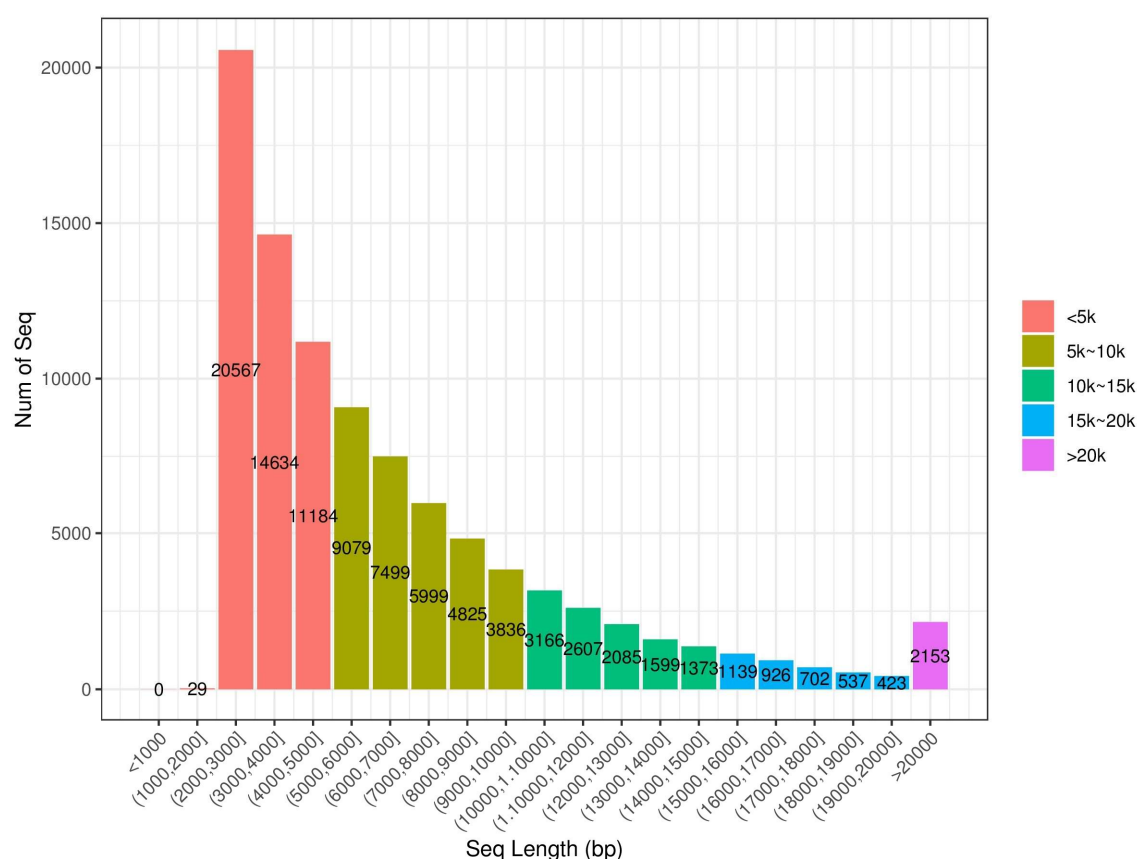


# Genomic Analysis of *Sphingopyxis* sp. USTB-05 for Biodegrading Cyanobacterial Hepatotoxins

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**Figure S1.** The length of the three generations of nanopore data is distributed in turn.



**Figure S2.** Comparison of genes related to hepatotoxin biodegradation in the genomes of *Sphingomonas morindae* sp. NBD5 and *Sphingopyxis* sp. USTB-05. The gray box in the figure indicate that the gene exists in two strains, and the white indicate that the gene does not exist in two strains.