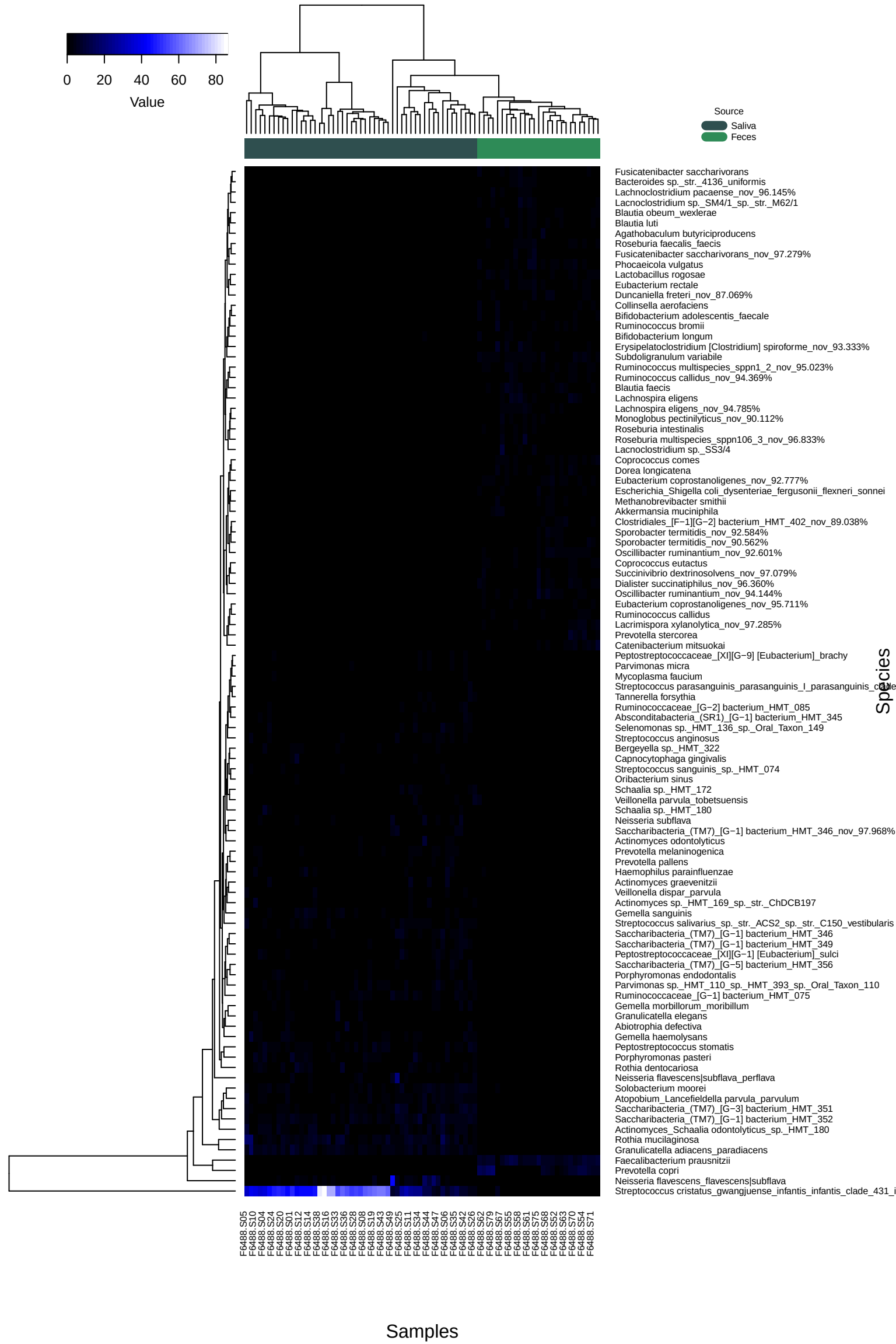
0
20
40
60
80
Value
Source
Saliva
Feces
Fusicatenibacter saccharivorans
Bacteroides sp._str._4136_uniformis
Lachnoclostridium pacaense_nov_96.145%
Lacnoclostridium sp._SM4/1_sp._str._M62/1
Blautia obeum_wexlerae
Blautia luti
Agathobaculum butyriciproducens
Roseburia faecalis_faecis
Fusicatenibacter saccharivorans_nov_97.279%
Phocaeicola vulgatus
Lactobacillus rogosae
Eubacterium rectale
Duncaniella freteri_nov_87.069%
Collinsella aerofaciens
Bifidobacterium adolescentis_faecale
Ruminococcus bromii
Bifidobacterium longum
Erysipelatoclostridium [Clostridium] spiroforme_nov_93.333%
Subdoligranulum variabile
Ruminococcus multispecies_sppn1_2_nov_95.023%
Ruminococcus callidus_nov_94.369%
Blautia faecis
Lachnospira eligens
Lachnospira eligens_nov_94.785%
Monoglobus pectinilyticus_nov_90.112%
Roseburia intestinalis
Roseburia multispecies_sppn106_3_nov_96.833%
Lacnoclostridium sp._SS3/4
Coprococcus comes
Dorea longicatena
Eubacterium coprostanoligenes_nov_92.777%
Escherichia_Shigella coli_dysenteriae_fergusonii_flexneri_sonnei
Methanobrevibacter smithii
Akkermansia muciniphila
Clostridiales_[F−1][G−2] bacterium_HMT_402_nov_89.038%
Sporobacter termitidis_nov_92.584%
Sporobacter termitidis_nov_90.562%
Oscillibacter ruminantium_nov_92.601%
Coprococcus eutactus
Succinivibrio dextrinosolvens_nov_97.079%
Dialister succinatiphilus_nov_96.360%
Oscillibacter ruminantium_nov_94.144%
Eubacterium coprostanoligenes_nov_95.711%
Ruminococcus callidus
Lacrimispora xylanolytica_nov_97.285%
Prevotella stercorea
Catenibacterium mitsuokai
Peptostreptococcaceae_[XI][G−9] [Eubacterium]_brachy
Parvimonas micra
Mycoplasma faucium
Streptococcus parasanguinis_parasanguinis_I_parasanguinis_clade
Tannerella forsythia
Ruminococcaceae_[G−2] bacterium_HMT_085
Absconditabacteria_(SR1)_[G−1] bacterium_HMT_345
Selenomonas sp._HMT_136_sp._Oral_Taxon_149
Streptococcus anginosus
Bergeyella sp._HMT_322
Capnocytophaga gingivalis
Streptococcus sanguinis_sp._HMT_074
Oribacterium sinus
Schaalia sp._HMT_172
Veillonella parvula_tobetsuensis
Schaalia sp._HMT_180
Neisseria subflava
Saccharibacteria_(TM7)_[G−1] bacterium_HMT_346_nov_97.968%
Actinomyces odontolyticus
Prevotella melaninogenica
Prevotella pallens
Haemophilus parainfluenzae
Actinomyces graevenitzii
Veillonella dispar_parvula
Actinomyces sp._HMT_169_sp._str._ChDCB197
Gemella sanguinis
Streptococcus salivarius_sp._str._ACS2_sp._str._C150_vestibularis
Saccharibacteria_(TM7)_[G−1] bacterium_HMT_346
Saccharibacteria_(TM7)_[G−1] bacterium_HMT_349
Peptostreptococcaceae_[XI][G−1] [Eubacterium]_sulci
Saccharibacteria_(TM7)_[G−5] bacterium_HMT_356
Porphyromonas endodontalis
Parvimonas sp._HMT_110_sp._HMT_393_sp._Oral_Taxon_110
Ruminococcaceae_[G−1] bacterium_HMT_075
Gemella morbillorum_moribillum
Granulicatella elegans
Abiotrophia defectiva
Gemella haemolysans
Peptostreptococcus stomatis
Porphyromonas pasteri
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
Rothia mucilaginosa
Granulicatella adiacens_paradiacens
Faecalibacterium prausnitzii
Prevotella copri
Neisseria flavescens_flavescens|subflava
Streptococcus cristatus_gwangjuense_infantis_infantis_clade_431_i
Species
F6488.S05
F6488.S10
F6488.S04
F6488.S24
F6488.S20
F6488.S01
F6488.S12
F6488.S14
F6488.S38
F6488.S16
F6488.S33
F6488.S36
F6488.S28
F6488.S08
F6488.S19
F6488.S43
F6488.S49
F6488.S25
F6488.S11
F6488.S34
F6488.S44
F6488.S47
F6488.S06
F6488.S35
F6488.S42
F6488.S26
F6488.S62
F6488.S79
F6488.S67
F6488.S55
F6488.S58
F6488.S61
F6488.S75
F6488.S68
F6488.S52
F6488.S63
F6488.S70
F6488.S54
F6488.S71
Samples