





Integrating computational and biophysical approaches within VITALITY enabling platform for Drug Discovery

Dr. Gian Marco Elisi

Campus Enrico Mattei Università degli Studi di Urbino 23 maggio 2024







Projects resulting in accessible pipelines for external partners

- Integration of biophysical experiments and binding free-energy calculations (GSK-3β-D₃ dual-acting compounds)
- Workflows for database screening integration in virtual screening (GSK-3β-D₃ dual-acting compounds)
- Fast shape screening approaches prioritizing ligand structure and conformation (GSK-3β-D₃ dual-acting compounds and *CysH* project)
- Virtual screening **prioritization through deep learning** (GSK-3β-D₃ dual-acting compounds)
- Screening of covalent compounds (CysH project)
- Evaluation of ligand residence time and interaction stability (CysH project)



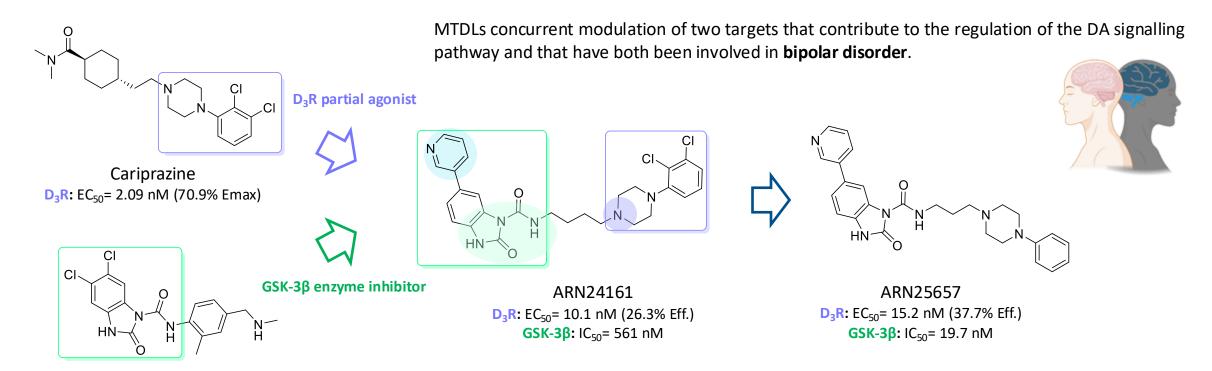
Establishing rapid workflows for DMTA cycles in academic and industrial collaborations







Design of dual-acting compounds targeting D3 receptor and GSK-3β enzymatic activity



- **GSK-3** β : IC₅₀= 8 nM
- Di Martino R. M. C. et al., ChemMedChem. **2020**, 15, 949-54

- Piperazine nitrogen is responsible for the salt-bridge formation with D110^{3.32}
- Benzo[d]imidazole-1-carboxamide portion as GSK-3β hinge binder (D133 e V135)
- 3-pyridil substituent increasing solubility and establishing a contact with K185

D133



K85





Integrating free-energy simulations and biophysical experiments



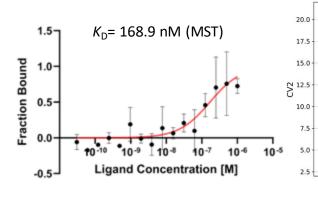




Design and synthesis of macrocyclic derivatives



Interrogation of public databases



MST binding assays

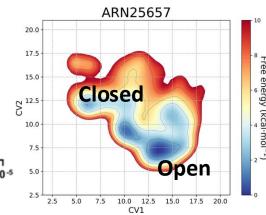


7.5 10.0 12.5 15.0 17.5 20.0 CV1

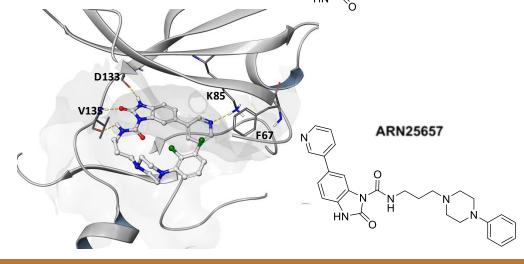
Open

ARN24161

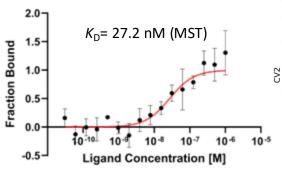
Closed



Design and synthesis of spirocyclic derivatives



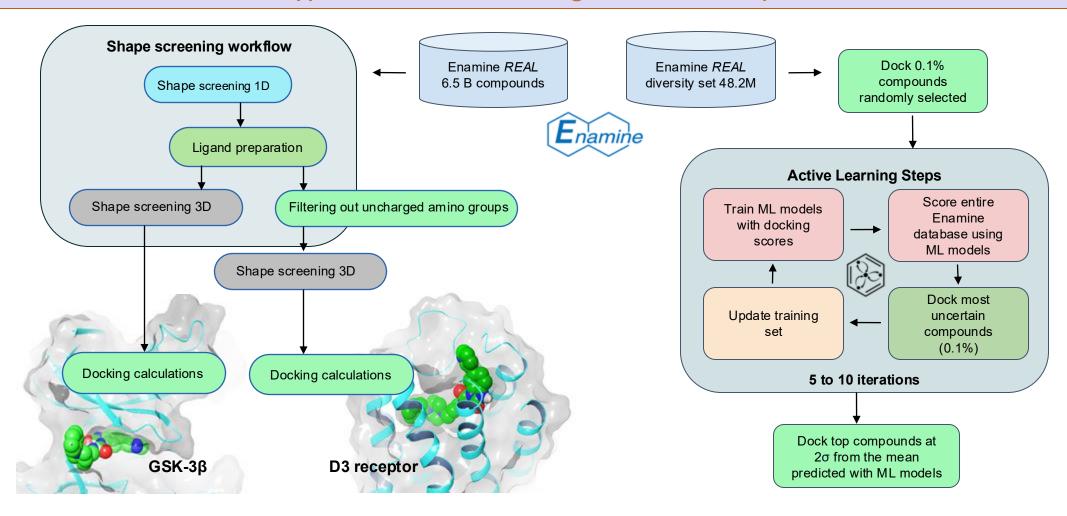
ARN24161







Approaches for virtual screening of commercial compounds



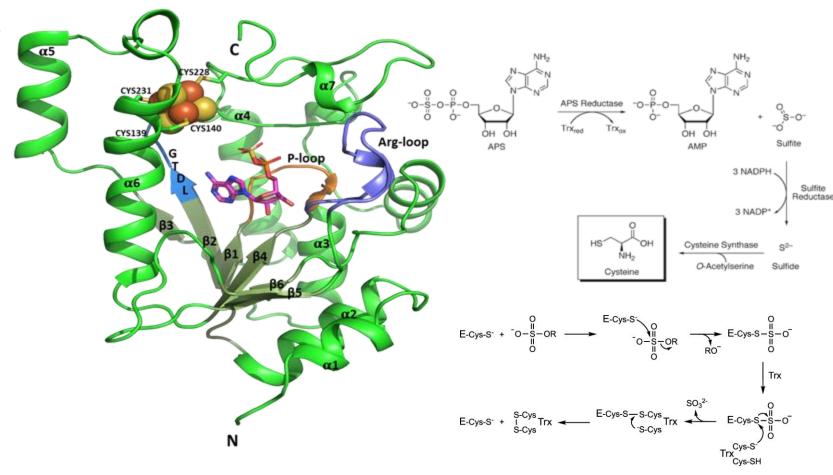






Combating Pseudomonas aeruginosa resistant infections

- Reduction of inorganic sulfate due ASPR activity is necessary for cysteine biosynthesis.
- Disruption of the CysH gene encoding for this enzyme has been correlated to decreased pathogen virulence and cysteine auxotrophy in mycobacteria.
- Mechanism is mediated by C256 located in the non-crystallized C-terminal portion.



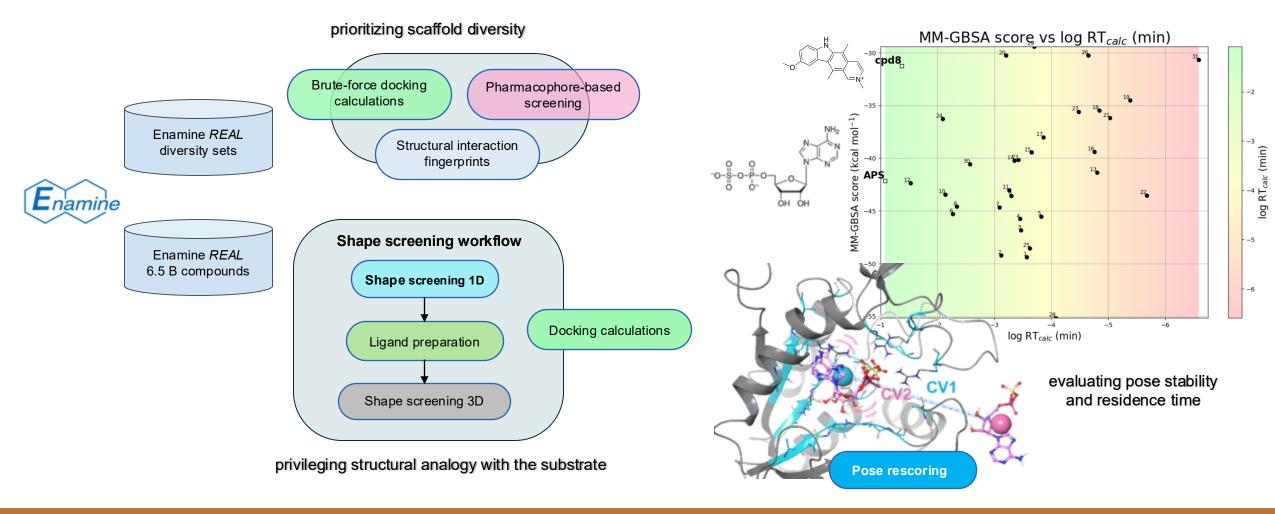
Chartron J. et al., J. Mol Biol. **2006**, 364, 152-69







Finding APSR enzymatic activity inhibitors through concurrent strategies





of covalent probes as chemical

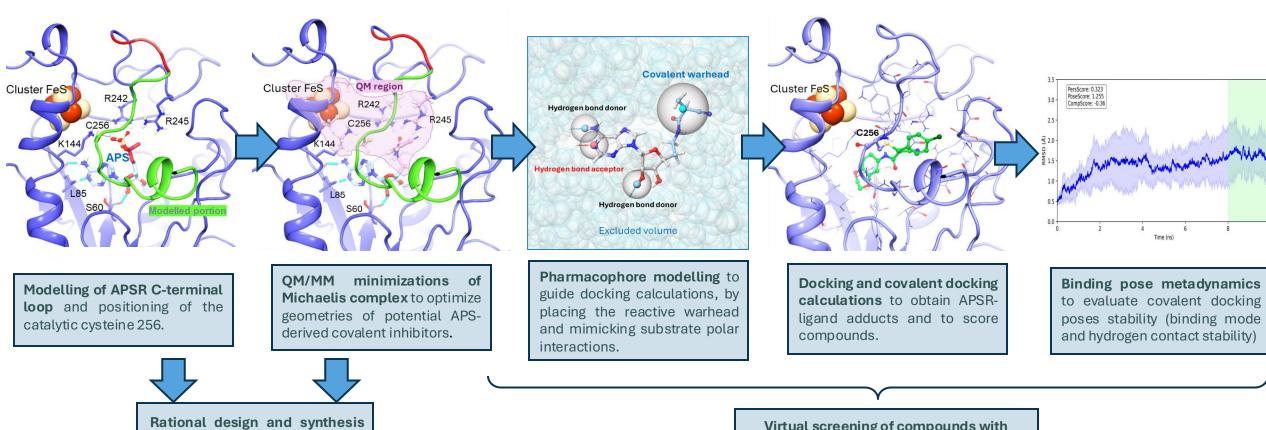
biology tools to retain APSR in a

closed conformation.





A computational workflow to identify covalent inhibitors of APS reductase enzymatic activity



Virtual screening of compounds with acrylamide, chloroacetamide, cyanoacrylamide and vinylsulfone warhead





Thanks for the attention