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__ʁacteroidia;_o__Bacteroidales;_f__Bacteroidales_S24–7_group;_g__norank_f__Bacteroidales_S24–7_group;_s__uncultured_Porphyromonadaceae
   p_Bacteroidetes; c_Bacteroidia;_o_Bacteroidales;_f_Prevotellaceae;_g_Prevotella_9;_s_uncultured_bacterium_g_Prevotella_9
   Bacteroidetes, Asacteroidia; o_Bacteroidales; f_Prevotellaceae; g_Prevotella_9; s_uncultured_bacterium_g_Prevotella_9
   p__Bacteroidetes;_c__Bacteroidates;_f__Prevotellaceae;_g__Prevotellaceae_NK3B31_group;_s__uncultured_bacterium_g__Prevotellaceae_NK3B31_group;_s__uncultured_bacterium_g__Prevotellaceae_NK3B31_group;_s__uncultured_bacterium_g__Prevotellaceae_NK3B31_group;_s__uncultured_bacterium_g__Prevotellaceae_NK3B31_group;_s__uncultured_bacterium_g__Prevotellaceae_NK3B31_group;_s__uncultured_bacterium_g__Prevotellaceae_NK3B31_group;_s__uncultured_bacterium_g__Prevotellaceae_NK3B31_group;_s__uncultured_bacterium_g__Prevotellaceae_NK3B31_group;_s__uncultured_bacterium_g__Prevotellaceae_NK3B31_group;_s__uncultured_bacterium_g__Prevotellaceae_NK3B31_group;_s__uncultured_bacterium_g__Prevotellaceae_NK3B31_group;_s__uncultured_bacterium_g__Prevotellaceae_NK3B31_group;_s__uncultured_bacterium_g__Prevotellaceae_NK3B31_group;_s__uncultured_bacterium_g__Prevotellaceae_NK3B31_group;_s__uncultured_bacterium_g__Prevotellaceae_NK3B31_group;_s__uncultured_bacterium_g__Prevotellaceae_NK3B31_group;_s__uncultured_bacterium_g__Prevotellaceae_NK3B31_group;_s__uncultured_bacterium_g__Prevotellaceae_NK3B31_group;_s__uncultured_bacterium_g__Prevotellaceae_NK3B31_group;_s__uncultured_bacterium_g__Prevotellaceae_NK3B31_group;_s__uncultured_bacterium_g__Prevotellaceae_NK3B31_group;_s__uncultured_bacterium_g__Prevotellaceae_NK3B31_group;_s__uncultured_bacterium_g__Prevotellaceae_NK3B31_group;_s__uncultured_bacterium_g__Prevotellaceae_NK3B31_group;_s__uncultured_bacterium_g__Prevotellaceae_NK3B31_group;_s__uncultured_bacterium_g__Prevotellaceae_NK3B31_group;_s__uncultured_bacterium_g__Prevotellaceae_NK3B31_group;_s__uncultured_bacterium_g__Prevotellaceae_NK3B31_group;_s__uncultured_bacterium_g__Prevotellaceae_NK3B31_group_g__Prevotellaceae_NK3B31_g__Prevotellaceae_NK3B31_g__Prevotellaceae_NK3B31_g__Prevotellaceae_NK3B31_g__Prevotellaceae_NK3B31_g__Prevotellaceae_NK3B31_g__Prevotellaceae_NK3B31_g__Prevotellaceae_NK3B31_g__Prevotellaceae_NK3B31_g__Prevotellaceae_NK3B31_g__Prevotellaceae_NK3B31_g__Prevotellaceae_NK3B31_g__Prevotellaceae_NK3B31_g__Prevotel
   p_Bacteroidetes; c___Bacteroidia; o__Bacteroidales;_f__Prevotellaceae;_g__Prevotellaceae_NK3B31_group;_s__uncultured_bacterium_g__Prevotellaceae_NK3B31_group;_s__uncultured_bacterium_g__Prevotellaceae_NK3B31_group;_s__uncultured_bacterium_g__Prevotellaceae_NK3B31_group;_s__uncultured_bacterium_g__Prevotellaceae_NK3B31_group;_s__uncultured_bacterium_g__Prevotellaceae_NK3B31_group;_s__uncultured_bacterium_g__Prevotellaceae_NK3B31_group;_s__uncultured_bacterium_g__Prevotellaceae_NK3B31_group;_s__uncultured_bacterium_g__Prevotellaceae_NK3B31_group;_s__uncultured_bacterium_g__Prevotellaceae_NK3B31_group;_s__uncultured_bacterium_g__Prevotellaceae_NK3B31_group;_s__uncultured_bacterium_g__Prevotellaceae_NK3B31_group;_s__uncultured_bacterium_g__Prevotellaceae_NK3B31_group;_s__uncultured_bacterium_g__Prevotellaceae_NK3B31_group;_s__uncultured_bacterium_g__Prevotellaceae_NK3B31_group;_s__uncultured_bacterium_g__Prevotellaceae_NK3B31_group;_s__uncultured_bacterium_g__Prevotellaceae_NK3B31_group;_s__uncultured_bacterium_g__Prevotellaceae_NK3B31_group;_s__uncultured_bacterium_g__Prevotellaceae_NK3B31_group;_s__uncultured_bacterium_g__Prevotellaceae_NK3B31_group;_s__uncultured_bacterium_g__Prevotellaceae_NK3B31_group;_s__uncultured_bacterium_g__Prevotellaceae_NK3B31_group;_s__uncultured_bacterium_g__Prevotellaceae_NK3B31_group;_s__uncultured_bacterium_g__Prevotellaceae_NK3B31_group;_s__uncultured_bacterium_g__Prevotellaceae_NK3B31_group;_s__uncultured_bacterium_g__Prevotellaceae_NK3B31_group;_s__uncultured_bacterium_g__Prevotellaceae_NK3B31_group;_s__uncultured_bacterium_g__Prevotellaceae_NK3B31_group;_s__uncultured_bacterium_g__Prevotellaceae_NK3B31_group_g__Prevotellaceae_NK3B31_group_g__Prevotellaceae_NK3B31_g__Prevotellaceae_NK3B31_g__Prevotellaceae_NK3B31_g__Prevotellaceae_NK3B31_g__Prevotellaceae_NK3B31_g__Prevotellaceae_NK3B31_g__Prevotellaceae_NK3B31_g__Prevotellaceae_NK3B31_g__Prevotellaceae_NK3B31_g__Prevotellaceae_NK3B31_g__Prevotellaceae_NK3B31_g__Prevotellaceae_NK3B31_g__Prevotellaceae_NK3B31_g__Prevotellaceae_NK
   p_Firmicutes; c_Negativicutes; o_Selenomonadales; f_Veillonellaceae; g_Megasphaera; s_Megasphaera_elsdenii_DSM_20460
   p_Firmicutes;_c_Clostridia;_to_Clostridiales;_f_Clostridiaceae_1;_g_Clostridium_sensu_stricto_1;_s_uncultured_bacterium_g_Clostridium_sensu_stricto_1
p_Firmicutes;_c_Bacilli;_o_Lactobacillaes;_f_Lactobacillaceae;_g_Lactobacillus;_s_Lactobacillus_amylovorus
   p_Firmicutes;_c_Negativicutes;_o_Selenomonadales;_f_Acidaminococcaceae;_g_Phascolarctobacterium;_s_unclassified_g_Phascolarctobacterium
   p_Firmicutes;_c_Clostridia_p_Costridiales;_f_Lachnospiraceae;_g_[Eubacterium]_rectale_group;_s_uncultured_organism_g_[Eubacterium]_rectale_group
  p_Firmicutes;_c_Clostri<mark>dia_p</mark>s_Otostridiales;_f_Ruminococcaceae;_g_Ruminococcaceae_UCG-002;_s_uncultured_bacterium_g_Ruminococcaceae_UCG-002
p_Firmicutes;_c_Negaţivicutes;_&_Selenomonadales;_f_Veillonellaceae;_g_norank_f_Veillonellaceae;_s_uncultured_rumen_bacterium_g_norank_f_Veillonellace
          Firmicutes; c_Negativicutes; o Selenomonadales; f_Veillonellaceae; g_Selenomonas; s_Selenomonas_bovis
p_Firmicutes;_c_Clostri<mark>dia; sò:_Clo</mark>stridiales;_f_Ruminococcaceae;_g_Faecalibacterium;_s_unclassified_g_Faecalibacterium
  p_Firmicutes; c_Baciti; o_Lactobacillales; f_Streptococcaceae; g_Streptococcus; s_Streptococcus_gallolyticus_subsp._macedonicus
   Firmicutes; c Wegatvicutes, o Selenomonadales; f Veillonellaceae; g Megasphaera; s unclassified g Megasphaera
p__Firmicutes;_c__pao@s_o__lactobacillales;_f__Lactobacillaceae;_g__Lactobacillus;_s__Lactobacillus_johnsonii
   p__Firmicutes;____Bacilli;_o__Lactobacillales;_f__Lactobacillaceae;_g__Lactobacillus;_s__Lactobacillus_reuteri
          _Proteobacteria_c_c.__$ammaproteobacteria;_o__Enterobacteriales;_f__Enterobacteriaceae;_g__Escherichia–Shigella;_s__Escherichia_coli_g__Escherichia–Shigella
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