

Sumaira Umbreen, Sabine Foro*
and Boris SchmidtClemens-Schöpf-Institut für Organische Chemie
und Biochemie, Technische Universität Darm-
stadt, Petersenstraße 22, D-64287 Darmstadt,
Germany

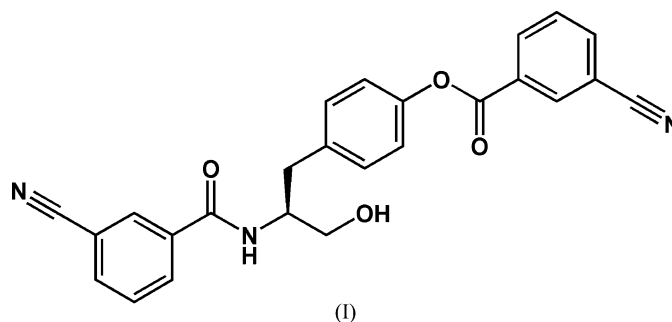
Correspondence e-mail: foro@tu-darmstadt.de

Key indicators

Single-crystal X-ray study
 $T = 299$ K
Mean $\sigma(\text{C}-\text{C}) = 0.005$ Å
 R factor = 0.038
 wR factor = 0.079
Data-to-parameter ratio = 8.4For details of how these key indicators were
automatically derived from the article, see
<http://journals.iucr.org/e>.**(S)-4-[2-(3-Cyanobenzamido)-3-hydroxy-
propyl]phenyl 3-cyanobenzoate**Non-planar molecules of the title compound, $\text{C}_{25}\text{H}_{19}\text{N}_4\text{O}_3$, are
linked by intermolecular $\text{N}-\text{H}\cdots\text{O}$, $\text{O}-\text{H}\cdots\text{N}$ and $\text{C}-$
 $\text{H}\cdots\text{O}$ hydrogen bonds to form a three-dimensional network.Received 17 May 2006
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Comment

The accumulation of β -amyloid peptide ($A\beta$) in the brain is thought to be a primary cause for the progression of Alzheimer's disease (Selkoe, 2001). Since $A\beta$ is generated from the cleavage of β -amyloid precursor protein (APP) by proteolytic enzymes, β - and γ -secretases (Sinha & Lieberburg, 1999), these two secretases represent potential therapeutic targets (Schmidt *et al.*, 2005). The identification of β -secretase (Vassar *et al.*, 1999) prompted us to develop effective inhibitors against this enzyme. β -Secretase belongs to an aspartyl protease family, similar to HIV protease. The majority of potent inhibitors of BACE are still peptide-based transition state analogues according to several reviews (Schmidt, 2003; Schmidt *et al.*, 2005, 2006). Hydroxyethylenes, statines, norstatines, bis-statines, hydroxyethylamines and hydroxyethylureas were employed. The hydroxyethylenes delivered the first highly potent inhibitors. The compound (*S*)-4-[2-(3-cyanobenzoamido)-3-hydroxypropyl]phenyl 3-cyanobenzoate (I), is an important precursor for hydroxyethylene amides. In the reaction of 3-cyanobenzoic acid with *L*-tyrosinol, we obtained compound (I) as the major and unexpected product. X-ray studies of the title compound, (I), have been carried out to obtain detailed structural information and the results are presented here.



The three intermolecular $\text{N}-\text{H}\cdots\text{O}$, $\text{O}-\text{H}\cdots\text{N}$ and $\text{C}-$
 $\text{H}\cdots\text{O}$ hydrogen bonds form a three-dimensional network.
Details of the hydrogen-bonding geometry are given in
Table 1. The dihedral angles $\text{C}8-\text{O}2-\text{C}9-\text{C}14$ and $\text{N}2-$
 $\text{C}18-\text{C}19-\text{C}24$ are 95.5 (3) and -34.2 (4) $^\circ$, respectively
indicating non-planarity in the molecule of (I).

Experimental

Ethyl-3-(3'-dimethylaminopropyl)carbodiimide hydrochloride (979 mg, 5.11 mmol) and *N*-hydroxybenzotriazole hydrate (828 mg, 6.13 mmol) were added to a solution of 3-cyanobenzoic acid (751 mg, 5.11 mmol) dissolved in CH₂Cl₂ (10 ml). The resulting mixture was stirred at ambient temperature for 5 min, then treated with *L*-tyrosinol (1.02 ml, 6.13 mmol) and triethylamine (1.42 ml, 10.22 mmol) for 6 h. CH₂Cl₂ (20 ml) was added, and the solution was washed with HCl (0.1 *N*, 5 × 30 ml), NaHCO₃ saturated solution (3 × 30 ml) and brine (1 × 30 ml), dried over Na₂SO₄ and concentrated to obtain the title compound, (I), as colourless crystals (1.5 g, 69%). Single crystals of (I) suitable for X-ray data collection were obtained by slow evaporation of a methanol/dichloromethane (2:8) solution.

Crystal data

C₂₅H₁₉N₃O₄ Z = 4
M_r = 425.43 *D_x* = 1.312 Mg m⁻³
 Monoclinic, *C*2 Mo *K*α radiation
a = 32.665 (9) Å μ = 0.09 mm⁻¹
b = 4.778 (1) Å *T* = 299 (2) K
c = 15.015 (4) Å Nneedle, colourless
 β = 113.19 (2)° 0.50 × 0.08 × 0.02 mm
V = 2154.1 (9) Å³

Data collection

Oxford Diffraction Xcalibur Diffraction, 2004
 diffractometer with Sapphire *T_{min}* = 0.974, *T_{max}* = 0.998
 CCD detector 7760 measured reflections
ω and φ scans 2438 independent reflections
 Absorption correction: analytical 1261 reflections with *I* > 2σ(*I*)
 (*CrysAlis RED*; Oxford) *R_{int}* = 0.052
 θ_{max} = 26.4°

Refinement

Refinement on *F*² H-atom parameters constrained
R [*F*² > 2σ(*F*²)] = 0.038 *w* = 1/[σ²(*F_o*²) + (0.0294*P*)²]
wR(*F*²) = 0.079 where *P* = (*F_o*² + 2*F_c*²)/3
S = 0.88 (Δ/σ)_{max} < 0.001
 2438 reflections Δρ_{max} = 0.15 e Å⁻³
 289 parameters Δρ_{min} = -0.14 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>
N2—H2N...O4 ⁱ	0.86	2.14	2.889 (4)	145
O3—H3O...N1 ⁱⁱ	0.82	2.19	3.007 (5)	173
C11—H11...O1 ⁱⁱⁱ	0.93	2.53	3.414 (4)	159

Symmetry codes: (i) *x*, *y* - 1, *z*; (ii) *x* + ½, *y* - ½, *z*; (iii) -*x* + ½, *y* + ½, -*z* + 1.

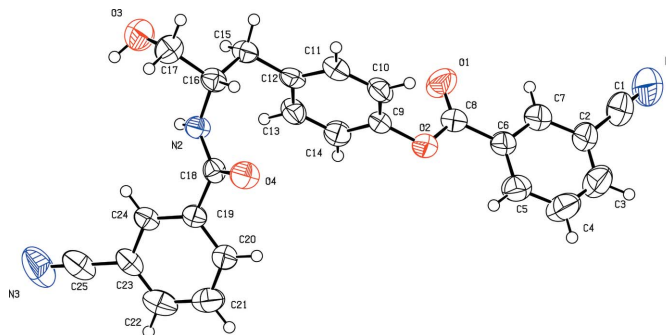


Figure 1 Molecular structure of (I), showing the atom labelling and displacement ellipsoids drawn at the 50% probability level.

H atoms were positioned with idealized geometry and were refined using a riding model, with C—H in the range 0.93–0.98 Å, O—H = 0.82 Å and N—H = 0.86 Å. *U_{iso}*(H) values were set equal to 1.2*U_{eq}* of the parent atom. In the absence of significant anomalous dispersion effects, Friedel pairs were merged and the Δ*f*' term set to zero. The absolute configuration was assigned according to the known absolute configuration of the educt *L*-tyrosinol.

Data collection: *CrysAlis CCD* (Oxford Diffraction, 2003); cell refinement: *CrysAlis RED* (Oxford Diffraction, 2004); data reduction: *CrysAlis RED*; program(s) used to solve structure: *SHELXS97* (Sheldrick, 1997); program(s) used to refine structure: *SHELXL97* (Sheldrick, 1997); molecular graphics: *PLATON* (Spek, 2003); software used to prepare material for publication: *SHELXL97*.

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