



INDIAN INSTITUTE OF TECHNOLOGY GUWAHATI
SHORT ABSTRACT OF THESIS

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Programme of Study : Ph.D.

Thesis Title: Cloning, expression, purification, biochemical, structure characterization and application of α -L-arabinofuranosidase (*PsGH43_12*) of family 43 glycoside hydrolase from *Pseudopedobacter saltans*

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SHORT ABSTRACT

Hemicellulose is the second most abundant polysaccharide present on earth. The presence of arabinose side chains in hemicellulose such as xylan interact with other polysaccharides and forms complex network resulting in the recalcitrant nature. α -L-Arabinofuranosidases hydrolyses the arabinose side chain and having immense industrial importance. *PsGH43_12* is the first α -L-arabinofuranosidase from *Pseudopedobacter saltans* belonging to family 43 glycoside hydrolase cloned and expressed. It showed maximum activity at pH 6.5 and 50°C with V_{max} 100.7 U/mg, K_m 2.17 mM, and k_{cat}/K_m 50.27 s⁻¹ mM⁻¹ against pNP- α -L-arabinofuranoside. *PsGH43_12* release α -L-araf substitution at O3 and both O2 and O3 positions and therefore, it was classified as type III α -L-arabinofuranosidase. The 3D structure of *PsGH43_12* developed by comparative modeling was compact and stable displaying a 5-bladed β -propeller fold at N-terminal and β -sandwich fold at C-terminal. Molecular docking analysis confirmed the involvement of active site residues, Asp71, Asp180 and Glu247 in the catalysis, which was also confirmed by the site-directed mutagenesis of these residues. SAXS analysis displayed that *PsGH43_12* is monomeric and a fully folded state in solution form. Guinier analysis gave the radius of gyration between (R_g) 2.8 ± 0.09 nm at 5 mg/ml protein concentration. *PsGH43_12* show synergistic behaviour with xylanase. Soaking in aqueous ammonia pretreated sugarcane bagasse on sequential enzymatic saccharification by *CtXyn11A*, *PsGH43_12* and *BoGH43* gave final TRS yield of 164.7 mg/g of ptd SB, with 73.5% hemicellulose conversion and 69.6% xylan to xylose conversion.