

Overexpression of Dihydroxyacetone Synthase 1 in Mut+ *Komagataella phaffii* (*Pichia pastoris*) Improves the Tolerance to Methanol and the Productivity of Recombinant Proteins

Comunicación oral

Biotecnología y bioprocesos

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Komagataella phaffii is a widely used methanol-inducible system for recombinant protein (rProt) production. However, methanol as a sole carbon source reduces the specific growth rate (μ) and promotes intracellular accumulation of formaldehyde, a toxic intermediate. To overcome these limitations and improve rProt productivity, we modified the methanol-assimilative pathway by overexpressing dihydroxyacetone synthase 1 (DAS1).

Three *K. phaffii* GS115 strains were engineered to overexpress DAS1 under different promoters: the strong constitutive glyceraldehyde-3-phosphate dehydrogenase promoter (P_{GAP}), the weak methanol-inducible peroxisomal malate dehydrogenase promoter (P_{MDH3}), and the strong methanol-inducible alcohol oxidase 1 promoter (P_{AOX1}). Each strain expressed *Candida antarctica* lipase B (CalB) as a model protein. Kinetic characterization was performed in deep-well plates using methanol concentrations ranging from 1 to 20 g·L⁻¹. CalB production was assessed in shake-flask and controlled bioreactor cultures.

PAOX1-DAS1 and P_{GAP} -DAS1 strains exhibited higher methanol tolerance, with inhibition constants (K_i) of 52 and 45 g·L⁻¹, respectively, compared to the control (K_i =

27 g·L⁻¹). Conversely, P_{MDH3}-DAS1 displayed increased methanol sensitivity (K_i = 17 g·L⁻¹). The maximum specific growth rate (μ_{max}) was highest in P_{GAP}-DAS1 (0.213 h⁻¹). Regarding CalB production, PGAP-DAS1 showed the greatest improvement across all tested conditions, achieving a 21% increase in volumetric productivity (U·L⁻¹·h⁻¹) during batch bioreactor cultivation compared to the control strain.

These findings demonstrate that DAS1 overexpression, particularly under a strong constitutive promoter, enhances methanol tolerance and recombinant protein productivity in *K. phaffii*. This strategy provides a promising approach to improve bioprocess performance when using methanol as both inducer and carbon source.

Palabras Clave

(1) *Komagataella phaffii*, (2) DAS1, (3) methanol assimilation, (4) recombinant protein, (5) bioprocess optimization.

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