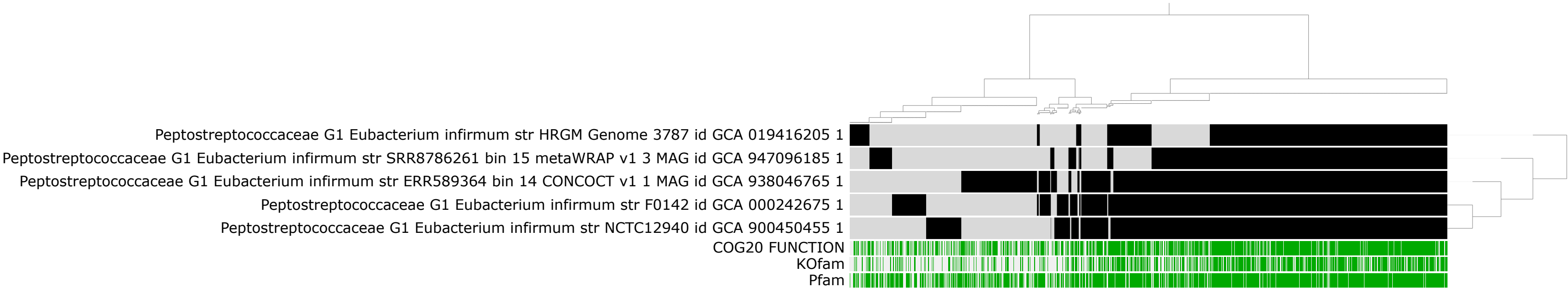
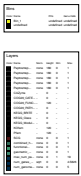


Peptostreptococcaceae G1 Eubacterium infirmum HMT 105

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



CAZyme

☐ UNKNOWN (2304) ☒ KNOWN (26)

COG20 CATEGORY

☒ KNOWN (1675) ☐ UNKNOWN (655)

COG20 FUNCTION

☒ KNOWN (1676) ☐ UNKNOWN (654)

COG20 PATHWAY

☐ UNKNOWN (1872) ☒ KNOWN (458)

KEGG BRITE

☒ KNOWN (1171) ☐ UNKNOWN (1159)

KEGG Class

☐ UNKNOWN (2113) ☒ KNOWN (217)

KEGG Module

☐ UNKNOWN (2113) ☒ KNOWN (217)

KOfam

☒ KNOWN (1172) ☐ UNKNOWN (1158)

Pfam

☒ KNOWN (1831) ☐ UNKNOWN (499)