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Publication date:
2017

Document license:
[Unspecified](#)

Citation for published version (APA):

Tung, T. T., Dao, T. T., Palmgren, M. B., Fuglsang, A. T., Christensen, S. B., & Nielsen, J. (2017). *LEGO-inspired drug design: Discovery of novel fungal Plasma membrane H⁺-ATPase (Pma1) inhibitors from small molecule libraries: An introduction of HFSA-SBS_DOS-RD strategy in drug discovery.*. Abstract from 253rd American Chemical Society National Meeting & Exposition, .



MEDI: Division of Medicinal Chemistry

374 - LEGO[®]-inspired drug design: Discovery of novel fungal Plasma membrane H⁺-ATPase (Pma1) inhibitors from small molecule libraries: An introduction of HFSA-SBS_DOS-RD strategy in drug discovery

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Abstract: Fungal plasma membrane H⁺-ATPase (Pma1) has recently emerged as a potential target for the discovery of new antifungal agents. This p-type pump which localized on the surface of fungal cells plays a crucial role in many physiological functions and processes inside the cell. Especially, by pumping proton to extracellular, this enzyme generates a transmembrane electrochemical gradient, as a consequence, fungi can uptake nutrients by secondary transport systems. Until now, only low resolution of protein structure has been reported, and notably there is no report of co-crystal structure of Pma1 with inhibitors. Therefore, we have identified the need for small molecule library of high quality for targeting Pma1. The LEGO[®]-inspired hypothesis encouraged us to first develop new strategy from the combination of hypothesis-based fragment selection and assembly (HFSA), specific biological relevance scaffold based diversity-oriented synthesis (SBS_DOS) and rational design (RD), so called HFSA-SBS_DOS-RD strategy in drug discovery and development process. Using HFSA-SBS_DOS-RD, our group successfully designed, synthesized, and performed SAR studies of novel compounds potent Pma1 inhibitors. An expeditious, high yield and scalable microwave-assisted synthesis was developed and applied for synthesis of library compounds. To our delight, our compound libraries were able to inhibit Pma1 activity and growth inhibitory activity of *C. albicans* and *S. cerevisiae* revealed the most promising example for future development of antifungal drugs on this target.