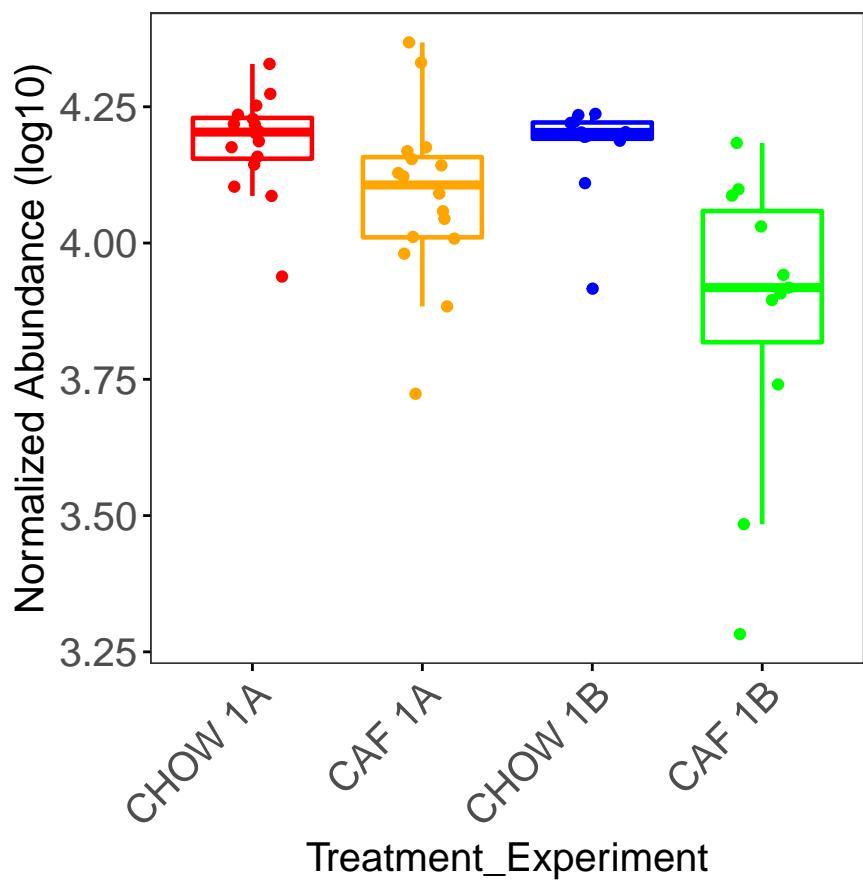
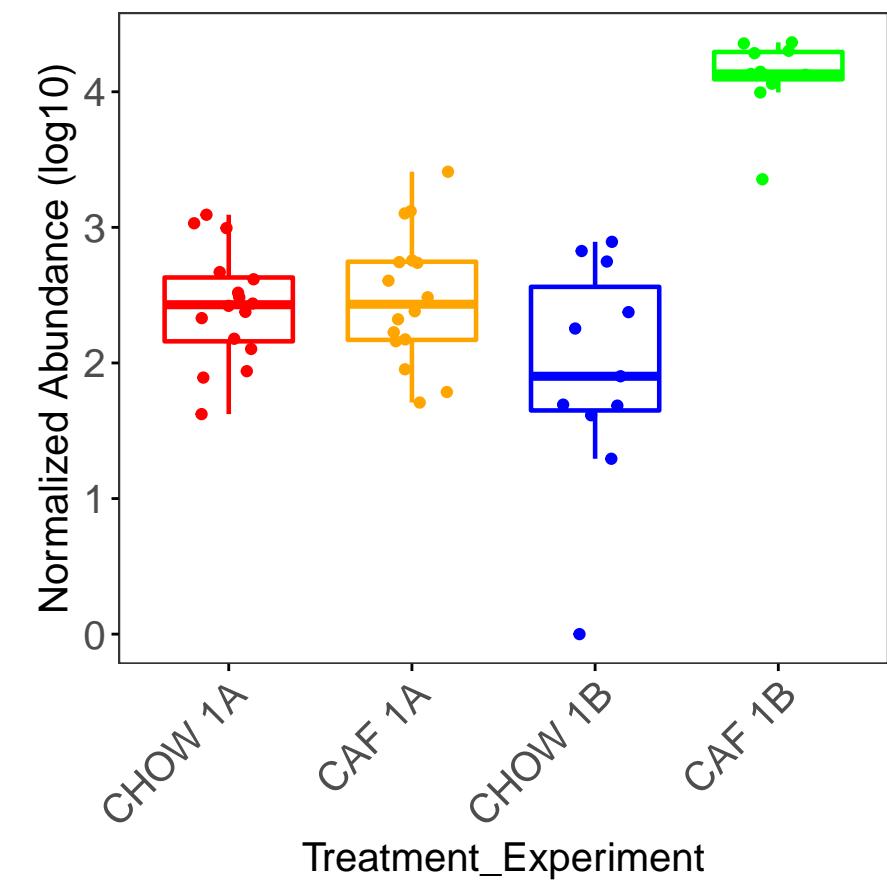


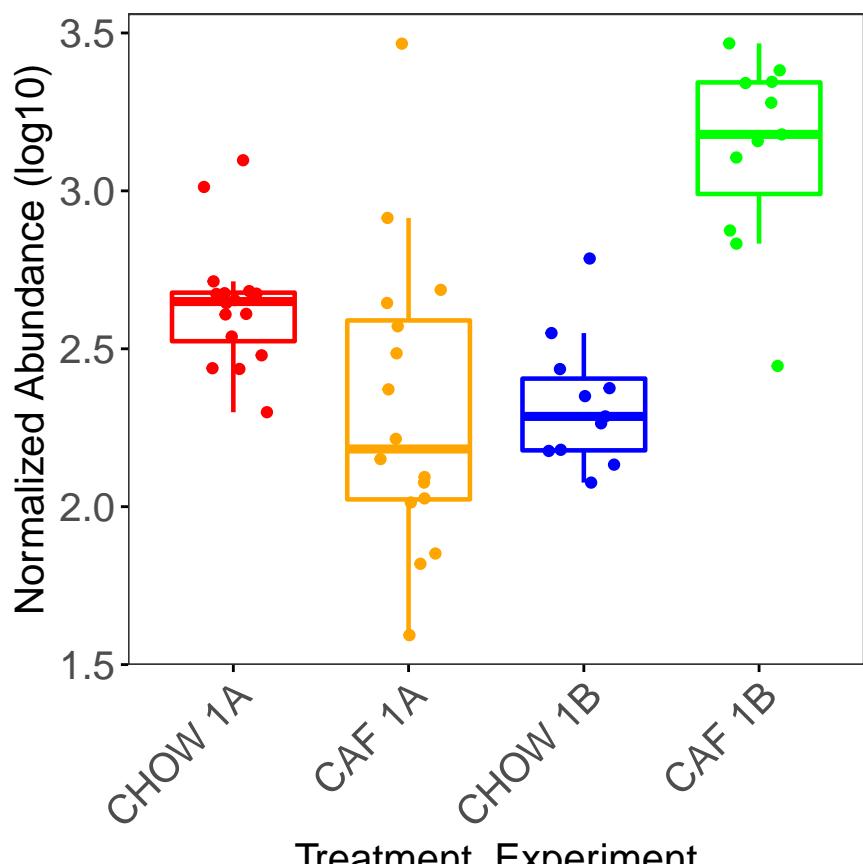
Bacteroides.D_2_Bacteroidia
Experiment 1A FDR = 0.06045
Experiment 1B FDR = 0.00038



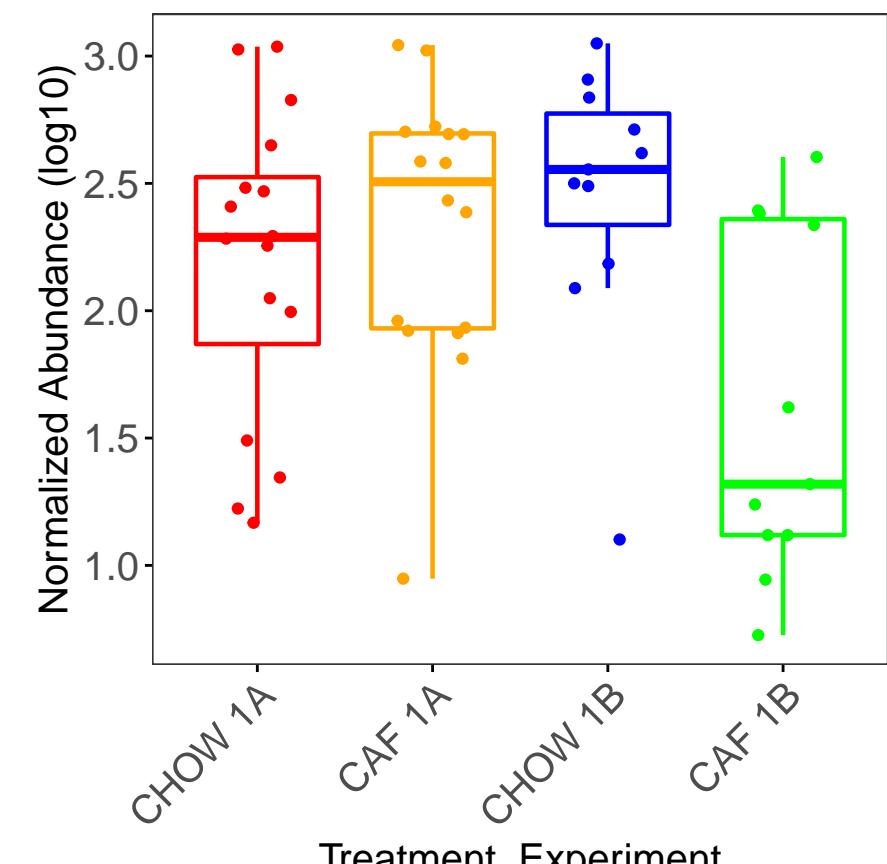
Firmicutes.D_2_Erysipelotrichia
Experiment 1A FDR = 0.94934
Experiment 1B FDR = 1e-05



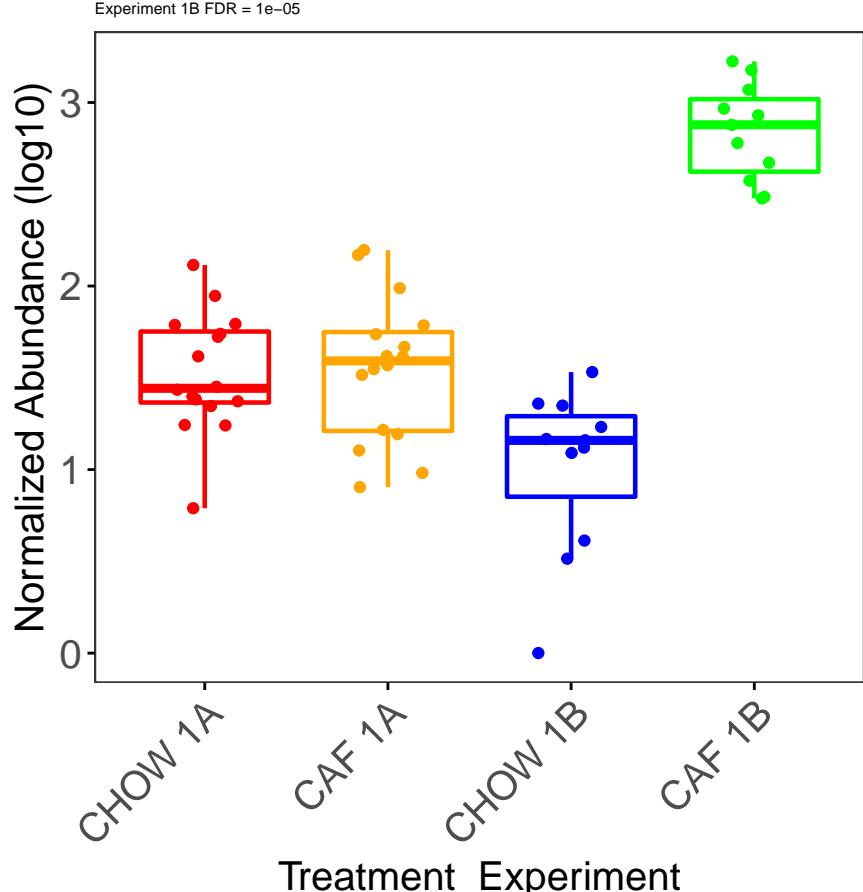
Proteobacteria.D_2_Gammaproteobacteria
Experiment 1A FDR = 0.06045
Experiment 1B FDR = 4e-05



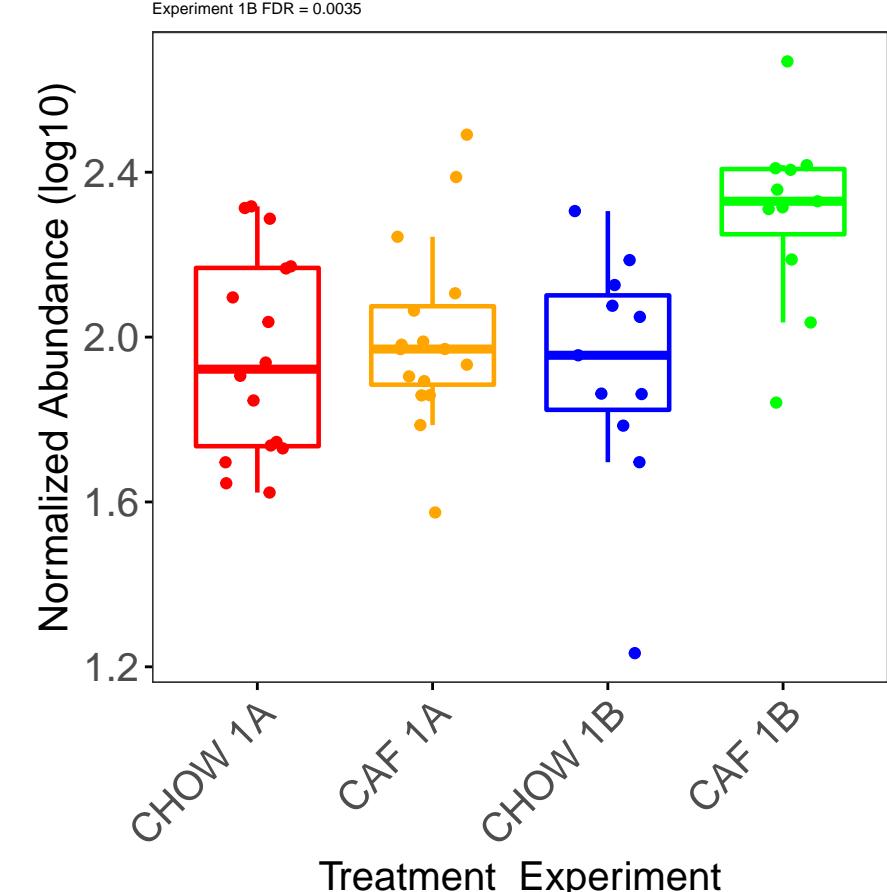
Tenericutes.D_2_Mollicutes
Experiment 1A FDR = 0.94017
Experiment 1B FDR = 0.01422

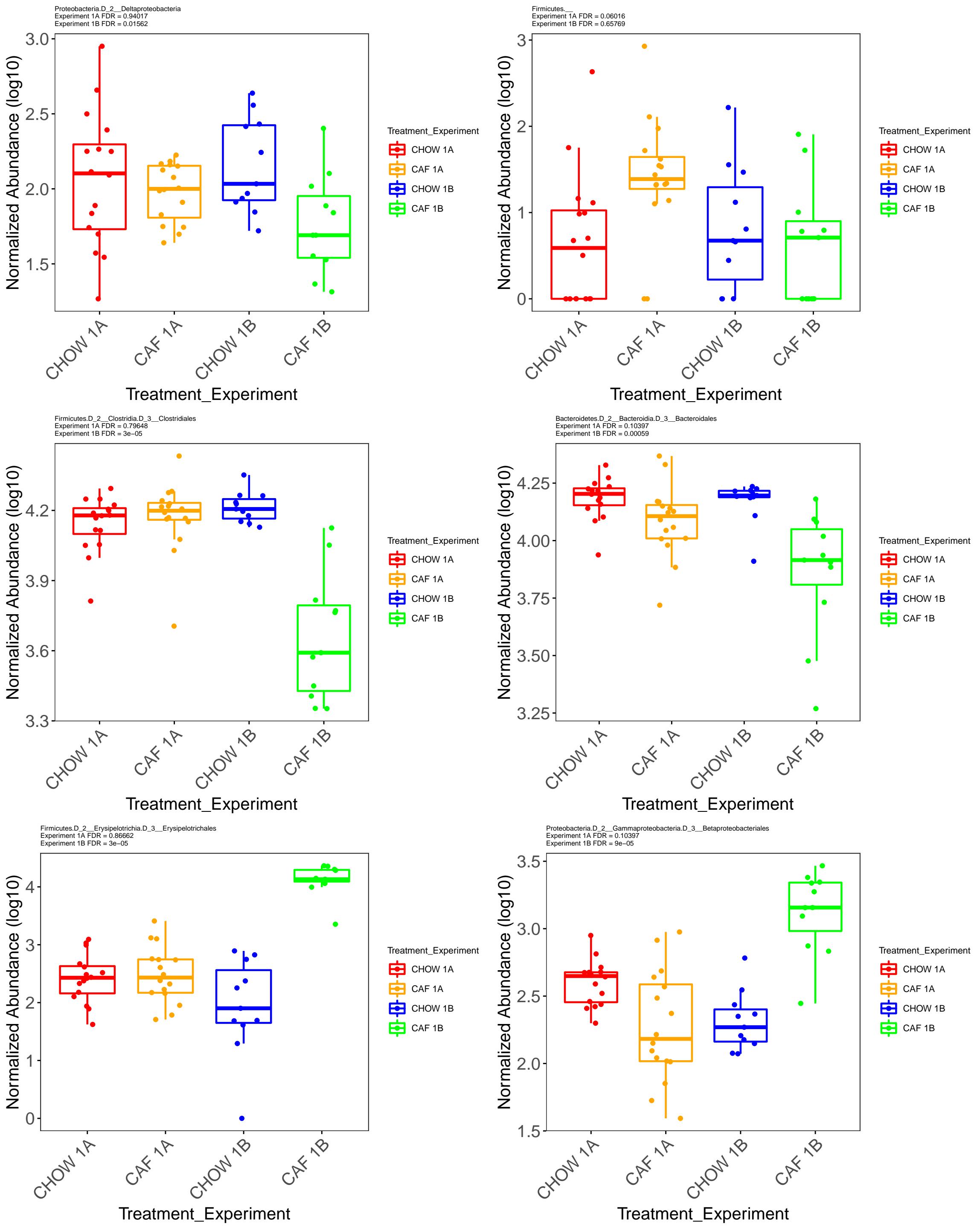


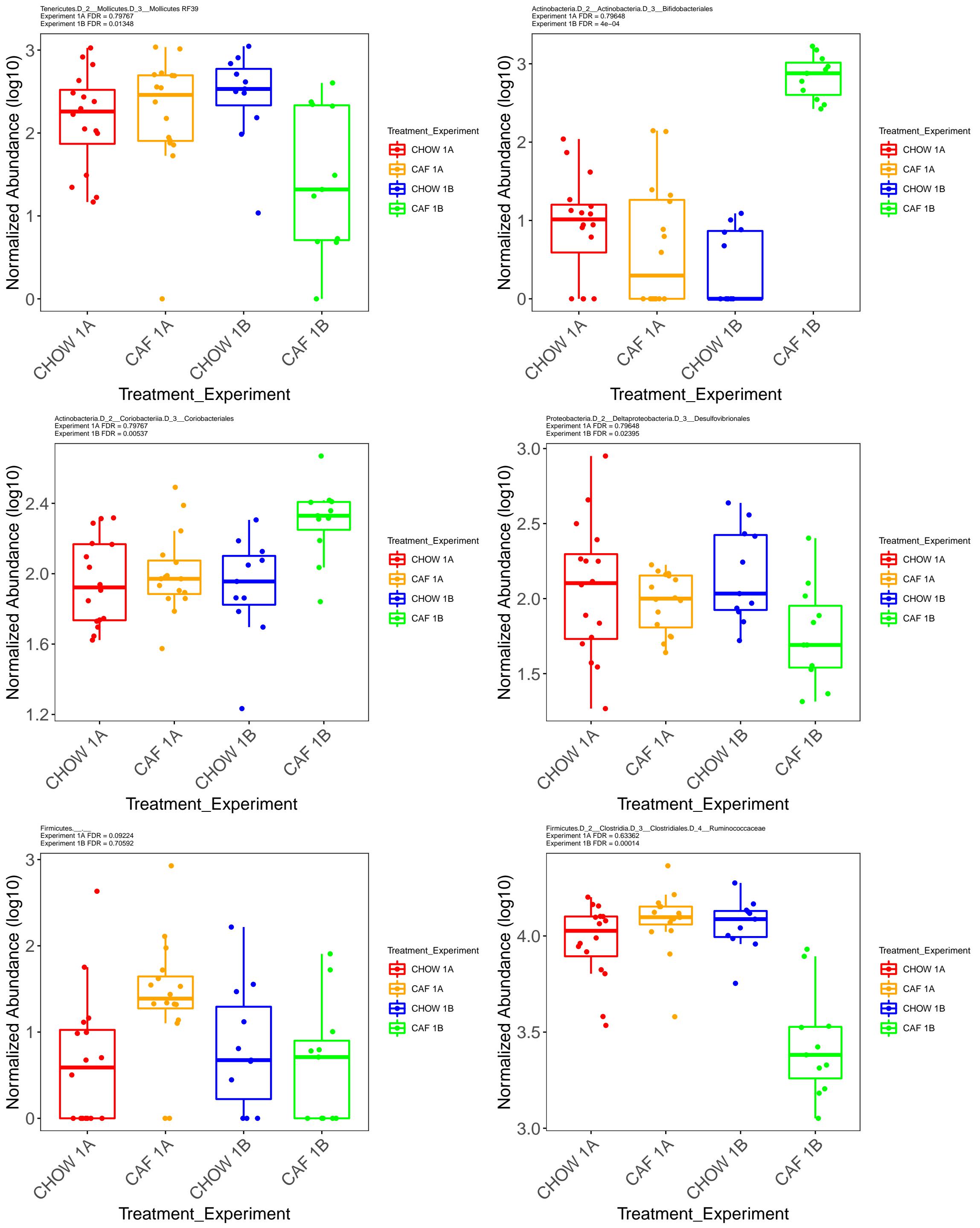
Actinobacteria.D_2_Actinobacteria
Experiment 1A FDR = 0.95556
Experiment 1B FDR = 1e-05

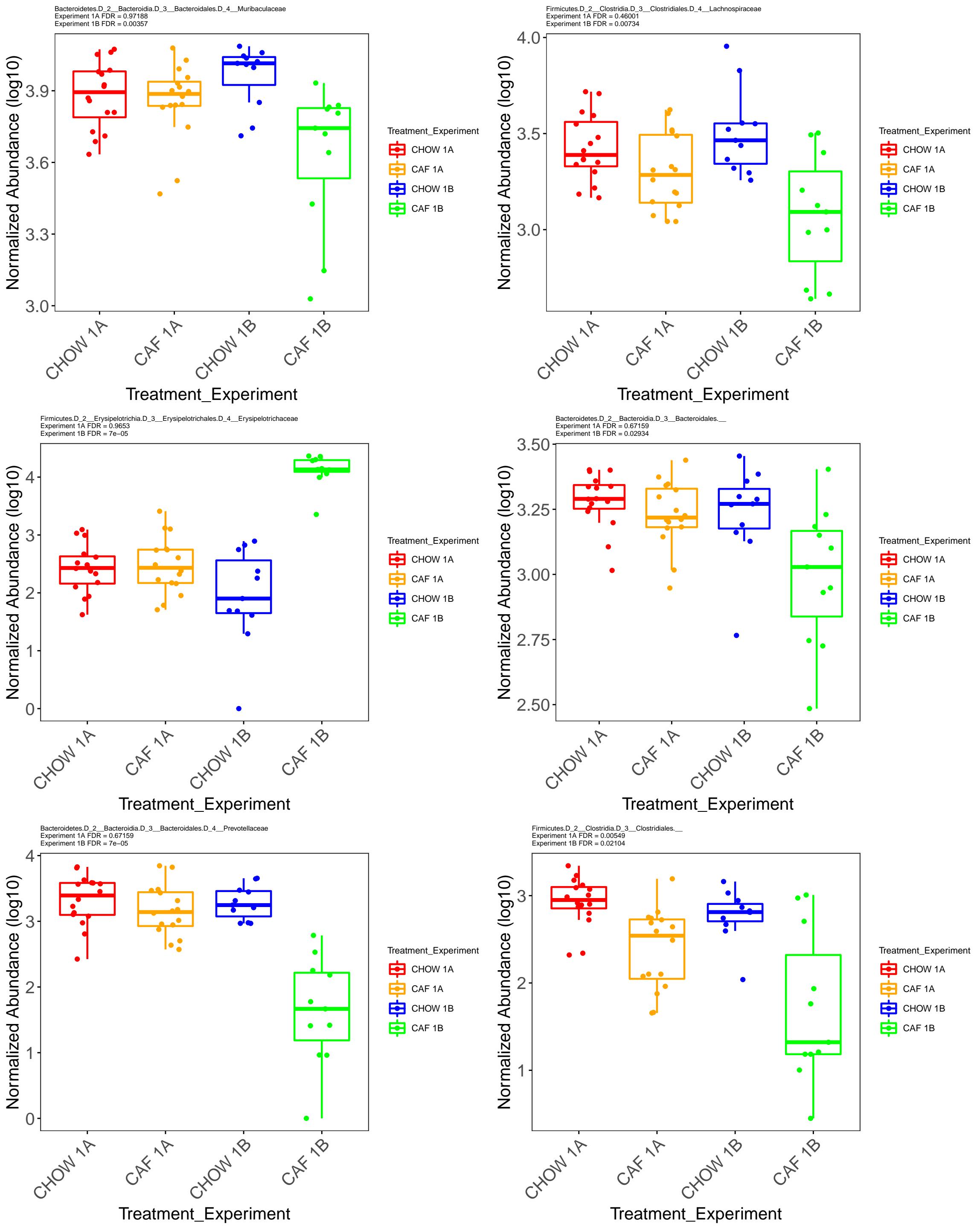


Actinobacteria.D_2_Coriobacteriia
Experiment 1A FDR = 0.94017
Experiment 1B FDR = 0.0035

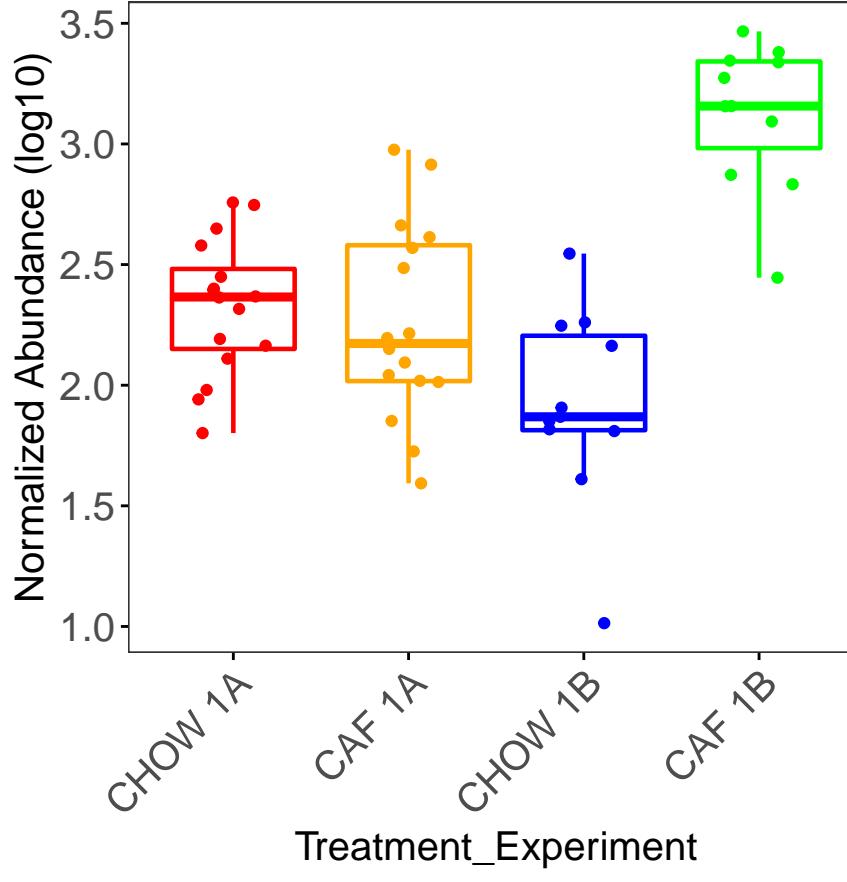




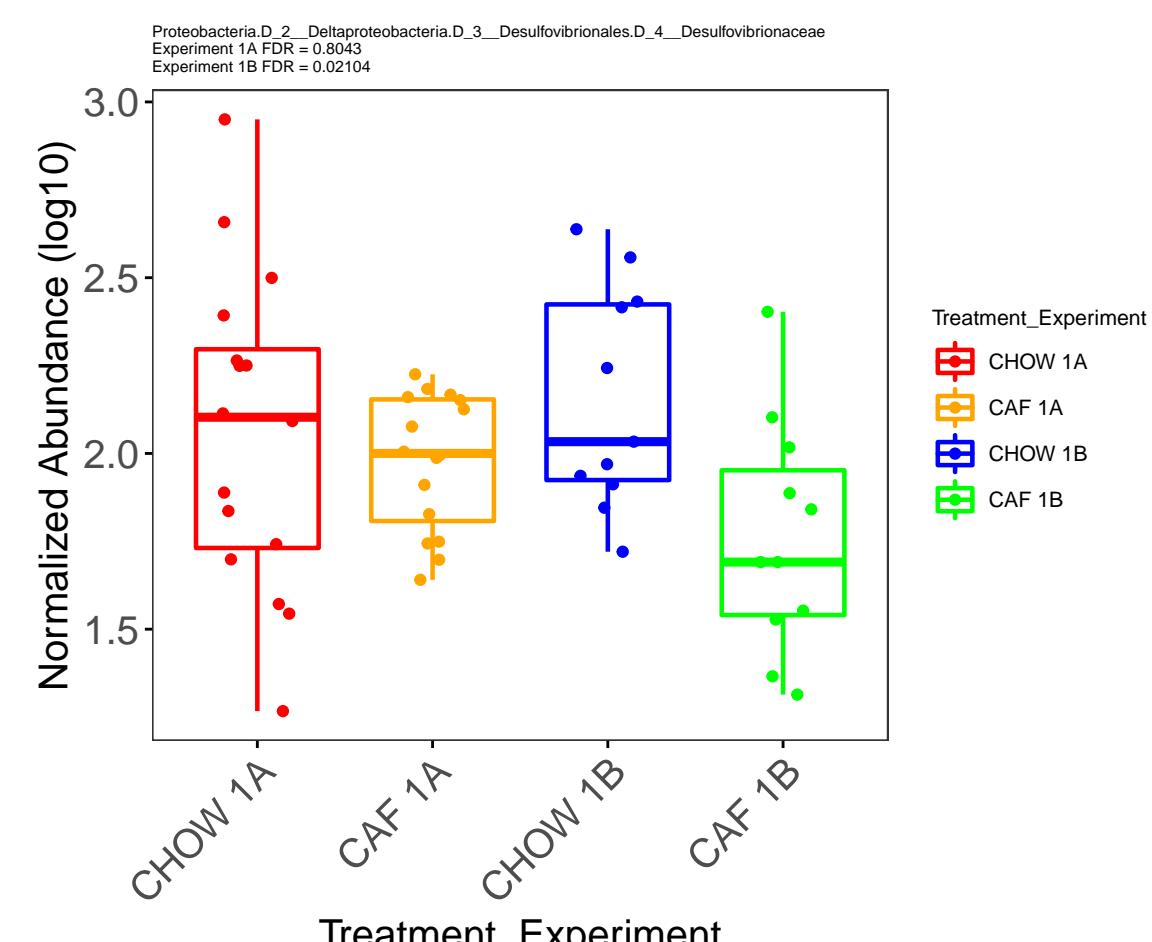
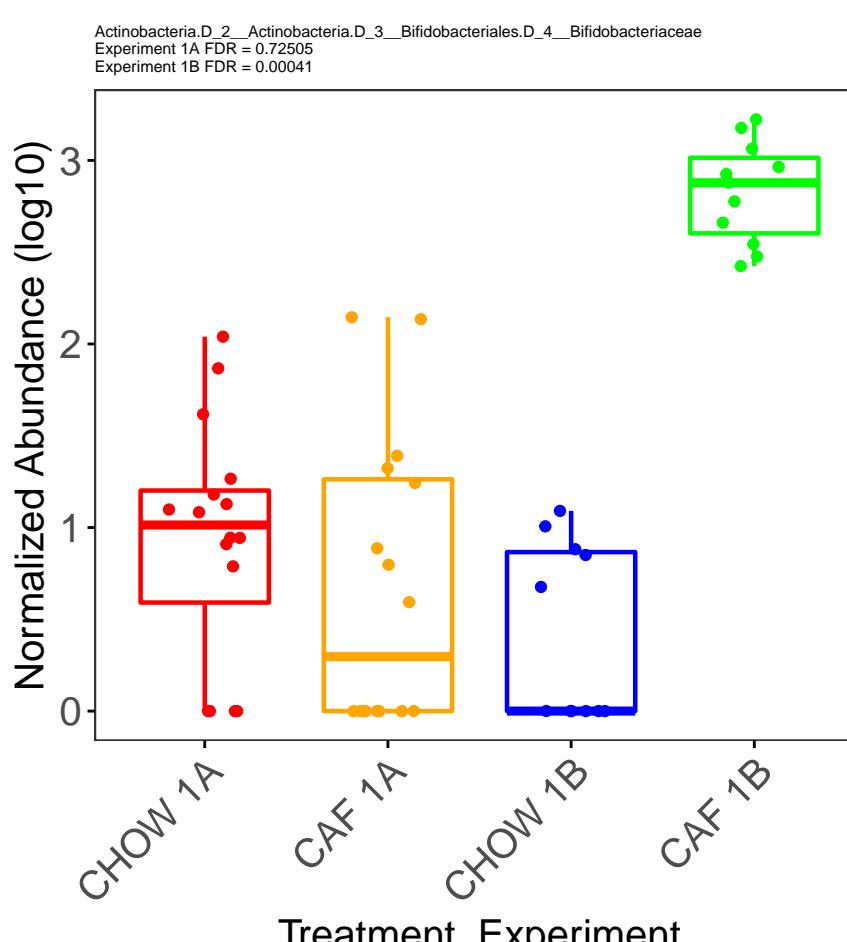
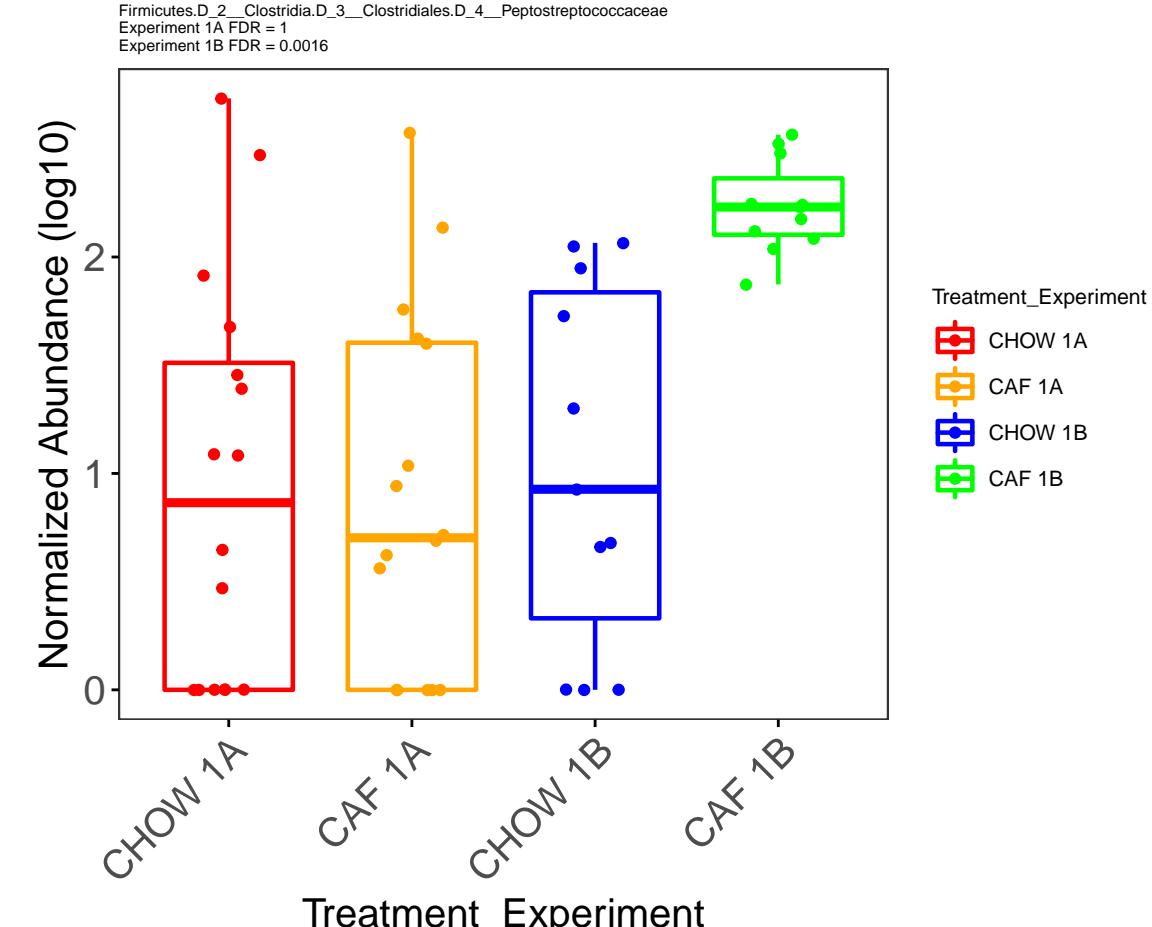
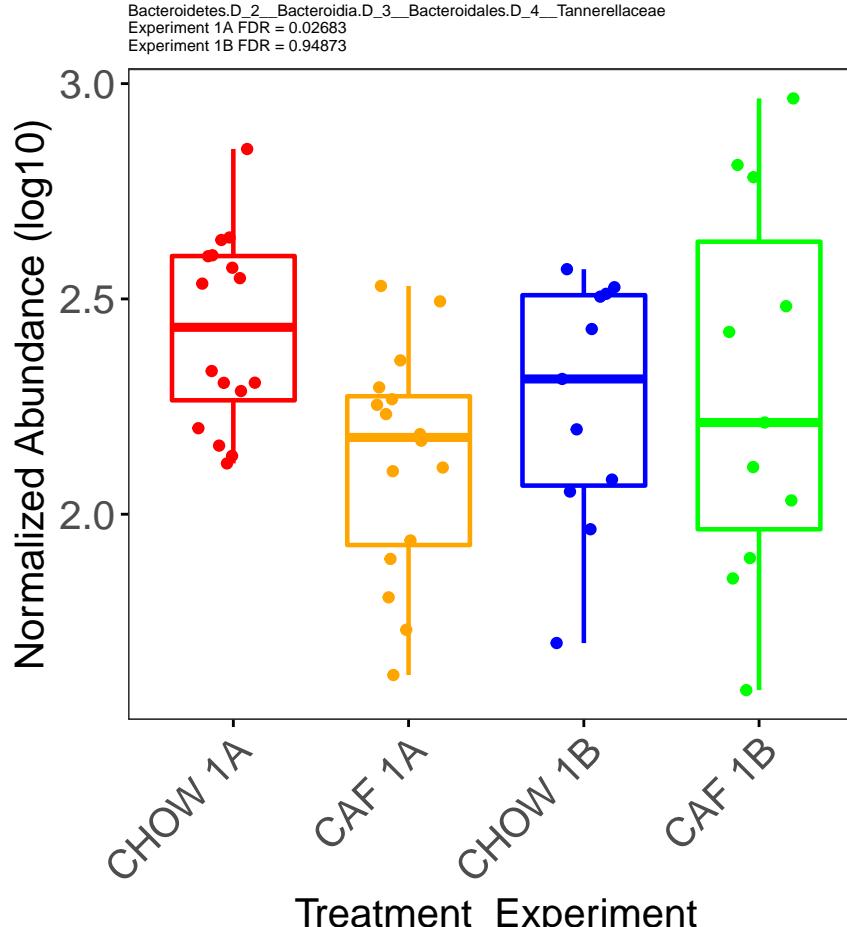
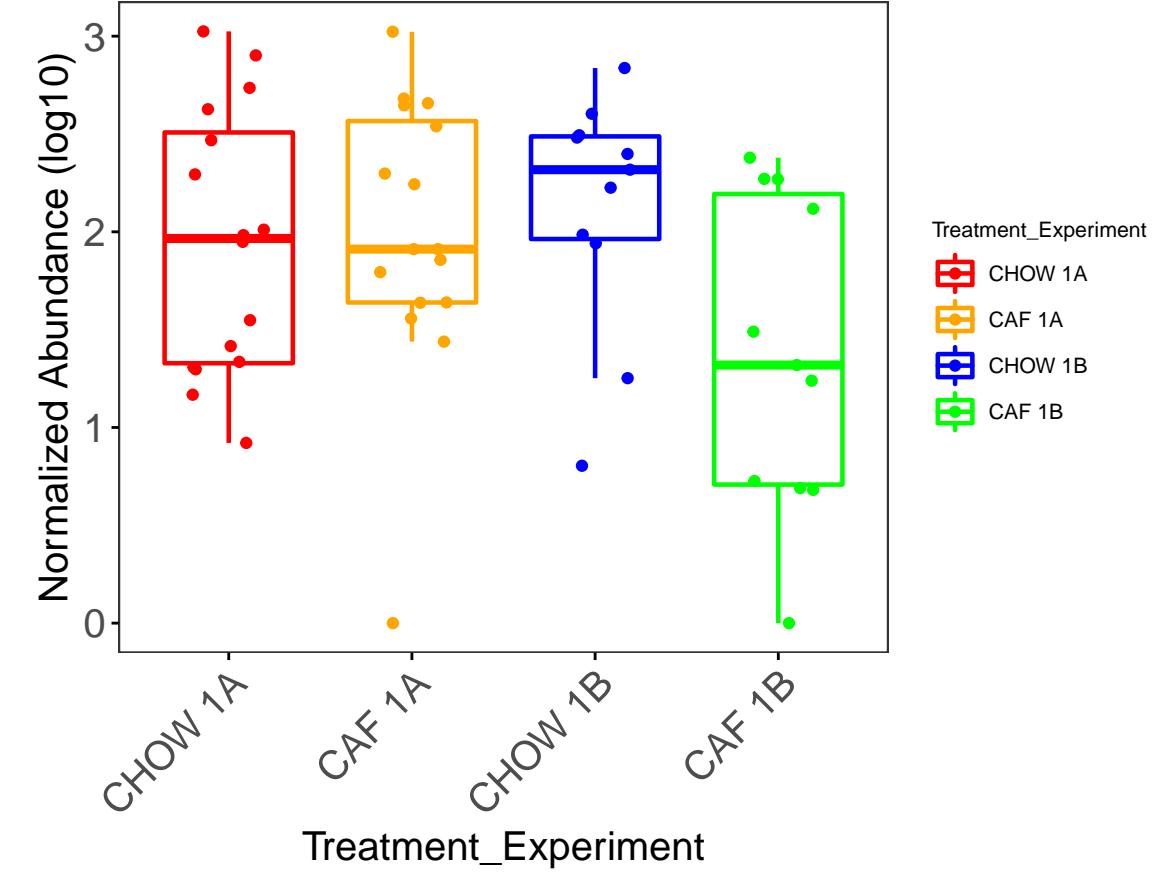


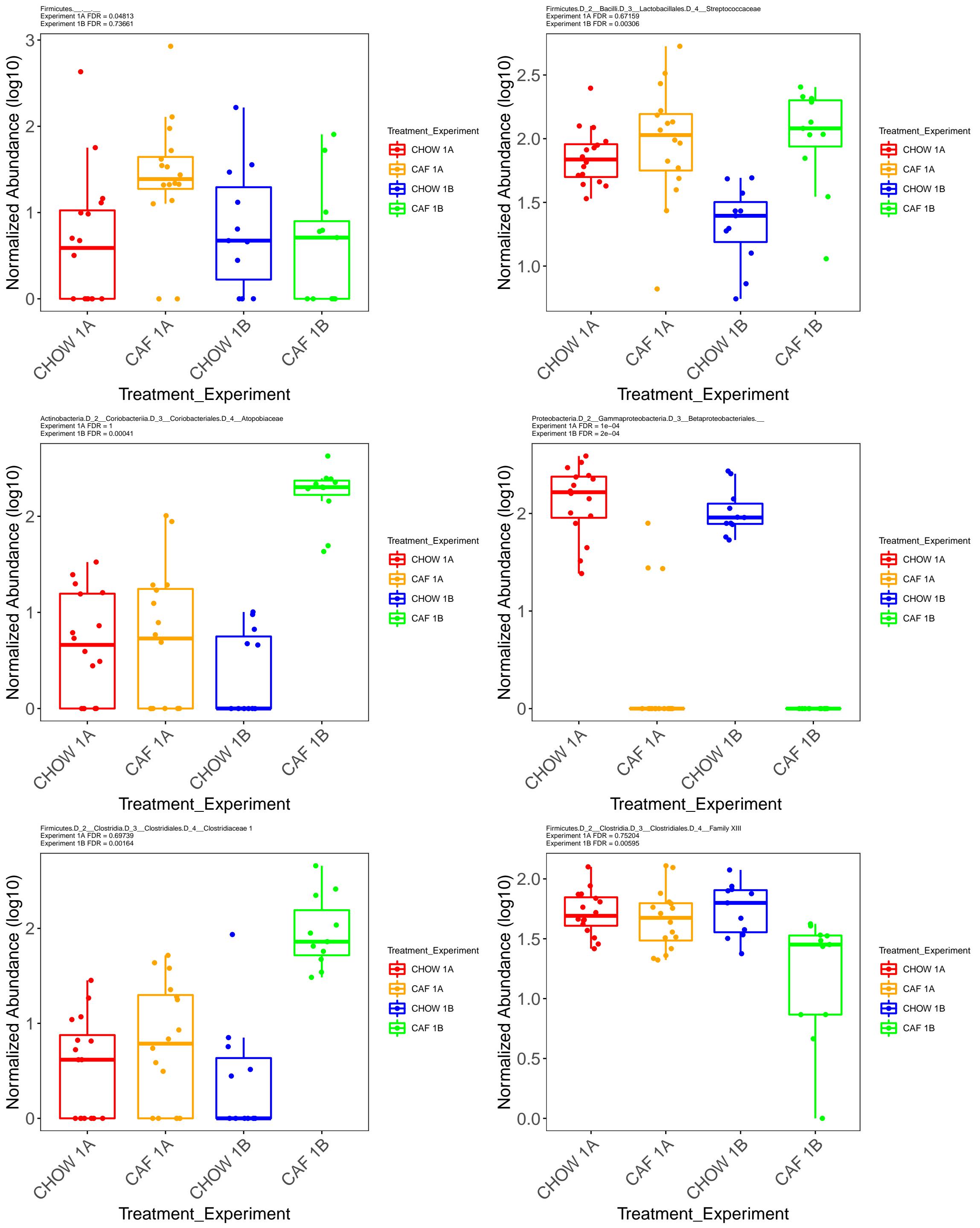


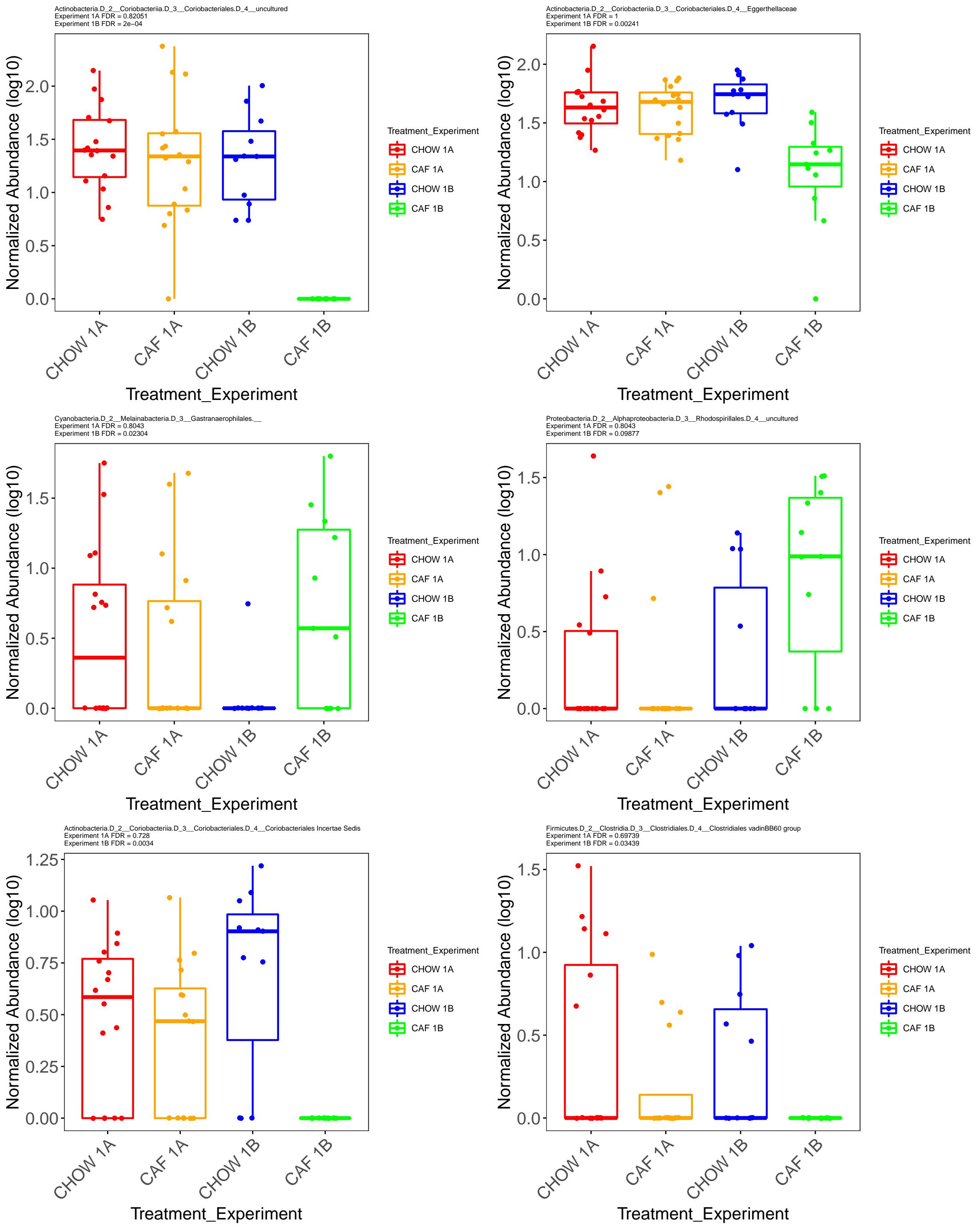
Proteobacteria.D_2_Gammaproteobacteria.D_3_Betaproteobacteriales.D_4_Burkholderiaceae
Experiment 1A FDR = 0.82051
Experiment 1B FDR = 9e-05

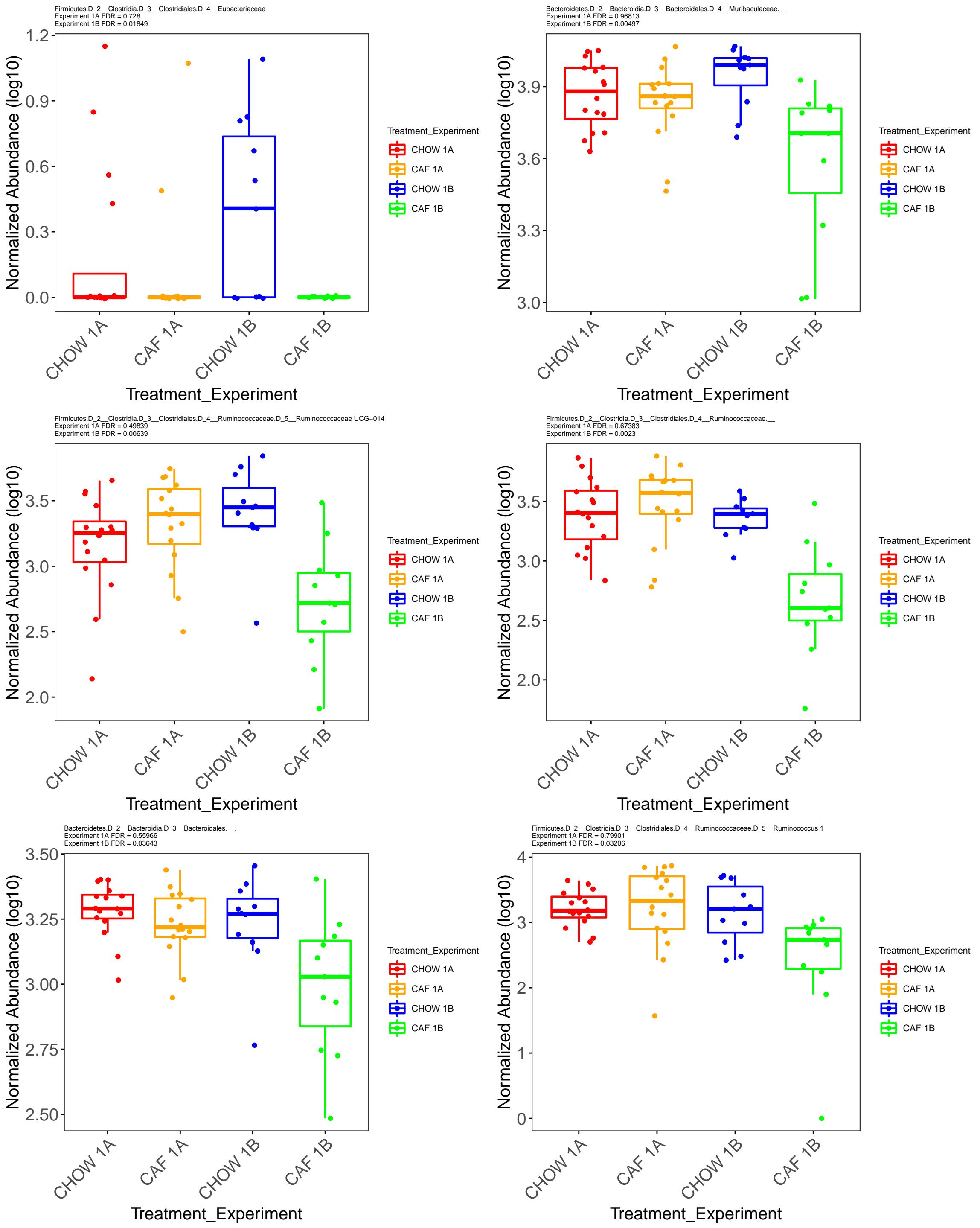


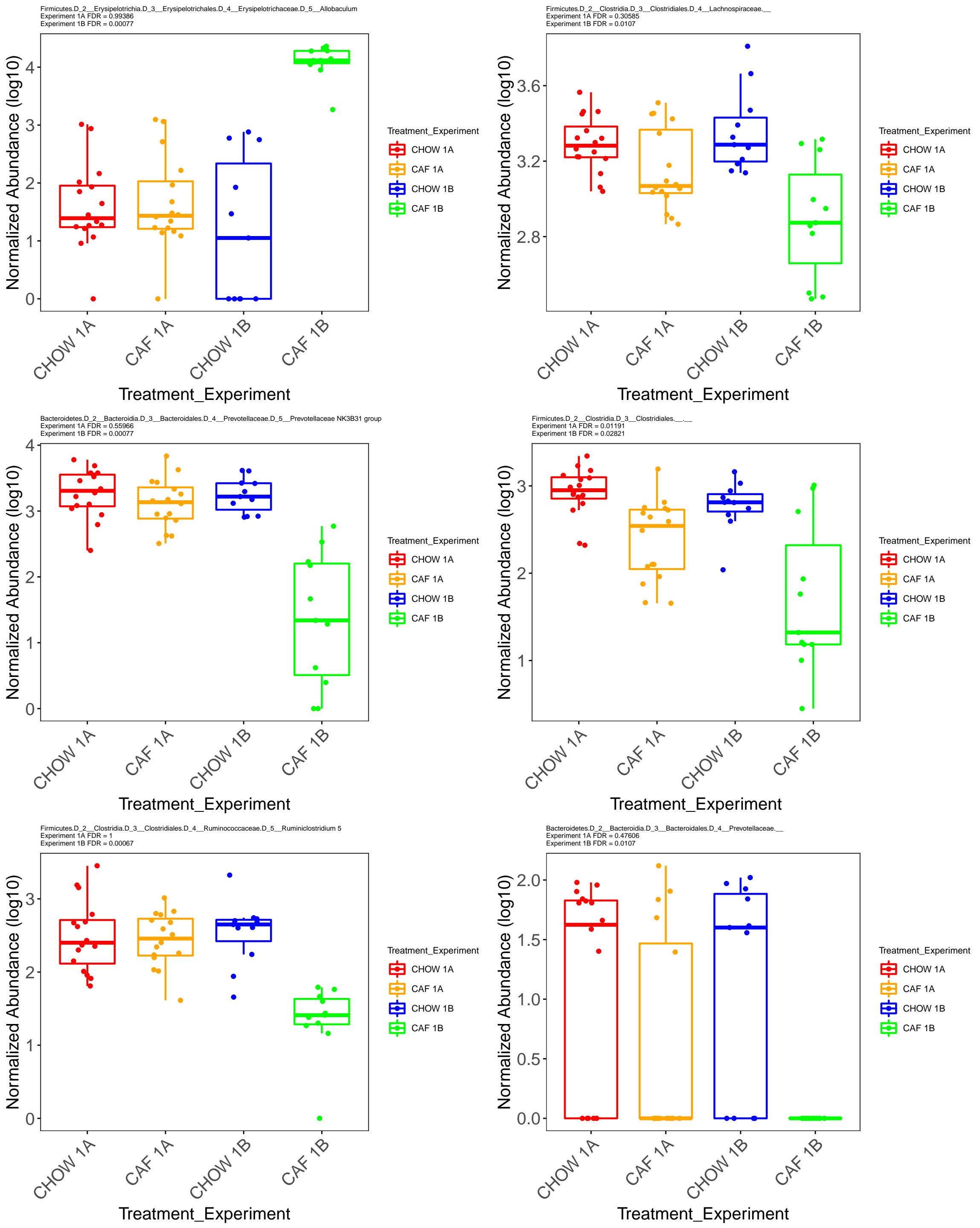
Tenericutes.D_2_Mollicutes.D_3_Mollicutes RF39.
Experiment 1A FDR = 0.85856
Experiment 1B FDR = 0.04013



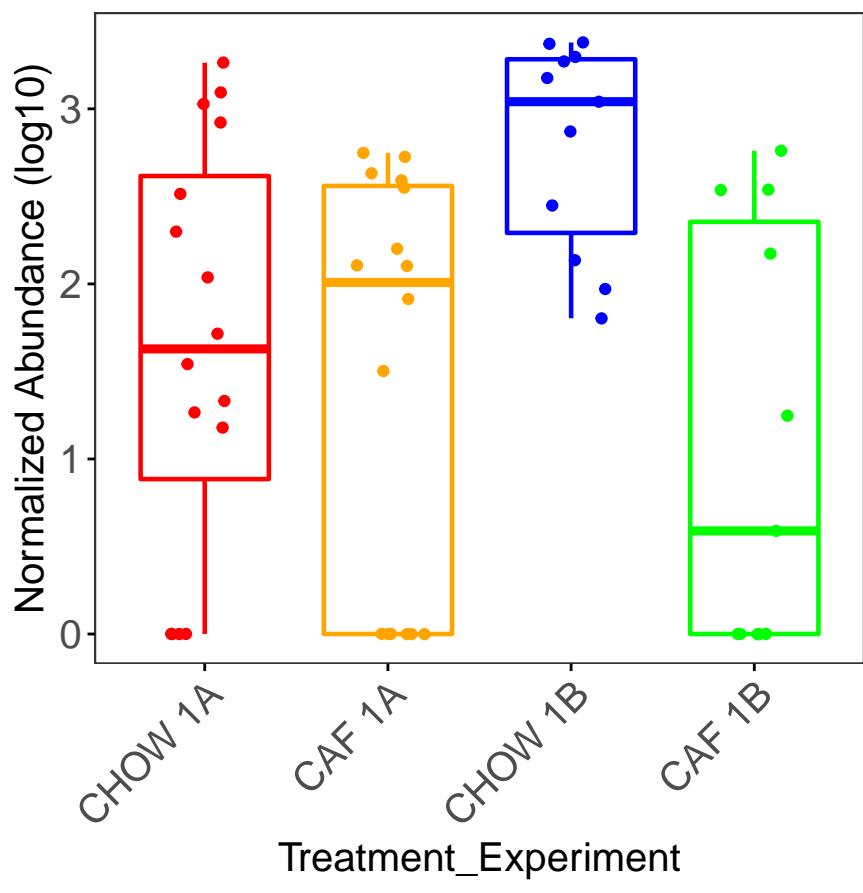




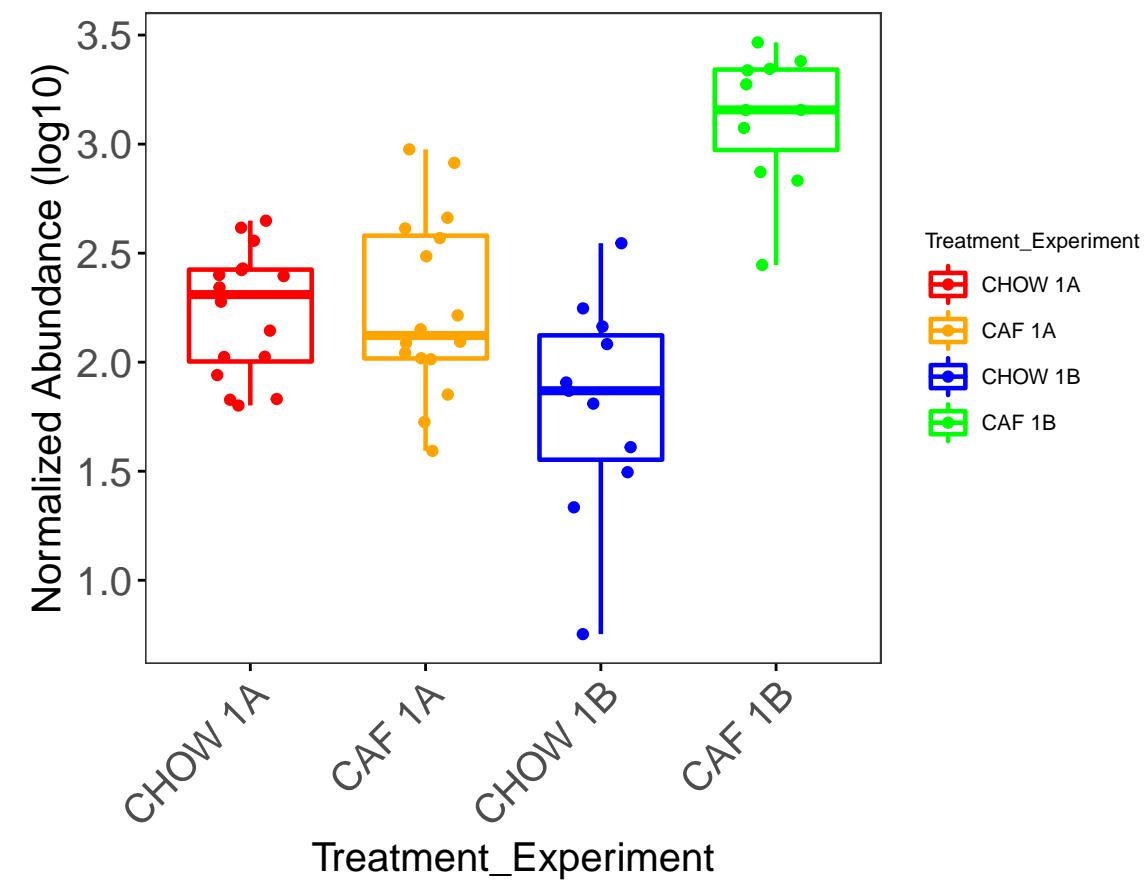




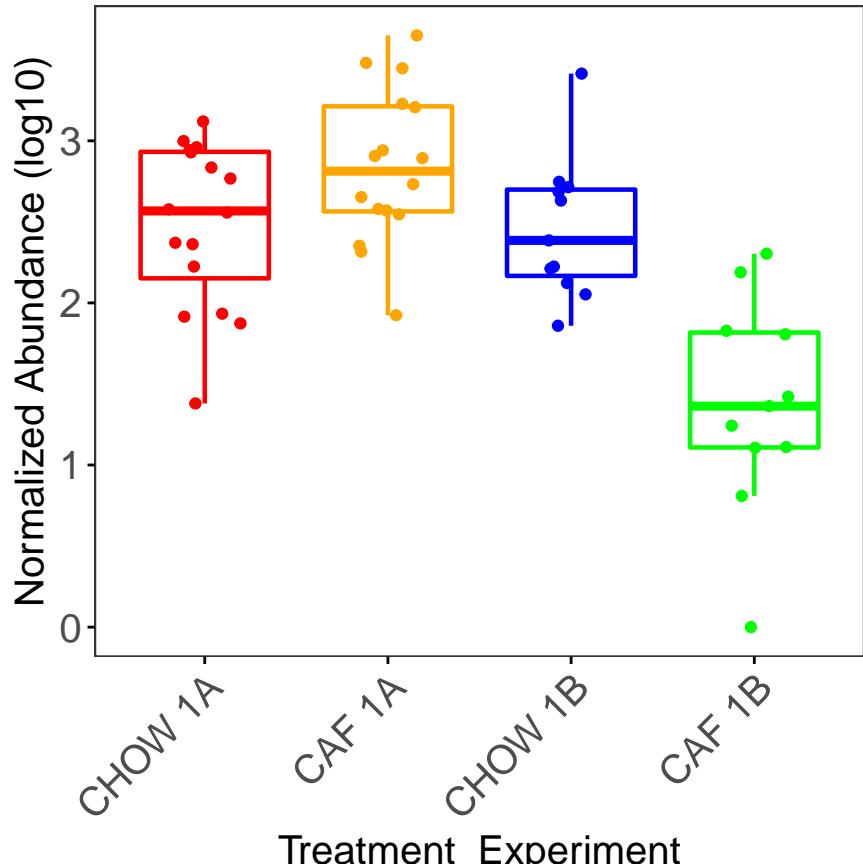
Firmicutes.D_2_Clostridia.D_3_Clostridiales.D_4_Ruminococcaceae.D_5_Ruminococcus 2
Experiment 1A FDR = 0.94822
Experiment 1B FDR = 0.01196



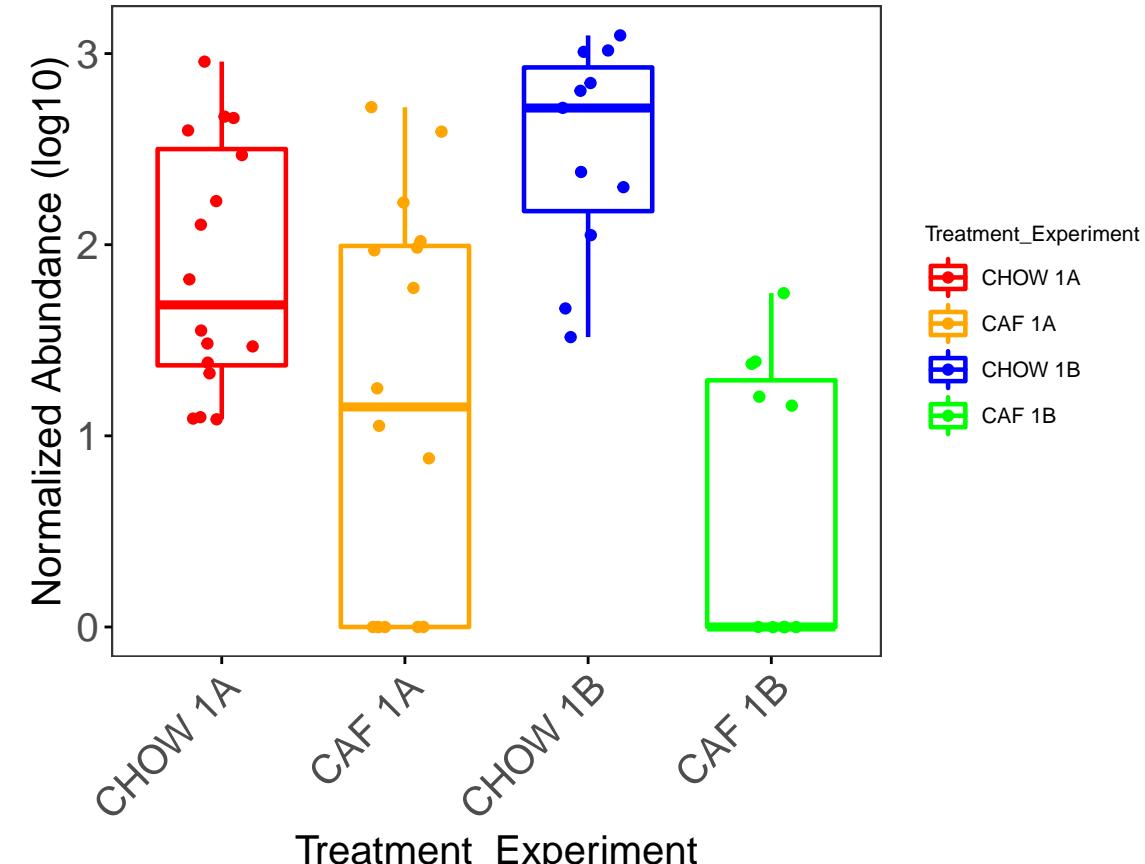
Proteobacteria.D_2_Gammaproteobacteria.D_3_Betaproteobacteriales.D_4_Burkholderiaceae.D_5_Parasutterella
Experiment 1A FDR = 1
Experiment 1B FDR = 6e-04



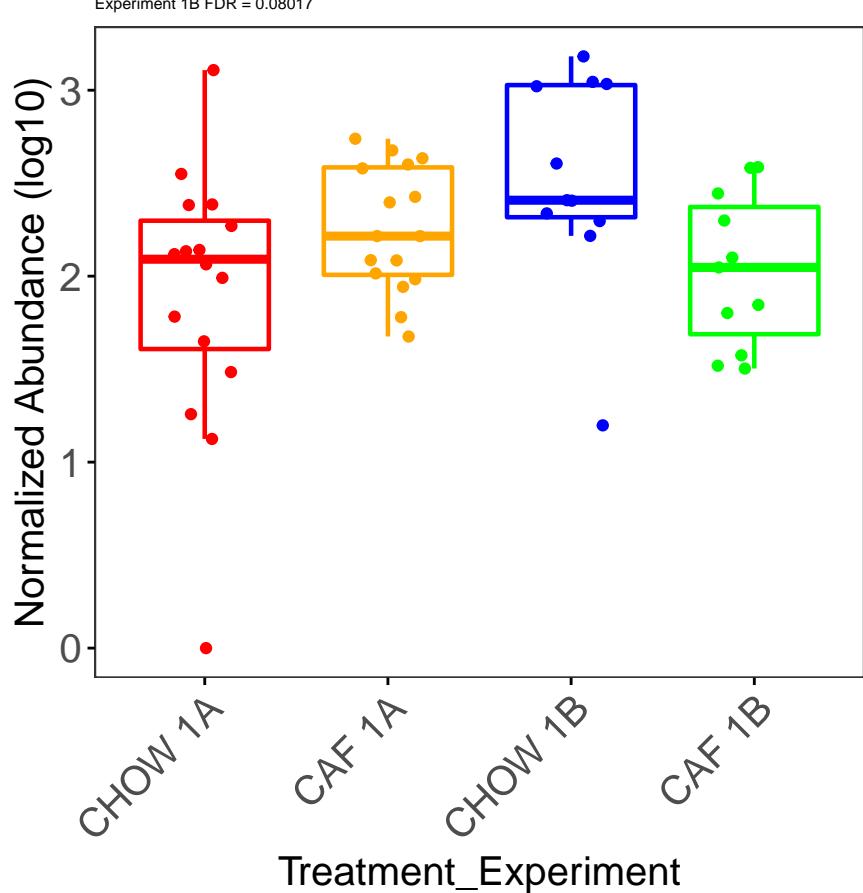
Firmicutes.D_2_Clostridia.D_3_Clostridiales.D_4_Ruminococcaceae UCG-013
Experiment 1A FDR = 0.47606
Experiment 1B FDR = 0.00153



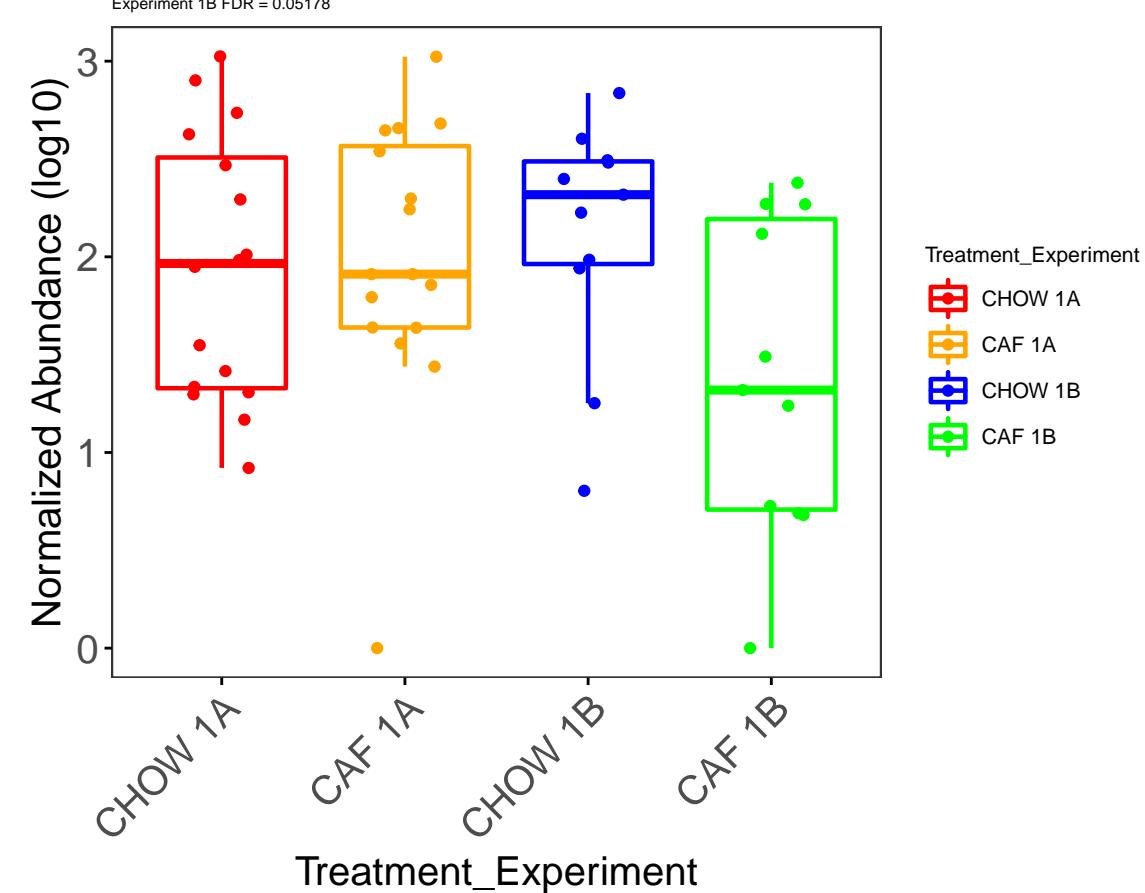
Firmicutes.D_2_Clostridia.D_3_Clostridiales.D_4_Ruminococcaceae.D_5_Ruminiclostridium 6
Experiment 1A FDR = 0.30585
Experiment 1B FDR = 0.00105

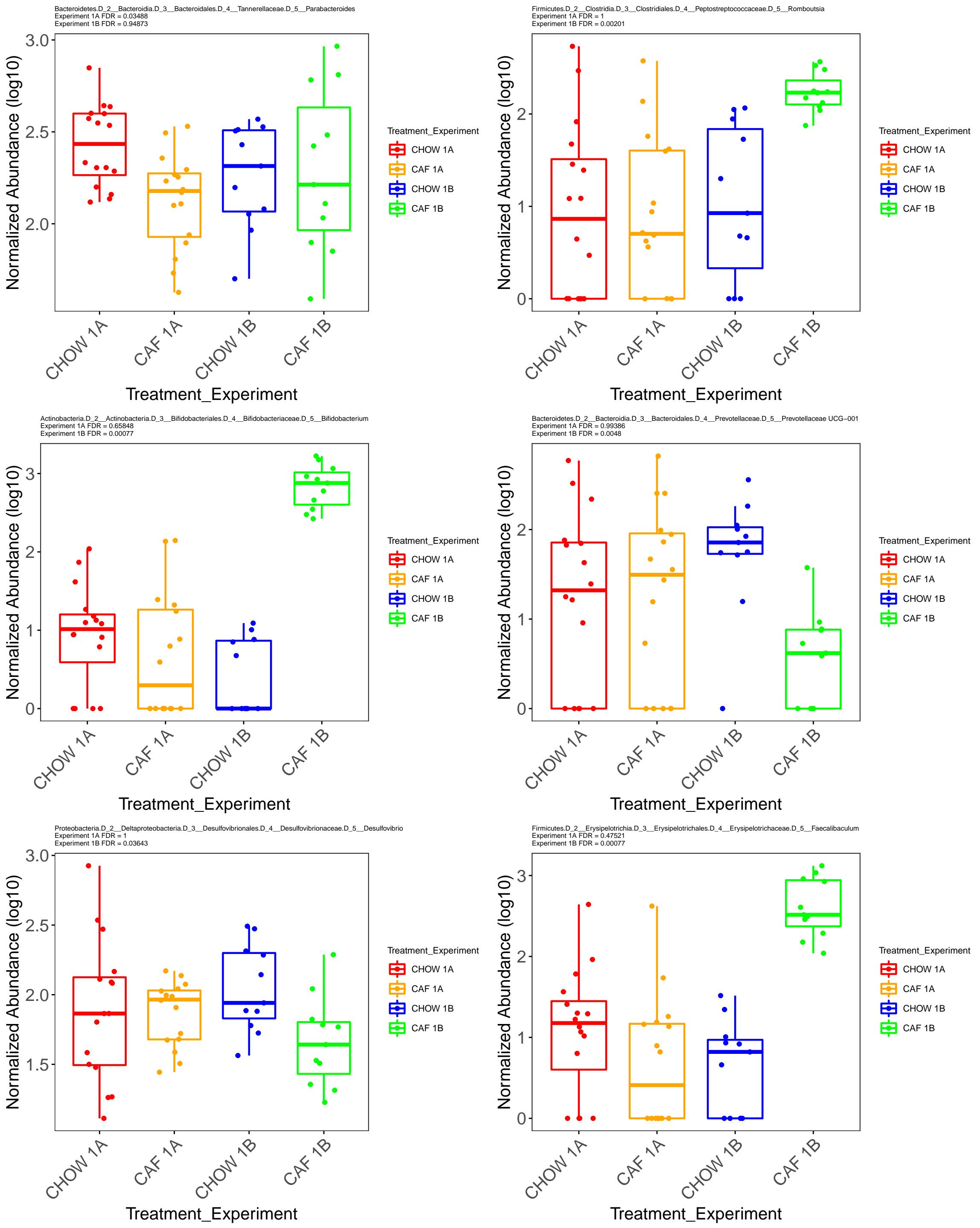


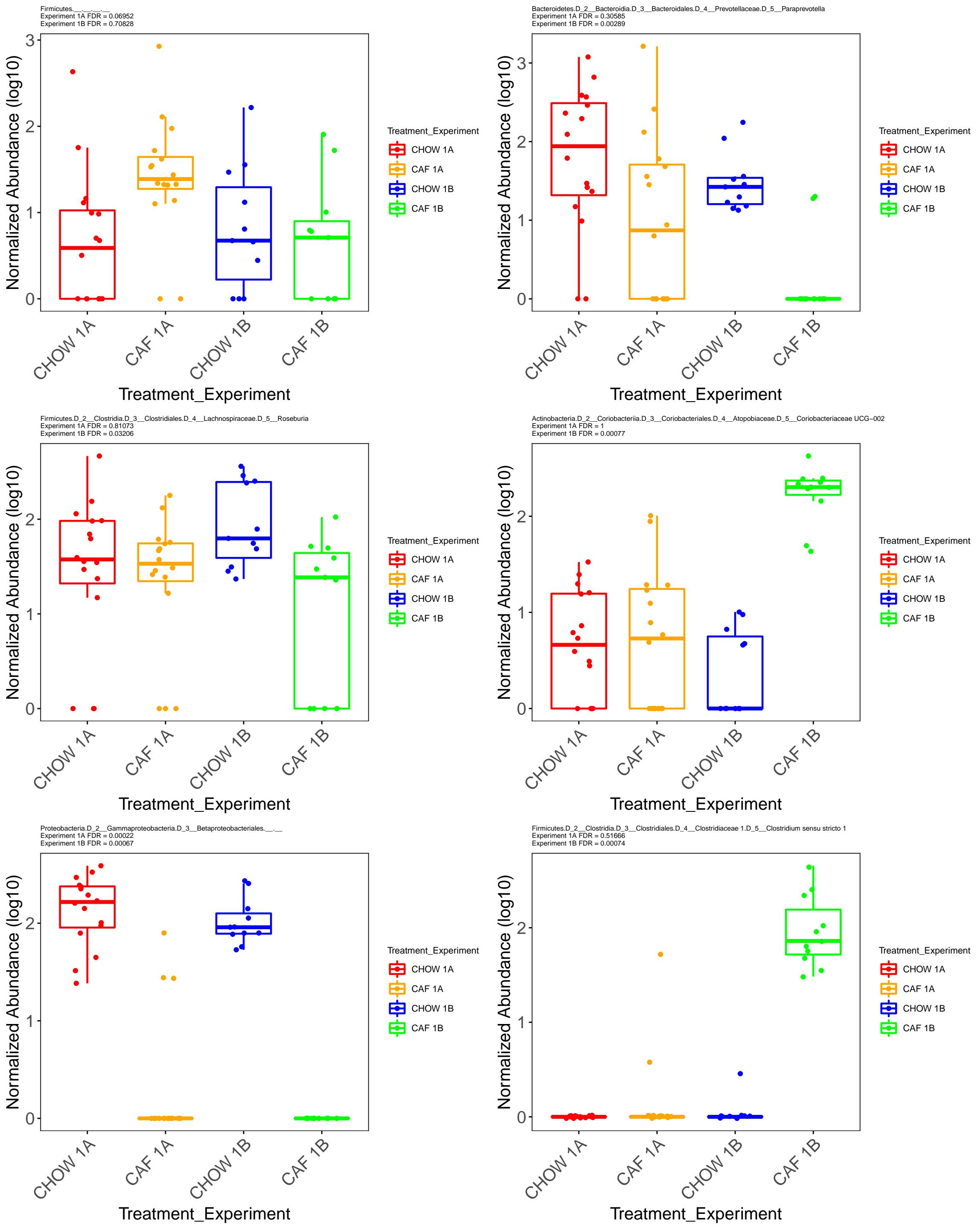
Firmicutes.D_2_Clostridia.D_3_Clostridiales.D_4_Lachnospiraceae.D_5_Lachnospiraceae NK4A136 group
Experiment 1A FDR = 0.47521
Experiment 1B FDR = 0.08017

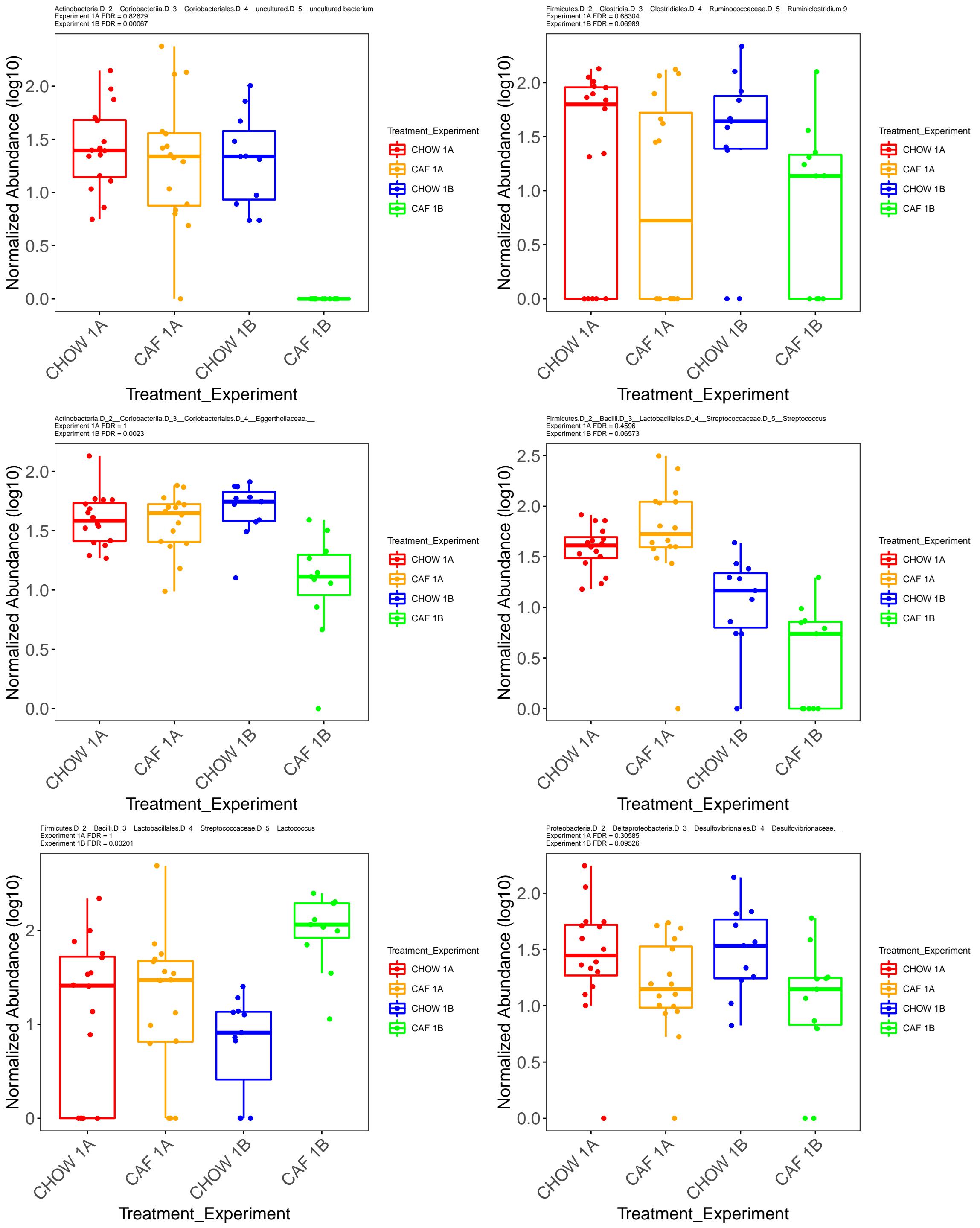


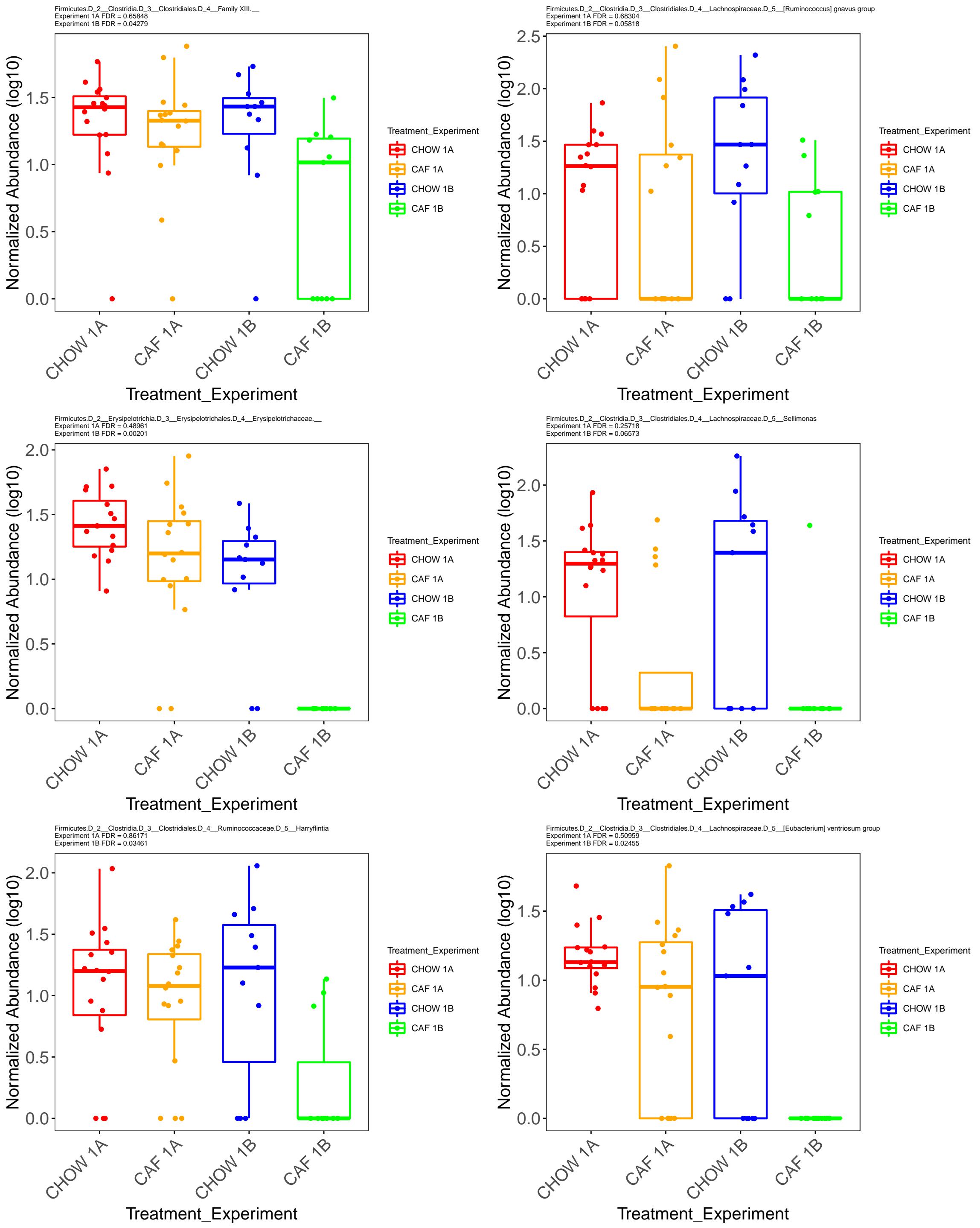
Tenericutes.D_2_Mollicutes.D_3_Mollicutes RF39
Experiment 1A FDR = 0.87697
Experiment 1B FDR = 0.05178



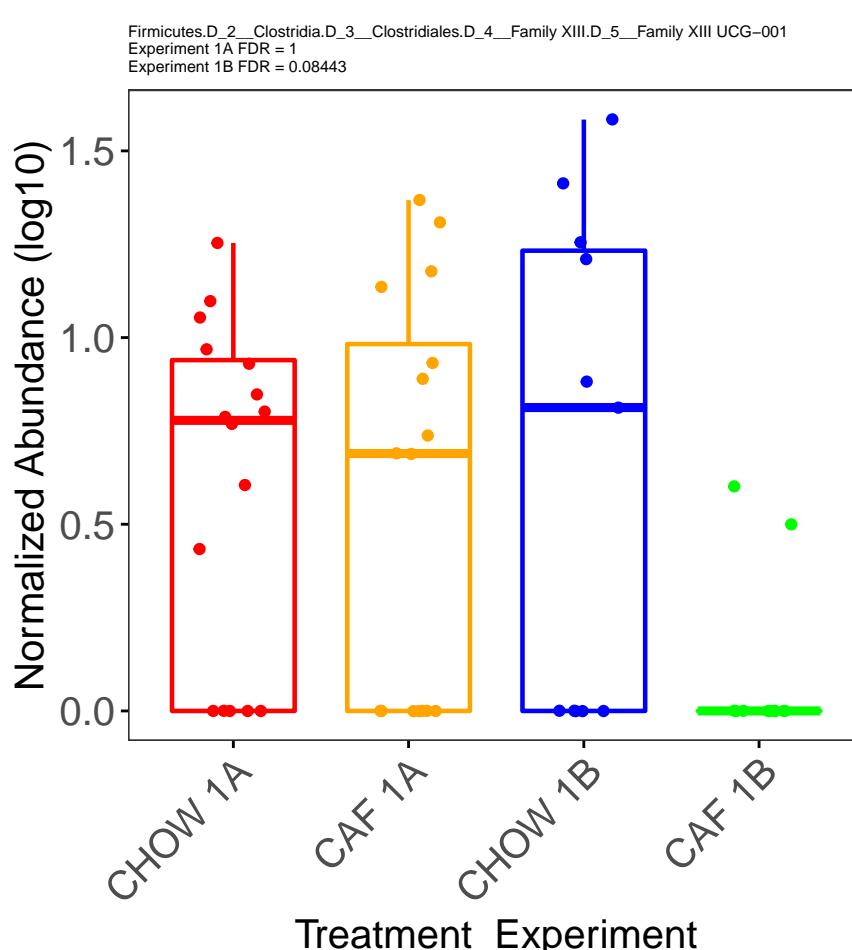
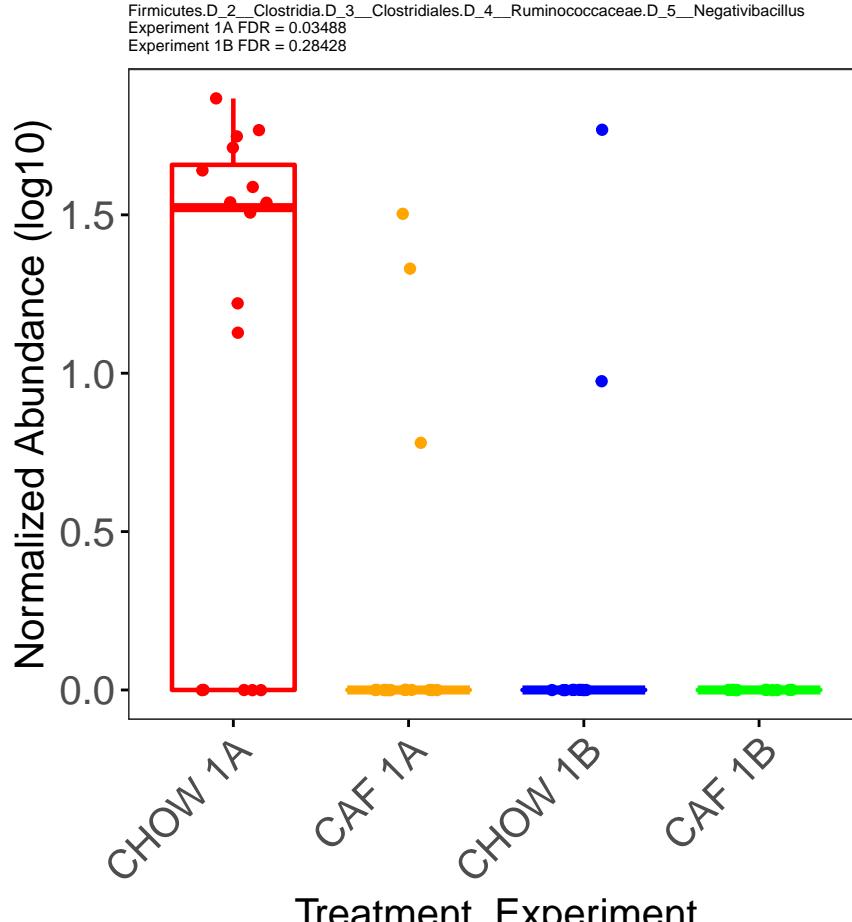
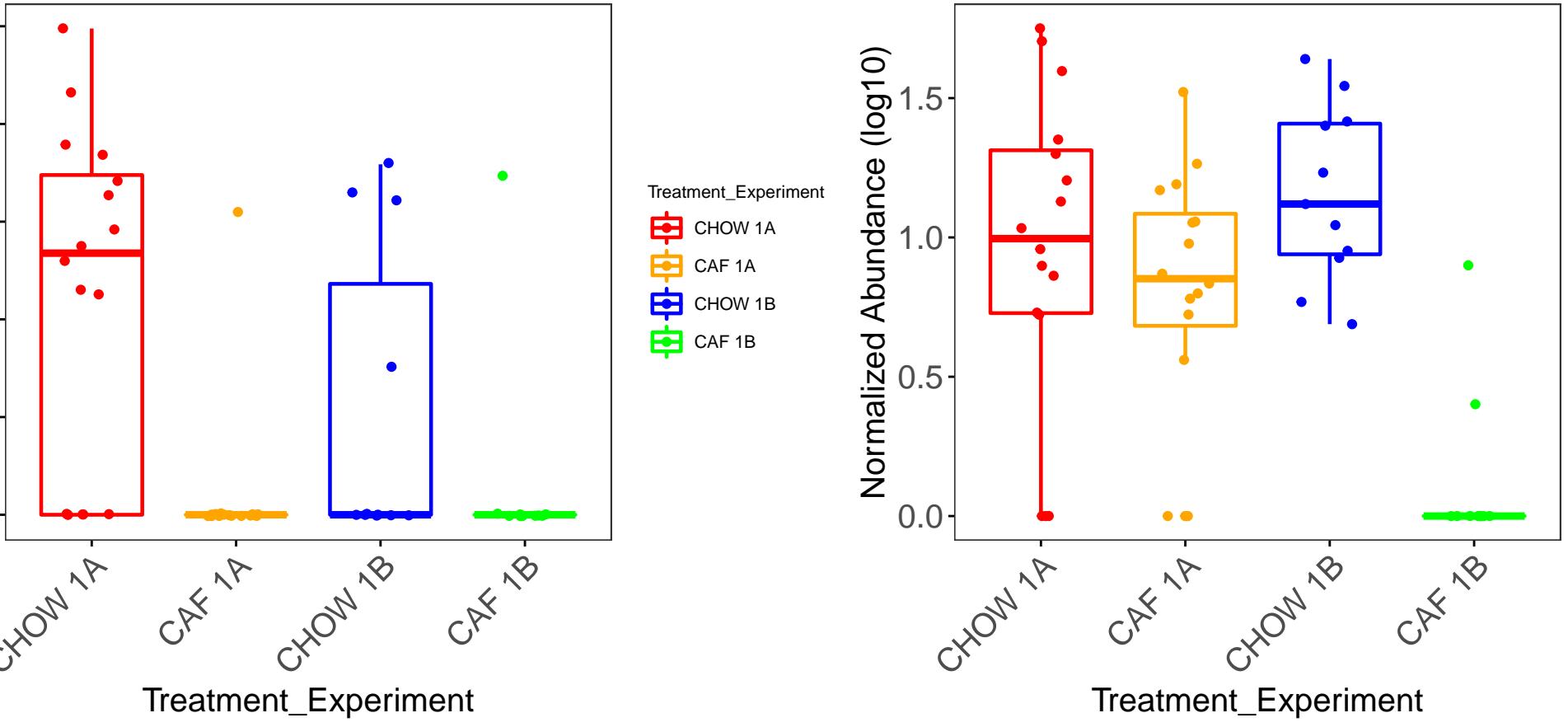




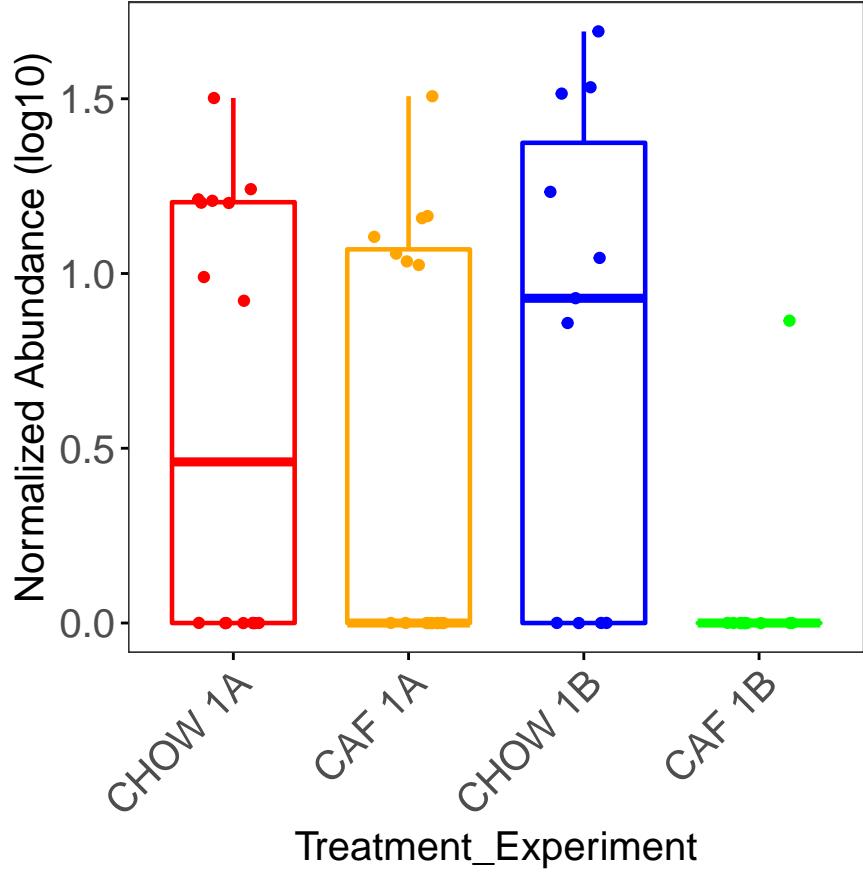




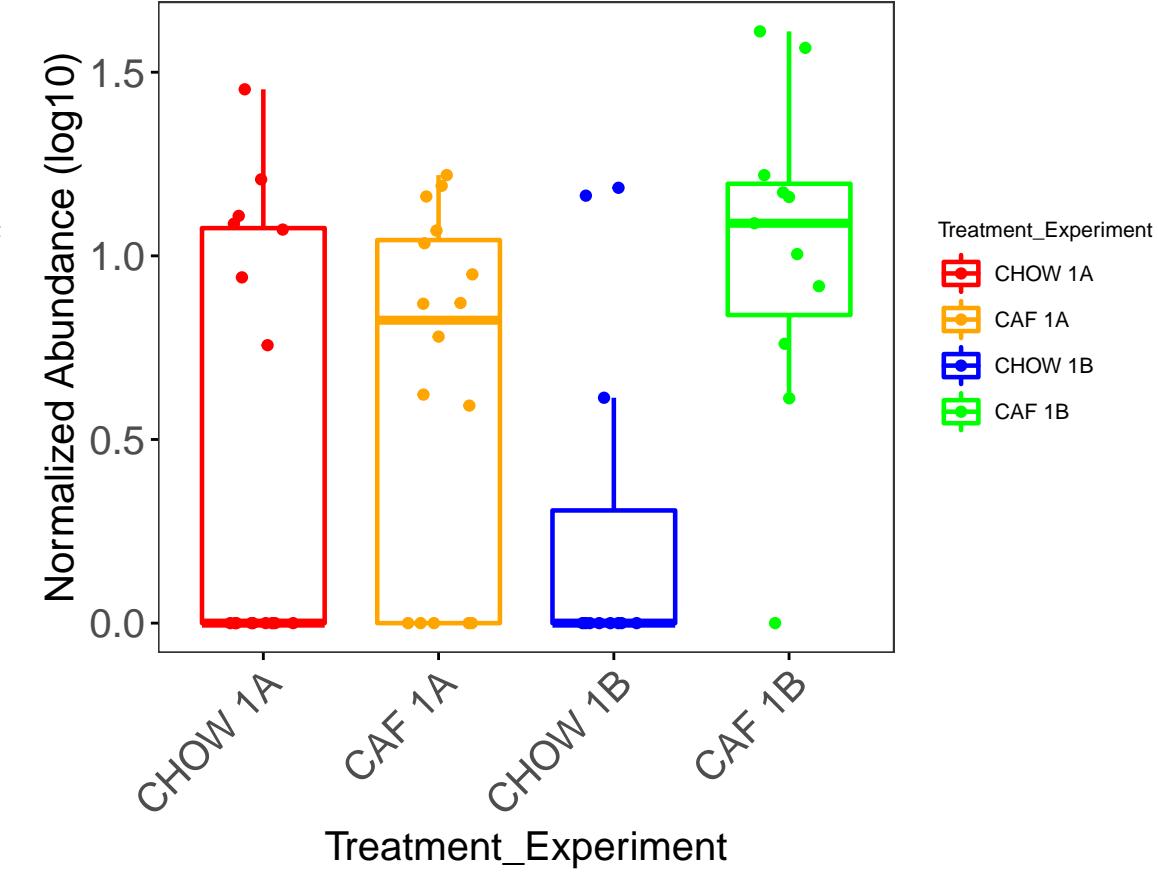
Proteobacteria.D_2_Gammaproteobacteria.D_3_Betaproteobacteriales.D_4_Burkholderiaceae.
Experiment 1A FDR = 0.01823
Experiment 1B FDR = 0.29501



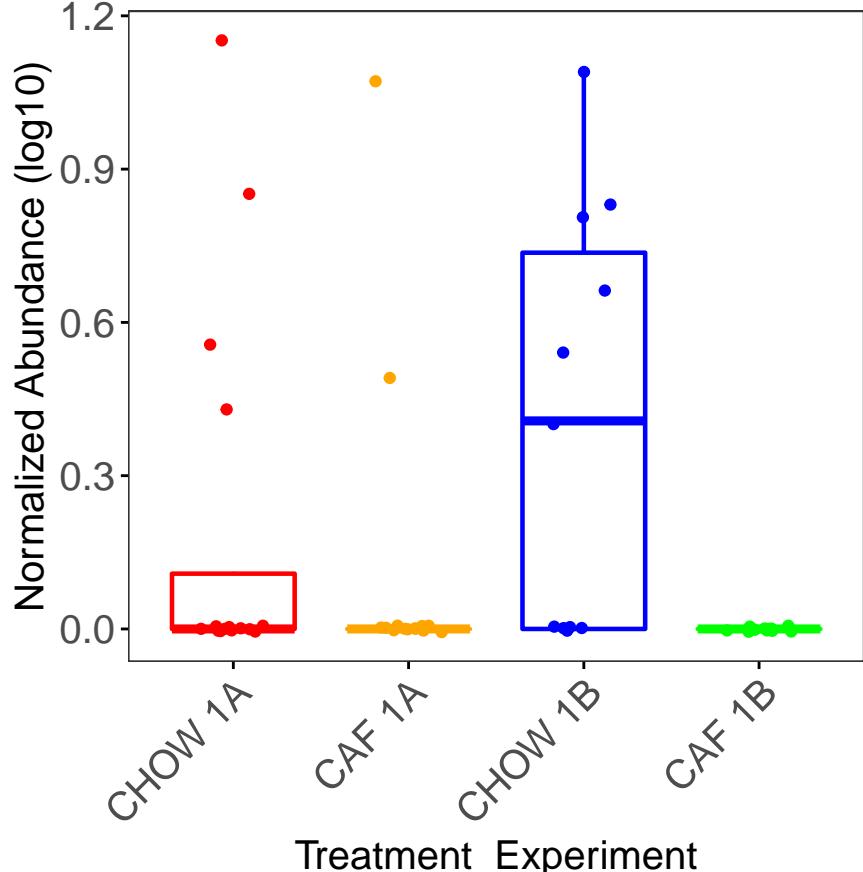
Firmicutes.D_2_Clostridia.D_3_Clostridiales.D_4_Ruminococcaceae.D_5_GCA-900066225
Experiment 1A FDR = 0.80513
Experiment 1B FDR = 0.02778



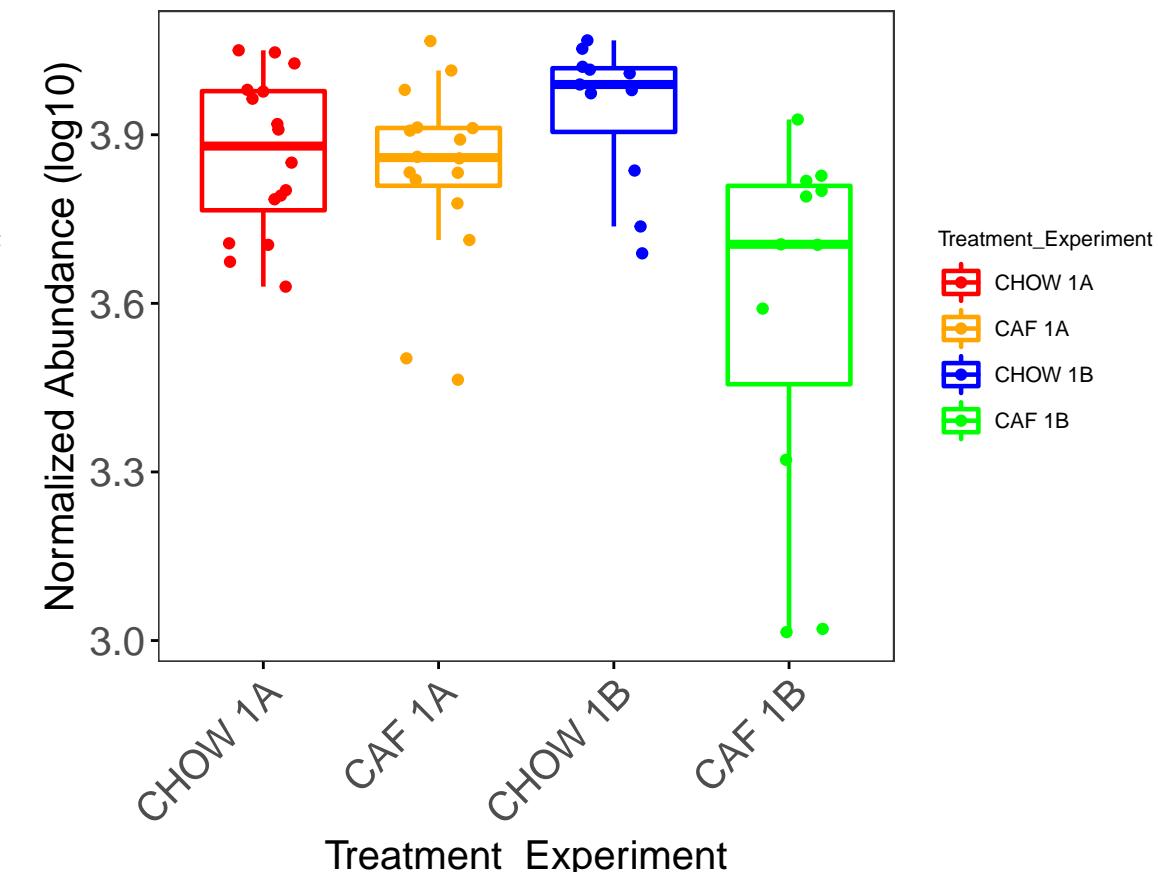
Firmicutes.D_2_Clostridia.D_3_Clostridiales.D_4_Lachnospiraceae.D_5_Tyzzerella
Experiment 1A FDR = 0.79004
Experiment 1B FDR = 0.02797



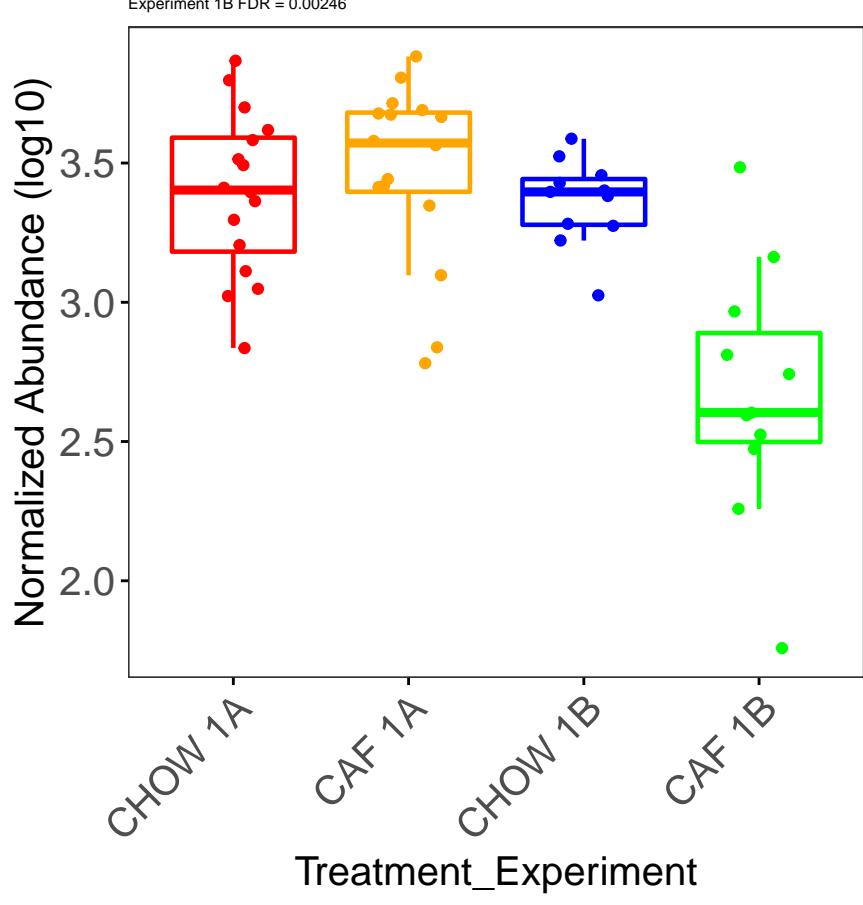
Firmicutes.D_2_Clostridia.D_3_Clostridiales.D_4_Eubacteriaceae.D_5_Anaerofustis
Experiment 1A FDR = 0.70708
Experiment 1B FDR = 0.02455



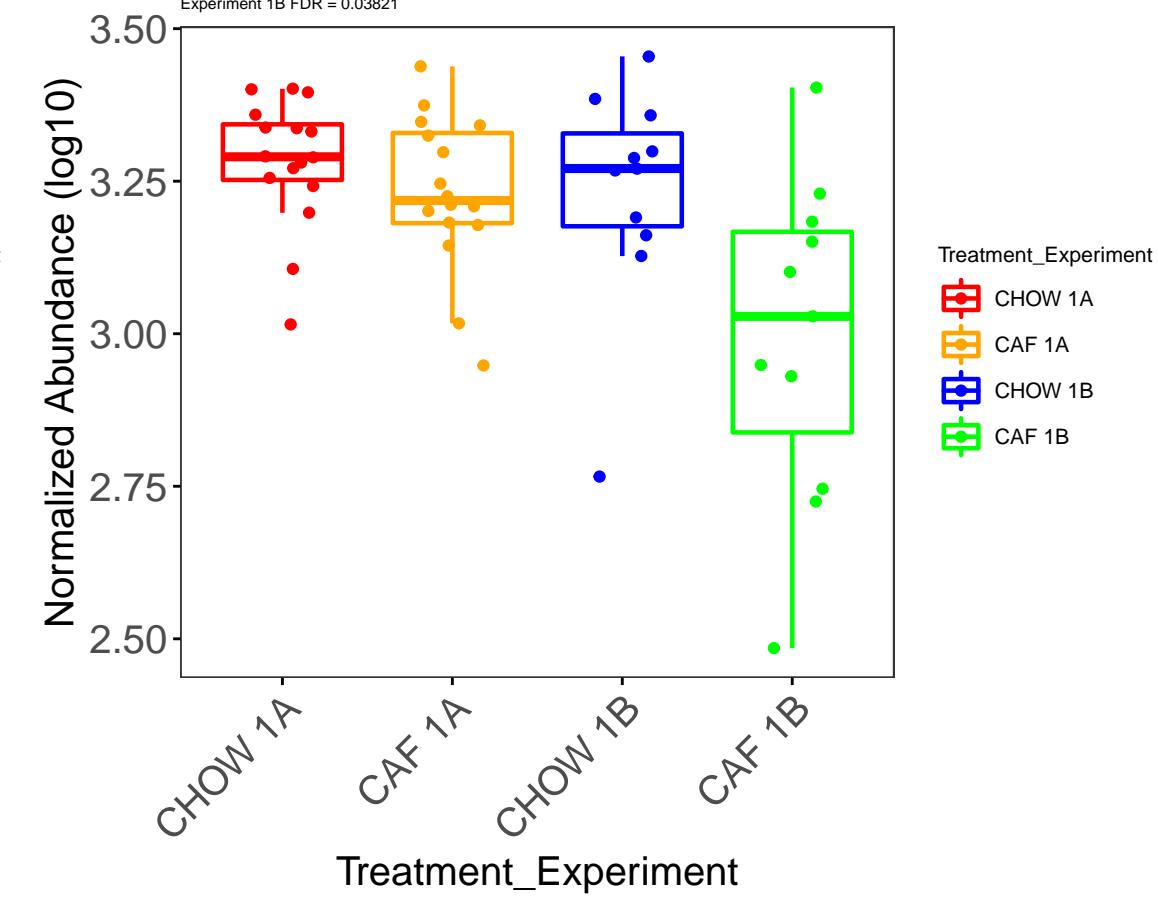
Bacteroidetes.D_2_Bacteroidia.D_3_Bacteroidales.D_4_Muribaculaceae.
Experiment 1A FDR = 0.97145
Experiment 1B FDR = 0.00565

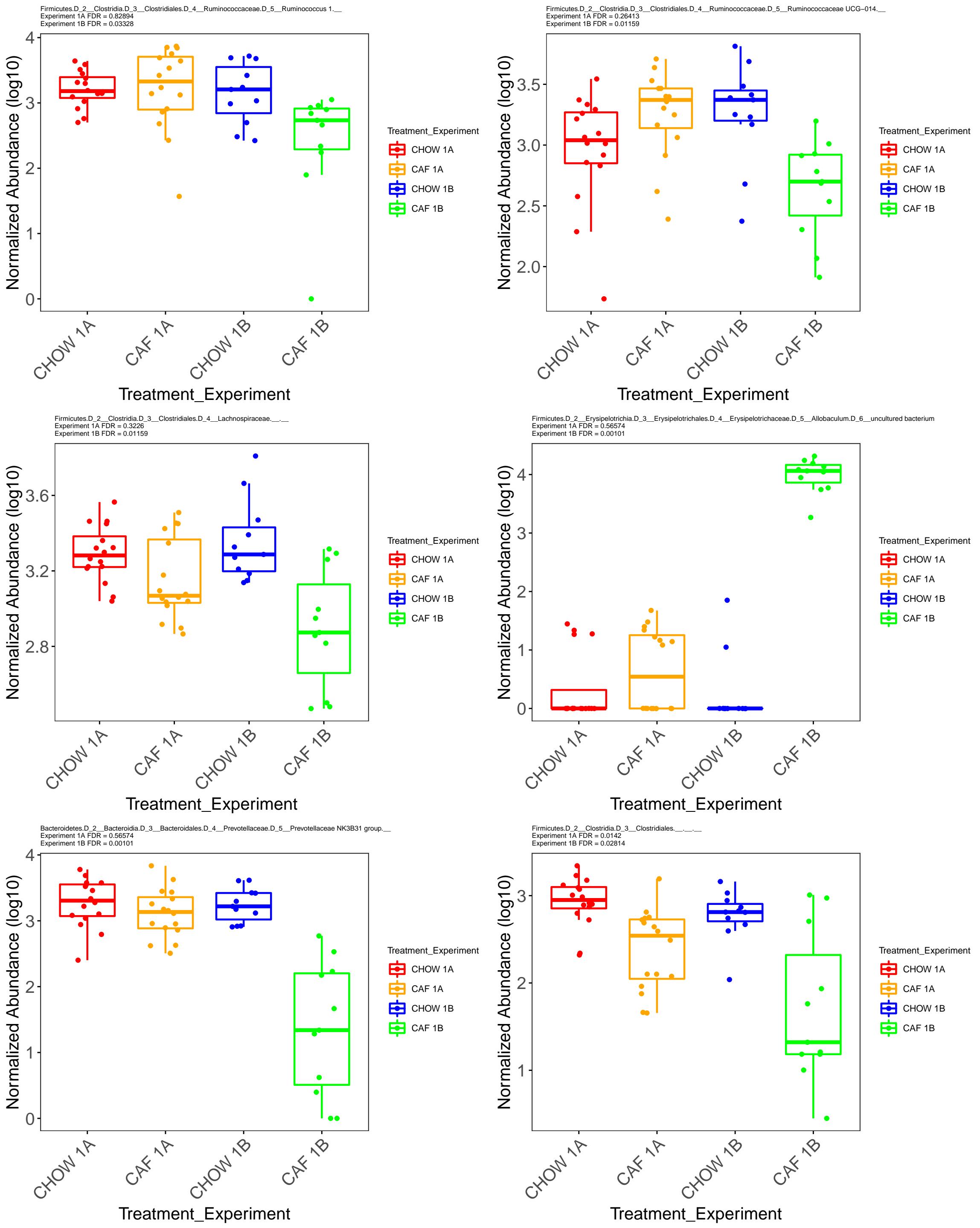


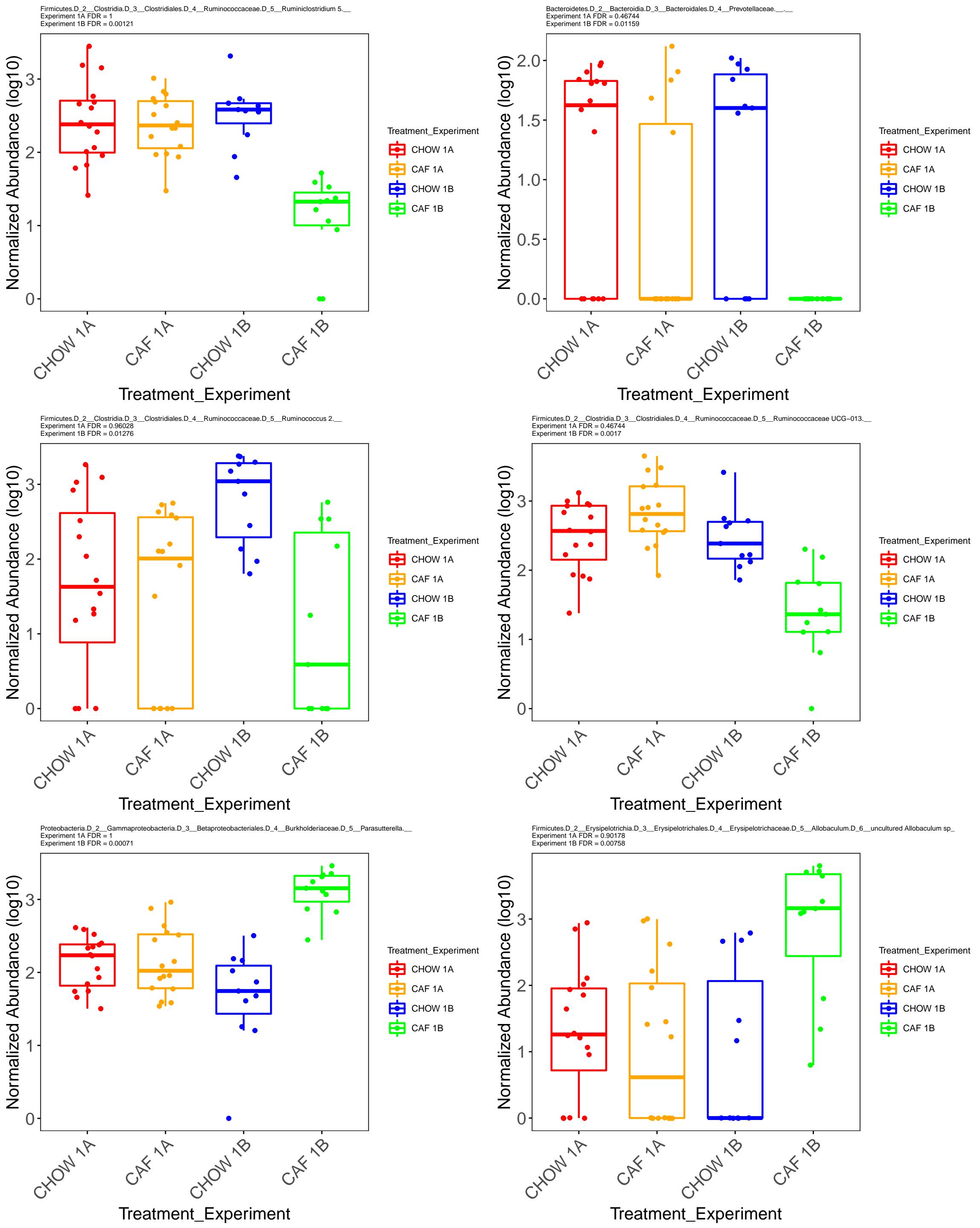
Firmicutes.D_2_Clostridia.D_3_Clostridiales.D_4_Ruminococcaceae.
Experiment 1A FDR = 0.68655
Experiment 1B FDR = 0.00246

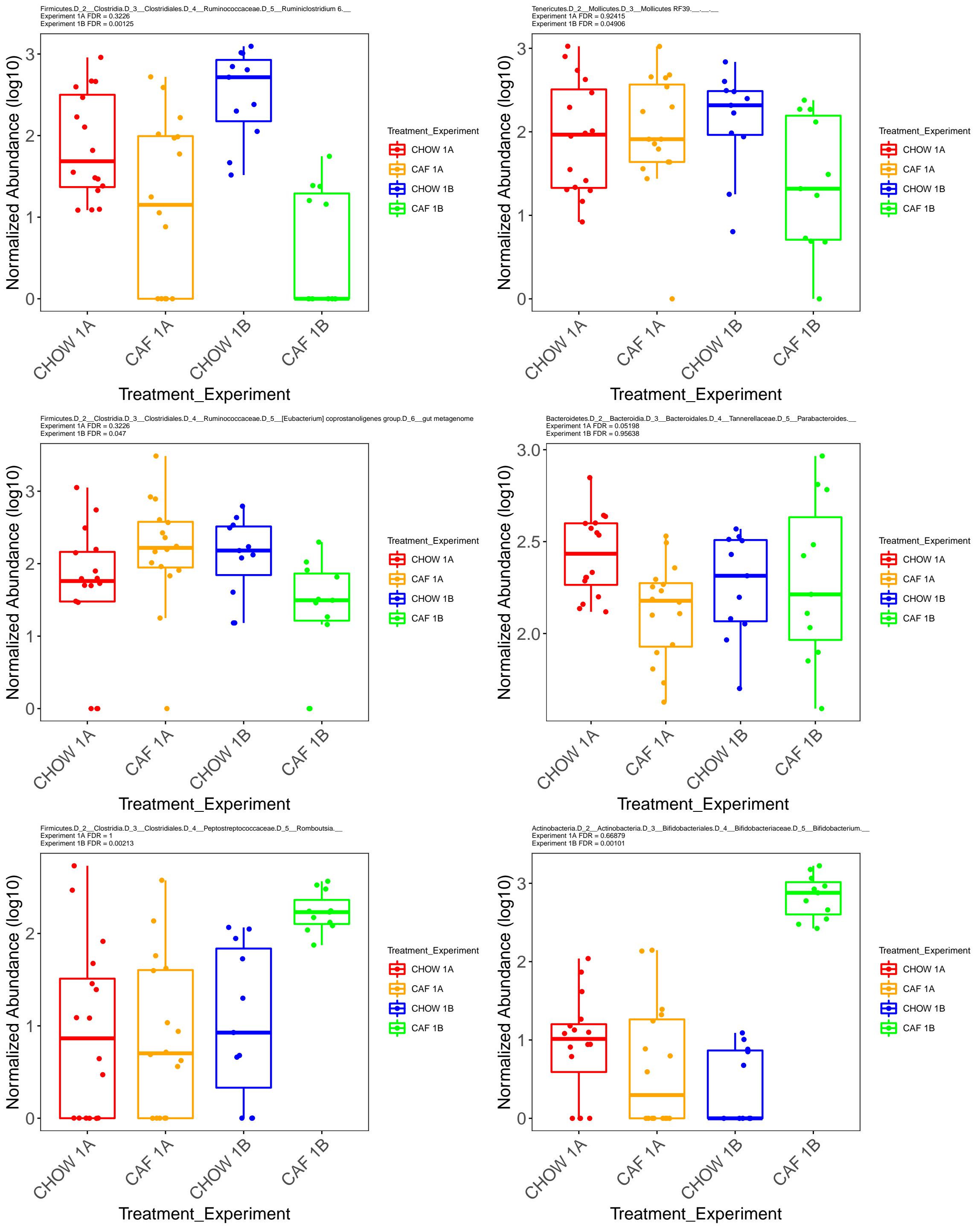


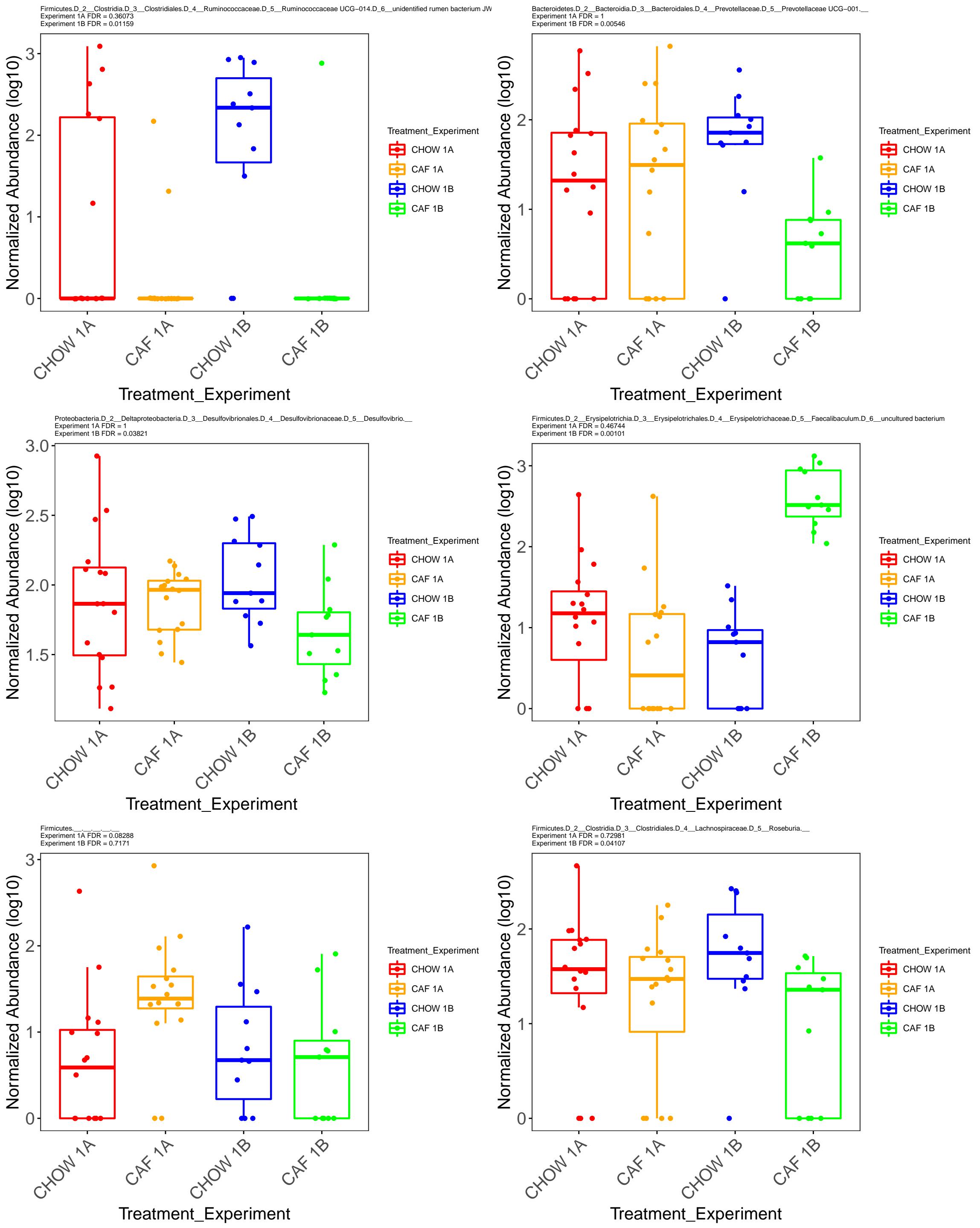
Bacteroidetes.D_2_Bacteroidia.D_3_Bacteroidales.
Experiment 1A FDR = 0.56574
Experiment 1B FDR = 0.03821



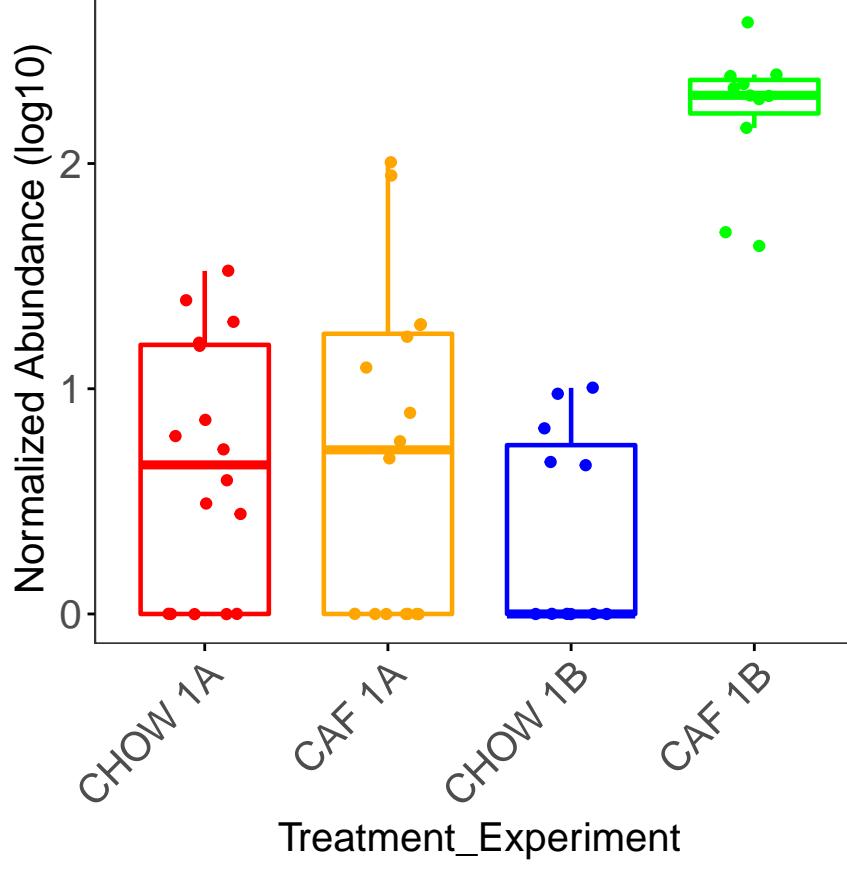




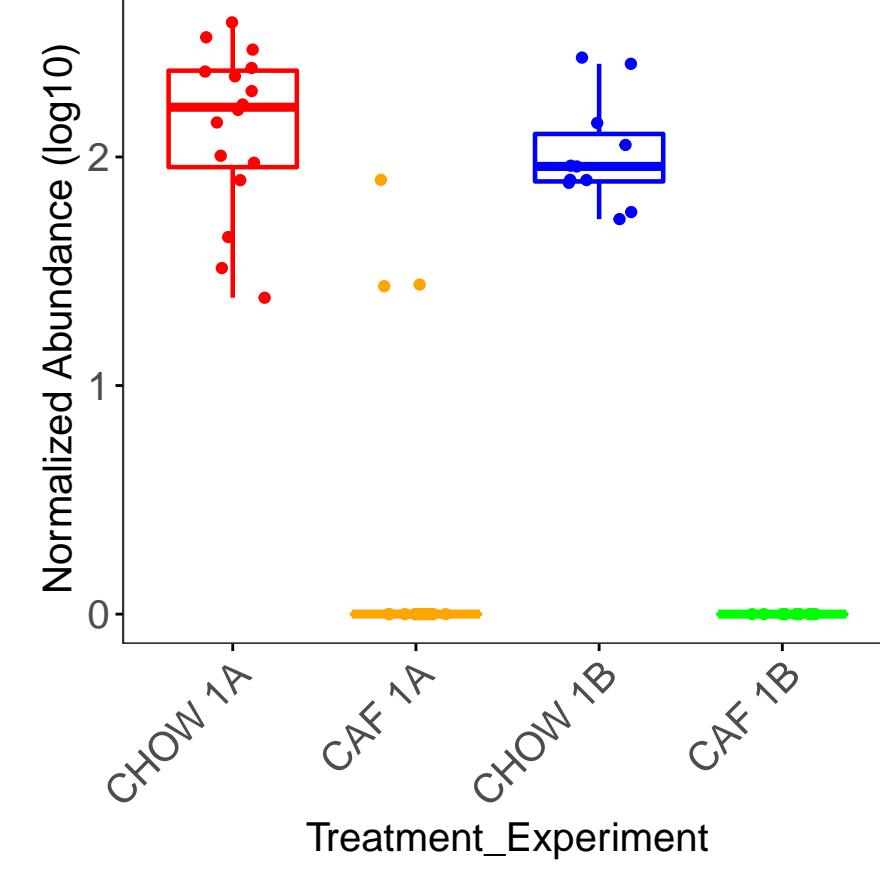




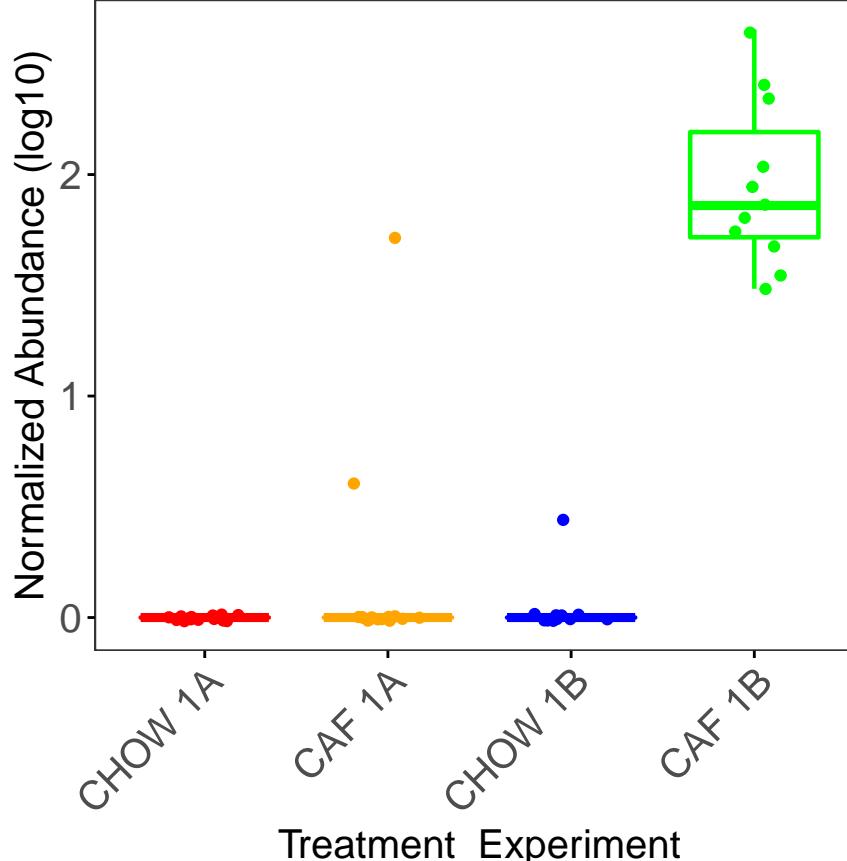
Actinobacteria.D_2_Coriobacteriia.D_3_Coriobacteriales.D_4_Atopobiaceae.D_5_Coriobacteriaceae UCG-002.D_6_uncultured bacterium
Experiment 1A FDR = 1
Experiment 1B FDR = 0.00101



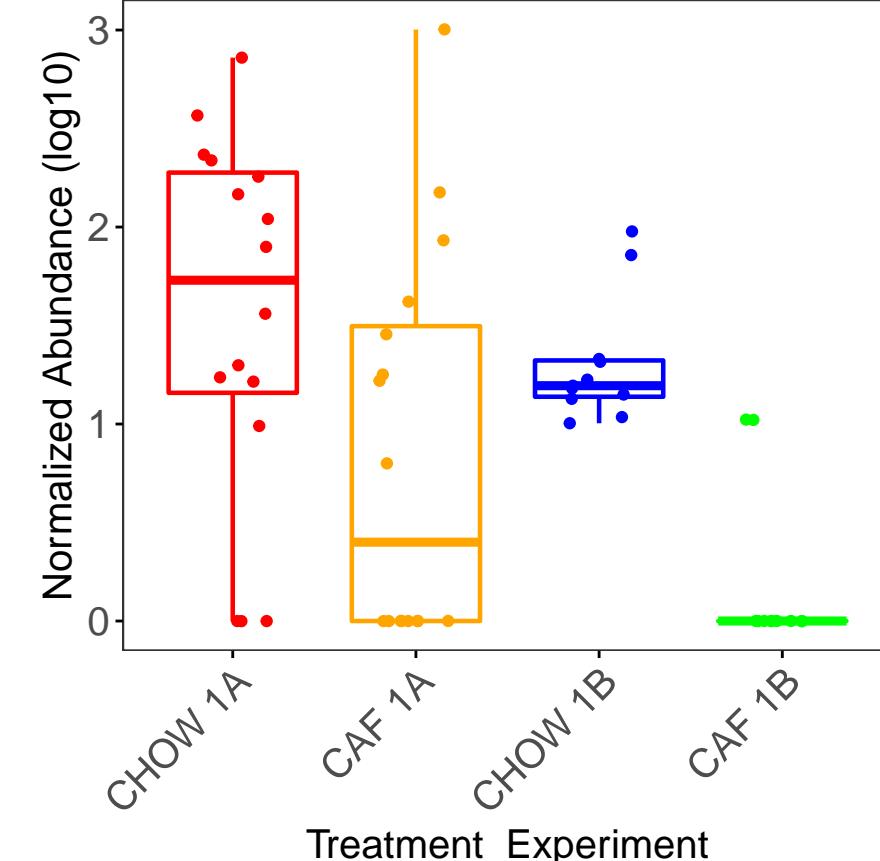
Proteobacteria.D_2_Gammaproteobacteria.D_3_Betaproteobacteriales.
Experiment 1A FDR = 0.00026
Experiment 1B FDR = 0.00101



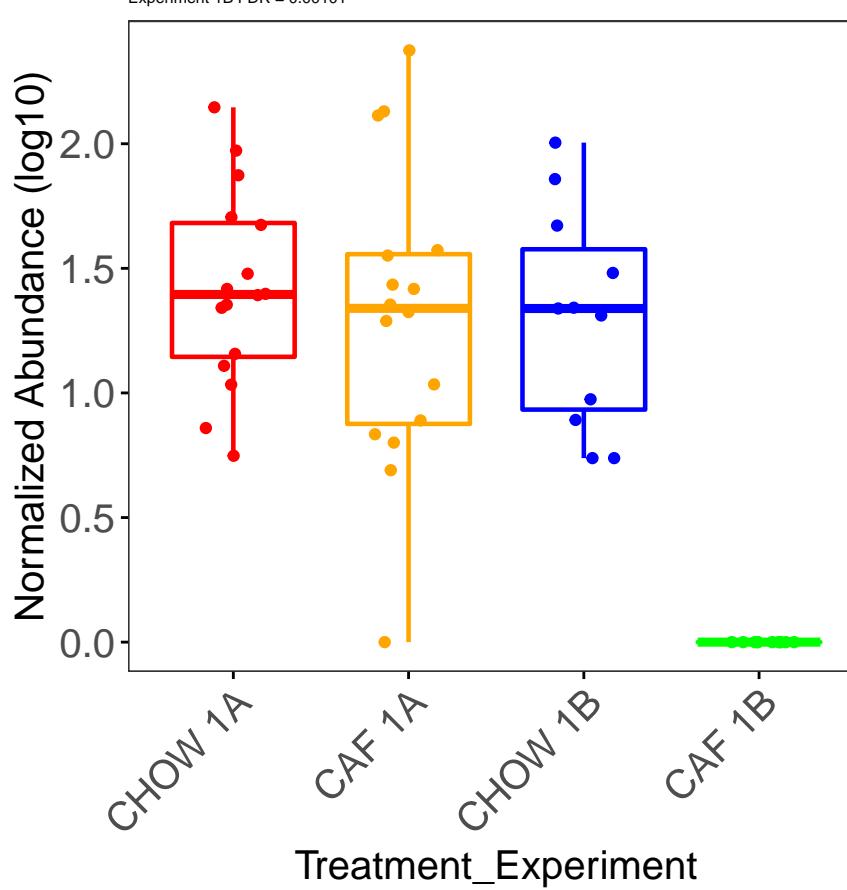
Firmicutes.D_2_Clostridia.D_3_Clostridiales.D_4_Clostridiaceae 1.D_5_Clostridium sensu stricto 1.
Experiment 1A FDR = 0.53496
Experiment 1B FDR = 0.00101



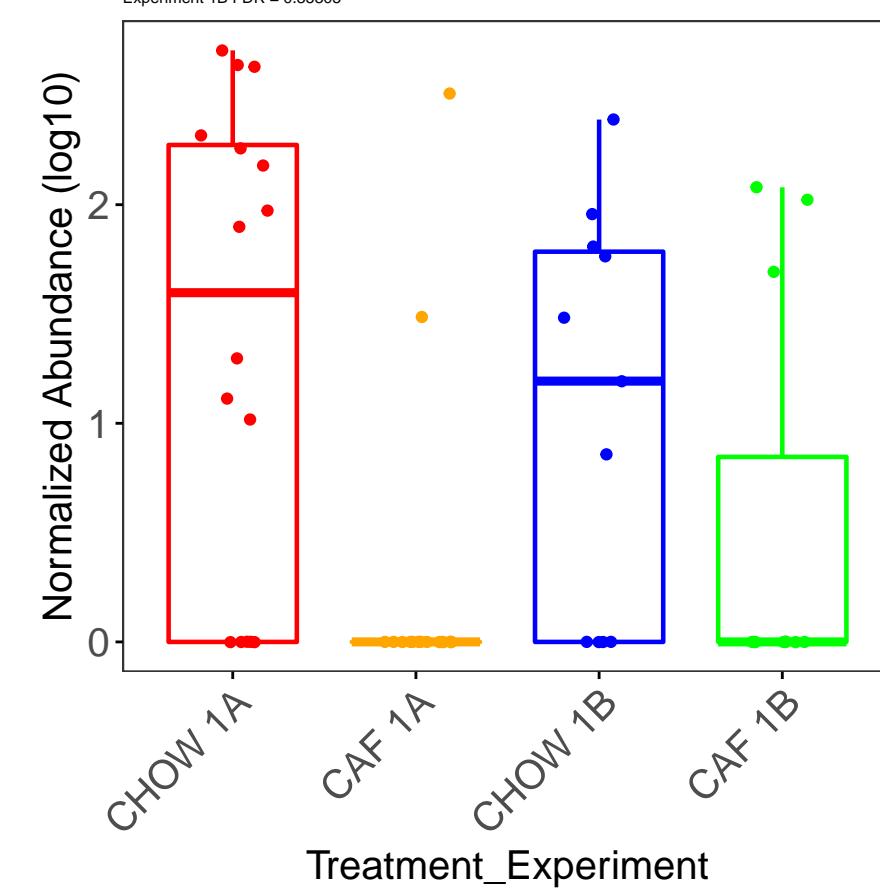
Bacteroidetes.D_2_Bacteroidia.D_3_Bacteroidales.D_4_Prevotellaceae.D_5_Paraprevotella.
Experiment 1A FDR = 0.3226
Experiment 1B FDR = 0.00101

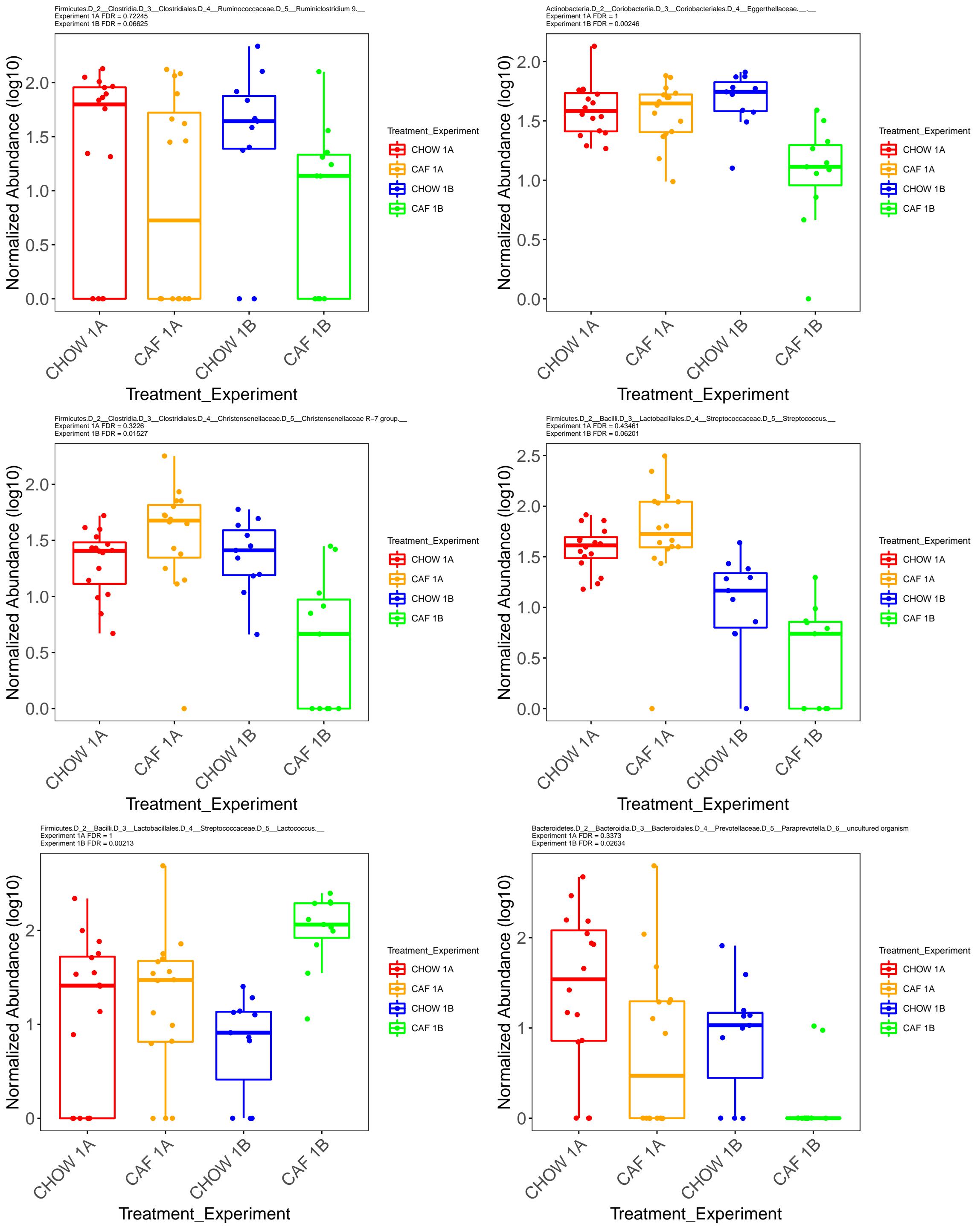


Actinobacteria.D_2_Coriobacteriia.D_3_Coriobacteriales.D_4_uncultured.D_5_uncultured bacterium.D_6_uncultured bacterium
Experiment 1A FDR = 0.89678
Experiment 1B FDR = 0.00101



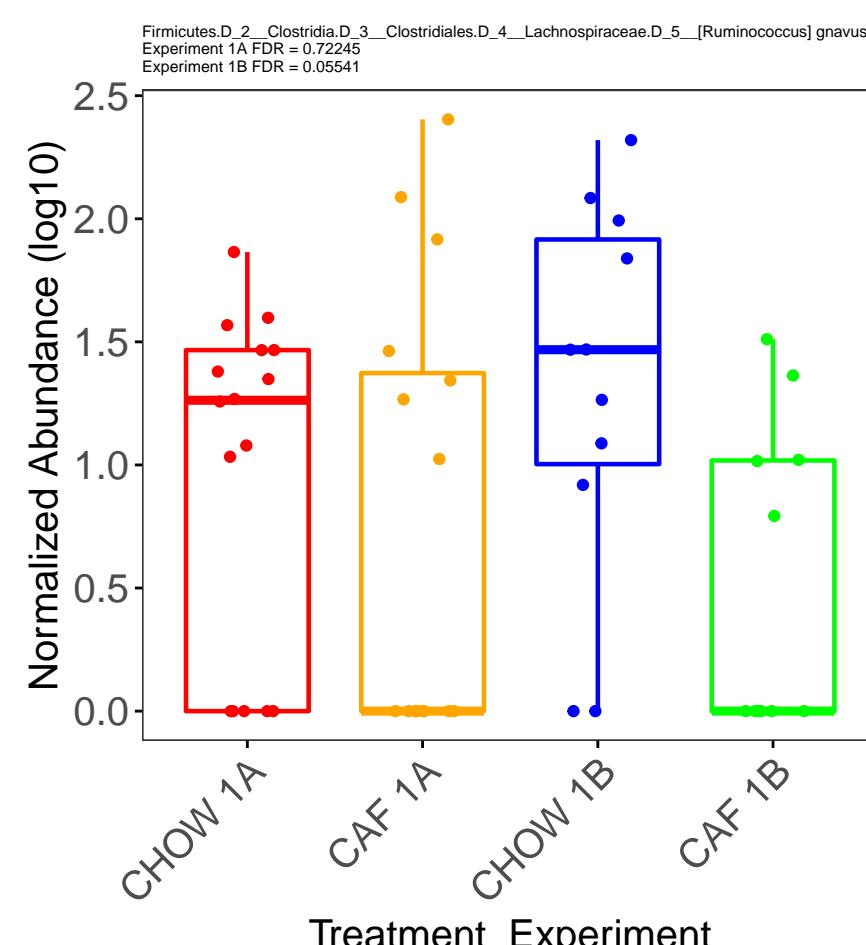
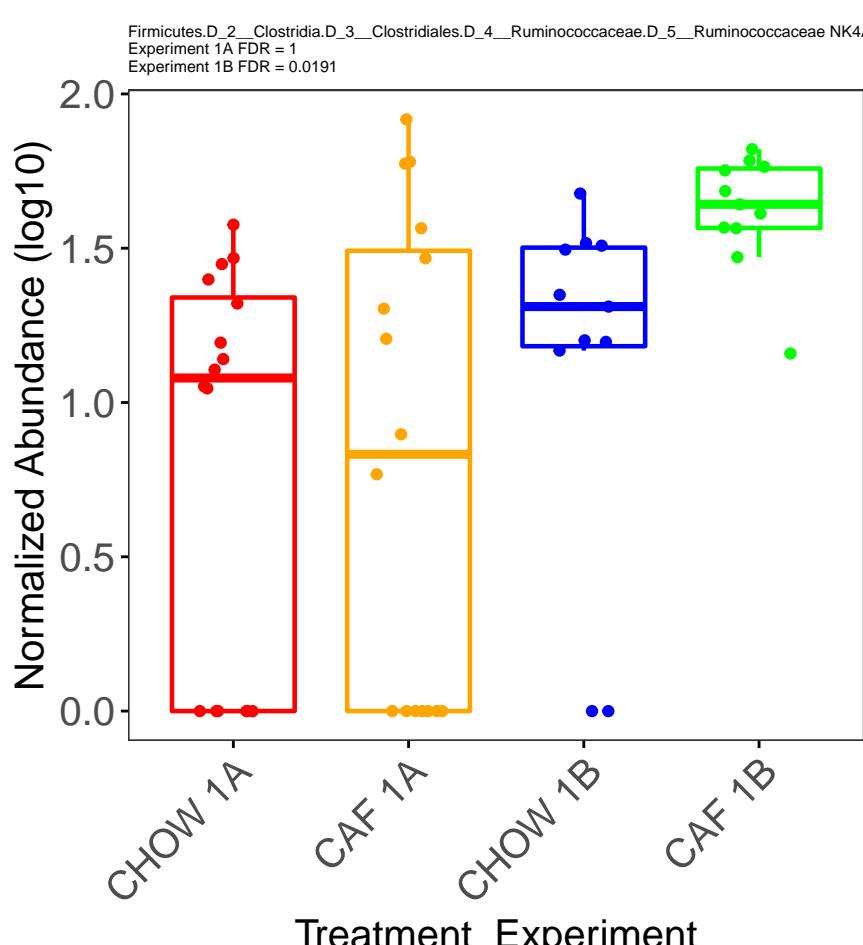
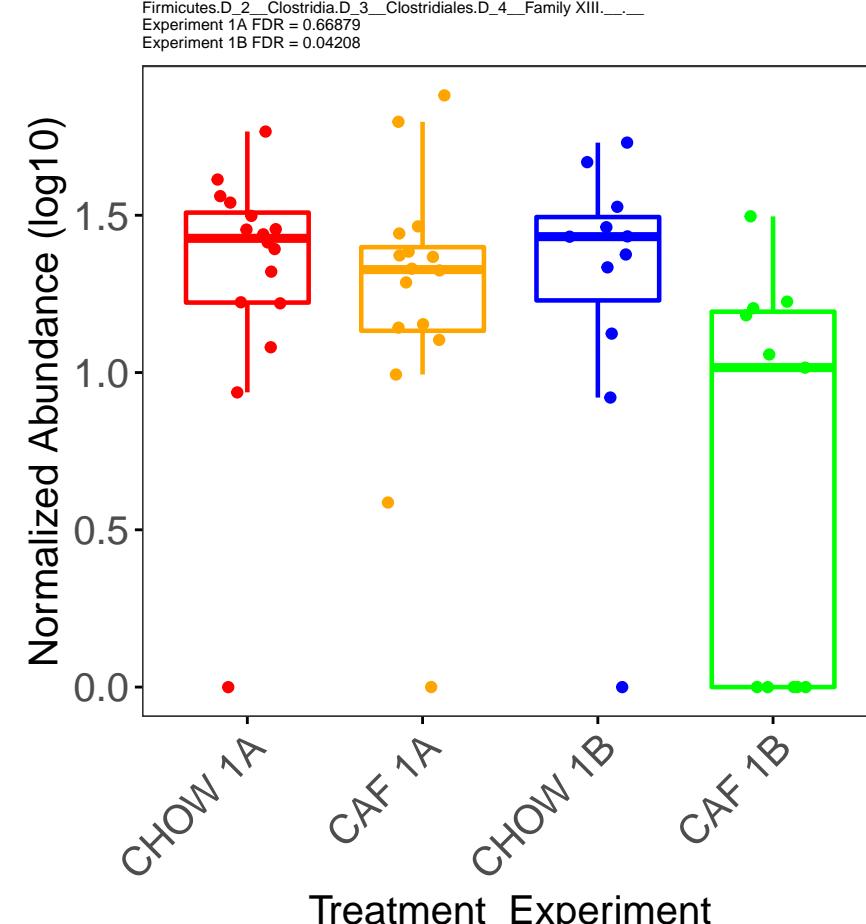
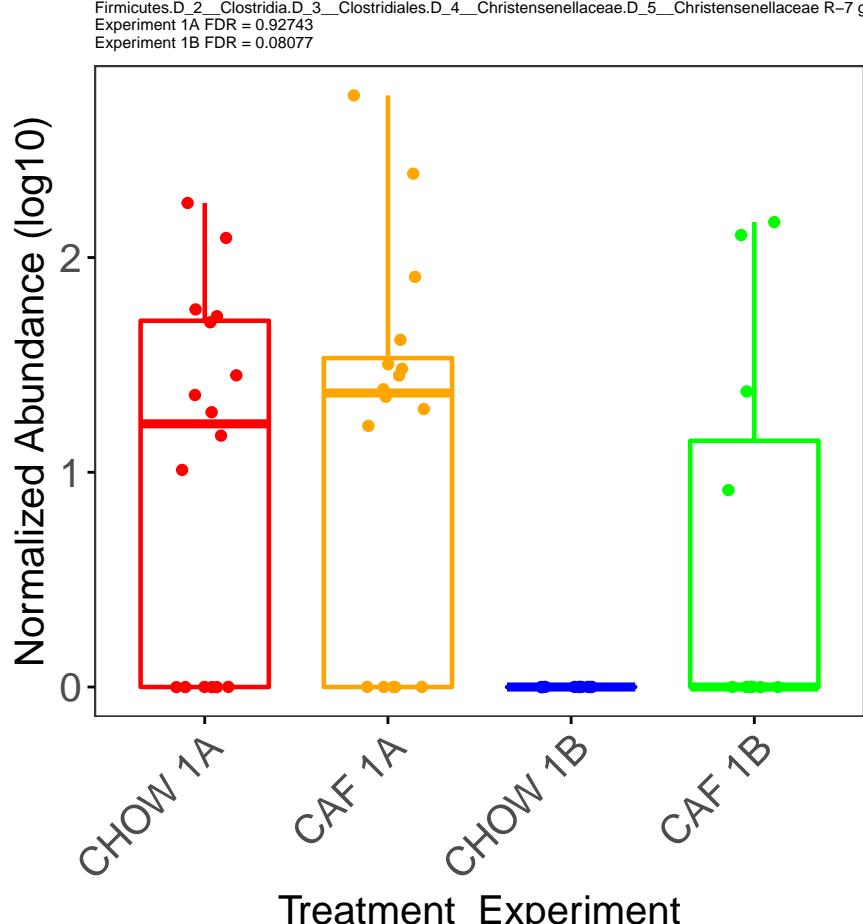
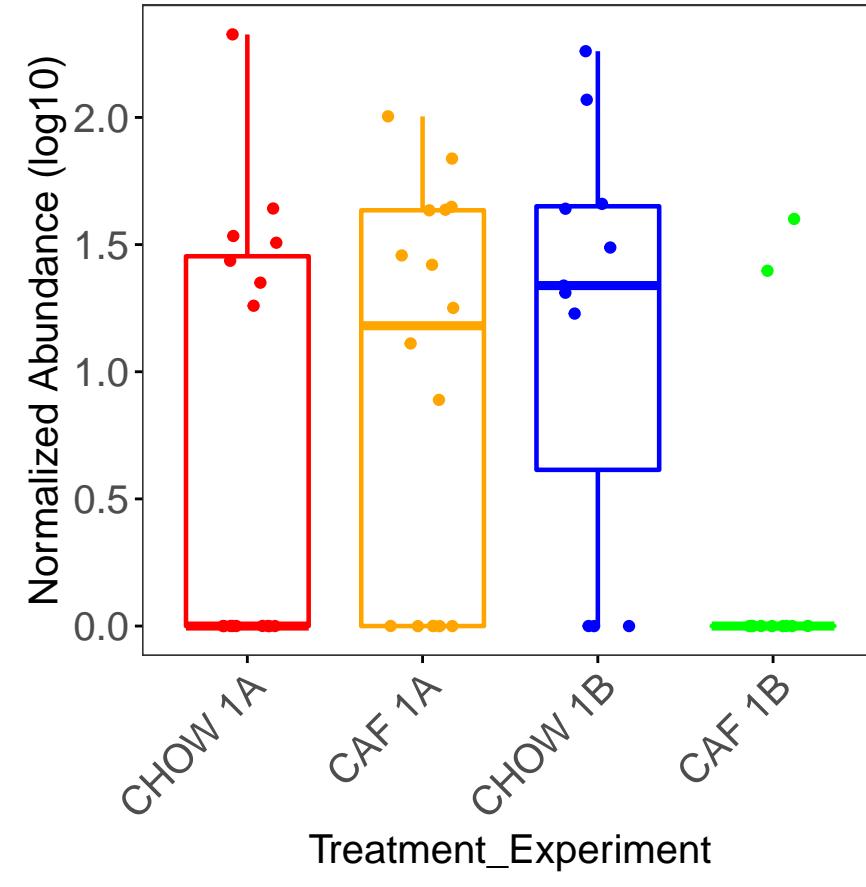
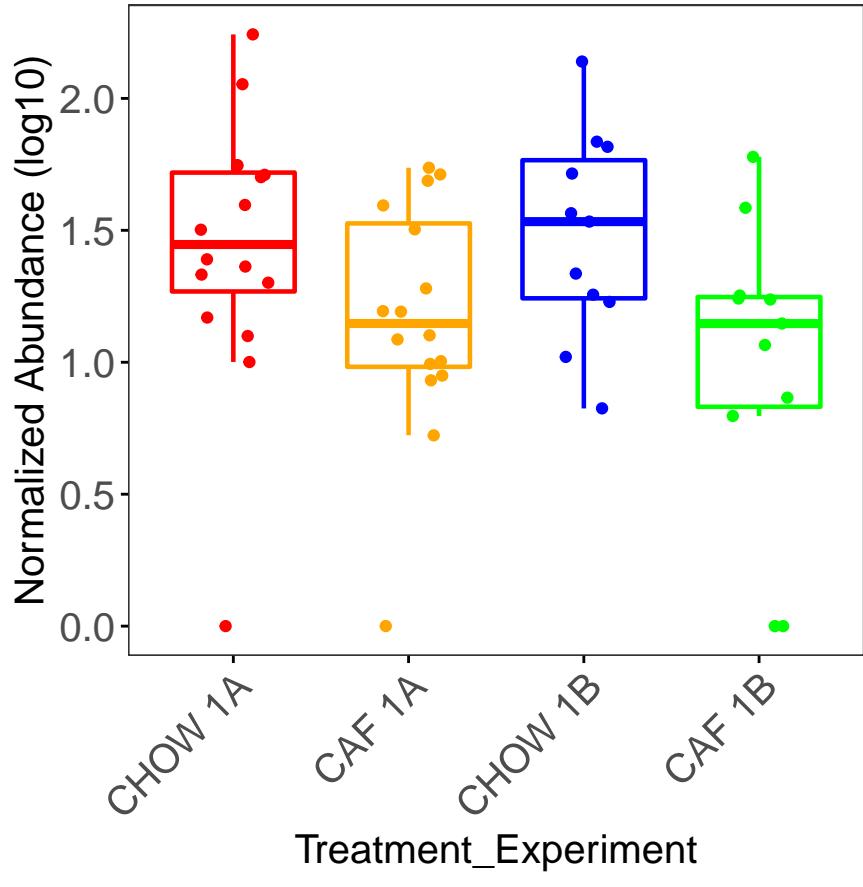
Firmicutes.D_2_Clostridia.D_3_Clostridiales.D_4_Ruminococcaceae.D_5_Ruminococcaceae UCG-005.D_6_uncultured bacterium adhufec108
Experiment 1A FDR = 0.06069
Experiment 1B FDR = 0.35305

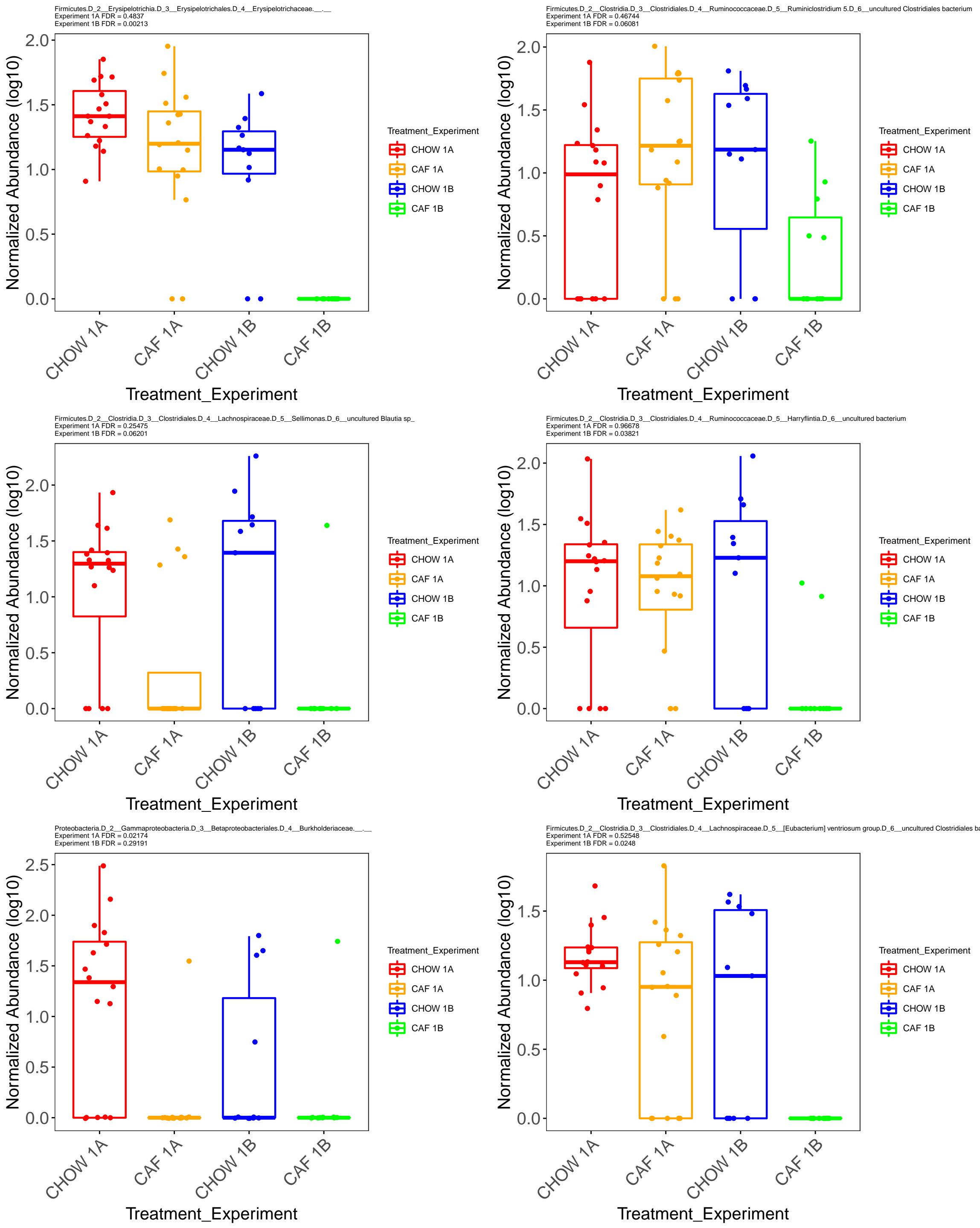


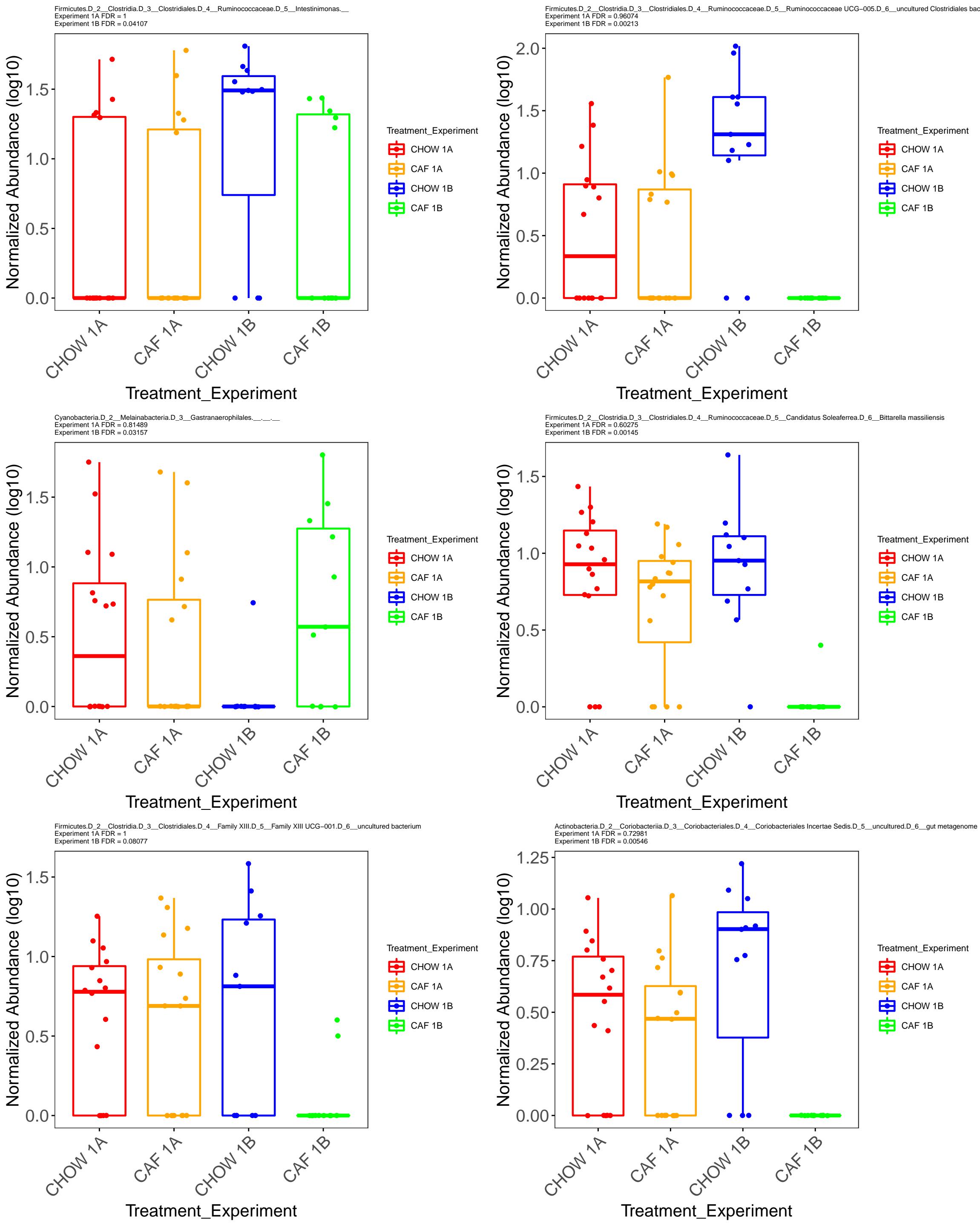


Proteobacteria.D_2_Delta-proteobacteria.D_3_Desulfovibrionales.D_4_Desulfovibrionaceae...
Experiment 1A FDR = 0.3226
Experiment 1B FDR = 0.09152

Firmicutes.D_2_Clostridia.D_3_Clostridiales.D_4_Lachnospiraceae.D_5_[Eubacterium] xylophilum group.D_6_uncultured bacterium
Experiment 1A FDR = 0.72981
Experiment 1B FDR = 0.04333

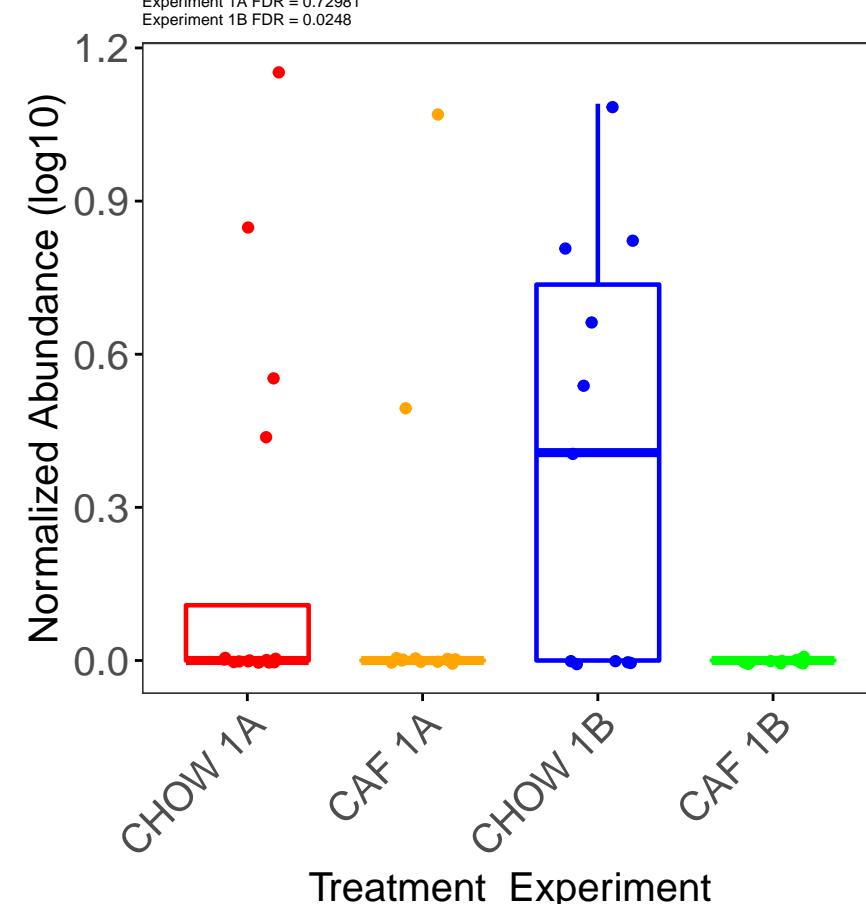
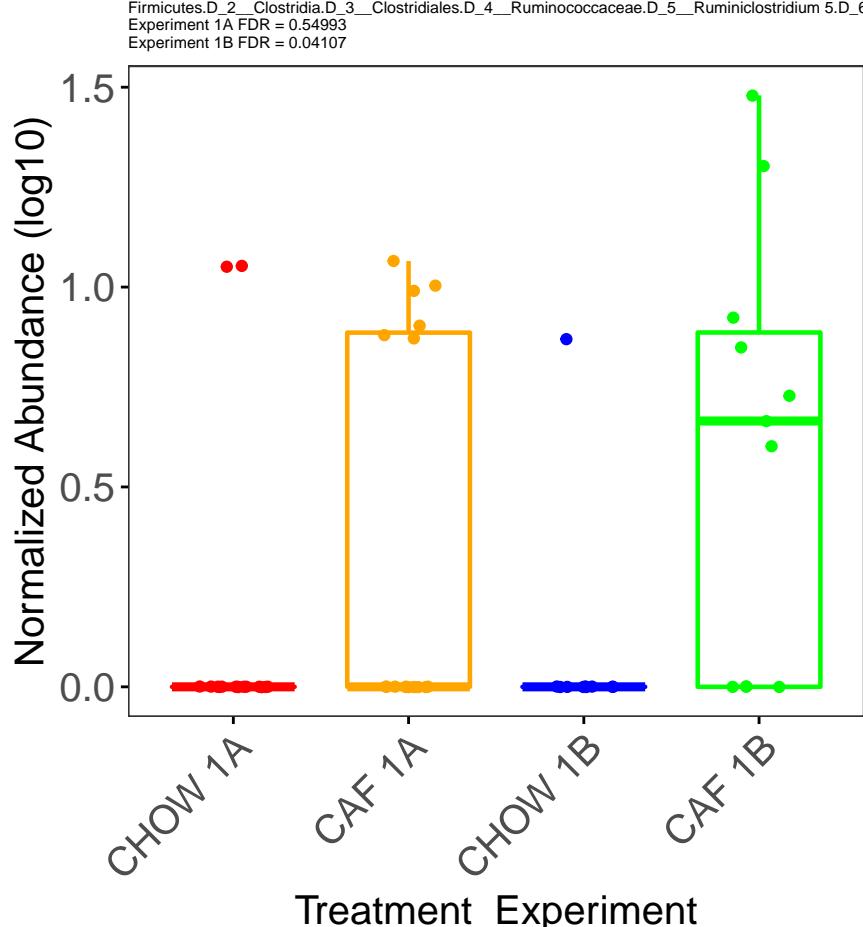
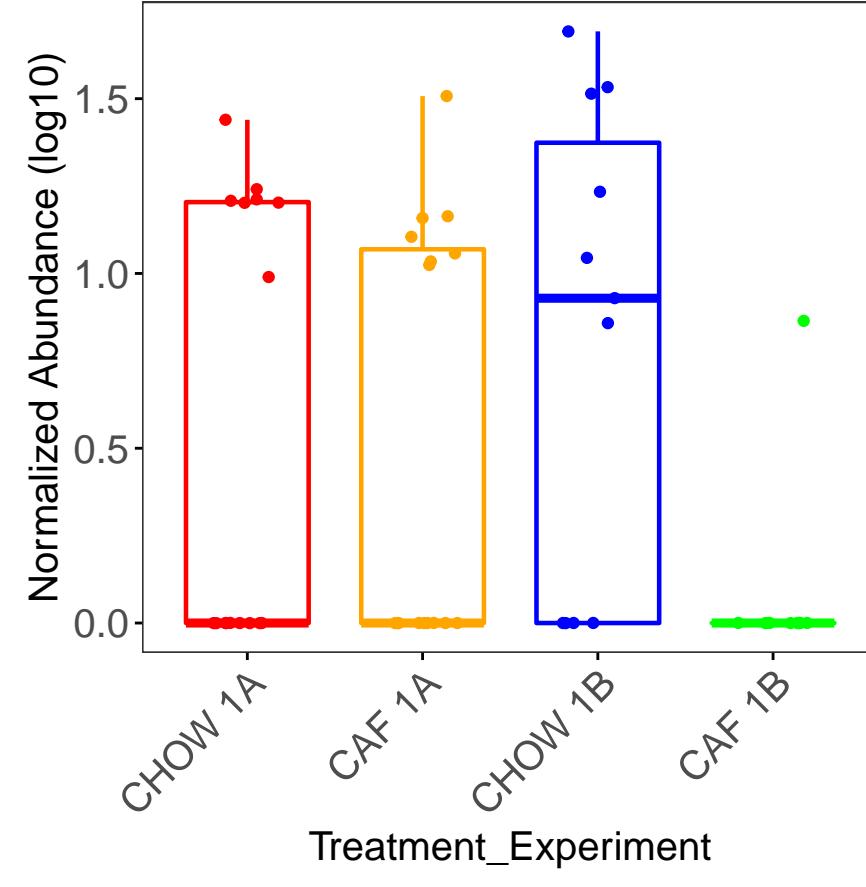
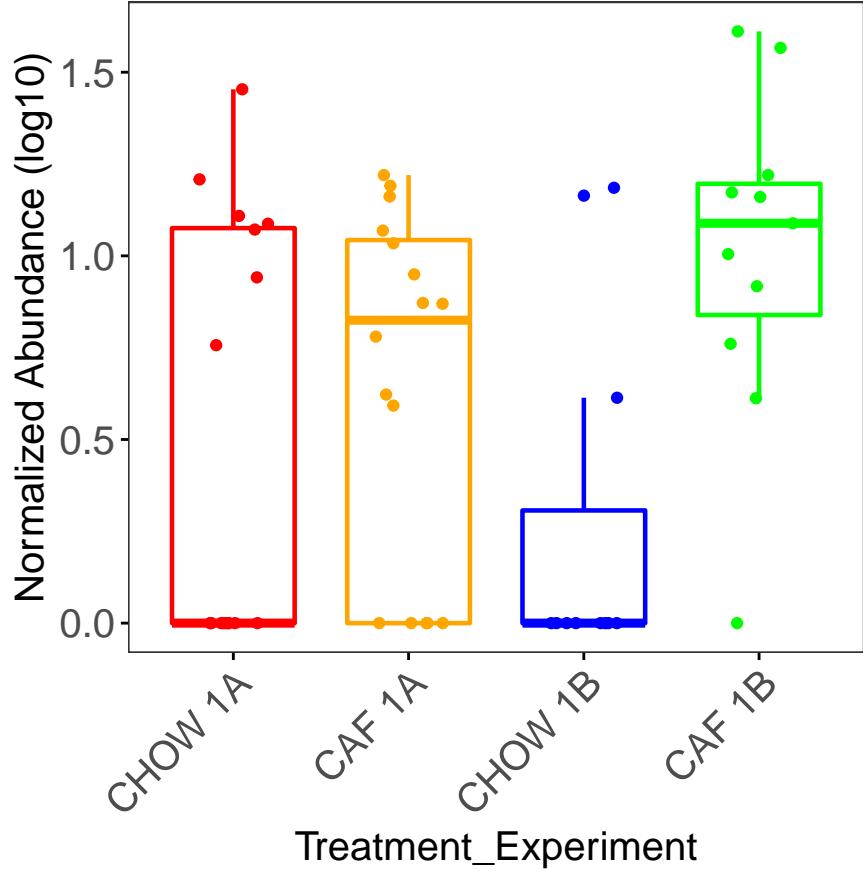






Firmicutes.D_2_Clostridia.D_3_Clostridiales.D_4_Lachnospiraceae.D_5_Tyzzerella.
Experiment 1A FDR = 0.80194
Experiment 1B FDR = 0.02775

Firmicutes.D_2_Clostridia.D_3_Clostridiales.D_4_Ruminococcaceae.D_5_GCA-900066225.D_6_uncultured Anaerotruncus sp.
Experiment 1A FDR = 0.92743
Experiment 1B FDR = 0.0274



Supplemental Figure 3 Abundance of the gut microbiota in the CAF vs CHOW groups when analyzed either before (1A) or after (1B) healthy dietary intervention. CHOW: chow-fed; CAF: cafeteria diet; FDR: false discovery rate