

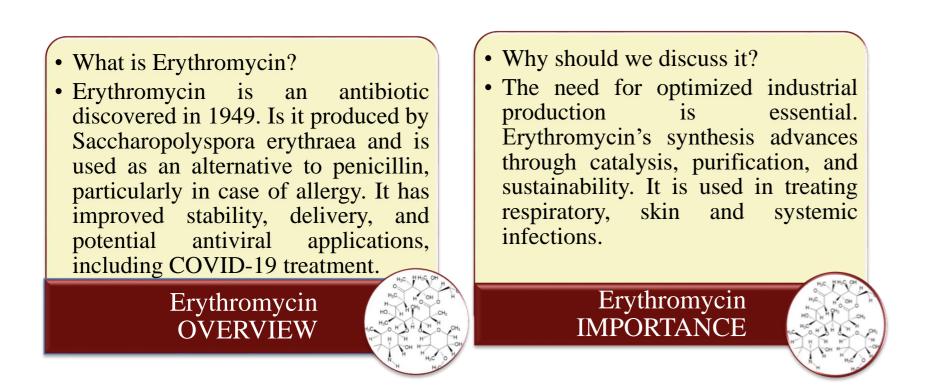
# **The 4th International Electronic Conference on Antibiotics**

21-23 May 2025 | Online

## Industrial Catalytic Production Process of Erythromycin

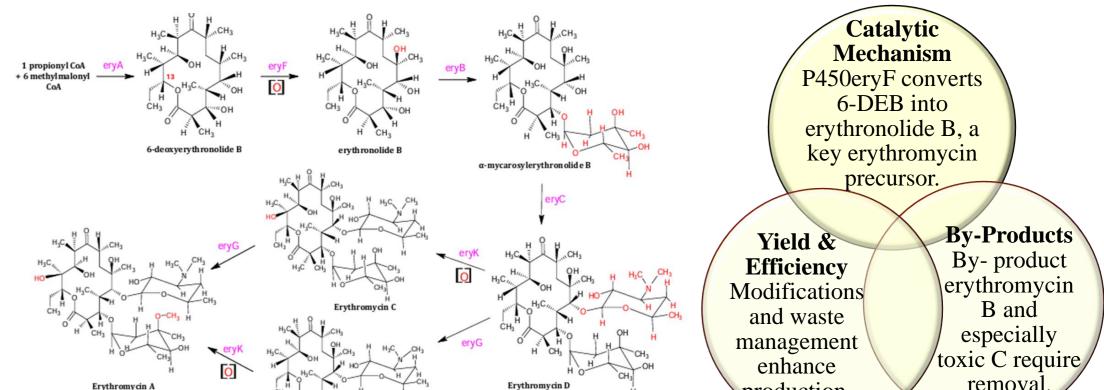
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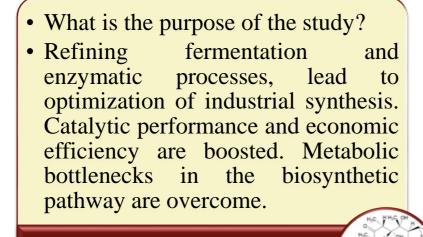
## **INTRODUCTION & AIM**



## **RESULTS & DISCUSSION**

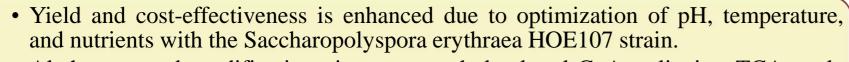
MDP





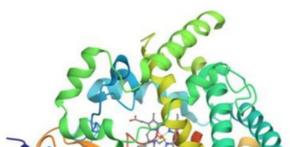
Erythromycin AIM

#### **METHOD**



- Al the targeted modifications increase methylmalonyl-CoA, relieving TCA cycle bottlenecks and all in all enhancing efficiency.
- Structural Analysis: Cytochrome's P450eryF's 3D structure-haem orientation and substrate pocket geometry essential for 6S-hudroxylation of 6 DEB are revealed through analytical techniques (Figures 1,2).
- Enzyme Optimization: Targeted modifications enhance catalytic turnover and specificity, improving production efficiency.
- Isolation erythromycin A is achieved by advanced crystallization methods (antisolvent, evaporative, reactive), reducing contaminants.
- This boosts purity and reduces processing costs. Purification

Sustainability is ensured through hydrolysis of fermentation residues that reduces chemical oxygen demand.

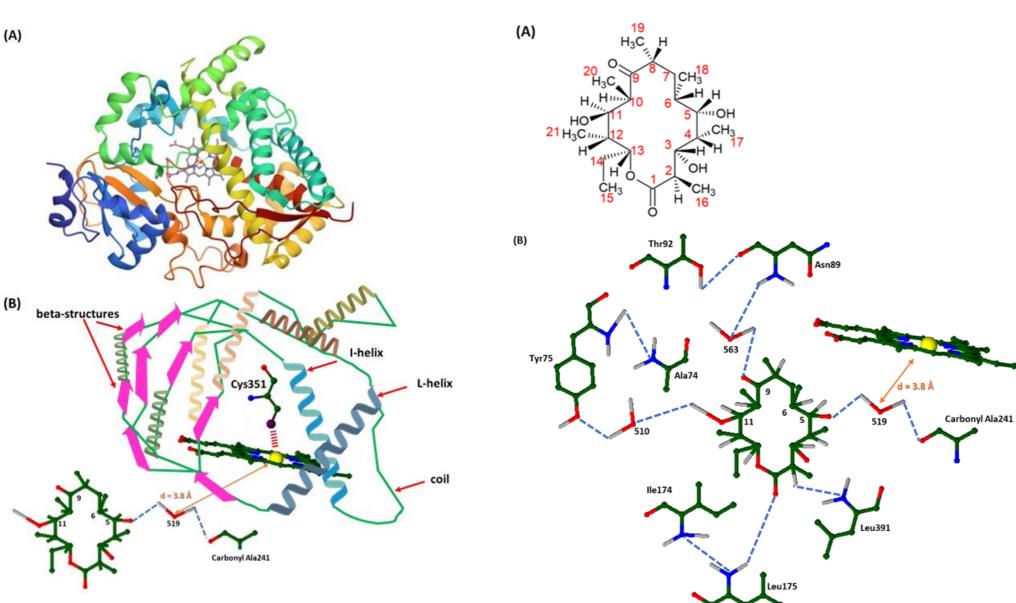


Production

Catalytic

System

Waste



production. removal. Figure 2: Catalytic process of erythromycin A via two different means (the oxidation reaction is displayed using [O]). The catalysts Erythromycin A is separated Challenges from the toxic by-products. of all reactions are depicted in pink, and all transformations are Three crystallization in marked in red. Purification methods-antisolvent, evaporative, and reactive. Genetic tailoring in Genetic Saccharopolyspora Engineering erythraea can deactivate Strategies genes that reduce erythromycin production.

#### Figure 3: Flow chart of erythromycin production.

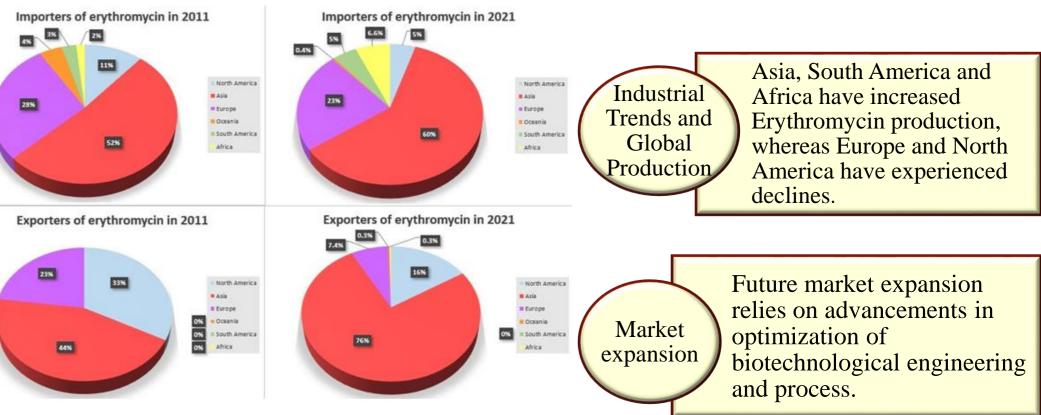


Figure 4: Importers and exporters of erythromycin throughout the years 2011–2021

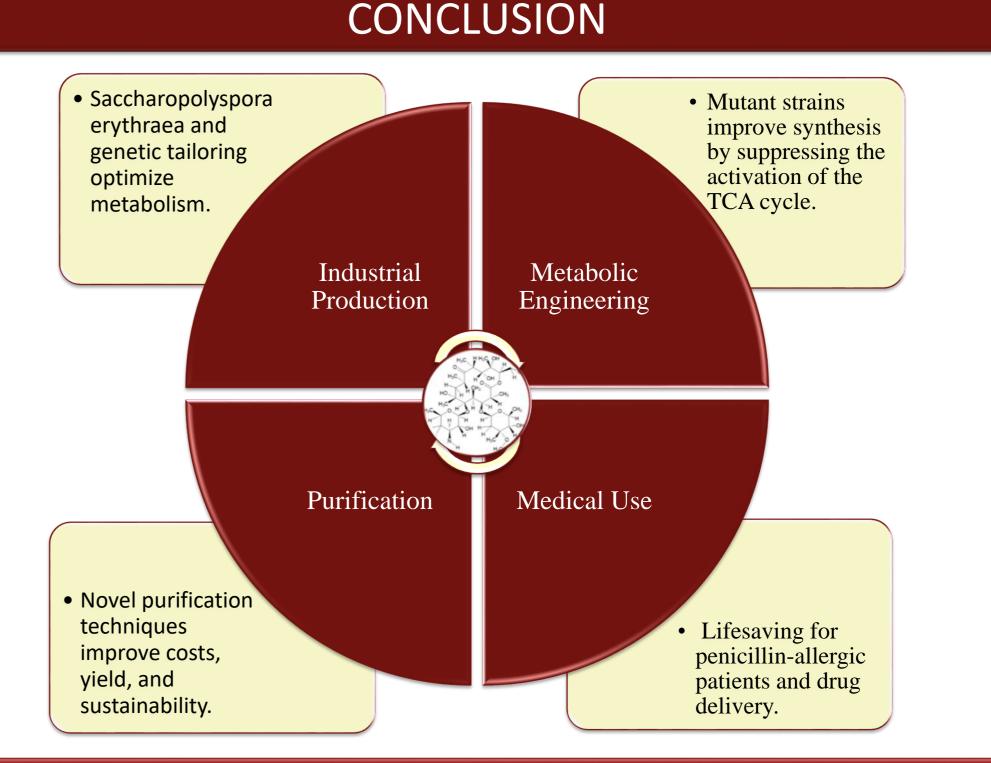


Figure 1: (A) Cytochrome P450eryF general 3D structure (downloaded by the RCSB PDB (Protein Data Bank)). (B) The alpha/beta structure of cytochrome P450eryF in detail. In figure (A), alpha helices are represented by ribbons of different colors on the right side, while beta-structures with coils on the left are depicted with orange arrows.

Figure 2: P450eryF substrate, 6-deoxyerythronolide, and numbering system. (B) 6-DEB in the active site. Red = oxygen, blue = nitrogen, green = carbon, yellow = iron (II). Cyan dashed lines show hydrogen bonds. Water molecules Wat510, Wat519, and Wat563 play roles, with Wat519 possibly being the proton donor in O2 cleavage (3.8Å from iron).

#### FUTURE WORK / REFERENCES Improve the efficiency of industrial production. Guengerich, F.P. Cytochrome P450 Enzymes. Am. Sci. 1993, 81, 440–447. Reduce by-products and waste. OMURA, T. Recollection of the Early Years of the Research on Cytochrome P450. Proc. Jpn. Acad. Ser. B

Expand medical applications

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