Isolation and Characterization of a ß-Glucuronidase of the Mollusk Bulimulus tenuissimus

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Efficient enzymes for the synthesis and degradation of glycosaminoglycans are important in natural biological systems of these glycoconjugates. Therefore, an understanding of the glycosidases structures and functions is imperative towards a better comprehension of their natural role. The aim of the present work was to study the enzyme involved in the metabolism of sulfated glycosaminoglycans in the mollusk Bulimulus tenuissimus. The mollusk tissues were homogenized at 4°C. in a 0.1 M sodium acetate buffer, pH 5.0, then centrifuged at 20,000 x g. The proteins in the supernatant were submitted to fractioning with increasing concentrations of ammonium sulfate (0-50% and 50-80%) getting the fractions F_1 and F_2 , respectively, with the best activity visualized in the fraction F_2 (50-80%), as verified by agarose gel electrophoresis carried in PDA buffer. A ß- glucuronidase (F_3) was isolated by gel filtration chromatography (Bio-gel A) - 1.5 m of fraction F_2 on flow of 1ml/3min. High Performance Liquid Chromatography (HPLC) revealed only a protein band, confirming a 1.8% yield after 170 fold purification, with molecular mass of 116 kDa determined by electrophoresis in polyacrylamide gel with SDS. The determination of a number of kinetic parameters ideal for pnitrophenyl- β -glucuronide catalysis by β - glucuronidase (F₃), demonstrated that its optimal activity is at pH 5.0 and temperature of 55°C for 4 hours, with an apparent K_m of 2.68 mM. A total of 0.88 µg of ß-glucuronidase is needed for the degradation of 0.09 µM p-N-ß-glucuronide. Future studies on the three dimensional structure and the amino acid sequence will enable a deeper discussion of the inhibition and stimulation of some compounds in the ß-glucuronidase activity of the mollusk Bulimulus tenuissimus. Supported by CNPq