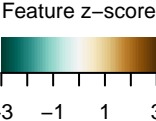
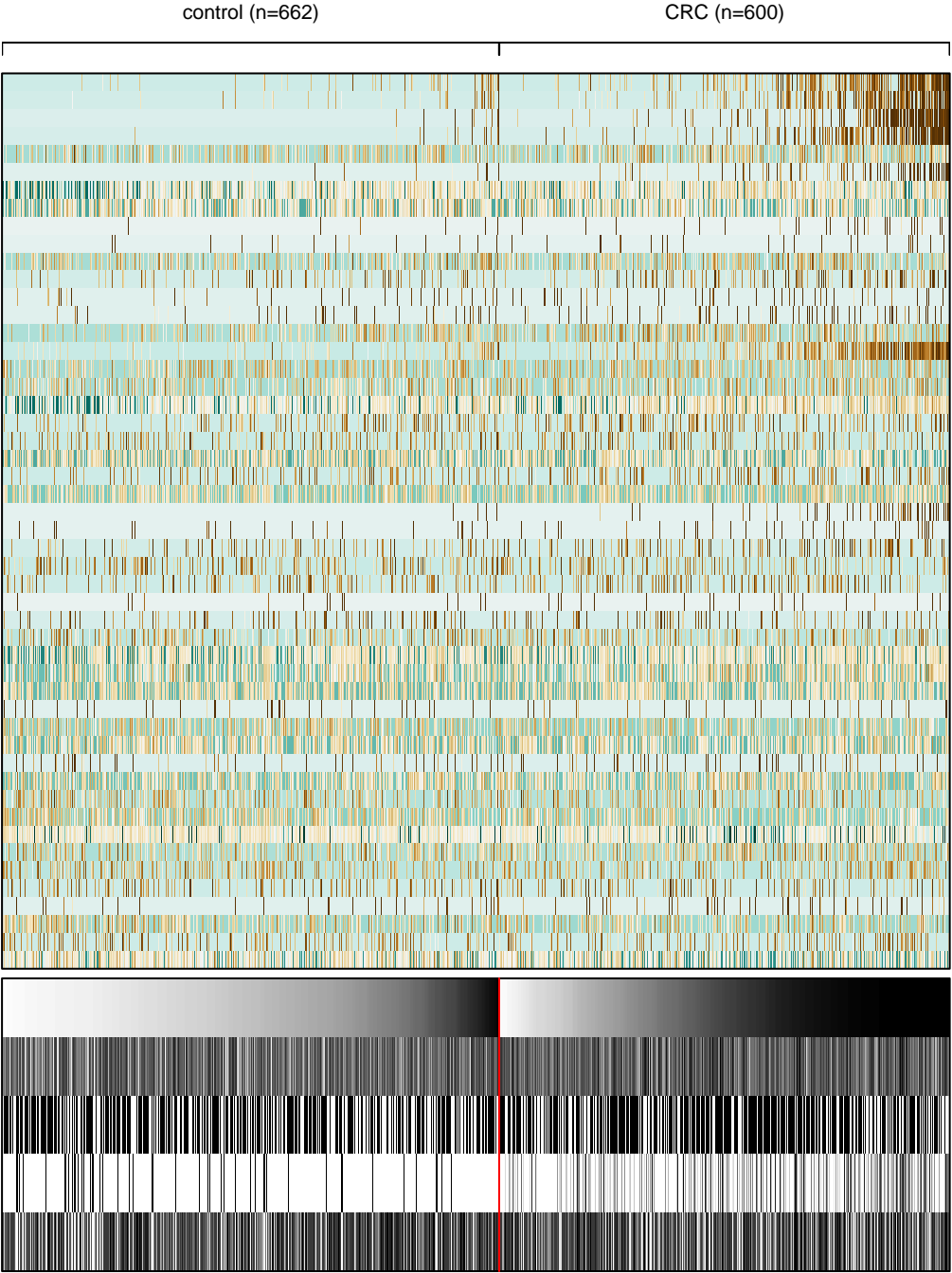
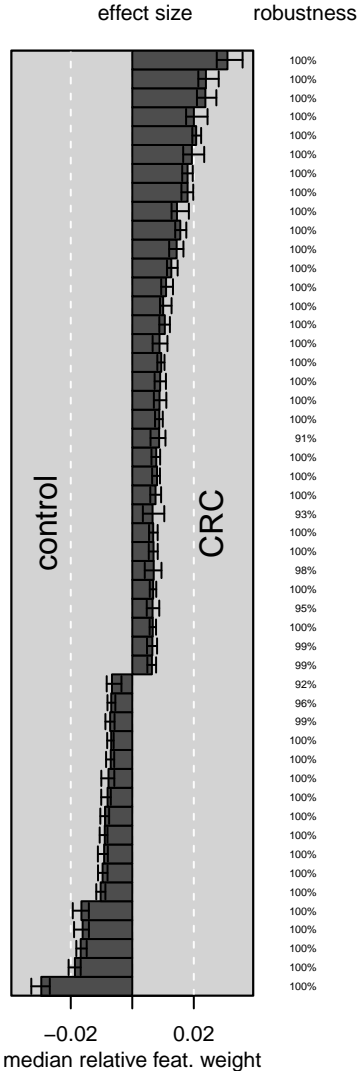


Feature Weights

Metagenomic Features



lasso model
(|W| = 50)



- Gemella_morbillorum
- Peptostreptococcus_stomatis
- Dialister_pneumosintes
- Fusobacterium_nucleatum
- Collinsella_massiliensis
- Porphyromonas_asaccharolytica
- Flavonifractor_plautii
- Streptococcus_parasanguinis
- Morganella_morganii
- Prevotella_bivia
- Streptococcus_mitis
- Bacteroides_nordii
- Enterobacter_cloacae_complex
- Prevotella_sp_CAG_520
- Anaerotruncus_colihominis
- Parvimonas_micra
- Eubacterium_siraeum
- Veillonella_parvula
- Ruthenibacterium_lactatiformans
- Eisenbergiella_massiliensis
- Klebsiella_pneumoniae
- Eubacterium_rectale
- Bacteroides_salysiae
- Ruminococcus_torques
- Porphyromonas_uenonis
- Klebsiella_oxytoca
- Ruminococcaceae_bacterium_D5
- Turicimonas_muris
- Allisonella_histaminiformans
- Citrobacter_braakii
- Ruminococcus_callidus
- Veillonella_atypica
- Collinsella_aerofaciens
- Collinsella_stercoris
- Coprococcus_comes
- Mitsuokella_jalaludinii
- Eubacterium_sp_CAG_38
- Roseburia_hominis
- Rothia_dentocariosa
- Eggerthella lenta
- Rothia_mucilaginosa
- Roseburia_intestinalis
- Faecalibacterium_prausnitzii
- Clostridium_innocuum
- Firmicutes_bacterium_CAG_110
- Klebsiella_varicola
- Granulicatella_elegans
- Haemophilus_painfluenzae
- Bacteroides_intestinalis
- Streptococcus_salivarius

