Supplementary figure S11

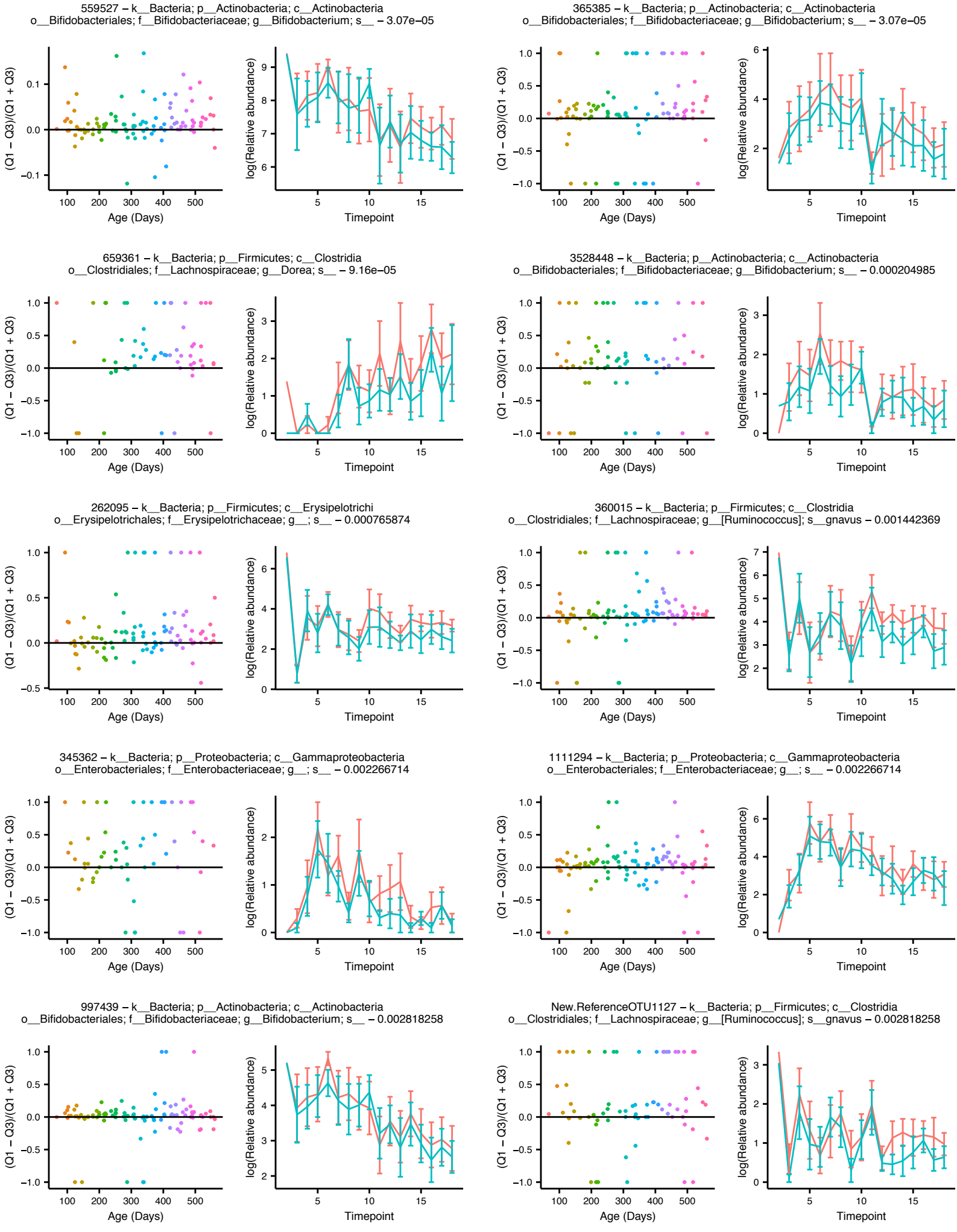
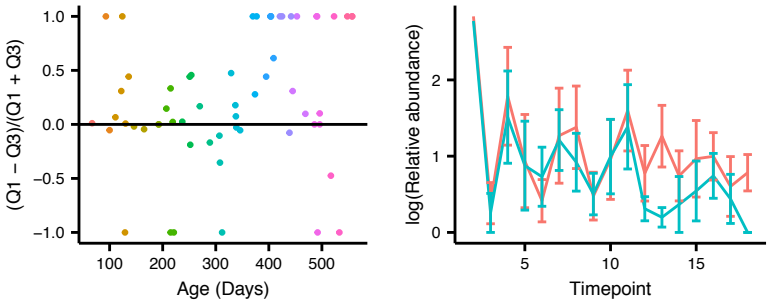
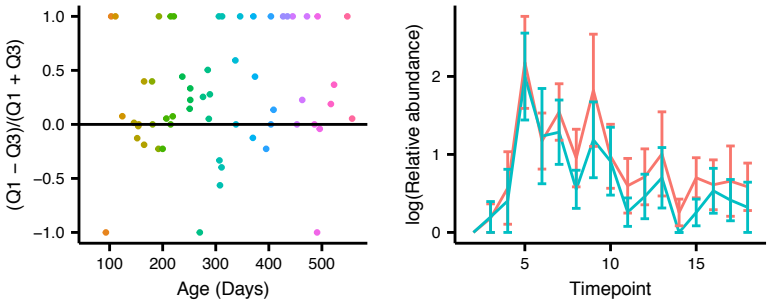


Figure Teddy_Q1_vs_Q3_supp

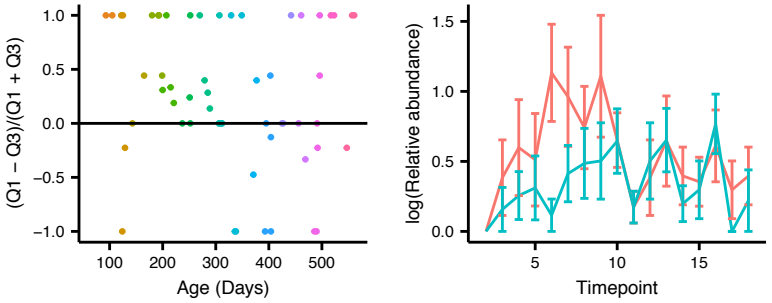
New.ReferenceOTU2021 – k__Bacteria; p__Firmicutes; c__Clostridia
o__Clostridiales; f__Lachnospiraceae; g__; s__ – 0.003342723



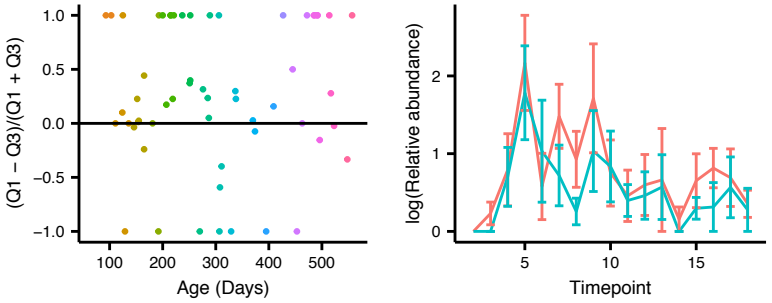
4333897 – k__Bacteria; p__Proteobacteria; c__Gammaproteobacteria
o__Enterobacteriales; f__Enterobacteriaceae; g__; s__ – 0.004022792



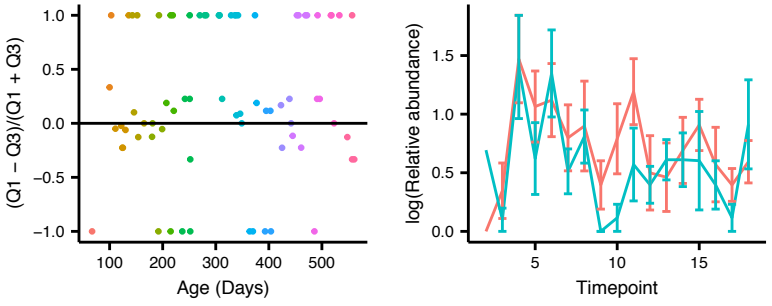
4481861 – k__Bacteria; p__Actinobacteria; c__Actinobacteria
o__Bifidobacteriales; f__Bifidobacteriaceae; g__Bifidobacterium; s__ – 0.004201506



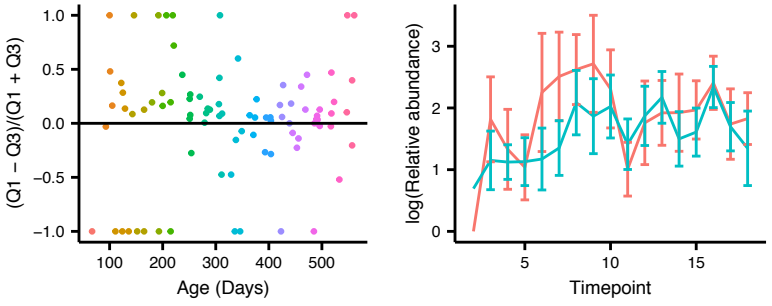
1109247 – k__Bacteria; p__Proteobacteria; c__Gammaproteobacteria
o__Enterobacteriales; f__Enterobacteriaceae; g__; s__ – 0.004446503



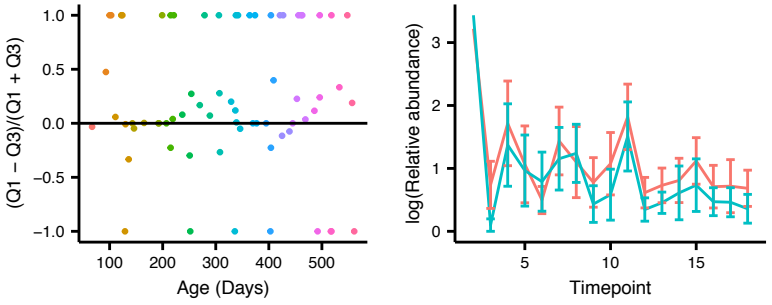
787709 – k__Bacteria; p__Actinobacteria; c__Actinobacteria
o__Actinomycetales; f__Actinomycetaceae; g__Actinomyces; s__ – 0.006104403



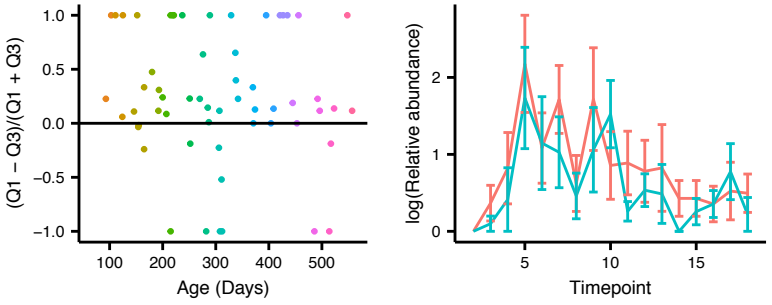
822770 – k__Bacteria; p__Actinobacteria; c__Actinobacteria
o__Bifidobacteriales; f__Bifidobacteriaceae; g__Bifidobacterium; s__ – 0.006989366



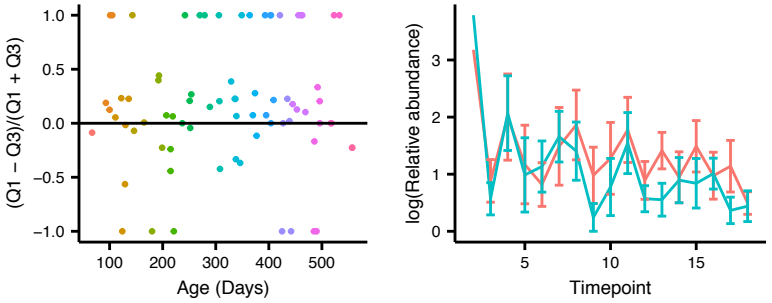
176704 – k__Bacteria; p__Firmicutes; c__Clostridia
o__Clostridiales; f__Lachnospiraceae; g__[Ruminococcus]; s__gnavus – 0.006989366



231787 – k__Bacteria; p__Proteobacteria; c__Gammaproteobacteria
o__Enterobacteriales; f__Enterobacteriaceae; g__; s__ – 0.006989366



3715618 – k__Bacteria; p__Firmicutes; c__Clostridia
o__Clostridiales; f__Lachnospiraceae; g__[Ruminococcus]; s__gnavus – 0.006989366



813217 – k__Bacteria; p__Proteobacteria; c__Gammaproteobacteria
o__Enterobacteriales; f__Enterobacteriaceae; g__; s__ – 0.012425239

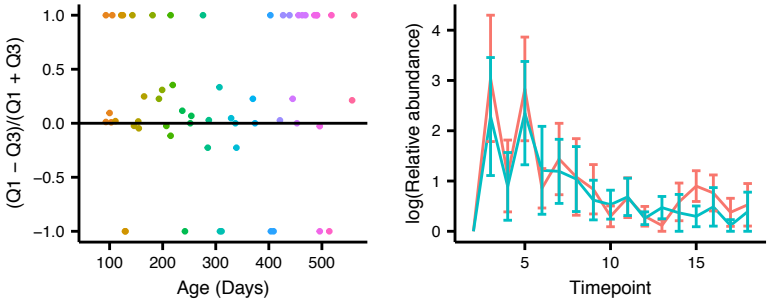
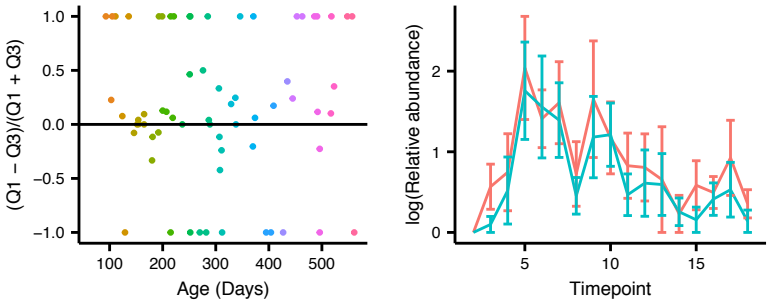
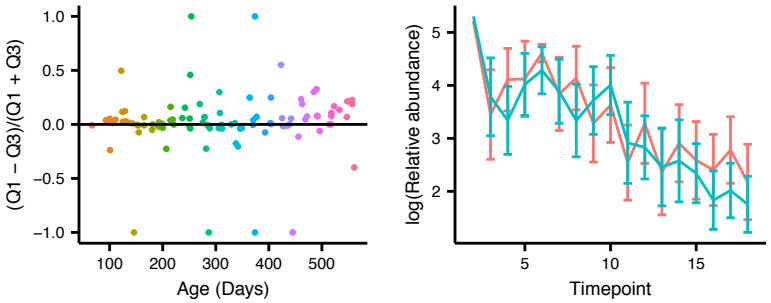


Figure Teddy_Q1_vs_Q3_supp

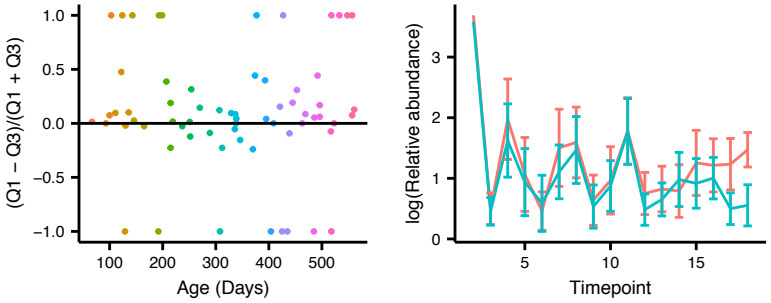
289709 – k__Bacteria; p__Proteobacteria; c__Gammaproteobacteria
o__Enterobacteriales; f__Enterobacteriaceae; g__; s__ – 0.018813852



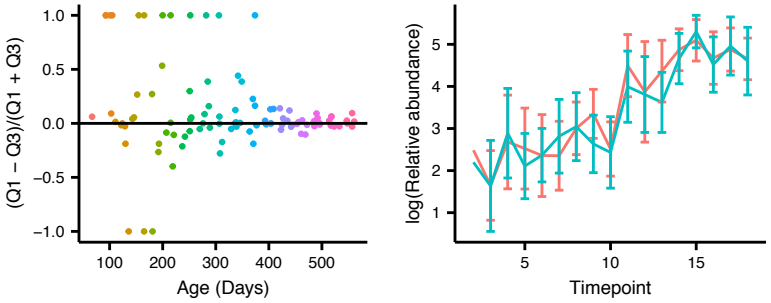
813479 – k__Bacteria; p__Actinobacteria; c__Actinobacteria
o__Bifidobacteriales; f__Bifidobacteriaceae; g__Bifidobacterium; s__ – 0.019154365



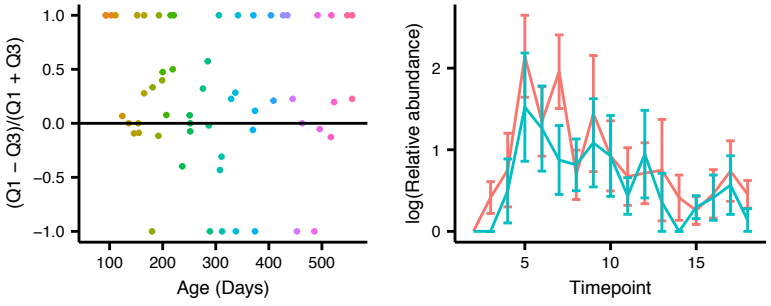
191999 – k__Bacteria; p__Firmicutes; c__Clostridia
o__Clostridiales; f__Lachnospiraceae; NA; NA – 0.019154365



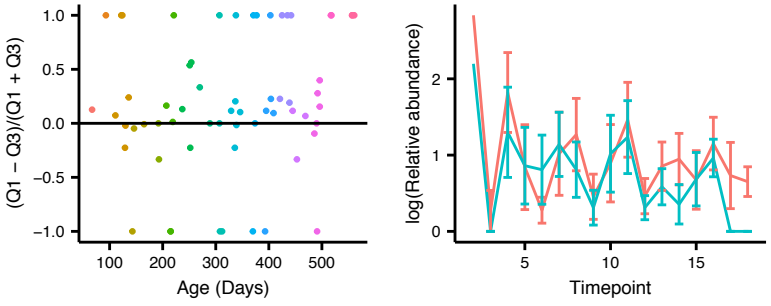
579608 – k__Bacteria; p__Firmicutes; c__Bacilli
o__Lactobacillales; f__Streptococcaceae; g__Streptococcus; s__ – 0.019271225



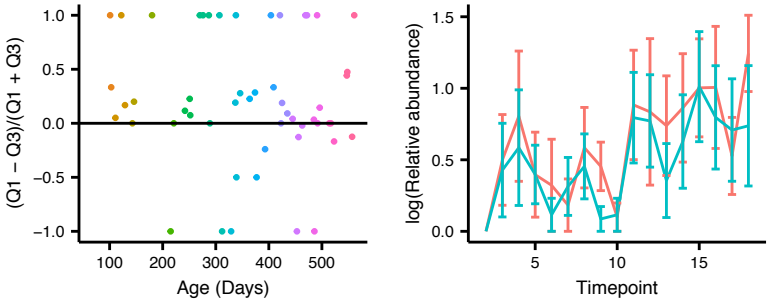
667570 – k__Bacteria; p__Proteobacteria; c__Gammaproteobacteria
o__Enterobacteriales; f__Enterobacteriaceae; g__; s__ – 0.022848523



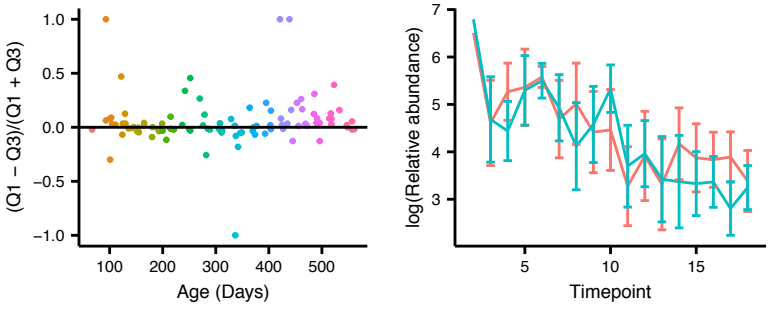
New.ReferenceOTU2576 – k__Bacteria; p__Firmicutes; c__Clostridia
o__Clostridiales; f__Lachnospiraceae; g__[Ruminococcus]; s__gnavus – 0.023776174



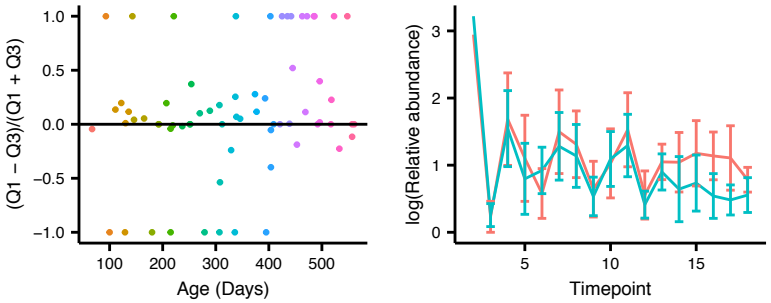
4306836 – k__Bacteria; p__Firmicutes; c__Bacilli
o__Lactobacillales; f__Streptococcaceae; g__Streptococcus; s__ – 0.026204072



235262 – k__Bacteria; p__Actinobacteria; c__Actinobacteria
o__Bifidobacteriales; f__Bifidobacteriaceae; g__Bifidobacterium; s__ – 0.026204072



New.ReferenceOTU2274 – k__Bacteria; p__Firmicutes; c__Clostridia
o__Clostridiales; f__Lachnospiraceae; g__[Ruminococcus]; s__gnavus – 0.026204072



591635 – k__Bacteria; p__Firmicutes; c__Clostridia
o__Clostridiales; f__Ruminococcaceae; g__Ruminococcus; s__ – 0.026759218

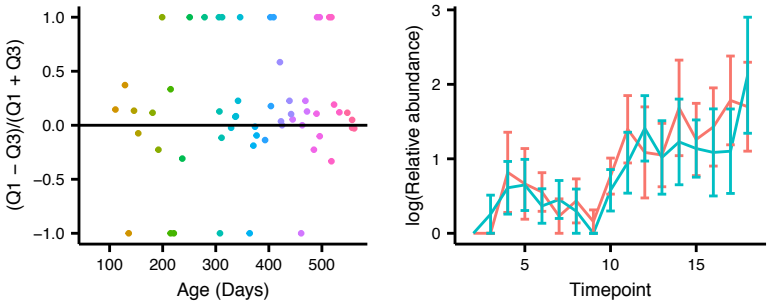
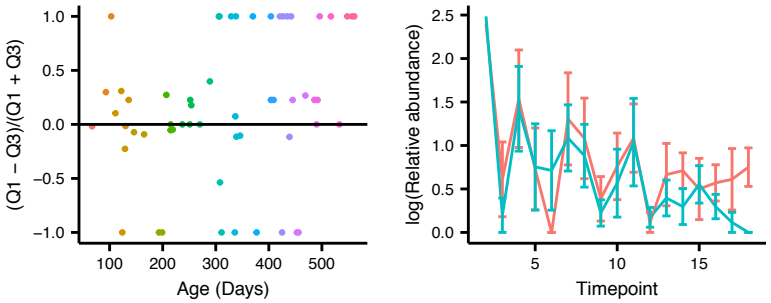


Figure Teddy_Q1_vs_Q3_supp

New.ReferenceOTU3135 – k__Bacteria; p__Firmicutes; c__Clostridia
o__Clostridiales; f__Lachnospiraceae; g__; s__ – 0.027901672



484304 – k__Bacteria; p__Actinobacteria; c__Actinobacteria
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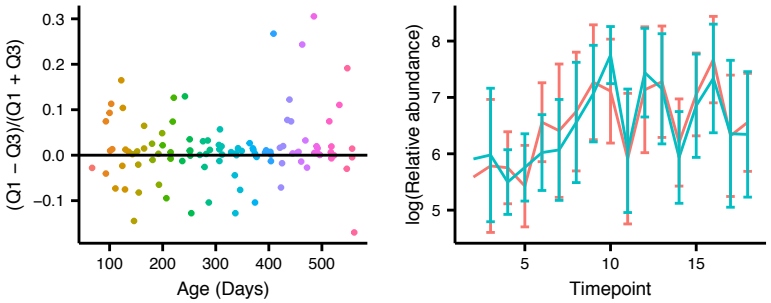
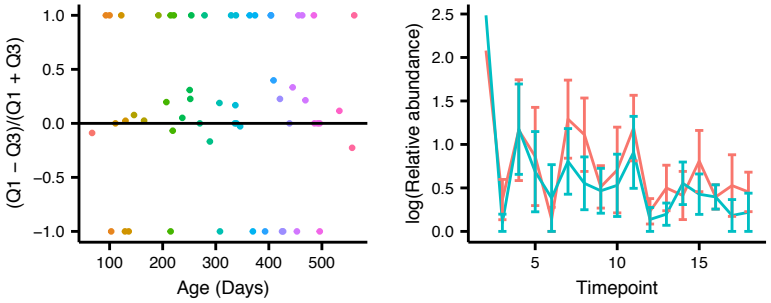
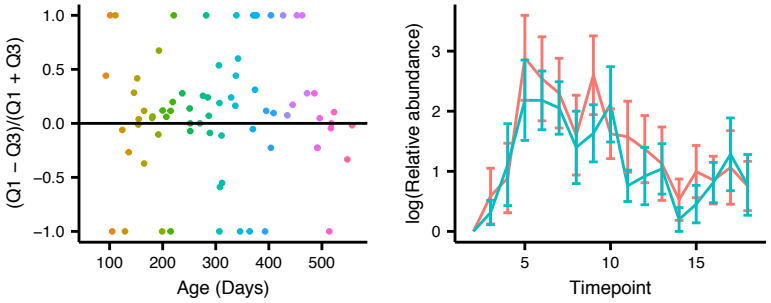


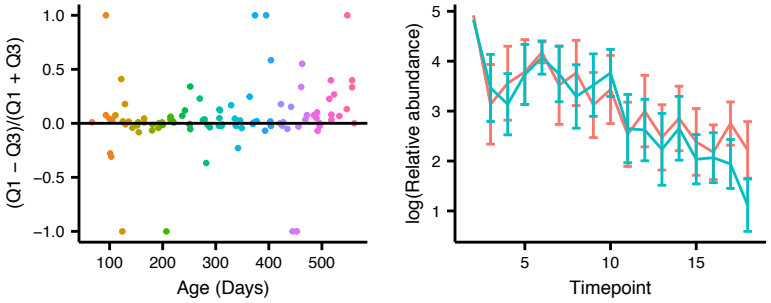
New.ReferenceOTU3990 – k__Bacteria; p__Firmicutes; c__Clostridia
o__Clostridiales; f__Lachnospiraceae; g__[Ruminococcus]; s__gnavus – 0.02931299



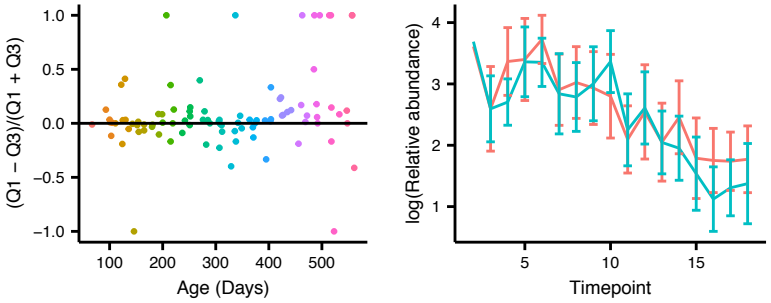
334459 – k__Bacteria; p__Proteobacteria; c__Gammaproteobacteria
o__Enterobacteriales; f__Enterobacteriaceae; g__; s__ – 0.0304457



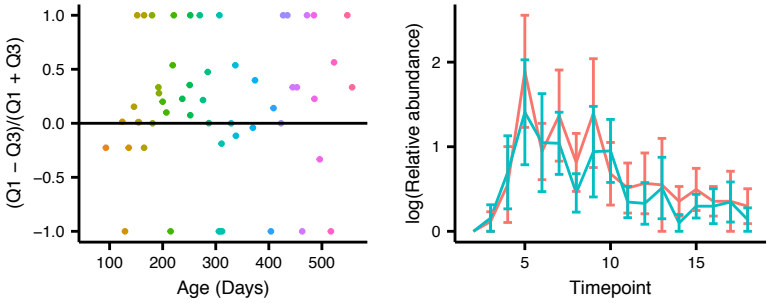
1142029 – k__Bacteria; p__Actinobacteria; c__Actinobacteria
o__Bifidobacteriales; f__Bifidobacteriaceae; g__Bifidobacterium; s__ – 0.031775561



370225 – k__Bacteria; p__Actinobacteria; c__Actinobacteria
o__Bifidobacteriales; f__Bifidobacteriaceae; g__Bifidobacterium; s__adolescentis – 0.034941842



4111715 – k__Bacteria; p__Proteobacteria; c__Gammaproteobacteria
o__Enterobacteriales; f__Enterobacteriaceae; g__; s__ – 0.040215536



514523 – k__Bacteria; p__Firmicutes; c__Clostridia
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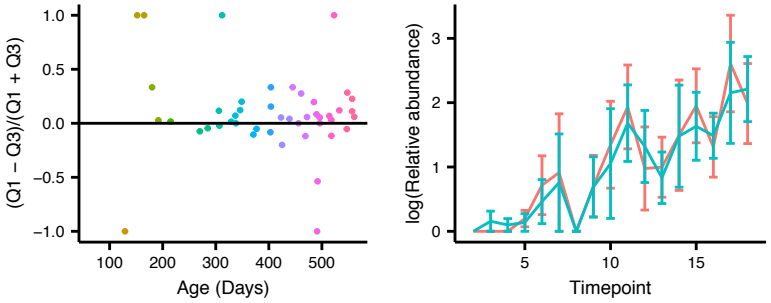
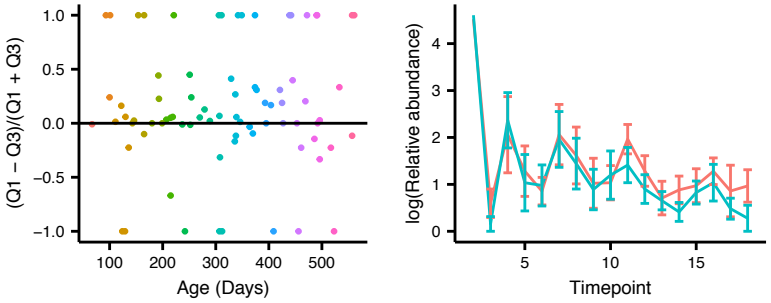


3376513 – k__Bacteria; p__Firmicutes; c__Clostridia
o__Clostridiales; f__Lachnospiraceae; g__[Ruminococcus]; s__gnavus – 0.047441106



182133 – k__Bacteria; p__Firmicutes; c__Clostridia
o__Clostridiales; f__Lachnospiraceae; g__Blautia; s__ – 0.000126849

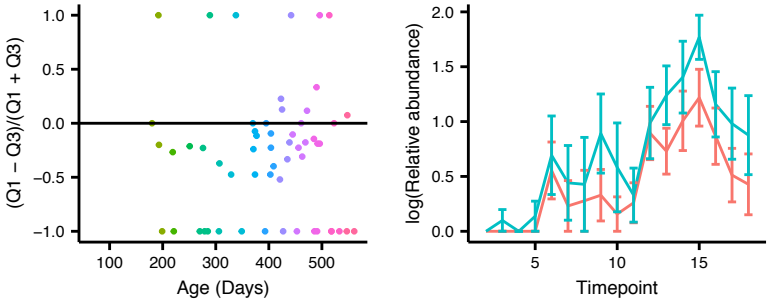


Figure Teddy_Q1_vs_Q3_supp

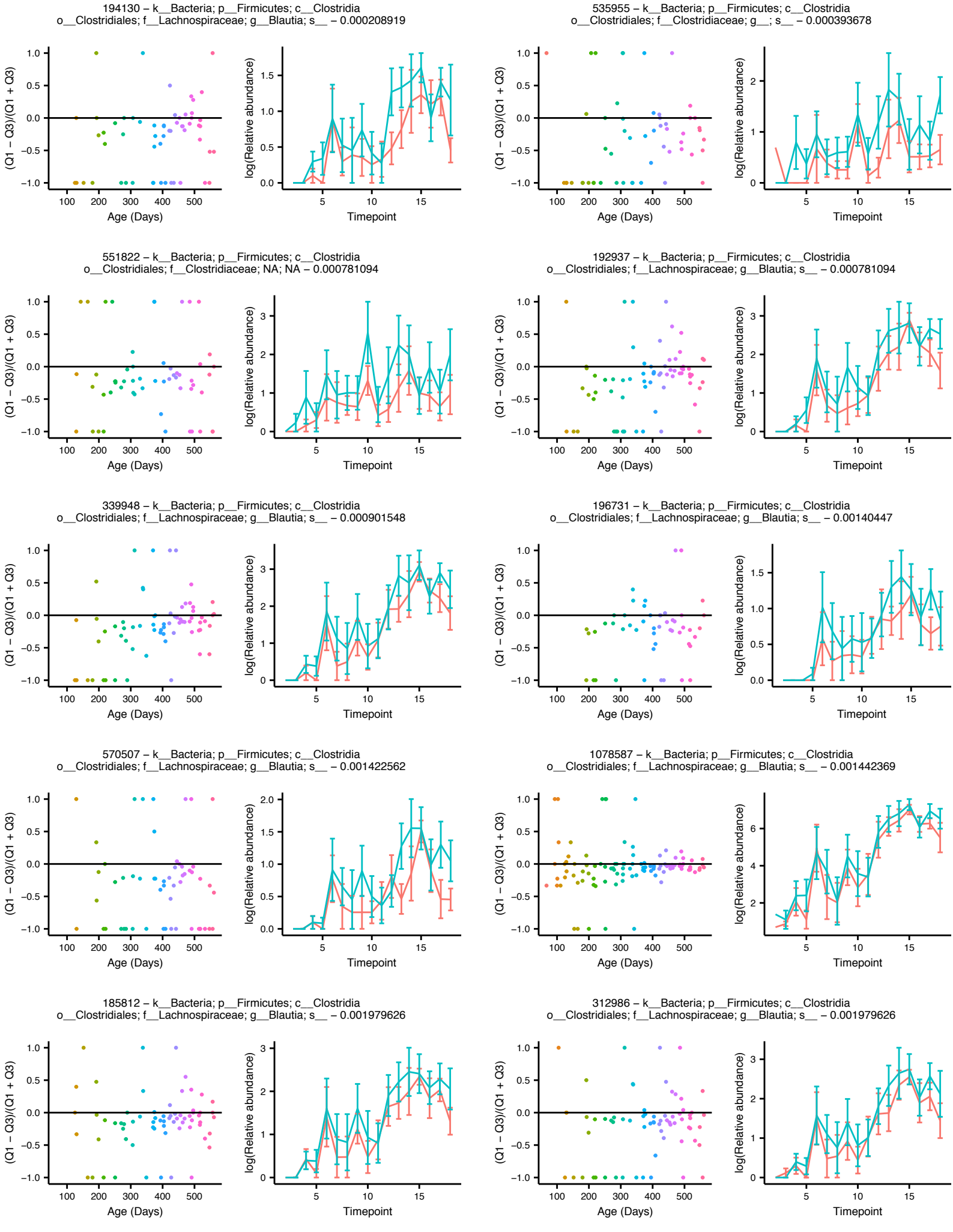


Figure Teddy_Q1_vs_Q3_supp

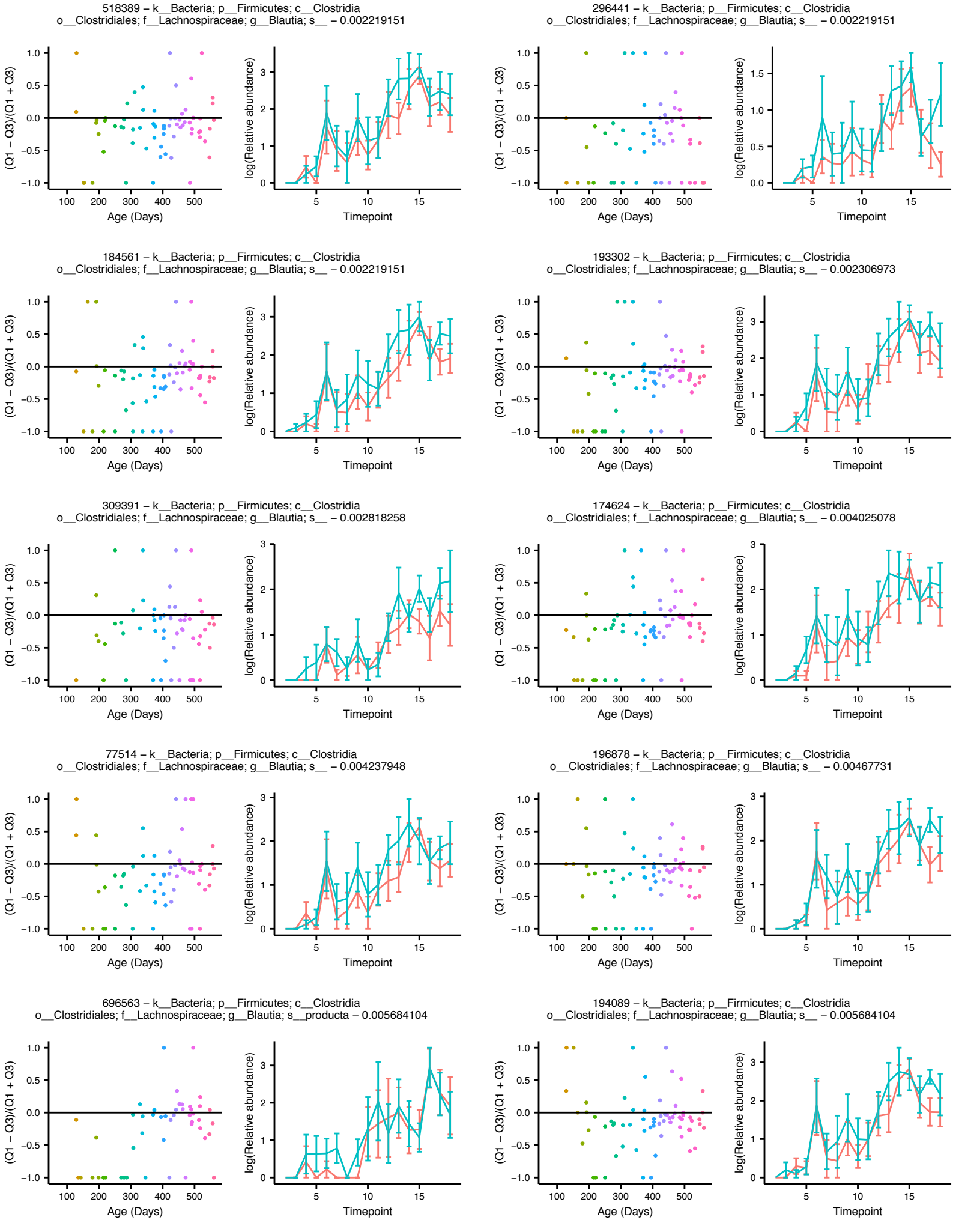
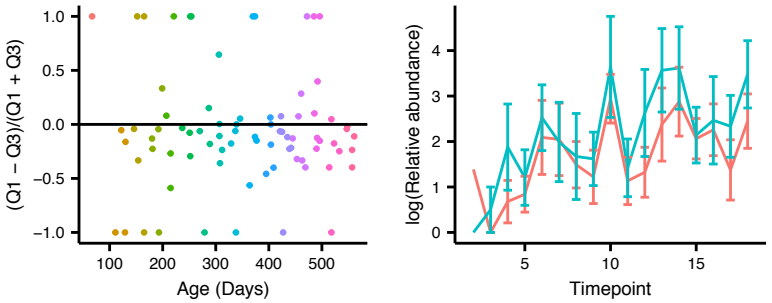
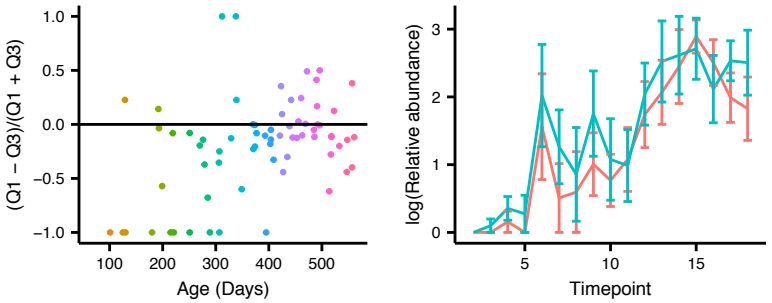


Figure Teddy_Q1_vs_Q3_supp

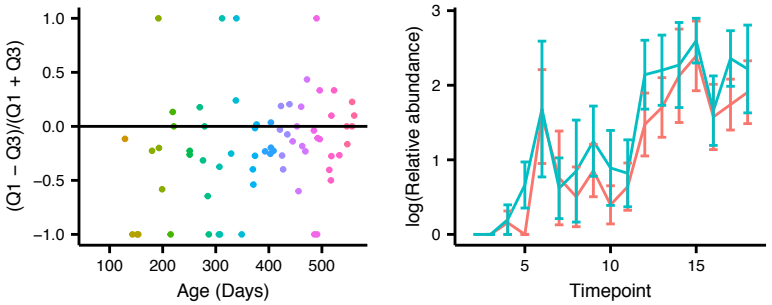
628226 – k__Bacteria; p__Firmicutes; c__Clostridia
o__Clostridiales; f__Clostridiaceae; g__; s__ – 0.005955668



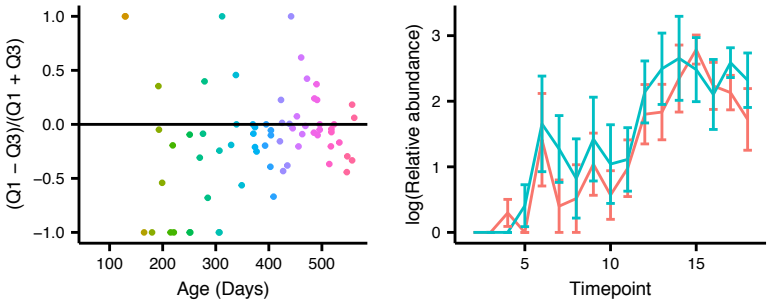
302683 – k__Bacteria; p__Firmicutes; c__Clostridia
o__Clostridiales; f__Lachnospiraceae; g__Blautia; s__ – 0.006104403



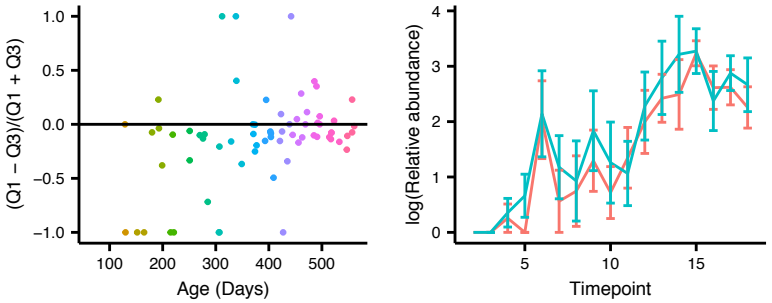
198059 – k__Bacteria; p__Firmicutes; c__Clostridia
o__Clostridiales; f__Lachnospiraceae; g__Blautia; s__ – 0.006989366



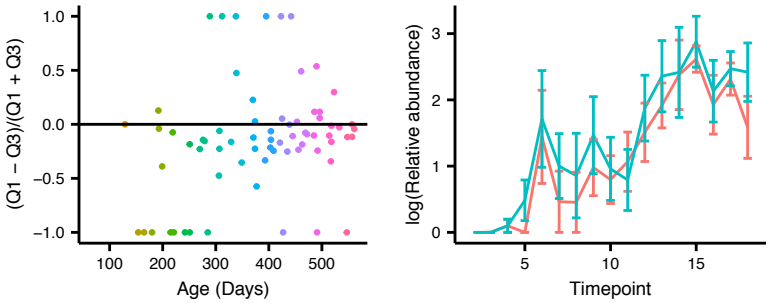
532203 – k__Bacteria; p__Firmicutes; c__Clostridia
o__Clostridiales; f__Lachnospiraceae; g__Blautia; s__ – 0.007074771



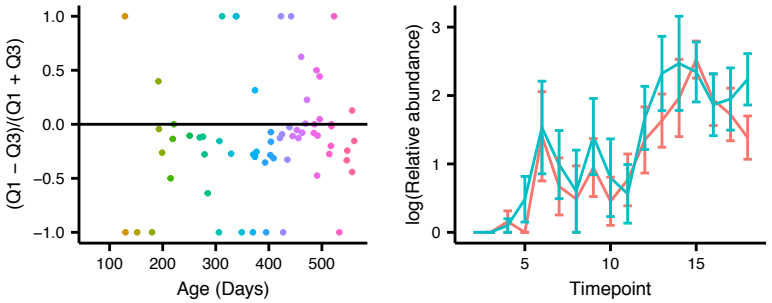
186883 – k__Bacteria; p__Firmicutes; c__Clostridia
o__Clostridiales; f__Lachnospiraceae; g__Blautia; s__ – 0.007765238



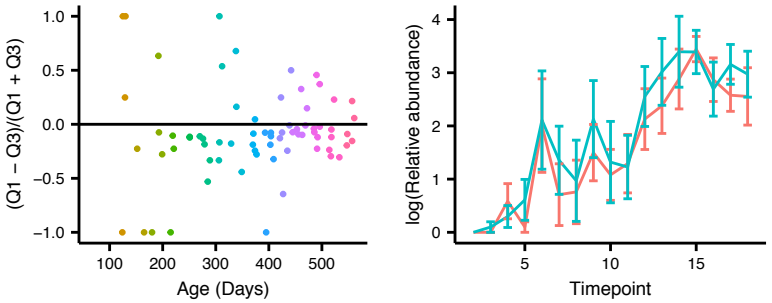
193744 – k__Bacteria; p__Firmicutes; c__Clostridia
o__Clostridiales; f__Lachnospiraceae; g__Blautia; s__ – 0.010001395



3924208 – k__Bacteria; p__Firmicutes; c__Clostridia
o__Clostridiales; f__Lachnospiraceae; g__Blautia; s__ – 0.010001395



362568 – k__Bacteria; p__Firmicutes; c__Clostridia
o__Clostridiales; f__Lachnospiraceae; g__Blautia; s__ – 0.011365948



68845 – k__Bacteria; p__Firmicutes; c__Clostridia
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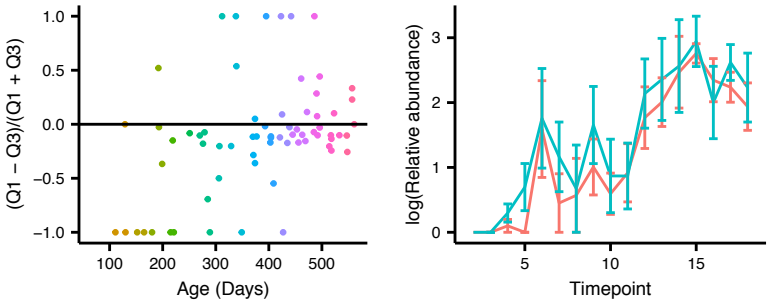

363348 – k__Bacteria; p__Firmicutes; c__Clostridia
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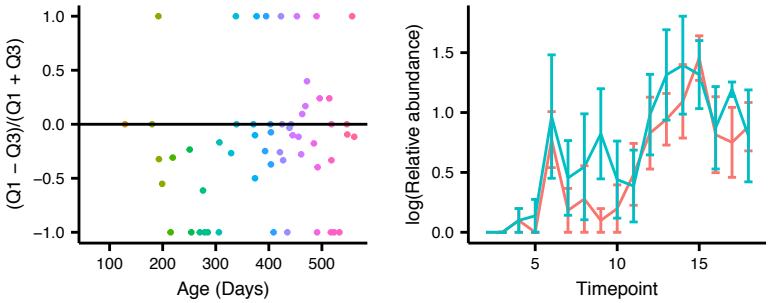
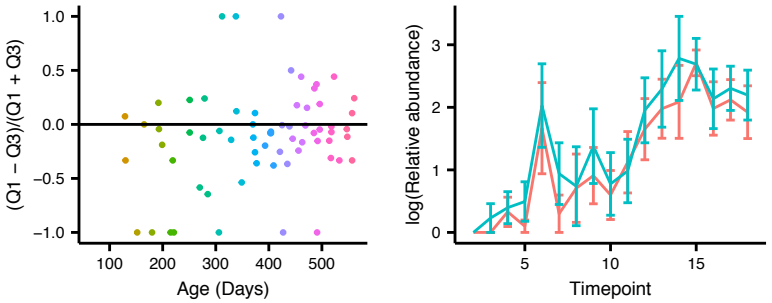
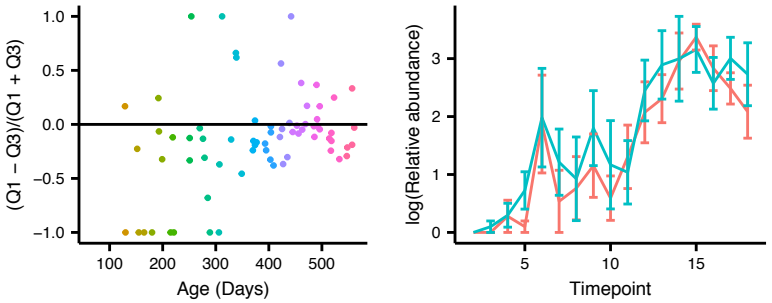


Figure Teddy_Q1_vs_Q3_supp

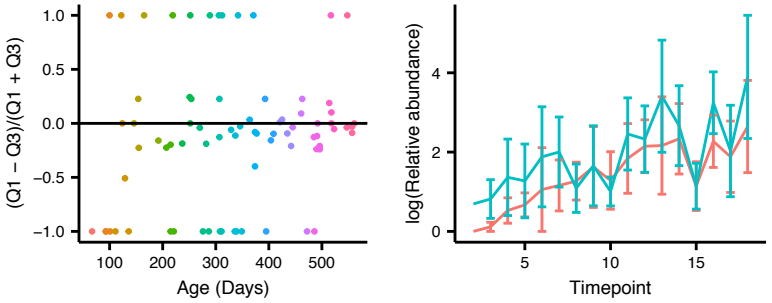
196942 – k__Bacteria; p__Firmicutes; c__Clostridia
o__Clostridiales; f__Lachnospiraceae; g__; s__ – 0.012425239



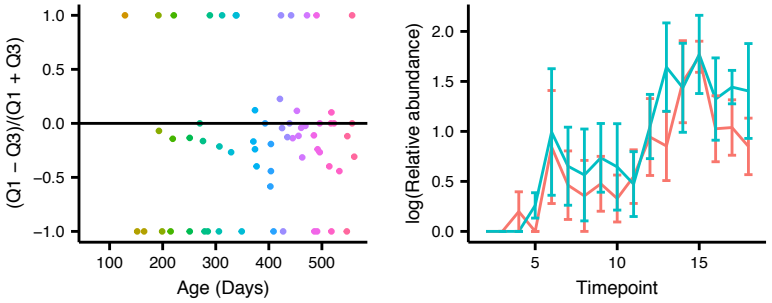
230741 – k__Bacteria; p__Firmicutes; c__Clostridia
o__Clostridiales; f__Lachnospiraceae; g__Blautia; s__ – 0.014346912



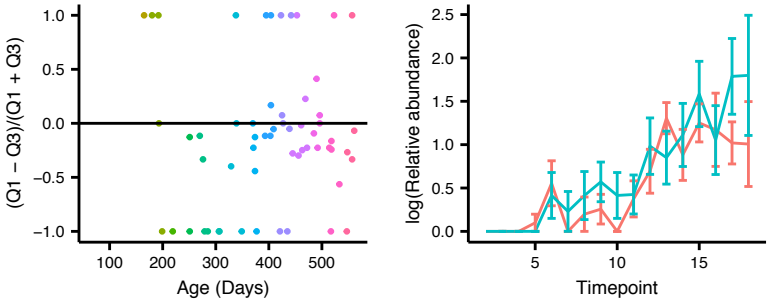
New.CleanUp.ReferenceOTU1267548 – k__Bacteria; p__Firmicutes; c__Clostridia
o__Clostridiales; f__Clostridiaceae; g__; s__ – 0.016238199



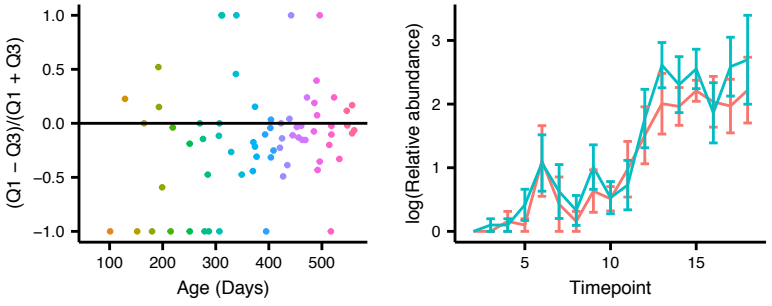
367790 – k__Bacteria; p__Firmicutes; c__Clostridia
o__Clostridiales; f__Lachnospiraceae; g__Blautia; s__ – 0.020590685



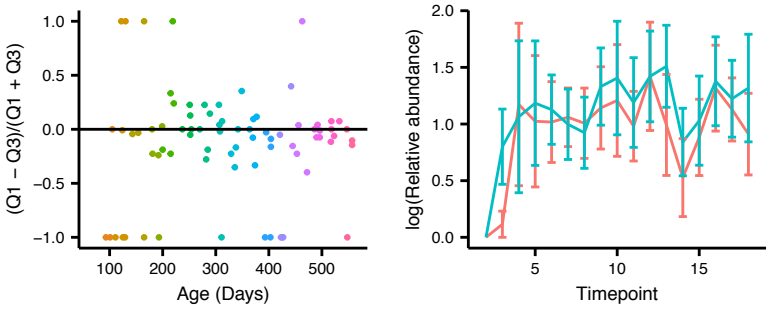
332929 – k__Bacteria; p__Firmicutes; c__Clostridia
o__Clostridiales; f__Lachnospiraceae; NA; NA – 0.021783656



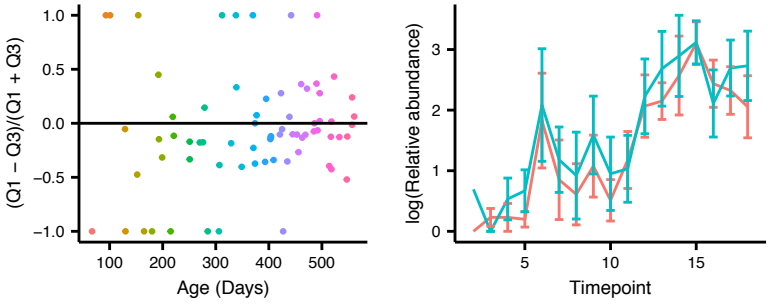
193484 – k__Bacteria; p__Firmicutes; c__Clostridia
o__Clostridiales; f__; g__; s__ – 0.026899706



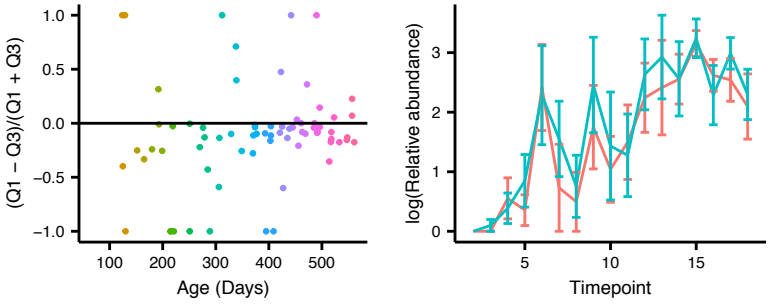
825808 – k__Bacteria; p__Actinobacteria; c__Actinobacteria
o__Bifidobacteriales; f__Bifidobacteriaceae; g__Bifidobacterium; s__ – 0.0304457



3014078 – k__Bacteria; p__Firmicutes; c__Clostridia
o__Clostridiales; f__Lachnospiraceae; g__Blautia; s__ – 0.0304457



3768338 – k__Bacteria; p__Firmicutes; c__Clostridia
o__Clostridiales; f__Lachnospiraceae; g__Blautia; s__ – 0.0304457



199147 – k__Bacteria; p__Firmicutes; c__Clostridia
o__Clostridiales; f__Lachnospiraceae; g__Blautia; s__ – 0.030896284