

S4 Table: List of oligonucleotides used in this study (1 of 3)

Primer	Sequence	Purpose
Rat1 FW 1	CGGAAAGATGGCACATGG	<i>dhp1</i> genotyping
Rat1 RV 3	ATGTTTCGTTGCTTGCTGTGC	<i>dhp1</i> genotyping
Dhp1-1 FW 1	CAACCCTTCAAGCCTTACGA	<i>dhp1-1>>ura4</i> genotyping, tagging <i>dhp1/dhp1-1</i>
Ura 5' RV 1	TAGCCAGTGGGATTTGTAGC	<i>dhp1-1>>ura4</i> genotyping
Din1 UP F	TCCAACCTGCATTACGTTG	<i>din1</i> genotyping, tagging <i>din1</i>
Din1 RV 1	CGACAGGATCAGCGTTTCG	<i>din1</i> genotyping
Rrp6 3' FW	GACCTCTCTATAGTTGCCTG	<i>rrp6</i> genotyping
Rrp6 3' RV	TACTTAGCGTTTCCTCGCAG	<i>rrp6</i> genotyping
Ago1_frw_check	ATGCGGTAATTTAAGAAAAGC	<i>ago1</i> Δ genotyping
Ago1_rev_check	TCTGTATTTCTATATTGGCG	<i>ago1</i> Δ genotyping
Ago1 Inside F		<i>ago1</i> genotyping
Ago1 Inside R		<i>ago1</i> genotyping
Clr4 UP	TTGCTCTGAAATTGAACACATCGAC	<i>clr4</i> genotyping
Clr4 DOWN	CCGAGTCGGAGTTACTTCCCTTTAGTC	<i>clr4</i> genotyping
Clr4 RV 3	CTTCCTGATATAGCACTTCCC	<i>clr4</i> Δ genotyping
Dcr1 FW 1	CCTATACTGATGGACATGGG	<i>dcr1</i> Δ genotyping, amplify <i>dcr1</i> Δ::kanMX cassette
Dcr1 RV 2	TTGCTGGAATACGGTTGCAG	<i>dcr1</i> Δ genotyping
Swi6 UP	GTGATGCTAGCCATTCTGTAC	<i>swi6</i> genotyping
Swi6 DOWN	AGAGTCGCTTTATGAGAAGC	<i>swi6</i> genotyping
Swi6 RV 3	CTGATGTCTTCATGTGTTGC	<i>swi6</i> Δ genotyping
Clr3 RV 2	CACGTGCTAACCATTACACC	<i>clr3</i> genotyping
Epe1 RV 2	GTGTGTTGTAGTATCCGGCT	<i>epe1</i> Δ genotyping
Rhn1 RV 2	CTGCACAACACATGGATGCG	<i>rhn1</i> Δ genotyping
Sir2 RV 3	GGTTGTGGTTTAGGTTCTCG	<i>sir2</i> genotyping
Kan2	ATCATTGGCAACGCTACCTT	<i>KanMX6</i> genotyping (5' RV)
Kan3	GCGCAATCACGAATGAATAA	<i>KanMX6</i> genotyping (3' FW)
Nat2	GGTGTCGGTGGTGAAGGACC	<i>NatN2</i> genotyping (5' RV)
Nat3	TGCCCTGCCCTAATCTCGA	<i>NatN2</i> genotyping (3' RV)
MT1	AGAAGAGAGAGTAGTTGAAG	Mating-type genotyping
MP	ACGGTAGTCATCGGTCTTCC	Mating-type genotyping
MM	TACGTTCACTAGACGTAGTG	Mating-type genotyping
Ura4 DS/E #1	GAGGGGATGAAAATCCCAT	<i>DS/E</i> genotyping, <i>ura4</i> ⁺ versus <i>DS/E</i> competitive PCR
Ura4 DS/E #2	TTCGACAACAGGATTACGACC	<i>DS/E</i> genotyping, <i>ura4</i> ⁺ versus <i>DS/E</i> competitive PCR
Ade6 #1	TGCGATGCACCTGACCAGGAAAGT	<i>DN/N</i> genotyping, <i>ade6</i> ⁺ versus <i>DN/N</i> competitive PCR
Ade6 #2	AGAGTTGGGTGTTGATTTGCTGA	<i>DN/N</i> genotyping, <i>ade6</i> ⁺ versus <i>DN/N</i> competitive PCR
JPO4	CGTGAGTATACAAACAATACTAGG	<i>otr::ura4</i> genotyping
JPO17	CTACTCTTCTCGATGATCCTGTAA	<i>otr::ura4</i> genotyping
Ade6 3' FW 1	GCGGACCATAGACATAACTG	<i>KΔ::ade6</i> ⁺ genotyping
K RV 2	GGTTTGTGTTAGCGCACTTTG	<i>KΔ::ade6</i> ⁺ genotyping

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Primer	Sequence	Purpose
Kan_Fwd	CGGATCCCCGGGTAAATTA	Amplify <i>KanMX6</i> Cassette
Kan_Rev	GAATTCGAGCTCGTTTAAAC	Amplify <i>KanMX6</i> Cassette
Nat-F2	CGGATCCCCGGGTAAATTA	Amplify <i>NatN2</i> Cassette
Nat2-S2	ATCGATGAATTCGAGCTCG	Amplify <i>NatN2</i> Cassette
Ade6 Full FW	TATGCCTTTTCGATGCCAATTG	Amplify <i>ade6+</i> Cassette
Ade6 Full RV	GAACATGGTCAACGGGG	Amplify <i>ade6+</i> Cassette
Clr3 FW 1	CGTCATTCACTCGTGTTTGT	Amplify <i>clr3Δ::kanMX</i> Cassette
Clr3 RV 1	TGGATGAGGGATTTCCCTCA	Amplify <i>clr3Δ::kanMX</i> Cassette
Clr4 F1	ACACAGATGGGTTAGACTCC	Amplify <i>clr4Δ::kanMX</i> Cassette
Clr4 R1	CAGGCAGATCCTTCCCTTGA	Amplify <i>clr4Δ::kanMX</i> Cassette
Dcr1 RV 1	GGAAGCTCAATCAATCGACG	Amplify <i>dcr1Δ::kanMX</i> Cassette
Epe1 FW 1	GCCGAATATAGCCTGCTTCA	Amplify <i>epe1Δ::kanMX</i> Cassette
Epe1 RV 1	CGAGTCAAAGTGGATTGATGC	Amplify <i>epe1Δ::kanMX</i> Cassette
K FW	CCGAATACCACGAAGTTATTG	Amplify K region fragment 1 (<i>KΔ::ade6+</i>)
Ade6-KΔ R	CAATTGGCATCGAAAGGCATAGTTCAG ACTGATCTCGCAATT	Amplify K region fragment 1 (<i>KΔ::ade6+</i>)
Ade6-KΔ F	AAACCCCGTTGACCATGTTTCGCTTTCC TCTCAGCATTGAG	Amplify K region fragment 3 (<i>KΔ::ade6+</i>)
K RV	GATCGCCAAAGTTGTATTGC	Amplify K region fragment 3 (<i>KΔ::ade6+</i>)
Rhn1 FW 1	TCGTTTGGCGATGCTAACAGC	Amplify <i>rhn1Δ::kanMX</i> cassette
Rhn1 RV 1	GCATATCCTCTACTTCTGCC	Amplify <i>rhn1Δ::kanMX</i> cassette
Swi6 5' FW	CTTGAGAGACTCAGCATTGC	Amplify <i>swi6Δ::NATN2</i> cassette
Swi6 RV 1	CATGAAGAGCAGTGCATGC	Amplify <i>swi6Δ::NATN2</i> cassette
Nat_Rat1_R	TTAATTAACCCGGGGATCCGATAGTAGC CATTCCCTGTTATTAT	<i>dhp1</i> tagging
Nat_Rat1_F	CGAGTCGAATTCATCGATGCGGGTTTG GGATATTGGAG	<i>dhp1</i> tagging
Rat1_dw_R	GGGGACATTCTACTGTTGTAG	<i>dhp1</i> tagging
Kan_Din_R	TTAATTAACCCGGGGATCCGAATTATTT GTCGTAAAACAAG	<i>din1</i> tagging
Kan_Din_F	CGAGCTCGAATTCATCGATAGTAGGGT TATACACTAGATGT	<i>din1</i> tagging
Din DOWN R	CAACCACGCCCTCCATATAC	<i>din1</i> tagging
Sir2 FW 1	ACGCATTCCAGTTCTCATGG	<i>sir2</i> deletion
Sir2 FW 2	CCTCCTACTACTCCTCAGAT	<i>sir2</i> tagging
NAT Sir2 FW	CGAGCTCGAATTCATCGATGGACTGAT CGATCTATCGAA	<i>sir2</i> deletion/tagging
NAT Sir2 RV 1	TTAATTAACCCGGGGATCCGATAGACTT GCGCGTACCCTT	<i>sir2</i> deletion
NAT Sir2 RV 2	TTAATTAACCCGGGGATCCGACAATA CCGTTTGTGTGC	<i>sir2</i> tagging
Sir2 RV 1	ACCTTTTGGCATGCTCGATG	<i>sir2</i> deletion/tagging

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Primer	Sequence	Purpose
q-Actin F	GGTTTCGCTGGAGATGATG	qRT-PCR: <i>act1+</i> (Endogenous Control)
q-Actin R	ATACCACGCTTGCTTTGAG	qRT-PCR: <i>act1+</i> (Endogenous Control)
q-Leu1 F1	ATTGCTCGTTTGGCTGCTTG	qChIP: <i>leu1+</i> (Endogenous Control)
q-Leu1 R1	TACAGTCTTGCGCCAAAGAC	qChIP: <i>leu1+</i> (Endogenous Control)
q-Ade6 F	ATGCTTATCCTACAACCTGAGACC	qPCR: <i>ade6+</i>
q-Ade6 R	TGAATTGAGAAGGGAAGACGAG	qPCR: <i>ade6+</i>
q-Cen(dg) F	AATTGTGGTGGTGTGGTAATAC	qPCR: Centromere (<i>dg</i>)
q-Cen(dg) R	GGGTTTCATCGTTTCCATTCAG	qPCR: Centromere (<i>dg</i>)
q-Mat F3	AGTAATTCTATGTACCCCGTGC	qPCR: Mating-type locus (<i>cenH</i>)
q-Mat R3	CGTCGTCACCAATCGATCTT	qPCR: Mating-type locus (<i>cenH</i>)
qUra4 5' F1	CTAGATTCTAGGATGCGTATTGG	qPCR: Spreading Assay "A"
qUra4 5' R1	CAGTTTGCTCATGGGTACAAC	qPCR: Spreading Assay "A"
Ura4 Stul F1	AATTGCCATACAGTGCCAGG	qPCR: Spreading Assay "B"
q-CenH <i>ade6</i> FW 2	GGTTAAATTGTGGTGGTGTGG	qPCR: Spreading Assay "B"
q-CenH <i>ade6</i> FW 1	TGACAAAGGTGCCGAATCAAC	qPCR: Spreading Assay "C"
q-CenH <i>ade6</i> RV 1	TTGCAACAGGCGGCCAAGC	qPCR: Spreading Assay "C"
qUra4 3' F1	CTGGCAATGTTACGAGGAATC	qPCR: Spreading Assay "D"
qUra4 3' R1	TTCAGAAGAACATTACTIONCGTGC	qPCR: Spreading Assay "D"
Dhp1 D55A FW	GACAACTTGTATTTGGCCATGAATGG TATTGTTTCATC	<i>dhp1</i> mutagenesis
Dhp1 D55A RV	GATGAACAATACCATTCATGGCCAAA TACAAGTTGTC	<i>dhp1</i> mutagenesis
Dhp1 E207Q FW	GTTCCCTGGTGAAGGTCAACATAAGAT CATGGAATT	<i>dhp1</i> mutagenesis
Dhp1 E207Q RV	AATTCCATGATCTTATGTTGACCTTCA CCAGGAAC	<i>dhp1</i> mutagenesis
Pst1- Dhp1 FW	ATCGCTGCAGCGCTACCATTGATGA GTCTG	Clone fragment containing <i>dhp1+</i> gene into pREP41
BamHI-Dhp1 RV	TCTAGGATCCCTCGATTGATGGATCT TATGCATG	Clone fragment containing <i>dhp1+</i> gene into pREP41