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Biosynthesis, molecular modeling and statistical optimization of xylanase from a mangrove associated actinobacterium *Streptomyces variabilis* (MAB3) using Box-Behnken design with its bioconversion efficacy

Muthusamy Sanjivkumar, Tamilselvan Silambarasan, Ramasamy Balagurunathan, Grasian Immanuel



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## Title page

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**Muthusamy Sanjivkumar<sup>1</sup>, Tamilselvan Silambarasan<sup>2</sup>, Ramasamy Balagurunathan<sup>2</sup> and Grasian Immanuel<sup>1\*</sup>**

1. MNP laboratory, Centre for Marine Science and Technology, M.S. University, Rajakamangalam - 629502, India.

2. Department of Microbiology, Periyar University, Salem – 636011, India.

\*Corresponding author: E-mail: gimmas@gmail.com

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