

Fragment screening as a tool for the search of human dihydroorotate dehydrogenase inhibitors

Nonato, M. C.^{1,2}, Purificação, A. D.^{1,2}, Godoi, B. F.^{1,2}, Santos, T.^{1,2}, Emery, F. S.^{1,2},

¹School of Pharmaceutical Sciences, University of São Paulo and ²Center for Research Advancement in Fragments and Targets

Avenida do café S/N Ribeirão Preto, S.P. Brazil 14040-903

cristy@fcfrp.usp.br

Keywords: dihydroorotate dehydrogenase, fragment screening, COVID-19

Our work aims at contributing to the development of an innovative therapy against COVID-19 based on the selective inhibition of the human enzyme dihydroorotate dehydrogenase (HsDHODH). HsDHODH takes part of the *de novo* pyrimidine biosynthetic pathway [1]. The fact that the virus relies on the host cell biochemical machinery to provide nucleosides for its viral replication cycle DHODH a potential target for the development of broad spectrum antivirals. Our objectives involve fragment synthesis, evaluation and characterization of DHODH fragment-based inhibitors as a strategy for the development of innovative treatment for viral diseases such as COVID-19. Inhibition assays against HsDHODH were carried out using 2,6-dichloroindophenol (DCIP) as the final electron acceptor. Co-crystallization assays were performed using vapour diffusion methods. Inhibitors were evaluated regarding their cytotoxicity and antiviral properties in Calu-3 cells. At 10 μ M, 8 non-cytotoxic nanomolar inhibitors were found to inhibit 100% of viral replication.

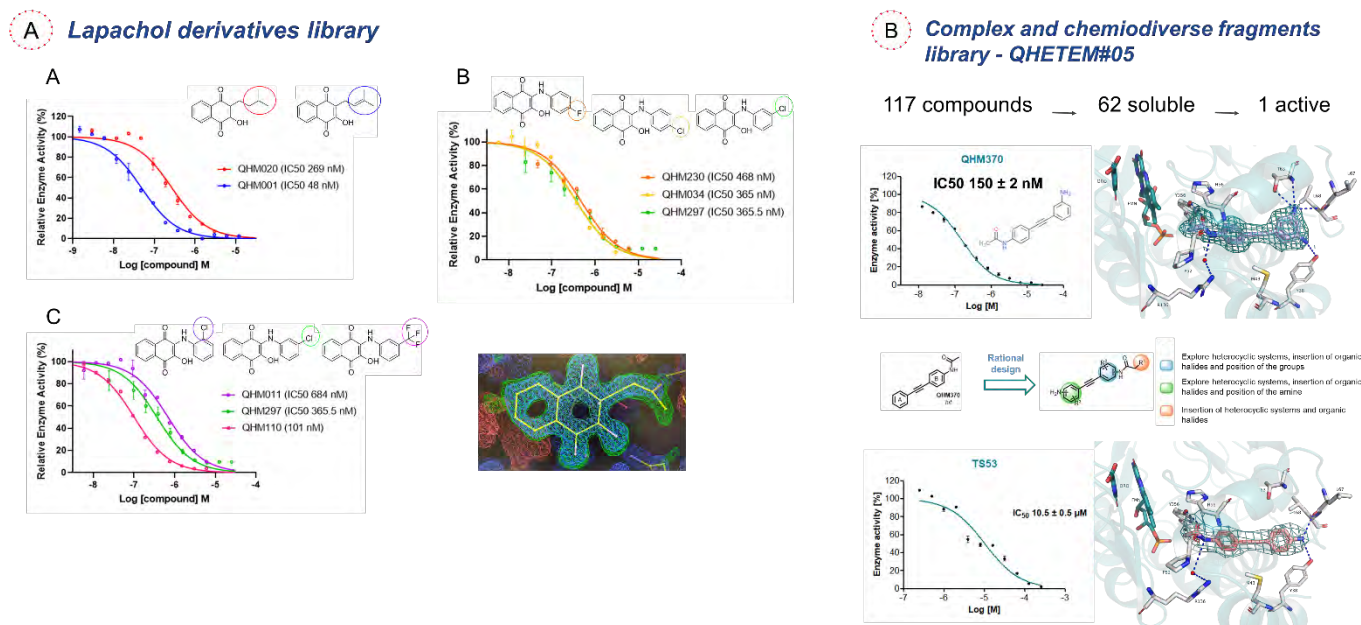


Figure 1. Screening and characterization of human DHODH inhibitors. Different libraries have been tested and potent inhibitors have been identified and characterized by inhibition assays and X-ray crystallography.

The search for new inhibitors for the HsDHODH enzyme is an important strategy in the fight against SARS-CoV-2, responsible for the COVID-19 pandemic with devastating consequences. In this scenario, the fragment screening (FS) is a great tool that can be exploited. In our work, we approached FS by screening different libraries and mapping the molecular basis of ligand interaction by X-ray crystallography. Our results show promising hits tools can be used for the discovery of new inhibitors for the HsDHODH.

[1] Reis RAG, Calil FA, Feliciano PR, Pinheiro MP, Nonato MC. (2017) Arch Biochem Biophys. **632**:175-191.

Acknowledgements: We would like to thank Fundação de Amparo à Pesquisa do Estado de São Paulo (FAPESP), Coordenação de Aperfeiçoamento de Pessoal de Nível Superior Conselho Nacional (CAPES) and Instruct-ERIC for funding. We also thank Manacá beamline at Sirius- Brazil and HZB Fragment Screening Facility for data collection.