

Corynebacterium 1

FDR: 1.145e-08
Coefficient: 8.34e-02

0.15

0.10

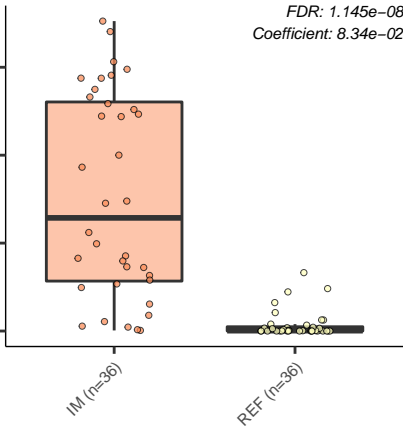
0.05

0.00

IM (n=36)

REF (n=36)

Diet



f_Bacillaceae

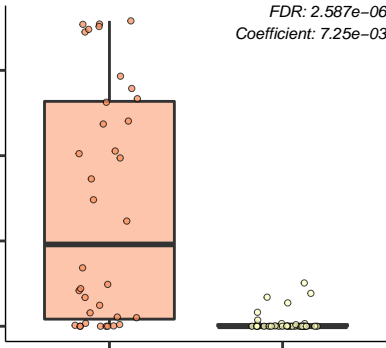
FDR: 2.587e-06
Coefficient: 7.25e-03

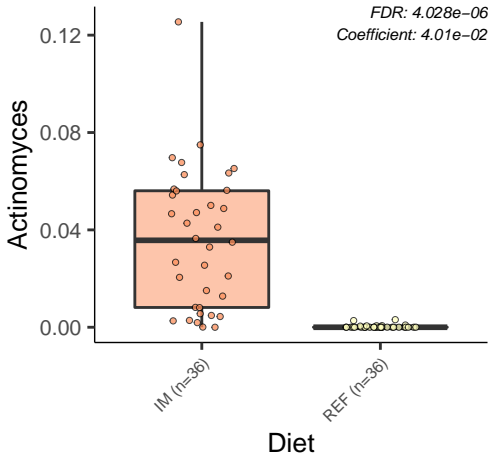
0.015
0.010
0.005
0.000

IM (n=36)

REF (n=36)

Diet





f_Butenbergiaceae

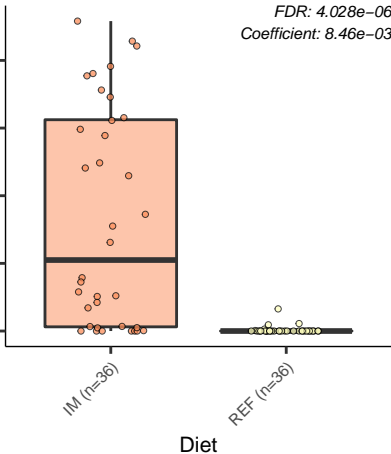
FDR: 4.028e-06
Coefficient: 8.46e-03

0.020
0.015
0.010
0.005
0.000

IM (n=36)

REF (n=36)

Diet



Gracilibacillus

FDR: 6.194e-06
Coefficient: 1.07e-03

0.002

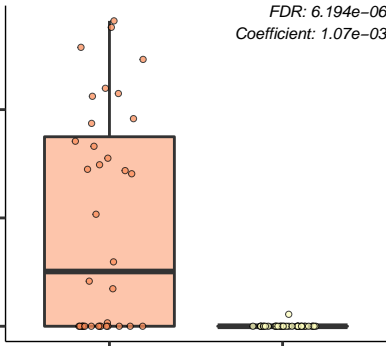
0.001

0.000

IM (n=36)

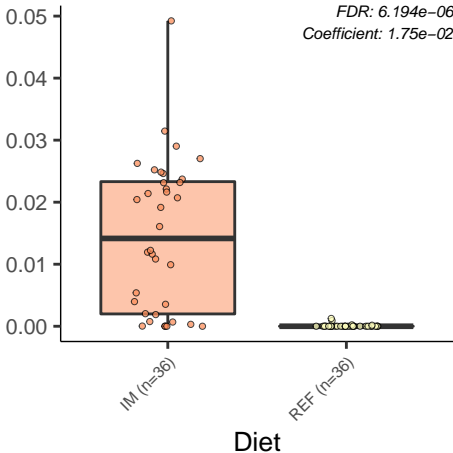
REF (n=36)

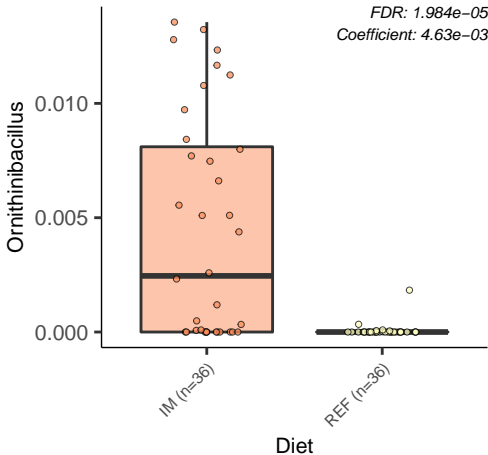
Diet

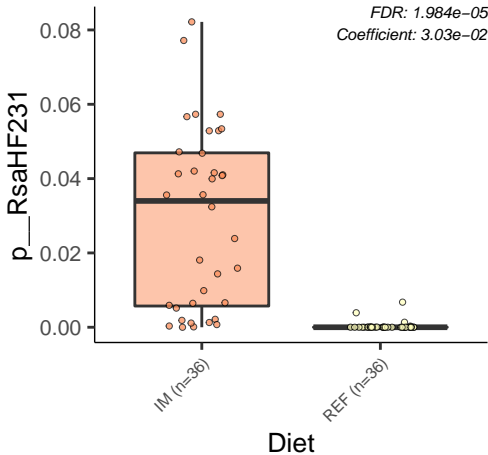


Microbacterium

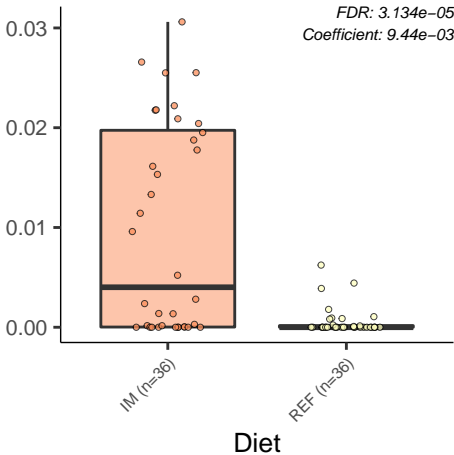
FDR: 6.194e-06
Coefficient: 1.75e-02





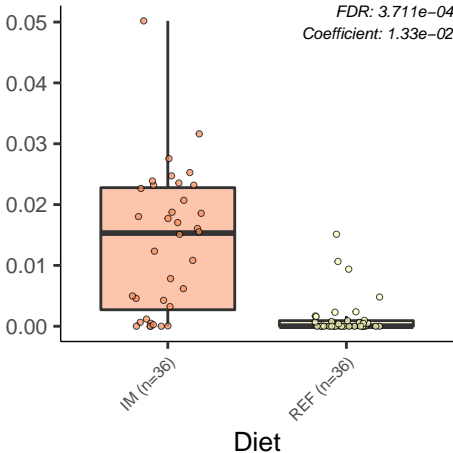


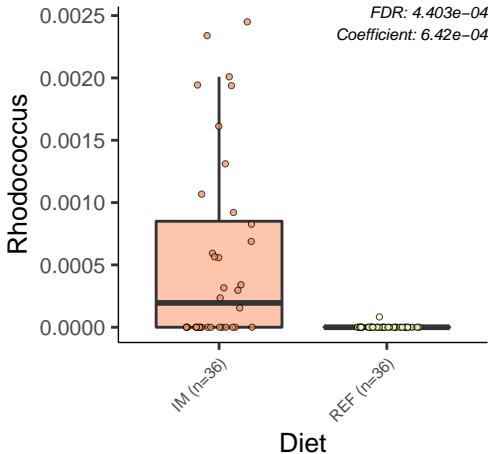
Lysinibacillus



Brevibacterium

FDR: 3.711e-04
Coefficient: 1.33e-02





Oceanobacillus

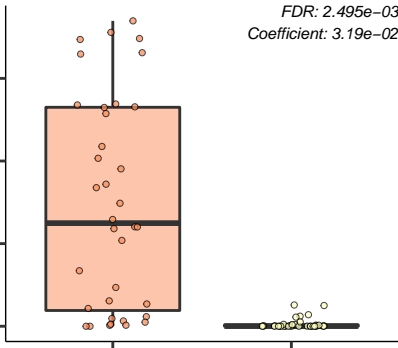
FDR: 2.495e-03
Coefficient: 3.19e-02

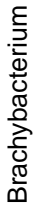
0.06
0.04
0.02
0.00

IM (n=36)

REF (n=36)

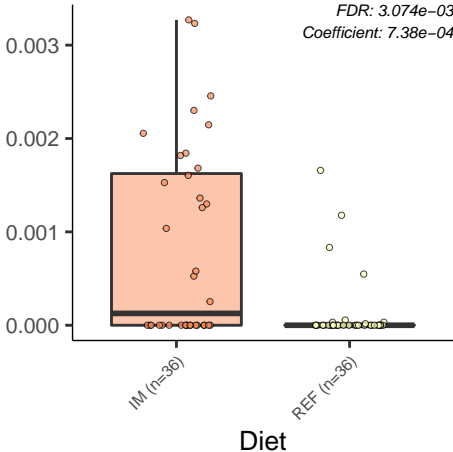
Diet





FDR: 3.074e-03

Coefficient: 7.38e-04



IM (n=36)

REF (n=36)

f_Lachnospiraceae

FDR: $8.569e-03$
Coefficient: $7.17e-03$

0.020

0.015

0.010

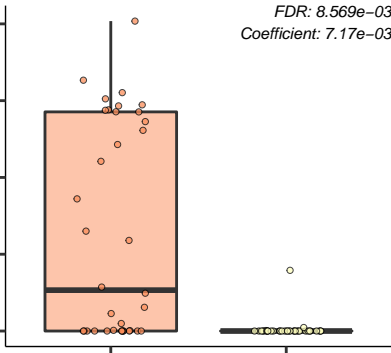
0.005

0.000

IM (n=36)

REF (n=36)

Diet



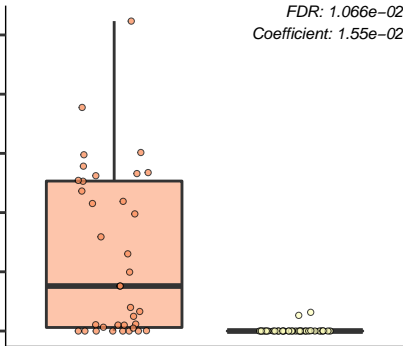
f__Enterococcaceae

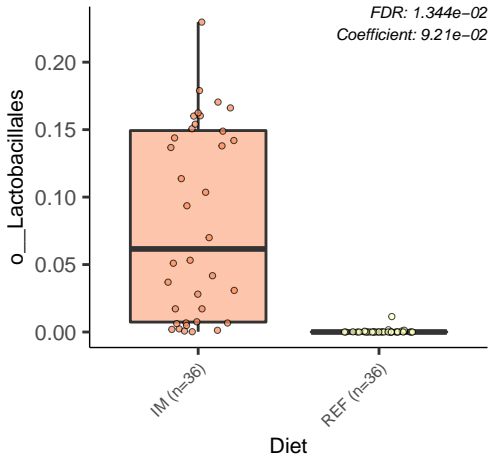
FDR: 1.066e-02
Coefficient: 1.55e-02

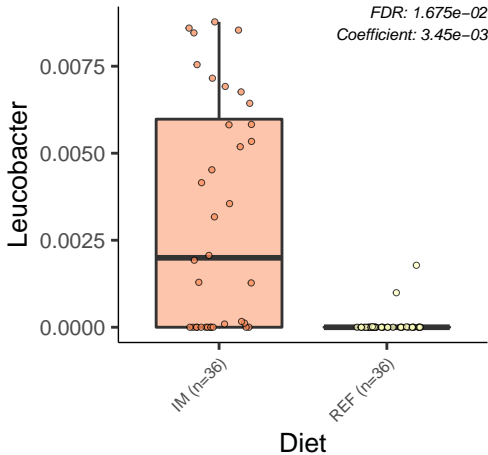
IM (n=36)

REF (n=36)

Diet







Tepidimicrobium

0.03

0.02

0.01

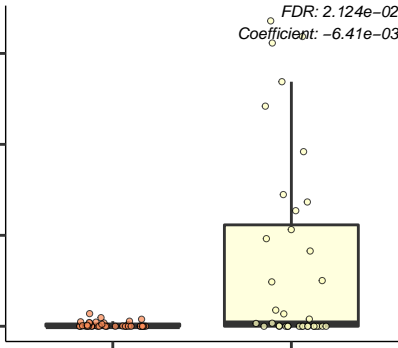
0.00

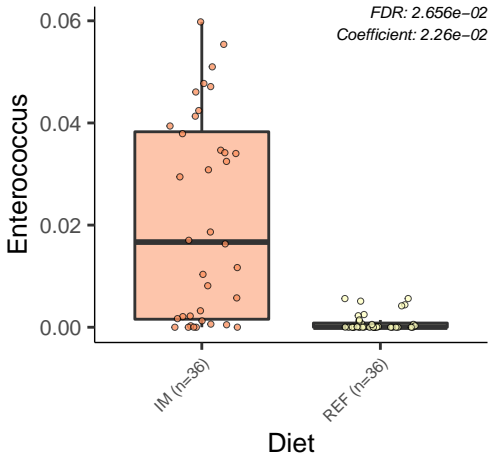
IM (n=36)

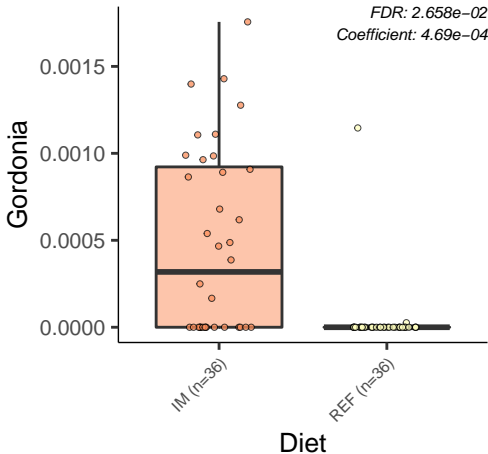
REF (n=36)

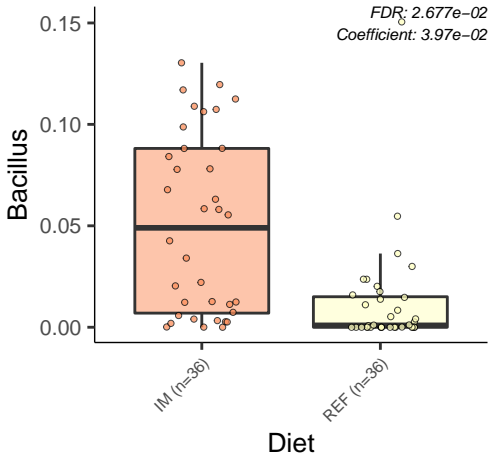
Diet

FDR: 2.124×10^{-2}
Coefficient: -6.41×10^{-3}

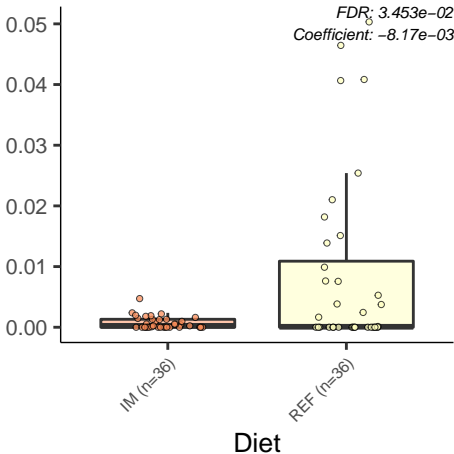


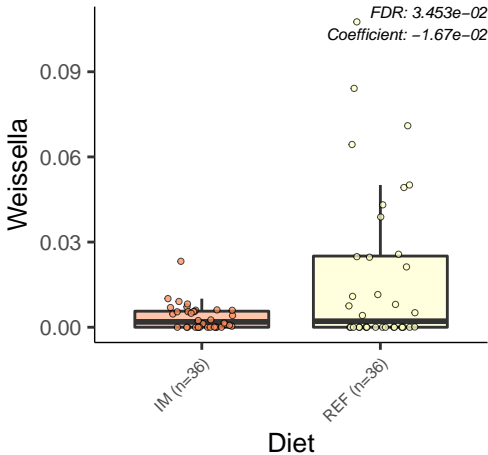


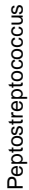




Geobacillus







0.02

0.01

0.00

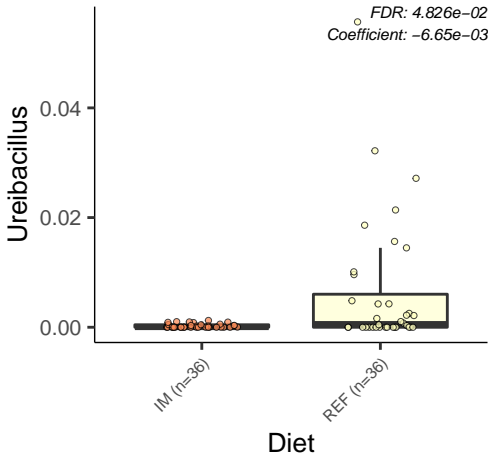
IM (n=36)

REF (n=36)

Diet

FDR: 3.472e-02
Coefficient: -4.15e-03

Coefficient: $-4.15e-03$



Photobacterium

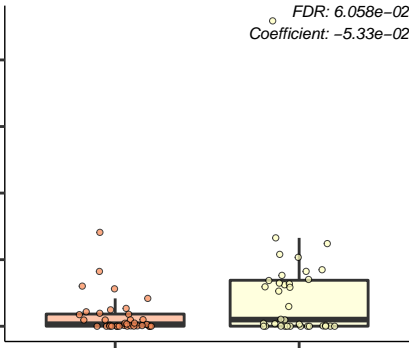
FDR: 6.058e-02
Coefficient: -5.33e-02

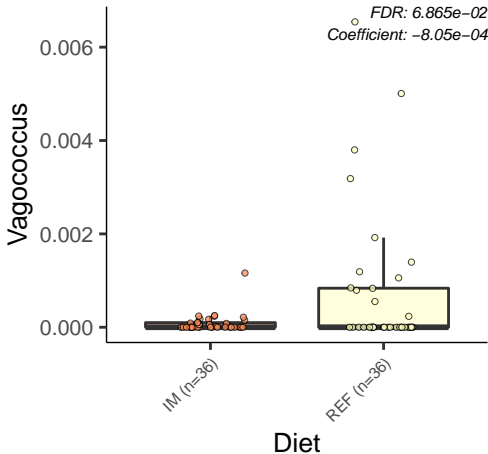
0.4
0.3
0.2
0.1
0.0

IM (n=36)

REF (n=36)

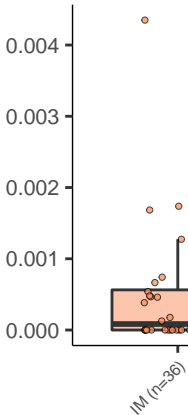
Diet





Erysipelatoclostridium

FDR: 8.516e-02
Coefficient: 5.84e-04



REF (n=36)

Diet

Brevinema andersonii

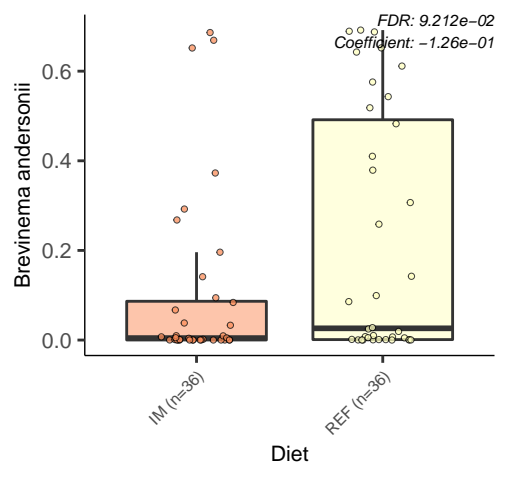
FDR: $9.212e-02$
Coefficient: $-1.26e-01$

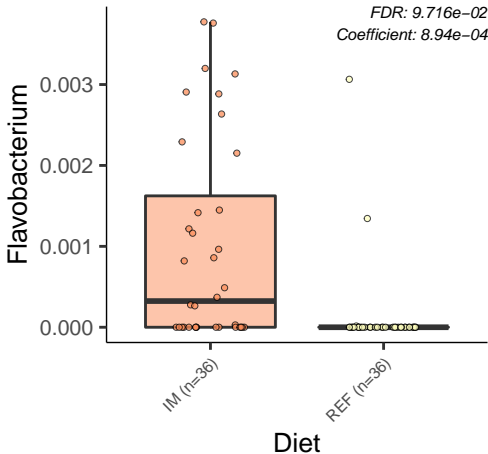
0.6
0.4
0.2
0.0

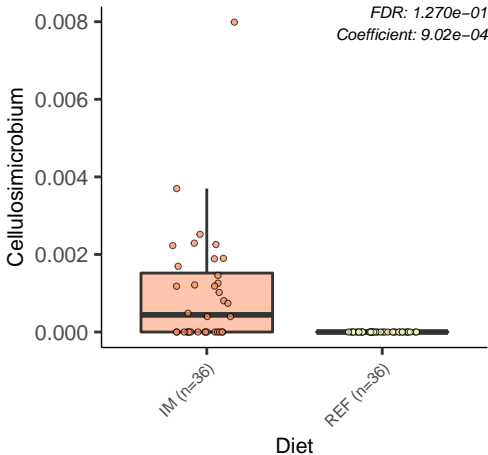
IM (n=36)

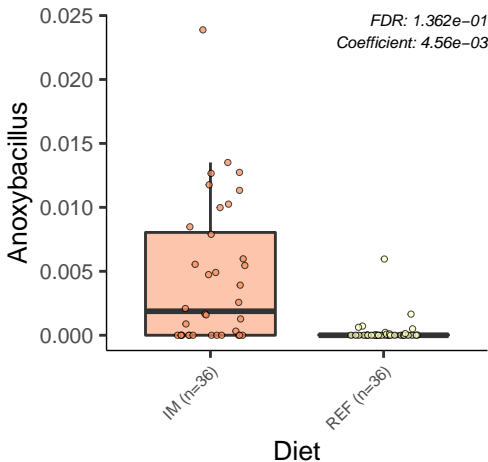
REF (n=36)

Diet

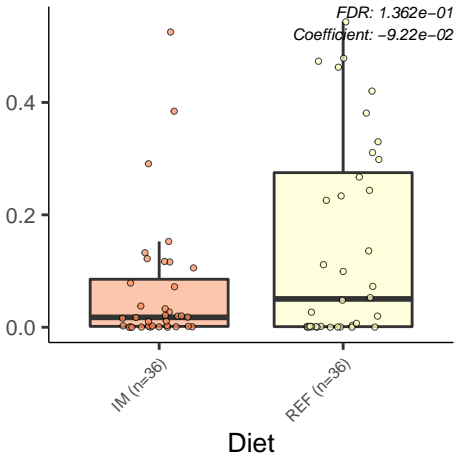


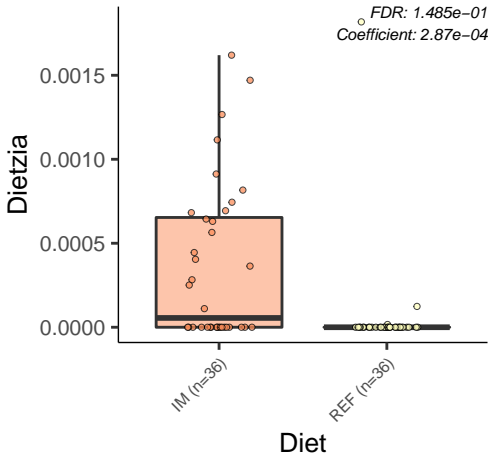






Mycoplasma





Streptococcus

FDR: $2.044e-01$
Coefficient: $-3.26e-03$

0.03

0.02

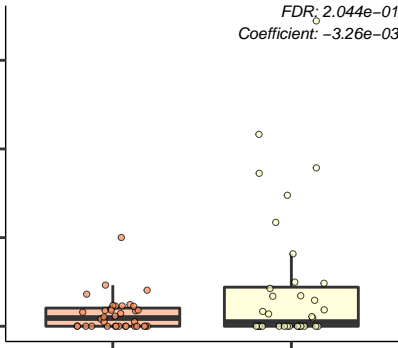
0.01

0.00

IM (n=36)

REF (n=36)

Diet



FDR: 2.473e-01
Coefficient: 1.71e-03

