**Table S1.** Primers used in the gene mapping, gene cloning, and RT-qPCR

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| --- | --- | --- |
| Primer names | Forward (5'–3') | Reverse (5'–3') |
| pInDel2 | GTGCTCGTACAATGTGCTTAGA | GGTTGATGCTTGAAGACTTGGA |
| pInDel3 | CCTCTAAGAACTTCCACCTC | CTTGGTCTGGTCCGTGTA |
| p500-V | GAATTCCCCTTCTCAGCTTCTTCCTTC (*EcoR*I site underlined) | GGATCCAGTTTTGTTATCCTCCGCCCT (*BamH*I site underlined) |
| pSlCHS3-V | GAATTCGCCCAAACTCTTGTCCCCGA (*EcoR*I site underlined) | GGATCCCAATGTAAGCCCAGCCCACT (*BamH*I site underlined) |
| p520-V | GAATTCAACCAACAAAAAGTGAAAAAAA (*EcoR*I site underlined) | GGATCCAGAAATAACAACAAGCCCATAA (*BamH*I site underlined) |
| pSlCHS3-O | TCTAGATGACATTAGTTGTTGAAGCACG (*Xba*I site underlined) | GGTACCAGCCAGATGAAATTATTTTACGA (*Kpn*I site underlined) |
| pN-Q | GCCTATGGATTACCTCTTGATGA | TCTGTGAGTTTTGCCTGTTTTTT |
| pRPL2-Q | CAGCGGATGTCGTGCTATGAT | GGGATGCTCCACTGGATTCA |
| p500-Q | CCCTTCTCAGCTTCTTCCTTC | AGTTTTGTTATCCTCCGCCCT |
| pSlCHS3-Q | CAAAAGGCCATCAAAGAATG | ACTGGGTCCACGGAACGTAA |
| p520-Q | TTTTTCTTATGGGCTTGTTGTT | CTCTTCAGATTCCGCTTCTTCT |