Table S2. List of oligonucleotides used in this study.

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| **Oligo** | **Sequence** | **Description** |
| AS131 | AAGGAATGTGCCTCGTCAAATT | *dg* FWD, qPCR, sRNA northern |
| AS132 | TGCTTCACGGTATTTTTTGAAATC | *dg* REV, qPCR, RT, sRNA northern |
| GJ173 | ATTGACGCCGGTGTTAGTGTAGGT | *fbp1* FWD, qPCR |
| GJ174 | TGACACGATGACCTGTGGTAAGCA | *fbp1* REV, qPCR |
| GSqPCR\_27 | AGACGGTTCAAACTCCCTCT | *tam14* FWD, qPCR |
| GSqPCR\_28 | ACGAGGAATCTTGGTAGGAGGA | *tam14* REV, qPCR |
| AS133 | GTATTTGGATTCCATCGGTACTATGG | *dh* FWD, qPCR |
| AS134 | ACTACATCGACACAGAAAAGAAAACAA | *dh* REV, qPCR |
| XW411 | CATATCAATTACAAACATATTGTTTGCTGC | *cenH-mat2*, qPCR |
| XW412 | GCTTCATGTTAAGTTGATTGGTGTAATTAG | *cenH-mat2*, qPCR |
| GSqPCR\_67 | ACAGGGCGTCAACCATGACA | *tlh1* FWD, qPCR |
| GSqPCR\_68 | TTGACGGCTTGCACACGACT | *tlh1* REV, qPCR |
| NI1234 | GAGGTAAAGCGAATGATTAGAGGT | 28S rRNA, qPCR |
| NI1235 | TTAAAGTTTGAGAATAGGTTGAGGAAA | 28S rRNA, qPCR |
| NI1226/1227 | GTTGTTGCAGTTAAAAAGCTCGTA | 18S rRNA, qPCR |
| NI1226/1227 | ATGACCAGTAAACACGCCTTG | 18S rRNA, qPCR |
| GSqPCR\_71 | CGTCCATACCGTGAGTGATACC | *gfp* REV, qPCR, RT |
| GSqPCR\_72 | CGGTCCAGTTTTGTTGCCAGAC | *gfp* FWD, qPCR |
| GSqPCR\_87 | ACTGAACGTACTCCGAGACC | *mat3* FWD (position 6), qPCR |
| GSqPCR\_88 | TCGCATACGTACTTCCTTTGA | *mat3* REV (position 6), qPCR |
| AS49 | CAACCCTCAGCTTTGGGTCTTG | *act1* FWD, qPCR |
| AS50 | TCCTTTTGCATACGATCGGCAATAC | *act1* REV, qPCR, RT |
| GSqPCR\_129 | CAAATCACCGCTTTTGAGATTAATCGT | *IR-L:ade6* (position 1) FWD, qPCR |
| GSqPCR\_130 | CAACAGGCGGCCAAGCAATTT | *IR-L:ade6* (position 1) REV, qPCR |
| GSqPCR\_137 | GCAGTTTAGACGGAAAAGTTTATGC | *ade6* FWD (position 2), qPCR |
| GSqPCR\_138 | ATTGAGAAGGGAAGACGAGCAGG | *ade6* REV (position 2), qPCR, qRT |
| GSqPCR\_125 | AGGACCCATACAACAACGGA | *RE-II* (position 3) FWD, qPCR |
| GSqPCR\_126 | TGCCAAGCAAGCAACTGTTTA | *RE-II* (position 3) REV, qPCR |
| GSqPCR\_97 | TGAGCATTTGGCTGAAGCACC | *mat2* FWD (position 4), qPCR |
| GSqPCR\_98 | CCAAATACACCAAGCCATAATACGCAC | *mat2* REV (position 4), qPCR |
| GSqPCR\_139 | GATGATAACGGATCTAGCTTCGCCA | *cenH* FWD (position 5), qPCR |
| GSqPCR\_140 | GCCTCTCTCTATATATCAGATATAAAGATGCG | *cenH* REV (position 5), qPCR, qRT |
| GSqPCR\_89 | GTGCTTCAGCCAAATGCTCA | *mat2* FWD, qPCR |
| GSqPCR\_90 | ACTGAACCCTGCTTATATGTAGTT | *mat2* REV, qPCR, RT |