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|  | **Seq ID** | **Primer F** | **Primer R** |
|
| 1 | T5A6F1 | 5’- GACTCTGCCACAGACGACTAC -3’ | 5’- CCAGAAGAGGGAGTGGTTGAC-3’ |
| 2 | T5AFX3 | 5’- GACGATGGGTCGCTAGATGG-3’ | 5’- GTCCGGGAACTCGATAGCTG-3’ |
| 3 | T5ALL5 | 5’- CCTTTACCCCTCCCGAACAG -3’ | 5’- AATCCTTACCCGCGGTTCTC-3’ |
| 4 | T5AKK7 | 5’- GTGTGGCAACTTACAGCGTG-3’ | 5’- GTGTAGCTAAGCGTGACCGT-3’ |
| 5 | T5A5N4 | 5’- GCGCCTACGTCACAATCAAG-3’ | 5’- TCGTAGTTCATCAGGTTCAGG-3’ |
| 6 | T5A212 | 5’- TGTCCAACCAGGCCATTACC -3’ | 5’- AGCACGAGGACGGAAAAGAG -3’ |
| 7 | 18Srna(inner control) | 5’-CTCGTAGTTGAACCTTGGGC-3’ | 5’-AACAAAATAGAACCGCGCG-3 |