

D_0__Bacteria;D_1__Proteobacteria;D_2__Gammaproteobacteria;D_3__Enterobacteriales;D_4__E
D_0__Bacteria;D_1__Actinobacteria;D_2__Coriobacteriia;D_3__Coriobacteriales;D_4__Coriobacte
D_0__Bacteria;D_1__Firmicutes;D_2__Clostridia;D_3__Clostridiales;D_4__Peptococcaceae;D_5__C
D_0__Bacteria;D_1__Firmicutes;D_2__Clostridia;D_3__Clostridiales;D_4__Lachnospiraceae;D_5__L
D_0__Bacteria;D_1__Firmicutes;D_2__Clostridia;D_3__Thermoanaerobacterales;D_4__Thermoana
D_0__Bacteria;D_1__Tenericutes;D_2__Mollicutes;D_3__Mollicutes RF9;D_4__uncultured bacteri
D_0__Bacteria;D_1__Actinobacteria;D_2__Coriobacteriia;D_3__Coriobacteriales;D_4__Coriobacte
D_0__Bacteria;D_1__Firmicutes;D_2__Clostridia;D_3__Clostridiales;D_4__Ruminococcaceae;D_5__C
D_0__Bacteria;D_1__Bacteroidetes;D_2__Flavobacteriia;D_3__Flavobacteriales;D_4__Flavobacteri
D_0__Bacteria;D_1__Firmicutes;D_2__Erysipelotrichia;D_3__Erysipelotrichales;D_4__Erysipelotric
D_0__Bacteria;D_1__Firmicutes;D_2__Clostridia;D_3__Clostridiales;D_4__Ruminococcaceae;D_5__C
D_0__Bacteria;D_1__Actinobacteria;D_2__Coriobacteriia;D_3__Coriobacteriales;D_4__Coriobacte
D_0__Bacteria;D_1__Firmicutes;D_2__Clostridia;D_3__Clostridiales;D_4__Lachnospiraceae;D_5__L
D_0__Bacteria;D_1__Firmicutes;D_2__Clostridia;D_3__Clostridiales;D_4__Lachnospiraceae;D_5__L
D_0__Bacteria;D_1__Fusobacteria;D_2__Fusobacteriia;D_3__Fusobacteriales;D_4__Fusobacteriac
D_0__Bacteria;D_1__Tenericutes;D_2__Mollicutes;D_3__Mollicutes RF9;Ambiguous_taxa;Ambigu
D_0__Bacteria;D_1__Actinobacteria;D_2__Coriobacteriia;D_3__Coriobacteriales;D_4__Coriobacte
D_0__Bacteria;D_1__Firmicutes;D_2__Clostridia;D_3__Clostridiales;D_4__Ruminococcaceae;D_5__C
D_0__Bacteria;D_1__Firmicutes;D_2__Clostridia;D_3__Clostridiales;D_4__Lachnospiraceae;D_5__L
D_0__Bacteria;D_1__Firmicutes;D_2__Clostridia;D_3__Clostridiales;D_4__Lachnospiraceae;D_5__L
D_0__Bacteria;D_1__Firmicutes;D_2__Clostridia;D_3__Clostridiales;D_4__Peptostreptococcaceae;
D_0__Bacteria;D_1__Bacteroidetes;D_2__Bacteroidia;D_3__Bacteroidales;D_4__Porphyromonada
D_0__Bacteria;D_1__Bacteroidetes;D_2__Bacteroidia;D_3__Bacteroidales;D_4__Porphyromonada
D_0__Bacteria;D_1__Bacteroidetes;D_2__Bacteroidia;D_3__Bacteroidales;D_4__Rikenellaceae;D_4
D_0__Bacteria;D_1__Bacteroidetes;D_2__Bacteroidia;D_3__Bacteroidales;D_4__Bacteroidales S24
D_0__Bacteria;D_1__Firmicutes;D_2__Bacilli;D_3__Lactobacillales;D_4__Enterococcaceae;D_5__E
D_0__Bacteria;D_1__Firmicutes;D_2__Clostridia;D_3__Clostridiales;D_4__Lachnospiraceae;D_5__L
D_0__Bacteria;D_1__Firmicutes;D_2__Clostridia;D_3__Clostridiales;D_4__Lachnospiraceae;D_5__L
D_0__Bacteria;D_1__Proteobacteria;D_2__Gammaproteobacteria;D_3__Aeromonadales;D_4__Su
D_0__Bacteria;D_1__Proteobacteria;D_2__Gammaproteobacteria;D_3__Enterobacteriales;D_4__E
D_0__Bacteria;D_1__Proteobacteria;D_2__Gammaproteobacteria;D_3__Enterobacteriales;D_4__E
D_0__Bacteria;D_1__Firmicutes;D_2__Clostridia;D_3__Clostridiales;D_4__Lachnospiraceae;D_5__L
D_0__Bacteria;D_1__Firmicutes;D_2__Clostridia;D_3__Clostridiales;D_4__Clostridiaceae 1;D_5__C

median_Heal	median_MS	mean_Health	mean_MS	MannW_pval	MannW_pvalue_fdr
0.00000	0.00000	0.00000	0.00037	0.00000	0.00013
0.00000	0.00000	0.00014	0.00000	0.00001	0.00134
0.00000	0.00000	0.00010	0.00001	0.00004	0.00279
0.00000	0.00000	0.00000	0.00054	0.00013	0.00626
0.00000	0.00000	0.00007	0.00000	0.00022	0.00837
0.00000	0.00000	0.00012	0.00023	0.00048	0.01324
0.00000	0.00000	0.00024	0.00000	0.00048	0.01324
0.00000	0.00000	0.00014	0.00000	0.00090	0.02128
0.00000	0.00000	0.00000	0.00007	0.00102	0.02128
0.00028	0.00009	0.00046	0.00020	0.00110	0.02128
0.07829	0.05122	0.09379	0.05925	0.00141	0.02228
0.00000	0.00000	0.00007	0.00000	0.00149	0.02228
0.00056	0.00028	0.00087	0.00044	0.00156	0.02228
0.00000	0.00000	0.00000	0.00001	0.00185	0.02228
0.00000	0.00000	0.00000	0.00060	0.00185	0.02228
0.00000	0.00000	0.00000	0.00005	0.00185	0.02228
0.00000	0.00000	0.00006	0.00001	0.00744	0.08452
0.00000	0.00000	0.00017	0.00000	0.00867	0.09295
0.00028	0.00009	0.00054	0.00028	0.01063	0.10361
0.00074	0.00042	0.00124	0.00063	0.01074	0.10361
0.00000	0.00000	0.00005	0.00001	0.02184	0.17777
0.00000	0.00000	0.00063	0.00017	0.02256	0.17777
0.00241	0.00501	0.00396	0.00607	0.02399	0.17777
0.00000	0.00000	0.00364	0.00135	0.02735	0.17777
0.00000	0.00000	0.00000	0.00017	0.02855	0.17777
0.00000	0.00000	0.00000	0.00000	0.02855	0.17777
0.00000	0.00000	0.00000	0.00000	0.02855	0.17777
0.00000	0.00000	0.00000	0.00000	0.02855	0.17777
0.00000	0.00000	0.00000	0.00042	0.02855	0.17777
0.00000	0.00000	0.00000	0.00011	0.02855	0.17777
0.00000	0.00000	0.00000	0.00013	0.02855	0.17777
0.00000	0.00009	0.00010	0.00016	0.04521	0.27267
0.00028	0.00005	0.00244	0.00162	0.04900	0.28655