Metagenomic Features

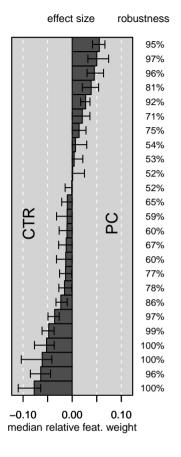
PC (n=57)

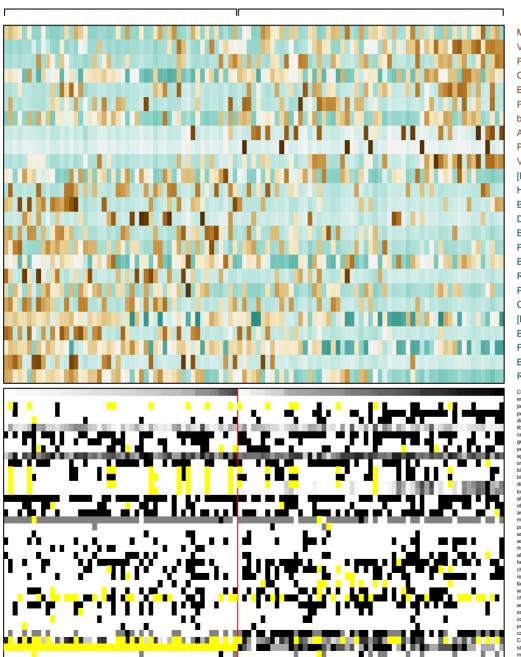
CTR (n=50)



-3 -1 1 3

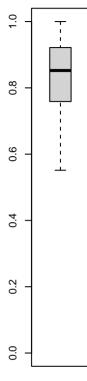
lasso_ll model (|W| = 25)





Methanobrevibacter smithii [r_03695] Veillonella atypica [r_01941] Firmicutes sp. [r_03641] Clostridium sp. [r_03622] Bacteroides finegoldii [r_03474] Firmicutes sp. [r_03629] bacterium LF-3 [r_03628] Alloscardovia omnicolens [r 02114] Prevotella species [m_12780] Veillonella species [m_13135] [Ruminococcus] gnavus [r_01594] Holdemanella biformis [m_12329] Bifidobacterium ruminantium [r_02702] Dorea sp. CAG:317 [r_07668] Bacteroides sp. CAG:144 [m_12596] Faecalibacterium species [m_12403] Bacteroides caecimuris [r_03476] Rikenellaceae sp. [r_03593] Paraprevotella clara [r_03698] Clostridium sp. CAG:217 [m_12270] [Eubacterium] rectale [r_03657] Bacteroides coprocola [r_11279] Faecalibacterium prausnitzii [r_06110] Bifidobacterium bifidum [r_03116] Romboutsia timonensis [r_09389]





proportion of weight shown