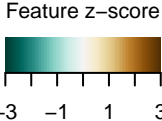
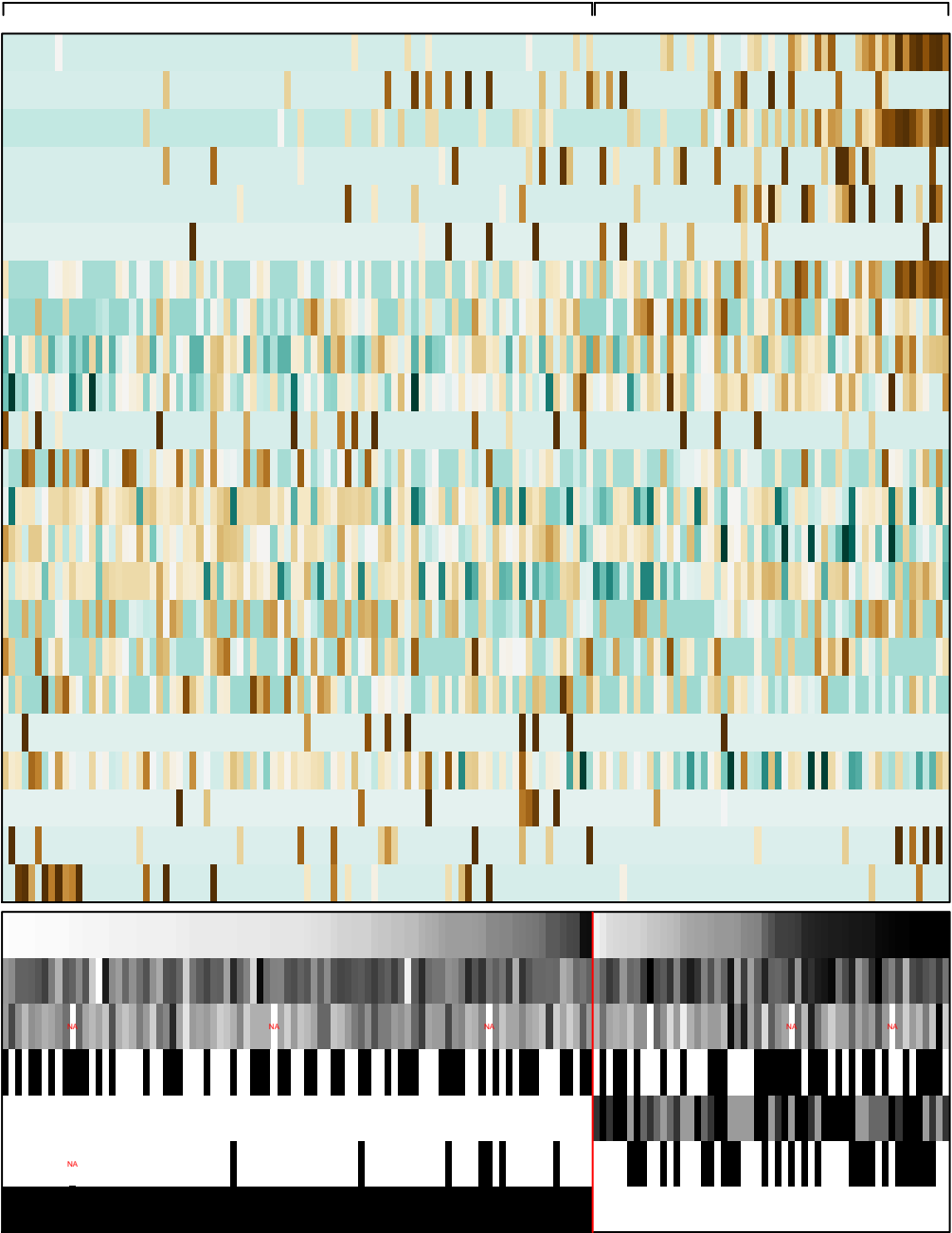
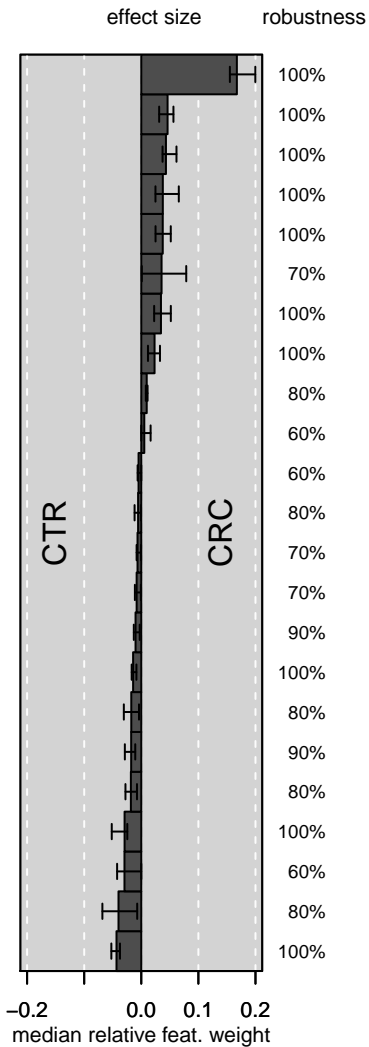


Feature Weights

Metagenomic Features



lasso model  
(|W| = 23)



- unclassified Fusobacterium [c:1482]
- Lactobacillus salivarius [h:1467]
- unclassified Fusobacterium [c:1481]
- Clostridium hylemonae [h:1607]
- Porphyromonas asaccharolytica [h:1056]
- Streptococcus agalactiae [h:1388]
- Peptostreptococcus stomatis [h:1530]
- Clostridium symbiosum [h:1600]
- Bacteroides fragilis [h:1090]
- Pseudoflavonifractor capillosus [h:1579]
- unnamed Erysipelotrichaceae bacterium 3\_1\_53 [u:1277]
- Butyrivibrio crossotus [h:1628]
- Eubacterium eligens [h:1627]
- unnamed Ruminococcus sp. 5\_1\_39BFAA [u:1620]
- Bacteroides caccae [h:1096]
- Phascolarctobacterium succinatutens [h:1659]
- Clostridium scindens [h:1606]
- Bifidobacterium angulatum [h:974]
- Enterococcus gallinarum [h:1362]
- Streptococcus salivarius [h:1377]
- Leuconostoc citreum [h:1431]
- Peptoniphilus harei [h:1493]
- Methanospaera stadmanae [h:94]

