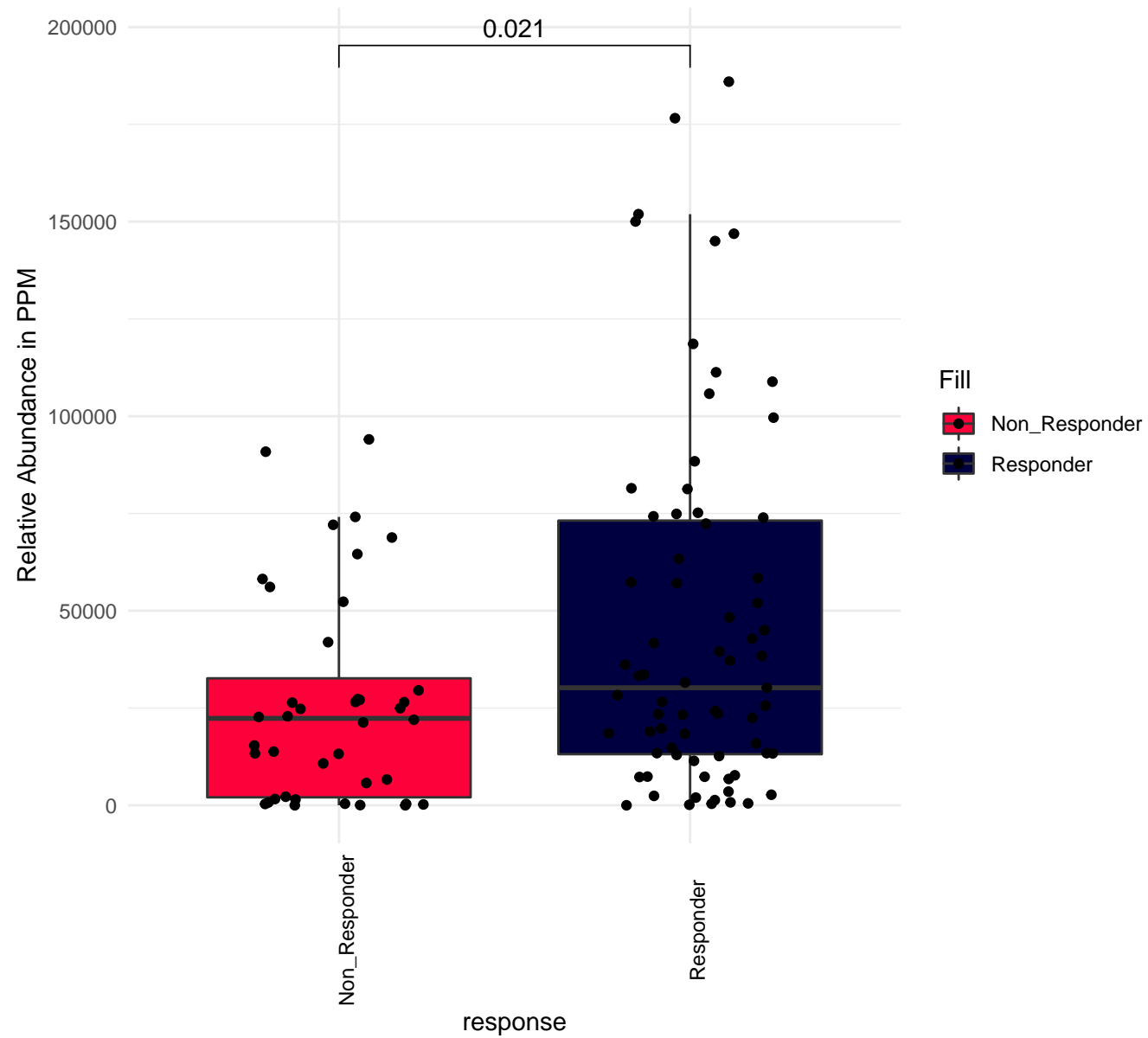


Relative Abundance Plot | Genus

g__Faecalibacterium

MannWhitneyWilcoxon | pval=0.0214 | padj_fdr=0.5787 | stat=1045 | Number of features assessed = 5

Log2FC=-1.567 | Positive l2fc means increased in Non_Responder



Relative Abundance Plot | Family

f__Ruminococcaceae

MannWhitneyWilcoxon | pval=0.0149 | padj_fdr=0.3133 | stat=1023 | Number of features assessed = 28

Log2FC=-1.173 | Positive l2fc means increased in Non_Responder

