

## CHAPTER V

### CONCLUSION

1. The purification of  $\beta$ -glucosidase from *Daldinia eschscholzii* required 4 steps including ammonium sulphate fractionation, SP-sepharose, Phenyl-sepharose, and Superdex-200. The enzyme  $\beta$ -glucosidase was obtained with 50.23 fold purification and 6.28% yield. The specific activity was 77.86 U/mg protein.
2. The enzyme had an apparent molecular weight of 64.2 kDa as analyzed by SDS-PAGE, and with a molecular weight of 65,747 Da estimated by MALDI-TOF spectrometry. An isoelectric point, pI, of 8.55, was obtained using isoelectric focusing.
3. The apparent  $K_m$  for PNPG as  $K_m$  of 1.52 mM, and  $V_{max}$  values of 3.21 mmole/min/mg.
4. Optimal activity with PNPG as the substrate was at pH 5.0 and 50°C. The enzyme was stable at pH 5.0 at temperatures up to 50 °C.
5. This enzyme was competitively inhibited by Glucose and glucono- $\delta$ -lactone, with  $K_i$  values of 0.79 mM, and 4.72  $\mu$ M, respectively.
6. The purified  $\beta$ -glucosidase was active against PNPG, cellobiose, sophorose, and gentiobiose, but did not hydrolyze lactose, sucrose, Avicel, and o-nitrophenyl- $\beta$ -D-galactopyranoside.
7. The activity of  $\beta$ -glucosidase was stimulated by  $Ca^{2+}$ ,  $Co^{2+}$ ,  $Mg^{2+}$ ,  $Mn^{2+}$ , glycerol, DMSO, DTT and EDTA and, strongly inhibited by  $Hg^{2+}$ .
8. The enzyme was chemically modified by EDC in the presence of glycine as nucleophile under various conditions to study role of carboxy groups in the catalytic mechanism of this enzyme.

9. The enzyme was inactivated by the binding of one mol of EDC per mol of the enzyme with a second-order rate constant of  $6.73 \times 10^{-2}$  mM/min. Glucono- $\delta$ -lactone as a competitive inhibitor, partly protected the active-site carboxy group against chemical modification, with  $k_d$  of  $70.74 \times 10^{-2}$  mM.
10. Transglycosylation activity of the enzyme toward alcohol was performed by using PNPG as glucosyl donor substrate. Glucoside products were separated and detected by TLC analysis.
11. The internal amino acid sequence of *D. eschscholzii*  $\beta$ -glucosidase has similarity to the sequence of the family 3  $\beta$ -glucosyl hydrolase.