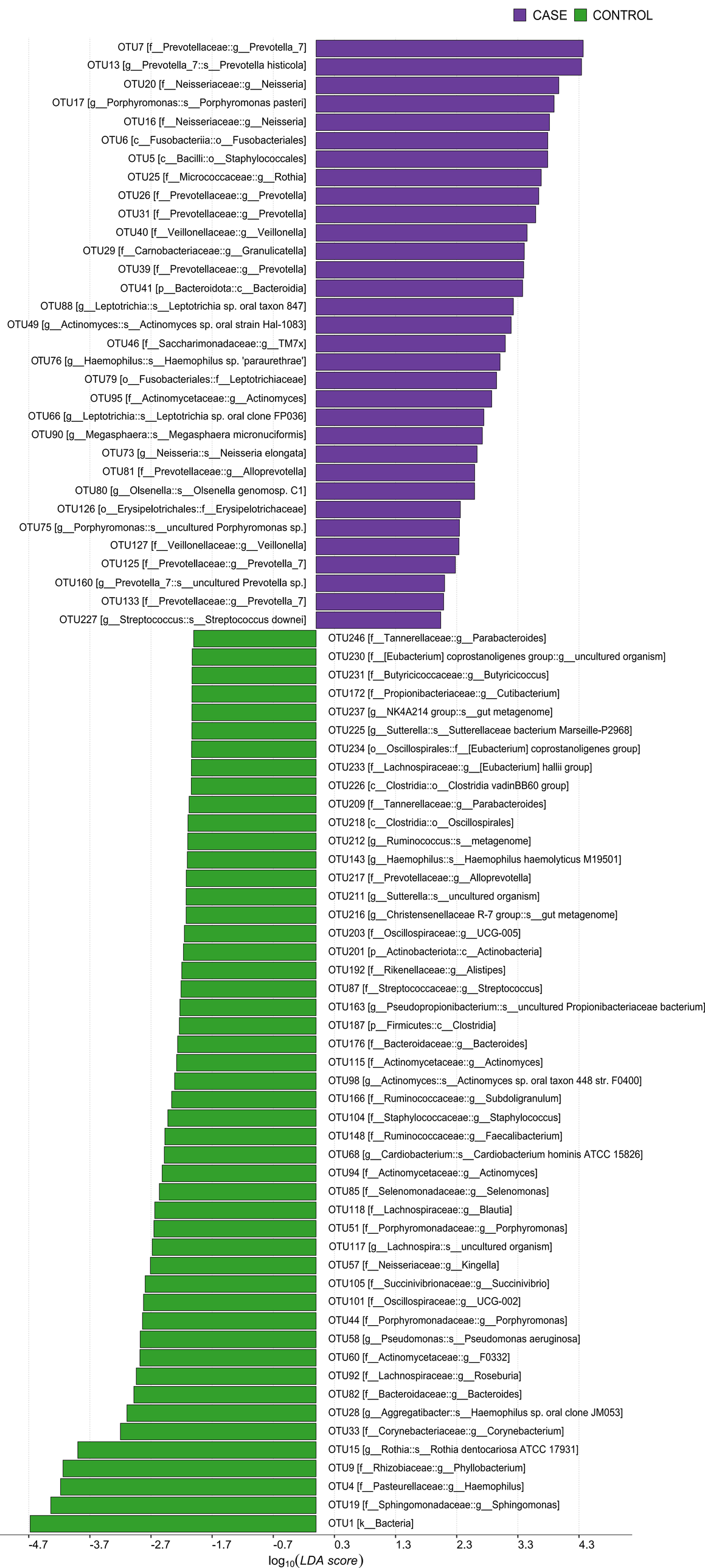
CASE
CONTROL
OTU7 [f__Prevotellaceae::g__Prevotella_7]
OTU13 [g__Prevotella_7::s__Prevotella histicola]
OTU20 [f__Neisseriaceae::g__Neisseria]
OTU17 [g__Porphyromonas::s__Porphyromonas pasteri]
OTU16 [f__Neisseriaceae::g__Neisseria]
OTU6 [c__Fusobacteriia::o__Fusobacteriales]
OTU5 [c__Bacilli::o__Staphylococcales]
OTU25 [f__Micrococcaceae::g__Rothia]
OTU26 [f__Prevotellaceae::g__Prevotella]
OTU31 [f__Prevotellaceae::g__Prevotella]
OTU40 [f__Veillonellaceae::g__Veillonella]
OTU29 [f__Carnobacteriaceae::g__Granulicatella]
OTU39 [f__Prevotellaceae::g__Prevotella]
OTU41 [p__Bacteroidota::c__Bacteroidia]
OTU88 [g__Leptotrichia::s__Leptotrichia sp. oral taxon 847]
OTU49 [g__Actinomyces::s__Actinomyces sp. oral strain Hal-1083]
OTU46 [f__Saccharimonadaceae::g__TM7x]
OTU76 [g__Haemophilus::s__Haemophilus sp. 'paraurethrae']
OTU79 [o__Fusobacteriales::f__Leptotrichiaceae]
OTU95 [f__Actinomycetaceae::g__Actinomyces]
OTU66 [g__Leptotrichia::s__Leptotrichia sp. oral clone FP036]
OTU90 [g__Megasphaera::s__Megasphaera micronuciformis]
OTU73 [g__Neisseria::s__Neisseria elongata]
OTU81 [f__Prevotellaceae::g__Alloprevotella]
OTU80 [g__Olsenella::s__Olsenella genomosp. C1]
OTU126 [o__Erysipelotrichales::f__Erysipelotrichaceae]
OTU75 [g__Porphyromonas::s__uncultured Porphyromonas sp.]
OTU127 [f__Veillonellaceae::g__Veillonella]
OTU125 [f__Prevotellaceae::g__Prevotella_7]
OTU160 [g__Prevotella_7::s__uncultured Prevotella sp.]
OTU133 [f__Prevotellaceae::g__Prevotella_7]
OTU227 [g__Streptococcus::s__Streptococcus downei]
OTU246 [f__Tannerellaceae::g__Parabacteroides]
OTU230 [f__[Eubacterium] coprostanoligenes group::g__uncultured organism]
OTU231 [f__Butyricicoccaceae::g__Butyricicoccus]
OTU172 [f__Propionibacteriaceae::g__Cutibacterium]
OTU237 [g__NK4A214 group::s__gut metagenome]
OTU225 [g__Sutterella::s__Sutterellaceae bacterium Marseille-P2968]
OTU234 [o__Oscillospirales::f__[Eubacterium] coprostanoligenes group]
OTU233 [f__Lachnospiraceae::g__[Eubacterium] hallii group]
OTU226 [c__Clostridia::o__Clostridia vadinBB60 group]
OTU209 [f__Tannerellaceae::g__Parabacteroides]
OTU218 [c__Clostridia::o__Oscillospirales]
OTU212 [g__Ruminococcus::s__metagenome]
OTU143 [g__Haemophilus::s__Haemophilus haemolyticus M19501]
OTU217 [f__Prevotellaceae::g__Alloprevotella]
OTU211 [g__Sutterella::s__uncultured organism]
OTU216 [g__Christensenellaceae R-7 group::s__gut metagenome]
OTU203 [f__Oscillospiraceae::g__UCG-005]
OTU201 [p__Actinobacteriota::c__Actinobacteria]
OTU192 [f__Rikenellaceae::g__Alistipes]
OTU87 [f__Streptococcaceae::g__Streptococcus]
OTU163 [g__Pseudopropionibacterium::s__uncultured Propionibacteriaceae bacterium]
OTU187 [p__Firmicutes::c__Clostridia]
OTU176 [f__Bacteroidaceae::g__Bacteroides]
OTU115 [f__Actinomycetaceae::g__Actinomyces]
OTU98 [g__Actinomyces::s__Actinomyces sp. oral taxon 448 str. F0400]
OTU166 [f__Ruminococcaceae::g__Subdoligranulum]
OTU104 [f__Staphylococcaceae::g__Staphylococcus]
OTU148 [f__Ruminococcaceae::g__Faecalibacterium]
OTU68 [g__Cardiobacterium::s__Cardiobacterium hominis ATCC 15826]
OTU94 [f__Actinomycetaceae::g__Actinomyces]
OTU85 [f__Selenomonadaceae::g__Selenomonas]
OTU118 [f__Lachnospiraceae::g__Blautia]
OTU51 [f__Porphyromonadaceae::g__Porphyromonas]
OTU117 [g__Lachnospira::s__uncultured organism]
OTU57 [f__Neisseriaceae::g__Kingella]
OTU105 [f__Succinivibrionaceae::g__Succinivibrio]
OTU101 [f__Oscillospiraceae::g__UCG-002]
OTU44 [f__Porphyromonadaceae::g__Porphyromonas]
OTU58 [g__Pseudomonas::s__Pseudomonas aeruginosa]
OTU60 [f__Actinomycetaceae::g__F0332]
OTU92 [f__Lachnospiraceae::g__Roseburia]
OTU82 [f__Bacteroidaceae::g__Bacteroides]
OTU28 [g__Aggregatibacter::s__Haemophilus sp. oral clone JM053]
OTU33 [f__Corynebacteriaceae::g__Corynebacterium]
OTU15 [g__Rothia::s__Rothia dentocariosa ATCC 17931]
OTU9 [f__Rhizobiaceae::g__Phyllobacterium]
OTU4 [f__Pasteurellaceae::g__Haemophilus]
OTU19 [f__Sphingomonadaceae::g__Sphingomonas]
OTU1 [k__Bacteria]
-4.7
-3.7
-2.7
-1.7
-0.7
0.3
1.3
2.3
3.3
4.3
log10(LDA score)