|  |  |  |  |
| --- | --- | --- | --- |
| **Sites of Primers** | **Primer Number** | **Sequences** | **Product****size (bp)** |
| 2370 | 1-1\_F | ATCCATCCCCTCCAAATGTTG |   |
| 2172 | 1-1\_R | GAGGATCCTGGTGACATGCTC | 199 |
| 1564 | 1\_F | CCGACAGAGATGCTGAAGAG |   |
| 1285 | 1\_R | CTTCTCTCTGCAGAGCGATC | 299 |
| 650 | 2\_F | CGCGGAAGTTAGGTCTCCAG |   |
| 426 | 2\_R | GTGACGAGGTAGGGGCAGAG | 225 |
| 522 | 3\_F | GGTAGGGAGGGGACAGGAGA |   |
| 322 | 3\_R | TTCGATTCGGGACCAGGTAG | 210 |
| -348 | 4\_F | AGGCTGATTTCTGGCAGAGG |   |
| -546 | 4\_R | GGTAGCCCCACTTCCCTTTC | 199 |
| -527 | 5\_F | GAAAGGGAAGTGGGGCTACC |   |
| -714 | 5\_R | GTCAGGGTTGGCAAAGAACC | 188 |
| -889 | 6\_F | CTCCTAGGGGAGGGCAGTTT |   |
| -1088 | 6\_R | AGTAGCCTGAACCCCGGAAA | 200 |
| -1061 | 7\_F |  AAAGCAGTTTTCCGGGGTTC |   |
| -1261 | 7\_R | GGGTCTCCAAGCCCAAACTA | 201 |
| -1566 | 8\_F | ATCACCTCTCCCCAGCATGT |   |
| -1759 | 8\_R | AGATGGGGGTGCTCACTTTG | 194 |
| -1999 | 9\_F | GGAGGCTGAGGCGGGTGGATCACG |   |
| -2201 | 9\_R | CAATGGCGCGATCTCGGCTCACC | 203 |
| -2887 | 10\_F | TCCCTGCCTGCTCATCTCAG |   |
| -3088 | 10\_R | TCCTATGTGCAAGGCACCAC | 202 |
| -3618 | 11\_F | CCCATGGTTTCCCATCACTT |   |
| -3822 | 11\_R | CTTGTCCCCAGACAGGGAAG | 205 |
| -4367 | 12\_F | TTAGAGACCACAACCTGCTTTC |   |
| -4573 | 12\_R | CCTGTAAAGGCCCAGGTGTC | 207 |
| -4906 | 13\_F | GACAGATGCTGAGCCCTTCC |   |
| -5109 | 13\_R | TTGGCATTTAGCAACGCTGT | 204 |