



# HCV protease inhibitor | BI-1388

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## Summary

BI-1388 is a highly selective inhibitor of NS3 protease with nanomolar potency across various HCV genotypes and against resistant mutants D168V and R155K.

## Chemical Structure

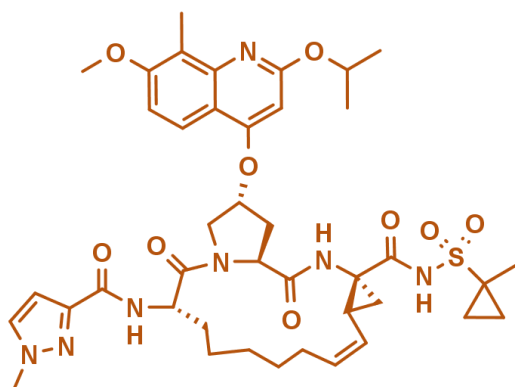


Figure 1: 2-D structure of BI-1388, an inhibitor of HCV NS3 protease

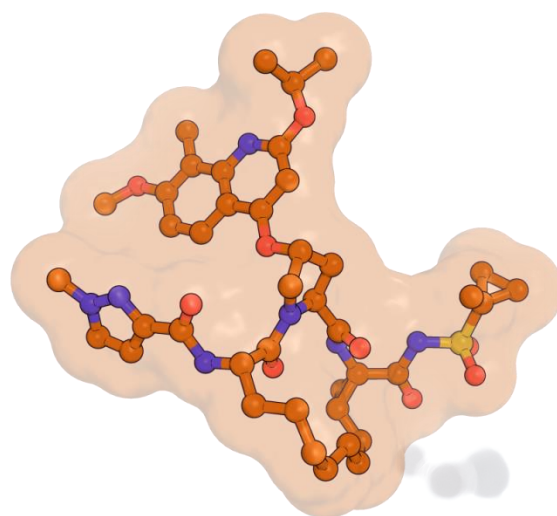


Figure 2: 3-D structure of BI-1388, an inhibitor of HCV NS3 protease

## Highlights

BI-1388 binds to the active site of NS3 that is located in the shallow and broad protein-protein interaction surface of the protease- and the helicase domain of the enzyme. It is a nanomolar to picomolar inhibitor of protease activity and of viral replication for various HCV genotypes and for resistant mutants D168V and R155K. BI-1388 was shown to be highly selective against other serine/cysteine proteases, and it is characterised by an exceptional liver partitioning.

## Target information

HCV NS3 protease is a 180-amino acid chymotrypsin-like serine protease. Its function is the auto-proteolytic cleavage of HCV viral polyprotein (~3000 aa) into individual, non-structural (NS) proteins with various functions. Thus it is an essential component of HCV replication and infectivity. The NS3 protein contains two functional domains: a serine protease- and a helicase domain. The active site of NS3 is located in the shallow and wide protein-protein interaction surface of these domains. BI-1230 and other known NS3 inhibitors cover significant parts of this interaction surface in addition to the active site. Boehringer Ingelheim was the first company to establish proof-of-concept in humans for an HCV NS3 protease inhibitor as a treatment of HCV infection <sup>4</sup>.



**Figure 3: BI-1388 bound to the active site of NS3 (PDB code: 4i31)<sup>3</sup>**

## *In vitro* activity

PROBE NAME	BI-1388
MW [Da]	820
IC <sub>50</sub> [nM] <sup>a</sup> , WT enzyme, genotype 1a	0.48
IC <sub>50</sub> [nM] <sup>a</sup> , WT enzyme, genotype 1b	1.1
IC <sub>50</sub> [nM] <sup>a</sup> , WT enzyme, genotype 2a	0.14
IC <sub>50</sub> [nM] <sup>a</sup> , WT enzyme, genotype 3a	12
IC <sub>50</sub> [nM] <sup>a</sup> , WT enzyme, genotype 4a	0.23
IC <sub>50</sub> [nM] <sup>a</sup> , WT enzyme, genotype 5a	0.24
IC <sub>50</sub> [nM] <sup>a</sup> , WT enzyme, genotype 6a	0.16
IC <sub>50</sub> [nM] <sup>a</sup> , R155K, genotype 1a	4.7
IC <sub>50</sub> [nM] <sup>a</sup> , D168V, genotype 1a	58
IC <sub>50</sub> [nM] <sup>a</sup> , A156T, genotype 1b	2.7
IC <sub>50</sub> [nM] <sup>a</sup> , D156V, genotype 1b	4.3
IC <sub>50</sub> [nM] <sup>a</sup> , D168A, genotype 1b	16
IC <sub>50</sub> [nM] <sup>a</sup> , D168V, genotype 1b	15
EC <sub>50</sub> [nM], WT, replicon assay, genotype 1a <sup>b</sup>	0.11
EC <sub>50</sub> [nM], R155K, replicon assay, genotype 1a <sup>b</sup>	1.0
EC <sub>50</sub> [nM], WT, replicon assay, genotype 1b <sup>b</sup>	0.11
EC <sub>50</sub> [nM], D168V, replicon assay, genotype 1b <sup>b</sup>	0.14

<sup>a</sup> Enzymatic assay, see ref. 3 for assay details

<sup>b</sup> Cell-based HCV RNA replication Luciferase reporter assay, see ref. 3 for assay details

## *In vitro* DMPK and CMC parameters

PROBE NAME / NEGATIVE CONTROL	BI-1388
Aqueous solubility @ pH 7 [ $\mu\text{g}/\text{ml}$ ]	3.4
CACO permeability @ pH 7.4 [ $*10^{-6} \text{ cm}/\text{s}$ ]	10
CACO efflux ratio	1.2
Microsomal stability: $t_{1/2}$ [min], RLM	138
Plasma protein binding human [%]	99.800

## *In vivo* DMPK parameters

Pharmacokinetic profile of BI-1388 in rat <sup>a</sup>

ROUTE		
i.v.	CL [ml/min/kg]	19
	T $_{1/2}$ (h)	2.6
	$V_{ss}$ [l/kg] <sup>b</sup>	0.97
p.o.	F(%)	14
	C <sub>max</sub> ( $\mu\text{M}$ )	1.3
	AUC <sub>0-inf</sub> ( $\mu\text{M}\cdot\text{h}$ )	1.4

<sup>a</sup> Dose = i.v., 4 mg/kg (70% PEG-400 and 30% water); p.o., 10 mg/kg (1% MP, 0.3% Tween-80, and Methocel 0.5%)

## Selectivity

BI-1388 has been shown to be highly selective against other serine/cysteine proteases.<sup>3</sup>

Activity of BI-1388 against other proteases

PROTEASE	IC <sub>50</sub> <sup>A</sup>
Caspase I	>10
Cathepsin B	>30
Cathepsin G	>10
Chymotrypsin	>25
Elastase, leucocyte	>30
MMP-1	>10
Trypsin	>10

<sup>a</sup>All values are in  $\mu\text{M}$ . Value given is the highest concentration tested, which in all cases gave <50% inhibition.

## Co-crystal structure of the BI probe compound and the target protein.

The X-ray structure of BI-1388 in complex with NS3 is available (PDB code: 4i31, 4i32, 4i33 – complexes with WT and resistant mutants D168V and R155K)

## Reference molecule(s)

For a recent review of HCV NS3 protease inhibitors see reference 6.

## Summary

BI-1388 is a nanomolar to picomolar inhibitor of HCV NS3 protease activity and of viral replication for various HCV genotypes and for resistant mutants D168V and R155K. BI-1388 was shown to be highly selective against other serine/cysteine proteases.

## References

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