doi: 10.4172/2168-9652.1000e134

T e enzymes of sulfatase family (EC 3.1.6) are ubiquitously present across all domains of life and are involved in catalyzing hydrolysis of sulfate ester bonds for a vast array of substrates [1,2]. T e substrates for these enzymes range from small cytosolic steriods, like estrogen sulfate, to complex cell surface carbohydrates, such as the glycosaminoglycans. 17 sulfatases have been identi fed in humans, with 14 of them known to be associated with speci fc catabolic activities [1,2]. Sulfatases have been linked with various pathophysiological conditions including, lysosomal storage disorders, developmental abnormalities, and bacterial pathogenesis [1,2]. Biological importance of sulfatases is further manifested by several inherited metabolic diseases in humans, including metachromatic leukodystrophy, mucopolysaccaridosis VI, ichthyosis, which are all known to occur due to de fciency of speci fc single sulfatases [3]. Moreover, the activities of all sulfatases are severely reduced in a rare inherited disorder named multiple sulfatase de fciency (MSD) [3]. Sulfatases from di ferent enterobacteria such as, Klebsiella, Salmonella,eaIBDC /T11 1 Tw T-0.00722

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