Abstract in English

 β -N-Acetylhexosaminidase (EC 3.2.1.52) is exoglycosidase, which exhibits the unique properties in the filamentous fungi. Enzyme from these organisms are dimeric, inducible and secreted extracelluary. It is expressed as preproprotein, consists of a signal sequence, a large propeptid and a catalytic subunit. Although the enzyme is widely distributed, its structure differs in varies organisms. Bacteria have only monomeric hexosaminidase. Human enzymes are dimeric as well as fungal, but only hexosaminidase from filamentous fungi have the catalytic subunit noncovalently associated with the propeptide. Propeptide is a essential for the enzyme activity. It exists a homologues model of the catalytic subunit of β -N-acetylhexosaminidase from *Penicillium oxalicum*, but the structure of the propeptide has not yet been solved.

The first part of this diploma thesis deals with the optimization of production and purification conditions. The second part deals with structural studies of β -N-acetylhexosaminidases from the filamentous fungi *Penicillium oxalicum CCF* 3438. These studies were carried out using chemical cross-linking and high resolution mass spectrometry. The combination of these methods revealed region of the noncovalent interaction of the catalytic subunit with the propeptide.