

Erratum

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**The isolation of specific genes
from the basidiomycete *Schizophyllum commune***Eunice H. Froeliger¹, Alfredo M. Muñoz-Rivas^{2,*}, Charles A. Specht², Robert C. Ullrich², and Charles P. Novotny¹Departments of ¹Microbiology and ²Botany, University of Vermont, Burlington, VT 05405, USA

Due to an unfortunate printing error, Fig. 1 on p. 550 in the above article was printed upside down. The figure is corrected below.

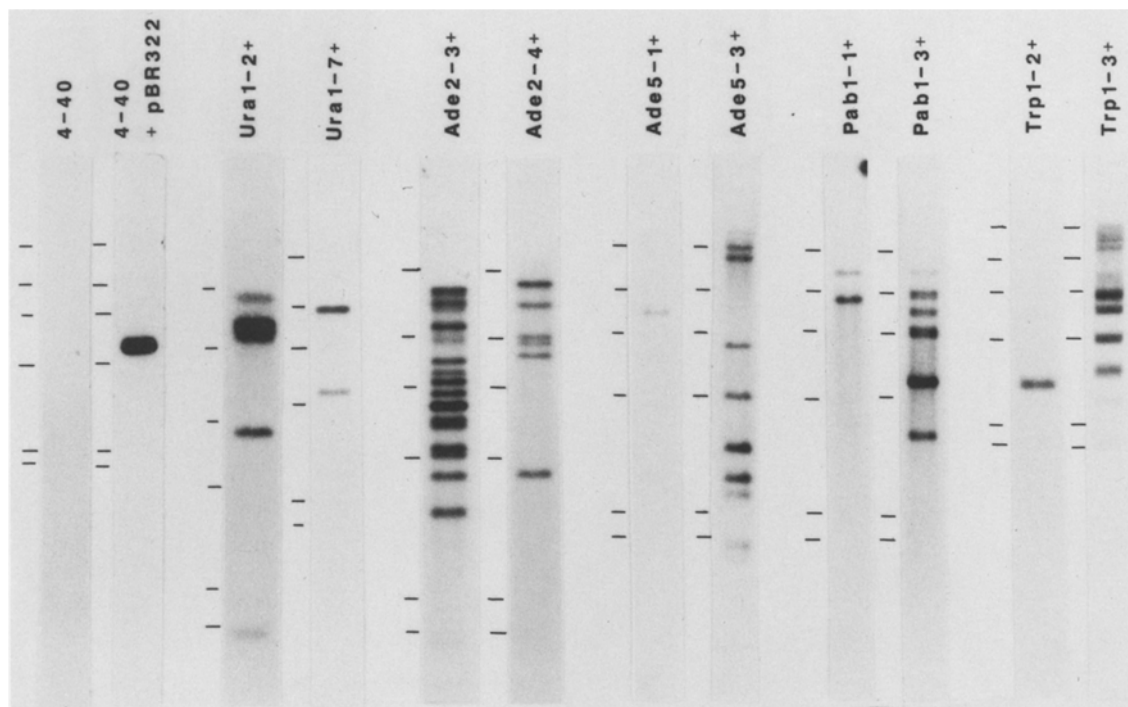


Fig. 1. Southern analysis of DNA from ten *S. commune* transformants and a nontransformed (control) strain. Analyses of two transformants are shown for each mutation used in transformation experiments. Genetic symbols are described in Materials and methods; for transformants, the second digit refers to a particular isolate. Samples of total DNA from transformants and the nontransformed strain, (HVD4-40) were digested with *EcoRI* and fractionated by electrophoresis in 0.85% agarose gels. Blots were hybridized with radiolabeled pBR322 DNA. Plasmid pBR322, restricted with *EcoRI*, was added to one sample of HVD4-40 DNA as a positive control for hybridization. Bars to the left of each lane indicate the positions of molecular weight standards which are, from top to bottom, 23.1, 9.4, 6.6, 4.4, 2.3, and 2.0 kilobases

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