

Targeting West Nile virus replicases: NS3 and heterodimers inhibitors

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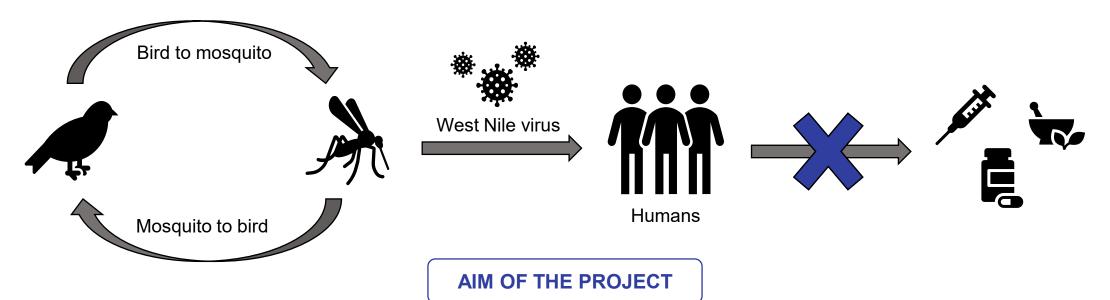


INTRODUCTION and AIM OF THE PROJECT



The **West Nile virus** (**WNV**) is a flavivirus of the Flaviviridae family. It circulates through a mosquito–bird–mosquito cycle and it is transmitted to humans through Culex mosquitoes.

Urgency for the **development of new WNV antiviral compounds**.



A direct approach targeting the active site of NS2B-NS3 protease.

Peptides/peptidomimetics as potential inhibitors.

An **indirect approach** targeting protein-protein interactions essential for NS3 activity.

Peptides to disfavour NS2B-NS3 heterodimer.



NS2B-NS3 ACTIVE SITE INHIBITORS



Tri and tetra-peptide inhibitors

Several peptides and peptidomimetics were synthesized as possible NS2B-NS3 inhibitors of WNV, tested and their IC_{50} value was evaluated.

Peptide name	Interaction	MW (g/mol)	Purity	IC ₅₀ (μΜ)
Nona-D-Arginine	NON – COVALENT	1422.73	87%	20.70 ± 3.60
PhAc-Lys-Lys-Arg-NH ₂		547.71	97%	130.4 ± 8.3
Aun-Lys-Lys-Arg-NH ₂		612.86	96%	34.43 ± 3.65
Palm-Lys-Lys-Arg-NH ₂		667.99	95%	29.43 ± 3.81
PhAc-Lys-Lys-Arg-(cyclic-dehydro)		514.67	96%	8.33 ± 0.33
Aun-Lys-Lys-Arg-(cyclic-dehydro)		579.83	98%	8.34 ± 0.78
Palm-Lys-Lys-Arg-(cyclic-dehydro)		634.95	97%	10.34 ± 1.71
PhAc-Lys-Lys-Arg-H	COVALENT	532.69	95%	3.14 ± 0.19
Aun-Lys-Lys-Arg-H		597.85	97%	7.05 ± 0.97



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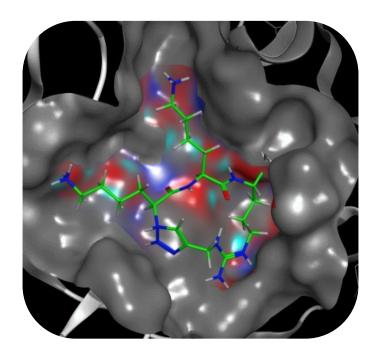
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Cyclic peptide inhibitor

Cyclization of the tripeptides to increase the inhibitory efficacy and to disfavor their degradation.



Superimposition of cyclic tripeptide on the active site pocket of NS2B-NS3 protease of WNV.



NS2B-NS3 ACTIVE SITE INHIBITORS



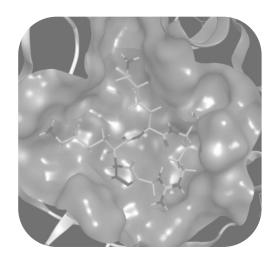
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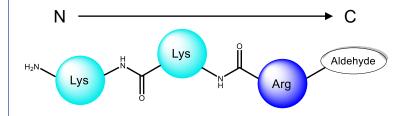
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Cyclization of the tripeptide to increase the inhibitory efficacy and to disfavor its degradation.

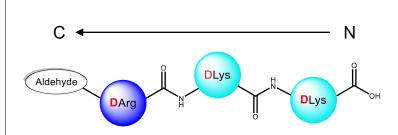


Retro-inverse peptide inhibitors

To increase the inhibitory efficacy and to disfavor its degradation.



Reference peptide



Retro-inverso peptide

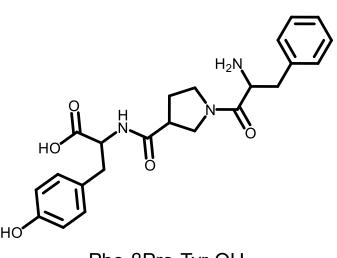


NS2B-NS3 ALLOSTERIC SITE INHIBITORS



We designed **different tripeptides** that mimic the structure of an already known reversible allosteric inhibitor (SID:852843) of the formation of the functional conformation of the NS3 protease of WNV and we studied these **possible allosteric inhibitors by docking**.

Active tripeptide allosteric inhibitor



Phe-βPro-Tyr-OH



$$IC_{50}$$
 = **48.30** ± 11.07 µM

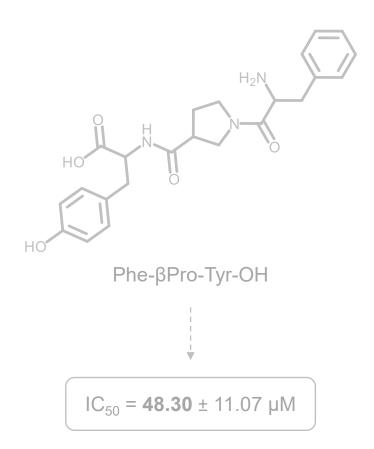


NS2B-NS3 ALLOSTERIC SITE INHIBITORS



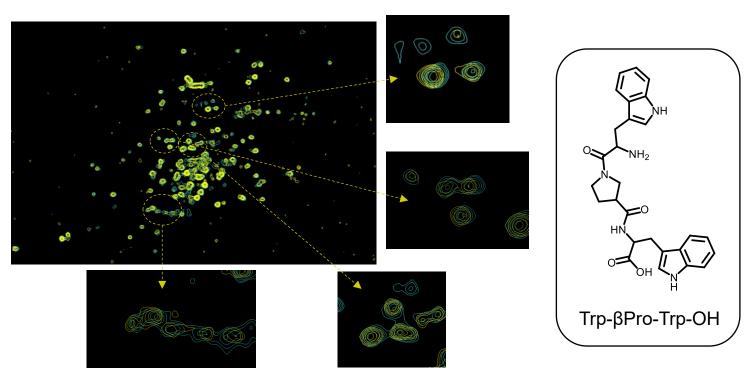
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Active tripeptide allosteric inhibitor



Protein-peptides interactions by NMR studies

2D NMR BEST-TROSY spectrum of the Trp-βPro-Trp peptide, which interacts with the NS2B-NS3 protease, as evidenced by Chemical Shift Perturbations.





AKNOWLEDGMENTS



Prof. **Cristina Peggion**: supervisor

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Prof. Mattia Sturlese

Caterina Zulian Thomas Stella



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University of Cagliari - Department of life and environmental sciences

Prof. Angela Corona Prof. Enzo Tramontano



INF-ACT

