
Deep Fixed Freeze V4 20 020 0598 Full

November 17, 2021 – Twenty-six morphometric measurements used in this study are used in the molecular phylogenetic analysis shown in Table 4. This analysis allows estimation of genome length in standard units. For each sample, the genomes identified in this study were added together to estimate relative length. Analysis of the data presented in Table 4 shows that all of these samples had relatively short genomes, averaging about 60 million nucleotides. These results are consistent with other studies showing that genome sizes are shrinking rapidly and that there is significant genetic diversity.

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