

7-Methoxy-4-morpholino-6-(3-morpholino-propoxy)quinazoline trihydrate

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Key indicators

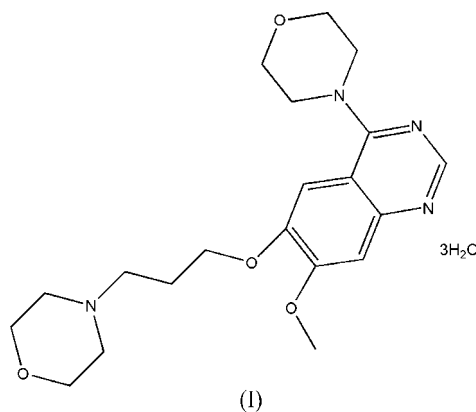
Single-crystal X-ray study
 $T = 293$ K
Mean $\sigma(\text{C}-\text{C}) = 0.007$ Å
 R factor = 0.080
 wR factor = 0.205
Data-to-parameter ratio = 15.9For details of how these key indicators were
automatically derived from the article, see
<http://journals.iucr.org/e>.

In the title compound, $\text{C}_{20}\text{H}_{28}\text{N}_4\text{O}_4 \cdot 3\text{H}_2\text{O}$, the two morpholine rings adopt chair conformations. The crystal packing is stabilized by intermolecular $\text{O}-\text{H} \cdots \text{N}$ and $\text{O}-\text{H} \cdots \text{O}$ hydrogen bonds involving the water molecules.

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Comment

Epidermal growth factor receptor (EGFR) has become one of the significant target proteins in drug discovery (Kamath & Buolamwini, 2006), since its excess always leads to a variety of cancers. Among the small molecular inhibitors of EGFR, the quinazoline ring system, including 4-anilinoquinazoline and its derivatives, has been widely investigated. Many small molecular inhibitors of EGFR are used as anticancer drugs (Speake *et al.*, 2005), such as gefitinib (Levin *et al.*, 2002) and erlotinib (Sorbera *et al.*, 2002). As part of our studies in this area, we have focused our attention on 4-aminoquinazoline derivatives, and now report the synthesis and structure of the title compound, (I) (Fig. 1).



In compound (I), the two morpholine rings adopt chair conformations. For the C17–C20/N4/O4 ring, N4 and O4 deviate from the mean plane of C17/C18/C19/C20 by 0.627 (6) and -0.641 (6) Å, respectively. Atoms N1 and O1 deviate from the plane of C1–C4 by 0.655 (7) and -0.624 (7) Å, respectively.

In the crystal structure of (I), the constituent species interact by way of $\text{O}-\text{H} \cdots \text{N}$ and $\text{O}-\text{H} \cdots \text{O}$ hydrogen bonds (Table 1 and Fig. 2), involving the water molecules.

Experimental

Potassium iodide (0.1 g) was added to a solution of 4-chloro-6-(3-chloropropoxy)-7-methoxyquinazoline (0.51 g, 1.8 mmol) in morpholine (10 ml) and the mixture was stirred at 343 K for 15 min.

Water (50 ml) was added to the reaction mixture, which was then extracted with chloroform (3 × 10 ml), washed with brine (10 ml), dried over anhydrous sodium sulfate and concentrated under reduced pressure. The crude product was crystallized from ethyl acetate, to afford the title compound as a crystalline solid. Crystals of (I) were obtained from ethyl acetate, by slow evaporation at room temperature (m.p. 372–373 K). Analysis calculated for C₂₀H₃₄N₄O₇: C 54.28, H 7.74, N 12.66%; found C 54.37, H 7.82, N 12.80%.

Crystal data

C₂₀H₂₈N₄O₄·3H₂O
M_r = 442.51
 Monoclinic, *P*2₁/*c*
a = 10.437 (2) Å
b = 10.438 (2) Å
c = 21.004 (4) Å
 β = 97.17 (3)°
V = 2270.3 (8) Å³
Z = 4
 Mo *K*α radiation
 μ = 0.10 mm⁻¹
T = 293 (2) K
 0.40 × 0.20 × 0.10 mm

Data collection

Enraf–Nonius CAD-4 diffractometer
 Absorption correction: ψ scan (North *et al.*, 1968)
*T*_{min} = 0.962, *T*_{max} = 0.990
 4457 measured reflections
 4457 independent reflections
 2267 reflections with *I* > 2σ(*I*)
 3 standard reflections every 200 reflections
 intensity decay: ?

Refinement

R[*F*² > 2σ(*F*²)] = 0.080
wR(*F*²) = 0.205
S = 1.01
 4457 reflections
 280 parameters
 23 restraints
 H-atom parameters constrained
 Δρ_{max} = 0.93 e Å⁻³
 Δρ_{min} = -0.42 e Å⁻³

Table 1

Hydrogen-bond geometry (Å, °).

<i>D</i> –H... <i>A</i>	<i>D</i> –H	H... <i>A</i>	<i>D</i> ... <i>A</i>	<i>D</i> –H... <i>A</i>
OW2–HW2B...OW3	0.85	2.22	2.787 (5)	125
OW3–HW3B...OW1	0.85	2.14	2.747 (5)	128
OW3–HW3A...N2	1.01	1.84	2.836 (5)	167
OW1–HW1A...OW2 ⁱ	0.85	2.15	2.715 (5)	123
OW1–HW1B...N1 ⁱⁱ	1.06	1.95	2.899 (5)	148
OW2–HW2A...O1 ⁱⁱⁱ	0.85	2.11	2.834 (5)	143

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

The H atoms bonded to C atoms were placed in calculated positions, with C–H = 0.96–0.97 Å, and refined as riding, with *U*_{iso}(H) = 1.2*U*_{eq}(C) or 1.5*U*_{eq}(methyl C). The water H atoms were located in a difference map and refined as riding in their as-found relative positions with *U*_{iso}(H) = 1.2*U*_{eq}(O). Atoms C5, C6 and C7 were restrained so that their *U*^{*ij*} components approximated to isotropic behavior.

Data collection: *CAD-4 Software* (Enraf–Nonius, 1989); cell refinement: *CAD-4 Software*; data reduction: *XCAD4* (Harms & Wocadlo, 1995); program(s) used to solve structure: *SHELXS97* (Sheldrick, 1997a); program(s) used to refine structure: *SHELXL97* (Sheldrick, 1997a); molecular graphics: *SHELXTL* (Sheldrick, 1997b); software used to prepare material for publication: *SHELXL97*.

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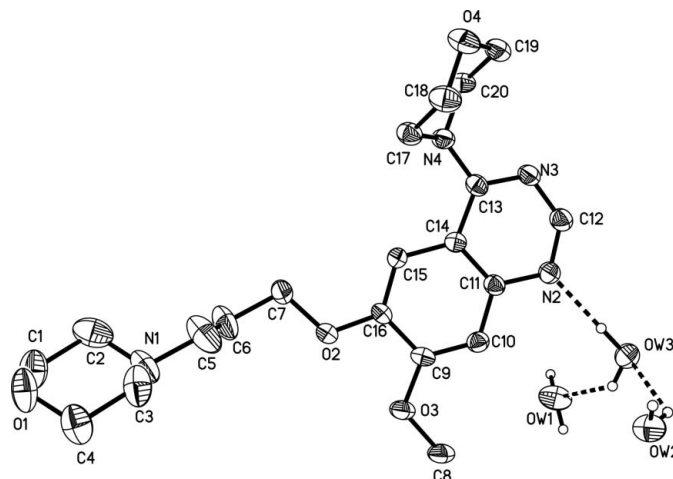


Figure 1

The molecular structure of (I) showing 40% displacement ellipsoids (arbitrary spheres for the water H atoms; other H atoms omitted for clarity). Dashed lines indicate hydrogen bonds.

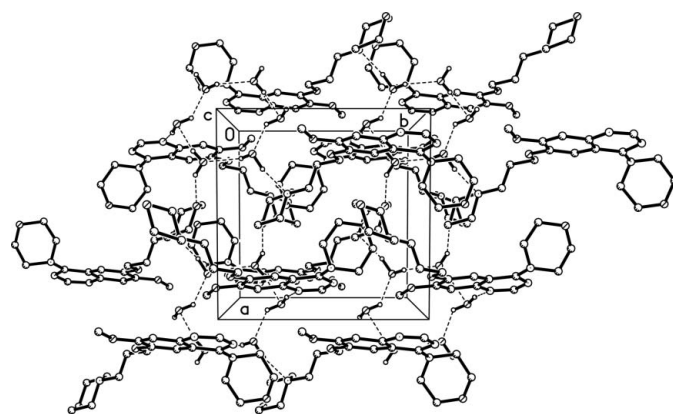


Figure 2

The packing of (I). Hydrogen bonds are shown as dashed lines, and the H atoms not involved in the hydrogen bonds have been omitted for clarity.

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