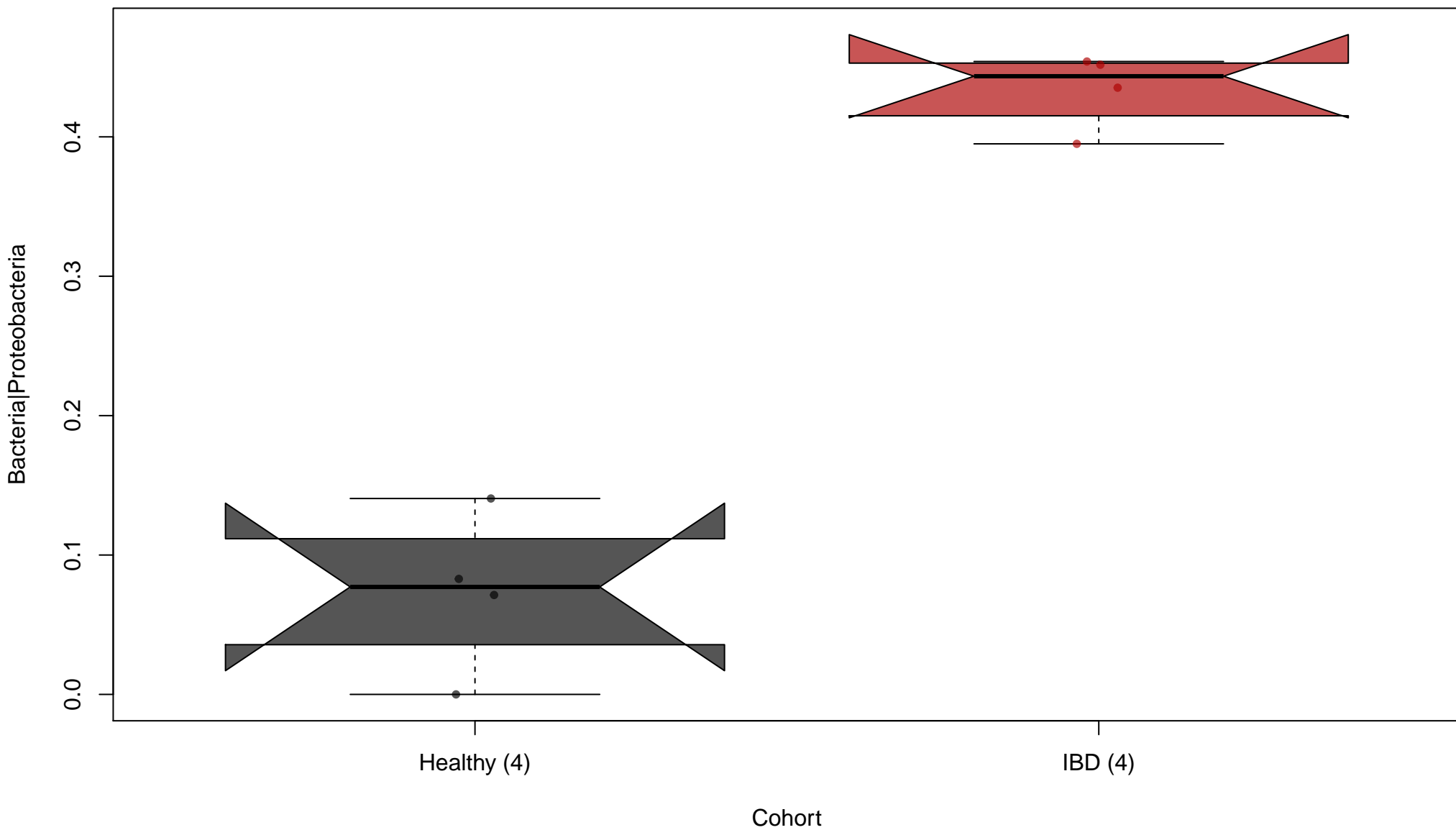
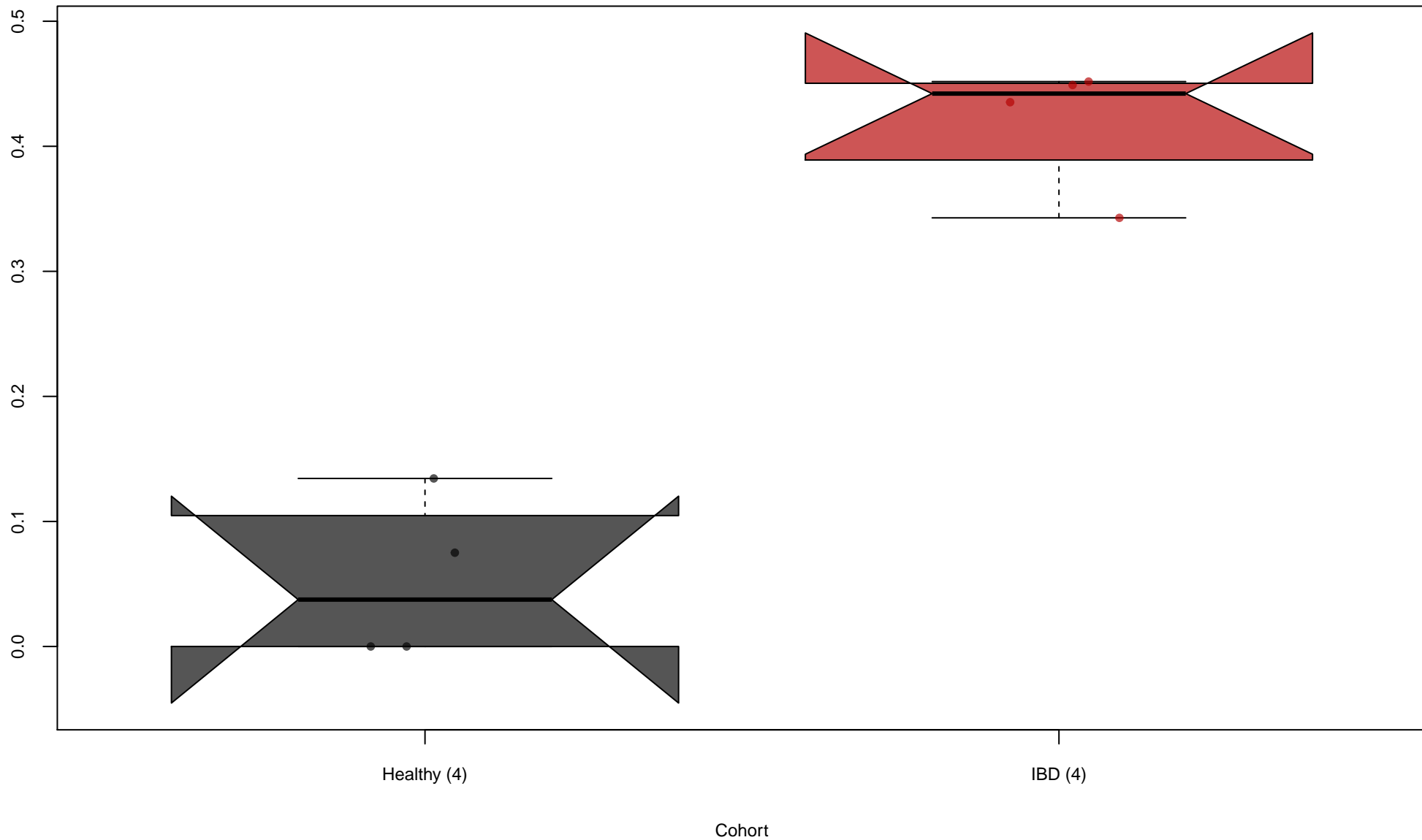


CohortIBD (0.361 sd 0.0311, $p=8.3e-05$, $q=0.0133$)



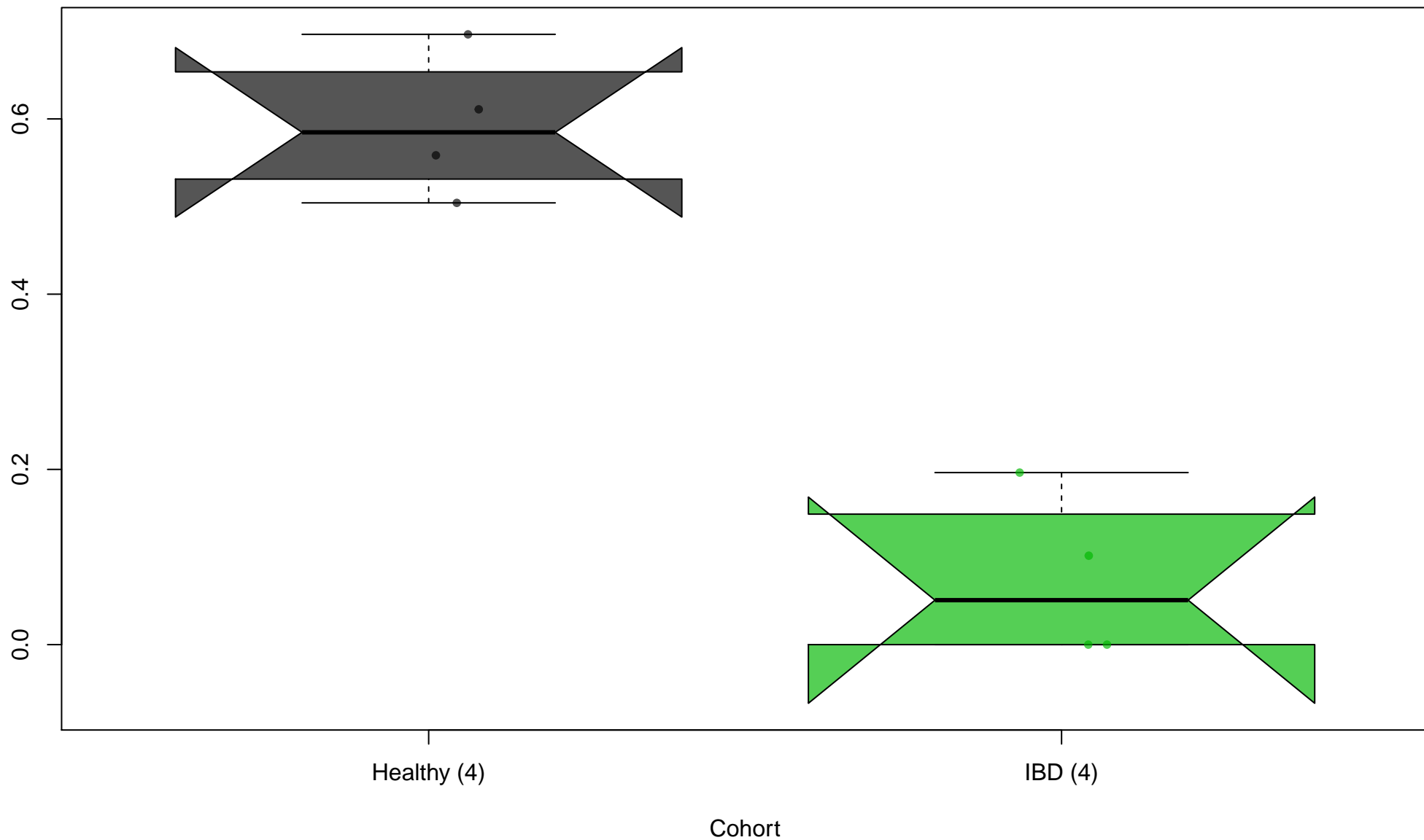
CohortIBD (0.368 sd 0.0409, p=0.000283, q=0.0226)

Bacteria|Proteobacteria|Gammaproteobacteria|Enterobacteriales|Enterobacteriaceae|Escherichia_Shigella|1532



Bacteria|Firmicutes|Clostridia|Clostridiales|Lachnospiraceae|Roseburia|1032

Cohort|IBD (-0.518 sd 0.0682 , $p=0.000628$, $q=0.0251$)



CohortIBD (-0.271 sd 0.0412 , $p=0.00121$, $q=0.0388$)

Bacteria|Firmicutes|Clostridia|Clostridiales|Ruminococcaceae|1156

