

Table S4.

Strain #	Genotype	Source
AH13	<i>MAT a/α lys2/lys2 ho::LYS2/ho::LYS2 leu2/leu2 his3/his3 ura3/ura3 trp1/trp1</i>	Hochwagen group
SAy771	<i>MAT a/α lys2/lys2 ho::LYS2/ho::LYS2 LEU2/leu2 ime4::HIS3/ime4::HIS3</i>	Agarwala, Blitzblau 2012
SAy821	<i>MAT a/α lys2/lys2 ho::LYS2/ho::LYS2</i>	Agarwala, Blitzblau 2012
SAy834	<i>MAT a/α lys2/lys2 ho::LYS2/ho::LYS2 ime1::HIS3MX6/ime1::HIS3mx6</i>	Agarwala, Blitzblau 2012
SAy841	<i>MAT a/α lys2/lys2 ho::LYS2/ho::LYS2 ndt80::LEU2/ndt80::LEU2</i>	Agarwala, Blitzblau 2012
SAy859	<i>MAT a/α lys2/lys2 ho::LYS2/ho::LYS2 ime2::KanMX/ime2::KanMX</i>	Agarwala, Blitzblau 2012
SAy914	<i>MAT a/α lys2/lys2 ho::LYS2/ho::LYS2 3xmyc-IME4/3xmyc-IME4</i>	Agarwala, Blitzblau 2012
SAy966	<i>MAT a/α lys2/lys2 ho::LYS2/ho::LYS2 ndt80::LEU2/ndt80::LEU2 ime4::HIS3/ime4::HIS3</i>	This study
SAy995	<i>MAT a/α lys2/lys2 ho::LYS2/ho::LYS2 leu2::hisG/LEU2 his3::hisG/HIS3 trp1::hisG/TRP1 P_{GAL1}-NDT80::TRP1/ndt80::LEU2, ura3::P_{GPD1}-GAL4(848).ER::URA3/URA3</i>	Agarwala, Blitzblau 2012
SAy1086	<i>MAT a/α lys2/lys2 ho::LYS2/ho::LYS2 ime4-D349A,W351A/ime4-D349A,W351A</i>	Agarwala, Blitzblau 2012
SAy1235	<i>MAT a/α lys2/lys2 ho::LYS2/ho::LYS2 LEU2/leu2 TRP1/trp1 3xHA-MUM2/3xHA-MUM2</i>	Agarwala, Blitzblau 2012
SAy1248	<i>MAT a/α lys2/lys2 ho::LYS2/ho::LYS2 ime4::P_{CUP1}-IME4::KanMX4/IME4 mum2::P_{CUP1}-MUM2::KanMX4/MUM2 slz1::P_{CUP1}-SLZ1::KanMX4/SLZ1</i>	Agarwala, Blitzblau 2012
SAy1254	<i>MAT a/α lys2/lys2 ho::LYS2/ho::LYS2 LEU2/leu2 TRP1/trp1 3xHA-SLZ1/3xHA-SLZ1</i>	Agarwala, Blitzblau 2012
SAy1280	<i>MAT a/α lys2/lys2 ho::LYS2/ho::LYS2 ndt80::LEU2/ndt80::LEU2 ime4-cat/ime4-cat</i>	This study
SAy1310	<i>MAT a/α lys2/lys2 ho::LYS2/ho::LYS2 ndt80::LEU2/ndt80::LEU2 mum2::HIS3/mum2::HIS3</i>	This study
SAy1383	<i>MAT a/α lys2/lys2 ho::LYS2/ho::LYS2 ime1::HIS3MX6/ime1::HIS3mx6 ime4::P_{CUP1}-IME4::KanMX4/IME4</i>	This study
SAy1332	<i>MAT a lys2 ho::LYS2 ime4::HIS3 HIS3 mrb1::KanMX</i>	This study
SAy1333	<i>MAT a lys2 ho::LYS2 ime4::HIS3 HIS3 leu2 mrb1::KanMX</i>	This study
SAy1345	<i>MAT a/α lys2/lys2 ho::LYS2/ho::LYS2 mrb36::KanMX/mrb36::KanMX</i>	This study

SAy1384	<i>MAT a/α lys2/lys2 ho::LYS2/ho::LYS2 ime1::HIS3MX6/ime1::HIS3mx6 mum2::P_{CUP1}-MUM2::KanMX4/MUM2</i>	This study
SAy1385	<i>MAT a/α lys2/lys2 ho::LYS2/ho::LYS2 ime1::HIS3MX6/ime1::HIS3mx6 slz1::P_{CUP1}-SLZ1::KanMX4/SLZ1</i>	This study
SAy1390¹	<i>MAT a/α lys2/lys2 ho::LYS2/ho::LYS2 leu2::hisG/ leu2::hisG his3::hisG/ leu2::hisG trp1::hisG/trp1::hisG P_{GAL1}-NDT80::TRP1/ndt80::LEU2, ura3::P_{GPD1}-GAL4(848).ER::URA3/URA3 ime4::HIS3/IME4 yLR149c-mut/ yLR149c</i>	This study
SAy1392¹	<i>MAT a/α lys2/lys2 ho::LYS2/ho::LYS2 leu2::hisG/ leu2::hisG his3::hisG/ leu2::hisG trp1::hisG/trp1::hisG P_{GAL1}-NDT80::TRP1/ndt80::LEU2, ura3::P_{GPD1}-GAL4(848).ER::URA3/URA3 ime4::HIS3/IME4 YSP3-mut/YSP3</i>	This study
SAy1394¹	<i>MAT a/α lys2/lys2 ho::LYS2/ho::LYS2 leu2::hisG/ leu2::hisG his3::hisG/ leu2::hisG trp1::hisG/trp1::hisG P_{GAL1}-NDT80::TRP1/ndt80::LEU2, ura3::P_{GPD1}-GAL4(848).ER::URA3/URA3 ime4::HIS3/IME4 MUS81-mut/MUS81</i>	This study
SAy1418¹	<i>MAT a/α lys2/lys2 ho::LYS2/ho::LYS2 leu2::hisG/ leu2::hisG his3::hisG/ leu2::hisG trp1::hisG/trp1::hisG P_{GAL1}-NDT80::TRP1/ndt80::LEU2, ura3::P_{GPD1}-GAL4(848).ER::URA3/URA3 ime4::HIS3/IME4 yNL195c-mut/ yNL195c</i>	This study
SAy1422	<i>S. mikatae MAT a/α HO+</i>	This study
SAy1424	<i>S. mikatae MAT a/α HO+ ndt80::hphMX/ndt80::hphMX</i>	This study
SAy1425	<i>S. mikatae MAT a/α HO+ ndt80::hphMX/ndt80::hphMX ime4::NatMX/ime4::NatMX</i>	This study
SAy1432	<i>MAT a/α lys2/lys2 ho::LYS2/ho::LYS2 leu2::hisG/ leu2::hisG his3::hisG/ leu2::hisG trp1::hisG/trp1::hisG ime1::KanMX/ime1::HisMX 2μm-URA3-pGPD</i>	This study
SAy1433	<i>MAT a/α lys2/lys2 ho::LYS2/ho::LYS2 leu2::hisG/ leu2::hisG his3::hisG/ leu2::hisG trp1::hisG/trp1::hisG ime1::KanMX/ime1::HisMX 2μm-URA3-pGPD:SLZ1</i>	This study
SAy1434	<i>MAT a/α lys2/lys2 ho::LYS2/ho::LYS2 leu2::hisG/ leu2::hisG his3::hisG/ leu2::hisG trp1::hisG/trp1::hisG ime1::KanMX/ime1::HisMX 2μm-URA3-pGPD:slz1-nlsΔ (Truncation of n-terminal 97 aa)</i>	This study
SAy1438	<i>MAT a/α lys2/lys2 ho::LYS2/ho::LYS2 leu2::hisG/ leu2::hisG his3::hisG/ leu2::hisG trp1::hisG/trp1::hisG ime1::KanMX/ime1::HisMX 2μm-URA3-pGPD:slz1-SV40 (n-terminal 97 aa replaced by SV40 NLS sequence)</i>	This study
SAy1439	<i>MAT a/α lys2/lys2 ho::LYS2/ho::LYS2 leu2::hisG/ leu2::hisG his3::hisG/ leu2::hisG trp1::hisG/trp1::hisG ime1::KanMX/ime1::HisMX 2μm-URA3-pGPD:slz1-A1 (n-terminal 97 aa replaced by A1 NLS sequence)</i>	This study
SAy1441	<i>MAT a/α lys2/lys2 ho::LYS2/ho::LYS2 leu2::hisG/ leu2::hisG his3::hisG/ leu2::hisG trp1::hisG/trp1::hisG ime1::KanMX/ime1::HisMX 2μm-URA3-pGPD:slz1-Nop1^{nterm} (n-terminal 97 aa replaced by n-terminal 82aa of Nop1)</i>	This study
SAy 1469	<i>MAT a/α lys2/lys2 ho::LYS2/ho::LYS2 leu2::hisG/ leu2::hisG his3::hisG/ leu2::hisG trp1::hisG/trp1::hisG P_{GAL1}-NDT80::TRP1/ndt80::KANMX 3xHA-MUM2/MUM2 3xMYC-IME4/IME4</i>	This study

SAy1479 ¹	<i>MAT a/α lys2/lys2 ho::LYS2/ho::LYS2 leu2/leu2 his3/his3 ura3/ura3 trp1/trp1 ysp3-mut/ysp3-mut</i>	This study
SAy1480 ¹	<i>MAT a/α lys2/lys2 ho::LYS2/ho::LYS2 leu2/leu2 his3/his3 ura3/ura3 trp1/trp1 ysp3-mut/ysp3-mut</i>	This study
SAy1481 ¹	<i>MAT a/α lys2/lys2 ho::LYS2/ho::LYS2 leu2/leu2 his3/his3 ura3/ura3 trp1/trp1 ylr149c-mut/ylr149c-mut</i>	This study
SAy1482 ¹	<i>MAT a/α lys2/lys2 ho::LYS2/ho::LYS2 leu2/leu2 his3/his3 ura3/ura3 trp1/trp1 ynl195c-mut/ynl195c-mut</i>	This study
SAy1491 ¹	<i>MAT a/α lys2/lys2 ho::LYS2/ho::LYS2 leu2/leu2 his3/his3 ura3/ura3 trp1/trp1 mei5-mut/ mei5-mut</i>	This study
SAy1492 ¹	<i>MAT a/α lys2/lys2 ho::LYS2/ho::LYS2 leu2/leu2 his3/his3 ura3/ura3 trp1/trp1</i>	This study
SAy1493 ¹	<i>MAT a/α lys2/lys2 ho::LYS2/ho::LYS2 leu2/leu2 his3/his3 ura3/ura3 trp1/trp1 zip1-mut/zip1-mut</i>	This study
SAy1494 ¹	<i>MAT a/α lys2/lys2 ho::LYS2/ho::LYS2 leu2/leu2 his3/his3 ura3/ura3 trp1/trp1 nam8-mut/nam8-mut</i>	This study

¹The “-mut” allele designation refers to the allele that has been mutated at the m⁶A site.