

Supplementary Material

Targeting the essential transcription factor HP1043 of *Helicobacter pylori*: a drug repositioning study

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1 Supplementary Tables

1.1 Table S1 Plasmids used in this study

Plasmid	Description	Source/Reference
pTrc::1043	Derivative of pTrcHisA expressing the HP1043 response regulator; Amp ^r	(Delany et al., 2002)
pGEMt-P1227 WT	pGEMt-Easy derivative, containing a 61 bp probe corresponding to the region from 1.290.664 to 1.290.725 of <i>H. pylori</i> G27 genome. The region corresponds to the promoter of <i>HPG27_RS06145</i> gene (<i>hp1227</i> according to <i>H. pylori</i> 26695 strain annotation)	(Pelliciari et al., 2017)

1.2 Table S2 Oligonucleotides used in this study

Primer	Sequence	Source/Reference
Php1227_EMSA_F	GCCAAAACGCCTAAAGCC	(Zannoni et al., 2021)
Php1227_EMSA_R	TTGAAAGCGCAATAACCGC	(Zannoni et al., 2021)
16SRTF	GGAGTACGGTCGCAAGATTAAA	(Pelliciari et al., 2017)
16SRTR	CTAGCGGATTCTCTCAATGTCAA	(Pelliciari et al., 2017)

1.3 Table S3 Docking comparison to literature

Docking box have dimension limits, so to include the whole structure, two docking boxes were designed, HP1043_site1 and Hp1043_site 2. Boxes overlap at dimer interface.

		HP1043	-DNA		HP1043 site1		HP1043 site2			
		binding	#		binding	#		binding	#	
		energy	cluster	Ki	energy	cluster	Ki	energy	cluster	Ki
				31.28			899.53			29.83
nifedipine	ZINC85205448	-6.15	25	uM	-8.25	9	nM	-6.17	14	uM
				277.87			419.39			12.41
nicardipine -	ZINC19796087	-8.94	10	nM	-8.70	13	nM	-6.69	13	uM
				707.68			6.29			1.90
	ZINC84400879	-8.39	10	nM	-7.10	2	uM	-7.81	12	uM
				88.97			10.36			9.70
nicoldinino	ZINC591993	-5.53	13	uM	-6.80	6	uM	-6.84	22	uM
insolutplife				497.04			5.33			4.67
	ZINC19632706	-8.60	4	nM	-7.19	12	uM	-7.27	21	uM
				55.67			22.10			6.41
nimodinino	ZINC19632713	-5.80	16	uM	-6.35	28	uM	-7.08	21	uM
mmoupme				205.41			53.39			1.26
	ZINC19632718	-5.03	11	uM	-5.83	11	uM	-8.05	8	uM
				63.63			4.03			16.44
nitrandinina	ZINC84403367	-5.72	24	uM	-7.36	26	uM	-6.53	39	uM
intrendipine				2.29			2.32			2.58
ZI	ZINC100001908	-7.70	10	uM	-7.69	93	uM	-7.62	46	uM
				1.27			4.15			1.85
lercondining	ZINC19685788	-8.04	7	uM	-7.34	4	uM	-7.82	1	uM
lereandiplite				41.87			243.13			333.09
	ZINC19685790	-10.07	1	nM	-9.02	1	nM	-8.84	1	nM
chrysin				353.26			416.11			201.10
cinysin	ZINC3872070	-8.80	83	nM	-8.70	34	nM	-9.14	9	nM
anigenin				192.69			762.31			960.09
apigeiiii	ZINC3871576	-9.16	65	nM	-8.35	12	nM	-8.21	11	nM
lutaolin				397.61			39.33			331.86
lucom	ZINC18185774	-8.73	88	nM	-10.10	5	nM	-8.84	17	nM
hesperetin				260.60			2.19			2.86
(yso2)	ZINC39092	-8.98	92	nM	-7.72	72	uM	-7.56	89	uM
Iro ama fanal				627.90			1.72			2.28
Kaempieroi	ZINC3869768	-8.46	16	nM	-7.86	10	uM	-7.70	47	uM
quercitin				371.41			1.17			121.28
(meletin)	ZINC3869685	-8.77	34	nM	-8.09	10	uM	-9.44	9	nM
muricotin				85.18			3.71			1.06
myriceun	ZINC3874317	-9.64	42	nM	-7.41	24	uM	-8.15	46	uM

1.4 Table S4 docking results

ligand	binding energy (kcal/mol)	run	Ki	
ZINC000022443609 HP1043-DNA	-23.84	38	3.37 aM	
ZINC000013492579_HP1043-DNA	-16.79	117	494.39 fM	Not for sale
ZINC000003938681_HP1043-DNA	-16.41	66	937.24 fM	DNA
ZINC000019363537_HP1043-DNA	-15.57	81	3.86 pM	
ZINC000001566899_HP1043-DNA	-15.41	98	5.06 pM	
ZINC000004214772_HP1043-DNA	-15.31	147	5.96 pM	DNA
ZINC000100088802_HP1043-DNA	-14.99	156	10.26 pM	DNA
ZINC000150338708 HP1043-DNA	-14.39	11	28.4 pM	Expensive
ZINC000013492553_HP1043-DNA	-14.3	190	33.17 pM	Not for sale
ZINC000028823338_HP1043-DNA	-14.14	124	43.44 pM	Not for sale
ZINC000004097448_HP1043-DNA	-14.05	43	50.02 pM	DNA
ZINC000004097404_HP1043-DNA	-13.99	79	55.55 pM	DNA
ZINC000222508879_HP1043-DNA	-13.99	75	55.58 pM	Not for sale
ZINC000028823530_HP1043-DNA	-13.93	60	61.88 pM	Not for sale
ZINC000013494070_HP1043-DNA	-13.92	42	63.08 pM	Not for sale
ZINC000019364225_HP1043-DNA	-13.84	70	71.87 pM	
ZINC000013494068_HP1043-DNA	-13.73	46	86.02 pM	Not for sale
ZINC000043206370_HP1043-DNA	-13.56	176	115.23 pM	DNA
ZINC000001612996_HP1043-DNA	-13.32	199	171.12 pM	DNA
ZINC000013494069_HP1043-DNA	-13.1	10	250.98 pM	Not for sale
ZINC000036701290_HP1043-DNA	-13.08	103	256.35 pM	
ZINC000028823491_HP1043-DNA	-12.92	139	339.8 pM	Not for sale
ZINC000087666889_HP1043-DNA	-12.92	22	335.64 pM	Not for sale
ZINC000098023177_HP1043-DNA	-12.89	92	356.1 pM	
ZINC000150598297_HP1043-DNA	-12.82	121	401.34 pM	Not for sale
ZINC000004097383_HP1043-DNA	-12.81	40	406.52 pM	
ZINC000028134101_HP1043-DNA	-12.79	153	421.91 pM	Not for sale
ZINC000028823512_HP1043-DNA	-12.79	70	420.99 pM	Not for sale
ZINC000087666882_HP1043-DNA	-12.79	140	420.91 pM	Not for sale
ZINC000072316335_HP1043-DNA	-12.7	128	487.81 pM	
ZINC000022448983_HP1043-DNA	-12.66	167	526.94 pM	
ZINC000028823490_HP1043-DNA	-12.56	47	616.87 pM	Not for sale
ZINC000028823510_HP1043-DNA	-12.53	9	652.73 pM	Not for sale
ZINC000003978083_HP1043-DNA	-12.46	143	737.86 pM	
ZINC000030690433_HP1043-DNA	-12.46	46	740.65 pM	Expansive

ZINC000052955754_HP1043-DNA	-12.46	31	736.06 pM	
ZINC000012503187_HP1043-DNA	-12.42	134	791.48 pM	
ZINC000014880002_HP1043-DNA	-12.39	15	827.22 pM	
ZINC000087666886_HP1043-DNA	-12.38	137	837.75 pM	Not for sale
ZINC000222508821_HP1043-DNA	-12.32	63	936.79 pM	Not for sale
ZINC000028823531_HP1043-DNA	-12.29	149	985.83 pM	Not for sale
ZINC000022443609_HP1043	-17.65	35	116.01 fM	
ZINC000019363537_HP1043	-13.68	21	94.13 pM	
ZINC000019363537_HP1043	-12.86	20	373.37 рМ	
ZINC000003934128_HP1043	-12.23	146	1.09 nM	
ZINC000003934128_HP1043	-11.18	102	6.41 nM	
ZINC000029059664_HP1043	-10.82	134	11.73 nM	Not for sale
ZINC000028824061 HP1043	-10.67	124	15.01 nM	Not for sale
ZINC000072126305 HP1043	-10.2	118	33.25 nM	Not for sale
ZINC000072126306_HP1043	-10.18	137	34.59 nM	Not for sale
ZINC000042804032_HP1043	-10.04	102	43.55 nM	Not for sale
ZINC00000004724_HP1043	-9.94	103	51.39 nM	
ZINC000044352037_HP1043	-9.91	178	54.83 nM	Not for sale
ZINC000029060547_HP1043	-9.7	127	76.97 nM	Not for sale
			1 0 1 5 1	

2 Reference

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- Zannoni, A., Pelliciari, S., Musiani, F., Chiappori, F., Roncarati, D., and Scarlato, V. (2021). Definition of the Binding Architecture to a Target Promoter of HP1043, the Essential Master Regulator of Helicobacter pylori. *Int. J. Mol. Sci. Artic.* doi:10.3390/ijms22157848.

3 Supplementary Figures



RMSD HP1043_DNA



Supplementary Figure S1. C α RMSD. On the top HP1043 bound to DNA complexes, bottom HP1043 free complexes.



RMSF HP1043_DNA

Supplementary Figure S2. RMSF per-residue. On the top HP1043 bound to DNA complexes, bottom HP1043 free complexes.

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Supplementary Figure S3. EMSAs in the presence of putative ligands. All EMSAs show the same amount of specific (P_{hp1227}) and non-specific (16S rRNA gene) DNA probes and the same samples order: lane a) DNA probes control, protein- and ligand-free; b) compound control at 1 mM (indicated by a "+") without the protein; c) DNA binding control in the presence of HP1043; lane d) to h) show samples containing a fixed amount of HP1043 monomeric protein (4 μ M) with a decreasing concentration of the ligand, respectively 1, 0.5, 0.2, 0.1, 0.05 mM; the absence of protein and compound is indicated by a "-"; the compound concentration is depicted as a grey triangle, while a white rectangle is used for the HP1043 protein fixed concentration. **A**) Oxcarbazepine did not cause a significant loss of the mobility shift if compared with the protein control (lane c). As well as **B**) Ribocilcib and **C**) Plerixafor, whose presence in the reaction mix did not affect the DNA binding. **D**) Dihydroergotamine interacted with both DNA probes at higher concentrations causing a shift in a protein-free reaction (lane b). Therefore, a putative inhibition was not assessable.