**Table S3**. Positive controls used for each primer pair. Concentration corresponds to the final concentration in the PCR mix. Lineage corresponds to *Plasmodium* mtDNA cytochrome *b* lineage [(Bensch et al., 2009)](https://www.zotero.org/google-docs/?6ZdU0Z).

|  |  |  |
| --- | --- | --- |
|  | Species | Concentration (ng/µl) |
| C+ *Culi01* | *Aedes koreicus* | 2 |
| *Aedes japonicus* | 2 |
| C+ *Aves02\_mix* | *Psittacus erithacus* | 0.148 |
| *Neopsephotus bourkii* | 0.148 |
| *Quelea quelea* | 0.148 |
| *Urocolius indicus* | 0.148 |
| C+ *Plas01* | Lineage AFR046 | NA |
| Lineage AEDVEX01 | NA |