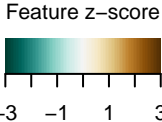
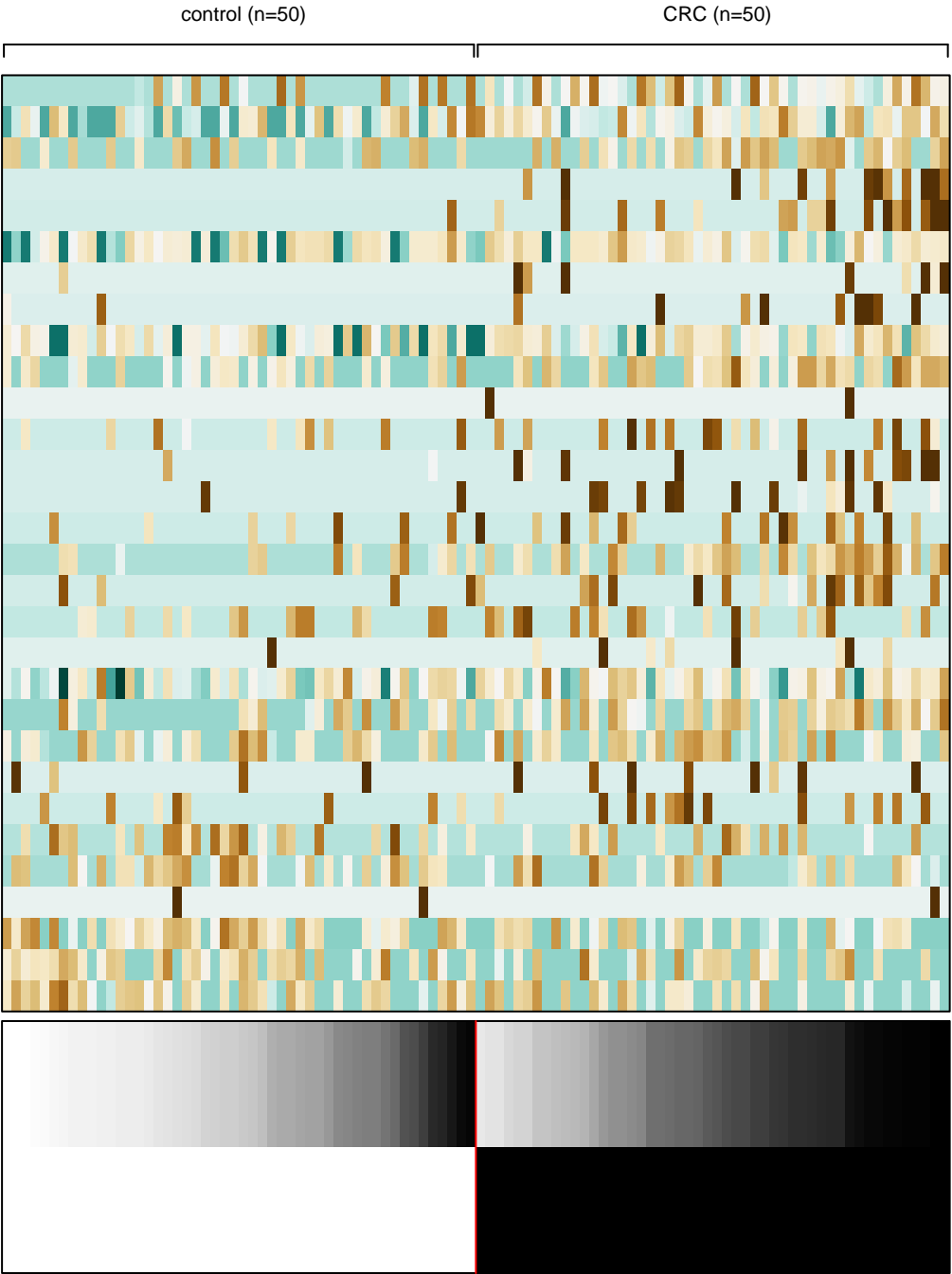
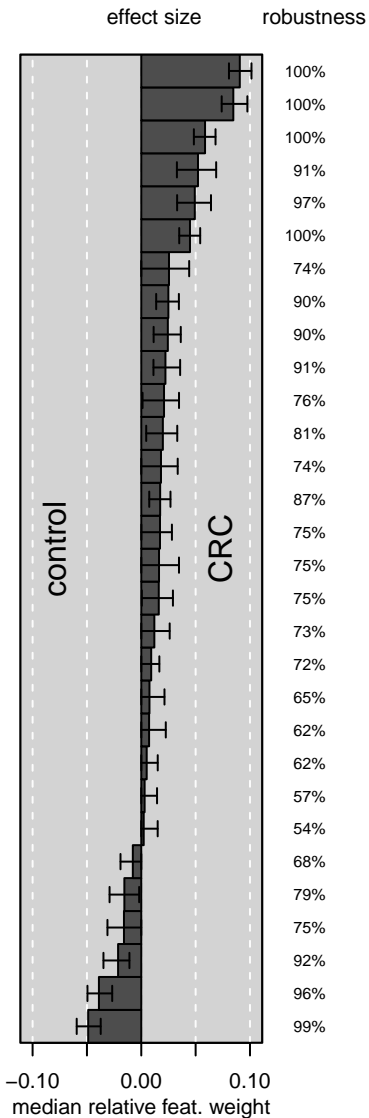


Feature Weights

Metagenomic Features



lasso model
(|W| = 30)



- Bacteroides_coprocola
- Bacteroides_fragilis
- Bacteroides_massiliensis
- Peptostreptococcus_stomatis
- Eisenbergiella_tayi
- Parabacteroides_merdae
- Solobacterium_moorei
- Fusobacterium_varium
- Bacteroides_caccae
- Alistipes_indistinctus
- Slackia_isoflavoniconvertens
- Firmicutes_bacterium_CAG_110
- Dialister_pneumosintes
- Phascolarctobacterium_succinatutens
- Clostridium_clostridioforme
- Clostridium_symbiosum
- Erysipelatoclostridium_ramosum
- Adlercreutzia_equolifaciens
- Prevotella_sp_CAG_5226
- Bacteroides_ovatus
- Hungatella_hathewayi
- Bifidobacterium_pseudocatenulatum
- Coprobacter_secundus
- Firmicutes_bacterium_CAG_95
- Gemella_haemolysans
- Eubacterium_hallii
- Phascolarctobacterium_sp_CAG_266
- Roseburia_intestinalis
- Haemophilus_parainfluenzae
- Eubacterium_sp_CAG_38

