**Table S2** Usage of initial and termination codons in five *Cyrtodactylus* species

|  |  |  |  |  |  |  |  |  |  |  |
| --- | --- | --- | --- | --- | --- | --- | --- | --- | --- | --- |
| Species/Gene | *Cyrtodactylus peguensis* | | *Cyrtodactylus thirakhupti* | | *Cyrtodactylus auribalteatus* | | *Cyrtodactylus chanhomeae* | | *Cyrtodactylus tigroides* | |
| Start | Stop | Start | Stop | Start | Stop | Start | Stop | Start | Stop |
| *ND1* | ATG | TAG | ATG | AGA | GTG | TAA | GTG | TAA | GTG | TAA |
| *ND2* | ATA | TAA | ATA | TAA | ATA | TAG | ATA | TAG | ATA | TAA |
| *COI* | ATG | AGG | ATG | AGA | ATG | AGA | ATG | AGA | ATG | TAA |
| *COII* | ATG | T++1 | ATG | T++1 | ATG | T++1 | GTG | TAA | ATG | T++1 |
| *ATPase8* | ATG | TAA | ATG | TAA | GTG | TAA | ATG | TAA | ATG | TAA |
| *ATPase6* | ATG | TAA | ATG | TAA | ATG | TAA | ATG | TAA | ATG | TAA |
| *COIII* | ATG | T++1 | ATG | TA+1 | ATG | TA+1 | ATG | TA+1 | ATG | T++1 |
| *ND3* | ATA | TAG | ATA | TA | ATG | TA+1 | ATA | TA+1 | ATG | TAG |
| *ND4L* | ATG | TAA | ATG | TAA | ATG | TAA | ATG | TAA | ATG | TAA |
| *ND4* | ATG | T++1 | ATG | T++1 | ATG | T++1 | ATG | TAG | ATG | T++1 |
| *ND5* | ATG | TAG | ATA | TAG | ATG | TAA | ATG | TAA | ATA | TAA |
| *ND6* | ATG | TAG | ATG | AGG | ATG | TAG | ATG | TAG | ATG | TAG |
| *CYTB* | ATG | TAG | ATG | TAG | ATG | TA+1 | ATG | TAA | ATG | TAG |

1TAA stop codon is completed by the addition of 3'A residues to mRNA