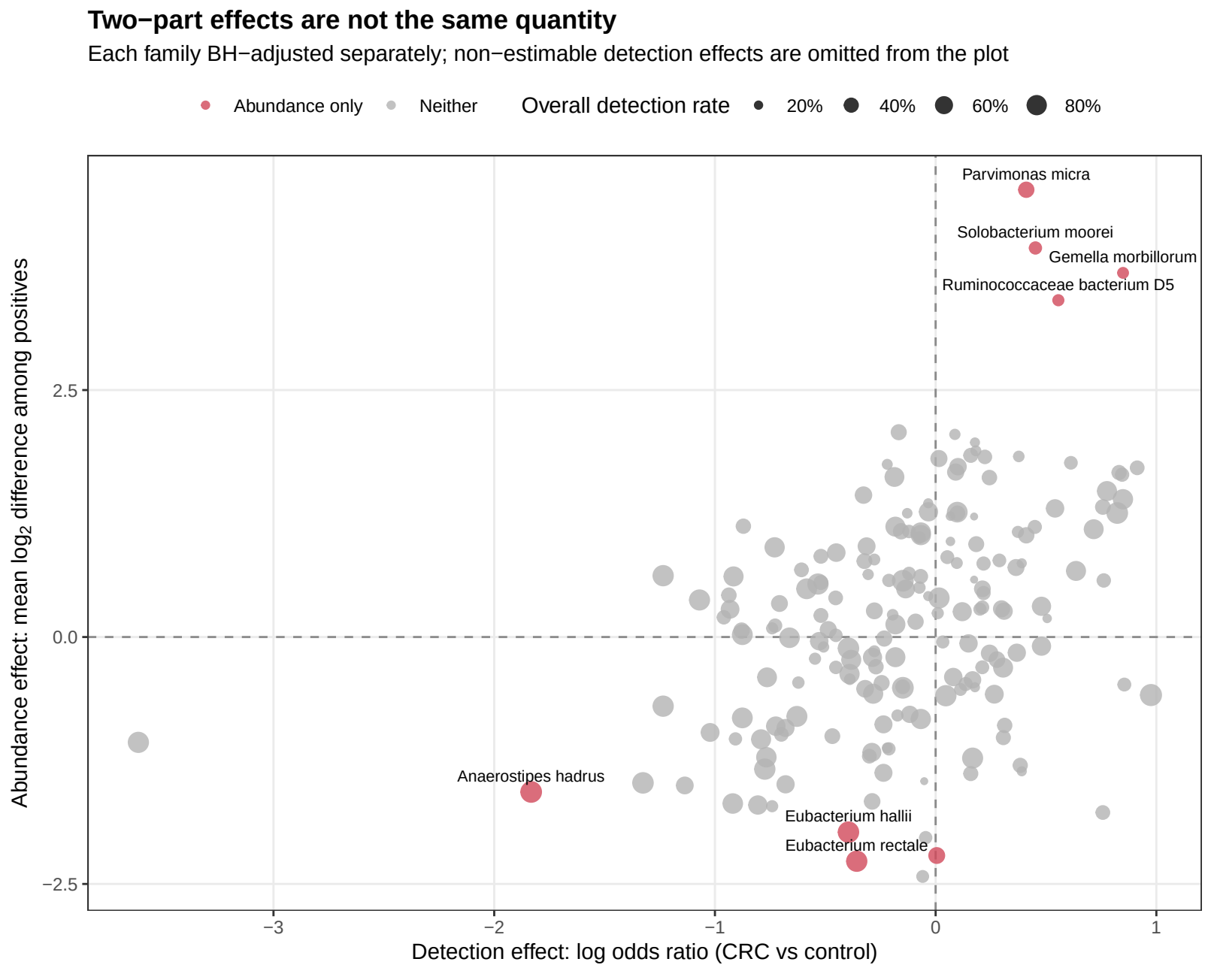
Two−part effects are not the same quantity
Each family BH−adjusted separately; non−estimable detection effects are omitted from the plot
Abundance only
Neither
Overall detection rate
20%
40%
60%
80%
Parvimonas micra
Solobacterium moorei
Gemella morbillorum
Ruminococcaceae bacterium D5
Anaerostipes hadrus
Eubacterium hallii
Eubacterium rectale
2.5
0.0
−2.5
−3
−2
−1
0
1
Abundance effect: mean log2 difference among positives
Detection effect: log odds ratio (CRC vs control)