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| **ID** | **Forward primer** | **Reverse primer** |
| *ACTB* | 5’–TGAGCTGCGTGTGGCTCC–3’ | 5’–GGCATGGGGGAGGGCATACC–3’ |
| *RPS11* | 5’-TGCACTACATCCGCAAGTACA-3’ | 5’-CACGTTGAAGCGCACTGTCT-3’ |
| *RPS14* | 5’-CGTGTGACTGGTGGGATGAA-3’ | 5’-TCCAGGGGTCTTGGTCCTAT-3’ |
| genomic *tax* | 5′-GTTGTATGAGTGATTGGCGGGGTAA-3′ | 5′-TGTTTGGAGACTGTGTACAAGGCG-3′ |
| *TAX* | 5’-ATCCCGTGGAGACTCCTCAA-3’ | 5’-CCAAACACGTAGACTGGGTATCC-3’ |
| *PBM\_TAX* first | 5'-TCACAAATTTCAAACCAAGGCC+-3' | 5'-AACATGGGGAGGAAATGGGT-3' |
| *PBM\_TAX* internal | 5'-CTACTCTCACACGGCCTCAT-3' | 5'-GGACACGTCAGGGCCTAG-3' |