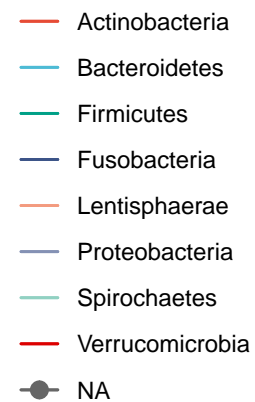


*bsh* gene



phylum



Proportion of MCBAs detected among tested

