



HCV protease inhibitor | BI-1230

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Summary

BI-1230 is a nanomolar inhibitor of HCV protease and of viral replication with good *in vivo* PK characteristics.

Chemical Structure

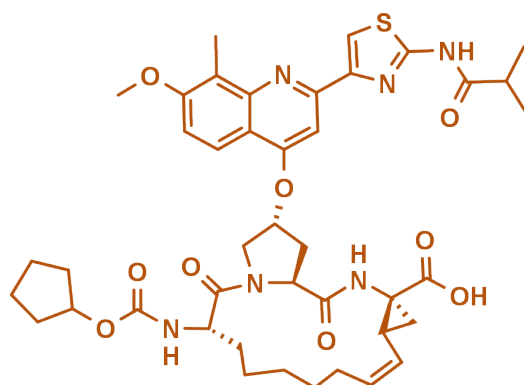


Figure 1: 2-D structure of BI-1230, an inhibitor of HCV NS3 protease

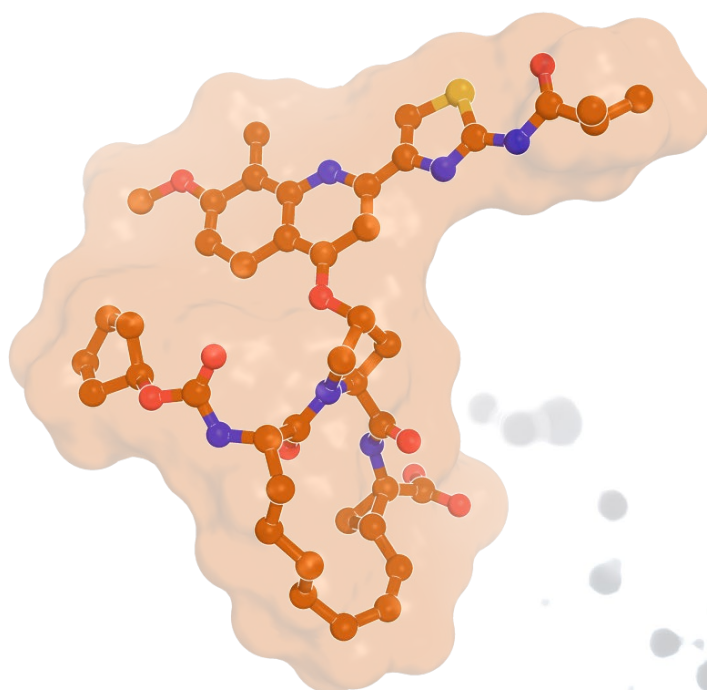


Figure 2: 3-D structure of BI-1230, an inhibitor of HCV NS3 protease

Highlights

BI-1230 is a single-digit nanomolar inhibitor of HCV NS3 protease activity and of viral replication. It binds to the active site of NS3 and was shown to be highly selective against other serine/cysteine proteases. BI-1230 shows good *in vivo* PK properties, including half-life and bioavailability. As such, this compound is a valuable tool for both *in vitro* and *in vivo* studies. We also offer [faldaprevir](#) from our infectious diseases pipeline for pre-clinical research purposes.

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.⁴

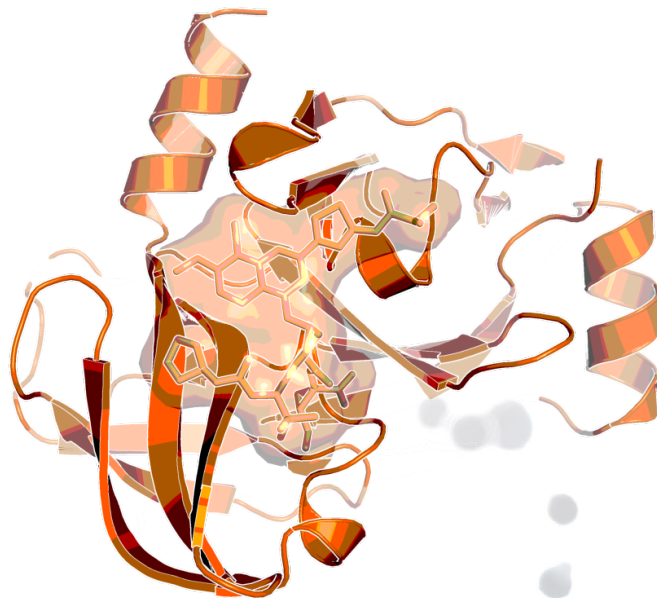


Figure 3: X-ray structure of HCV NS3 protease with faldaprevir, a close analog of BI-1230 (PDB code: 3p8n)

In vitro activity

PROBE NAME / NEGATIVE CONTROL	BI-1230	BI-1675
MW [Da]	817	554.6
IC ₅₀ [nM] ^a	6.7	4870
EC ₅₀ [nM], replicon assay, genotype 1a ^b	4.6	n.d.
EC ₅₀ [nM], replicon assay, genotype 1b ^b	<1.8	n.d.

^a Enzymatic assay, NS3-NS4A heterodimer, fluorogenic substrate, 60 min incubation

^b Cell-based HCV RNA replication Luciferase reporter assay, genotype background 1a and 1b, Huh7 cells, 72 h incubation

In vitro DMPK and CMC parameters

PROBE NAME / NEGATIVE CONTROL	BI-1230	BI-1675
Aqueous solubility @ pH 7 [µg/ml]	20	n.d.
Caco permeability @ pH 7.4 [$*10^{-6}$ cm/s]	8.7	n.d.
Caco efflux ratio	0.3	n.d.
Microsomal stability [% Q _H]	<24	n.d.
Plasma protein binding human [%]	99.9	n.d.

In vivo DMPK parameters

In vivo DMPK parameters of BI-1230 in rat and dog.^a

ROUTE		RAT	DOG
<i>i.v.</i>	CL [ml/min/kg]	15	1.9
	Mean residence time after iv dose (h)	2.3	3.4
	V _{ss} [l/kg]	2.05	0.39
<i>p.o.</i>	T _{1/2} (h)	2.1	5.1
	t _{max} (h)	1.8	1.7
	C _{max} (nM)	405	7370
	AUC _{0-inf} (nM*h)	2550	49700
	F [%]	42	92

^a Dose = *i.v.*, 2 mg/kg; *p.o.*, 5 mg/kg

Negative control

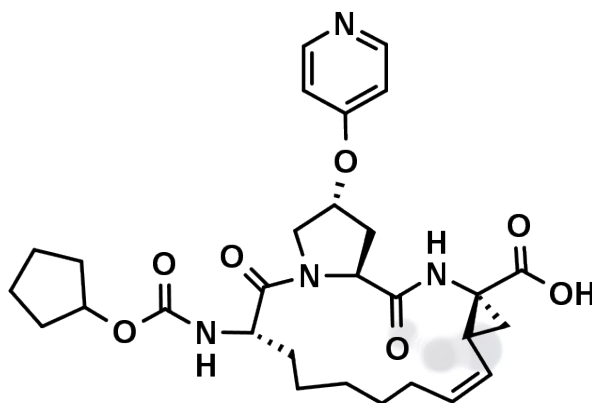


Figure 4: BI-1675, negative control

Selectivity

BI-1230 is highly selective against other serine/cysteine proteases.

SELECTIVITY DATA AVAILABLE	BI-1230	BI-1675
SafetyScreen44™ with kind support of 	Yes	Yes
Invitrogen®	No	No
DiscoverX®	No	No
Dundee	No	No

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

An X-ray structure of BI-1230 in complex with NS3 is not available. However, the structure with the highly related [faldaprevir](#) (BI 201335) was solved.

Reference molecule(s)

For a recent review of HCV NS3 protease inhibitors see reference 5.

Supplementary data

2D structures can be downloaded free of charge from [openMe](#).

References

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