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Comparative metabolic profiling of halotolerant bacterial strains and identification of novel species-specific metabolites

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The salt stress response of four representative halotolerant bacterial species (*Halomonas hydrothermalis* VITP9, *Bacillus aquimaris* VITP4, *Planococcus maritimus* VITP21 and *Virgibacillus dokdonensis* VITP14) isolated from a previously unexplored solar saltern in Kumta, along the Arabian Sea coast in Karnataka, India was analyzed using comparative metabolomics approach. Chemometric analysis of ¹H NMR spectra revealed salt-dependent increase in the levels of metabolites, mainly from the aspartate and glutamate family, that are directed from the glycolytic pathway, pentose phosphate pathway and citric acid cycle. The composition of the metabolites was found to be different with respect to the species and the type of growth medium. Two dimensional NMR data revealed accumulation of two rare diaminoacids, N ϵ -acetyl- α -lysine and N δ -acetylornithine apart from other well known compatible solutes. Metabolite profiles of species capable of synthesizing N ϵ -acetyl- α -lysine and N δ -acetylornithine suggested their biosynthesis from lysine and ornithine using aspartate and glutamate as their precursors, respectively. One of the species, *Planococcus maritimus* VITP21 was found to accumulate an unusual sugar, (2-acetamido-2-deoxy- α -D-glucopyranosyl)-(1 \rightarrow 2)- β -D-fructofuranose is not previously reported for its natural synthesis by any other organism. The protective effects of N ϵ -acetyl- α -lysine and (2-acetamido-2-deoxy- α -D-glucopyranosyl)-(1 \rightarrow 2)- β -D-fructo furanose along with other commonly occurring bacterial osmolytes, ectoine, proline, sucrose, trehalose and glycine betaine on protein stability and activity were evaluated with a few of possible biotechnological application.

Biography

Jayaraman G obtained his PhD degree at the National Tsing Hua University (Taiwan) in Structural Biology in 1998 and continued as a Post-doctoral Fellow with funding from National Health Research Institute, Taiwan. Though continuing to explore the structure-function relation of snake venom proteins, he has expanded his interest in understanding the adaptative features of halotolerant organisms. Currently, he is a professor at VIT University (India) and has more than 75 research articles published in international journals.

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