**Table S2. *S. cerevisiae* strains**

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| Straina | Genotype | Source |
| NH2221 | *MAT***a** *leu2 arg4-Nsp hoΔ::LYS2 ura3 DMC1::URA3 dmc1::LEU2 mek1-as lys4Δ::hphMX4**MATα leu2 arg4-Nsp hoΔ::LYS2 ura3 DMC1::URA3 dmc1::LEU2 mek1-as lys4Δ::hphMX4**ndt80Δ::natMX4**ndt80Δ::natMX4* | This work |
| NH716 | *MATα leu2::hisG his4-X::LEU2(NgoMIV+ ori) hoΔ::hisG ura3(Δpst-sma)**MAT***a** *leu2::hisG HIS4::LEU2(BamH + oriI) hoΔ::hisG ura3(Δpst-sma)* | [1] |
| NH792 | NH716 only *dmc1Δ::kanMX6* | [2] |
| NH2188 | NH716 only *ndt80Δ::natMX4* | [2] |
| NH1054 | NH716 only *sae2Δ/com1Δ::kanMX6* | this work |
| NH1055 | NH716 only *spo11Δ::natMX4* | this work |
| NH2337 | NH716 only *spo11Δ::natMX4 ndt80Δ::hphMX4* | this work |
| NH2179b | NH716 only  *MAT leu2::hisG his4-X::LEU2(NgoMIV+ ori) zip1Δ::kanMX6* *trp1-5’Δ::hphMX4* *MAT leu2::hisG his4-X::LEU2(NgoMIV+ ori)* *MAT***a** *leu2::hisG HIS4::LEU2(BamH + oriI)* | this work |
| NH2234c | NH2179 only *ndt80Δ::natMX4* | this work |
| NH2228b | NH2179 only *zip1Δ::natMX4*  | this work |
| NH2230b | NH2228 only *kanMX6::pCLB2-3HA-SGS1* | this work |
| NH2242b | NH2230 only *zip2Δ::KlURA3* | this work |
| NH2243b | NH2230 only *zip3Δ::KlURA3* | this work |
| NH2249b | NH2230 only *spo11Δ::KlURA3* | this work |
| NH2265b | NH2230 only *msh5Δ::KlURA3* | this work |
| NH2266b | NH2230 only *sae2Δ::KlURA3* | this work |
| NH2267b | NH2230 only *mer3Δ::KlURA3* | this work |
| NH2271b | NH2230 only *spo16Δ::KlURA3* | this work |
| NH2305b | NH2230 only *zip4Δ::KlURA3* | this work |
| NH2297 c | same as NH716 only *zip1Δ::kanMX6 trp1-5’Δ::hphMX4* | this work |
| NH2298c | same as NH716 only *zip1Δ::natMX4* *trp1-5’Δ::hphMX4* | this work |
| NH2299c | same as NH716 only *zip1Δ::natMX4* *trp1-5’Δ::hphMX4 kanMX6::pCLB2-3HA-SGS1* | this work |
| NH2300b | same as NH716 only *zip1Δ::kanMX6* *trp1-5’Δ::hphMX4 mus81Δ::natMX4* | this work |
| NH2322 | same as NH716 only *cdc7-as3-9myc* | this work |
| NH2309-6-3 | *MAT leu2 arg4-Nsp lys2 hoΔ::LYS2 cdc7-as3-9myc mcm5-bob1 ura3 trp1-5’Δ::hphMX4* | this work |
| NHY957 | *MAT***a** *CENIII::ADE2 leu2::hisG his4-B* *trp5-S CYH2 met13-B LYS5* *can1 ura3(ΔSma-Pst)**MATα CENIII LEU2 HIS4* *TRP1 cyh2 MET13 lys5-P* *CAN1 ura3(ΔSma-Pst)* *CENVIII::URA3 thr1-a cup1* *ade2* *ho::hisG TRP1* *CENVIII THR1 CUP1 ade2 ho::hisG TRP1* | [3] |
| NH2241b | *MAT***a** *CENIII::ADE2 leu2::hisG his4-B* *trp5-S CYH2 met13-B LYS5* *can1 ura3(ΔSma-Pst)**MATα CENIII LEU2 HIS4* *TRP1 cyh2 MET13 lys5-P* *CAN1 ura3(ΔSma-Pst)* *CENVIII::URA3 thr1-a cup1* *ade2* *ho::hisG TRP1* *zip1Δ::natMX4**CENVIII THR1 CUP1 ade2 ho::hisB trp1-5’Δ::hphMX4::TRP1 zip1Δ::natMX4* | this work |
| NH144 | *MAT***a** *leu2ΔhisG his4-x ARG4 ura3 lys2 hoΔ::LYS2* *MATα leu2-K HIS4 arg4-Nsp ura3 lys2 hoΔ::LYS2*  | [4] |
| NH774d | NH144 only *sml1Δ::HA mec1(∆1632-2368)::kanMX6* | this work |
| NH742d | NH144 only *tel1(Δ2381-2787)::kanMX6* | this work |
| NH2072 | NH144 only *CDC7-3FLAG::kanMX6 ndt80Δ::hphMX4* | this work |
| NH2082 | NH144 only *cdc7-as-3FLAG::kanMX6 ndt80Δ::hphMX4* | this work |
| NH2309-6-3 | *MATα leu2 arg4-Nsp lys2 hoΔ::LYS2 cdc7-myc-as3 mcm5-bob1 ura3 trp1-5’Δ::hphMX4* | this work |
| NH2311 | *MAT***a** *leu2::hisG ho::LYS2 lys2 ura3 his3::hisG trp1::hisG::ZIP1::TRP1 zip1::LEU2* *MATα leu2::hisG ho::LYS2 lys2 ura3 his3::hisG trp1::hisG::ZIP1::TRP1 zip1::LEU2*  *SPO11-PROTEIN A::S.pombe his5*+ *SPO11-PROTEIN A::S.pombe his5+* | this work |
| NH2312 | NH2311 only *TRP1 zip1::LEU2* | this work |
| NH2313 | NH2311 only *zip1-4A* | this work |
| NH2314 | NH2311 only *ZIP1-4D* | this work |
| NH520/pLW3 | NH144 only *dmc1Δ::LEU2* *mek1::kanX6/2μ URA3 GST-mek1-as* | [5] |
| NH566::pLT11::pXC6 | *MAT***a** *leu2 HIS4 hoΔ::LYS2 arg4-Nsp hop1::LEU2 mek1Δ::LEU2 ura3::HOP1::URA3* *MATα leu2 his4 hoΔ::LYS2 arg4-Nsp hop1::LEU2 mek1Δ::LEU2 ura3* *ade2::mek1-K199R-ADE2**ade2* | this work |
| BR5892-8A::pB211\*b | *MATα leu2-3,112 trp1-289 ura3-1 thr1-4::MATa THR1 ade2-1 CTF10-myc-kanMX6 zip1Δ::URA3 spo1∆::ADE2* | this work |
| NH2247\*b | *MAT***a** *leu2-3,112 his4-260 trp1-289 thr1-4 arg4 lys2 ade2-1 ura3 CTF19-myc-kanMX6**MAT leu2-3,112 his4-260 trp1-289 thr1-4 ARG4 lys2 ade2-1 ura3 CTF19-myc-kanMX6**zip1Δ::URA3 zip2Δ::LEU2**zip1Δ::URA3 zip2Δ::LEU2* | this work |
| BR5894 ECM11-myc-kan\*b | *MAT***a** *leu2-3,112 his4-260 trp1-289 thr1-4 ade2-1 ECM11 zip1Δ::URA3* *MATα leu2-3,112 his4-260 trp1-289 thr1-4 ade2-1 ECM11-myc-kan zip1Δ::URA3* | this work |

aAll strains are derived from the SK1 strain background, except those marked with an asterisk, which are derived from the BR strain background.

bStrains contain a single copy of either pRS304, p382 (*ZIP1*), p382-4A (*zip1-4A*) or p382-4D (*zip1-4D*) integrated at *trp1*.

c Strains are homozygous for either pRS304, p382 (*ZIP1*), p382-4A (*zip1-4A*) or p382-4D (*zip1-4D*) integrated at *trp1.*

dThe *MEC1* deletion removes the PI3/PI4 kinase domain located between 2074-2316. The *TEL1* deletion removes the PI3/PI4 kinase domain between 2459-2704.

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