

Poster Presentation

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## *Kluyveromyces lactis* *SSO1* and *SEB1* genes are functional in *Saccharomyces cerevisiae* and enhance production of secreted proteins when overexpressed

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### Background

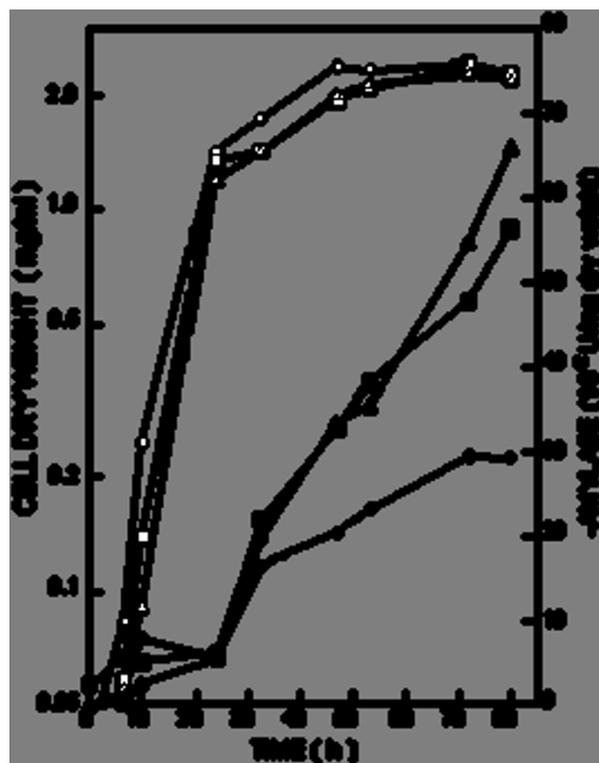
The *SEB1/SBH1* and the *SSO* genes encode components of the protein secretory machinery functioning at the opposite ends, ER translocation and exocytosis, respectively, of the secretory pathway of *Saccharomyces cerevisiae*. Overexpression of these genes can rescue temperature-sensitive (ts) growth defect of many *sec* mutants impaired in protein secretion. Their overexpression in wild-type yeast enhances production of secreted proteins in *S. cerevisiae*, which suggests that they may be rate-limiting factors in this process.

### Results

*Kluyveromyces lactis* homologs of *S. cerevisiae* *SEB* and *SSO* genes were isolated by multicopy suppression of *Saccharomyces* mutations [1]. *KISSO1* and *KISEB1* are up to 70% identical with the *S. cerevisiae* homologs at the amino acid level and can functionally replace them. These single copy genes were able to complement the ts growth defect of *sso2-1* and *seb1Δ seb2Δ sem1Δ* strains, respectively. In addition, *KISSO1* multicopy suppressed both sporulation defects of *S. cerevisiae* mutants *sso1Δ/sso1Δ* or *mso1Δ/mso1Δ* and ts growth defect of exocyst mutant *sec15-1*. Furthermore, *KISSO1* and *KISEB1* enhanced production of a secreted protein (Fig.1) similarly to *SSO* and *SEB1* genes of *S. cerevisiae* [2,3] when overexpressed.

### Conclusion

The single copy genes *KISSO1* and *KISEB1* are both structurally and functionally conserved in evolution with their duplicated *S. cerevisiae* homologs.



**Figure 1**  
Increased production of secreted  $\alpha$ -amylase by overexpression of *KISSO1* (▲) and *KISEB1* (■) in comparison to the control strain carrying empty vector (●). The cell growth is presented with open symbols.

## References

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