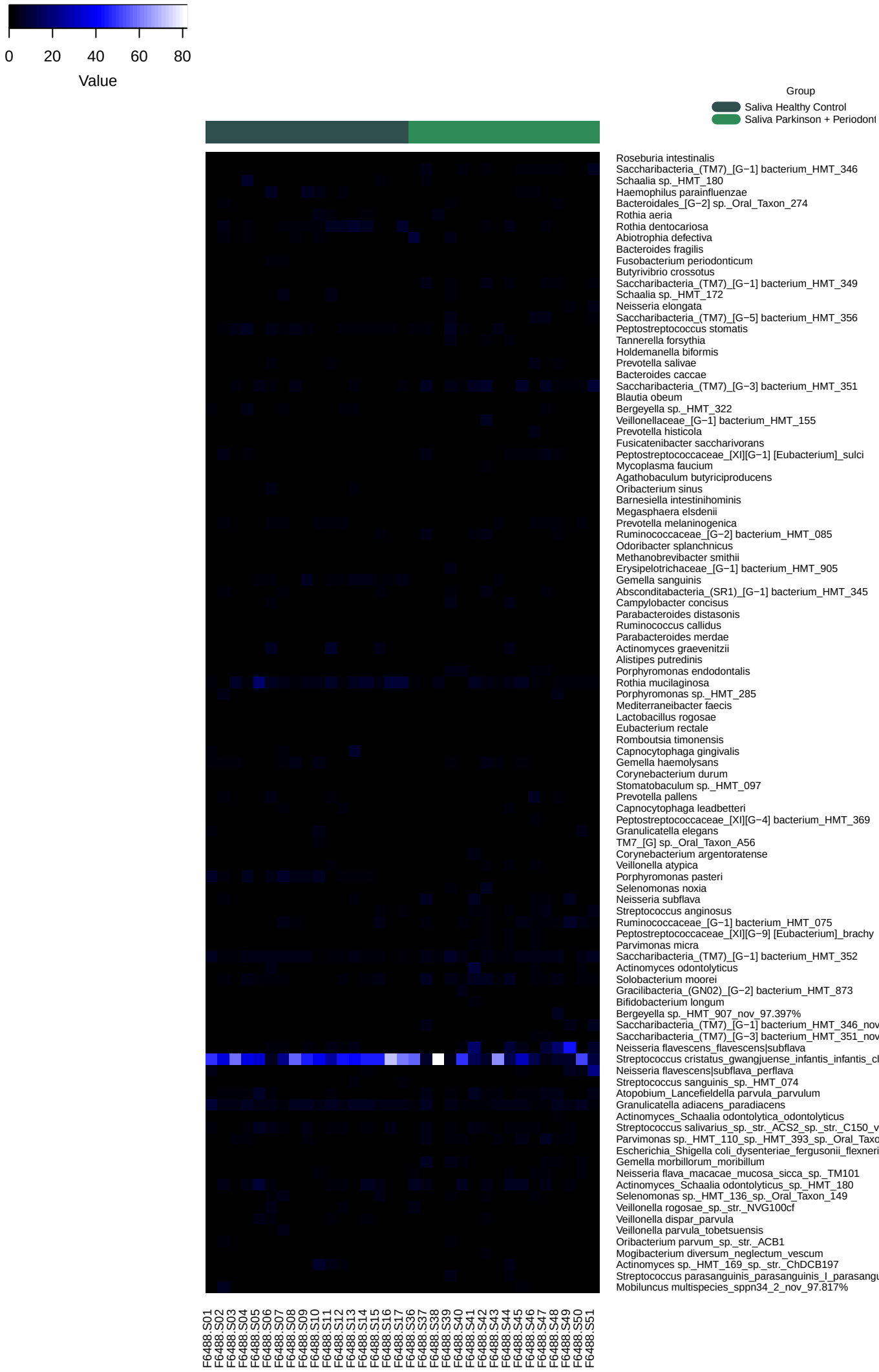
0
20
40
60
80
Value
Group
Saliva Healthy Control
Saliva Parkinson + Periodont
Roseburia intestinalis
Saccharibacteria_(TM7)_[G−1] bacterium_HMT_346
Schaalia sp._HMT_180
Haemophilus parainfluenzae
Bacteroidales_[G−2] sp._Oral_Taxon_274
Rothia aeria
Rothia dentocariosa
Abiotrophia defectiva
Bacteroides fragilis
Fusobacterium periodonticum
Butyrivibrio crossotus
Saccharibacteria_(TM7)_[G−1] bacterium_HMT_349
Schaalia sp._HMT_172
Neisseria elongata
Saccharibacteria_(TM7)_[G−5] bacterium_HMT_356
Peptostreptococcus stomatis
Tannerella forsythia
Holdemanella biformis
Prevotella salivae
Bacteroides caccae
Saccharibacteria_(TM7)_[G−3] bacterium_HMT_351
Blautia obeum
Bergeyella sp._HMT_322
Veillonellaceae_[G−1] bacterium_HMT_155
Prevotella histicola
Fusicatenibacter saccharivorans
Peptostreptococcaceae_[XI][G−1] [Eubacterium]_sulci
Mycoplasma faucium
Agathobaculum butyriciproducens
Oribacterium sinus
Barnesiella intestinihominis
Megasphaera elsdenii
Prevotella melaninogenica
Ruminococcaceae_[G−2] bacterium_HMT_085
Odoribacter splanchnicus
Methanobrevibacter smithii
Erysipelotrichaceae_[G−1] bacterium_HMT_905
Gemella sanguinis
Absconditabacteria_(SR1)_[G−1] bacterium_HMT_345
Campylobacter concisus
Parabacteroides distasonis
Ruminococcus callidus
Parabacteroides merdae
Actinomyces graevenitzii
Alistipes putredinis
Porphyromonas endodontalis
Rothia mucilaginosa
Porphyromonas sp._HMT_285
Mediterraneibacter faecis
Lactobacillus rogosae
Eubacterium rectale
Romboutsia timonensis
Capnocytophaga gingivalis
Gemella haemolysans
Corynebacterium durum
Stomatobaculum sp._HMT_097
Prevotella pallens
Capnocytophaga leadbetteri
Peptostreptococcaceae_[XI][G−4] bacterium_HMT_369
Granulicatella elegans
TM7_[G] sp._Oral_Taxon_A56
Corynebacterium argentoratense
Veillonella atypica
Porphyromonas pasteri
Selenomonas noxia
Neisseria subflava
Streptococcus anginosus
Ruminococcaceae_[G−1] bacterium_HMT_075
Peptostreptococcaceae_[XI][G−9] [Eubacterium]_brachy
Parvimonas micra
Saccharibacteria_(TM7)_[G−1] bacterium_HMT_352
Actinomyces odontolyticus
Solobacterium moorei
Gracilibacteria_(GN02)_[G−2] bacterium_HMT_873
Bifidobacterium longum
Bergeyella sp._HMT_907_nov_97.397%
Saccharibacteria_(TM7)_[G−1] bacterium_HMT_346_nov_97.968%
Saccharibacteria_(TM7)_[G−3] bacterium_HMT_351_nov_97.072%
Neisseria flavescens_flavescens|subflava
Streptococcus cristatus_gwangjuense_infantis_infantis_clade_431_i
Neisseria flavescens|subflava_perflava
Streptococcus sanguinis_sp._HMT_074
Atopobium_Lancefieldella parvula_parvulum
Granulicatella adiacens_paradiacens
Actinomyces_Schaalia odontolytica_odontolyticus
Streptococcus salivarius_sp._str._ACS2_sp._str._C150_vestibularis
Parvimonas sp._HMT_110_sp._HMT_393_sp._Oral_Taxon_110
Escherichia_Shigella coli_dysenteriae_fergusonii_flexneri_sonnei
Gemella morbillorum_moribillum
Neisseria flava_macacae_mucosa_sicca_sp._TM101
Actinomyces_Schaalia odontolyticus_sp._HMT_180
Selenomonas sp._HMT_136_sp._Oral_Taxon_149
Veillonella rogosae_sp._str._NVG100cf
Veillonella dispar_parvula
Veillonella parvula_tobetsuensis
Oribacterium parvum_sp._str._ACB1
Mogibacterium diversum_neglectum_vescum
Actinomyces sp._HMT_169_sp._str._ChDCB197
Streptococcus parasanguinis_parasanguinis_I_parasanguinis_clade
Mobiluncus multispecies_sppn34_2_nov_97.817%
Species
F6488.S01
F6488.S02
F6488.S03
F6488.S04
F6488.S05
F6488.S06
F6488.S07
F6488.S08
F6488.S09
F6488.S10
F6488.S11
F6488.S12
F6488.S13
F6488.S14
F6488.S15
F6488.S16
F6488.S17
F6488.S36
F6488.S37
F6488.S38
F6488.S39
F6488.S40
F6488.S41
F6488.S42
F6488.S43
F6488.S44
F6488.S45
F6488.S46
F6488.S47
F6488.S48
F6488.S49
F6488.S50
F6488.S51
Samples