

## **Pichia pastoris as a host to overexpress the thermostable T1 lipase from Geobacillus zalihae**

### **ABSTRACT**

*Pichia pastoris* was known to be a good expression system in producing various heterologous proteins. The gene encoding thermostable T1 lipase from *Geobacillus zalihae* was cloned into pPICZ $\hat{I}$ +B and expressed in *P. pastoris* strains (GS115, X-33 and KM71H) under regulation of alcohol oxidase promoter. The expression of the gene in *Escherichia coli* system showed low expression level. Therefore, this study would highlight on the overexpression of the T1 lipase in yeast extracellularly. Recombinant X-33/pPICZ $\hat{I}$ +B/T1-2 (XPB2), GS115/pPICZ $\hat{I}$ +B/T1-5 (GPB5) and KM71H/pPICZ $\hat{I}$ +B/T1-7 (KPB7) were chosen for optimization in shake flask. Optimization strategies showed that these recombinants preferred YPTM medium with initial induction OD<sub>600nm</sub> = 7 cell biomass and 2% (v/v) methanol to provide optimal expression conditions. Hyper-resistant transformants at 3000 ug/mL zeocin gave better expression than 100 ug/mL zeocin selection. Time course study by using different inocula age showed that OD<sub>600nm</sub> = 7 expressed lipase at the highest level. The highest expression level was attained with GPB5 (88 U/ml), XPB2 (81 U/ml) and KPB7 (51 U/ml). Western Blot analysis confirmed that the molecular mass of recombinant T1 lipase was 45 kDa. In conclusion, thermostable T1 lipase was successfully overexpressed by using secretory *P. pastoris* system with two-fold higher than *E. coli* system.

**Keyword:** Expression; *Geobacillus zalihae*; *Pichia pastoris* thermostable lipase