \mathrm{O}$
$M_{r}=441.45$
Triclinic, $P \overline{1}$
$a=8.7035(7) \AA$
$b=9.7426(8) \AA$
$c=14.5212(9) \AA$
$\alpha=70.652(4)^{\circ}$
$\beta=89.493(1)^{\circ}$
$\gamma=77.484(3)^{\circ}$
$V=1131.52(15) \AA^{\circ}$

$$
\begin{aligned}
& M_{r}=441.45 \\
& \text { Triclinic. } P \overline{1}
\end{aligned}
$$

$$
Z=2
$$

$$
\text { Triclinic, } P 1
$$

$$
a=8.7035(7) \AA
$$

$$
b=9.7426 \text { (8) A }
$$

$$
c=14.5212(9) \AA
$$

$$
\alpha=70.652(4)^{\circ}
$$

$$
\gamma=77.484(3)^{\circ}
$$

$$
V=1131.52(15) \AA^{3}
$$



Figure 2
The molecular packing, showing the hydrogen-bonding interactions (dashed lines). C -bound H atoms have been omitted for clarity.

## Data collection

Rigaku R-AXIS RAPID
3972 independent reflections
3264 reflections with $I>2 \sigma(I)$
diffractometer
$\omega$ scans
Absorption correction: multi-scan
(ABSCOR; Higashi, 1995)
$T_{\text {min }}=0.956, T_{\text {max }}=0.969$
8371 measured reflections
$R_{\text {int }}=0.058$
$\theta_{\text {max }}=25.2^{\circ}$
$h=-10 \rightarrow 10$
$k=-11 \rightarrow 11$
8371 measured reflections
$l=-17 \rightarrow 17$

## Refinement

Refinement on $F^{2}$
$R\left[F^{2}>2 \sigma\left(F^{2}\right)\right]=0.063$
$w R\left(F^{2}\right)=0.186$
$S=1.08$
3972 reflections
318 parameters
H atoms treated by a mixture of independent and constrained refinement

$$
\begin{aligned}
& \begin{array}{l}
w=1 /\left[\sigma^{2}\left(F_{\mathrm{o}}^{2}\right)+(0.0909 P)^{2}\right. \\
\quad \quad+0.4785 P] \\
\quad \text { where } P=\left(F_{\mathrm{o}}^{2}+2 F_{\mathrm{c}}^{2}\right) / 3 \\
(\Delta / \sigma)_{\max }=0.001 \\
\Delta \rho_{\max }=0.48 \mathrm{e} \AA^{-3} \\
\Delta \rho_{\min }=-0.44 \mathrm{e} \AA^{-3} \\
\text { Extinction correction: } S H E L X L 97 \\
\text { Extinction coefficient: } 0.028(5)
\end{array}
\end{aligned}
$$

Table 1
Selected geometric parameters ( $\left({ }^{\circ},{ }^{\circ}\right)$.

| F1-C1 | $1.282(6)$ | $\mathrm{N} 1-\mathrm{H} 3$ | $0.86(3)$ |
| :--- | :--- | :--- | :--- |
| F2-C1 | $1.320(7)$ | $\mathrm{N} 2-\mathrm{C} 8$ | $1.328(3)$ |
| F3-C1 | $1.307(6)$ | $\mathrm{N} 2-\mathrm{C} 9$ | $1.341(3)$ |
| N1-C8 | $1.362(3)$ | $\mathrm{N} 3-\mathrm{C} 9$ | $1.319(3)$ |
| N1-C6 | $1.409(3)$ | $\mathrm{N} 3-\mathrm{C} 10$ | $1.371(3)$ |
|  |  |  |  |
| C13-O3-C17 | $116.34(17)$ | $\mathrm{C} 9-\mathrm{N} 3-\mathrm{C} 10$ | $115.86(19)$ |
| C14-O4-C16 | $117.31(19)$ | $\mathrm{F} 1-\mathrm{C} 1-\mathrm{C} 2$ | $114.1(3)$ |
| C8-N1-C6 | $126.92(18)$ | $\mathrm{F} 3-\mathrm{C} 1-\mathrm{C} 2$ | $113.4(4)$ |
| C8-N1-H3 | $119.6(16)$ | $\mathrm{F} 2-\mathrm{C} 1-\mathrm{C} 2$ | $111.1(4)$ |
| C6-N1-H3 | $112.5(16)$ |  |  |

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

| $D-\mathrm{H} \cdots A$ | $D-\mathrm{H}$ | $\mathrm{H} \cdots A$ | $D \cdots A$ | $D-\mathrm{H} \cdots A$ |
| :--- | :--- | :--- | :--- | :--- |
| $\mathrm{~N} 1-\mathrm{H} 3 \cdots \mathrm{O} 1$ | $0.86(3)$ | $2.12(3)$ | $2.965(3)$ | $169(2)$ |
| $\mathrm{O} 2-\mathrm{H} 2 \cdots \mathrm{~N}^{\mathrm{i}}$ | 0.82 | 1.93 | $2.745(3)$ | 173 |
| $\mathrm{O} 1-\mathrm{H} 1 \cdots \mathrm{O}^{2 i}$ | 0.82 | 1.90 | $2.690(3)$ | 162 |

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

All H atoms were initially located in a difference Fourier map. All methyl H atoms were then constrained to an ideal geometry, with

## organic papers

$\mathrm{C}-\mathrm{H}$ distances of $0.96 \AA$ and $U_{\mathrm{iso}}(\mathrm{H})=1.5 U_{\text {eq }}(\mathrm{C})$, but each group was allowed to rotate freely about the $\mathrm{C}-\mathrm{C}$ bond. The aromatic and hydroxyl H atoms were placed in geometrically idealized positions and constrained to ride on their parent atoms, with $\mathrm{C}-\mathrm{H}=0.93 \AA$, $\mathrm{O}-\mathrm{H}=0.82 \AA$, and $U_{\text {iso }}(\mathrm{H})=1.2 U_{\text {eq }}(\mathrm{C})$ or $1.5 U_{\text {eq }}(\mathrm{O})$. The amine H atom was refined isotropically. The $\mathrm{CF}_{3}$ group is disordered and the major and minor orientations have refined occupancies of 0.70 (1) and 0.30 (1), respectively. The atomic displacement parameters of one ethanol solvent molecule ( $\mathrm{O} 2 / \mathrm{C} 20 / \mathrm{C} 21$ ) are significantly higher than those of the other solvent molecule, indicating a possible partial disorder of the former residue.
Data collection: RAPID-AUTO (Rigaku, 1998); cell refinement: RAPID-AUTO; data reduction: CrystalStructure (Rigaku/MSC, 2002); program(s) used to solve structure: SHELXS97 (Sheldrick, 1997); program(s) used to refine structure: SHELXL97 (Sheldrick, 1997); molecular graphics: ORTEP3 (Farrugia, 1997); software used to prepare material for publication: SHELXL97.

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