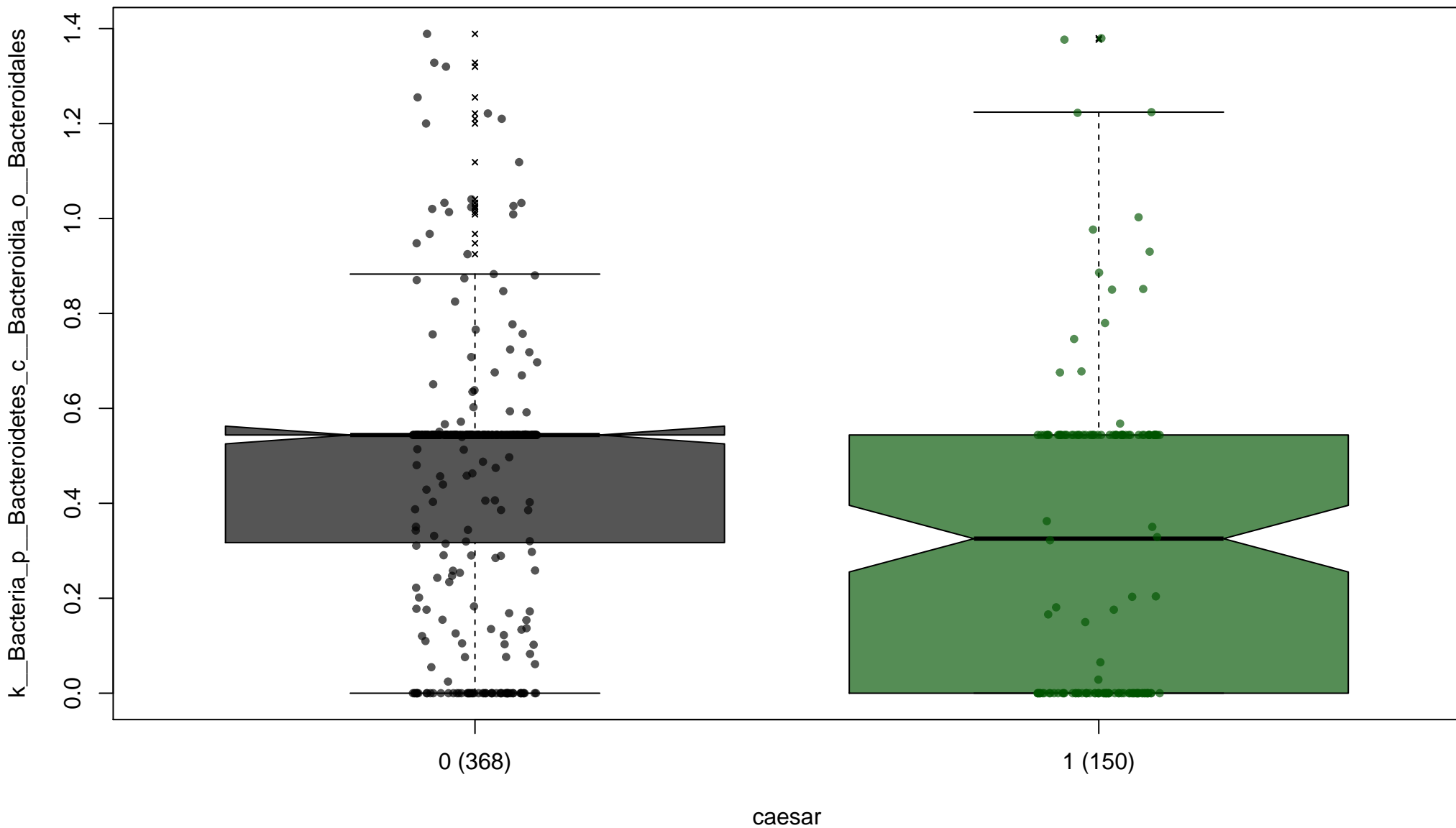
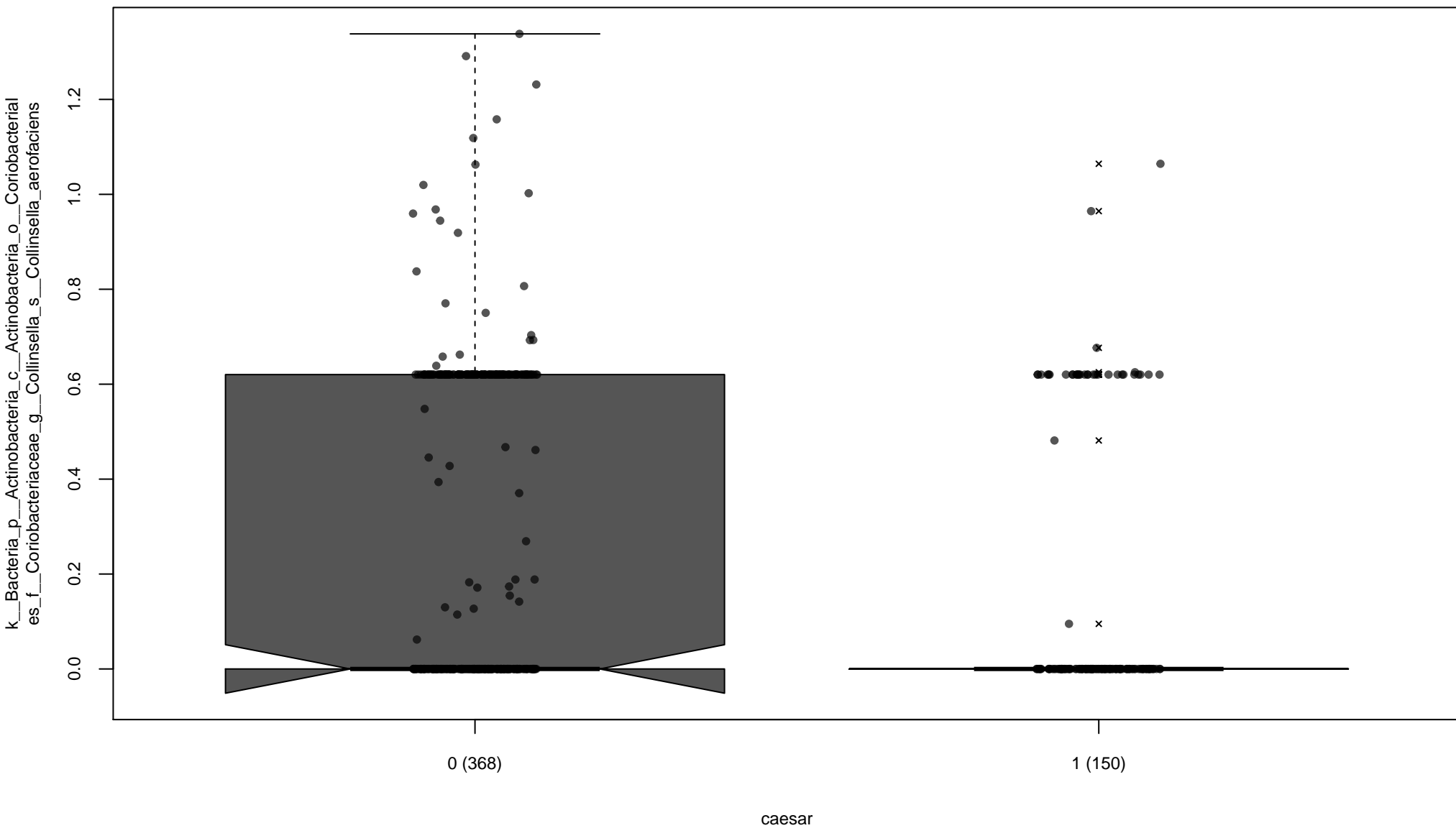


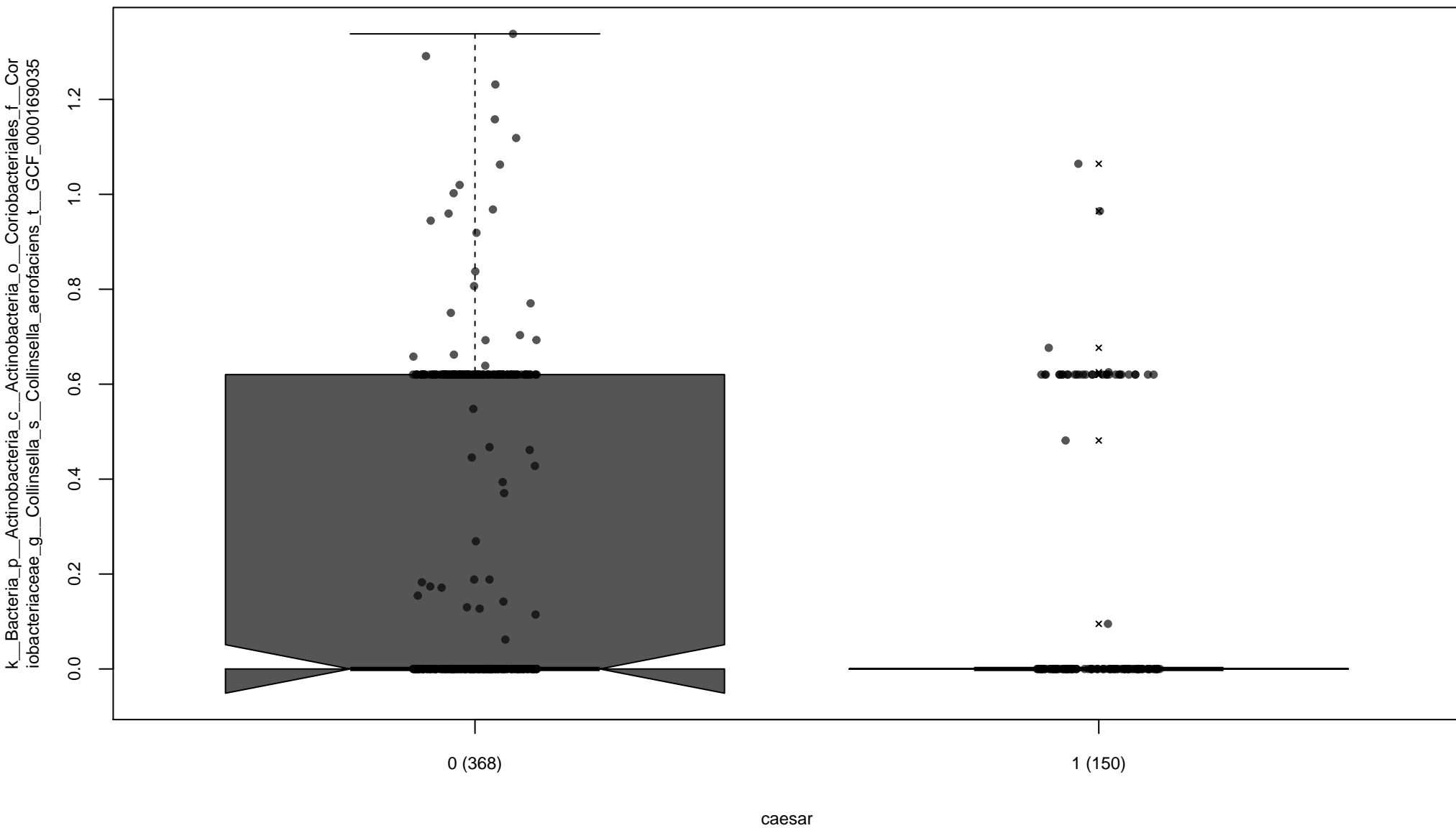
caesar1 (-0.123 sd 0.0264, p=6.01e-06, q=7.34e-05)



caesar1 (-0.121 sd 0.0366, p=0.00114, q=0.0117)

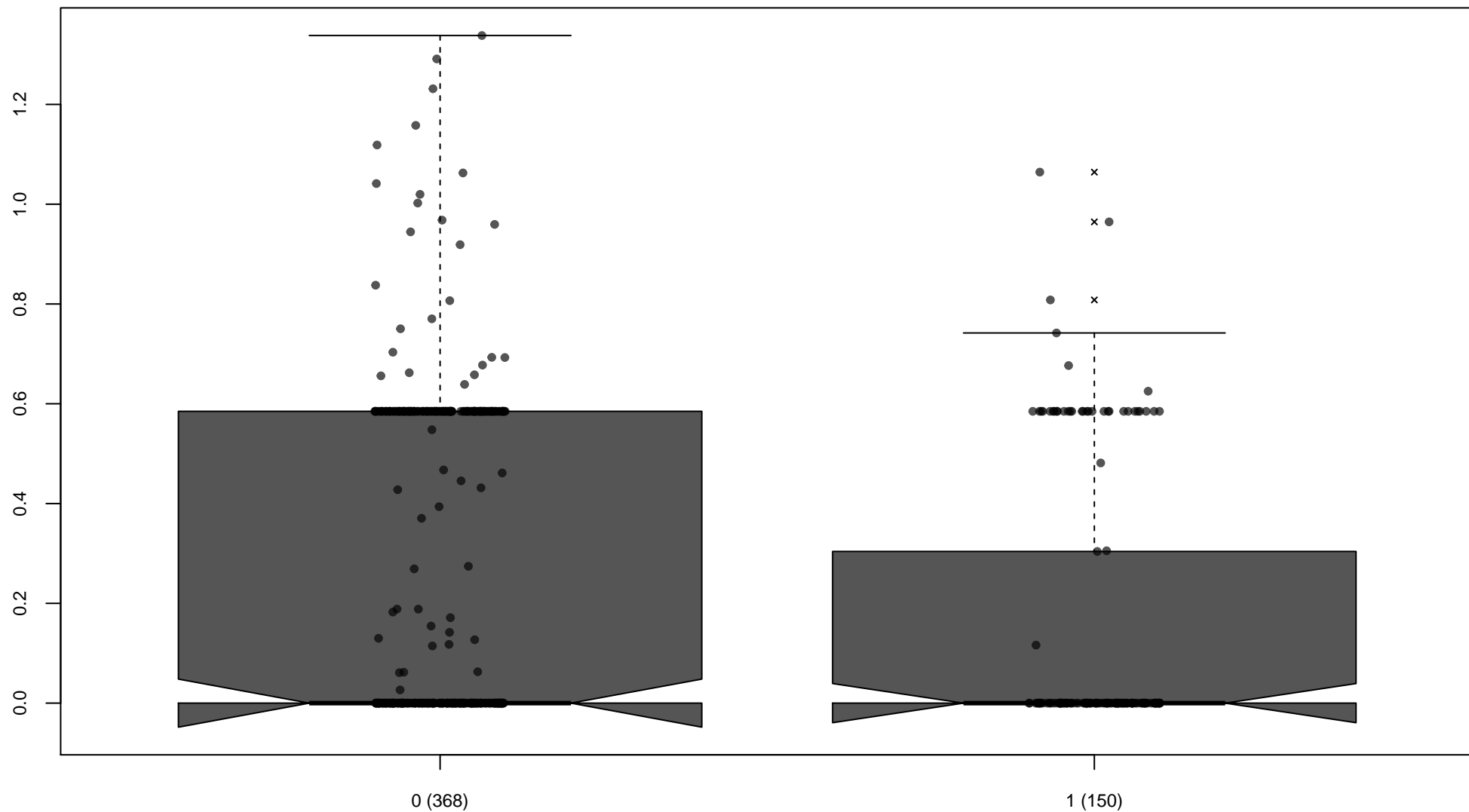


caesar1 (-0.121 sd 0.0366, p=0.00114, q=0.0117)



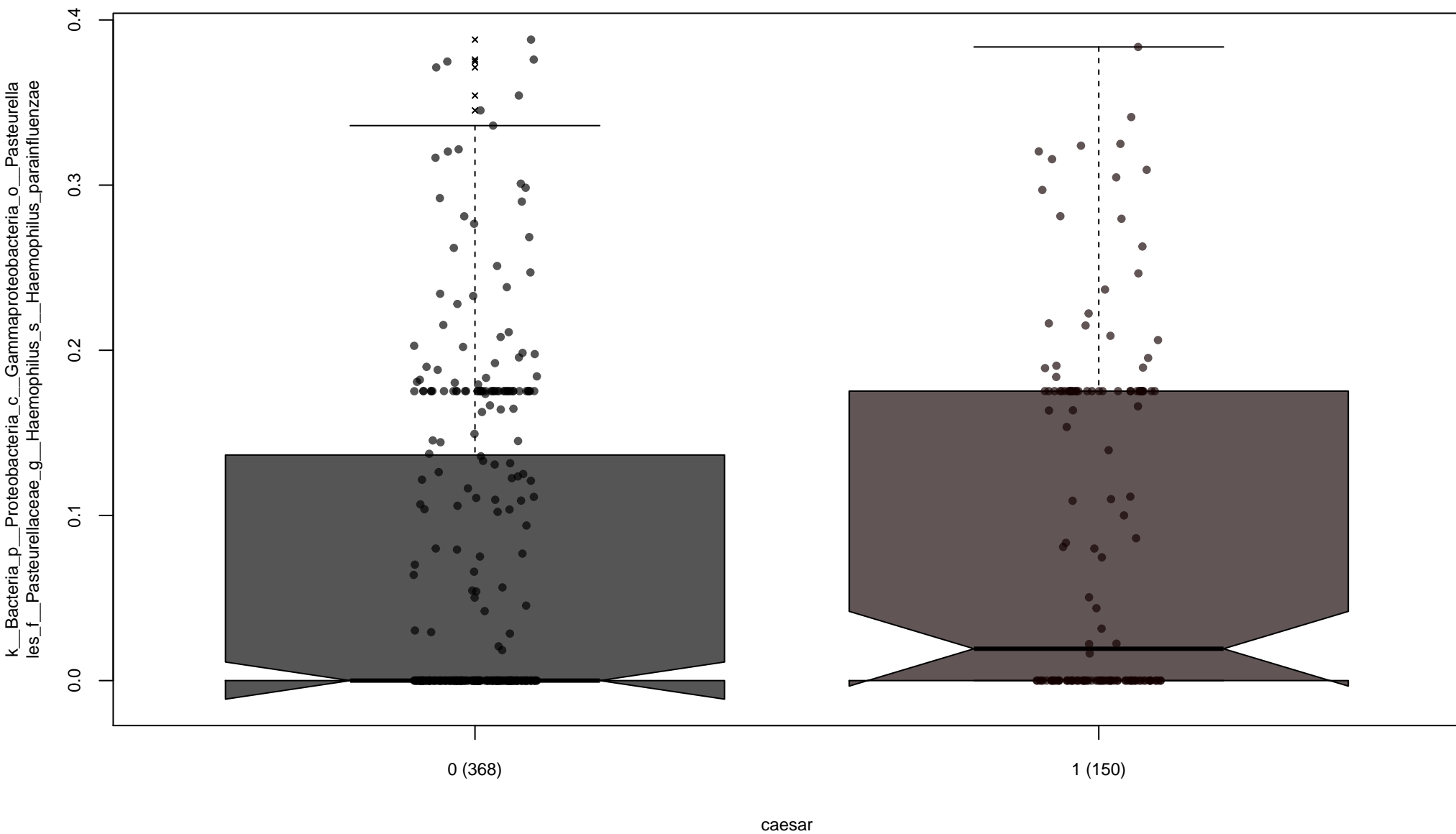
k__Bacteria_p__Actinobacteria_c__Actinobacteria_o__Coriobacteriales_f__Coriobacteriaceae_g__Collinsella

caesar1 (-0.114 sd 0.0348, p=0.00118, q=0.0121)



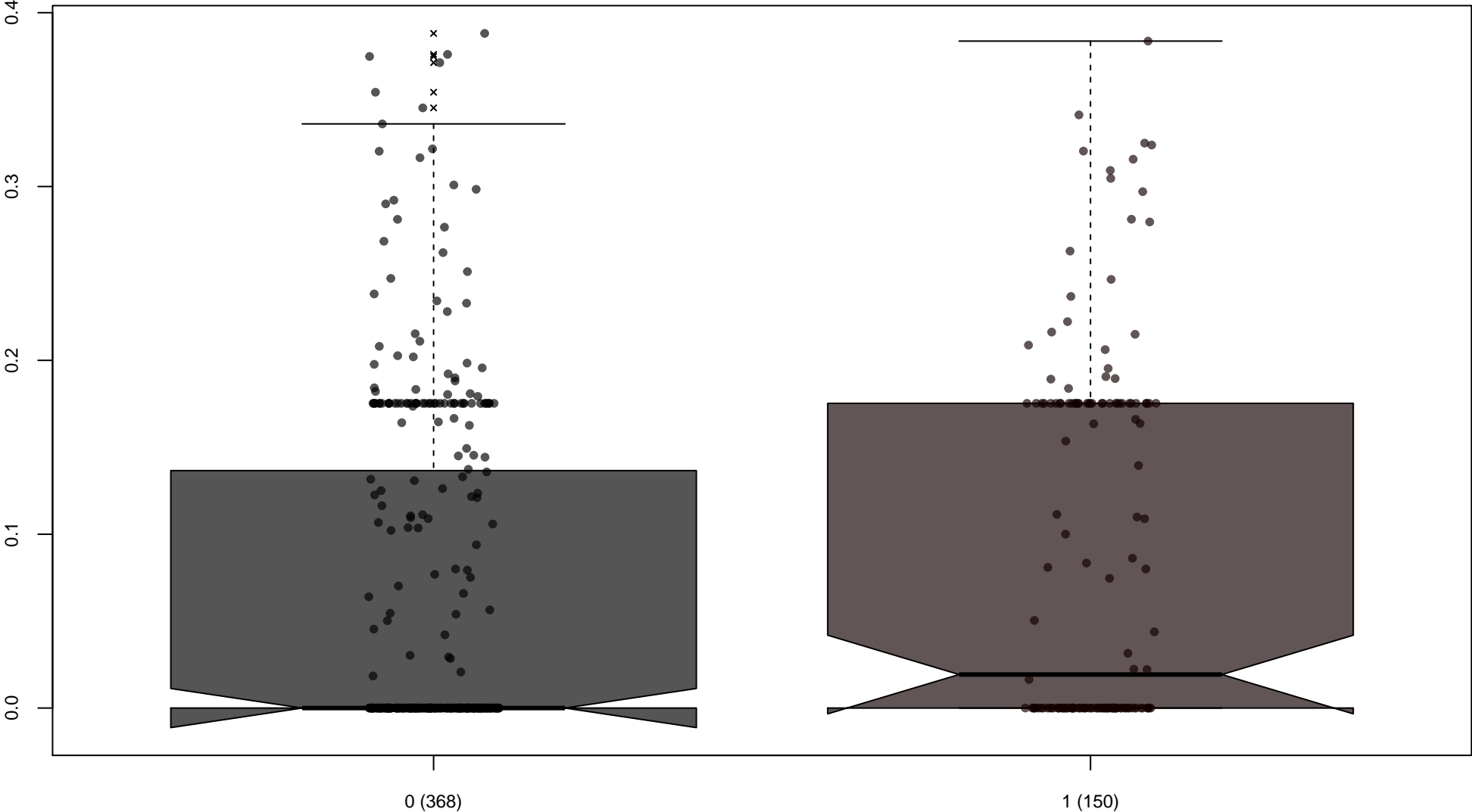
caesar

caesar1 (0.0311 sd 0.00991, p=0.00194, q=0.0195)



caesar1 (0.0311 sd 0.00992, p=0.00197, q=0.0198)

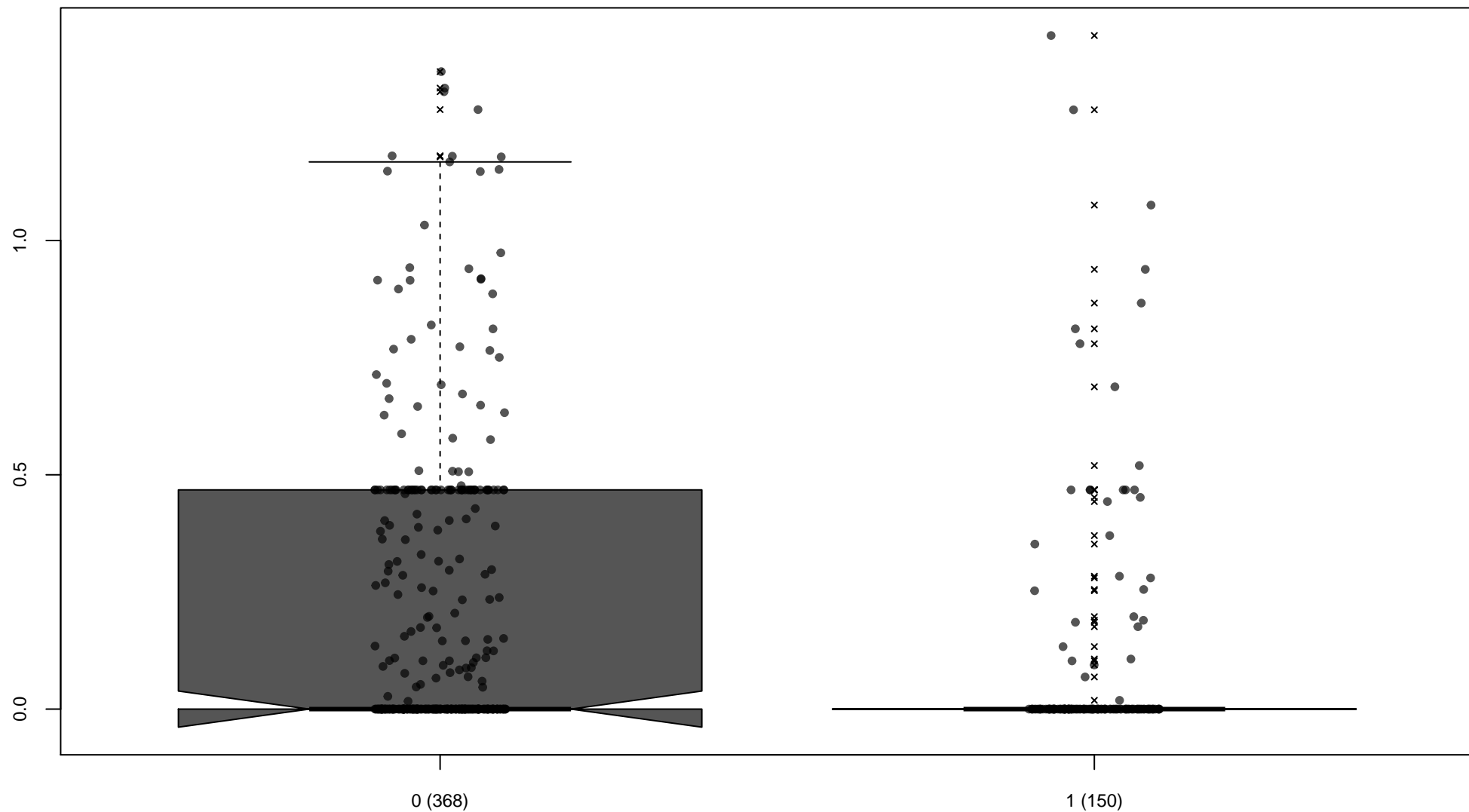
k_Bacteria_p_Proteobacteria_c_Gammaproteobacteria_o_Pasteurellales_f_Pasteurellaceae
e.g_Haemophilus_s_Haemophilus_parainfluenzae_t_Haemophilus_parainfluenzae_unclassified



caesar

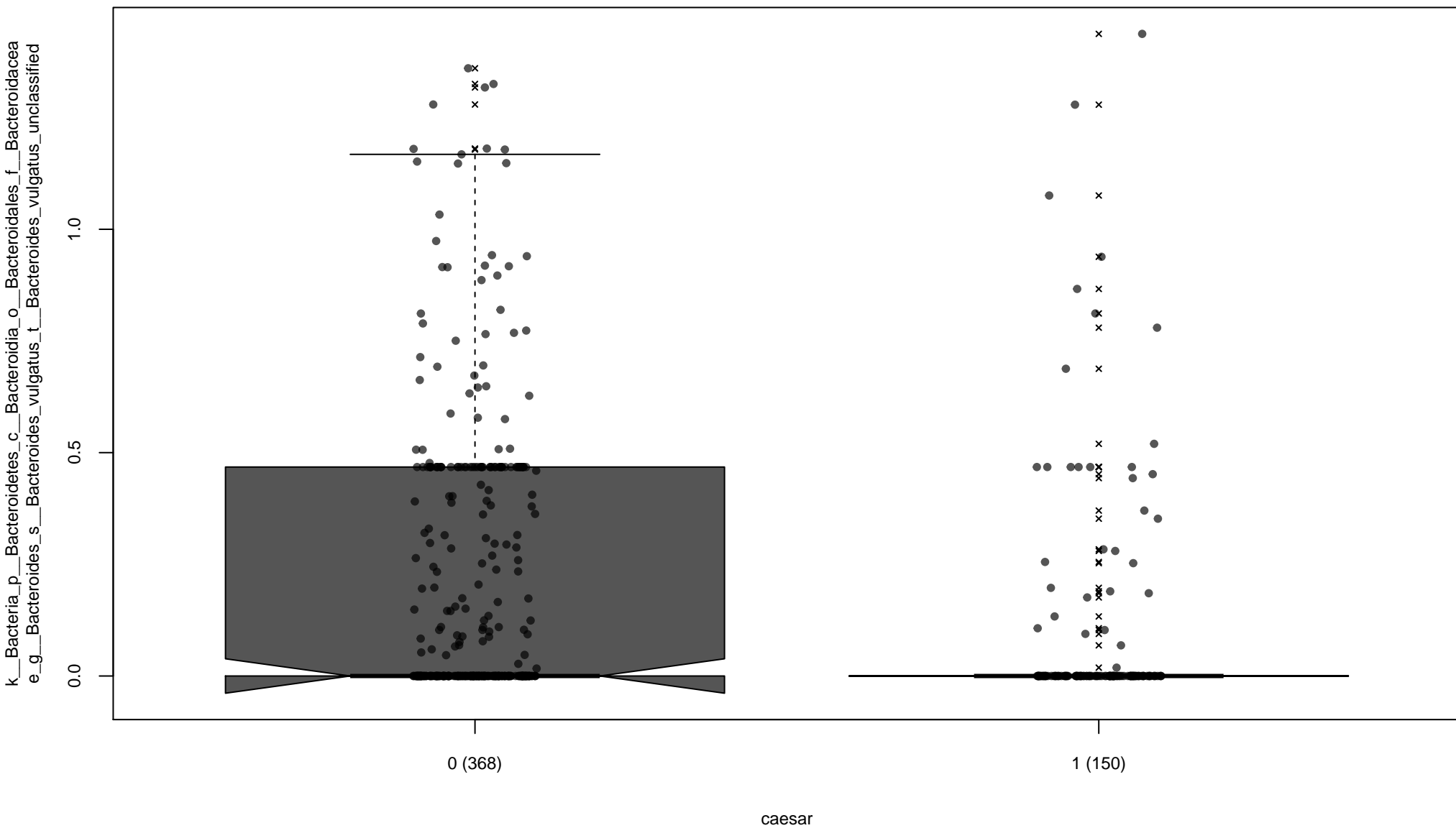
caesar1 (-0.101 sd 0.0324, p=0.00211, q=0.021)

k_Bacteria_p__Bacteroidetes_c__Bacteroidia_o__Bacteroidales_f__Bacteroidaceae_g__Bacteroides_s__Bacteroides_vulg



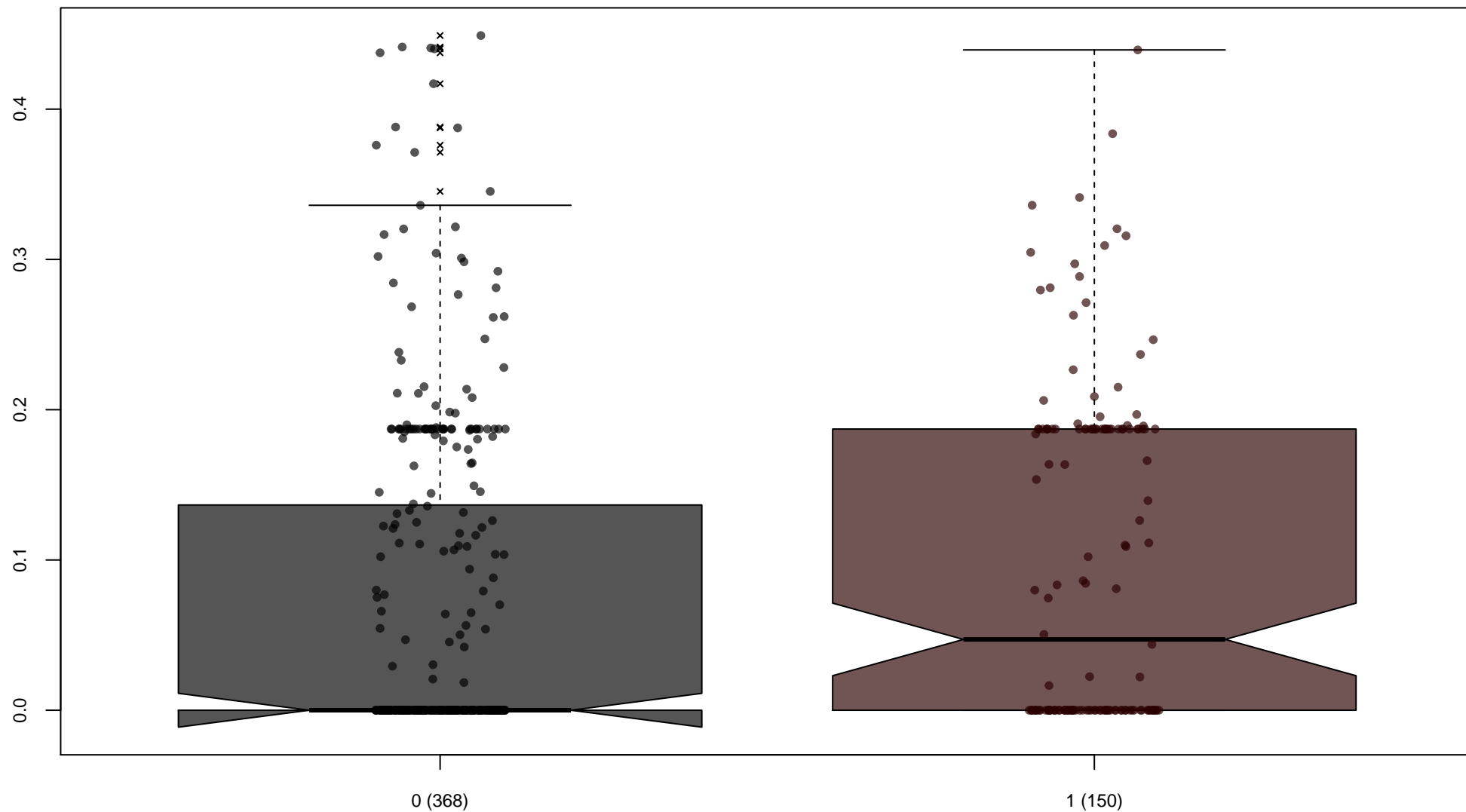
caesar

caesar1 (-0.101 sd 0.0324, p=0.00211, q=0.021)



caesar1 (0.0337 sd 0.0109, p=0.00234, q=0.023)

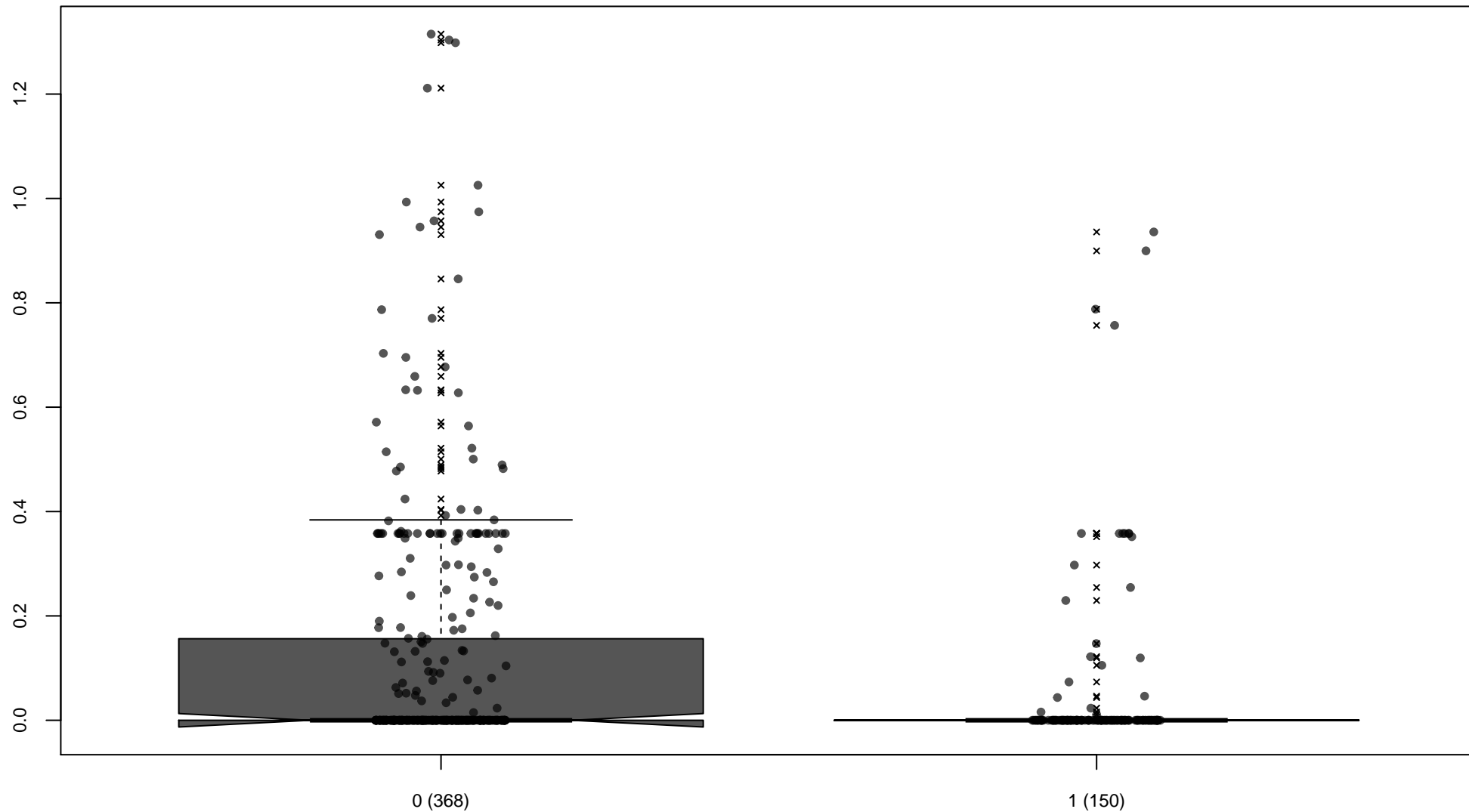
k_Bacteria_p__Proteobacteria_c__Gammaproteobacteria_o__Pasteurellales_f__Pasteurellaceae



caesar

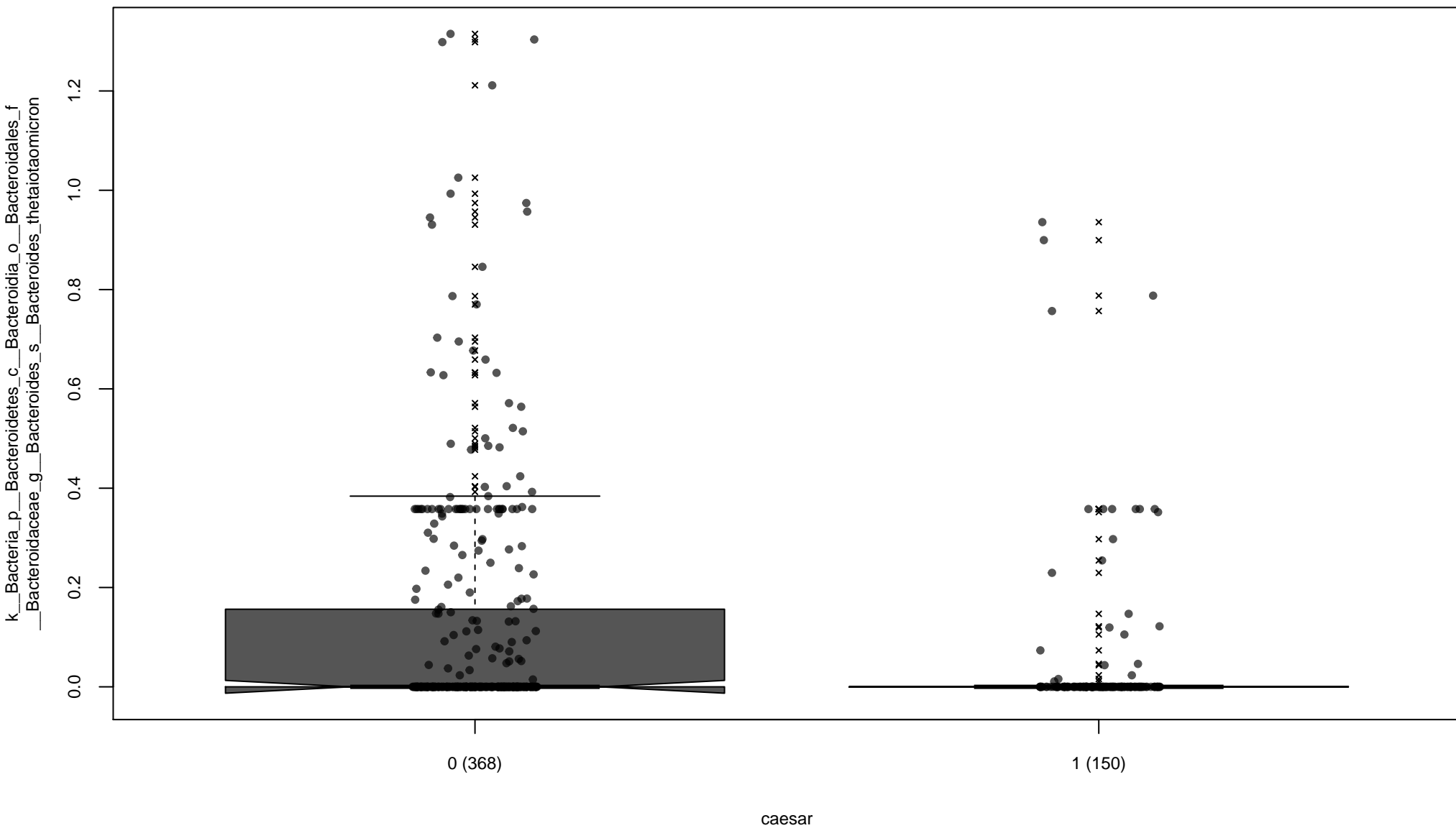
caesar1 (-0.0729 sd 0.0245, p=0.00325, q=0.0317)

k__Bacteria_p__Bacteroidetes_c__Bacteroidia_o__Bacteroidales_f__Bacteroidaceae_g__Bacteroides_thetaiotaomicron_t__Bacteroides_thetaiotaomicron_unclassified

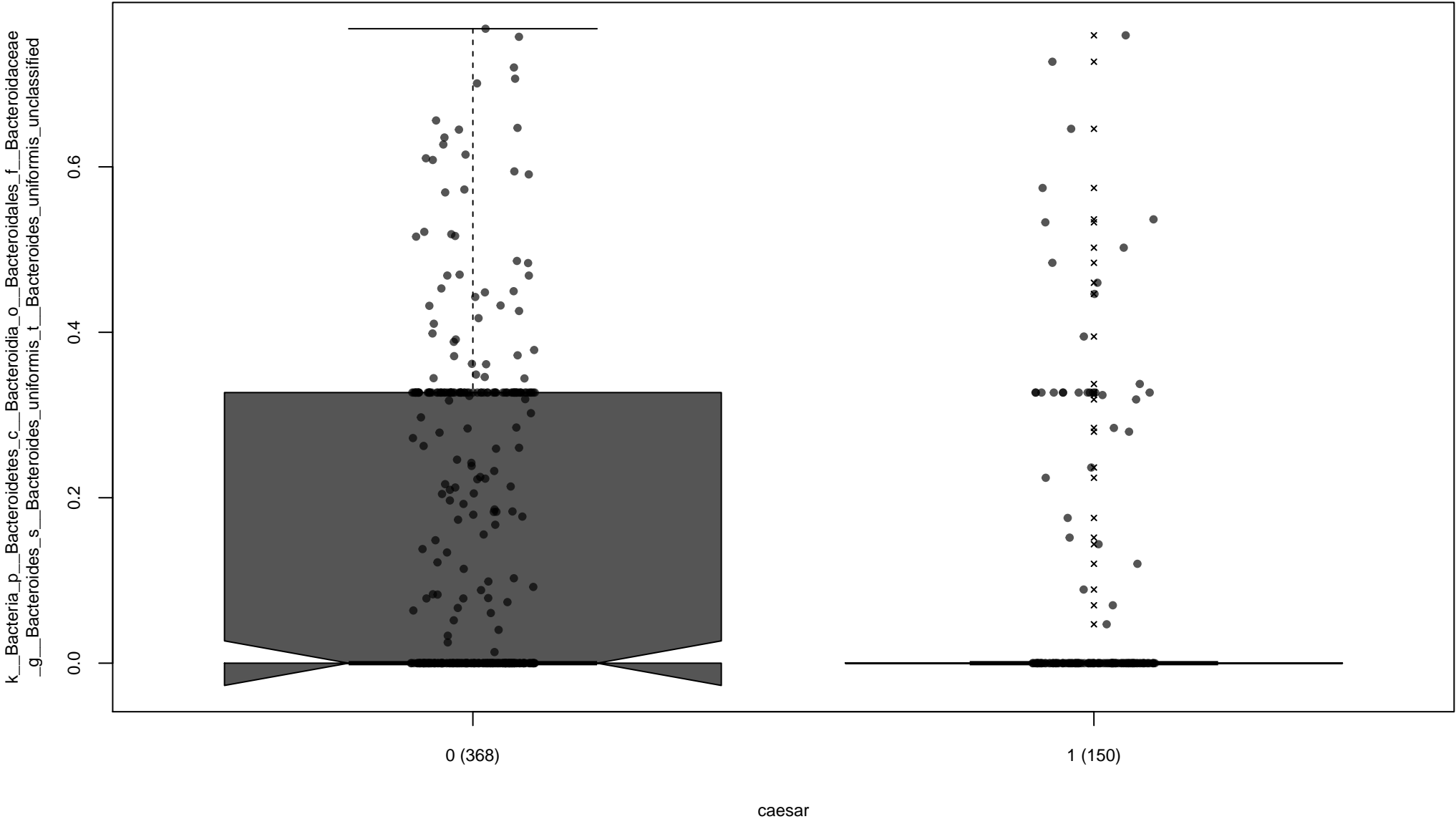


caesar

caesar1 (-0.0723 sd 0.0244, p=0.00346, q=0.0336)



caesar1 (-0.0595 sd 0.0201, p=0.00352, q=0.0341)



caesar1 (-0.0607 sd 0.0218, p=0.00599, q=0.0569)

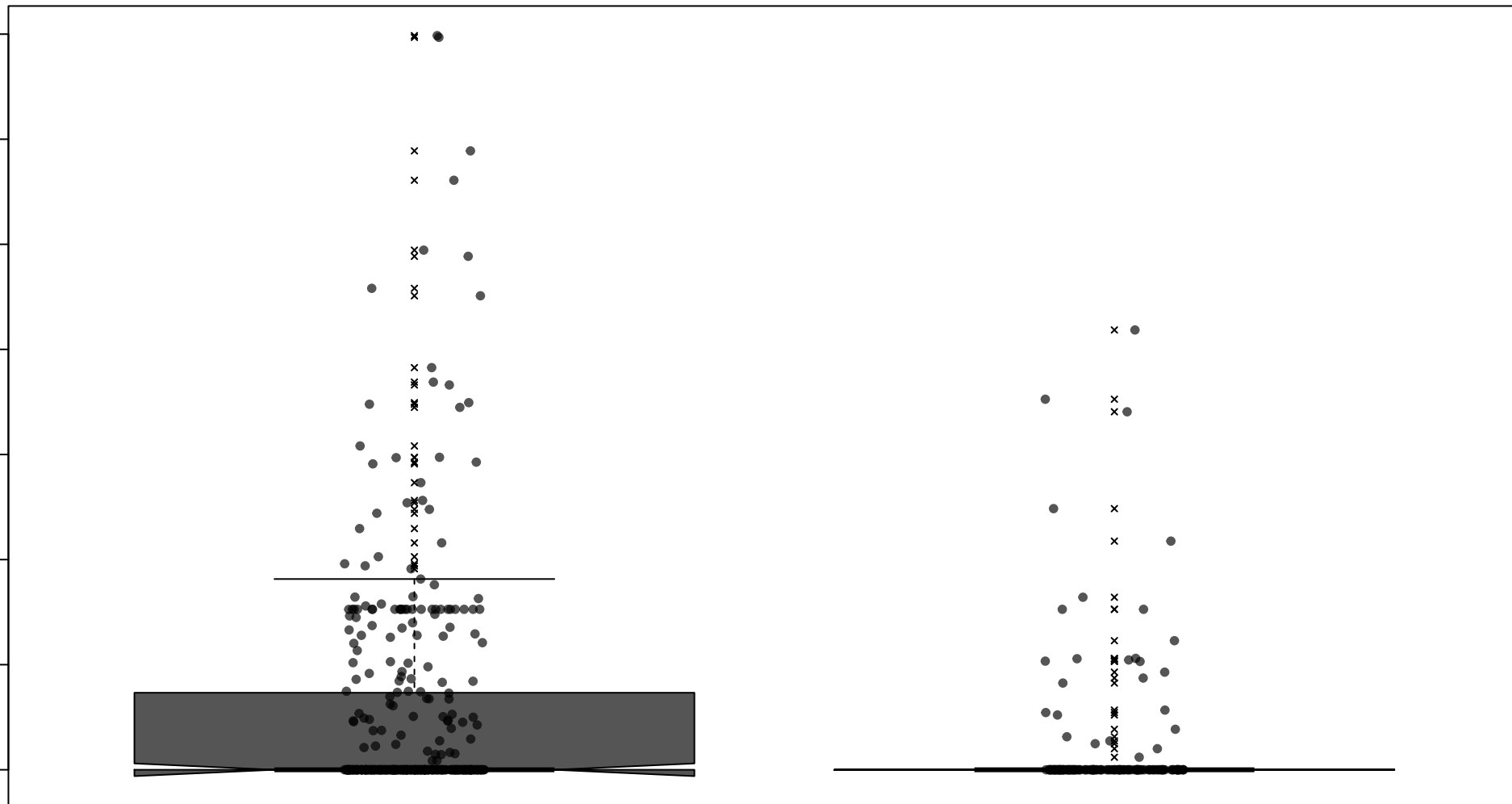
k_Bacteria_p_Bacteroidetes_c_Bacteroidia_o_Bacteroidales_f_Bacteroidaceae_g_Porphyromonadaceae_g_Parabacteroides_s_Parabacteroides_distasonis_t_Parabacteroides_distasonis_u

1.4
1.2
1.0
0.8
0.6
0.4
0.2
0.0

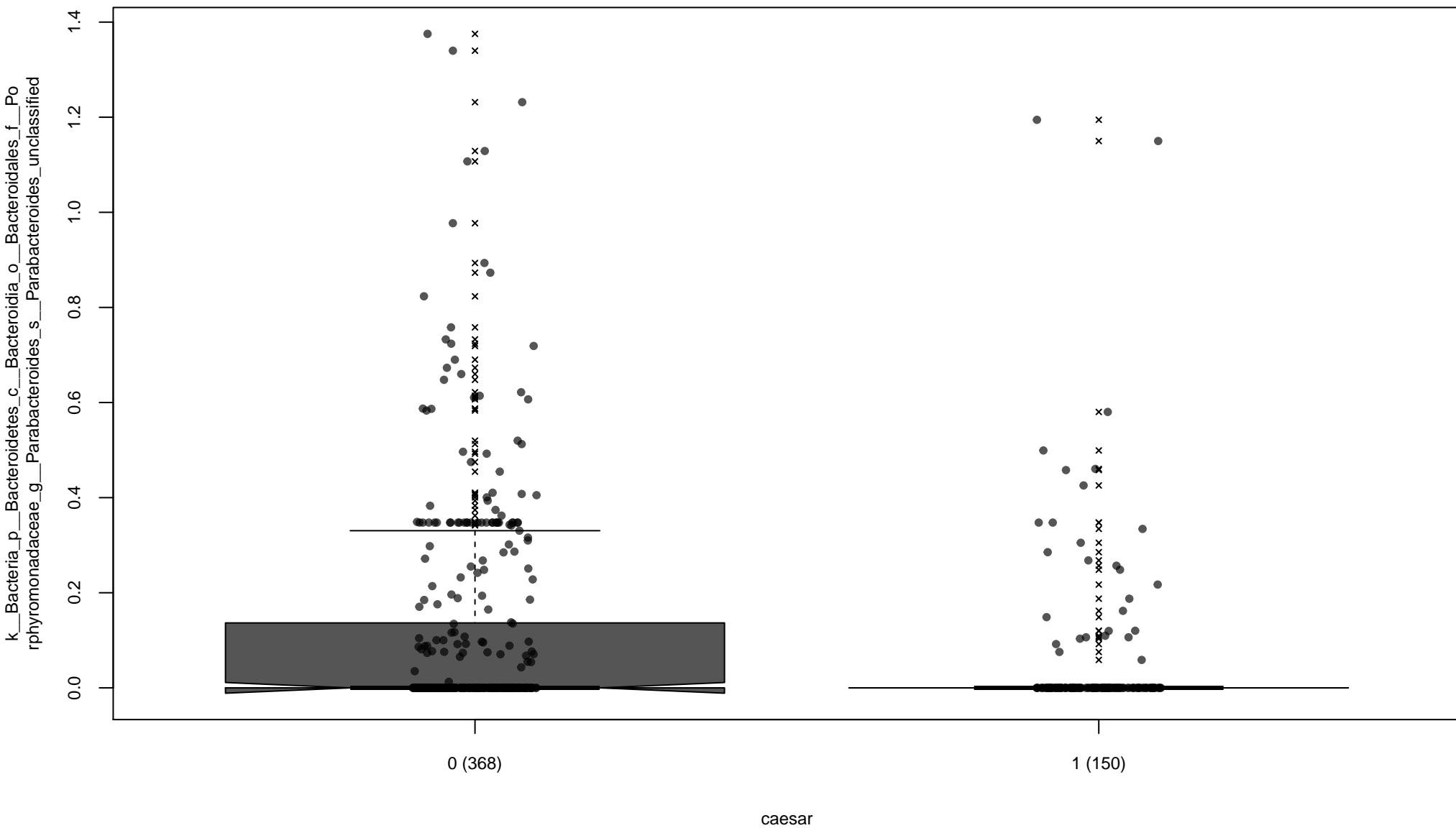
0 (368)

1 (150)

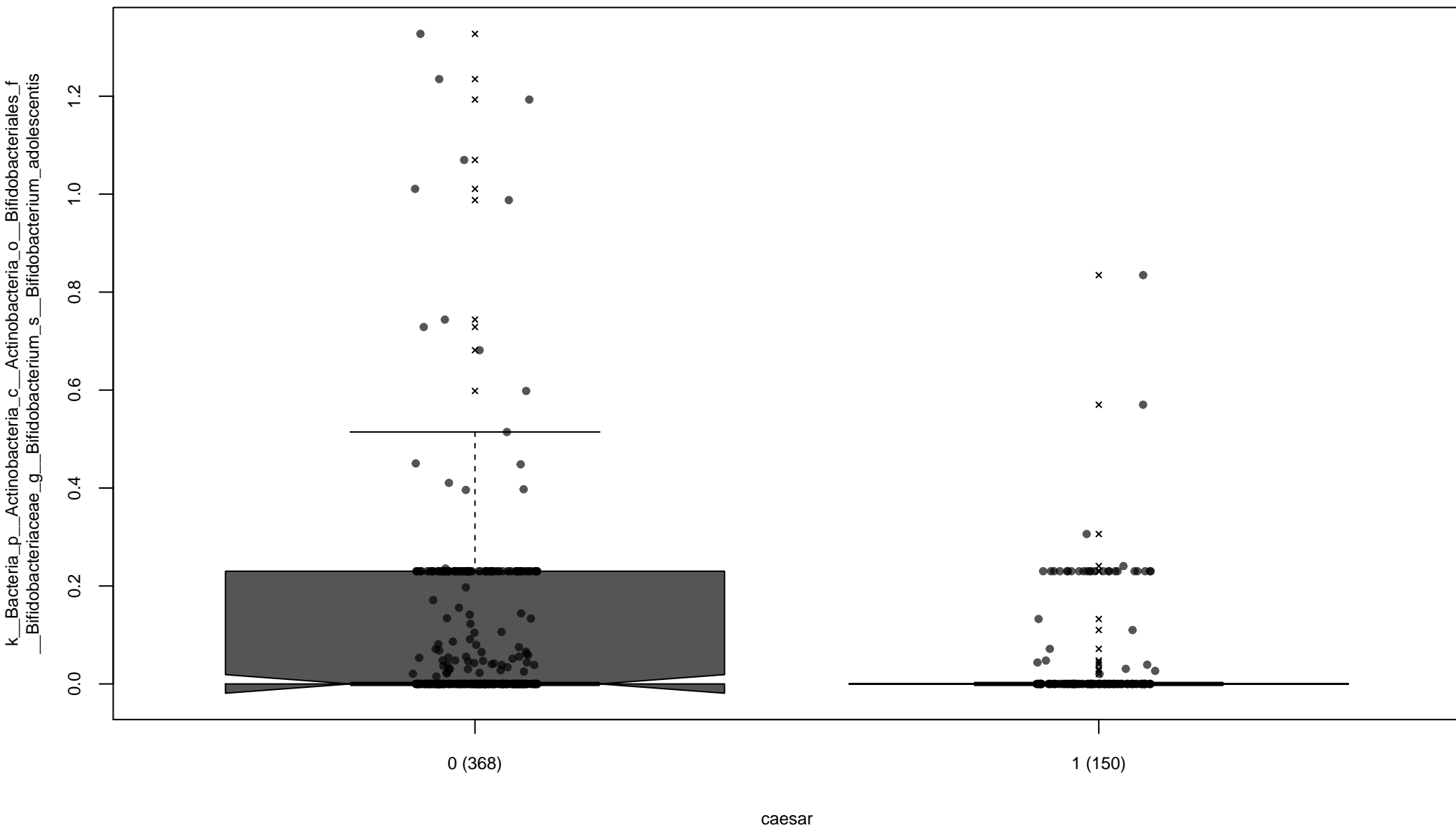
caesar



caesar1 (-0.0607 sd 0.0227, p=0.00806, q=0.0756)

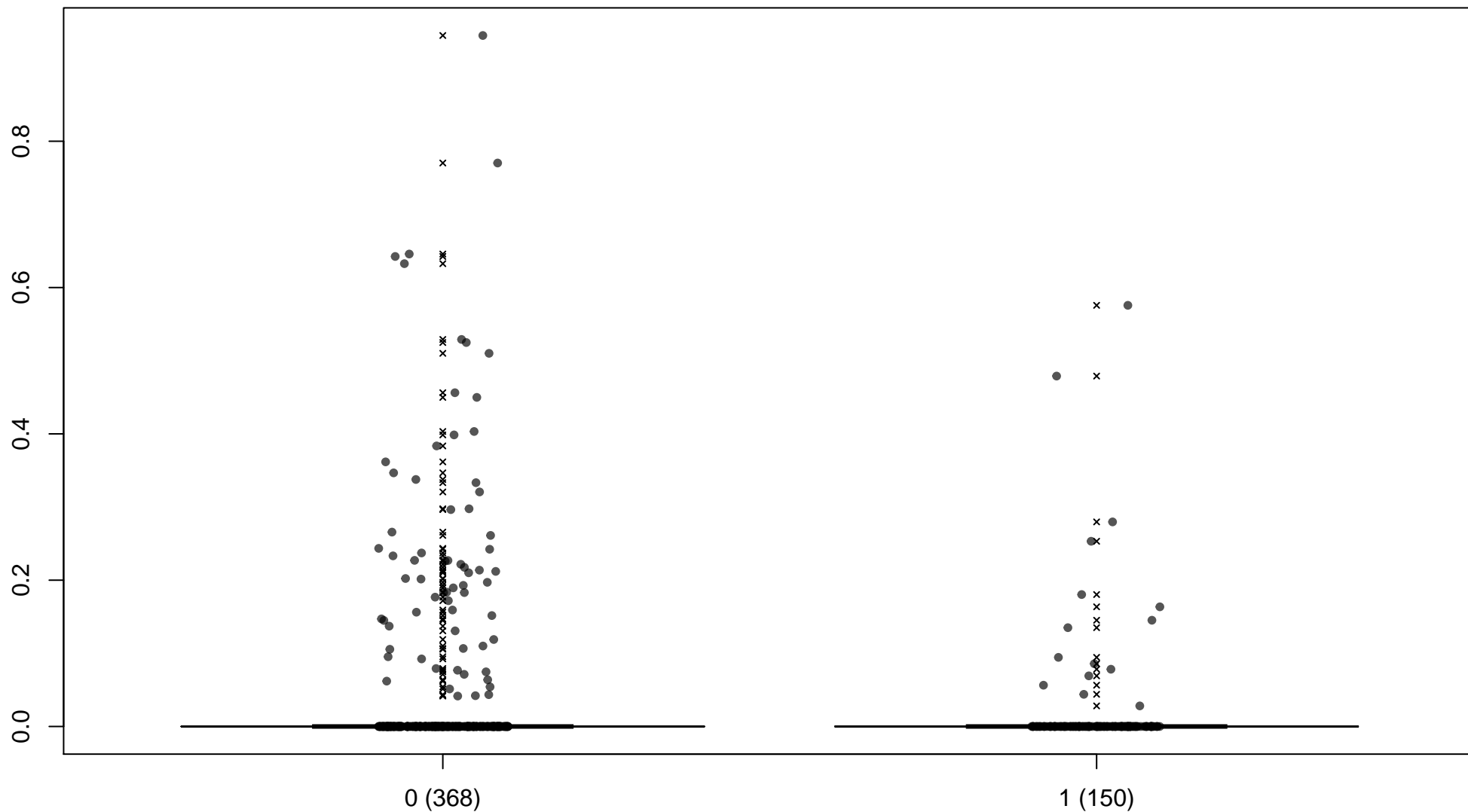


caesar1 (-0.0443 sd 0.0182, p=0.0159, q=0.143)



caesar1 (-0.0288 sd 0.012, p=0.0177, q=0.156)

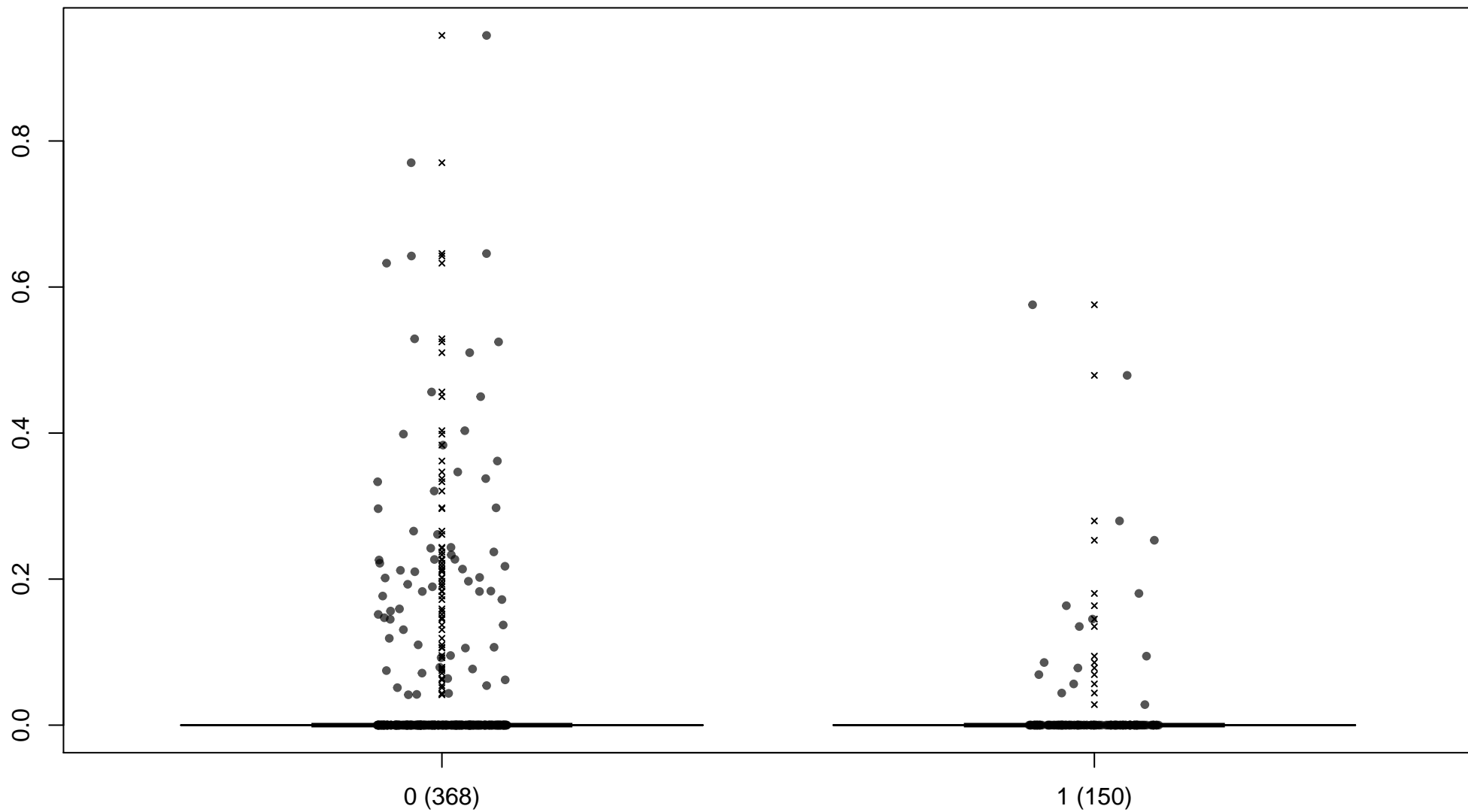
k_Bacteria_p__Proteobacteria_c__Deltaproteobacteria



caesar

k__Bacteria_p__Proteobacteria_c__Deltaproteobacteria_o__Desulfovibrionales

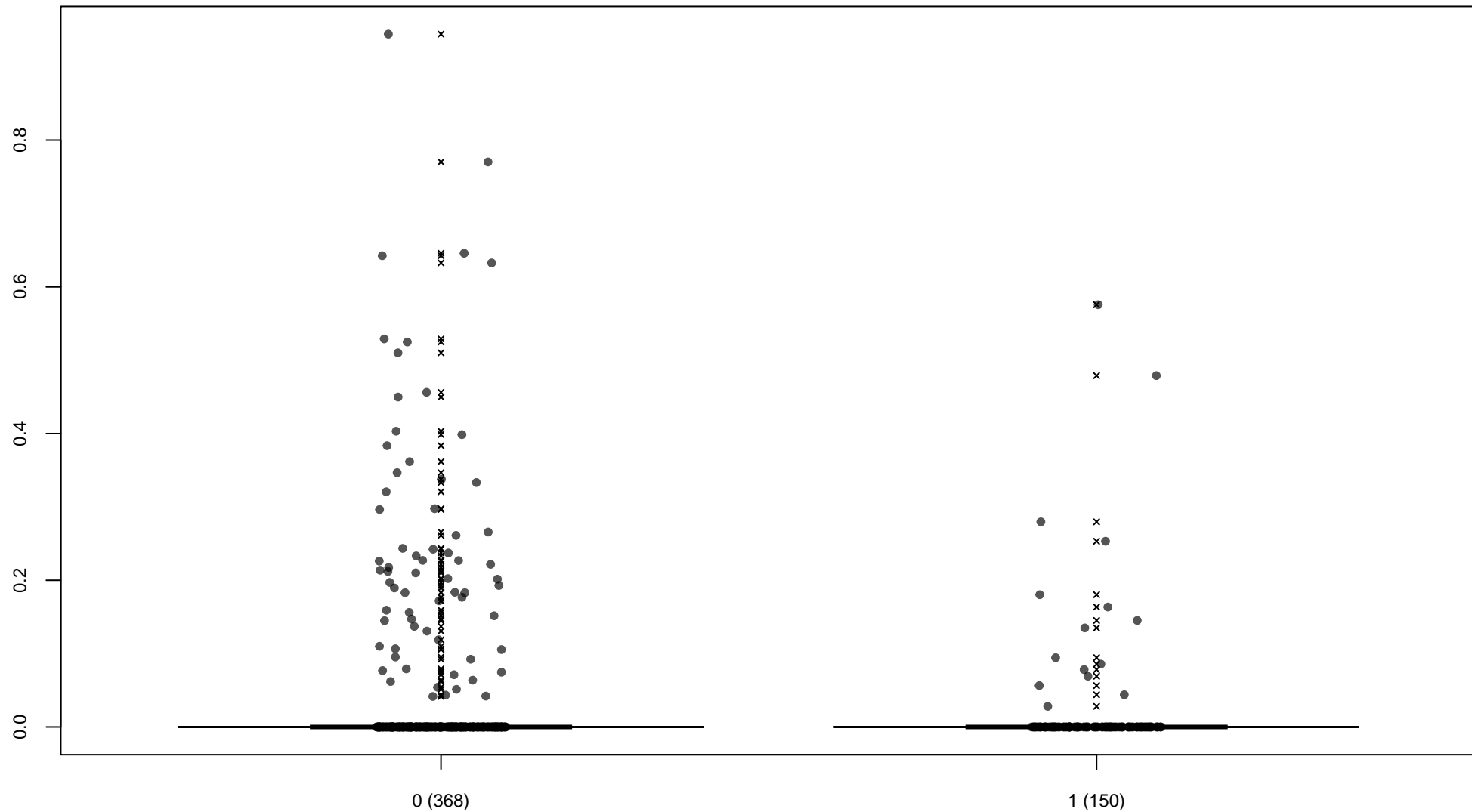
caesar1 (-0.0288 sd 0.012, p=0.0177, q=0.156)



caesar

caesar1 (-0.0288 sd 0.012, p=0.0177, q=0.156)

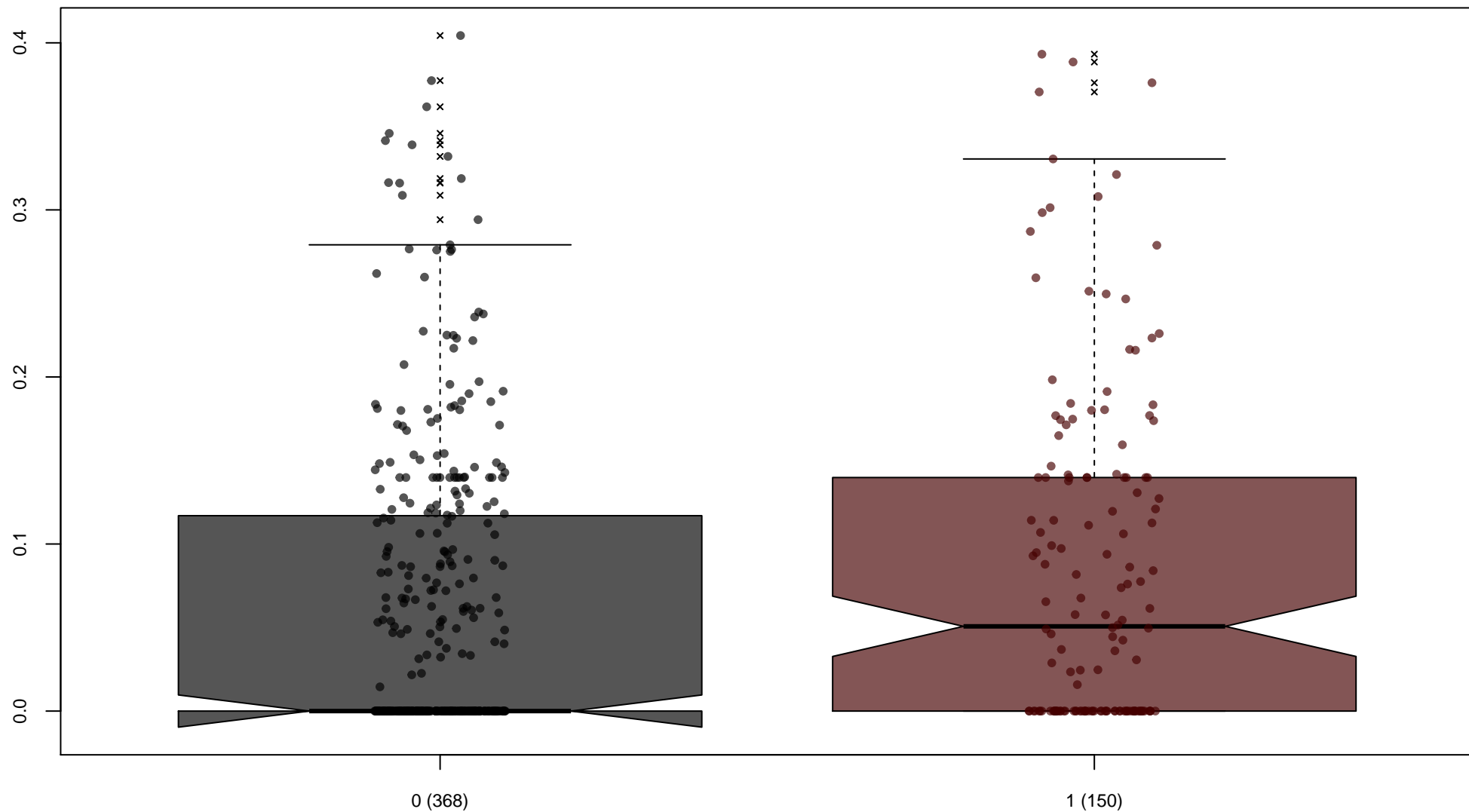
k_Bacteria_p__Proteobacteria_c__Deltaproteobacteria_o__Desulfovibrionales_f__Desulfovibrionaceae



caesar

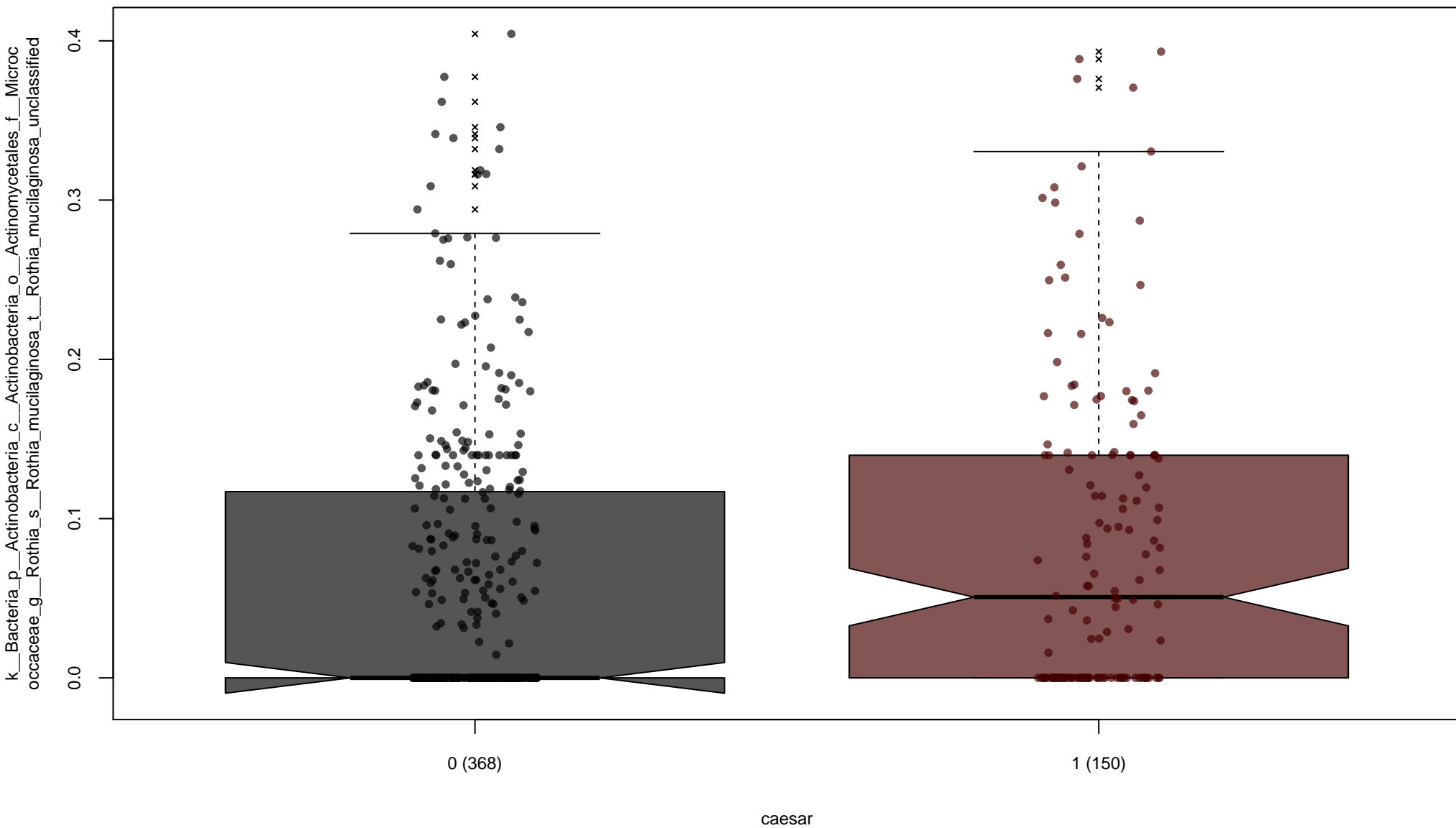
caesar1 (0.0212 sd 0.00923, p=0.0226, q=0.196)

k_Bacteria_p_Actinobacteria_c_Actinobacteria_o_Actinomycetales_f_Micrococaceae_g_Rothia_s_Rothia_mucilaginosa



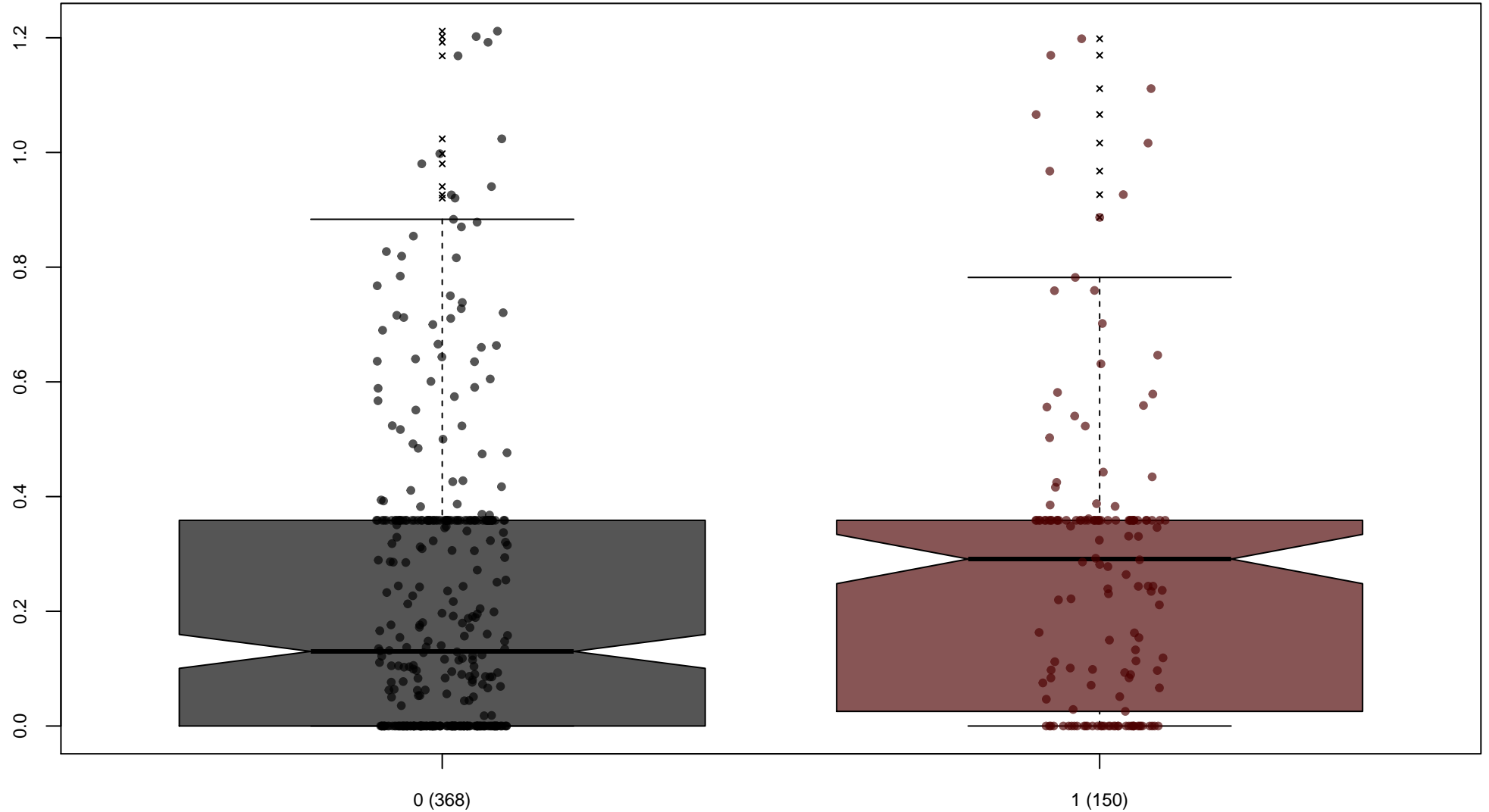
caesar

caesar1 (0.0212 sd 0.00923, p=0.0226, q=0.196)



caesar1 (0.0586 sd 0.0256, p=0.0233, q=0.201)

k__Bacteria_p__Firmicutes_c__Negativicutes_o__Selenomonadales_f__Veillonellaceae_g__Veillonella



caesar