

Multicopy suppressor screen

Dominique Loqué

Carnegie Institution

Wolf Frommer

Carnegie Institution

Method Article

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Abstract

Introduction

To identify mutations that restore the activity of a defective ammonium transporter, natural suppressors or suppressors generated by mutagenesis can be identified.

Procedure

1) Select suppressor mutants from the 31019b yeast strain⁽¹⁾ carrying the high copy number plasmid pDR-AtAMT1;1Thr460Ala by growing at 28°C for 8 days on solid YNB medium supplemented with 3% glucose and 2 mM ammonium chloride. 2) Pick suppressors and amplify in liquid YNB medium supplemented with 3% glucose and 2 mM ammonium. 3) Isolate plasmid from yeast by CTAB extraction and use for *E. coli* transformation. 4) Amplify plasmids in *E. coli*, isolate and sequence.

Anticipated Results

When an inactive transporter **Trans** is subjected to a multi-copy suppressor screen, point mutations (single base changes, primarily) may occur spontaneously in the sequence of **Trans**. Such a spontaneous mutation occurs first in a single plasmid molecule, yielding a single **Trans-Mut** sequence. If **Trans-Mut** does not contribute to transport and growth, the cell will die and the sequence will be lost. If **Trans-Mut** does improve transport, it will initially function in $(\text{Trans})_2(\text{Trans-Mut})_1$ complexes, as **Trans** is present in vast excess. If activity is sufficient to lead to growth, **Trans-Mut** will replicate sufficiently to be observed by plasmid DNA sequencing. Reversion to the wild-type sequence in a single plasmid molecule would yield $(\text{Trans})_2(\text{wt})_1$ and $(\text{Trans})_1(\text{wt})_2$ complexes at the outset ($(\text{wt})_3$ would be a sufficiently small portion of the population that it would not contribute to growth). The absence of wild-type revertants from the suppressor screens suggests that these complexes are inactive.

References

(1) Marini, A. M., Soussi-Boudekou, S., Vissers, S. & André, B. A family of ammonium transporters in *Saccharomyces cerevisiae*. *Mol. Cell. Biol.* **17**, 4282-4293 (1997).