ROC curve

AUC–ROC
0.881
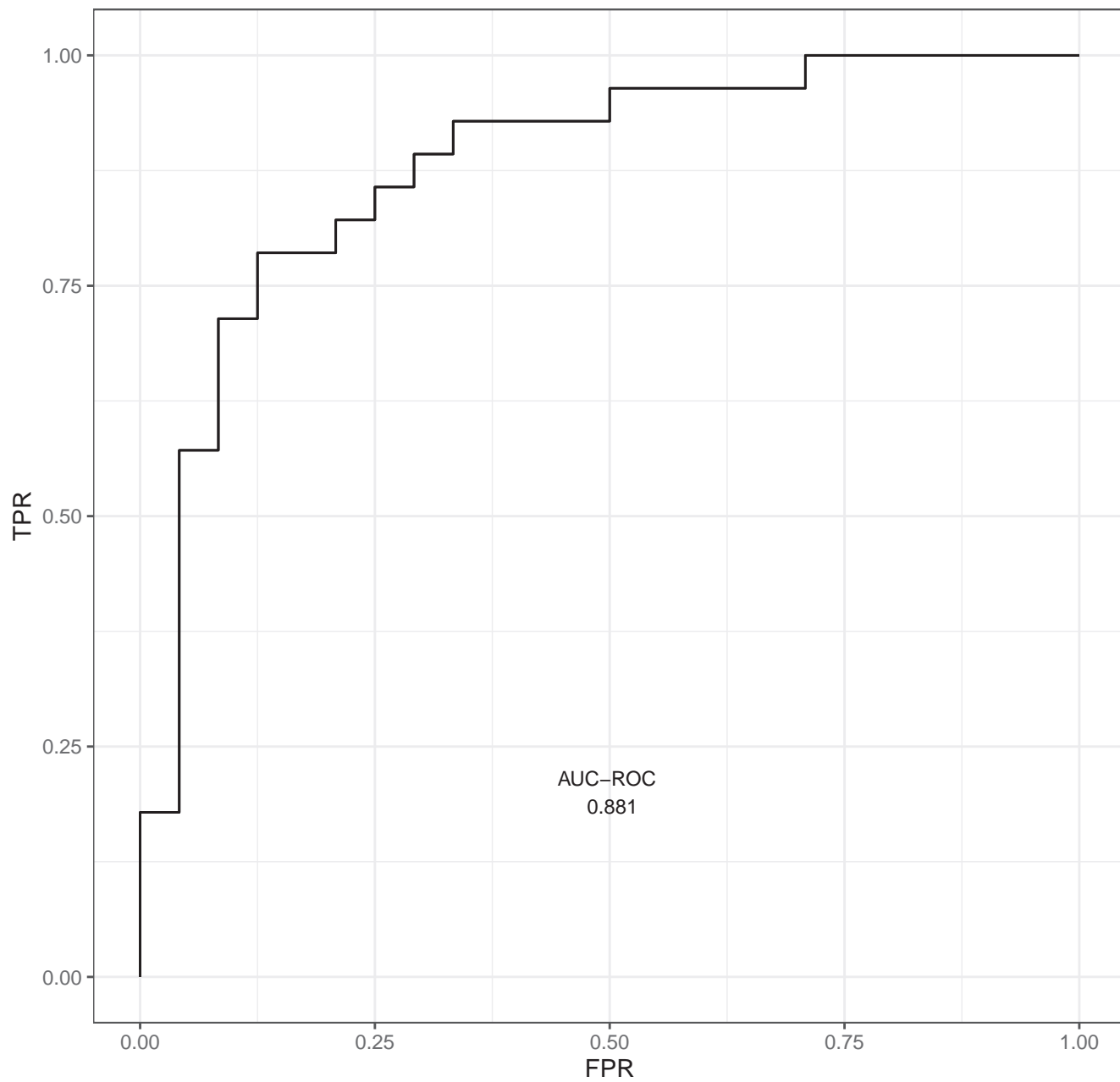
Cross validation in balance selection

Parasutterella
Campylobacter
Enterobacteriaceae
Hungatella
Enterococcus
Bilophila
Staphylococcus
Lactobacillus
Anaerococcus
Parabacteroides
DTU089
Fenollaria
Peptoniphilus
Pseudomonas
Acinetobacter
Ruminococcus
Dialister
Veillonella
Sutterella
Streptococcus
Bacteroides
Butyricicoccus
Lachnoclostridium
Colidextribacter
X.Clostridium..innocuum.group
Bifidobacterium
Akkermansia
Eisenbergiella
Phascolarctobacterium
Finegoldia
UBA1819

0   20   40   60
% of times included in a balance

Group of . . .
DEN
NUM