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| --- | --- | --- |
| **Primer** | **Sequence** | **Usage** |
| Acf\_Rec\_NF | GCCATTTAAACTTAAGGACATTCAAGAGCAAAAGGAAAACACCAAACATGGAAGTGCATACCAATCAGGAC | recombineering |
| Acf\_Rec\_NR | TTCTTGCCCTCCTTCTGATTCAGGTCGAATCCTTCCCGCTTGCAAATGGGCTTGTCGTCGTCATCCTTGTA | recombineering |
| Acf\_Rec\_CF | TGCAGCTACCGTTTAGGCCTAGCGATATGAACGGGGAAGTCAAAGCTTGCGAAGTGCATACCAATCAGGAC | recombineering |
| Acf\_Rec\_CR | GTAGACTAACTAATTTACACGATAGCTGGTGGAGATCAGCGTCCGGCTCACTTGTCGTCGTCATCCTTGTA | recombineering |
| UAS\_ChIP\_f1 | TCCGAGCGGAGACTCTAGC | ChIP-qPCR, UAS |
| UAS\_ChIP\_r1 | TTTGCTTGTTTGAATTGAATTGTC | ChIP-qPCR, UAS |
| UAS\_ChIP\_f2 | TAGGGAATTGGGAATTCGAG | ChIP-qPCR, 5' of LacZ |
| UAS\_ChIP\_r2 | TTAGCAGGCTCTTTCGATCC | ChIP-qPCR, 5' of LacZ |
| Spt4\_fw | GCTCCGATTCATAAGCCCAG | ChIP-qPCR, negative control region |
| Spt4\_rv | GCCTCTTTCGGAGCAGCTTT | ChIP-qPCR, negative control region |
| LacZ\_RT\_f1 | CGCTAGAGTCGACCAATTCC | RT-qPCR |
| LacZ\_RT\_r1 | GGCAACGAAAATCACGTTCT | RT-qPCR |
| RpII-140\_fw | GCGCTATGGGTAAGCAAGCT | RT-qPCR, control |
| RpII-140\_rv | TCACAAGTGGCTTCATCGGA | RT-qPCR, control |
| Acf1\_crispr\_LHS6 | TTAGCGCAGCGTCCACCTAA | CRISPR/Cas9, gRNA |
| Acf1\_crispr\_RHS2 | GGGCGCCTACCAGCTAAACG | CRISPR/Cas9, gRNA |

**Supplementary Table 3**