**Table S9.** Base composition and skewness of mitogenome of *Elaeidobius kamerunicus*.

|  |  |  |  |  |
| --- | --- | --- | --- | --- |
| Feature | Length | A+T% | AT-skew | GC-skew |
| Whole genome | 17729 | 73.5 | 0.085 | -0.224 |
| PCGs | 11112 | 72.1 | -0.13 | -0.065 |
| tRNAs | 1458 | 77.1 | 0.02 | 0.108 |
| rRNAs | 2043 | 77.7 | -0.075 | 0.316 |
| Control region | 2588 | 75.8 | 0.102 | -0.329 |