

Supplementary Materials

Novel Small-Molecule Inhibitors of the SARS-CoV-2 Spike Protein Binding to Neuropilin 1

Anja Kolarič ¹, Marko Jukič ^{1,2,*} and Urban Bren ^{1,2,*}

¹ Faculty of Chemistry and Chemical Engineering, University of Maribor, Smetanova 17, SI-2000 Maribor, Slovenia

² Faculty of Mathematics, Natural Sciences and Information Technologies, University of Primorska, Glagoljaška 8, SI-6000 Koper, Slovenia

* Correspondence: marko.jukic@um.si; Tel: +386 2 2294 428 (M.J.) and urban.bren@um.si; Tel: + 386 2 22 94 421 (U.B.)

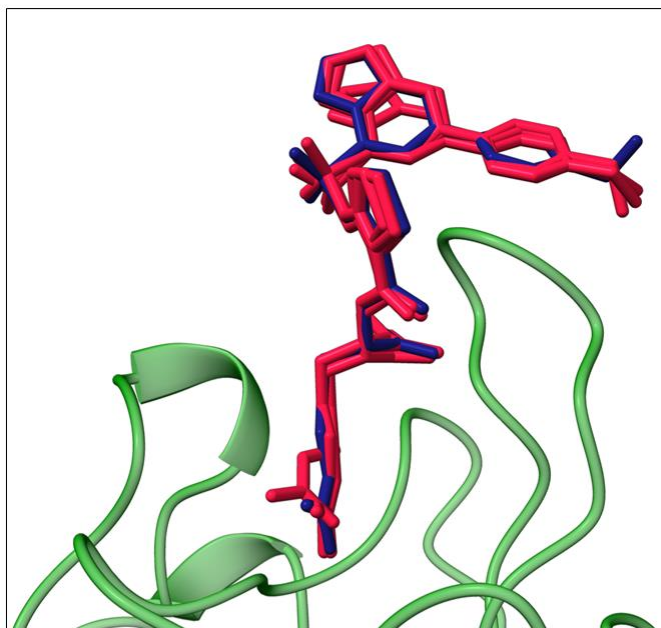


Figure S1. Spatial comparison of the natively present EG01377 conformation and its calculated docked poses for the molecular docking validation. The co-crystallized conformation is represented in blue, while its re-produced docked poses are depicted in red. NRP1 is colored in green.

GOLD configuration file

AUTOMATIC SETTINGS

autoscale = 2

POPULATION

popsiz = auto

select_pressure = auto

n_islands = auto

maxops = auto

niche_siz = auto

GENETIC OPERATORS

pt_crosswt = auto

allele_mutatewt = auto

migratewt = auto

FLOOD FILL

radius = 10

origin = 0 0 0

do_cavity = 1

floodfill_atom_no = 0

floodfill_center = cavity_from_ligand 17
atoms

DATA FILES

param_file = DEFAULT

set_ligand_atom_types = 1

set_protein_atom_types = 0

tordist_file = DEFAULT

make_subdirs = 0

save_lone_pairs = 0

fit_points_file = fit_pts.mol2

read_fitpts = 0

FLAGS

internal_ligand_h_bonds = 0

flip_free_corners = 0

match_ring_templates = 0

flip_amide_bonds = 0

flip_planar_n = 1 flip_ring_NRR

flip_ring_NHR

flip_pyramidal_n = 0

rotate_carboxylic_oh = flip

use_tordist = 1

postprocess_bonds = 1

rotatable_bond_override_file = DEFAULT

solvate_all = 1

TERMINATION

early_termination = 1

n_top_solutions = 3

rms_tolerance = 1

CONSTRAINTS

force_constraints = 0

COVALENT BONDING

covalent = 0

clean_up_option delete_all_log_files

output_file_format = MACCS

SAVE OPTIONS

save_score_in_file = 1

save_protein_torsions = 0

clean_up_option save_top_n_solutions 3

clean_up_option

delete_redundant_log_files

clean_up_option

delete_all_initialised_ligands

clean_up_option delete_empty_directories

clean_up_option delete_rank_file

FITNESS FUNCTION SETTINGS

initial_virtual_pt_match_max = 3

relative_ligand_energy = 1

gold_fitfunc_path = plp

score_param_file = DEFAULT

CONSTRAINTS

constraint protein_h_bond 10.0000

0.005000 1215 1305 1167 411 754 755

Table S1. ZINC15 [1] name and MolPort ID (vendor) of top 20 compounds obtained from molecular docking.

COMPOUND	ZINC15 NAME	MolPort ID
1	ZINC000390834636	MolPort-039-314-631
2	ZINC000065529168	MolPort-019-672-384
3	ZINC000070040863	MolPort-020-084-791
4	ZINC000024451999	MolPort-009-171-237
5	ZINC000020732290	MolPort-006-814-244
6	ZINC000057997476	MolPort-009-073-388
7	ZINC000015721546	MolPort-002-516-537
8	ZINC000012428013	MolPort-002-749-663
9	ZINC000003295631	MolPort-004-259-418
10	ZINC000095995534	MolPort-028-795-075
11	ZINC000058026820	MolPort-009-102-148
12	ZINC000048326459	MolPort-009-073-421
13	ZINC000008584635	MolPort-007-627-531
14	ZINC000003271207	MolPort-004-257-684
15	ZINC000003319242	MolPort-004-262-387
16	ZINC000008708441	MolPort-005-566-437
17	ZINC000012769255	MolPort-005-749-385
18	ZINC000426477237	MolPort-042-631-013
19	ZINC000036360945	MolPort-008-317-429
20	ZINC000009422900	MolPort-005-913-364
21 (EG00229) [2]		MolPort-046-416-868
22 [3]		MolPort-003-000-781

REFERENCES

1. ZINC. Available online: <https://zinc15.docking.org/> (accessed Oct 6, 2021)
2. Jarvis, A.; Allerston, C. K.; Jia, H.; Herzog, B.; Garza-Garcia, A.; Winfield, N.; Ellard, K.; Aqil, R.; Lynch, R.; Chapman, C.; Hartzoulakis, B.; Nally, J.; Stewart, M.; Cheng, L.; Menon, M.; Tickner, M.; Djordjevic, S.; Driscoll, P. C.; Zachary, I.; Selwood, D. L. Small molecule inhibitors of the neuropilin-1 vascular endothelial growth factor A (VEGF-A) interaction. *J. Med. Chem.* 2010, 53, 2215-2226. DOI: 10.1021/jm901755g.
3. Perez-Miller, S.; Patek, M.; Moutal, A.; Duran, P.; Cabel, C. R.; Thorne, C. A.; Campos, S. K.; Khanna, R. Novel compounds targeting neuropilin receptor 1 with potential to interfere with SARS-CoV-2 virus entry. *ACS Chem. Neurosci.* 2021, 12, 1299-1312. DOI: 10.1021/acscchemneuro.0c00619.