

Table S2. Strains in columns 1) and 2) are strains YHE1223 and YHE1407 with the corresponding plasmid from column 2) of Table S3 replacing pJ533. Strains in column 3) were derived from strain YHE1223 by integrative recombination of *SUP35NMspecies+C* replacing *sup35::kanMX*. Strains in column 4) were derived from strain HJK001 by integration of the corresponding *SUP35NMspecies+C* construct, followed by meiotic crosses.

Species source of Sup35NM	1) SUP35 deletion strain with LEU2 CEN P _{SUP35} NM+C	2) For cytoduction: <i>sup35Δ</i> strain with LEU2 CEN P _{SUP35} NM+C	3) SUP35 integrated strain	4) For cytoduction: SUP35 integrated strain
<i>S. cerevisiae</i>	YHE1227	YHE1409	HJK088	YAZ013
<i>C. albicans1</i>	YAG1	YHE1430	HJK089	YHE1387
<i>C. albicans2</i>	YAG2	YHE1431		
<i>K. lactis</i>	YAG5	YHE1435	HJK092	YHE1425
<i>D. hansenii</i>	YAG9	YHE1432		YHE1385
<i>C. maltosa</i>	YAG11	YHE1434	HJK122	YHE1389
<i>A. nidulans</i>	YAG6	YHE1459		
<i>C. glabrata</i>	YAG10			
<i>N. crassa</i>	YAG13			
<i>M. grisea</i>	YAG3	YHE1412		
<i>A. fumigatus</i>	YAG4	YHE1411		
<i>S. pombe</i>	YAG7	YHE1436	HJK110	YHE1383
<i>A. gossypii</i>	YAG12	YHE1433	HJK111	YHE1391
<i>C. neoformans</i>	YAG14	YHE1410		
<i>U. maydis</i>	YHE1228	YHE1413		