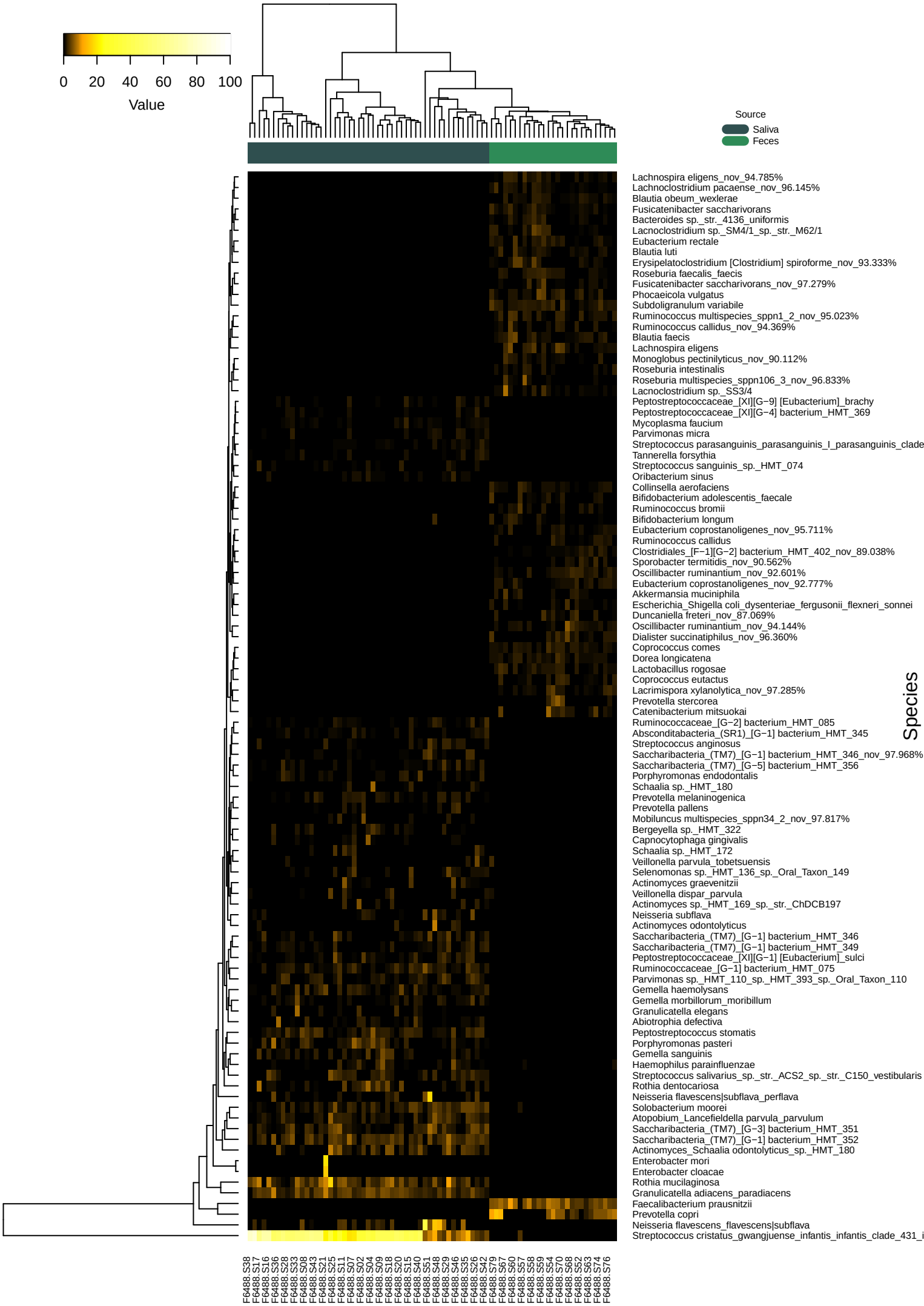
0
20
40
60
80
100
Value
Source
Saliva
Feces
Lachnospira eligens_nov_94.785%
Lachnoclostridium pacaense_nov_96.145%
Blautia obeum_wexlerae
Fusicatenibacter saccharivorans
Bacteroides sp._str._4136_uniformis
Lacnoclostridium sp._SM4/1_sp._str._M62/1
Eubacterium rectale
Blautia luti
Erysipelatoclostridium [Clostridium] spiroforme_nov_93.333%
Roseburia faecalis_faecis
Fusicatenibacter saccharivorans_nov_97.279%
Phocaeicola vulgatus
Subdoligranulum variabile
Ruminococcus multispecies_sppn1_2_nov_95.023%
Ruminococcus callidus_nov_94.369%
Blautia faecis
Lachnospira eligens
Monoglobus pectinilyticus_nov_90.112%
Roseburia intestinalis
Roseburia multispecies_sppn106_3_nov_96.833%
Lacnoclostridium sp._SS3/4
Peptostreptococcaceae_[XI][G−9] [Eubacterium]_brachy
Peptostreptococcaceae_[XI][G−4] bacterium_HMT_369
Mycoplasma faucium
Parvimonas micra
Streptococcus parasanguinis_parasanguinis_I_parasanguinis_clade
Tannerella forsythia
Streptococcus sanguinis_sp._HMT_074
Oribacterium sinus
Collinsella aerofaciens
Bifidobacterium adolescentis_faecale
Ruminococcus bromii
Bifidobacterium longum
Eubacterium coprostanoligenes_nov_95.711%
Ruminococcus callidus
Clostridiales_[F−1][G−2] bacterium_HMT_402_nov_89.038%
Sporobacter termitidis_nov_90.562%
Oscillibacter ruminantium_nov_92.601%
Eubacterium coprostanoligenes_nov_92.777%
Akkermansia muciniphila
Escherichia_Shigella coli_dysenteriae_fergusonii_flexneri_sonnei
Duncaniella freteri_nov_87.069%
Oscillibacter ruminantium_nov_94.144%
Dialister succinatiphilus_nov_96.360%
Coprococcus comes
Dorea longicatena
Lactobacillus rogosae
Coprococcus eutactus
Lacrimispora xylanolytica_nov_97.285%
Prevotella stercorea
Catenibacterium mitsuokai
Ruminococcaceae_[G−2] bacterium_HMT_085
Absconditabacteria_(SR1)_[G−1] bacterium_HMT_345
Streptococcus anginosus
Saccharibacteria_(TM7)_[G−1] bacterium_HMT_346_nov_97.968%
Saccharibacteria_(TM7)_[G−5] bacterium_HMT_356
Porphyromonas endodontalis
Schaalia sp._HMT_180
Prevotella melaninogenica
Prevotella pallens
Mobiluncus multispecies_sppn34_2_nov_97.817%
Bergeyella sp._HMT_322
Capnocytophaga gingivalis
Schaalia sp._HMT_172
Veillonella parvula_tobetsuensis
Selenomonas sp._HMT_136_sp._Oral_Taxon_149
Actinomyces graevenitzii
Veillonella dispar_parvula
Actinomyces sp._HMT_169_sp._str._ChDCB197
Neisseria subflava
Actinomyces odontolyticus
Saccharibacteria_(TM7)_[G−1] bacterium_HMT_346
Saccharibacteria_(TM7)_[G−1] bacterium_HMT_349
Peptostreptococcaceae_[XI][G−1] [Eubacterium]_sulci
Ruminococcaceae_[G−1] bacterium_HMT_075
Parvimonas sp._HMT_110_sp._HMT_393_sp._Oral_Taxon_110
Gemella haemolysans
Gemella morbillorum_moribillum
Granulicatella elegans
Abiotrophia defectiva
Peptostreptococcus stomatis
Porphyromonas pasteri
Gemella sanguinis
Haemophilus parainfluenzae
Streptococcus salivarius_sp._str._ACS2_sp._str._C150_vestibularis
Rothia dentocariosa
Neisseria flavescens|subflava_perflava
Solobacterium moorei
Atopobium_Lancefieldella parvula_parvulum
Saccharibacteria_(TM7)_[G−3] bacterium_HMT_351
Saccharibacteria_(TM7)_[G−1] bacterium_HMT_352
Actinomyces_Schaalia odontolyticus_sp._HMT_180
Enterobacter mori
Enterobacter cloacae
Rothia mucilaginosa
Granulicatella adiacens_paradiacens
Faecalibacterium prausnitzii
Prevotella copri
Neisseria flavescens_flavescens|subflava
Streptococcus cristatus_gwangjuense_infantis_infantis_clade_431_i
Species
F6488.S38
F6488.S17
F6488.S16
F6488.S36
F6488.S28
F6488.S33
F6488.S08
F6488.S43
F6488.S21
F6488.S25
F6488.S11
F6488.S07
F6488.S02
F6488.S04
F6488.S09
F6488.S18
F6488.S20
F6488.S15
F6488.S40
F6488.S51
F6488.S48
F6488.S29
F6488.S46
F6488.S35
F6488.S26
F6488.S42
F6488.S79
F6488.S67
F6488.S60
F6488.S57
F6488.S58
F6488.S59
F6488.S54
F6488.S70
F6488.S68
F6488.S52
F6488.S63
F6488.S74
F6488.S76
Samples