

Supplementary Figures

Kyphosid fish microbiome adaptations to sulfated dietary polysaccharides

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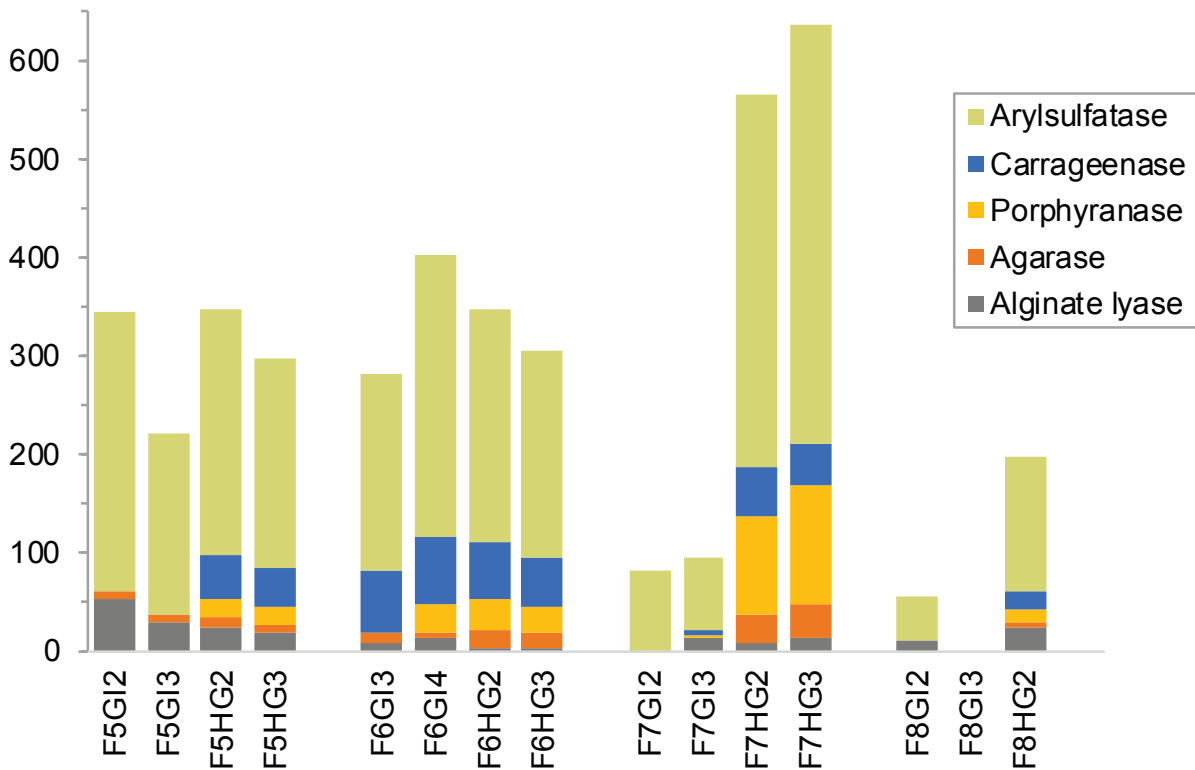
Supplementary Figure S1. Fish gut sample distribution of algal degradative enzyme keywords.

Prokka annotation keywords for assembled contigs have been normalized for total number of predicted proteins in each fish sample. Full data and significance calculations are provided in Supplementary Dataset S1

Supplementary Figure S2. Separation distances for CAZy and SulfAtlas class pairs.

A) fish metagenomes. B) Terrestrial ruminant metagenomes. Complete data is presented in Supplementary Dataset 3.

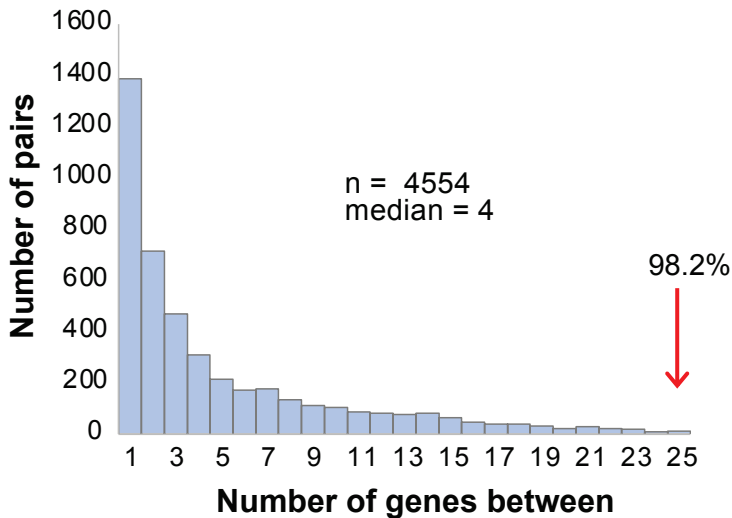
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Supplementary Figure S2. Separation distances for CAZy and SulfAtlas class pairs.

A) fish metagenomes. B) Terrestrial ruminant metagenomes. Complete data is presented in **Supplementary Dataset 3**.

A



B

