

KRAS switch I/II pocket inhibitor

BI-2852

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Summary

The *in vitro* tool compound BI-2852 is a potent nanomolar inhibitor of the KRAS switch I/II pocket and directly inhibits both the active and inactive forms of KRAS.

Chemical Structure

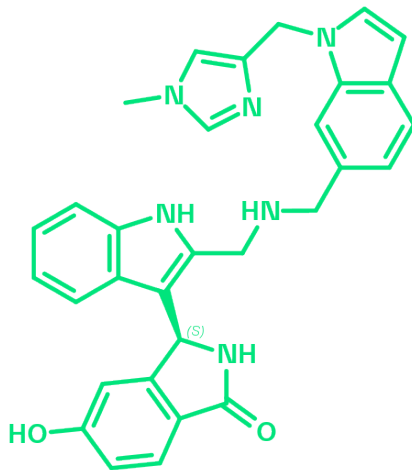


Figure 1: 2D structure of BI-2852

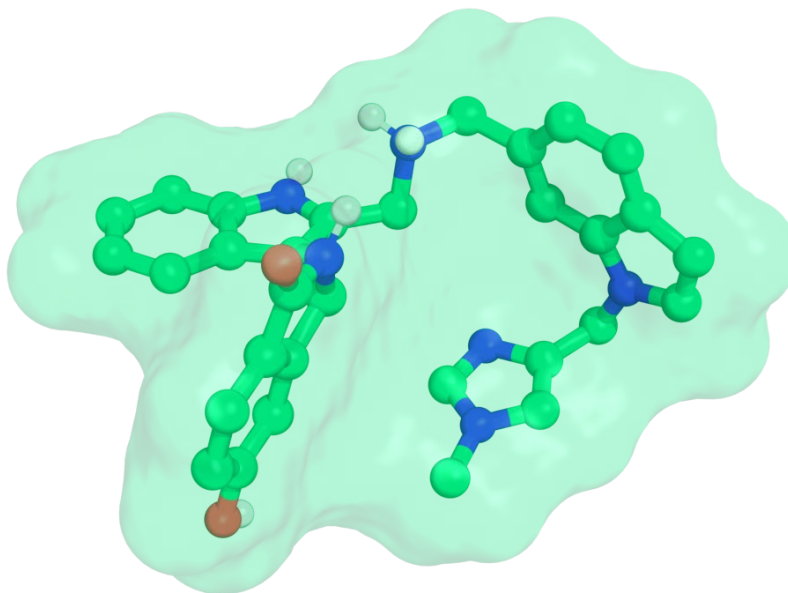


Figure 2: BI-2852, 3D conformation

Highlights

BI-2852 is a potent nanomolar inhibitor of the KRAS switch I/II pocket. It directly targets and inhibits active GTP-bound KRAS. This compound is suitable for *in vitro* experiments and may be an excellent tool for testing KRAS-related biological hypotheses. pERK modulation and antiproliferative effects were observed in KRAS^{mut} cells (NCI-H358) treated with BI-2852^{5,6}.

Target information

The three human *RAS* genes, *KRAS*, *NRAS* and *HRAS* encode four different RAS proteins (KRAS-4A, KRAS-4B, NRAS and HRAS) which belong to the protein family of small GTPases. The RAS proteins function as molecular switches between active GTP-bound and inactive GDP-bound conformations. RAS is the most frequently mutated oncogene in human cancers (~27%) with activating mutations mainly in codons 12, 13 and 61. The main mutation in codon 12 causes RAS activation by interfering with GAP binding and GAP-stimulated GTP hydrolysis. *KRAS* mutations rates are high in pancreatic (~90%), colorectal (~45%) and lung adenocarcinomas (~35%)^{1,2}. *KRAS* could serve as an excellent drug target for many cancers, but direct inhibition of oncogenic RAS has proven to be challenging¹⁻⁶.

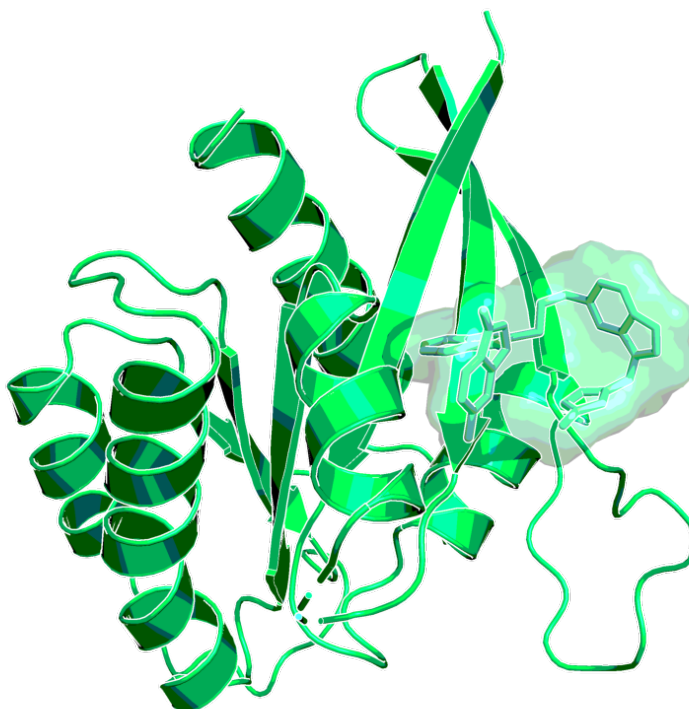


Figure 3: Complex of KRAS with BI-2852

In vitro activity

BI-2852 binds to KRAS^{G12D} with a K_D of 740 nM (ITC) and inhibits GTP-KRAS^{G12D} binding to effectors like SOS1, CRAF and PI3K α with an IC₅₀ of 490, 770 and 500 nM. BI-2852 displays an IC₅₀ of 490 nM in a GTP-KRAS^{G12D}::SOS1 AlphaScreen (AS) assay leading to low micromolar inhibition (EC₅₀ of 5.8 μ M) of pERK (in H358 cell line)⁵.

PROBE NAME / NEGATIVE CONTROL	BI-2852	BI-2853
MW [Da, free base] ^a	516.6	516.6
ITC (K _D) GCP-KRAS ^{G12D} [μ M] ^b	0.74	n.a.
ITC (K _D) GCP-KRAS ^{wt} [μ M] ^b	7.5	n.a.
ITC (K _D) GDP-KRAS ^{G12D} [μ M] ^b	2.0	n.a.
ITC (K _D) GDP-KRAS ^{wt} [μ M] ^b	1.1	n.a.
AS (IC ₅₀) GTP-KRAS ^{G12D} ::SOS1 [nM] ^b	490	4400
AS (IC ₅₀) GTP-KRAS ^{G12D} ::CRAF [nM] ^b	770	n.a.
AS (IC ₅₀) GTP-KRAS ^{G12D} ::PI3K α [nM] ^b	500	n.a.
AS (IC ₅₀) GDP-KRAS ^{G12D} ::SOS1 [nM] ^b	260	2500
AS (IC ₅₀) GTP-KRAS ^{wt} ::SOS1 [nM] ^b	490	n.a.
EC ₅₀ pERK H358 cells (2 h) [μ M] ^b	5.8	>50
EC ₅₀ H358 cells (low serum) [μ M] ^b	6.7	n.a.

^a For the salt form you will get, please refer to the label on the vial and for the molecular weight of the salt, please refer to the FAQs

^b For assay conditions please refer to reference 5

In vitro DMPK and CMC parameters

PROBE NAME / NEGATIVE CONTROL	BI-2852	BI-2853
logP @ pH 11	2.6	2.6
Solubility @ pH 6.8 [μ g/mL]	18	21
Caco-2 permeability AB @pH7.4 [$\times 10^{-6}$ cm/s]	5.0	<1.15
Caco-2 efflux ratio	2.1	n.a.

Microsomal stability (human/mouse/rat) [% Q _H]	91 / 93 / 90	92 / 95 / 86
Hepatocyte stability (human/mouse/rat) [% Q _H]	12 / 21 / 25	12 / 69 / 52
Plasma Protein Binding (human/mouse/rat) [%]	98.8 / 99.5 / 98.5	98.7 / 99.1 / 98.6
CYP 3A4 (IC ₅₀) [μM]	4.4	n.a.
CYP 2C8 (IC ₅₀) [μM]	8.4	n.a.
CYP 2C9 (IC ₅₀) [μM]	4.8	n.a.
CYP 2C19 (IC ₅₀) [μM]	11.0	n.a.
CYP 2D6 (IC ₅₀) [μM]	15.0	n.a.

In vivo DMPK parameters

No data available, BI-2852 is an *in vitro* tool compound.

In vivo pharmacology

No data available, BI-2852 is an *in vitro* tool compound.

Negative control

BI-2853 is the less active enantiomer of BI-2853. It shows no effect on cells and is around 10-fold less potent in the AS assays.

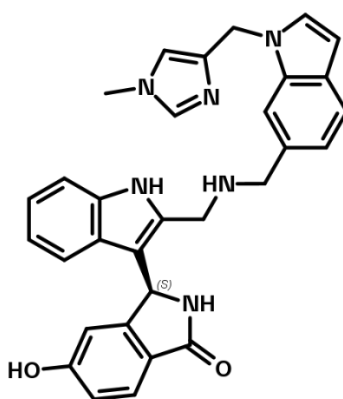


Figure 4: BI-2853 which serves as a negative control

Selectivity

SELECTIVITY DATA AVAILABLE	BI-2852	BI-2853
SafetyScreen44™ with kind support of  eurofins	Yes	Yes
Invitrogen®	No	No
DiscoverX®	No	No
Dundee	No	No

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

The X-ray crystal structure of target in complex with BI-2852 is available (PDB code: 6GJ8)^{5,6}.

Supplementary data

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

References

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