

Glycoengineering of Yeast Cells for Humanized N-glycans

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Due to its capacity to glycosylate proteins, yeasts have been considered for the production of glycoproteins derived from higher eukaryotes. However, yeast *N*-glycosylation pathway is only partially homologous to the pathway in human cells. Here we report the remodeling of glycosylation pathway for biosynthesis of humanized *N*-glycans in the methylotrophic yeast *Hansenula polymorpha*, which is rapidly gaining favor as a promising host for the production of recombinant proteins (1). Especially, *H. polymorpha* shows some advantages over the traditional yeast *Saccharomyces cerevisiae* in the production of recombinant glycoproteins for human therapeutic use. Most *N*-linked oligosaccharide species attached to *H. polymorpha*-derived glycoproteins have core-sized structures ($\text{Man}_{8-12}\text{GlcNAc}_2$) without highly immunogenic terminal α -1,3-linked mannose residues. Moreover, the outer chains of *H. polymorpha* *N*-linked glycans were shown to be extended mostly in 1,2-mannose linkages and to have very short 1,6 extensions, mainly composed of single α -1,6-linked mannose (2). As a first step toward humanizing *H. polymorpha* *N*-glycosylation pathway, the *H. polymorpha och2* mutant strain having a defect in the outer chain initiation on the core oligosaccharide $\text{Man}_8\text{GlcNAc}_2$ was developed and further engineered with the targeted expression of *Aspergillus saitoi* α -1,2-mannosidase in the ER. The engineered *H. polymorpha och2* strain was shown to produce the human high mannose-type $\text{Man}_5\text{GlcNAc}_2$ oligosaccharide as a major *N*-glycan (3). As an alternative approach, the remodeling of core oligosaccharide assembly pathway was carried out with additional deletion of the *H. polymorpha* ALG3 gene (*HpALG3*), encoding Dol-P-Man:- $\text{Man}_5\text{GlcNAc}_2$ -PP-Dol mannosyltransferase. The engineered double deletion (*alg3och2*) mutant strain expressing *A. saitoi* α -1,2-mannosidase generated mainly the trimannosyl-core form glycan ($\text{Man}_3\text{GlcNAc}_2$), which is an intermediate for further maturation to hu-

man-like complex *N*-glycans. These results demonstrate the potential of *H. polymorpha* to be developed as a host for the production of therapeutic glycoproteins with homogeneous complex *N*-glycan structures.

References

1. Kang H. A. and G. Gellissen (2005) *Hansenula polymorpha*. In production of recombinant proteins (ed. Gellissen G.), pp. 111-142. Wiley/VCH.
2. Kim, M.W., Rhee, S.K., Kim, J.Y., Shimma, Y., Chiba, Y., Jigami, Y., and Kang, H.A. (2004) Characterization of *N*-linked oligosaccharides assembled on secretory recombinant glucoseoxidase and cell wall mannoproteins from the methylotrophic yeast *Hansenula polymorpha*. *Glycobiol.* 14, 243-251.
3. Kim, M.W., Heo J. H., Rhee, S.K, and Kang, H.A. (2004) A novel *Hansenula polymorpha* gene coding for alpha 1,6-mannosyltransferase and process for the production of recombinant glycoproteins with *Hansenula polymorpha* mutant strain deficient in the same gene. PCT/KR2004/001819.