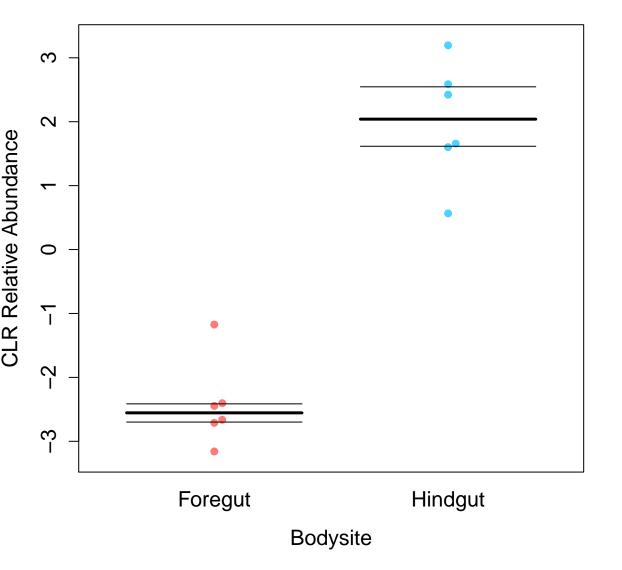
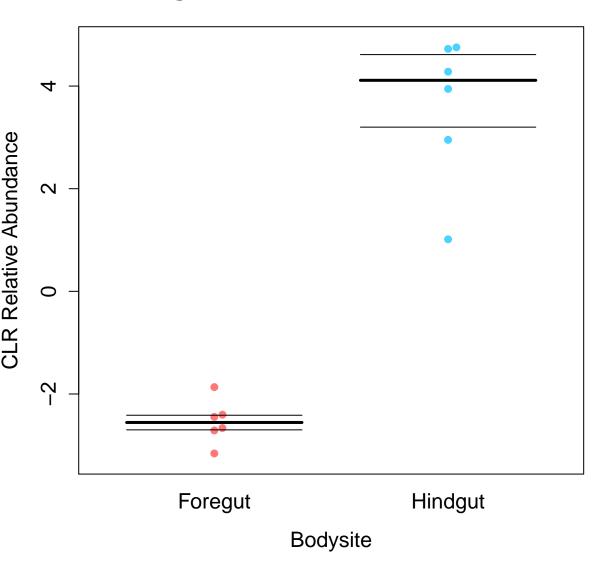
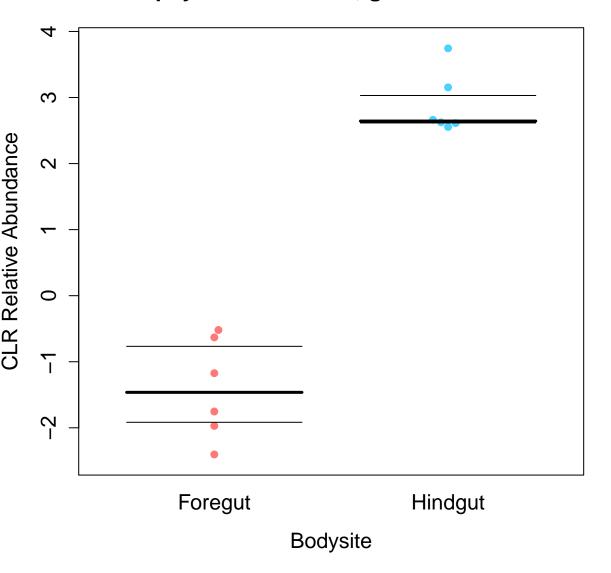
f__Coriobacteriaceae; g__Adlercreutzia



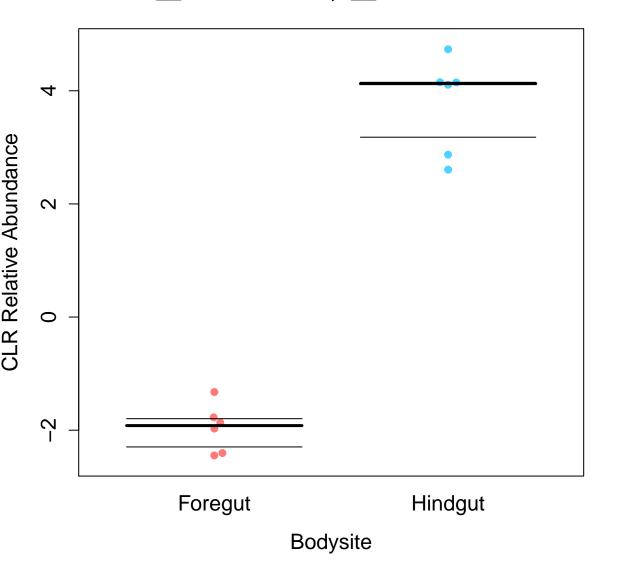
g__Bacteroides; s__uniformis



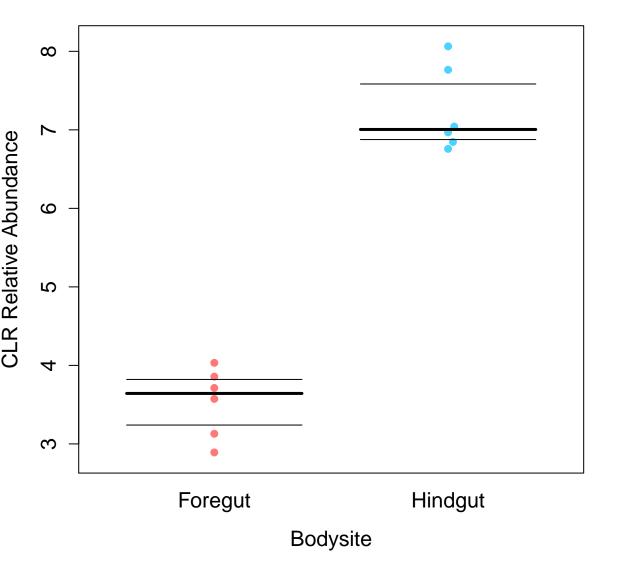
f__Porphyromonadaceae; g__Parabacteroides



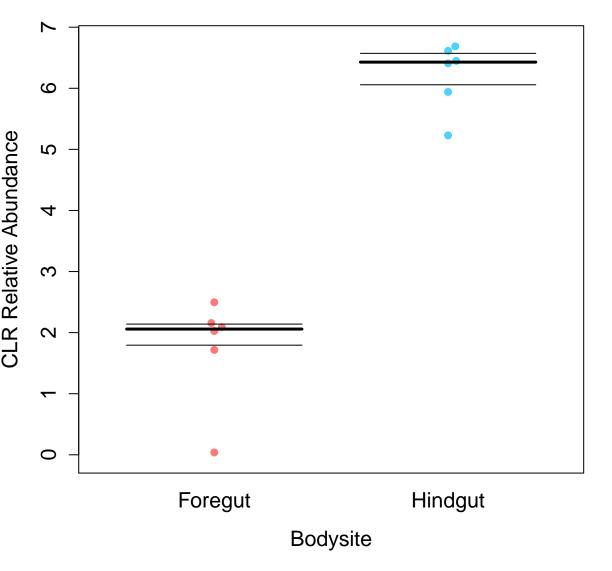
o__Bacteroidales; f__Rikenellaceae



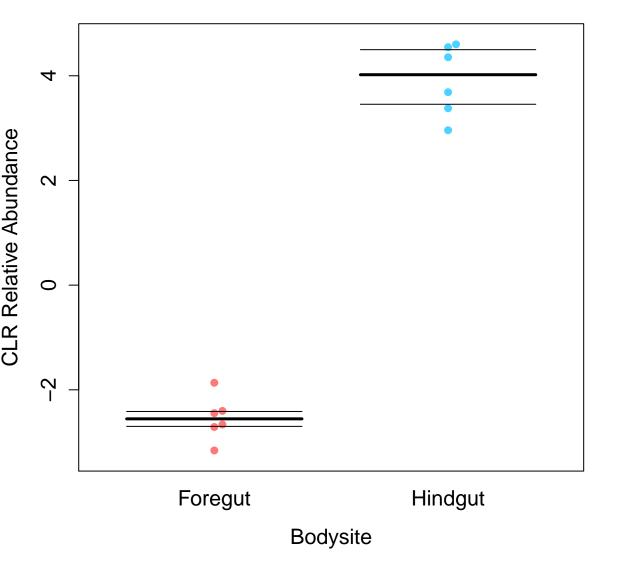
o__Bacteroidales; f__S24-7



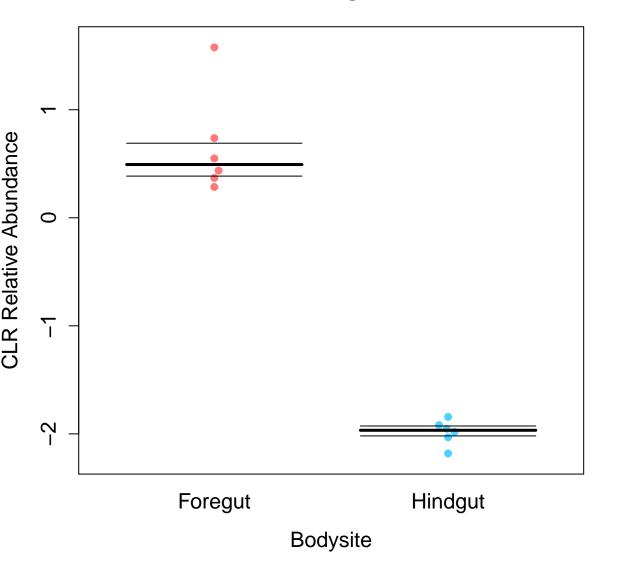




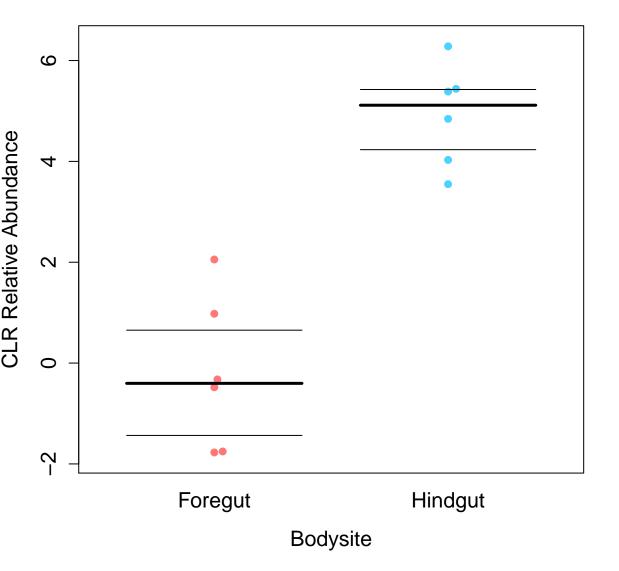
o__Erysipelotrichales; f__Erysipelotrichaceae



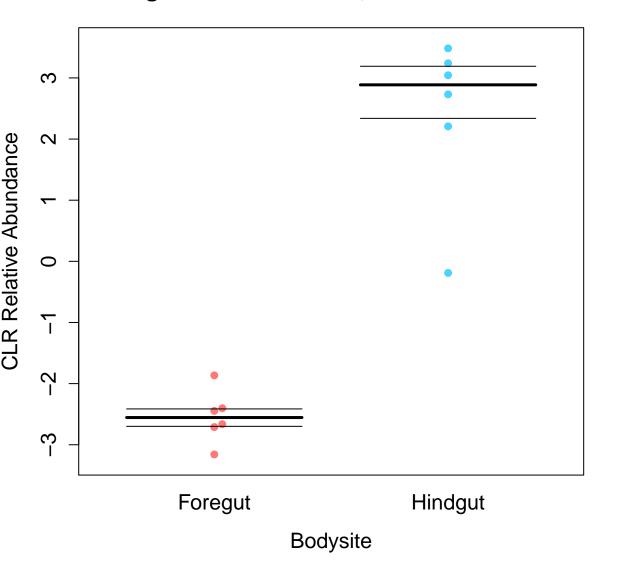
f__Pasteurellaceae; g__Actinobacillus



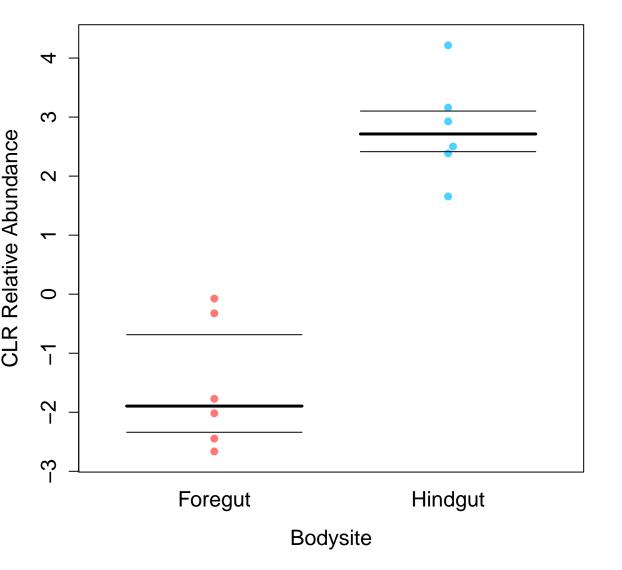




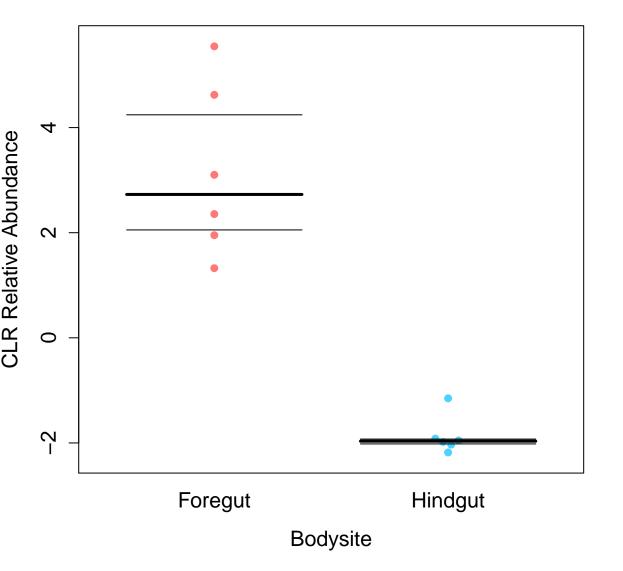
g__Parabacteroides; s__distasonis



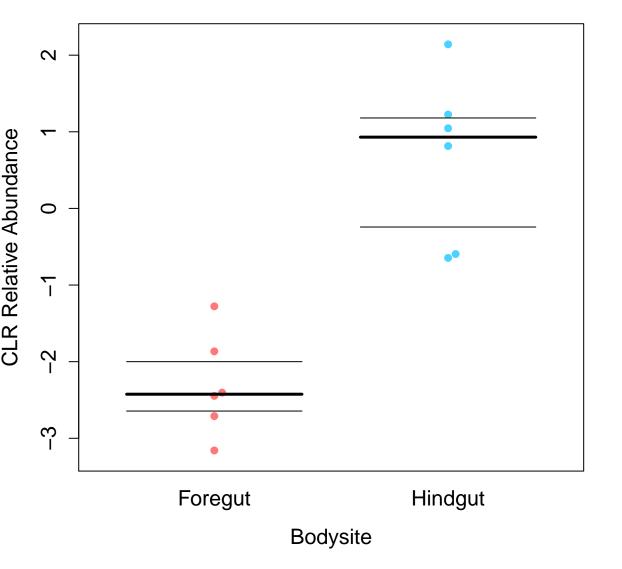
f__Lachnospiraceae; g__Roseburia



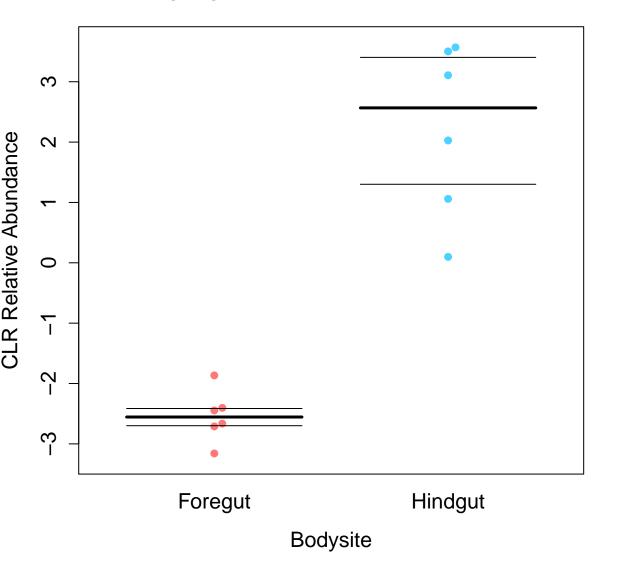
c__Bacilli; o__Lactobacillales



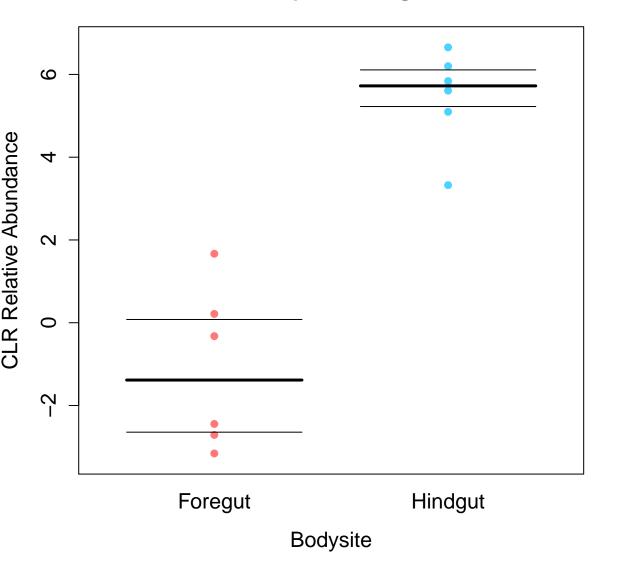
f__Helicobacteraceae; g__Flexispira



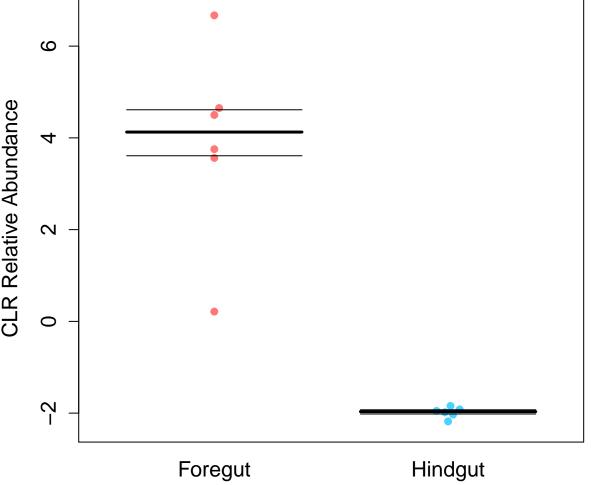
c__Alphaproteobacteria; o__Rickettsiales



f__Lachnospiraceae; g__Dorea

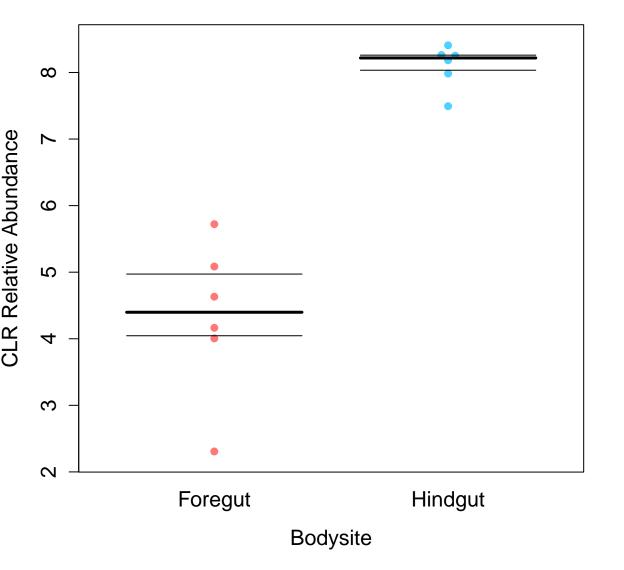


c__Bacilli; o__Bacillales

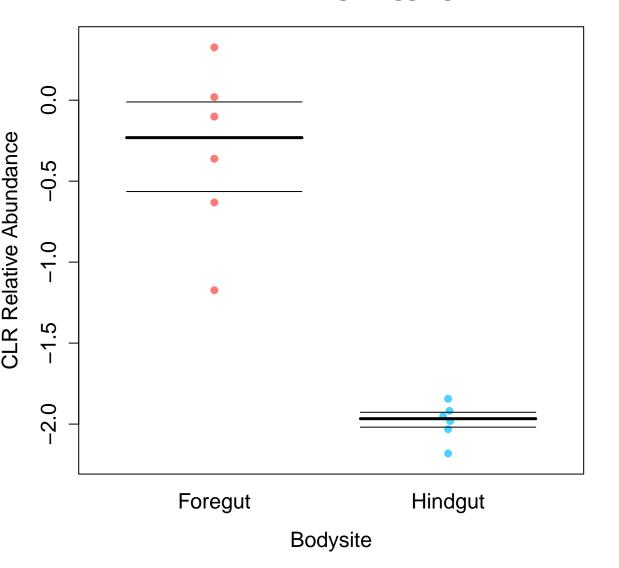


Bodysite

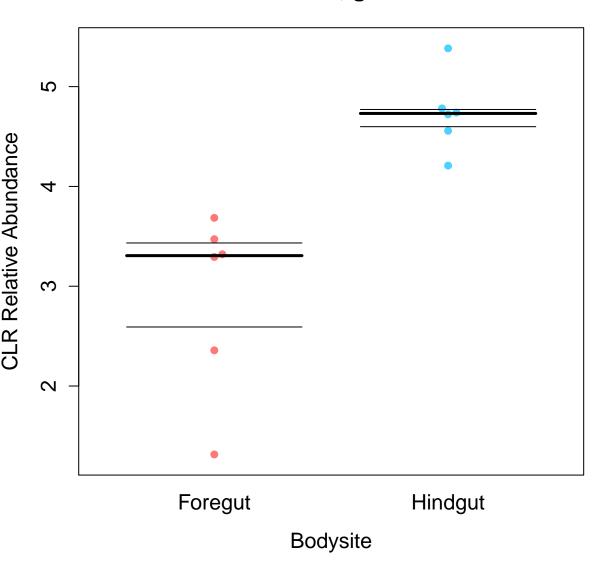
o__Clostridiales; f__Ruminococcaceae



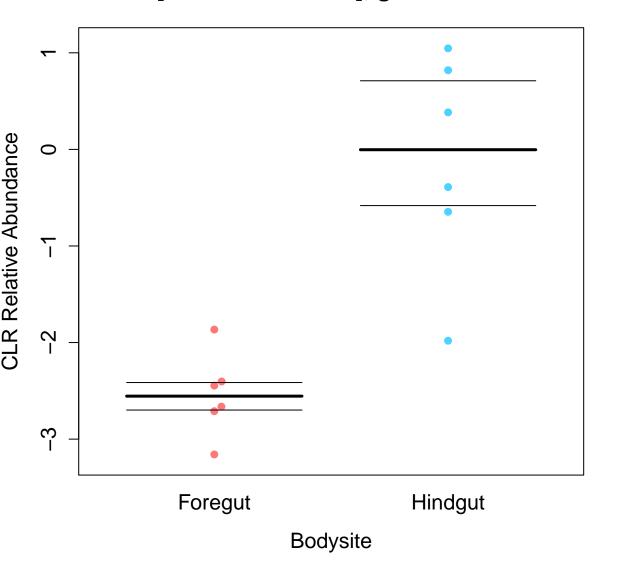




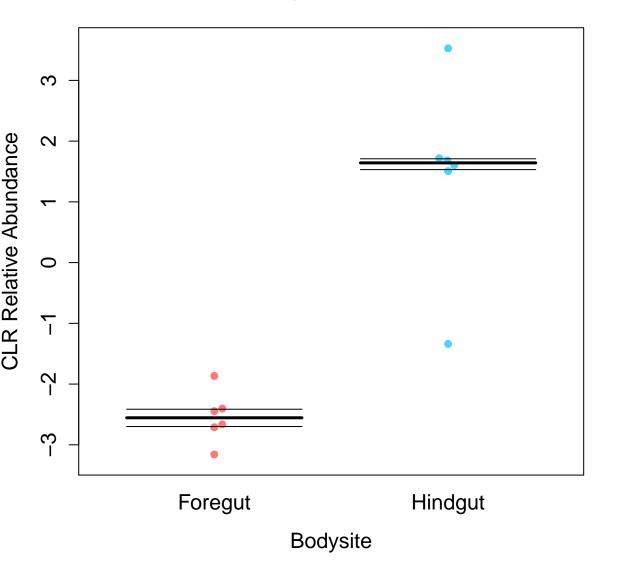
f__Clostridiaceae; g__Clostridium



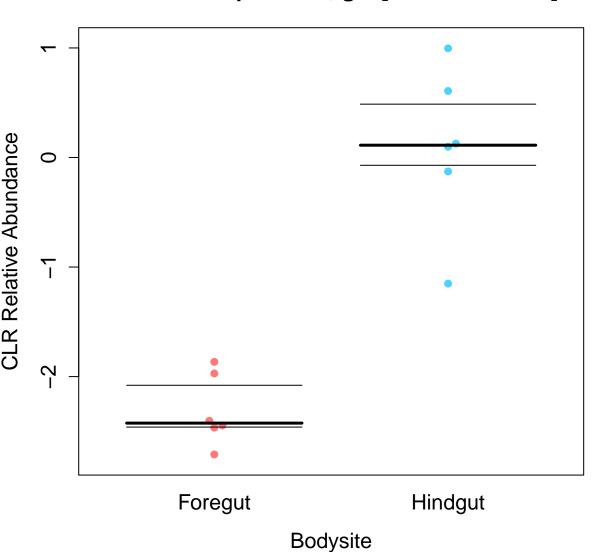
f__[Odoribacteraceae]; g__Odoribacter



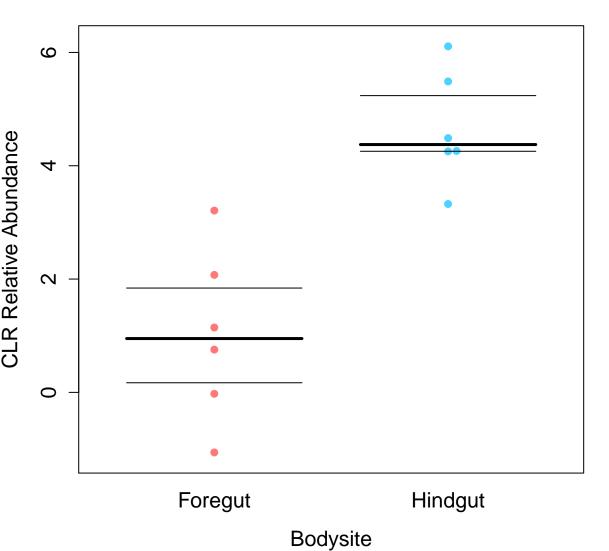
o__Clostridiales; f__Dehalobacteriaceae



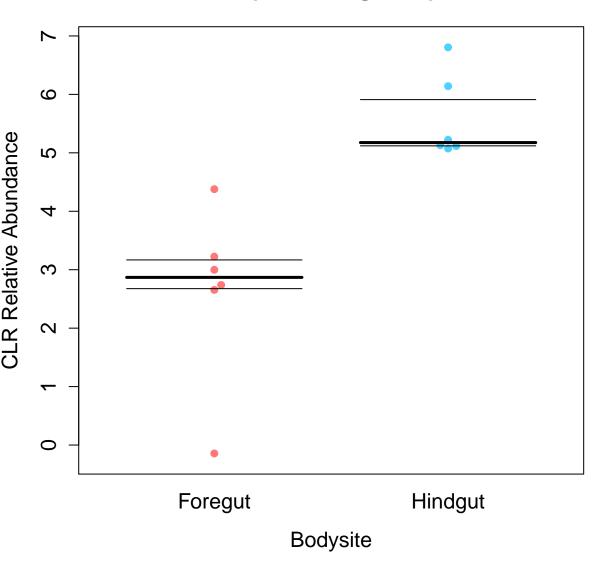
f_Lachnospiraceae; g_[Ruminococcus]



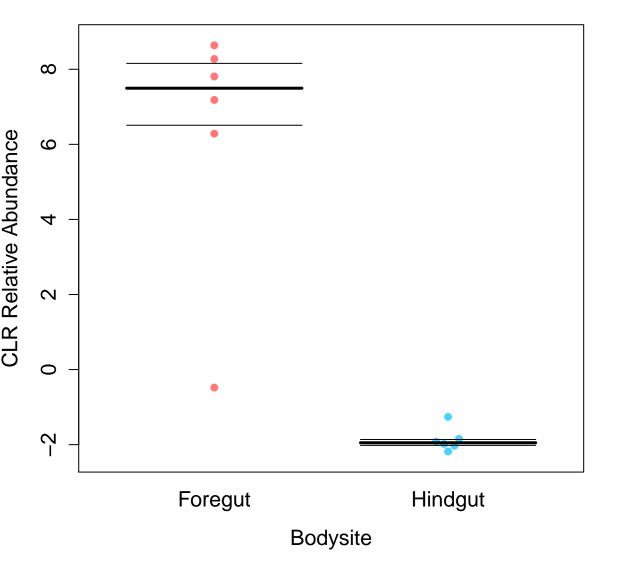




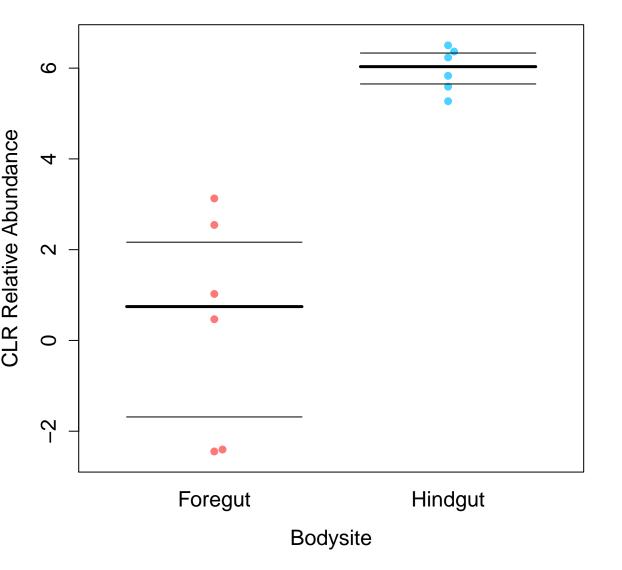
f__Lachnospiraceae; g__Coprococcus



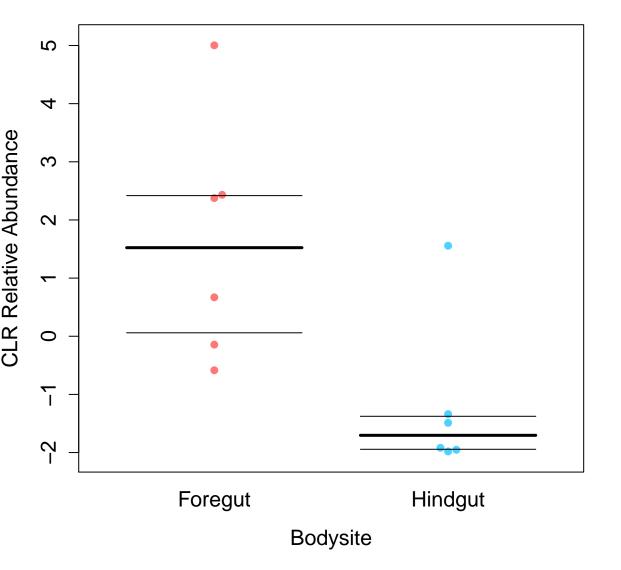
f__Moraxellaceae; g__Acinetobacter



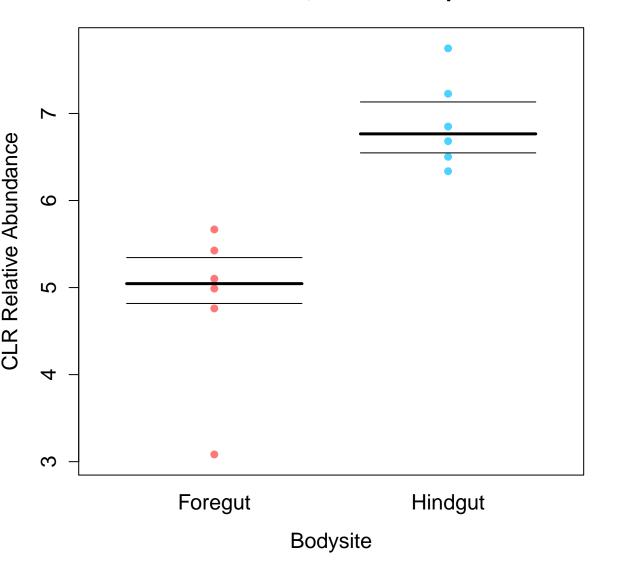
f__Bacteroidaceae; g__Bacteroides



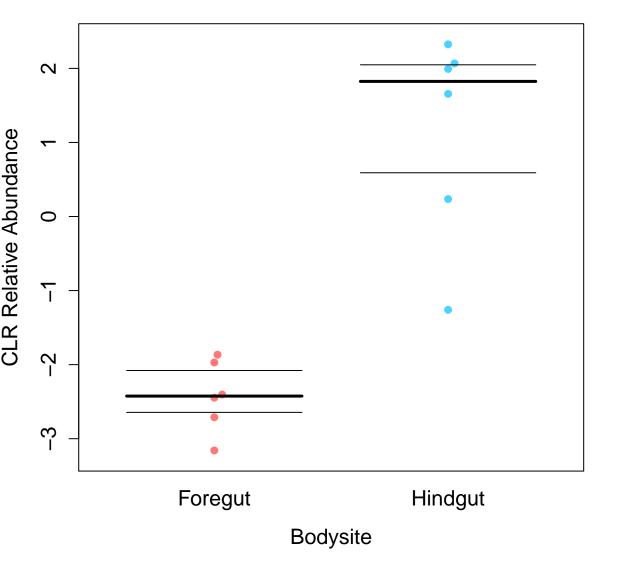
g__Streptococcus; s__alactolyticus



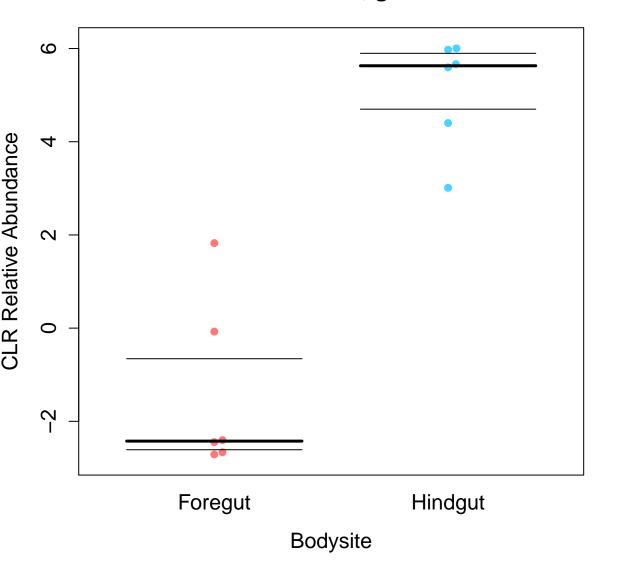
o__Clostridiales; f__Lachnospiraceae



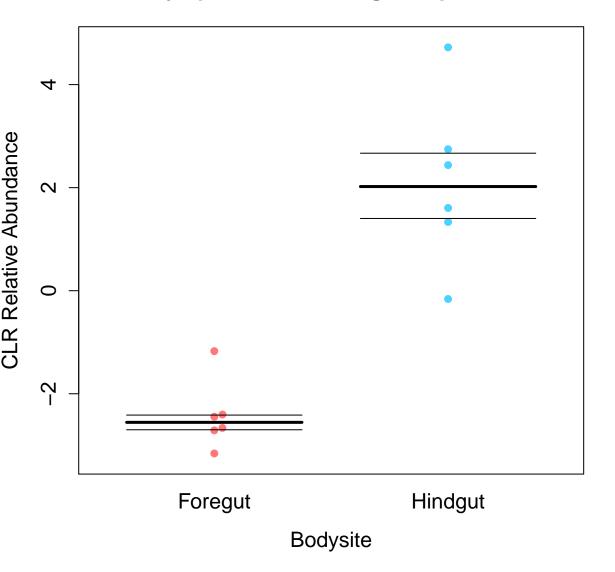
o__Clostridiales; f__Peptococcaceae



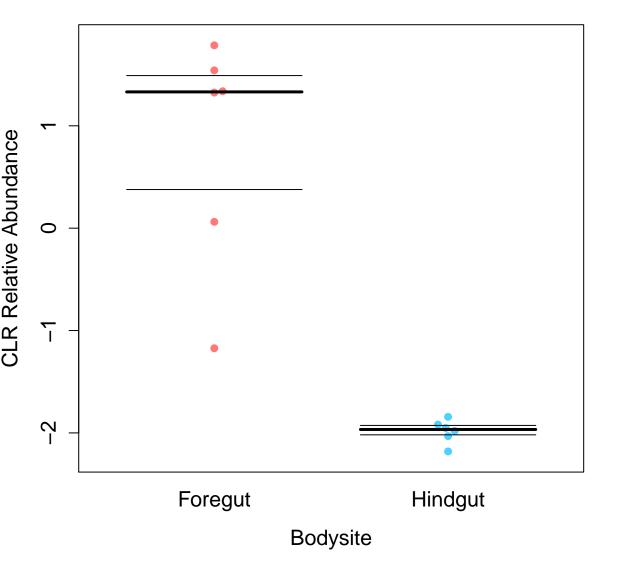
f__Ruminococcaceae; g__Clostridium



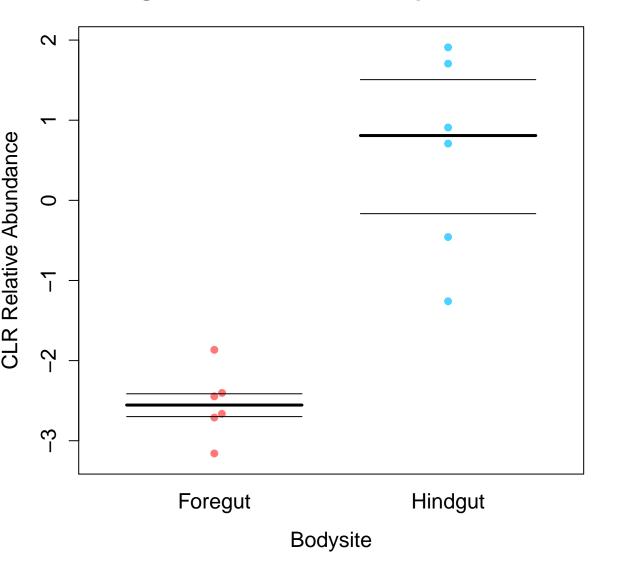
f__Erysipelotrichaceae; g__Coprobacillus



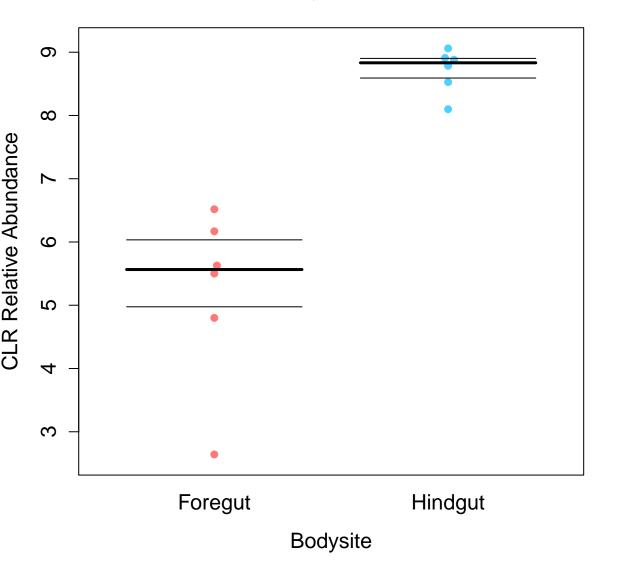
g__Aggregatibacter; s__pneumotropica

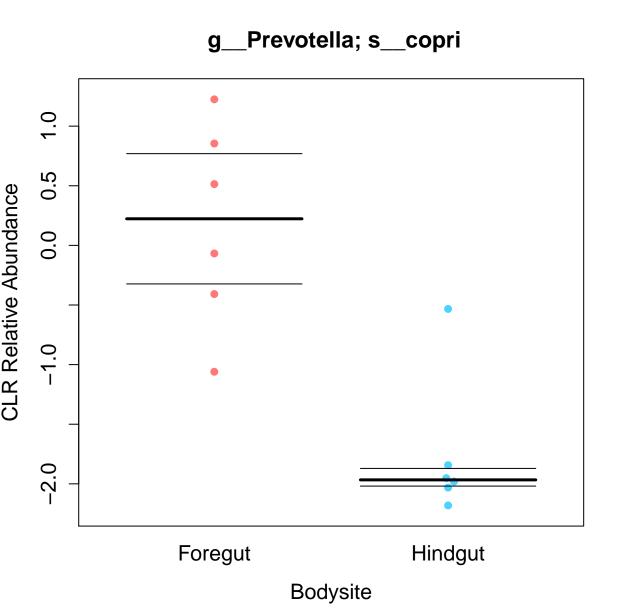


g__Faecalibacterium; s__prausnitzii

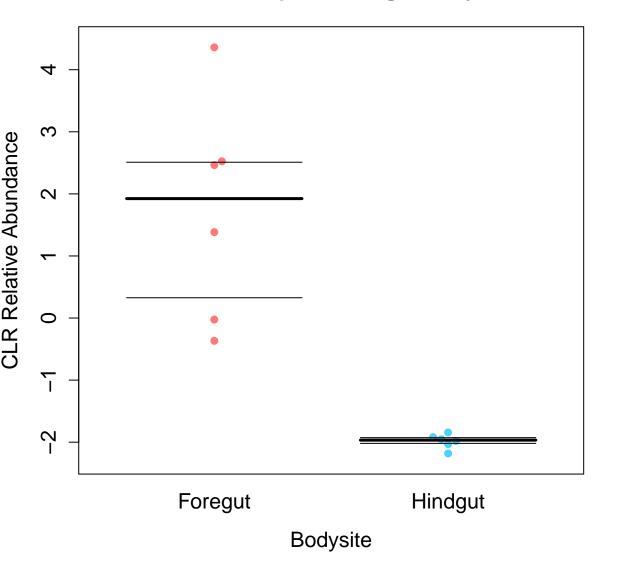


c__Clostridia; o__Clostridiales

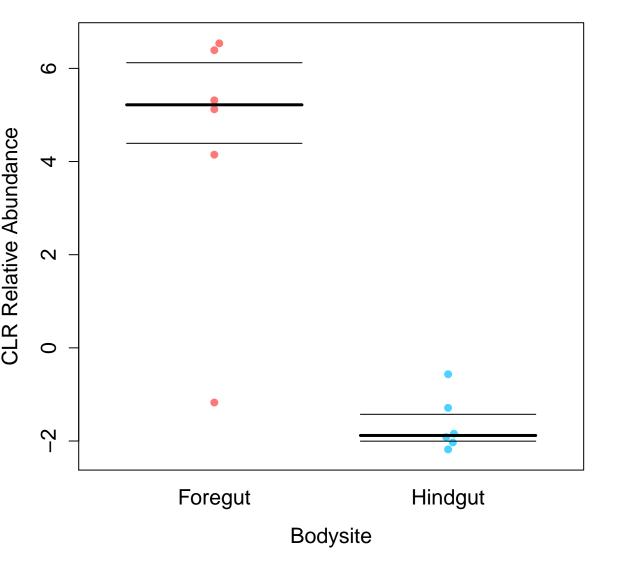




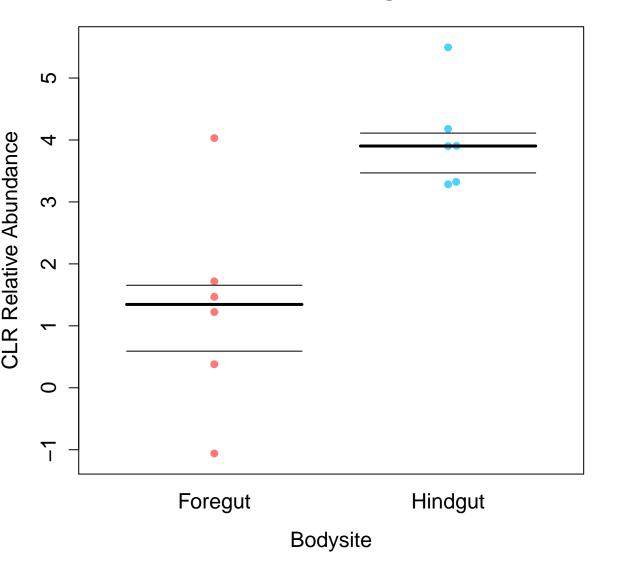
f__Lachnospiraceae; g__Moryella



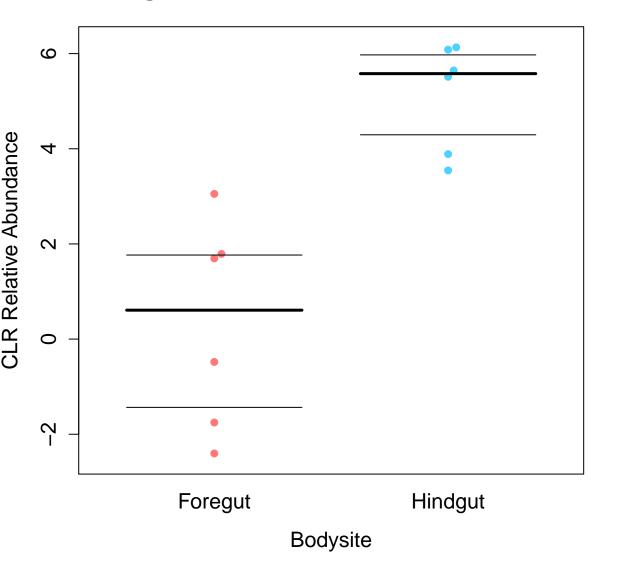
o__Bacillales; f__Planococcaceae



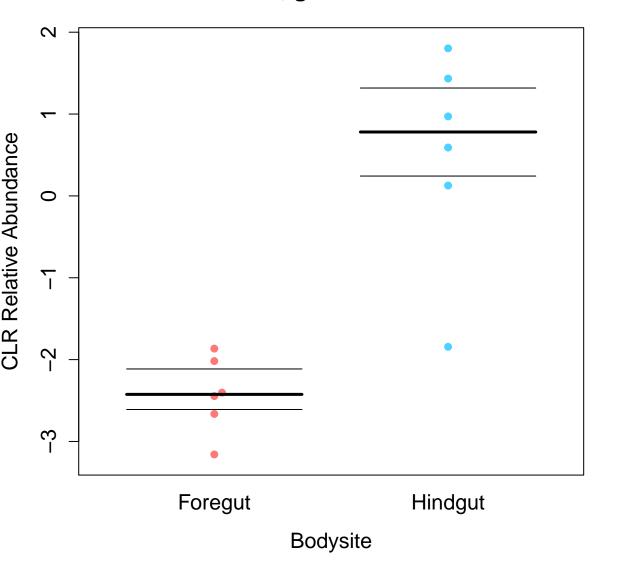
o__Clostridiales; f__[Mogibacteriaceae]



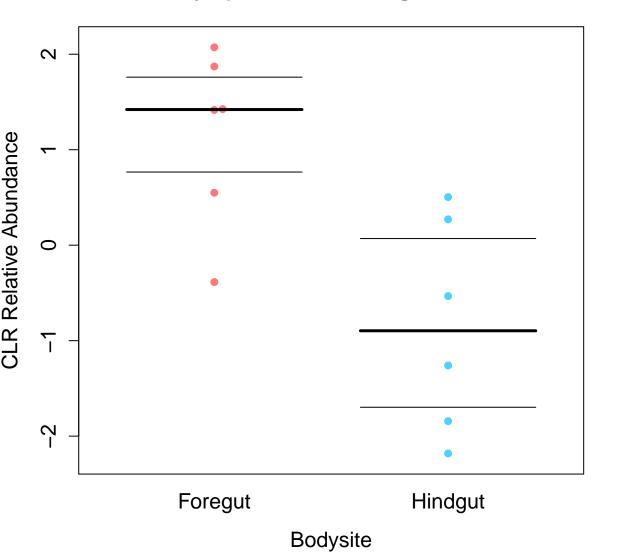
g__Ruminococcus; s__flavefaciens



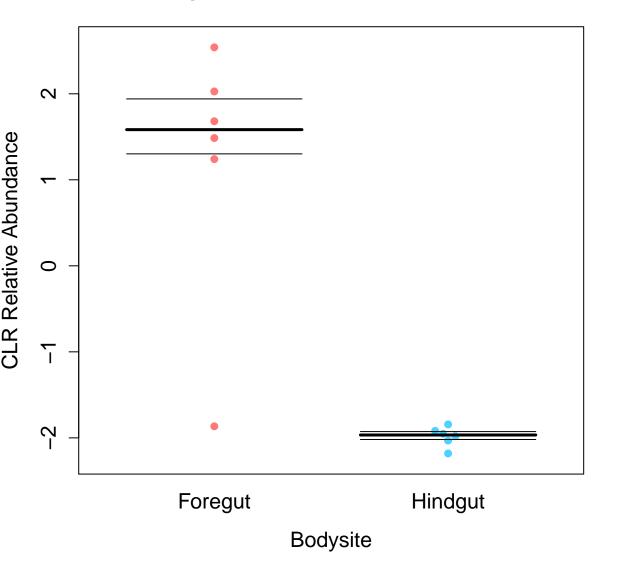
f__Veillonellaceae; g__Phascolarctobacterium



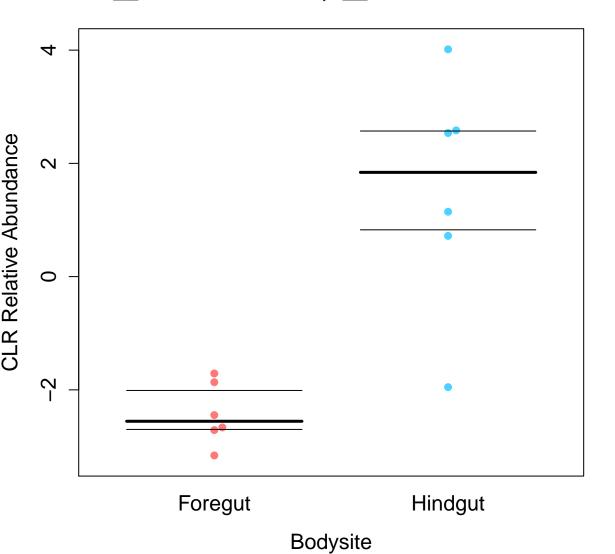
f__Erysipelotrichaceae; g__Bulleidia



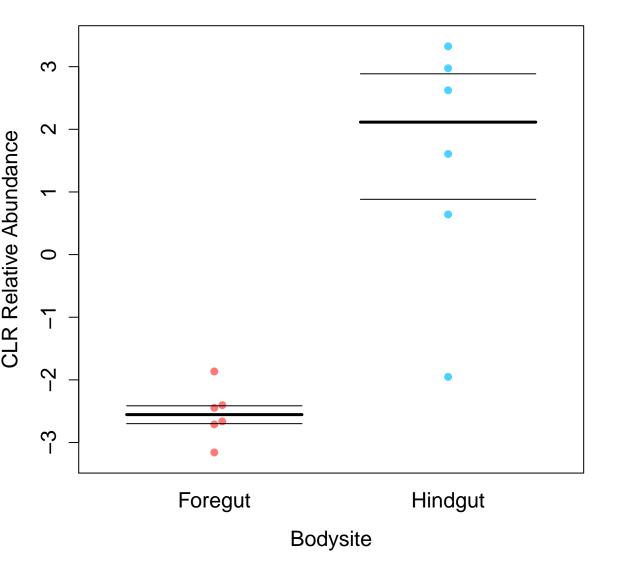
c__Gammaproteobacteria; o__Pseudomonadales



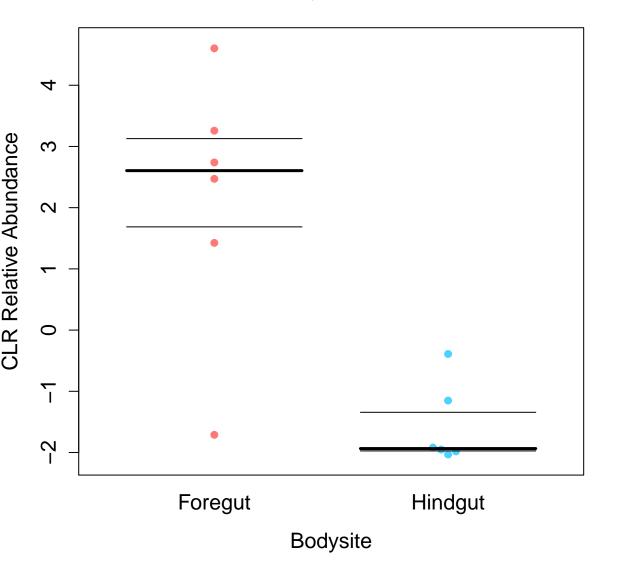
o__Bifidobacteriales; f__Bifidobacteriaceae



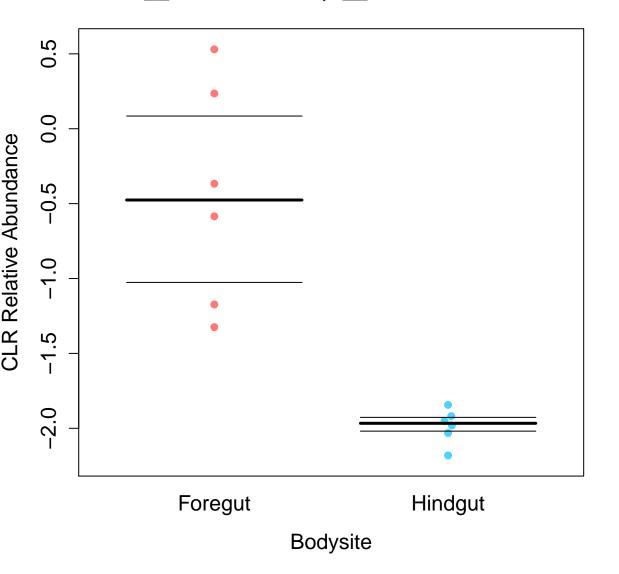




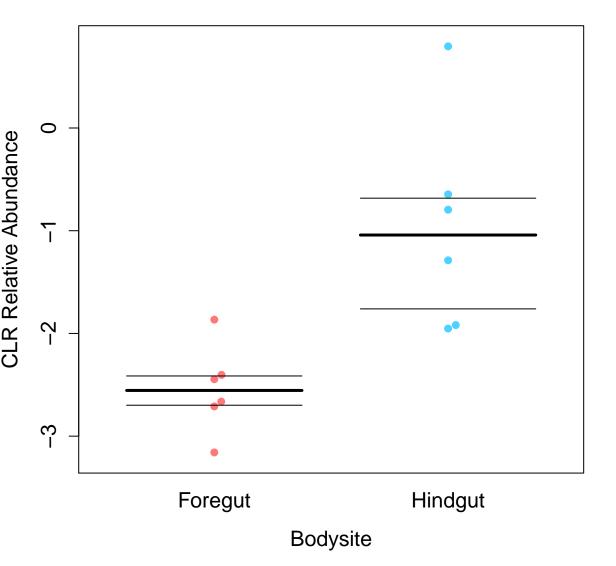
o__Lactobacillales; f__Leuconostocaceae



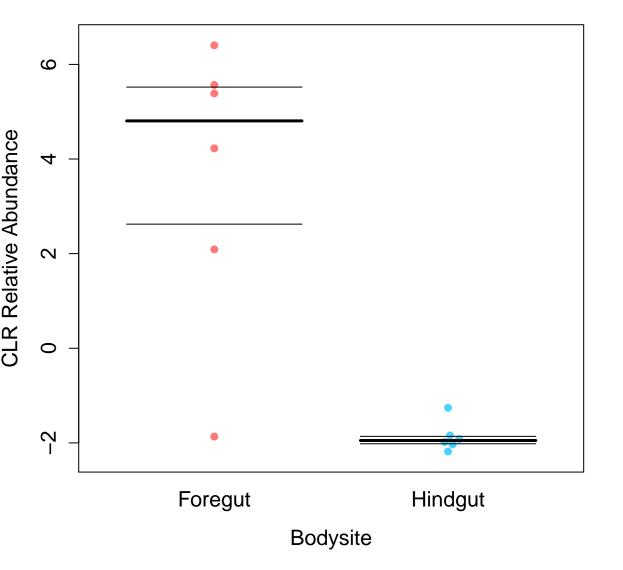
o__Pasteurellales; f__Pasteurellaceae



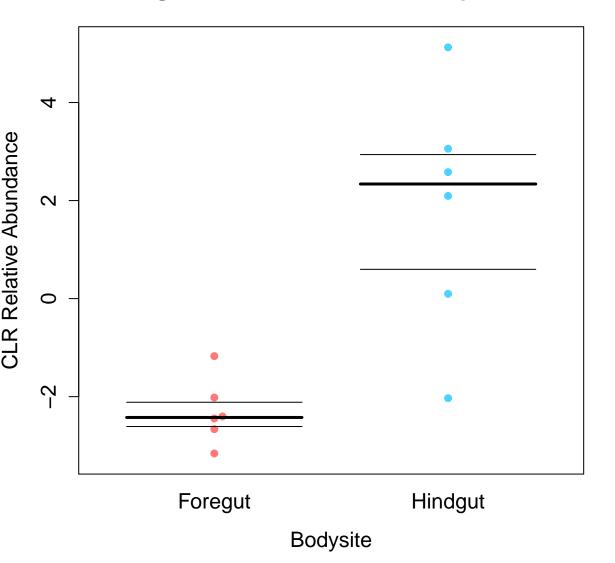


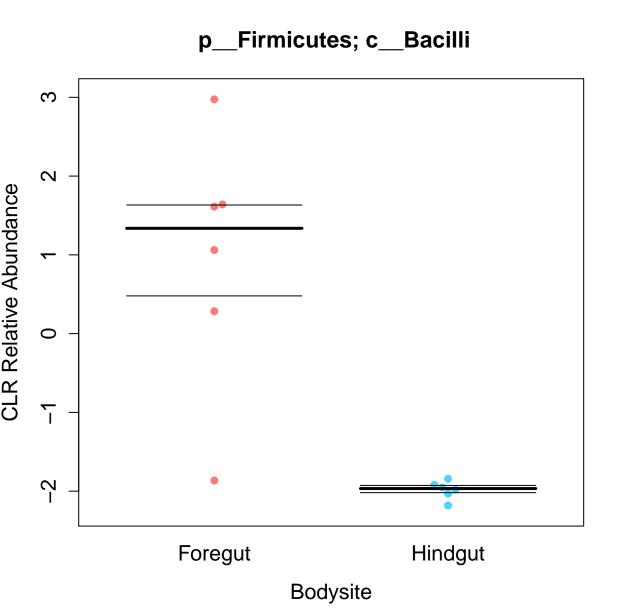


g__Acinetobacter; s__guillouiae

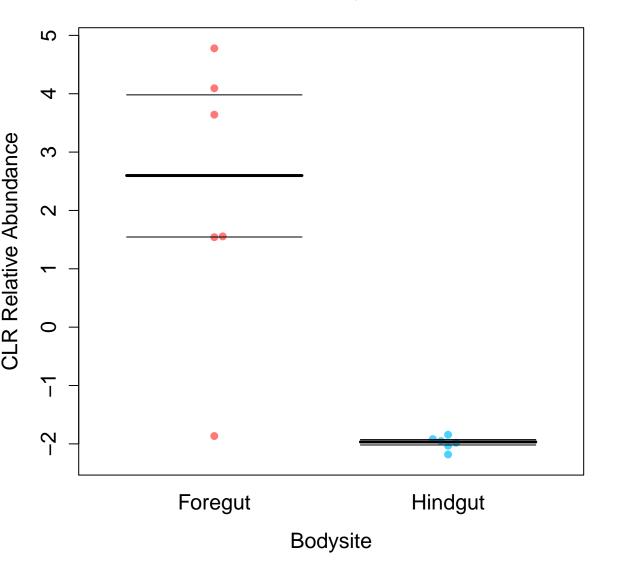


g__Akkermansia; s__muciniphila

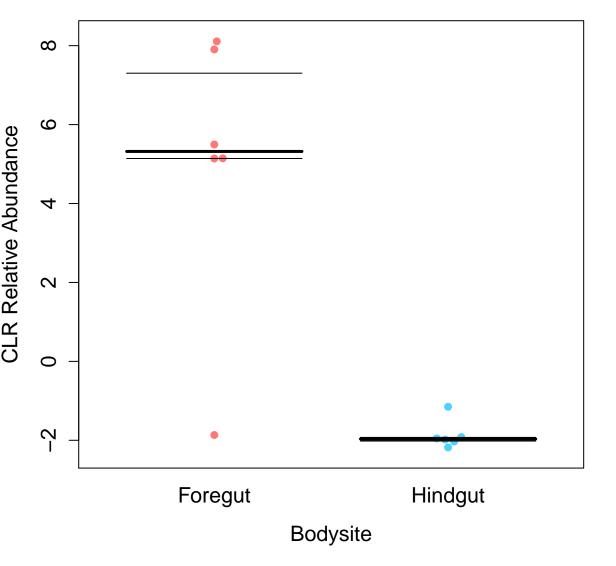




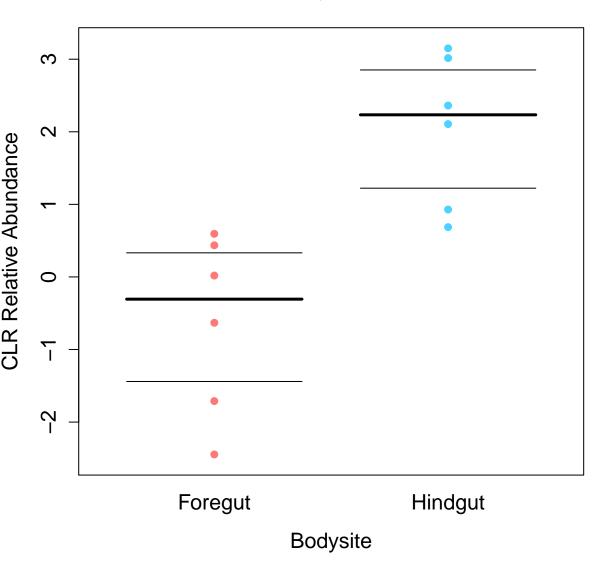
o__Pseudomonadales; f__Moraxellaceae



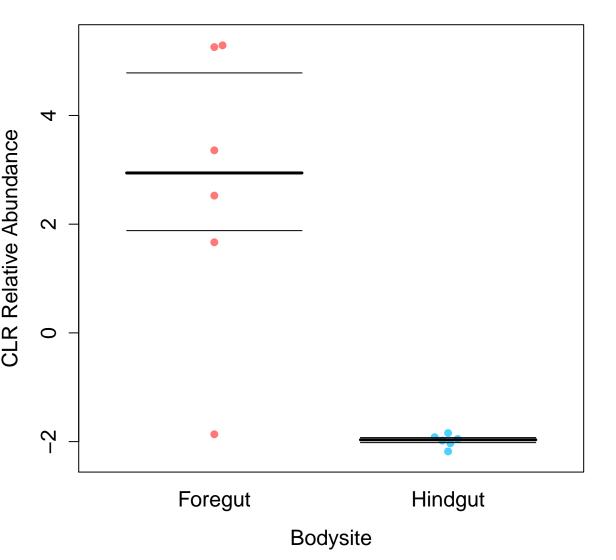
g__Acinetobacter; s__johnsonii



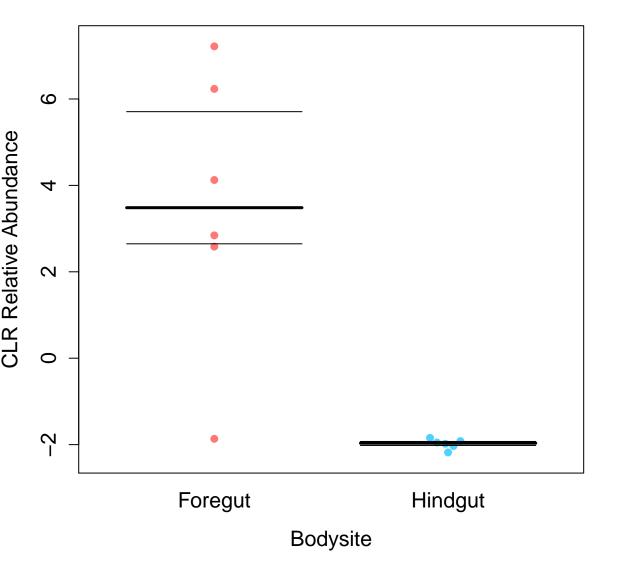
o__Clostridiales; f__Veillonellaceae



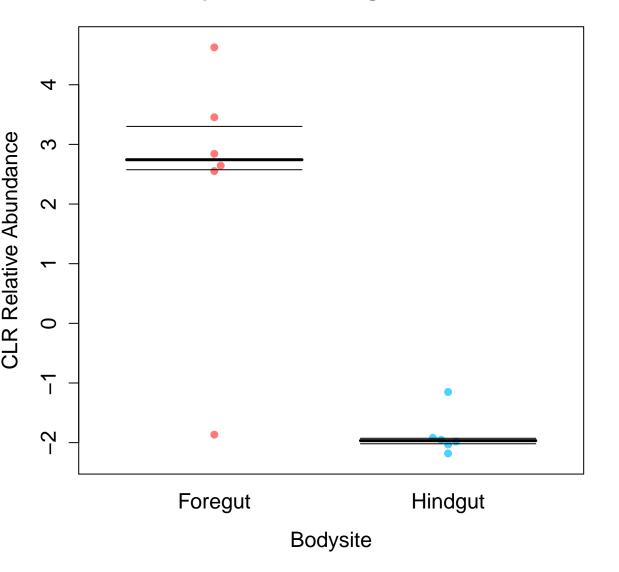




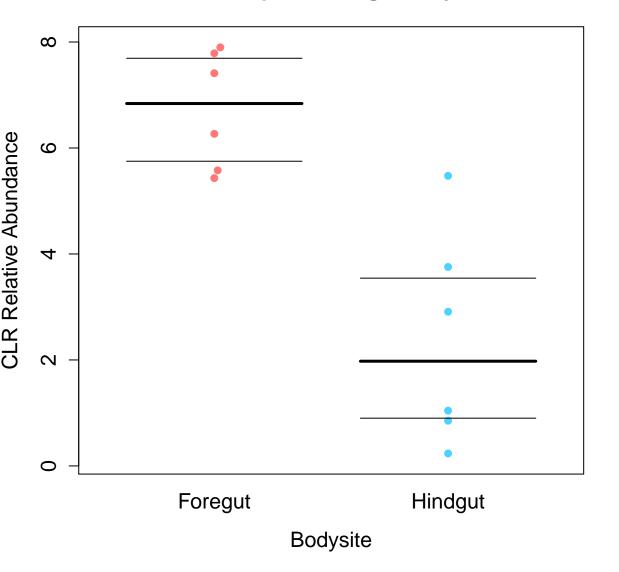




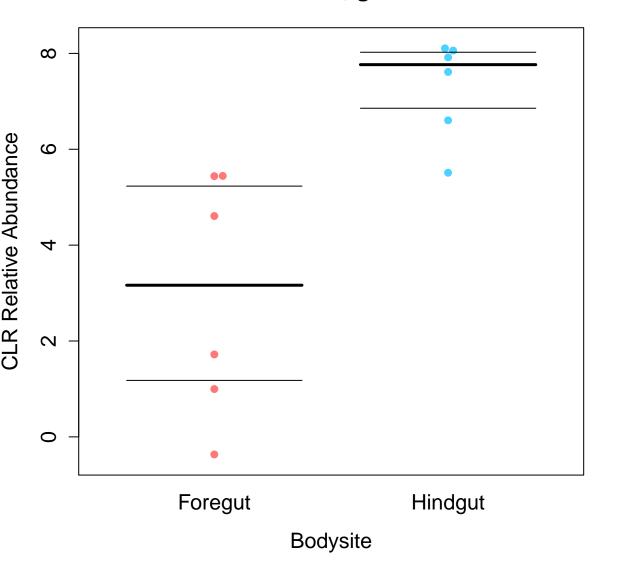
f_Streptococcaceae; g_Lactococcus



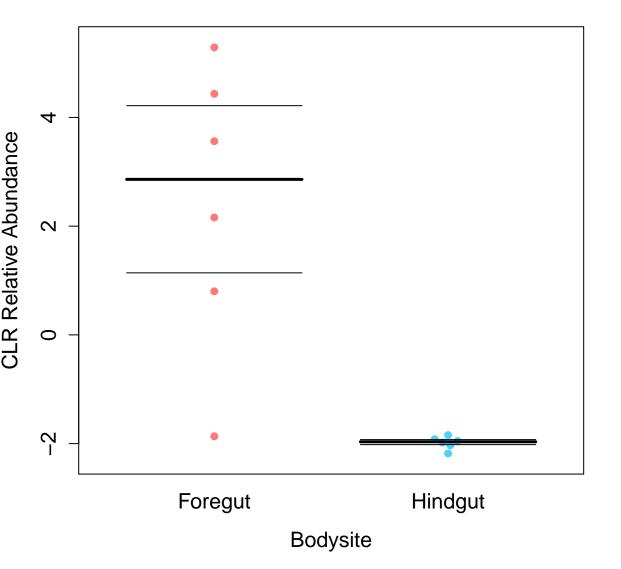
f__Lachnospiraceae; g__Butyrivibrio

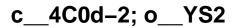


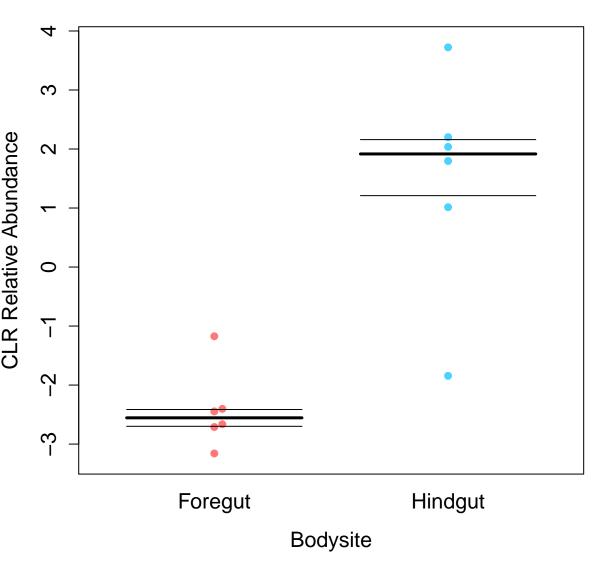
f__Ruminococcaceae; g__Ruminococcus



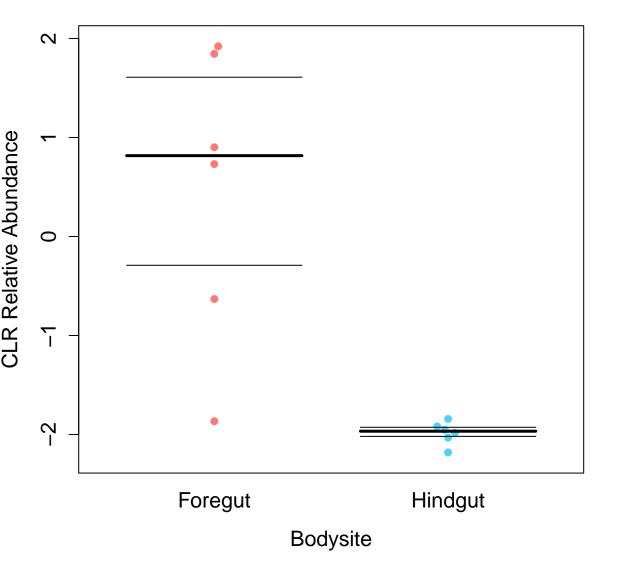
o__Aeromonadales; f__Aeromonadaceae



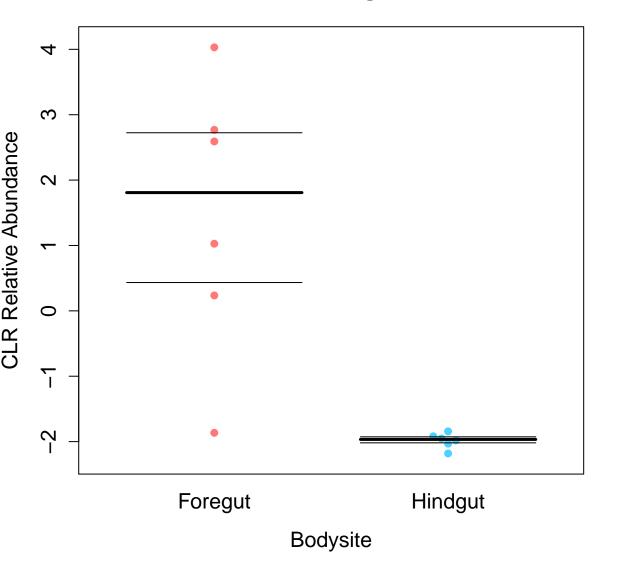




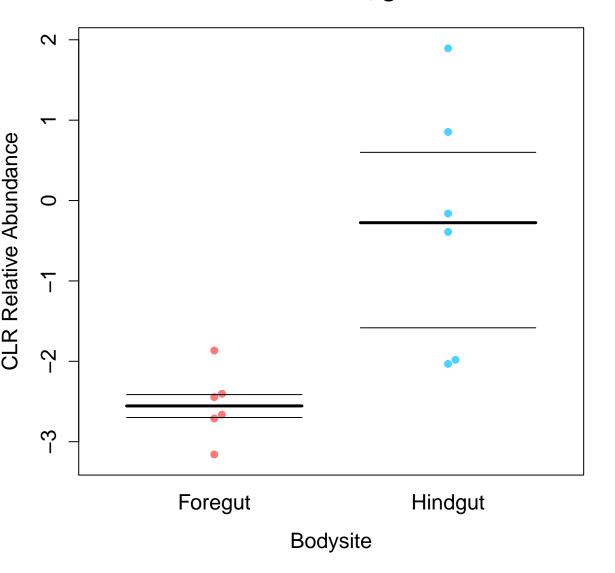
f__Planococcaceae; g__Rummeliibacillus



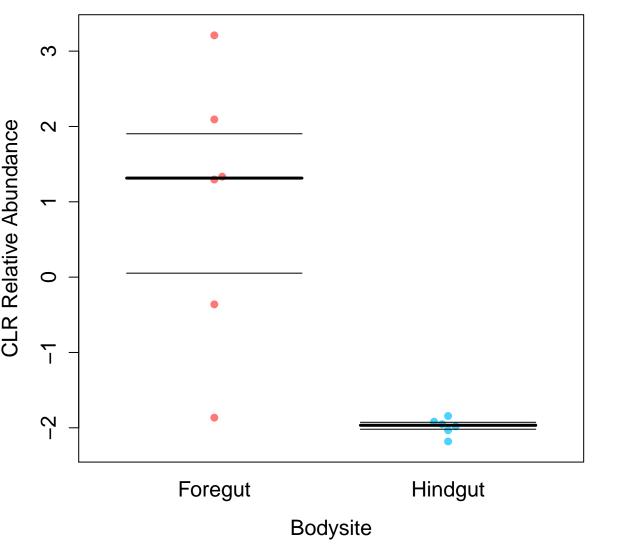
f_Planococcaceae; g_Solibacillus



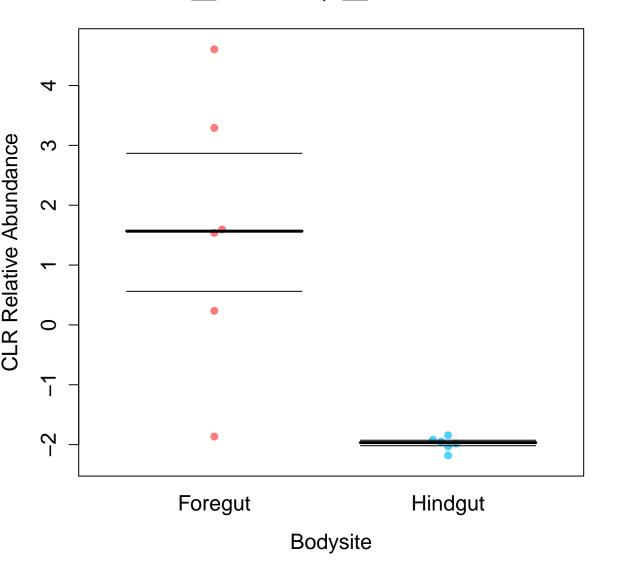
f__Verrucomicrobiaceae; g__Akkermansia



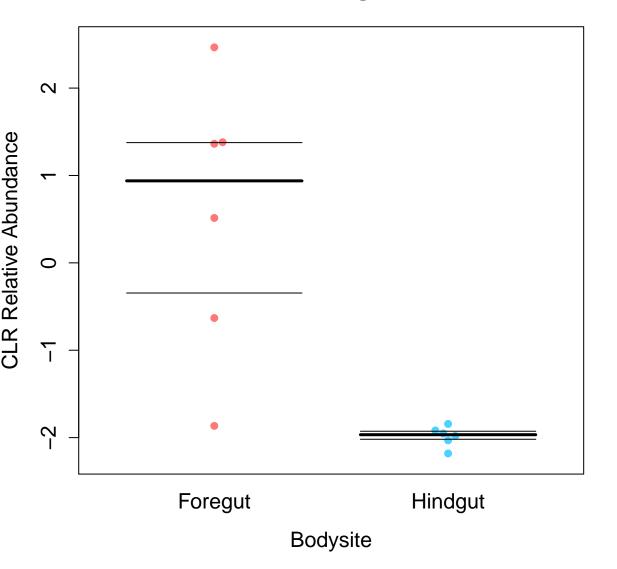
p__Proteobacteria; c__Gammaproteobacteria



o__Bacillales; f__Bacillaceae

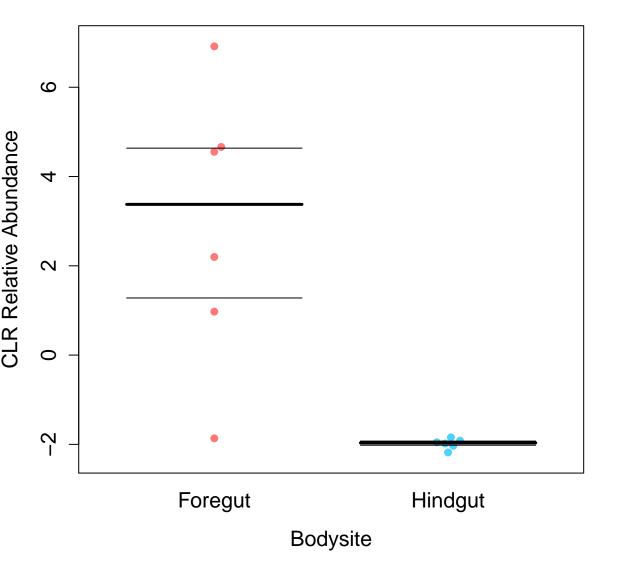




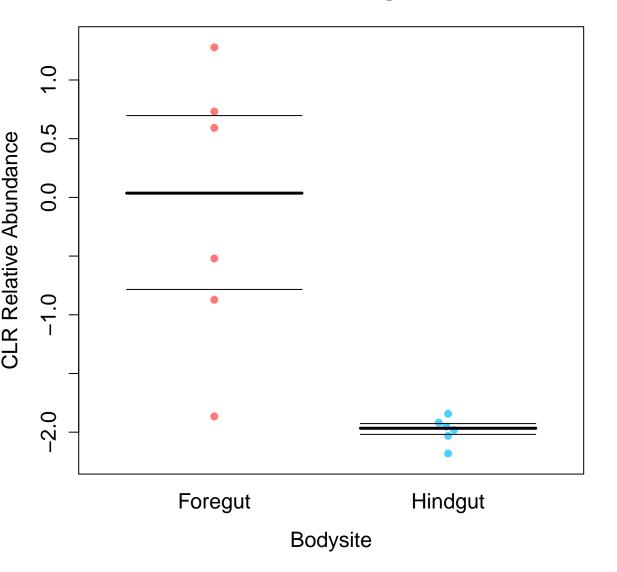


o__Clostridiales; f__Clostridiaceae 3.5 **CLR Relative Abundance** 2.5 Hindgut **Foregut Bodysite**

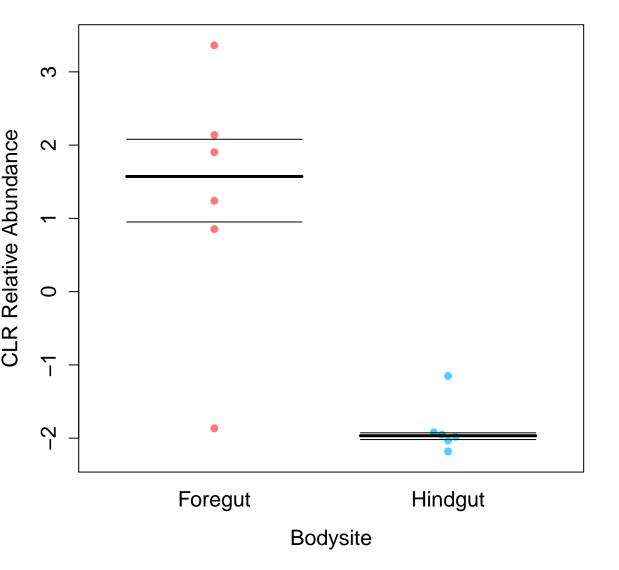




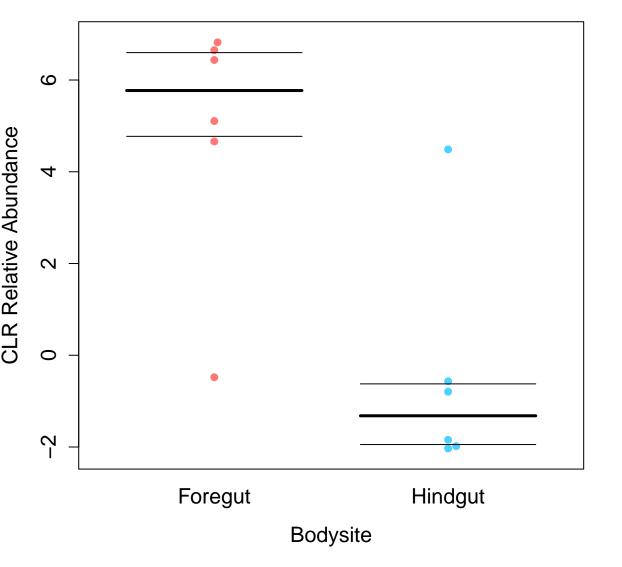




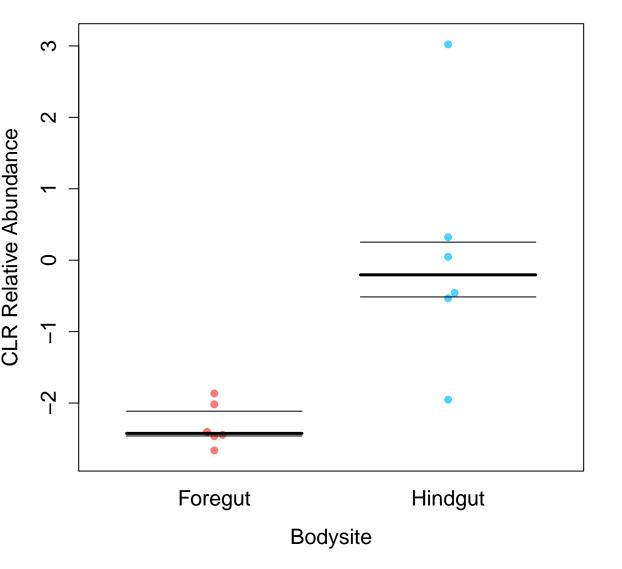




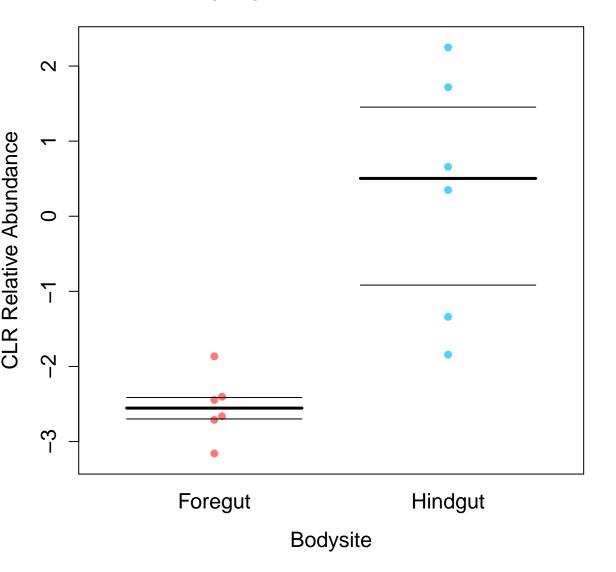
o__Enterobacteriales; f__Enterobacteriaceae



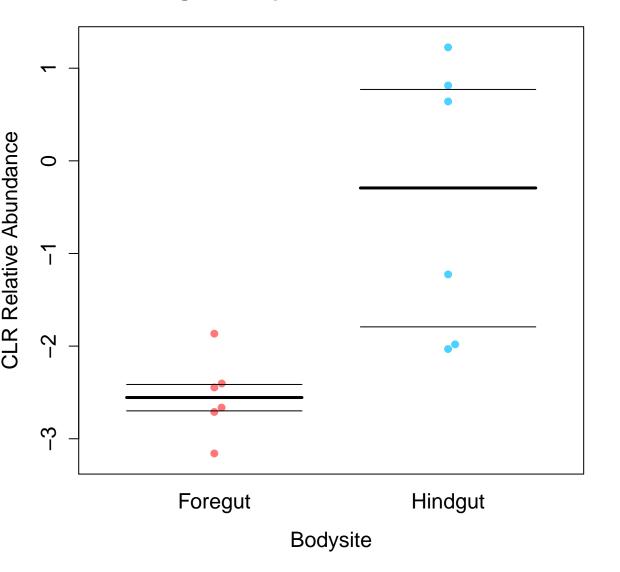
f__Lachnospiraceae; g__Anaerostipes



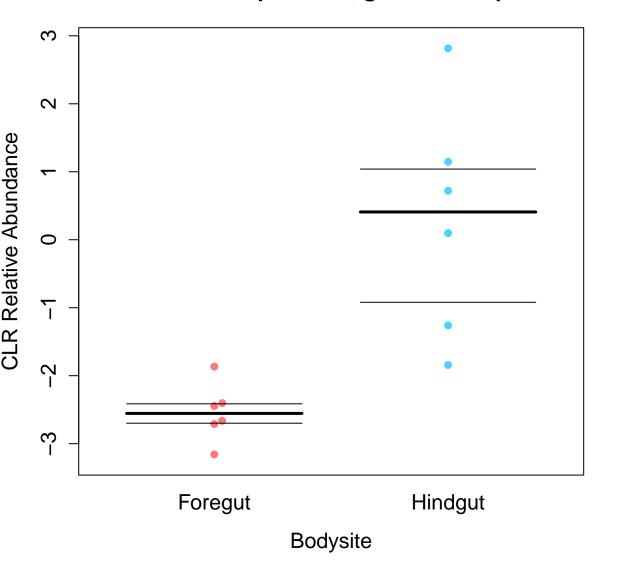
c__Alphaproteobacteria; o__RF32



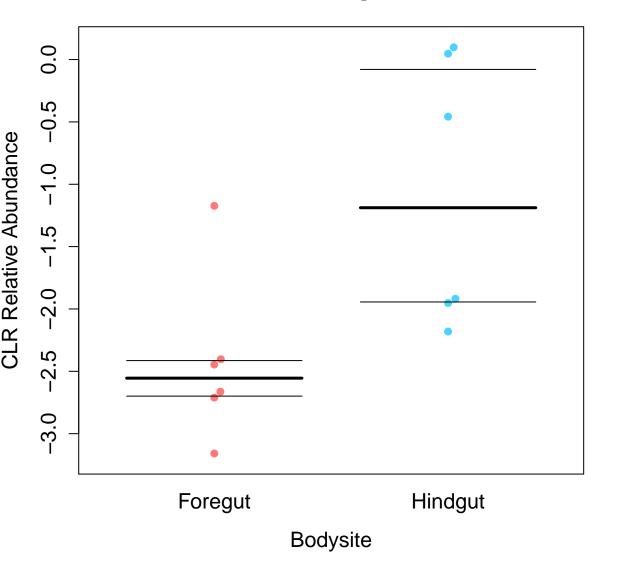
g__Alistipes; s__indistinctus



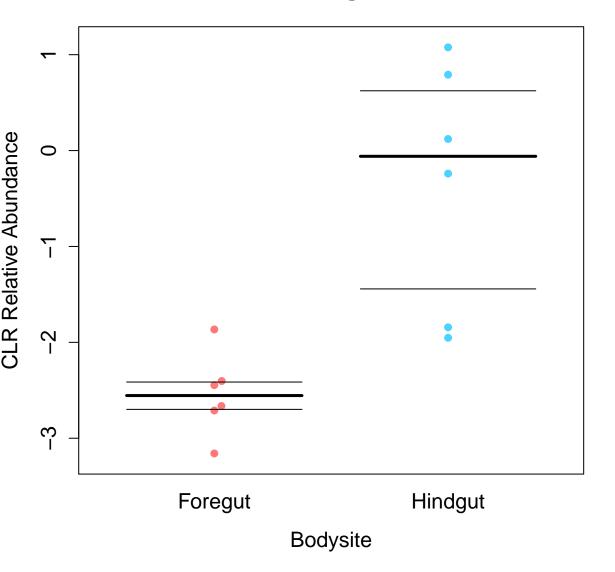
f__Lachnospiraceae; g__Lachnospira



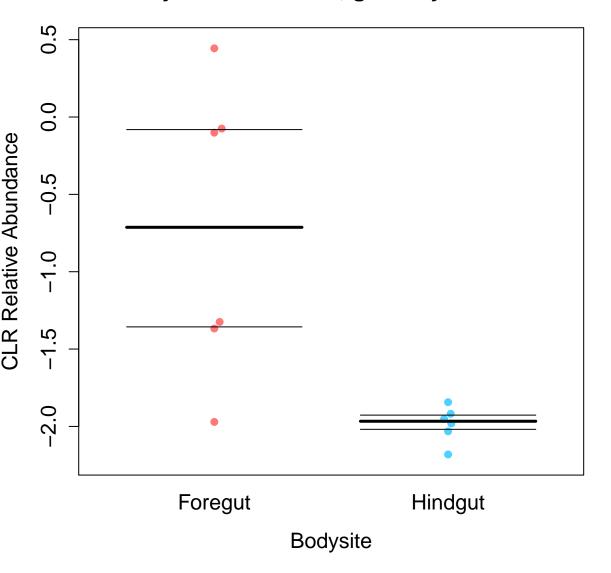
f__Ruminococcaceae; g__Anaerotruncus



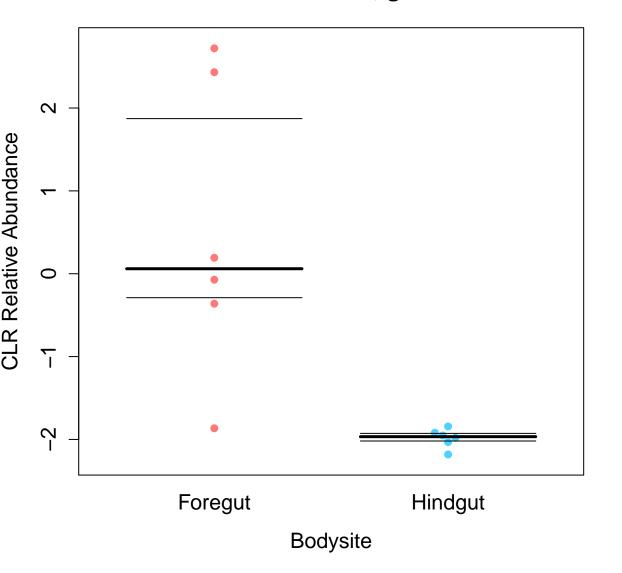
f__Ruminococcaceae; g__Faecalibacterium







f__Planococcaceae; g__Kurthia



f__Lachnospiraceae; g__Syntrophococcus

