**Supplementary Table 2: qPCR primers**

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| **Primer Name** | **5’-Sequence-3’** | **Reference** |
| tRNA-K-CTT-qF | CTAGCTCAGTCGGTAGAGCATGA | (Aldrich and Maggert, 2014; Paredes and Maggert, 2009b) |
| tRNA-K-CTT-qR | CCAACGTGGGGCTCGAAC |
| 18S-qF | AGCCTGAGAAACGGCTACCA |
| 18S-qR | AGCTGGGAGTGGGTAATTTACG |
| 28S-qF | AATGGATGTGATGCCAATGTA | (Greil and Ahmad, 2012) |
| 28S-qR | TTCAGTGGATCGCAGTATGG |
| 5.8S-qF | GCTCATGGGTCGATGAAGAA | This study |
| 5.8S-qR | GGACTGCGATATGCGTTCA |
| Gapdh-qF | TAAATTCGACTCGACTCACGGT | DRSC FlyPrimerBank |
| Gapdh-qR | CTCCACCACATACTCGGCTC |
| R1 qPCR2 fw | TAGAGCTTGTAGTGGTCGAG | This study |
| R1 qPCR2 rv | ATGGGTCGTCGGCATGATCT |
| R2 qPCR fw | ATGAACTGTTATCGCGTCCG |
| R2 qPCR rv | AAACGCGTGGGTTGACTAAC |