|  |  |  |  |  |  |  |
| --- | --- | --- | --- | --- | --- | --- |
| Locus | Repeat motif | Primer Sequences (5’-3’) | Ta (ºC) | Size range (bp) | NA | GB |
| NZ | AAC | F: GCCCCTACGCGTAAAACATGR: TGAAACACACCATGGAATGCA | 55 | 129 - 201 | 29 | MF490249 |
| N3 | AC | F: AAAGGCCATCAATCCCAAGR: ATCGTGCTAGGGCTCGACTA | 55 | 160 - 226 | 38 | MF490250 |
| 7T | AAC | F: CGTGTTGCTCCTCACATTTCR: GGTCCTAGATGGGAGGCATA | 55 | 297 - 363 | 27 | MF490251 |
| UT | AC | F: TGTACAAATAACAGAGGCACCAAR: AGTGTCACATTCATCTGGATAGC | 55 | 135 - 207 | 36 | MF490252 |
| QL | AC | F: ACTCAAGAGAATGGCGATGAR: TGGTCCATATTGGTTGGAAA | 55 | 167 - 235 | 39 | MF490253 |
| 8D | AC | F: GAGGTGCCTAAGGTGGAACAR: TGCTCAACAAAACACCGAAA | 55 | 211 - 293 | 41 | MF490254 |