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| **Primer name** | **Primer sequence** | **Annealing position**  |
| 1F | ACCAGTTGGAAGGAAAAGCTCT | 29733 |
| 2F | AGACGGCTCATTTGTGGGTT | 911 |
| 3F | CAACAGCAGCAAACACTGGG | 20496 |
| 4F | GACAACGGAAACATCGCACC | 28574 |
| 5F | GCACACCCTAACTGCGTCAT | 34377 |
| 6F | TGGACCCCTAGCGGAAGTTA | 23291 |
| 7F | ATGGCATGTCTAGCGCTCTC | 10532 |
| 8F | GTGGGCACTGCTAAGAGTGT | 32412 |
| 9F | TCAATCGGCTTTGAACGCAC | 726 |
| 10F | CTGAACGTTTCGGTCTTGCC | 31485 |
| 11F | GAATATCCACGCTGGCGAGA | 474 |
| 12F | TTGGGGTCGTCCTCACATTG | 15809 |
| 13F | AATTGAAGCACATCGGGGGA | 26835 |
| 14F | AGCAAGGAAACTGACTGGCA | 2625 |
| 15F | GCAGCGCTTGCGATTAGTAT | 27206 |
| 16F | GTCCGATGTGTGGTCACGAA | 29208 |
| 17F | TCAACCATTGGGCAGACGAA | 33405 |
| 26F | AAGAGCGGTGTCCTCGAAAG | 32230 |