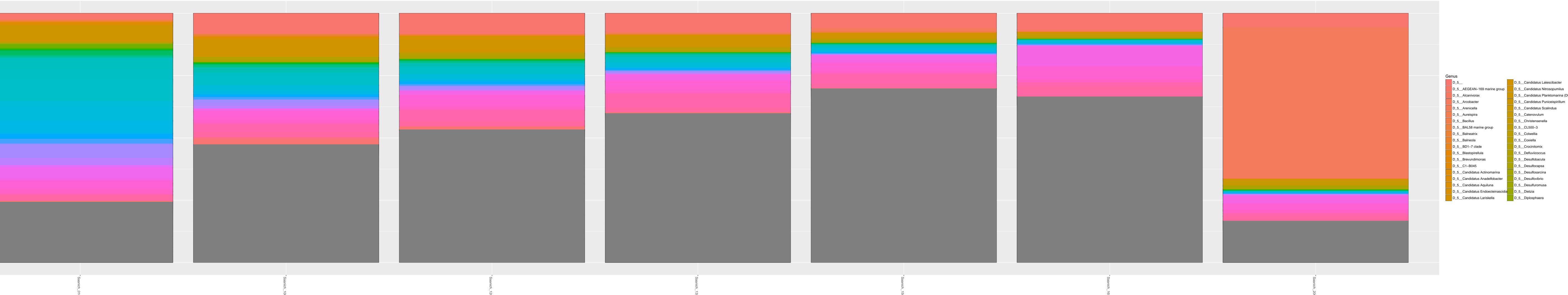
Phylum Abundance Through Saanich Inlet Depth



## D\_5\_\_Rhodobium D\_5\_Sulfurospirillum D\_5\_uncultured Flavobacteriia bacterium D\_5\_AEGEAN-169 marine group D\_5\_Candidatus Nitrosopumilus D\_5\_uncultured Flavobacterium sp. D\_5\_Candidatus Planktomarina (DC5-80-3 lineage) D\_5\_Erythrobacter D\_5\_Nitrosococcus D\_5\_Pelagicoccus D\_5\_Rhodopirellula D\_5\_Sulfurovum D\_5\_uncultured gamma proteobacterium D\_5\_Candidatus Puniceispirillum D\_5\_\_Filomicrobium D\_5\_Nitrosomonas D\_5\_Peredibacter D\_5\_\_Rickettsia D\_5\_Synechococcus D\_5\_Nitrospina D\_5\_Persicirhabdus D\_5\_Candidatus Scalindua D\_5\_\_Flammeovirga D\_5\_\_Robiginitalea D\_5\_\_Tenacibaculum D\_5\_uncultured marine archaeon D\_5\_uncultured marine bacterium D\_5\_\_Fluviicola D\_5\_\_Nocardioides D\_5\_\_Photobacterium D\_5\_\_Roseibacillus D\_5\_Christensenella D\_5\_Formosa D\_5\_NS2b marine group D\_5\_Phycisphaera D\_5\_Roseobacter clade NAC11-7 lineage D\_5\_Thiobacillus D\_5\_uncultured marine microorganism D\_5\_Francisella D\_5\_NS3a marine group D\_5\_Pir4 lineage D\_5\_Roseobacter clade OCT lineage D\_5\_Thiorhodospira D\_5\_uncultured Marinimicrobia bacterium D\_5\_Magnetococcus D\_5\_FS140-16B-02 marine group D\_5\_Magnetospira D\_5\_uncultured Planctomycetes Sva0503 D\_5\_\_Rubritalea D\_5\_NS4 marine group D\_5\_Planctomyces D\_5\_\_Fusibacter D\_5\_Rubrivirga D\_5\_uncultured Polyangiaceae bacterium D\_5\_NS5 marine group D\_5\_Polaribacter D\_5\_Glaciecola D\_5\_SAR92 clade D\_5\_Turneriella D\_5\_uncultured proteobacterium D\_5\_marine group D\_5\_Octadecabacter D\_5\_Porticoccus D\_5\_Halioglobus D\_5\_\_marine metagenome D\_5\_Ulvibacter D\_5\_uncultured SAR86 cluster bacterium D\_5\_uncultured D\_5\_unidentified marine bacterioplanktor D\_5\_Halomonas D\_5\_Marinicella D\_5\_Proteobacteria bacterium JGI 0000113-D06 D\_5\_SEEP-SRB1 D\_5\_\_Desulfocapsa D\_5\_Hellea D\_5\_Marinifilum D\_5\_OM27 clade D\_5\_Pseudoalteromonas D\_5\_Shewanella D\_5\_uncultured actinobacterium D\_5\_\_JL-ETNP-F27 D\_5\_\_Victivallis D\_5\_Marinimicrobia bacterium JGI OTU-2 D\_5\_OM43 clade D\_5\_Pseudomonas D\_5\_Simiduia D\_5\_uncultured alpha proteobacterium D\_5\_ZD0417 marine group D\_5\_Candidatus Anadelfobacter D\_5\_Desulfovibrio D\_5\_Kangiella D\_5\_uncultured archaeon D\_5\_\_Marinobacterium D\_5\_OM60(NOR5) clade D\_5\_Pseudospirillum D\_5\_Sneathiella D\_5\_\_Legionella D\_5\_OM75 clade D\_5\_Psychrilyobacter D\_5\_uncultured bacterium D\_5\_Marinoscillum D\_5\_Sphingobium

D\_5\_Opitutus D\_5\_Psychromonas

D\_5\_Owenweeksia D\_5\_Reichenbachiella

D\_5\_Leisingera

D\_5\_Lentibacter

D\_5\_Methylobacterium

D\_5\_Methylotenera

D\_5\_Sulfurimonas

D\_5\_uncultured Bacteroidetes bacterium

D\_5\_uncultured Bacteroidetes/Chlorobi group bacterium

D\_5\_Spirochaeta

D\_5\_uncultured delta proteobacterium