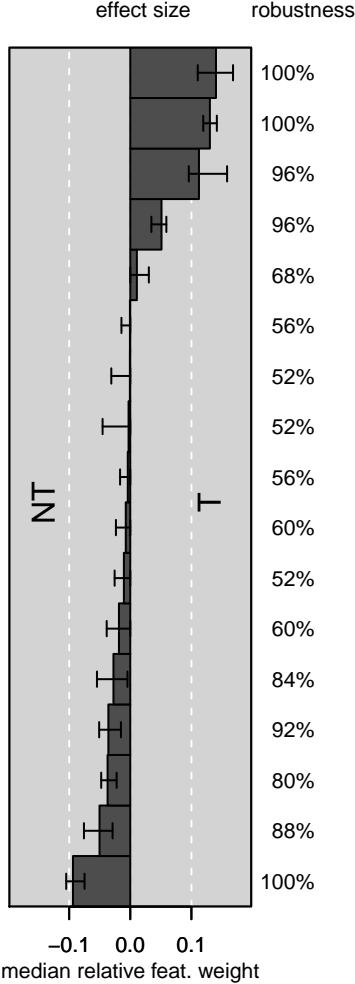
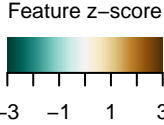
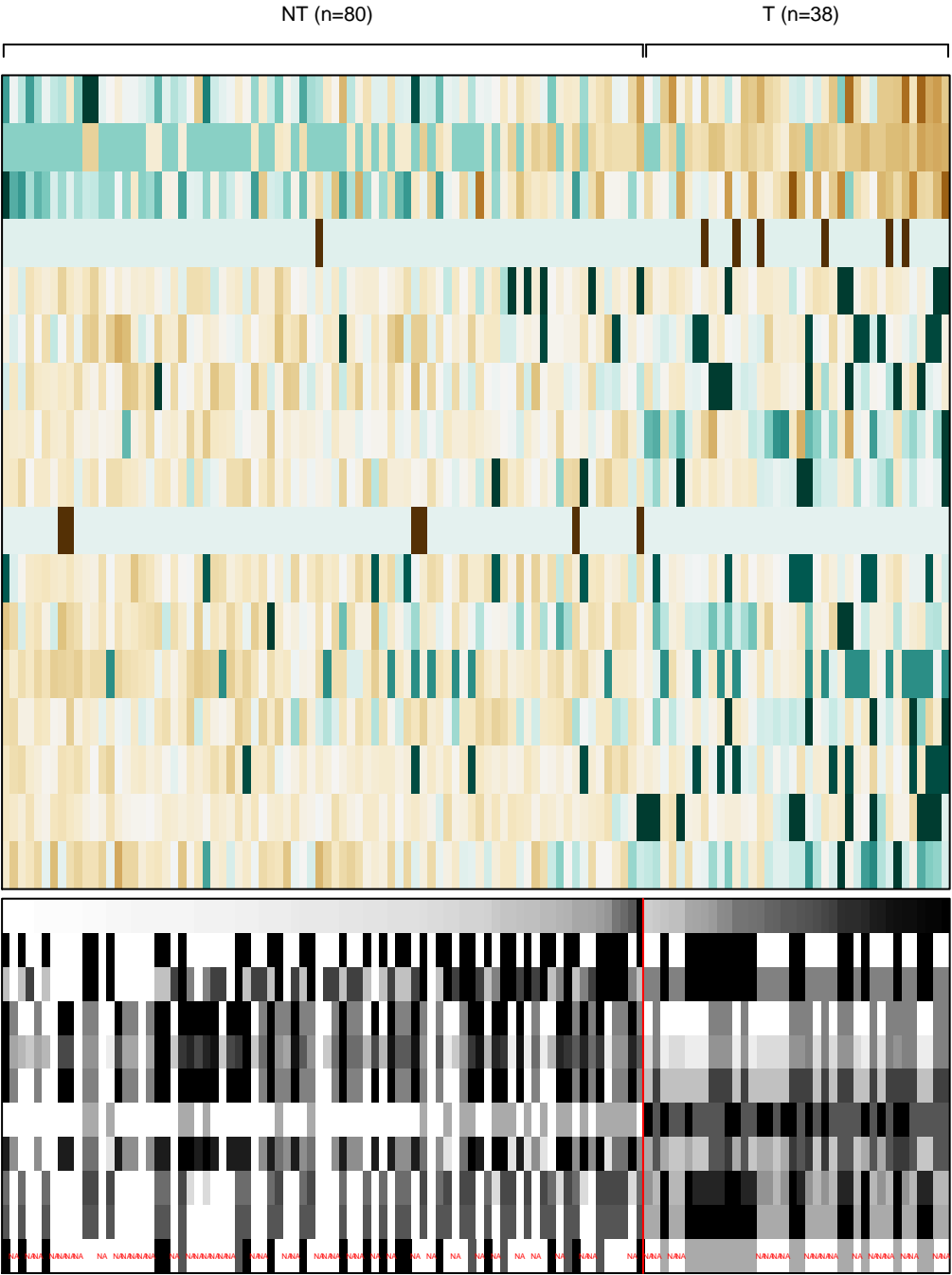


Feature Weights

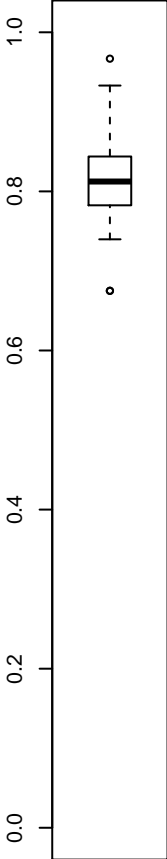


Metagenomic Features



lasso model  
(|W| = 17)

- OTU-OTU\_112:o\_\_Bacteroidales
- OTU-OTU\_170:Parabacteroides
- OTU-OTU\_18:Lactobacillus
- OTU-OTU\_483:Sphingomonas
- OTU-OTU\_123:f\_\_Lachnospiraceae
- OTU-OTU\_3869:f\_\_Lachnospiraceae
- OTU-OTU\_1768:f\_\_Lachnospiraceae
- OTU-OTU\_11:Anaerotruncus
- OTU-OTU\_129:Alistipes
- OTU-OTU\_665:Ruminococcus
- OTU-OTU\_106:f\_\_Porphyromonadaceae
- OTU-OTU\_73:Butyrivibrio
- OTU-OTU\_108:f\_\_Lachnospiraceae
- OTU-OTU\_409:f\_\_Lachnospiraceae
- OTU-OTU\_1002:f\_\_Ruminococcaceae
- OTU-OTU\_134:f\_\_Lachnospiraceae
- OTU-OTU\_79:o\_\_Clostridiales



- Classification result
- Mouse Line
- Genotype
- Age
- Age Genotype
- Age Phenotype
- Tissue
- Age Tissue
- Mouse Line Tissue
- Mouse Line Phenotype
- Responder