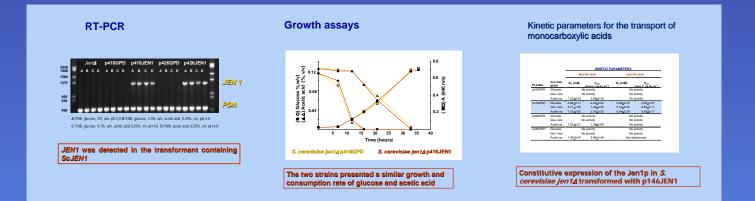
Exploring genetic tools for the overexpression of the lactate Jen1p of Saccharomyces cerevisiae: constitutive expression in S. cerevisiae and heterologous expression in Pichia pastor

Isabel Soares-Silva, Raquel P. Andrade, Dorit Schuller and Margarida Casal Centro de Ciências do Ambiente, Departamento de Biologia, Universidade do Minho, 4710-057 Braga Codex, Portugal 🖃 mcasal@bio uminho pt

Objective

In S. cerevisiae the active transport of lactate and pyruvate is dependent on the expression of JEN1 (1). JEN1 is the only S. cerevisiae member of the Sialate-Proton Symporters subfamily (TC#2.A.1.12) belonging to the Major Facilitator Superfamily (2). However members of other phylogenic subfamilies can be expected to transport monocarboxylic acids such as the five MCP Monocarboxylate Porters, the FNT Acetate:H* Symporter YHL008c or even the SSU1 Putative Transporter of Unknown Mechanism. To the data the possibilities whether Jen1p has regulatory (or sensor) or transport function haven't been discarded. The purpose of our work is to demonstrate non-ambiguously that Jen1p is a monocarboxylate transporter. Therefore the ScJEN1 gene was cloned in Pichia pastoris to produce significant amounts of active protein allowing heterologous reconstitution of lactate active transport in isolated membrane vesicles. The JEN1 gene was also overexpressed in S. cerevisiae (at a lower efficiency however) to characterize the kinetic properties of Jen1p at the cell level.

Constitutive Expression in Saccharomyes cerevisiae



heterologous expression in Pichia pastoris

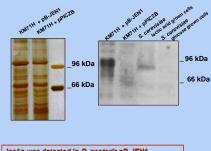
KM71H + pB-JEN1-I KM71H + pPICZB YPD MGY MM YPD MGY MM JEN1 rRNA

Northern-blot analysis

20 μg of total RNA extracted from transformant P. pastoris cells grown in YPD or MYG or incubated for 24 h in MM media were used.

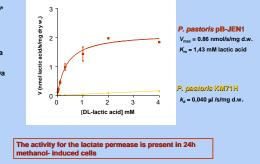
JEN1 mRNA was only detected in the transformant containing a copy of ScJEN1 after incubation in MM medium

Western-blot analysis



Jen1p was detected in P. pastoris pB-JEN1

Kinetic parameters of lactic acid uptake in *P. pastoris* KM71H and *P. pastoris* KM71H pBJEN1



Final remarks

- > Constitutive expression in S. cerevisiae was achieved
- >A 6-fold increase was obtained in Jen1p V_{max} in P. pastoris and only a 2-fold increase in S. cerevisiae
- >Jen1p was heterologous expressed in P. pastoris
- > JEN1 is a fully functional lactate permease

References

- ssal, M., Palva, S., Andrade, R. P., Gancedo, C., and Leão, C. (1999) The lactate-proton symport of Saccharomyces revisiae is encoded by JEPIJ. J Bacteriol 181:2620-2623 H Hertogh, B., Carvajal, F., Talla, E., Dujon, B., Barer, P., and Goffeau, A. (2002) Funct. Integr. Genomics 2, 154-170