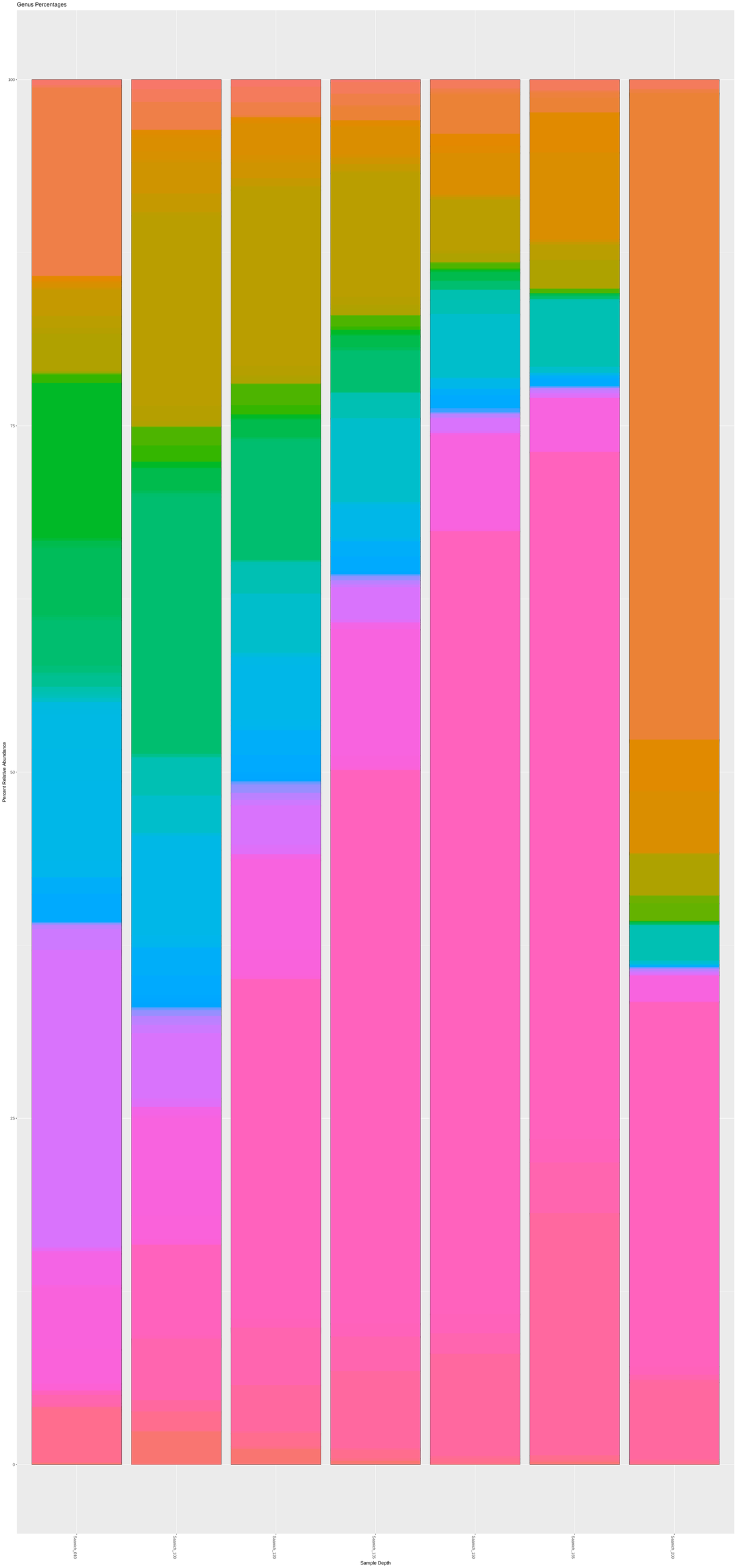
Genus Percentages



Genus

- Acidimicrobiales_unclassified
- AEGEAN~169_marine_group
- AEGEAN~245_ge
- Alcanivorax
- Alphaproteobacteria_unclassified
- Alteromonadaceae_ge
- Alteromonadaceae_unclassified
- Alteromonadales_unclassified
- Amylibacter
- Anaerolineaceae_unclassified
- Archaea_unclassified
- Arcobacter
- Arctic97B~4_marine_group_ge
- AT~s3~44_ge
- Aurantimonas
- Aureispira
- Bacillales_unclassified
- Bacilli_unclassified
- Bacteria_unclassified
- Bacteriovoracaceae_unclassified
- Bacteroidetes_BD2~2_ge
- Bacteroidetes_unclassified
- Bacteroidetes_VC2.1_Bac22_ge
- BAL58_marine_group
- Balneatrix
- Balneola
- BD2~11_terrestrial_group_ge
- BD7~11_ge
- Betaproteobacteria_unclassified
- Blri41_ge
- Blfdi19_ge
- Bradymonadales_ge
- Campylobacterales_unclassified
- Candidatus_Actinomarina
- Candidatus_Aquiluna
- Candidatus_Endoecteinascidia
- Candidatus_Magasanikbacteria_ge
- Candidatus_Nitrosopelagicus
- Candidatus_Nitrosopumilus
- Candidatus_Pacebacteria_ge
- Candidatus_Pelagibacter
- Candidatus_Peribacteria_ge
- Candidatus_Puniceispirillum
- Candidatus_Scalindua
- Candidatus_Tenderia
- Candidatus_Thioglobus
- Candidatus_Uhrbacteria_ge
- Celdivibrionaceae_unclassified
- Celdivibrionales_unclassified
- Chesapeake~Delaware_Bay_ge
- Chlamydiales_unclassified
- Chloroplast_ge
- CK~1C2~67_ge
- CL500~3
- Clostridiales_unclassified
- Clostridiales_unclassified
- Coxiella
- Crocinitomix
- Cryomorphaceae_unclassified
- CrystalBog021C3_ge
- Cytophagales_unclassified
- Deltaproteobacteria_unclassified
- Desulfatiglans
- Desulfobacteraceae_unclassified
- Desulfobacula
- Desulfovibrionales_unclassified
- Diplosphaera
- Draconibacteriaceae_unclassified
- E01~9C~26_marine_group_ge
- Eel~36e1D6_ge
- EF100~94H03_ge
- Emcibacter
- Erythrobacter
- Erythrobacteraceae_unclassified
- Erythrobacteraceae_unclassified
- Flammeovirgaceae_unclassified
- Flammeovirgaceae_unclassified
- Flavicella
- Flavobacteriaceae_unclassified
- Flavobacteriales_unclassified
- Fluviicola
- Formosa
- Frigidibacter
- FS140~16B~02_marine_group
- Gammaproteobacteria_unclassified
- Glaciecola
- Gracilibacteria_ge
- Haliaeaceae_unclassified
- Helilea
- Hoeflea
- Ilumatobacter
- JL~ETNP~F27
- JL~ETNP~Y6_ge
- JTB255_marine_benthic_group_ge
- Kangiella
- Kiloo9A_clade_ge
- Latescibacteria_ge
- Lentimonas
- Lentisphaera
- Lentisphaerae_unclassified
- Litoreibacter
- Litoricola
- Luminiphilus
- LWSR~14_ge
- Magnetococcus
- Magnetospira
- Maricaulis
- Maricurvus
- marine_group
- Marine_Group_I_ge
- Marine_Group_I_unclassified
- Marine_Group_II_ge
- Marine_Methylotrophic_Group_1_ge
- Marinimicrobia_(SAR406_clade)_ge
- Marinobacterium
- Marinoscillum
- MB11C04_marine_group_ge
- MD2902~B12_ge
- MD3~55_ge
- Methylococcales_unclassified
- Methylotenera
- Microbacteriaceae_unclassified
- Milano~WF1B~03_ge
- ML635J~21_ge
- MSBL7
- MSBL8_ge
- MVP~94_ge
- Neorhizobium
- NitSAr0613~695_ge
- Nitrospina
- Nocardioides
- NS11~12_marine_group_ge
- NS2b_marine_group
- NS3a_marine_group
- NS4_marine_group
- NS5_marine_group
- NS7_marine_group_ge
- NS9_marine_group_ge
- Oceanospirillales_unclassified
- Octadecabacter
- ODP1230B30.02_sediment_group_ge
- Oleiphilus
- Oleispira
- Oligoflexaceae_ge
- OM1_clade_ge
- OM182_clade_ge
- OM190_ge
- OM27_clade
- OM43_clade
- OM60(NOR5)_clade
- OM75_clade
- OPB35_soil_group_ge
- Opitutae_unclassified
- Owenweeksia
- P3OB~42_ge
- Paenibacillaceae_unclassified
- Paraglaciecola
- Parcubacteria_ge
- Parcubacteria_unclassified
- PAUC34f_ge
- Pelagibacterium
- Pelolinea
- PeM15_ge
- Peredibacter
- Persicirhabdus
- Piscirickettsiaceae_unclassified
- Pla3_lineage_ge
- Planctomyces
- Planctomycetaceae_unclassified
- Planctomycetes_unclassified
- Planococcaceae_unclassified
- Polaribacter
- Polaribacter_4
- Porticoccus
- possible_genus_03
- Proteobacteria_unclassified
- PS1_clade_ge
- Pseudoalteromonadaceae_unclassified
- Pseudoalteromonas
- Pseudofulvibacter
- Pseudohongiella
- Psychromonas
- Puniceicoccaceae_unclassified
- Rhizobiales_unclassified
- Rhodobacteraceae_unclassified
- Rhodobium
- Rhodospirillaceae_unclassified
- Rhodospirillales_unclassified
- Rickettsiaceae_unclassified
- Rickettsiales_unclassified
- Roseibacillus
- Rubritalea
- S25~593_ge
- Salinimonas
- Saprospiraceae_unclassified
- SAR11_clade_ge
- SAR11_clade_unclassified
- SAR116_clade_ge
- SAR116_clade_unclassified
- SAR202_clade_ge
- SAR324_clade(Marine_group_B)_ge
- SAR86_clade_ge
- SAR92_clade
- SBR1093_ge
- SHWN~night2_ge
- Sphingobacteriales_unclassified
- Sphingomonadales_unclassified
- Spirochaeta_2
- Spongibacteraceae_unclassified
- SPOTSOCT00m83_ge
- Subgroup_6_unclassified
- Sulfurimonas
- Sulfurospirillum
- SUP05_cluster_ge
- SUP05_cluster_unclassified
- Surface_1_unclassified
- Surface_3_ge
- Surface_4_ge
- Sva0996_marine_group_ge
- Synechococcus
- Tllo_ge
- Tenacibaculum
- Thermomarinilinea
- Thiorhodospira
- Thiotrichales_unclassified
- Tistrella
- TM6_(Dependentiae)_ge
- Ulvibacter
- uncultured
- uncultured_ge
- unknown_unclassified
- Verrucomicrobia_unclassified
- Vibrio
- Vibrionaceae_unclassified
- WCHB1~41_ge
- Woesearchaeota_(DHVEG~6)_ge
- ZD0417_marine_group