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| **S2 Table.** Oligonucleotides used in these experiments |
| **Oligo** | **Sequence** | **Description** |
| DG\_90 | CAGCTCCATTGCGTATCATGTCCTTTGCTATCGCGTGTGCTGGTAGGATTGGCGTCTTTCCGGAACCTGAATACGATCCC | Repair oligo to introduce *pol3-01* mutation via Cas9 |
| DG\_91 | TTCATTGAAGATACTAGGAAAATTGCATTTGCCGATCCAGTAGTGATGGCTTTCGCAATAGCTACCACGAAGCCGCCTTTAAAATTCCCGGATTCCGC | Repair oligo to introduce *pol2-4* mutation via Cas9 |
| DG\_253 | ACGTCTAGCTGAGCATGTGA | Sense oligo in Y region of *MAT*⍺. Anneals 327 bp upstream of HO cut site |
| DG\_254 | TACGTTAAACATGAATCGA | Reverse oligo in Taf2. Anneals 1869 bp downstream of HO cut site |
| DG\_285 | GATGTCTGGGTTTTGTTTGGGATGCAATTTATTGCTTctcgagTCGGCTTCACAATTTGTTTTTCCACTTTTCTAACAGG | 1 kb deletion donor - inserts *Xho*I site at *MAT*⍺ after HO-induced DSB and resection |
| DG\_287 | AAAATGCAGCACGGAATATGGGACTACTTCGCGCAACctcgagAGTATAATTTTATAAACCCTGGTTTTGGTTTTGTAGA | 100% homology donor – inserts *Xho*I site at *MAT* after HO-induced DSB |
| DG\_310 | TCTACAAAACCAAAACCAGGGTTTATAAAATTATACTctcgagGTTGCGCGAAGTAGTCCCATATTCCGTGCTGCATTTT | Reverse complement of DG\_287 to create dsDNA template |
| DG\_344 | ATCTGAGTTACTGTCTGTTTTCCTTCTGCTCGCTGAAGAATGGCACGCGGACAAAATGCACTCGAGGCACGGAATATGGGACTACTTCGCGCAACAGTATAATAGGAAACCCGTTTCTTCTGACGTAAGGGTGCGCACACGCGGACAAAATGCAGCAGTTTCAGAGCTATGCTGGAAACA | *MAT*⍺.80-nt retron donor sequence to repair *MAT* and insert *Xho*I insertion |
| DG\_373 | AcAATGCAGCgCGGAATATaGGACTACTgCGCGCAACctcgagAGTATAATTTTATAAACCCTGGTTTTGGTTTTGTAGA | Inserts *Xho*I site at *MAT*⍺ after HO-induced DSB. Mismatches in homology every 9th bp on the 5’ side |
| DG\_374 | AAAATGCAGCACGGAATATGGGACTACTTCGCGCAACctcgagAGTATAATgTTATAAACtCTGGTTTTaGTTTTGTAtA | Inserts *Xho*I site at *MAT* after HO-induced DSB. Mismatches in homology every 9th bp on the 5’ side |
| DG\_396 | CGGCATCCTGCATTGAATCTGAGTTACTGTCTGTTTTCCTAGGAAACCCGTTTCTTCTGACGTAAGGGTGCGCAATGAGTTTACGTTCGAGGCGGTTTCAGAGCTATGCTGGAAACAGCATAGCAAGTTGAAAT | gBlock to clone into pZS165. gRNA to target 5bp insertion of *lys5* in JKM179No retron donor sequence |
| DG\_397 | CGGCATCCTGCATTGAATCTGAGTTACTGTCTGTTTTCCTTTCAAGAGGATATACTCGCGGATGAGTTTACGTTCGAGGCATTAATGAGAACTTTGCCATTGGCGTCTCAAGCCAGAATCAGGAAACCCGTTTCTTCTGACGTAAGGGTGCGCAATGAGTTTACGTTCGAGGCGGTTTCAGAGCTATGCTGGAAACAGCATAGCAAGTTGAAAT | target 5bp insertion of *lys5* in JKM17980-nt retron donor sequence to restore Lys5 to prototrophy |