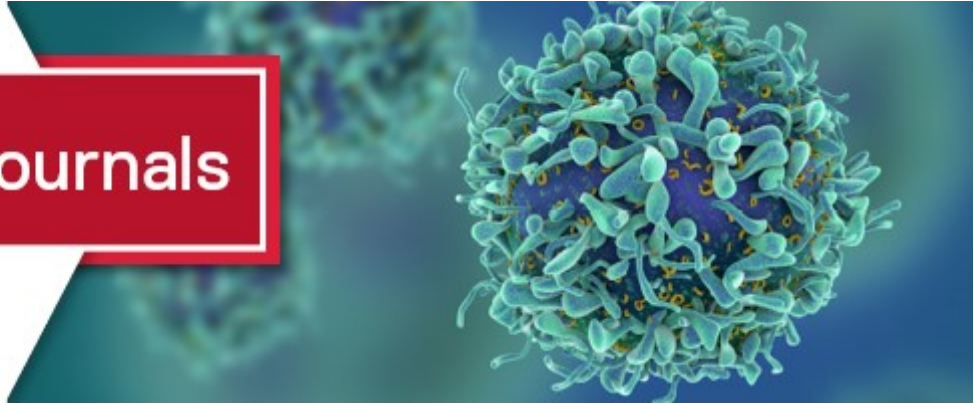
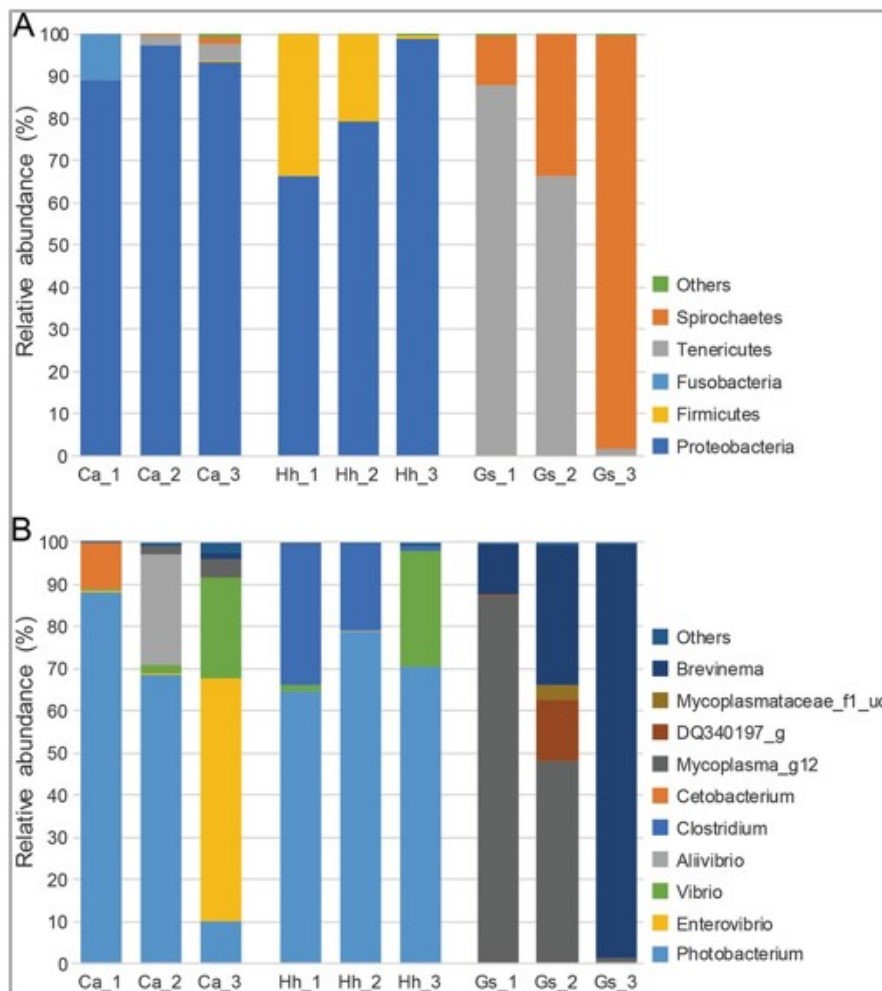


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**FIG 1.** Bar chart representing the taxonomic composition of the gut microbiota in deep-sea fish, based on 16S rRNA gene amplicon sequence analysis. (*16S rRNA Gene Amplicon Sequencing of Gut Microbiota in Three Species of Deep-Sea Fish in Suruga Bay, Japan*)

- **16S rRNA Gene Amplicon Sequencing of Gut Microbiota in Three Species of Deep-Sea Fish in Suruga Bay, Japan**
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