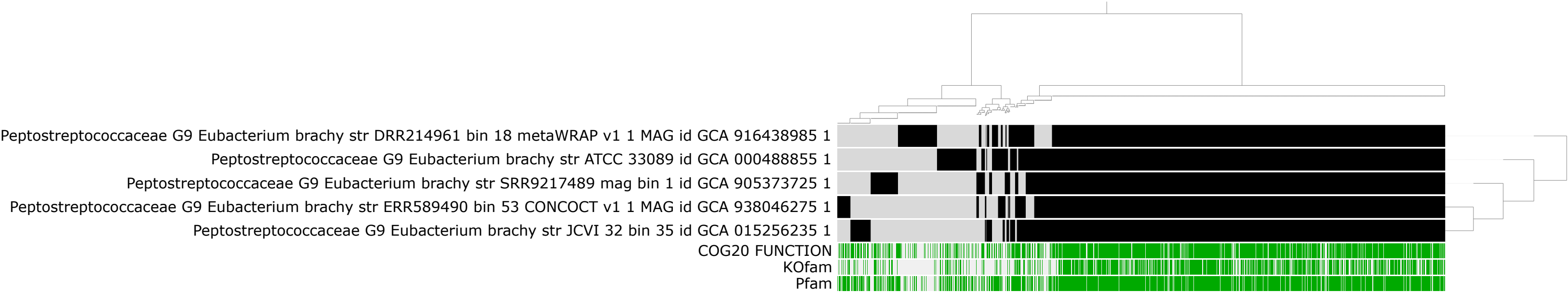
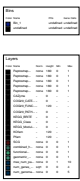


Peptostreptococcaceae G9 Eubacterium brachy HMT 557

Items order: Presence absence (D: Euclidean; L: Ward) | Current view: gene_cluster_presence_absence | Samples order: gene_cluster frequencies



CAZyme

☐ UNKNOWN (1700) ☒ KNOWN (13)

COG20 CATEGORY

☒ KNOWN (1228) ☐ UNKNOWN (485)

COG20 FUNCTION

☒ KNOWN (1228) ☐ UNKNOWN (485)

COG20 PATHWAY

☐ UNKNOWN (1319) ☒ KNOWN (394)

KEGG BRITE

☒ KNOWN (895) ☐ UNKNOWN (818)

KEGG Class

☐ UNKNOWN (1552) ☒ KNOWN (161)

KEGG Module

☐ UNKNOWN (1552) ☒ KNOWN (161)

KOfam

☒ KNOWN (899) ☐ UNKNOWN (814)

Pfam

☒ KNOWN (1328) ☐ UNKNOWN (385)