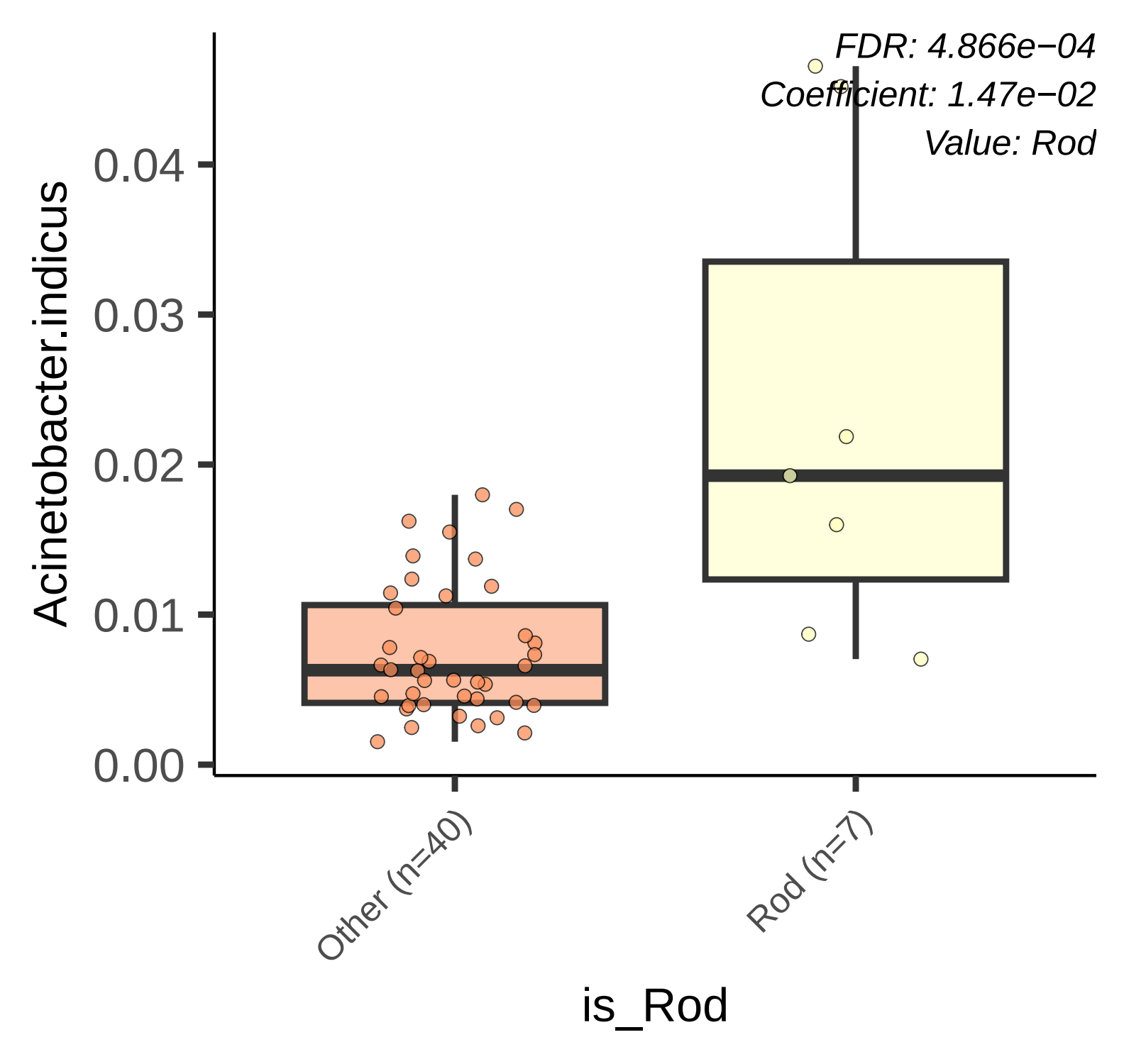
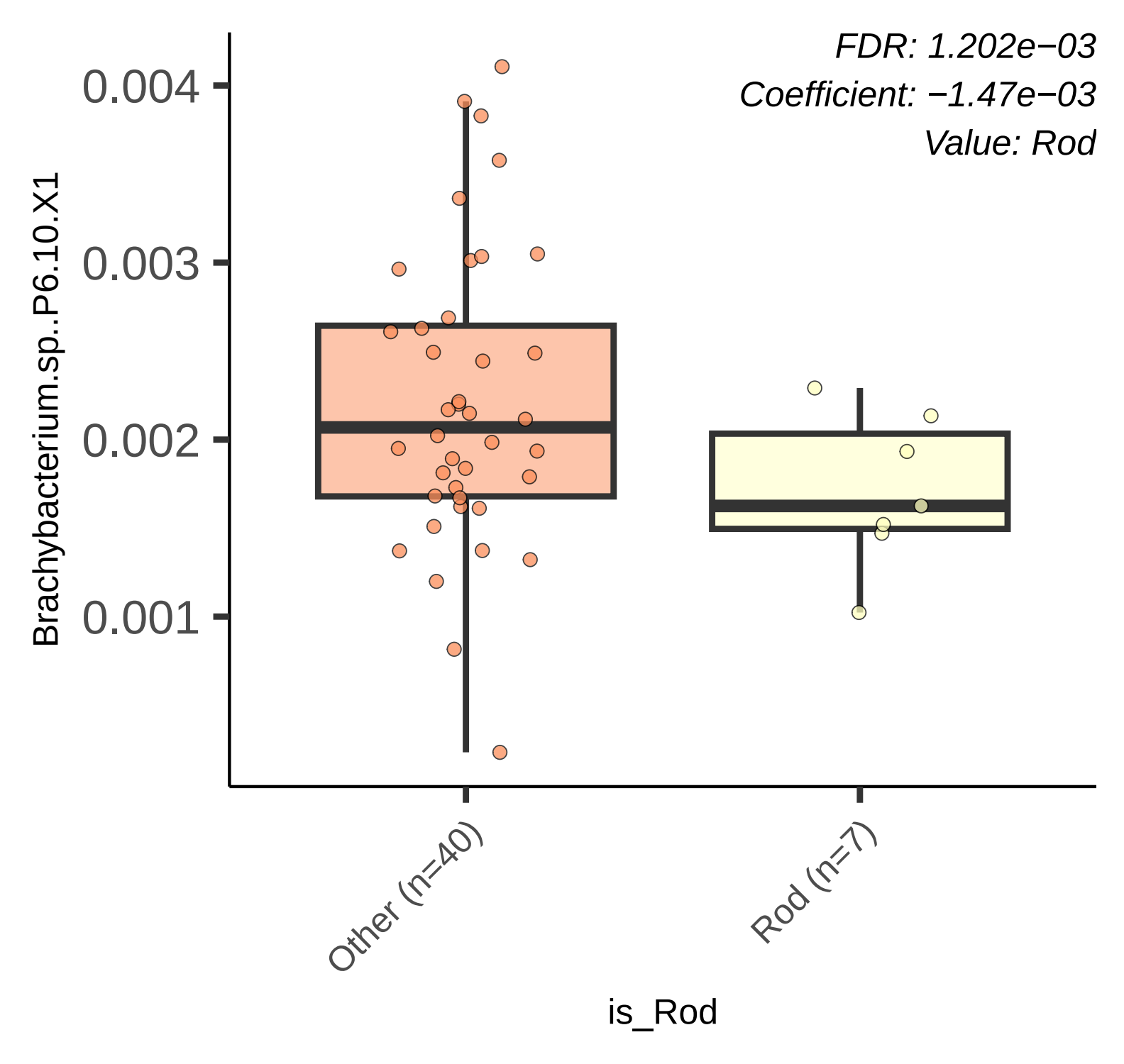
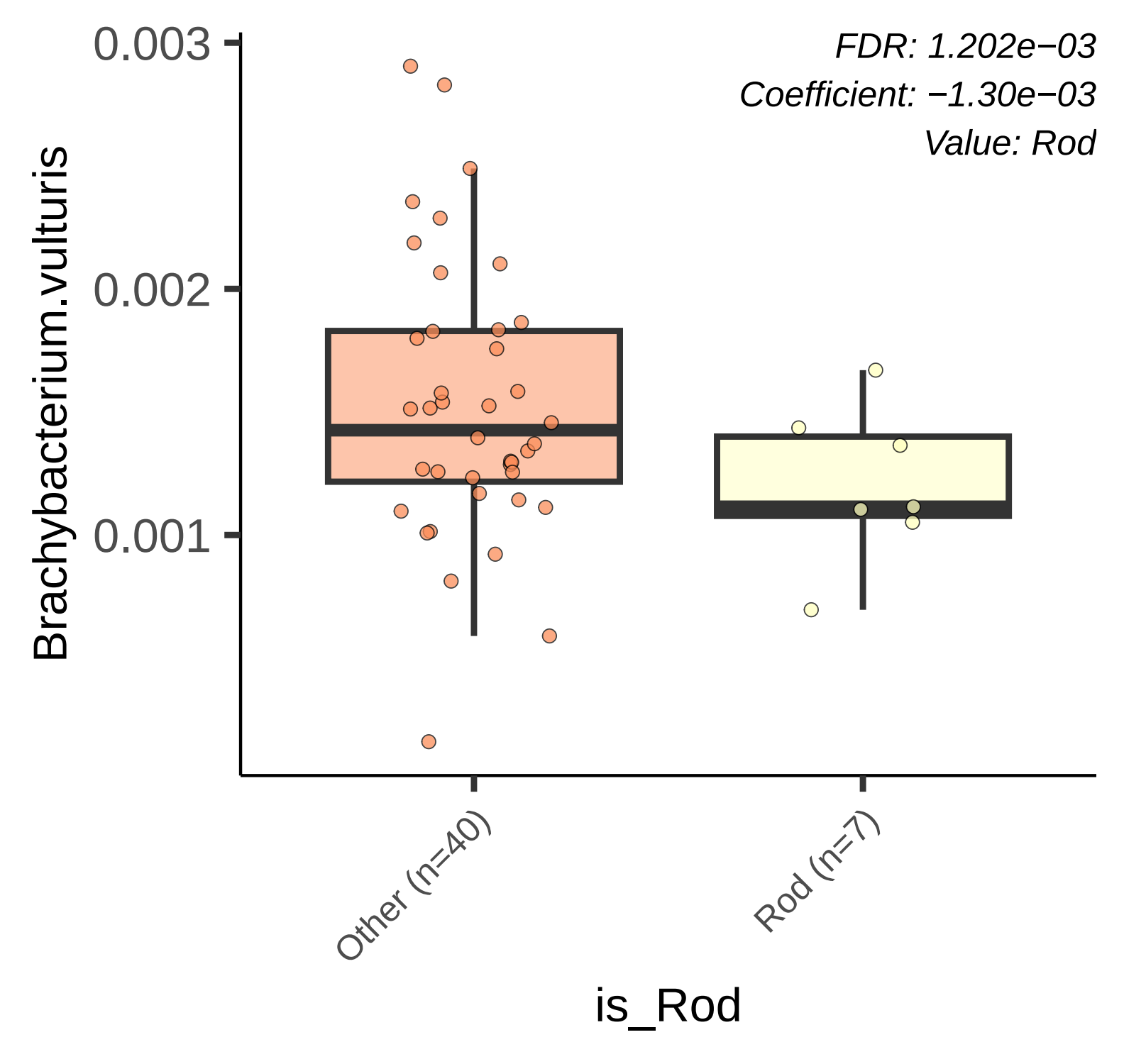
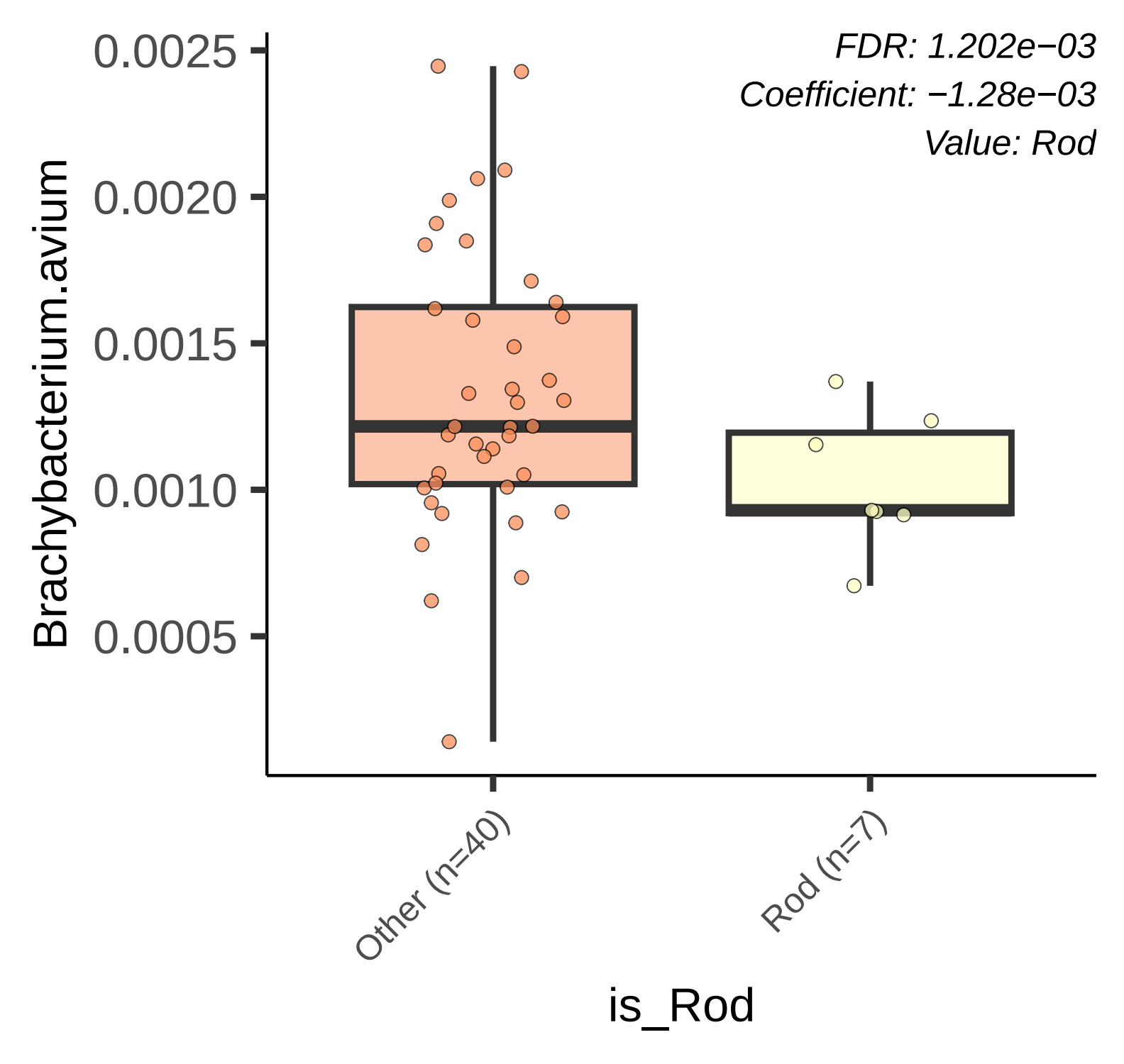
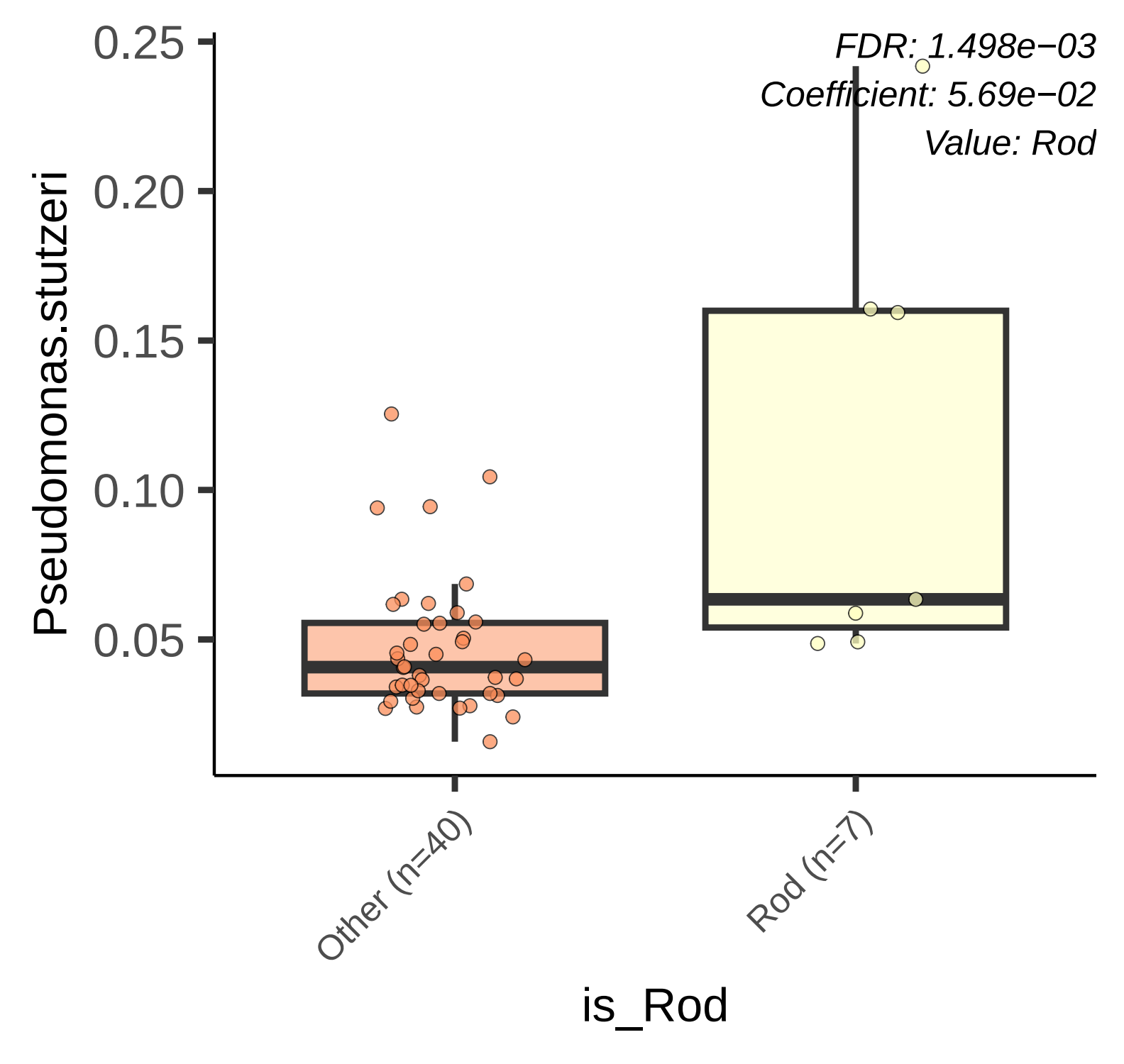
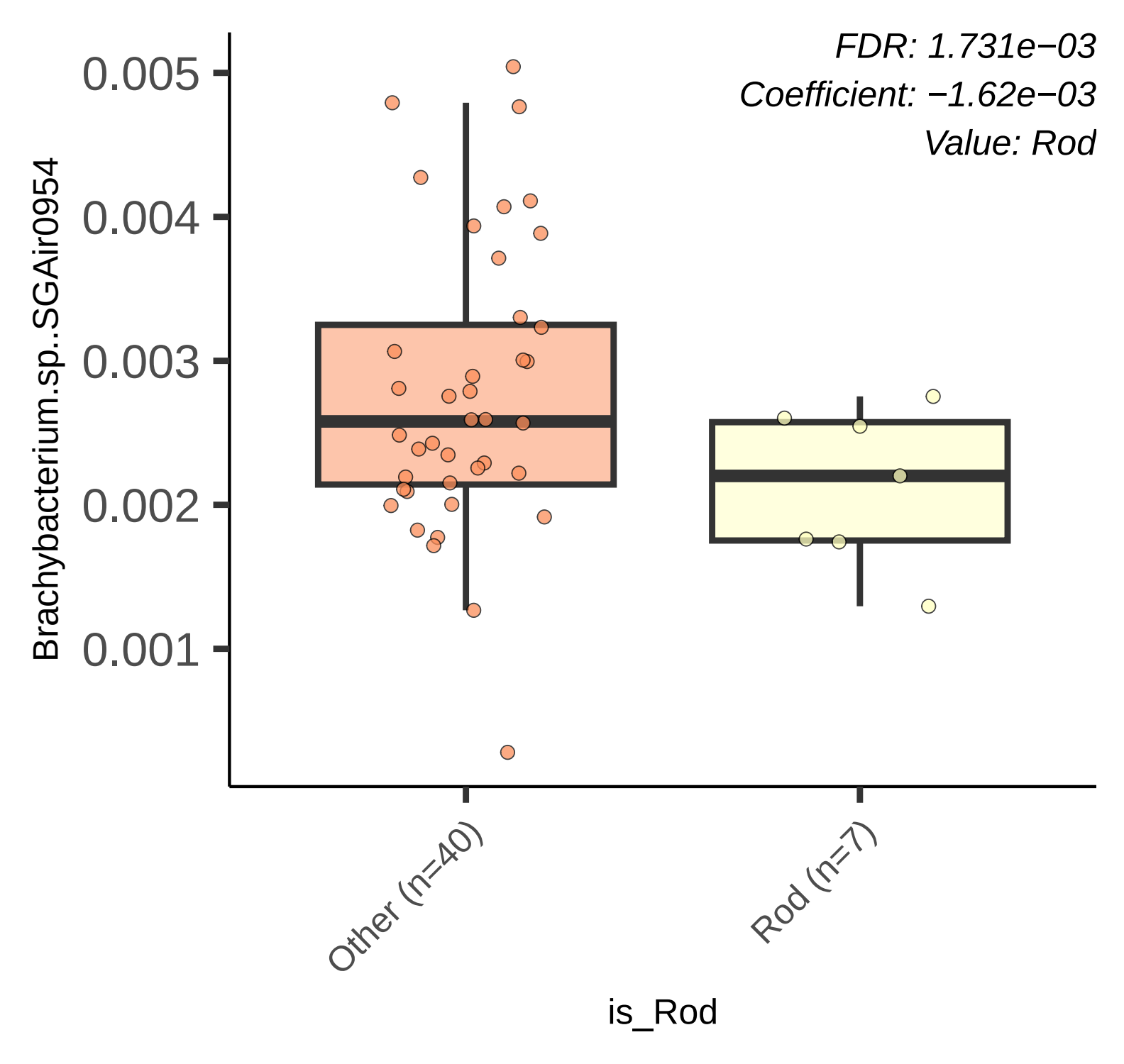

FDR: 4.866e−04
Coefficient: 1.47e−02
Value: Rod
Acinetobacter.indicus
0.04
0.03
0.02
0.01
0.00
Other (n=40)
Rod (n=7)
is_Rod

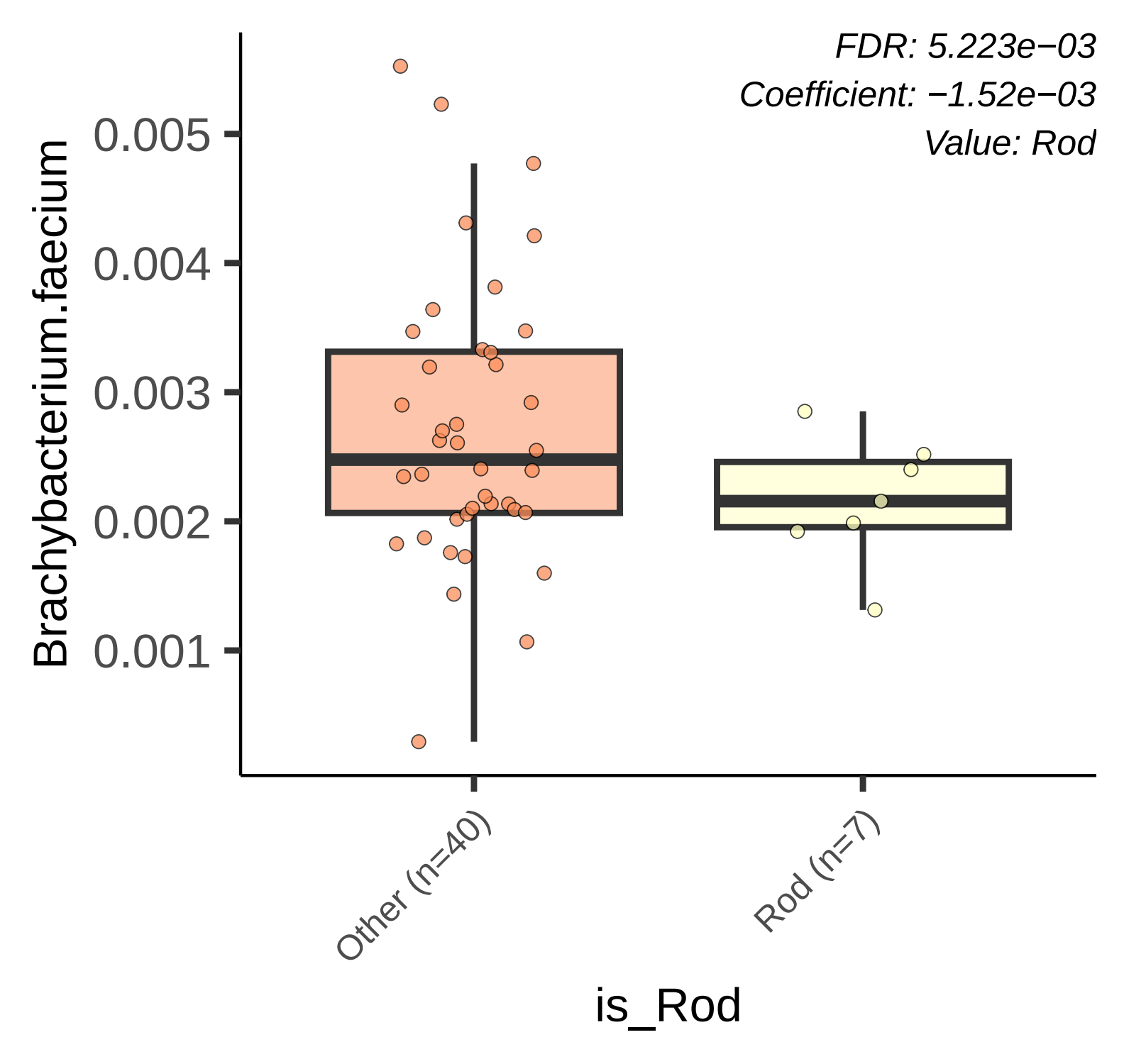

FDR: 1.202e−03
Coefficient: −1.47e−03
Value: Rod
Brachybacterium.sp..P6.10.X1
0.004
0.003
0.002
0.001
Other (n=40)
Rod (n=7)
is_Rod

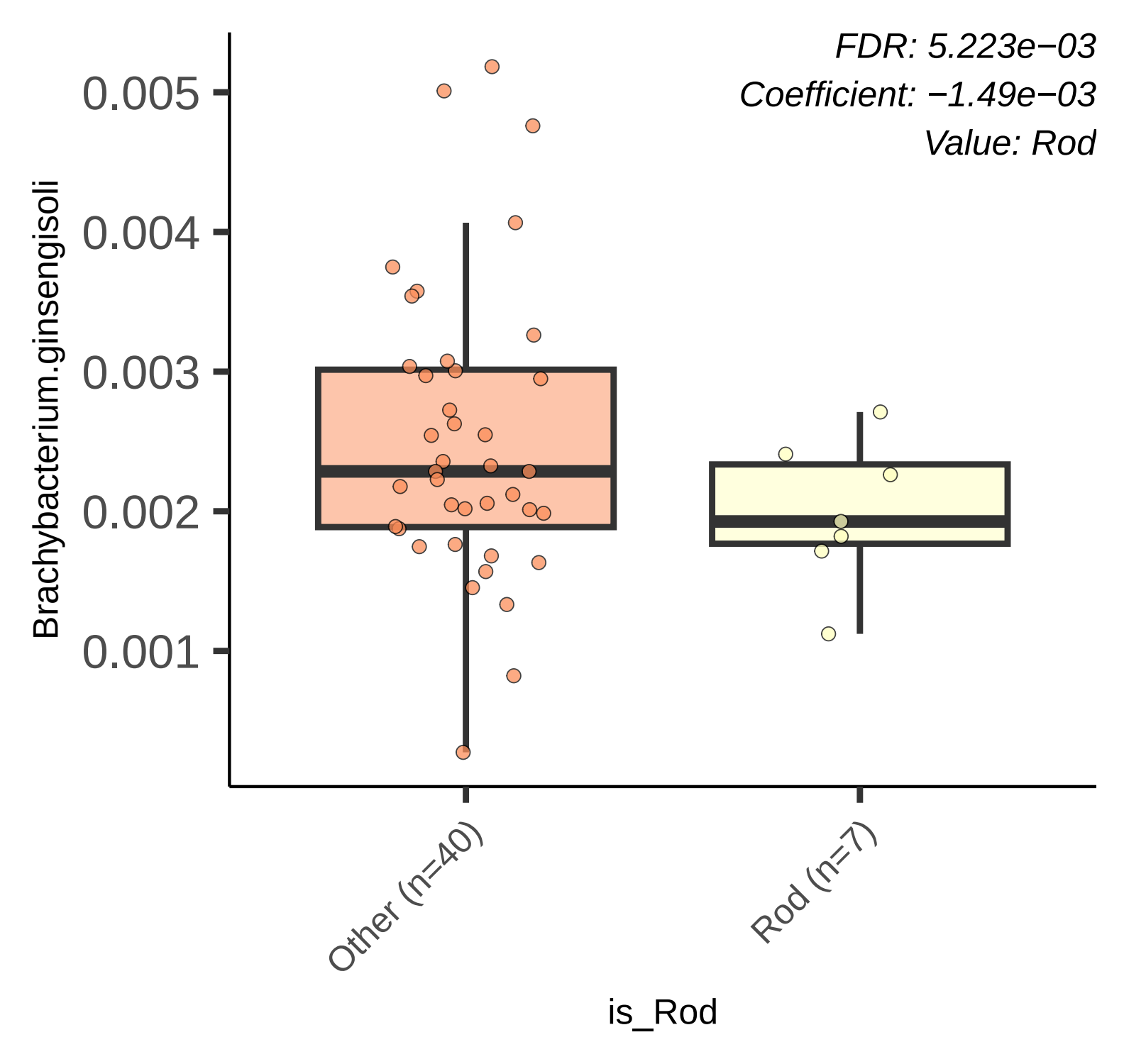

FDR: 1.202e−03
Coefficient: −1.30e−03
Value: Rod
0.003
0.002
0.001
Brachybacterium.vulturis
Other (n=40)
Rod (n=7)
is_Rod

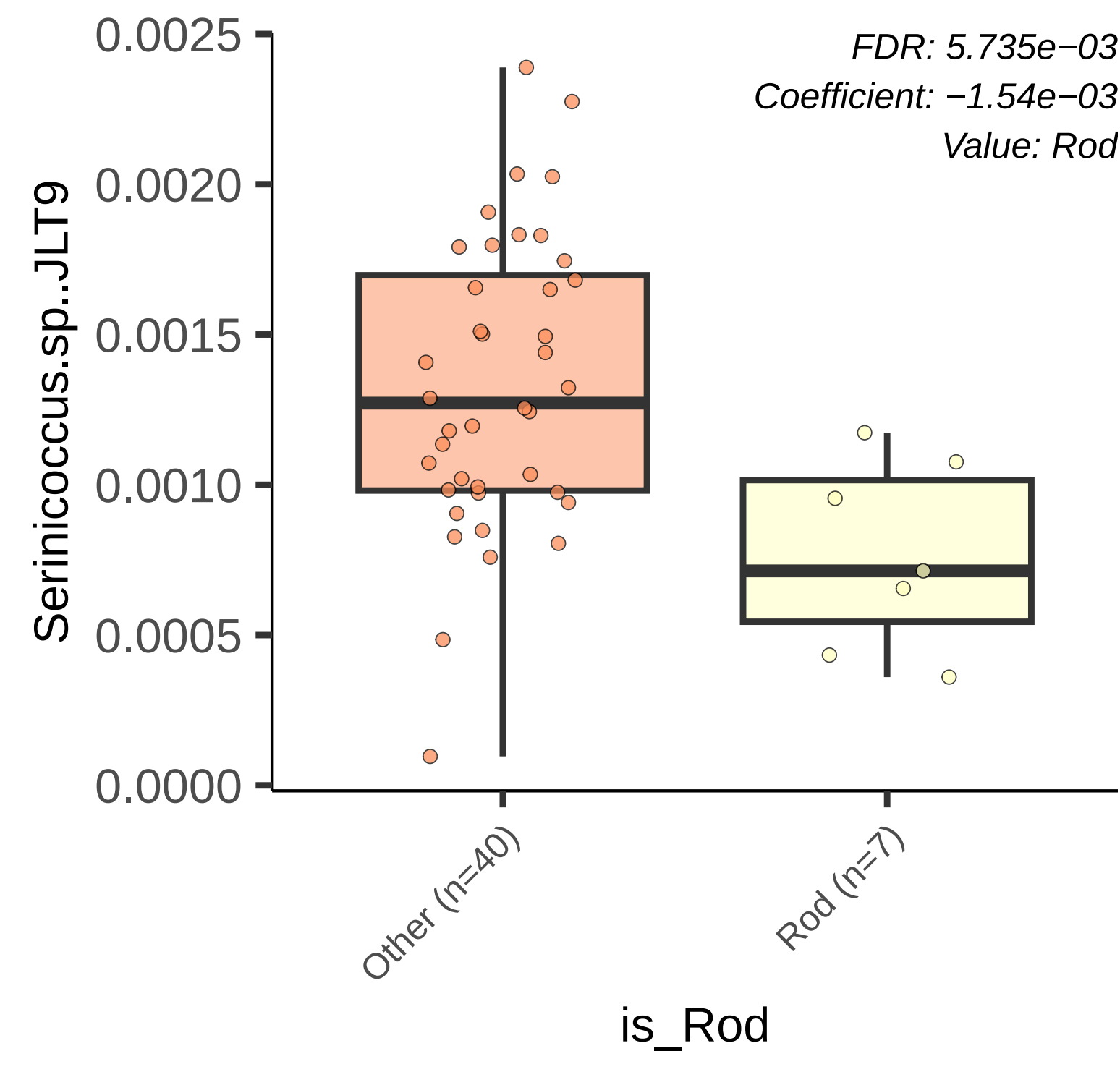

FDR: 1.202e−03
Coefficient: −1.28e−03
Value: Rod
Brachybacterium.avium
0.0025
0.0020
0.0015
0.0010
0.0005
Other (n=40)
Rod (n=7)
is_Rod

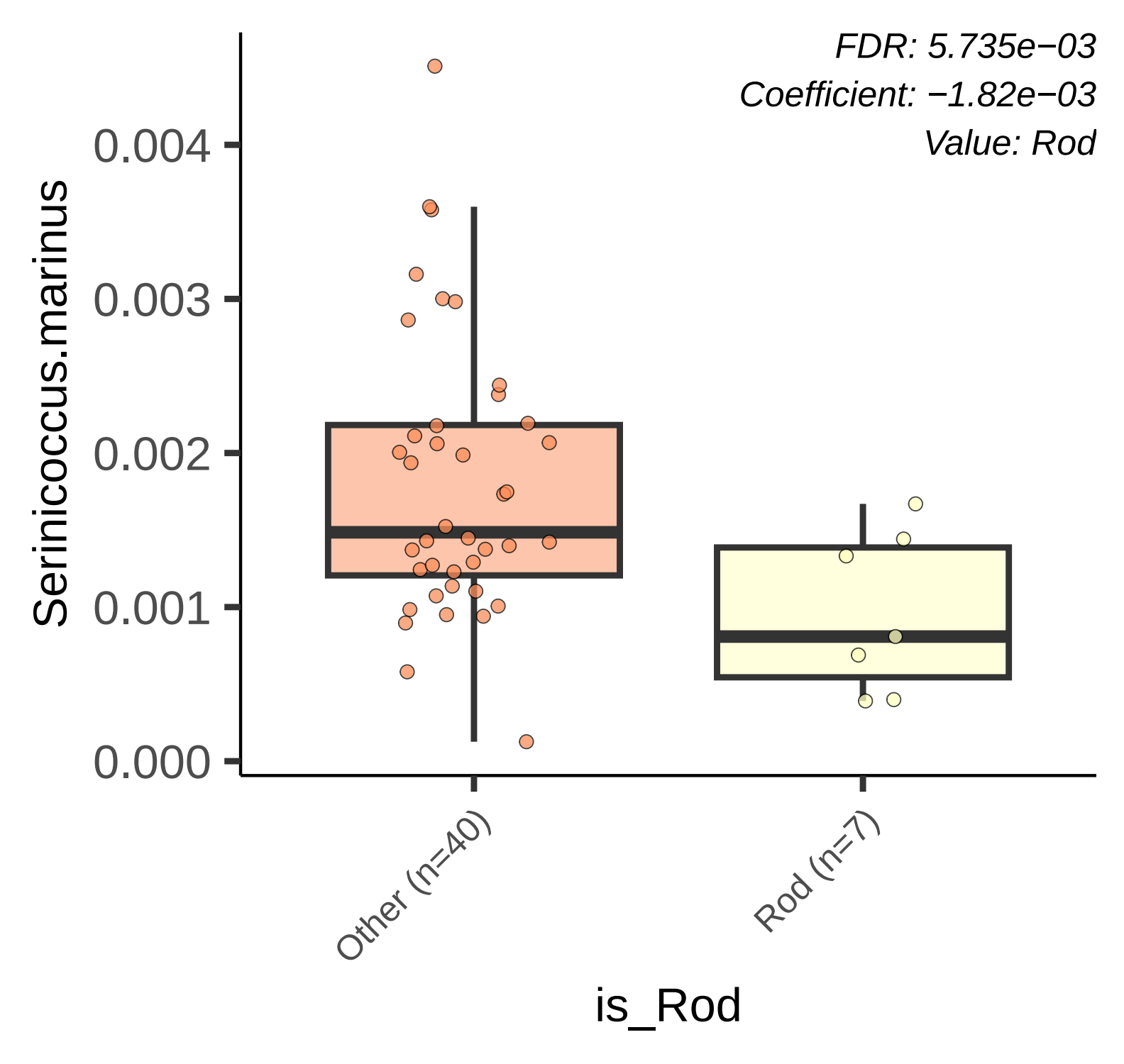

FDR: 1.498e−03
Coefficient: 5.69e−02
Value: Rod
Pseudomonas.stutzeri
0.25
0.20
0.15
0.10
0.05
Other (n=40)
Rod (n=7)
is_Rod

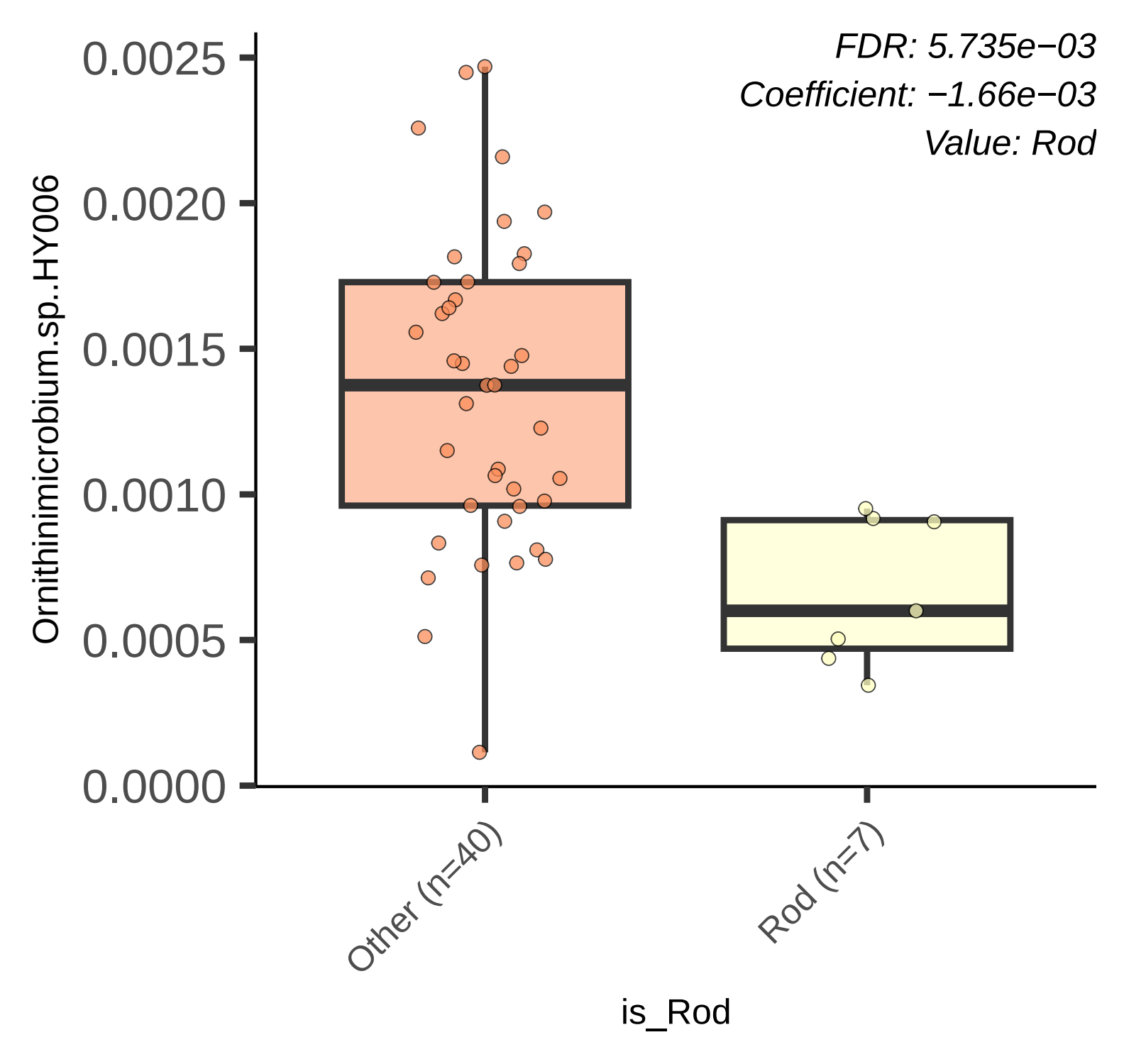

FDR: 1.731e−03
Coefficient: −1.62e−03
Value: Rod
0.005
0.004
0.003
0.002
0.001
Brachybacterium.sp..SGAir0954
Other (n=40)
Rod (n=7)
is_Rod

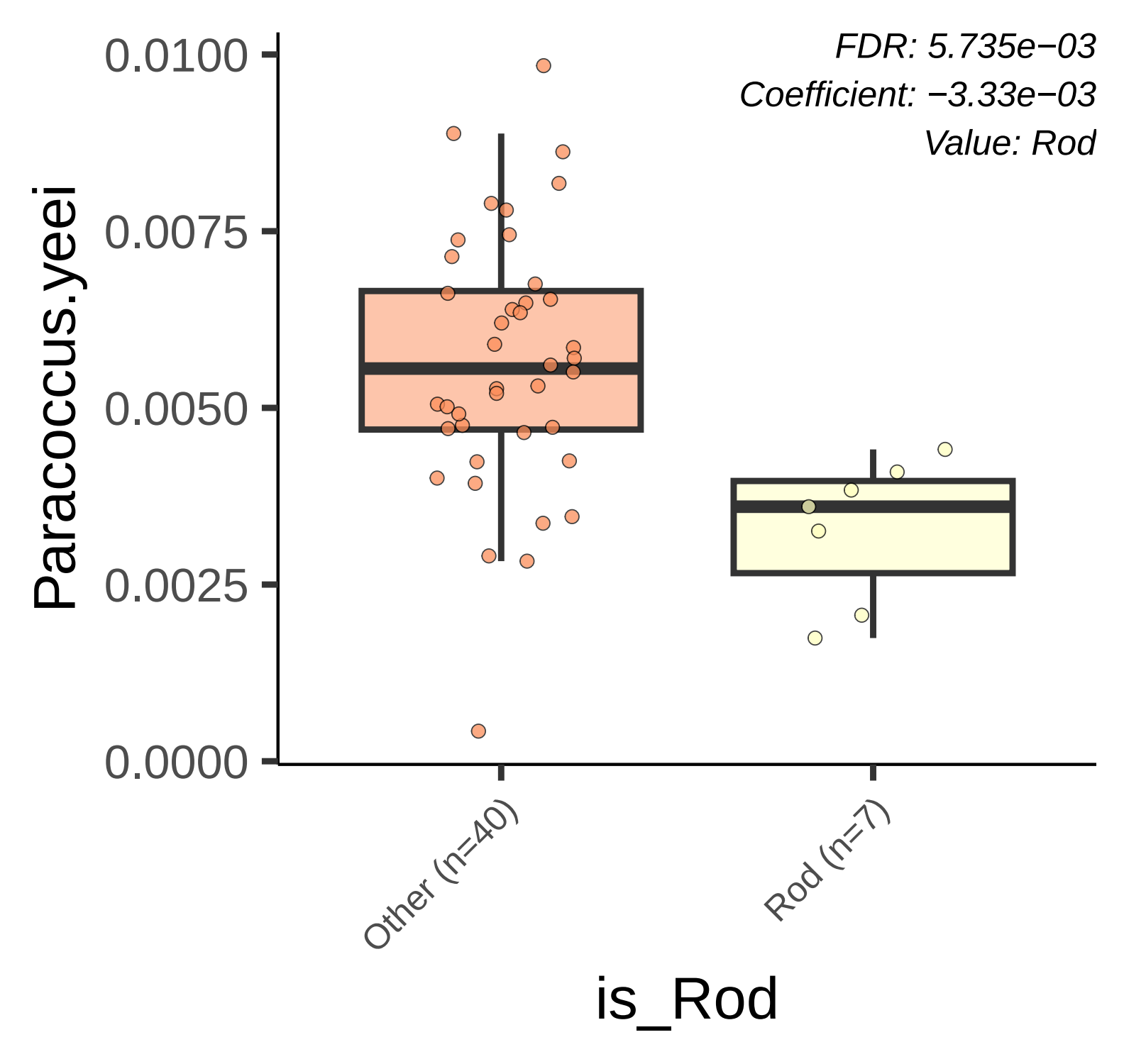

FDR: 5.223e−03
Coefficient: −1.52e−03
Value: Rod
Brachybacterium.faecium
0.005
0.004
0.003
0.002
0.001
Other (n=40)
Rod (n=7)
is_Rod

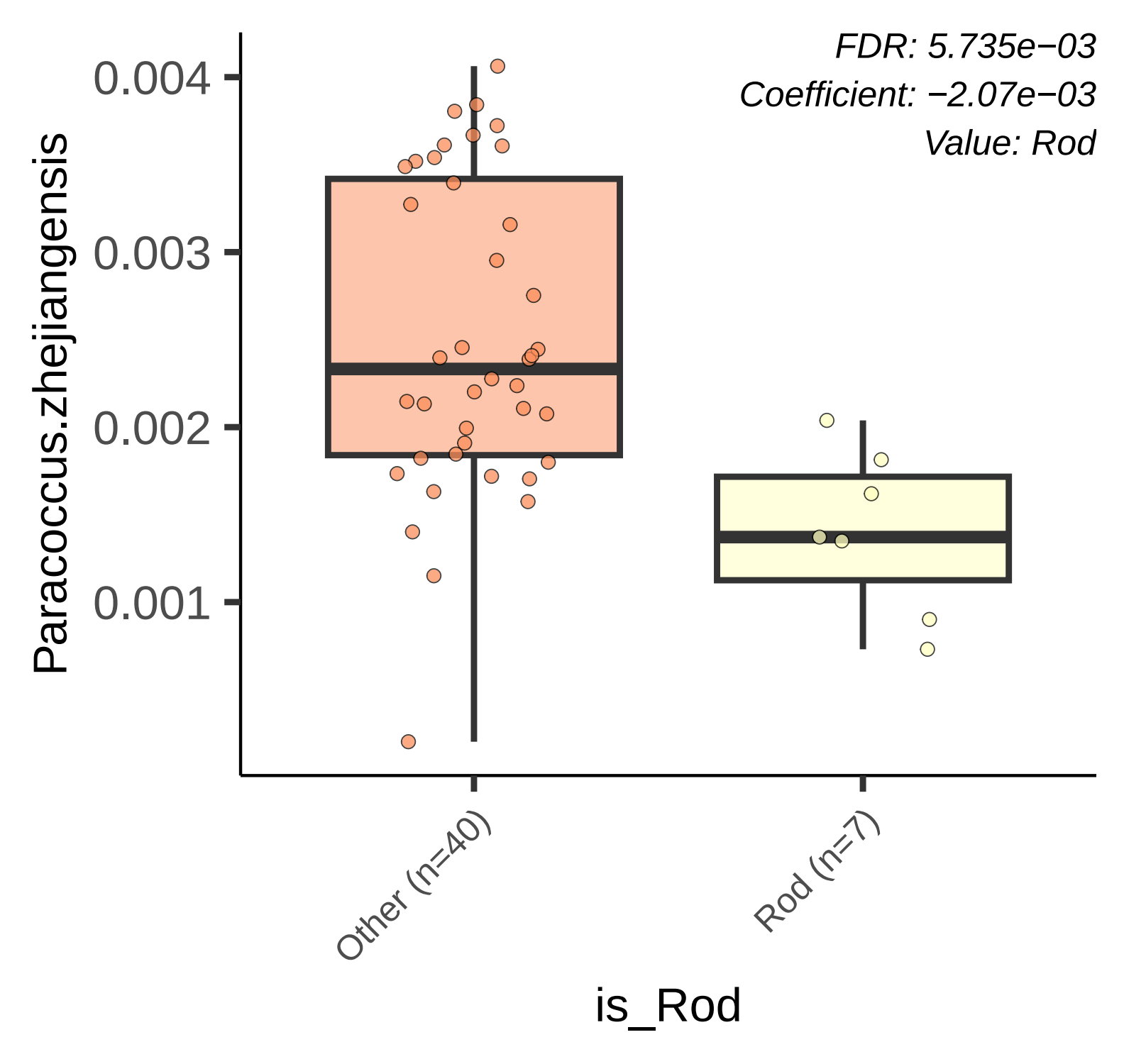

FDR: 5.223e−03
Coefficient: −1.49e−03
Value: Rod
Brachybacterium.ginsengisoli
0.005
0.004
0.003
0.002
0.001
Other (n=40)
Rod (n=7)
is_Rod

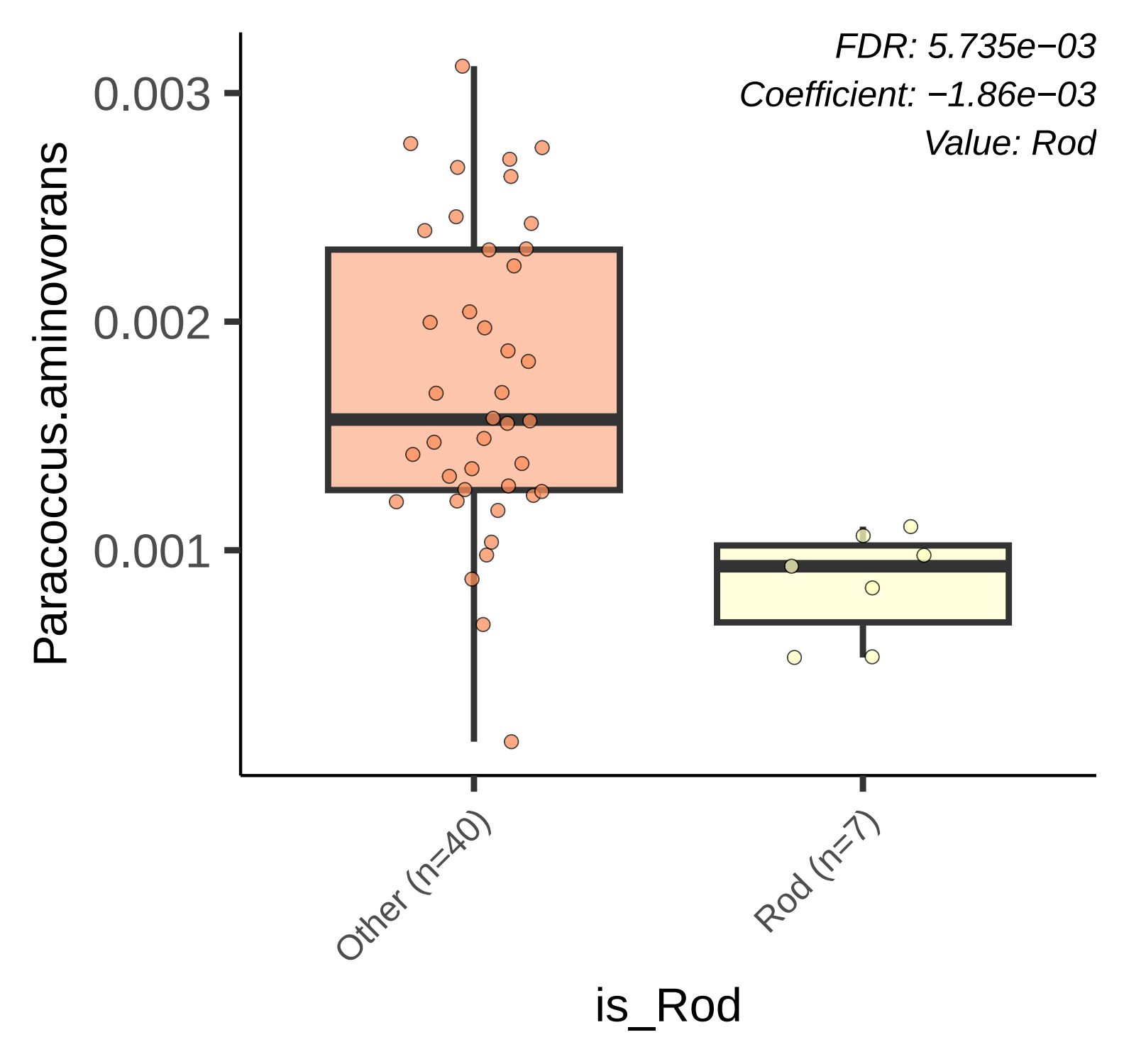

FDR: 5.735e−03
Coefficient: −1.54e−03
Value: Rod
Serinicoccus.sp..JLT9
0.0025
0.0020
0.0015
0.0010
0.0005
0.0000
Other (n=40)
Rod (n=7)
is_Rod

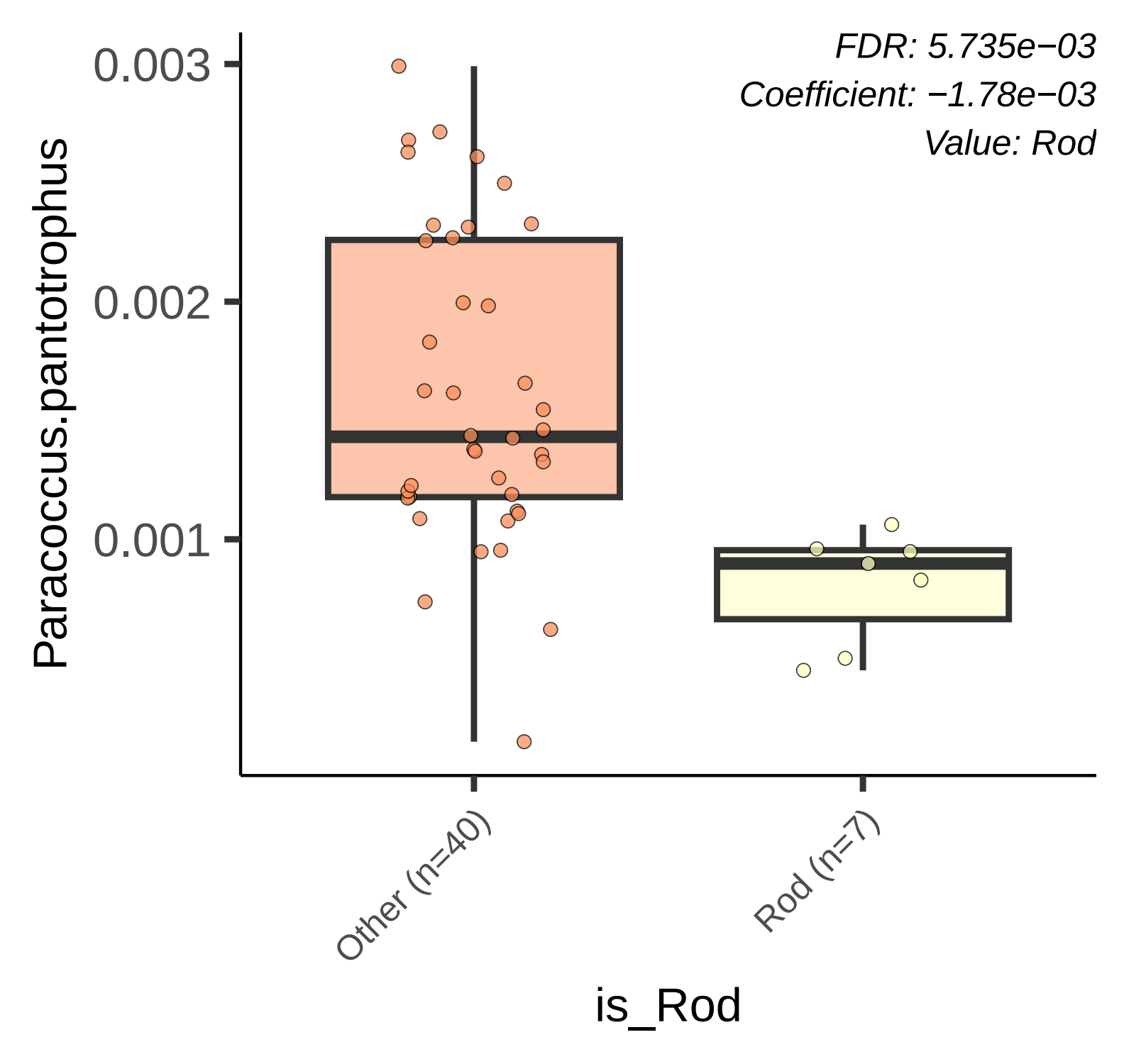

FDR: 5.735e−03
Coefficient: −1.82e−03
Value: Rod
Serinicoccus.marinus
0.004
0.003
0.002
0.001
0.000
Other (n=40)
Rod (n=7)
is_Rod

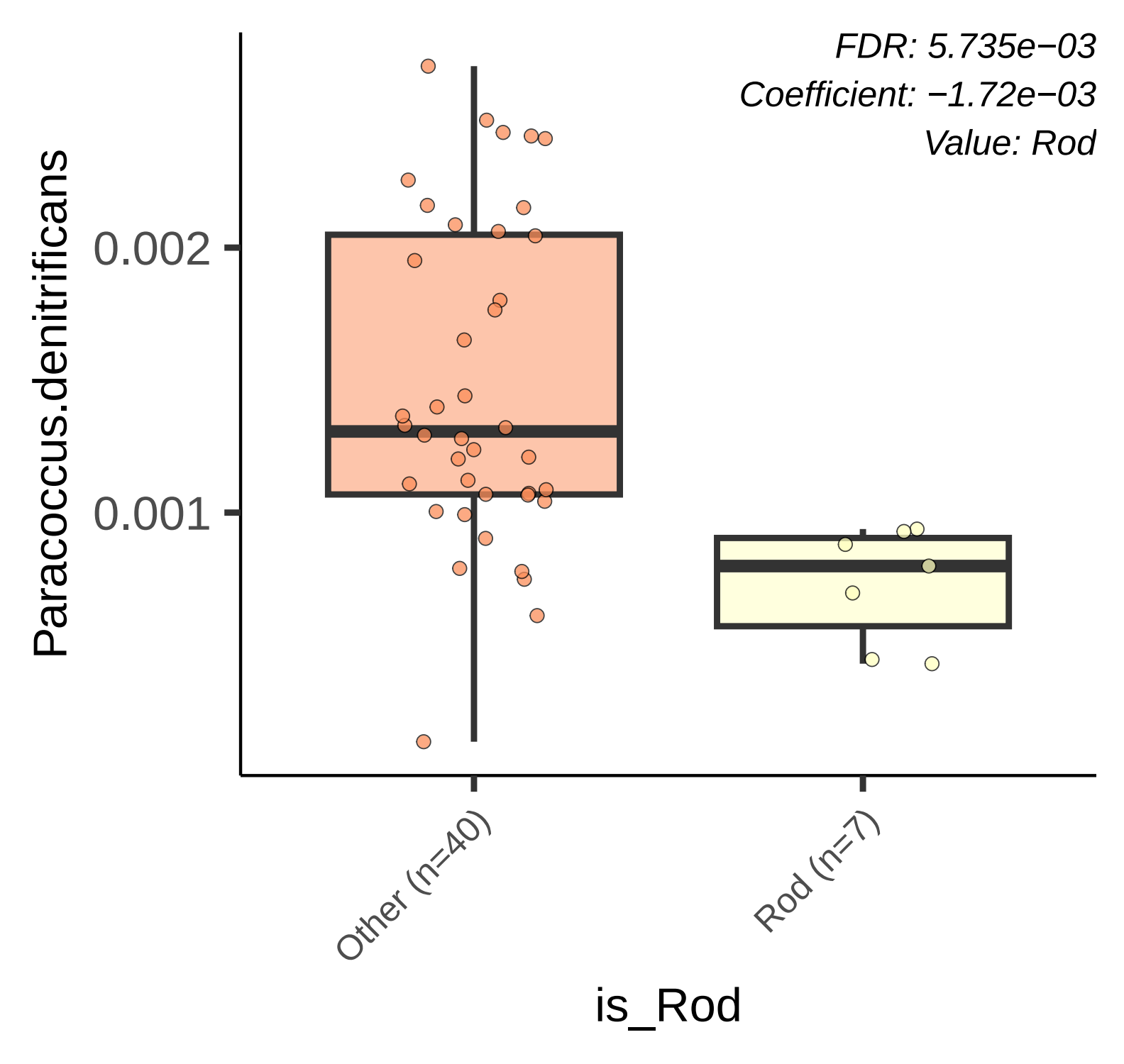

FDR: 5.735e−03
Coefficient: −1.66e−03
Value: Rod
Ornithinimicrobium.sp..HY006
0.0025
0.0020
0.0015
0.0010
0.0005
0.0000
Other (n=40)
Rod (n=7)
is_Rod

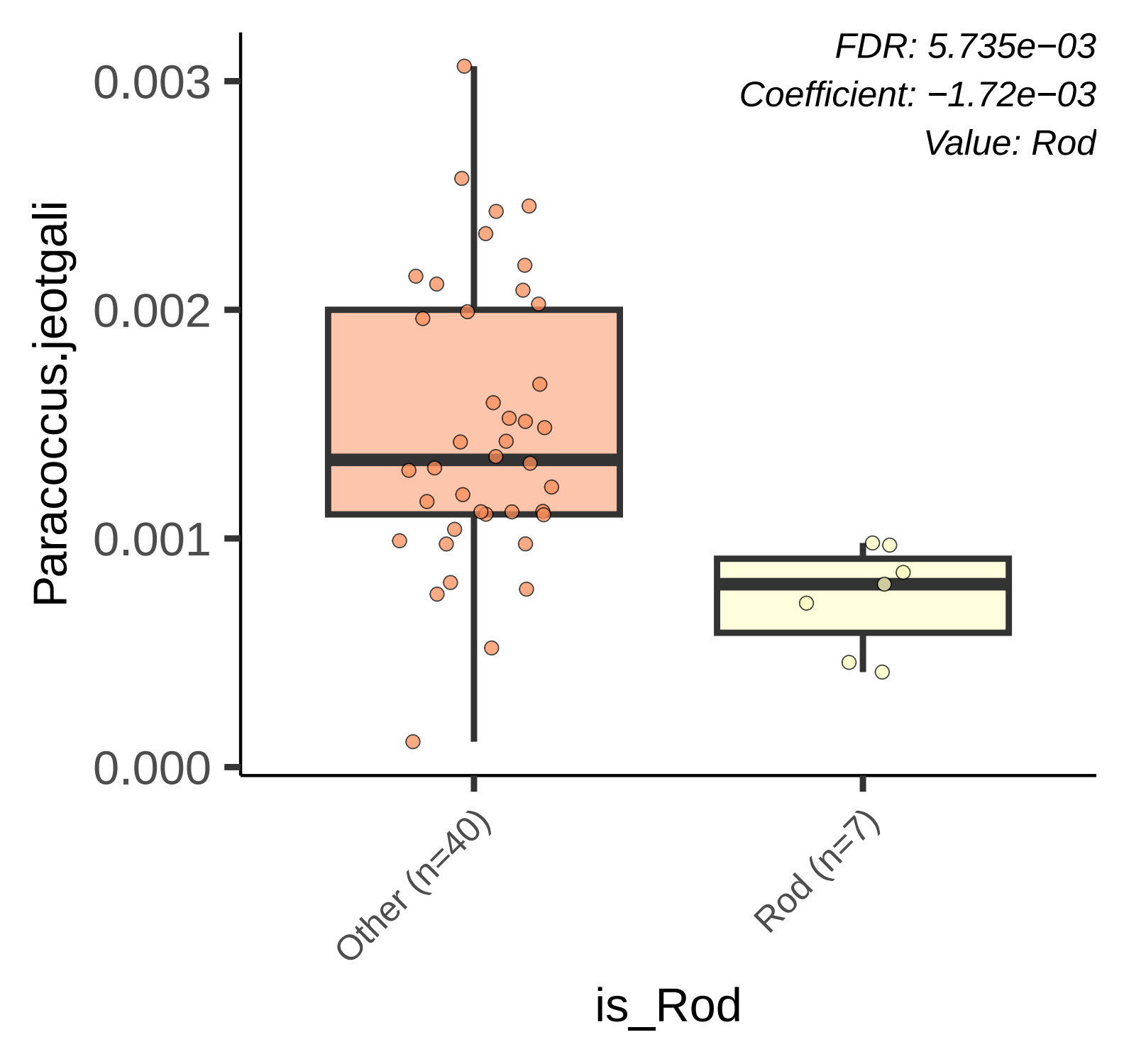

FDR: 5.735e−03
Coefficient: −3.33e−03
Value: Rod
Paracoccus.yeei
0.0100
0.0075
0.0050
0.0025
0.0000
Other (n=40)
Rod (n=7)
is_Rod

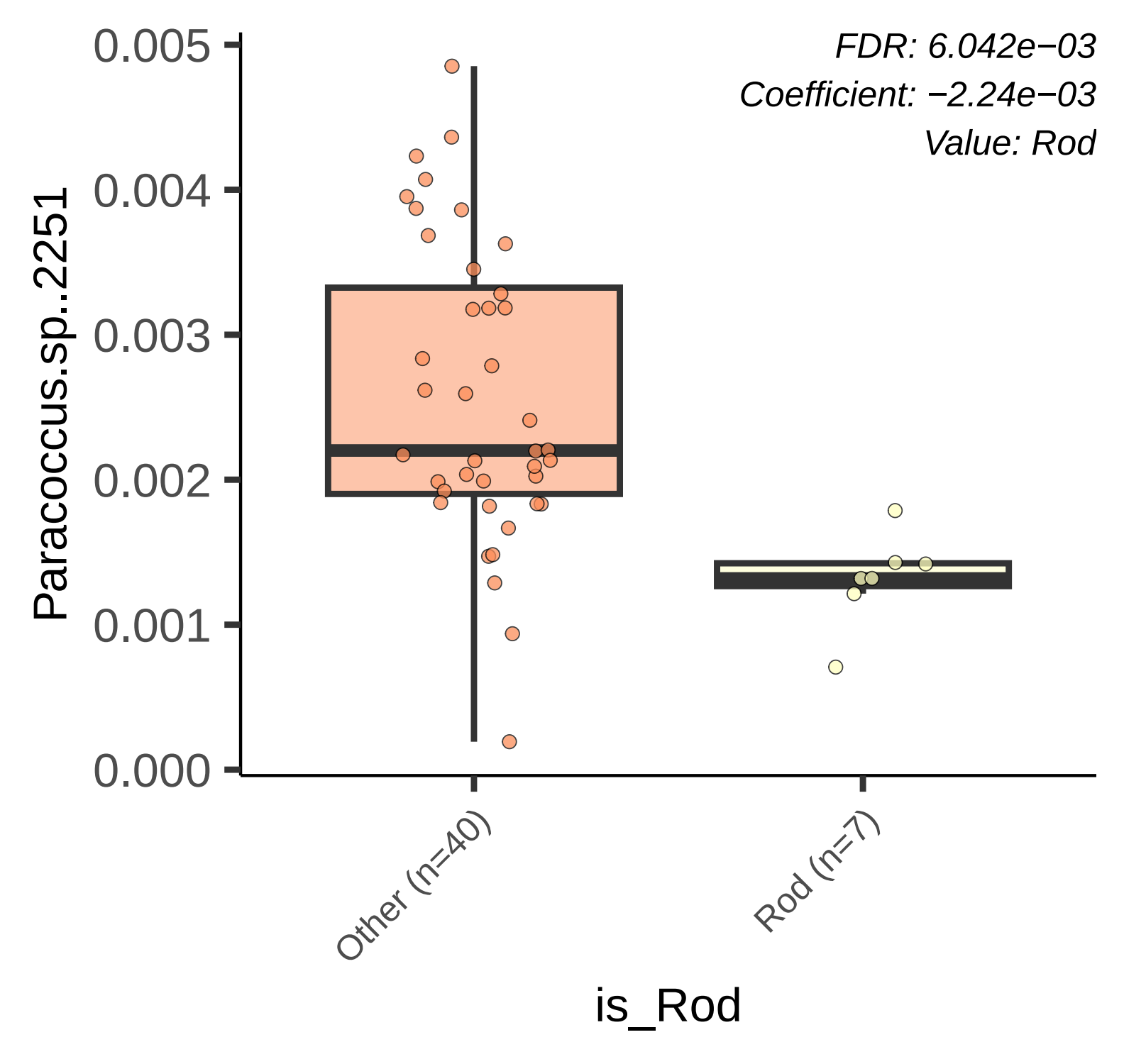

FDR: 5.735e−03
Coefficient: −2.07e−03
Value: Rod
Paracoccus.zhejiangensis
0.004
0.003
0.002
0.001
Other (n=40)
Rod (n=7)
is_Rod

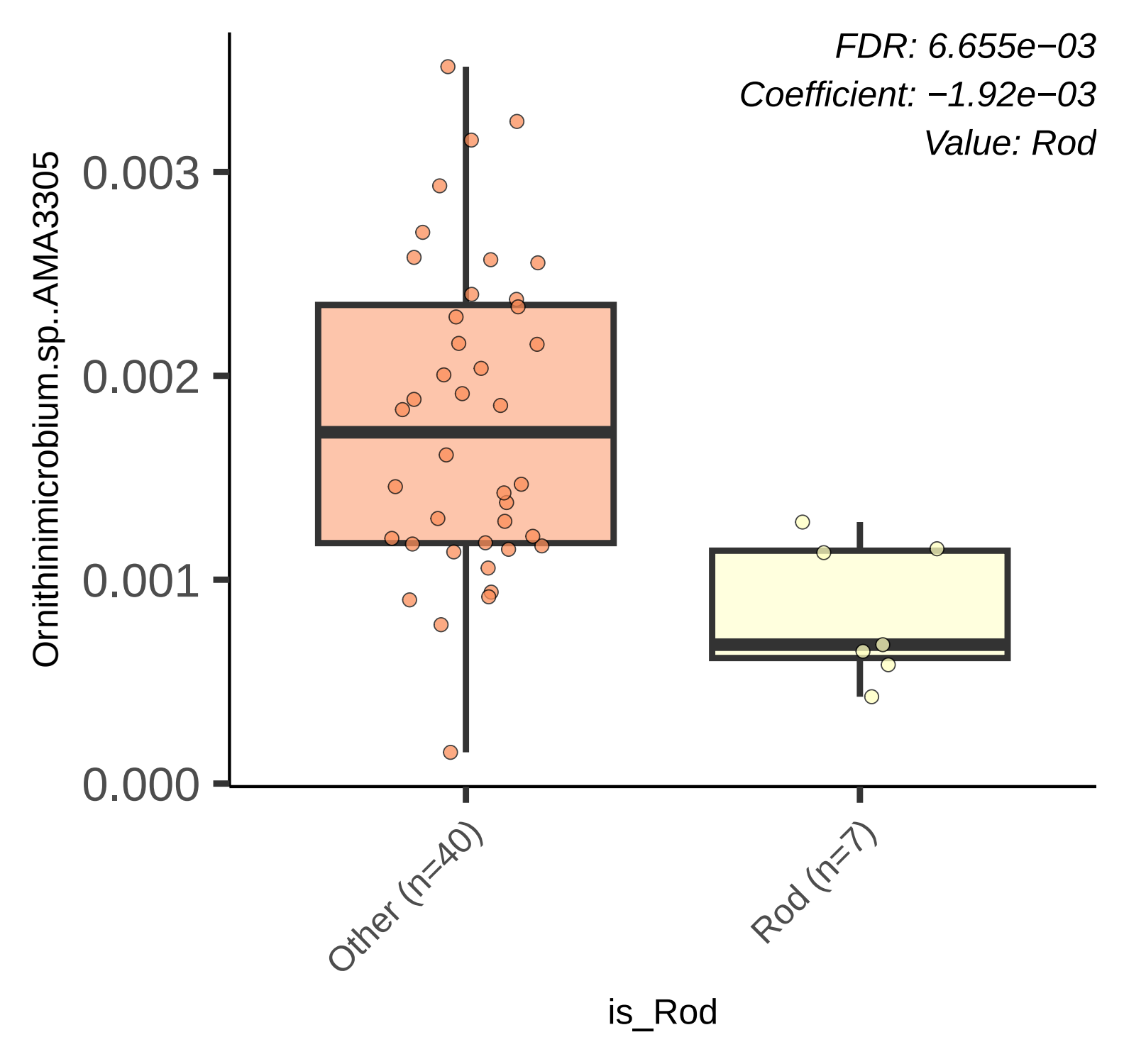

FDR: 5.735e−03
Coefficient: −1.86e−03
Value: Rod
0.003
0.002
0.001
Paracoccus.aminovorans
Other (n=40)
Rod (n=7)
is_Rod

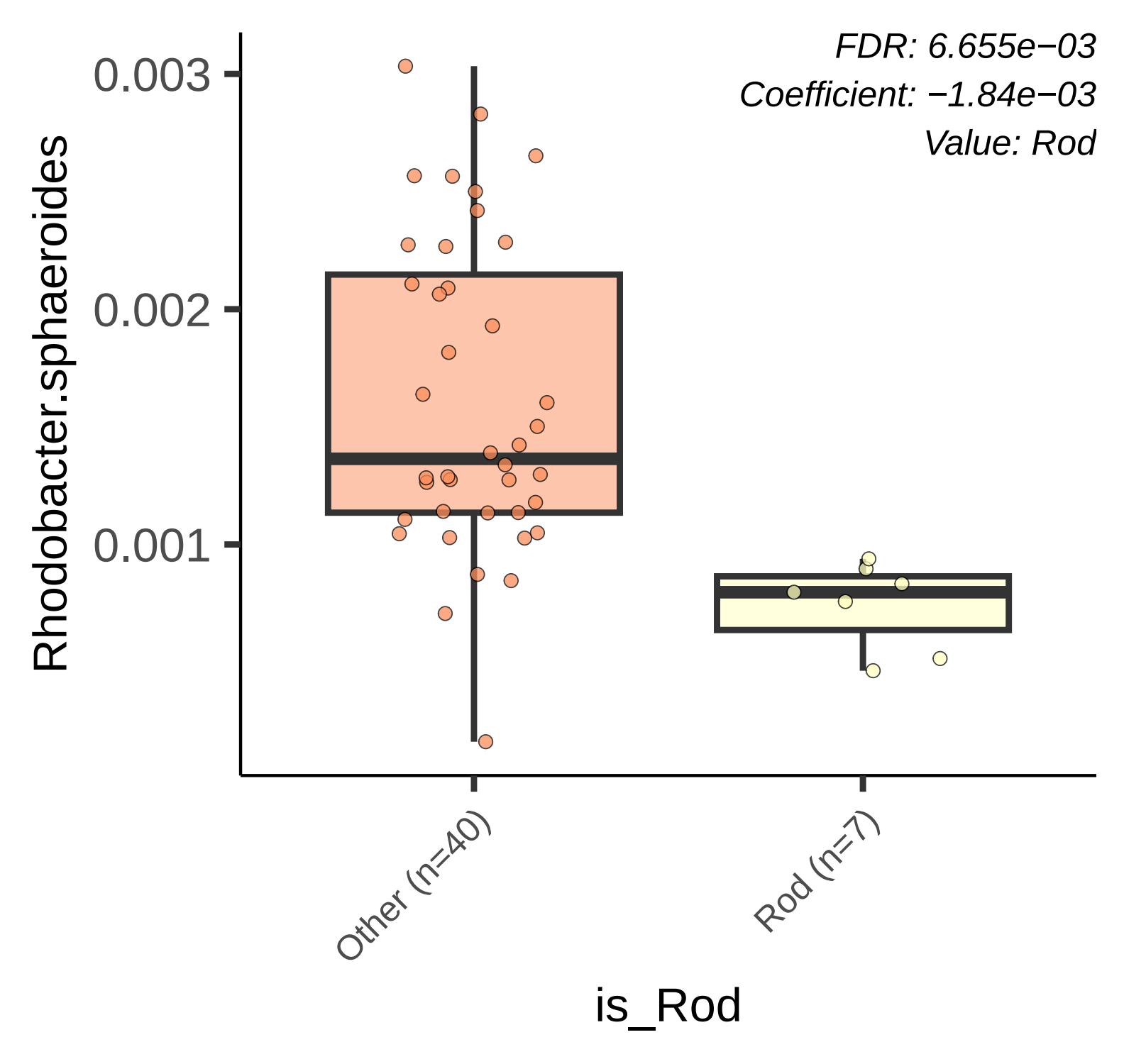

FDR: 5.735e−03
Coefficient: −1.78e−03
Value: Rod
0.003
0.002
0.001
Paracoccus.pantotrophus
Other (n=40)
Rod (n=7)
is_Rod

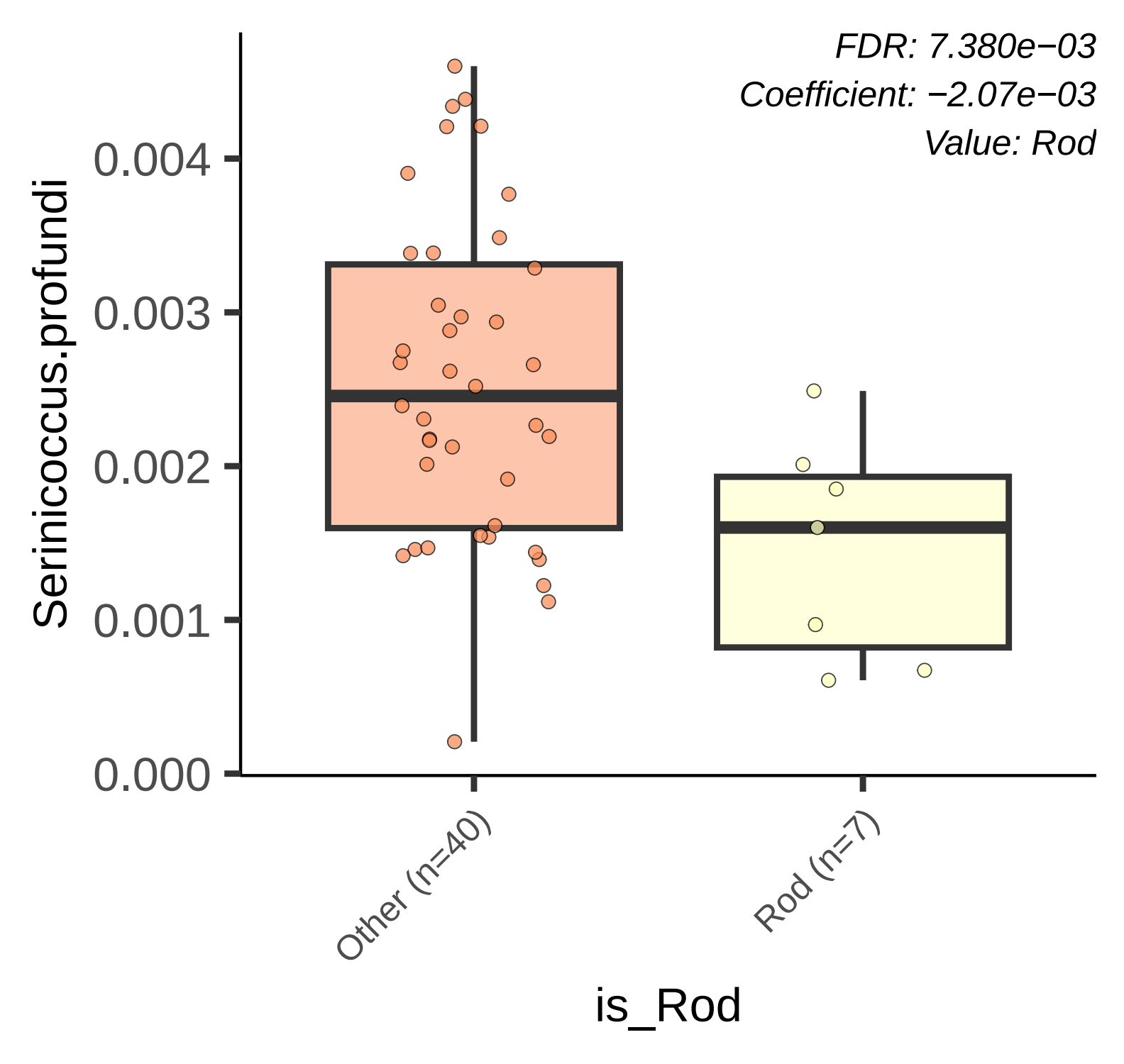

FDR: 5.735e−03
Coefficient: −1.72e−03
Value: Rod
Paracoccus.denitrificans
0.002
0.001
Other (n=40)
Rod (n=7)
is_Rod

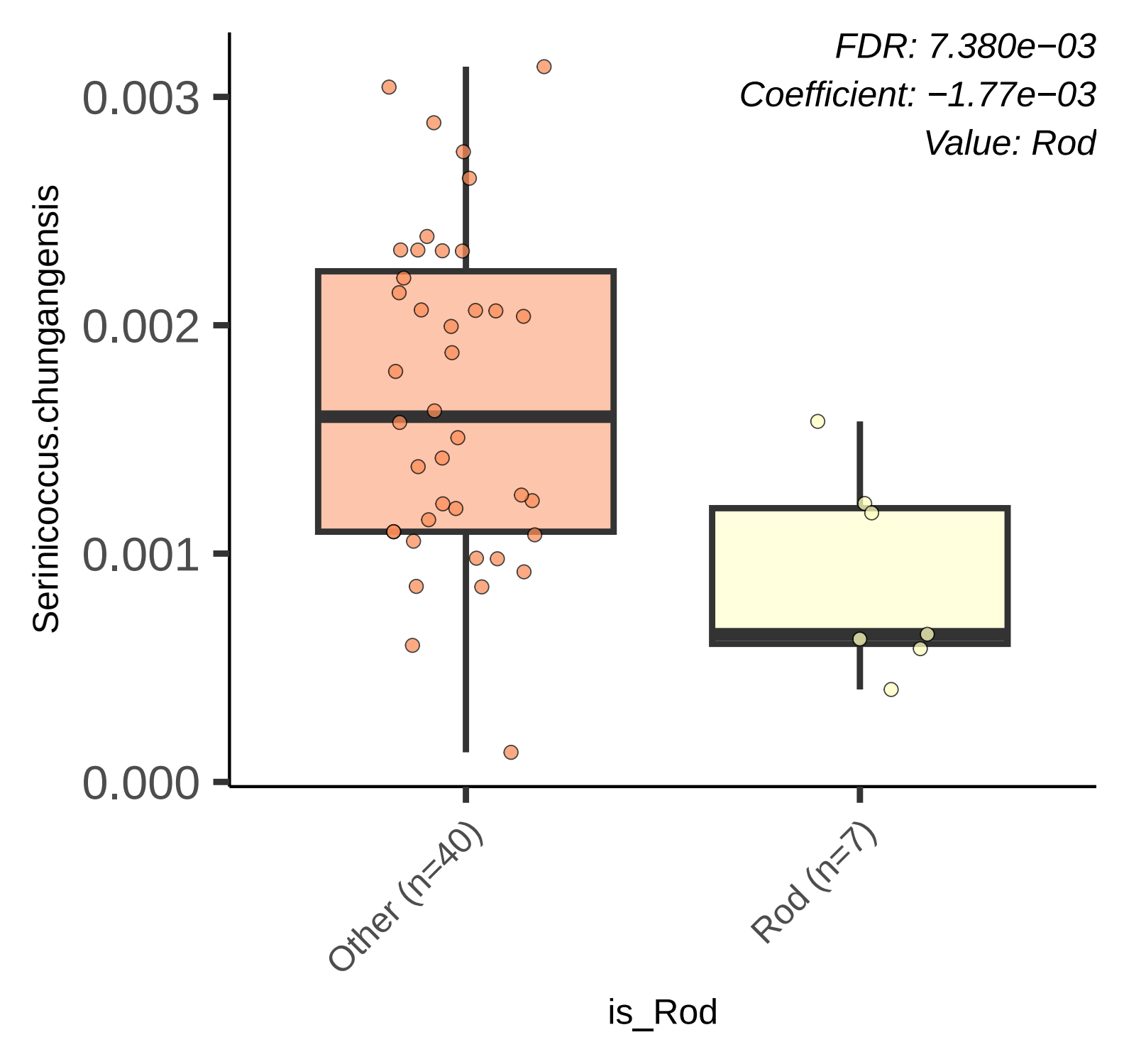

FDR: 5.735e−03
Coefficient: −1.72e−03
Value: Rod
Paracoccus.jeotgali
0.003
0.002
0.001
0.000
Other (n=40)
Rod (n=7)
is_Rod

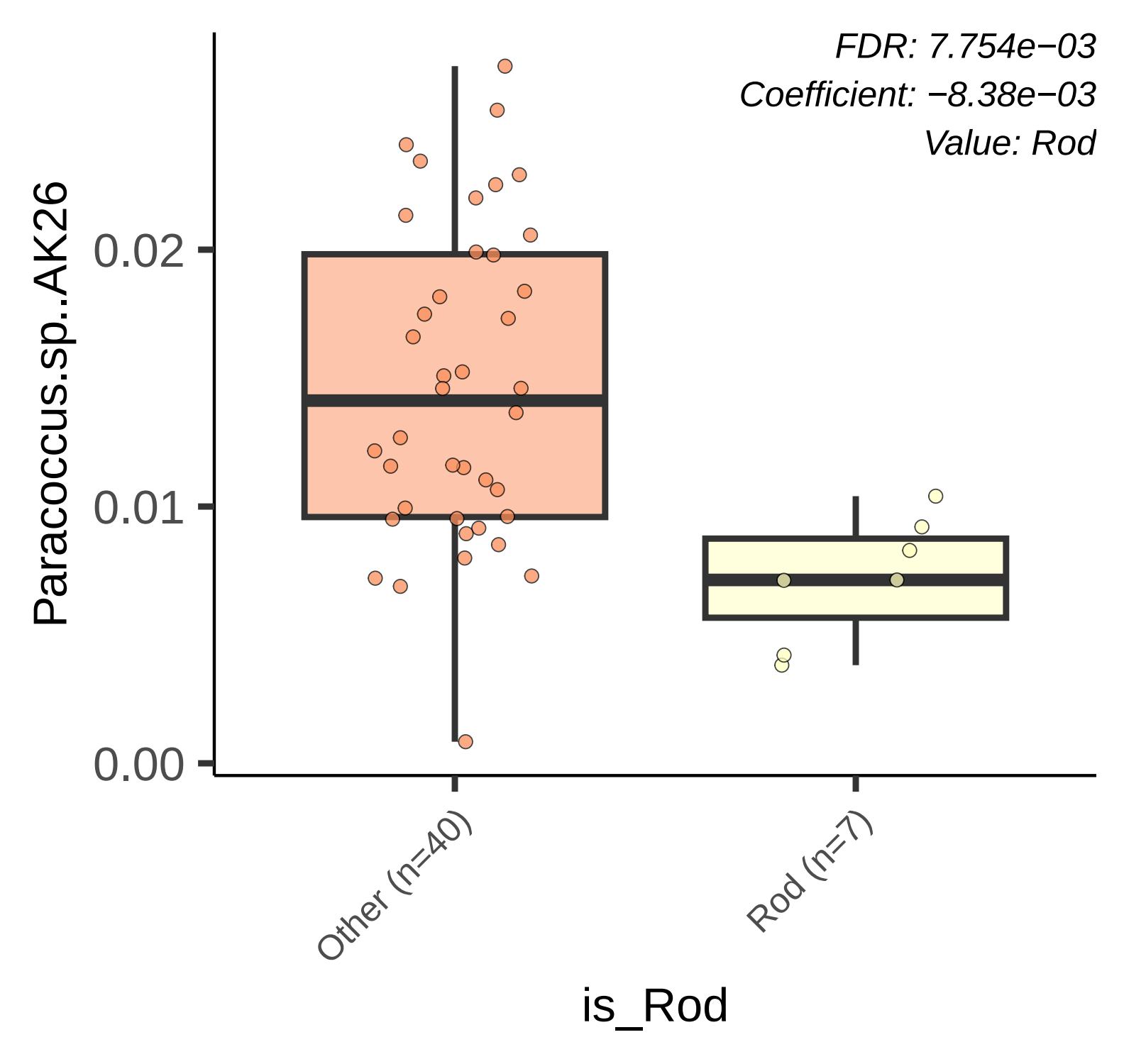

FDR: 6.042e−03
Coefficient: −2.24e−03
Value: Rod
Paracoccus.sp..2251
0.005
0.004
0.003
0.002
0.001
0.000
Other (n=40)
Rod (n=7)
is_Rod

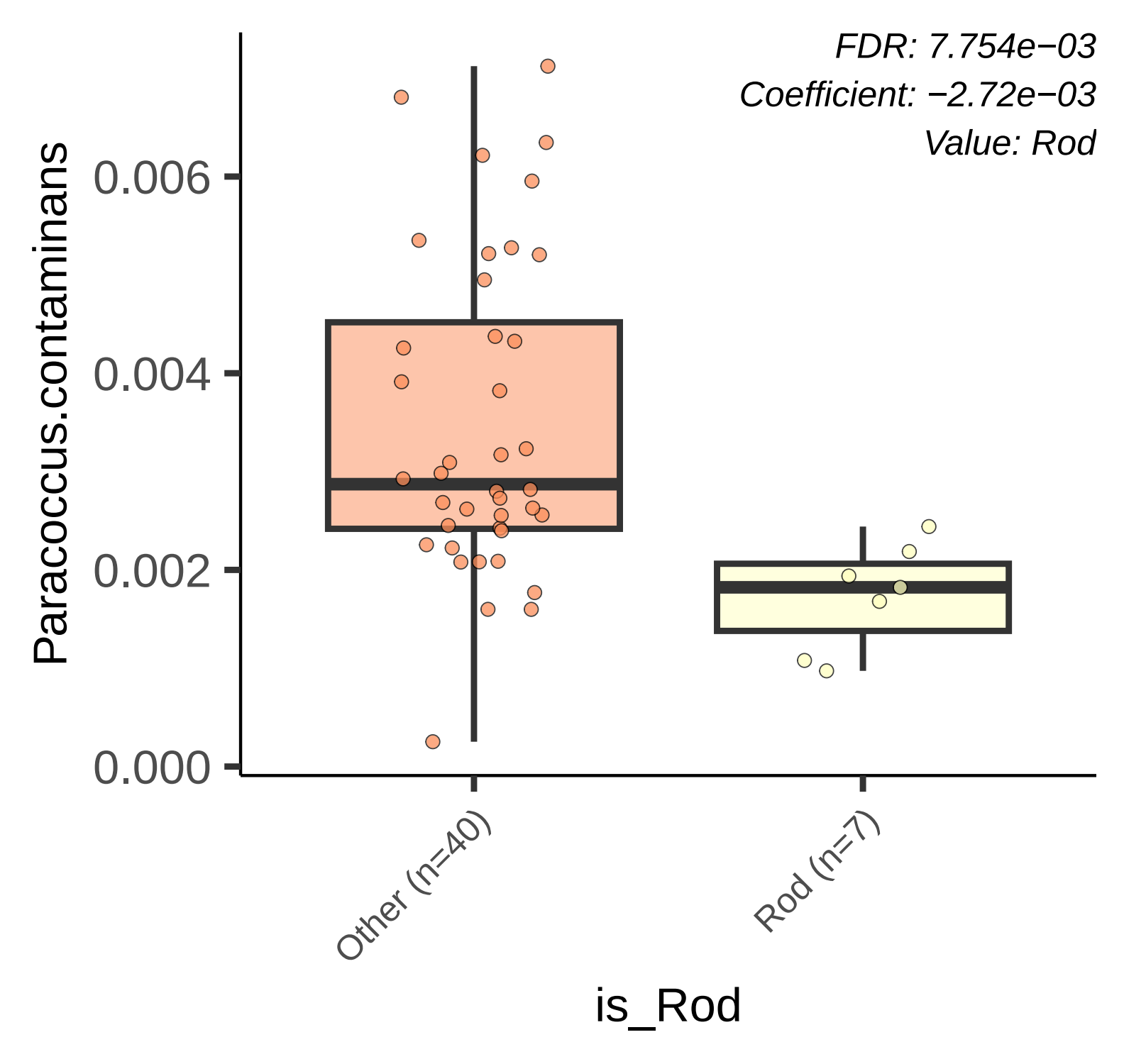

FDR: 6.655e−03
Coefficient: −1.92e−03
Value: Rod
Ornithinimicrobium.sp..AMA3305
0.003
0.002
0.001
0.000
Other (n=40)
Rod (n=7)
is_Rod

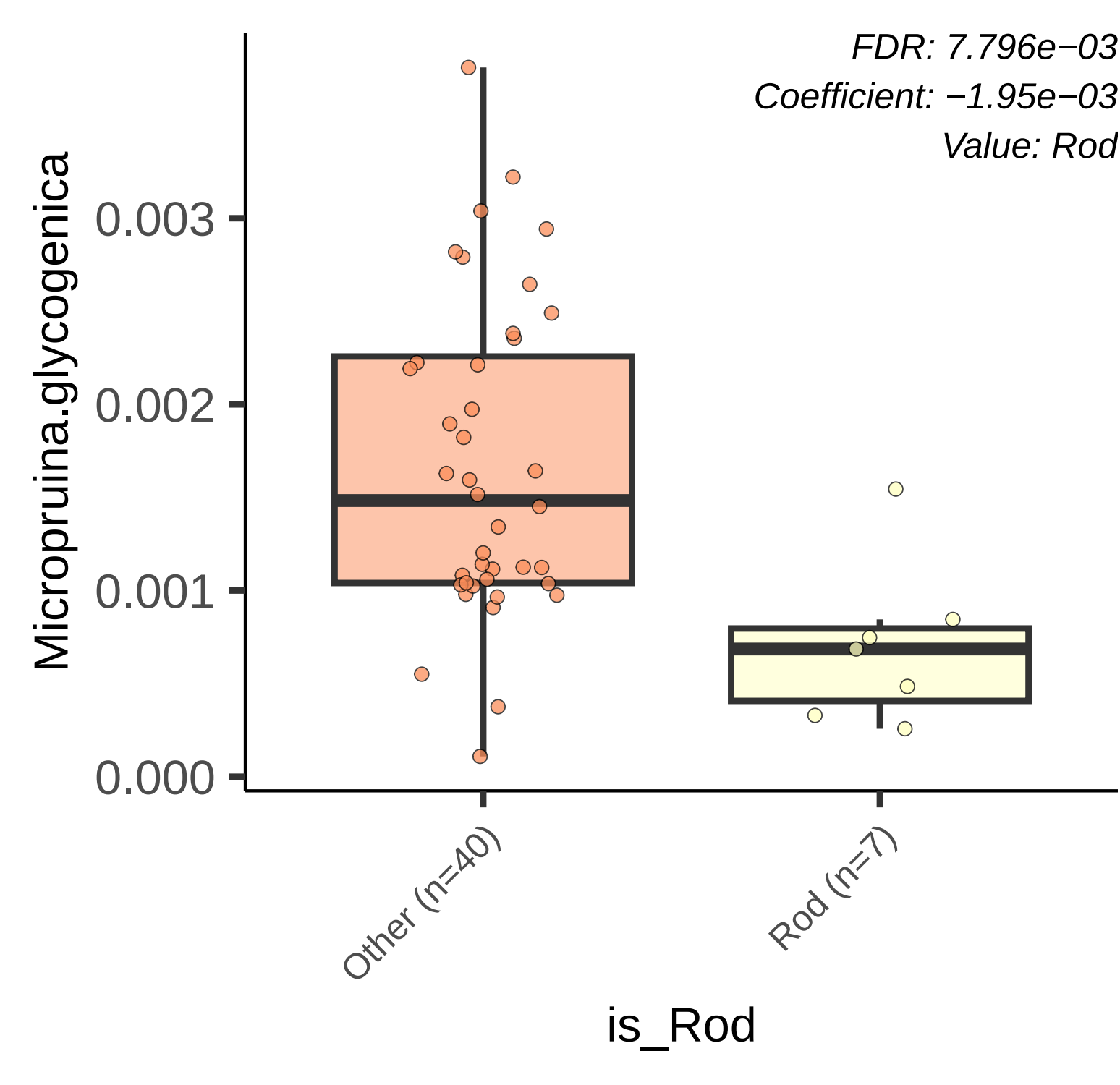

FDR: 6.655e−03
Coefficient: −1.84e−03
Value: Rod
Rhodobacter.sphaeroides
0.003
0.002
0.001
Other (n=40)
Rod (n=7)
is_Rod

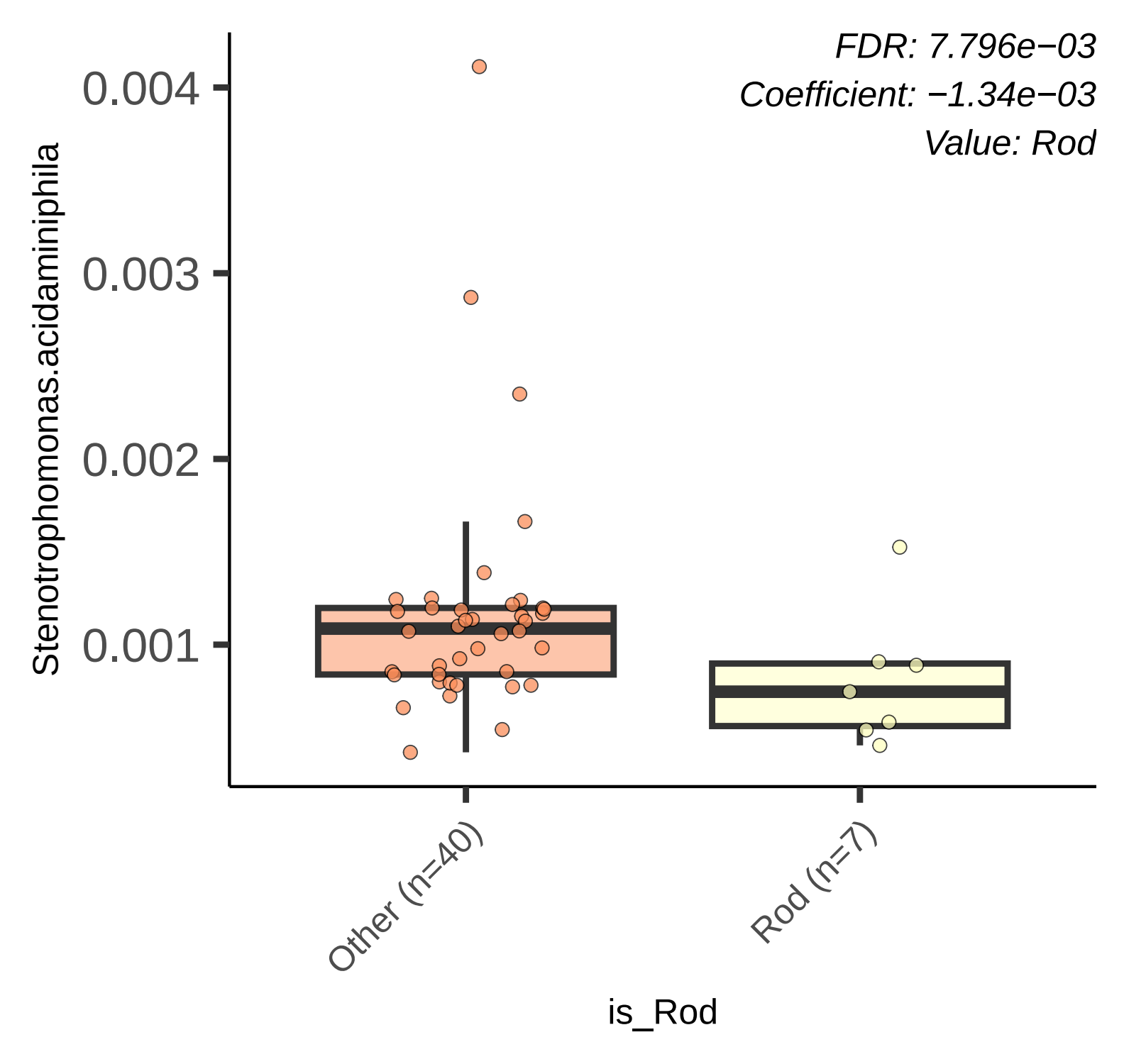

FDR: 7.380e−03
Coefficient: −2.07e−03
Value: Rod
Serinicoccus.profundi
0.004
0.003
0.002
0.001
0.000
Other (n=40)
Rod (n=7)
is_Rod

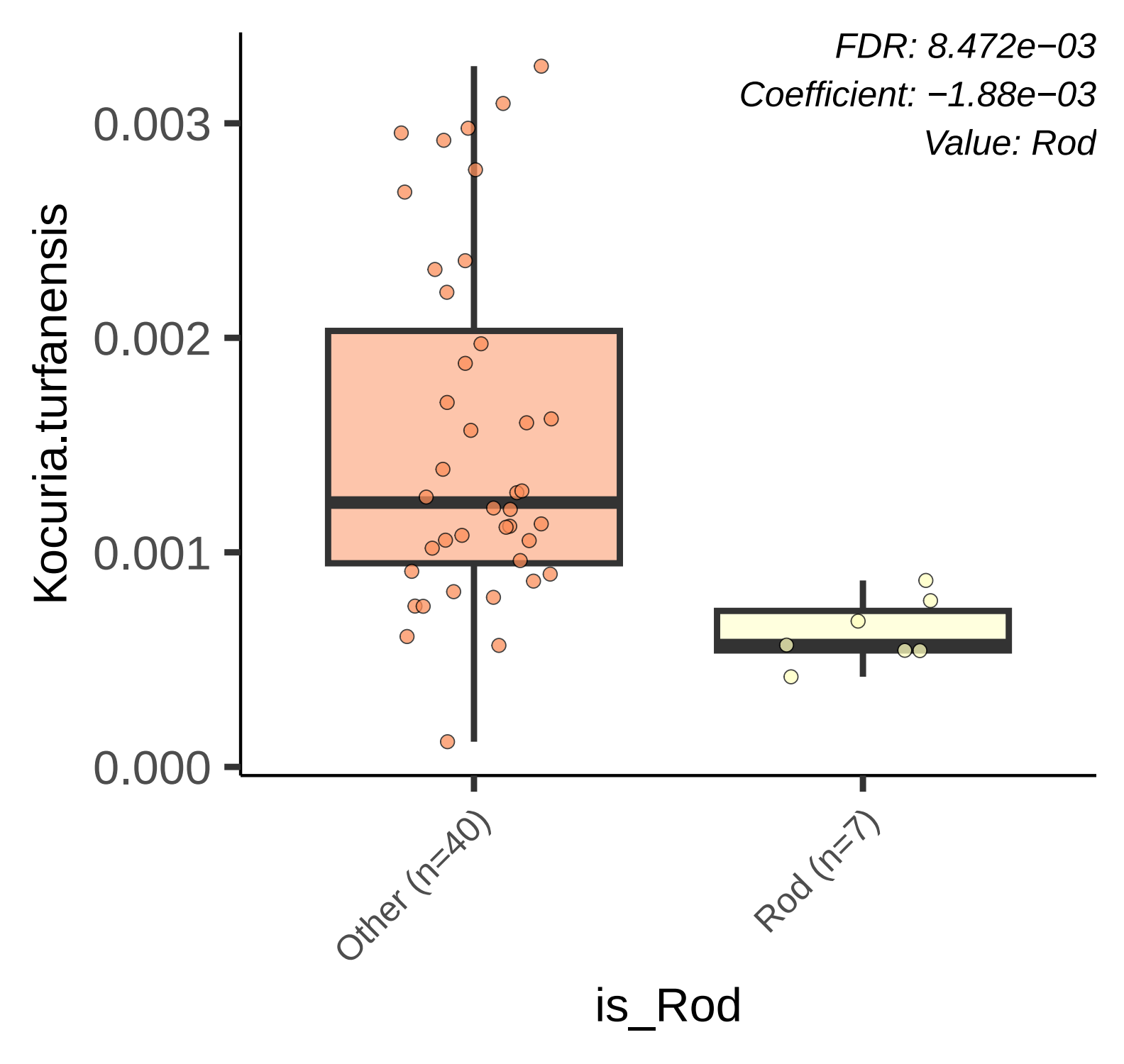

FDR: 7.380e−03
Coefficient: −1.77e−03
Value: Rod
Serinicoccus.chungangensis
0.003
0.002
0.001
0.000
Other (n=40)
Rod (n=7)
is_Rod

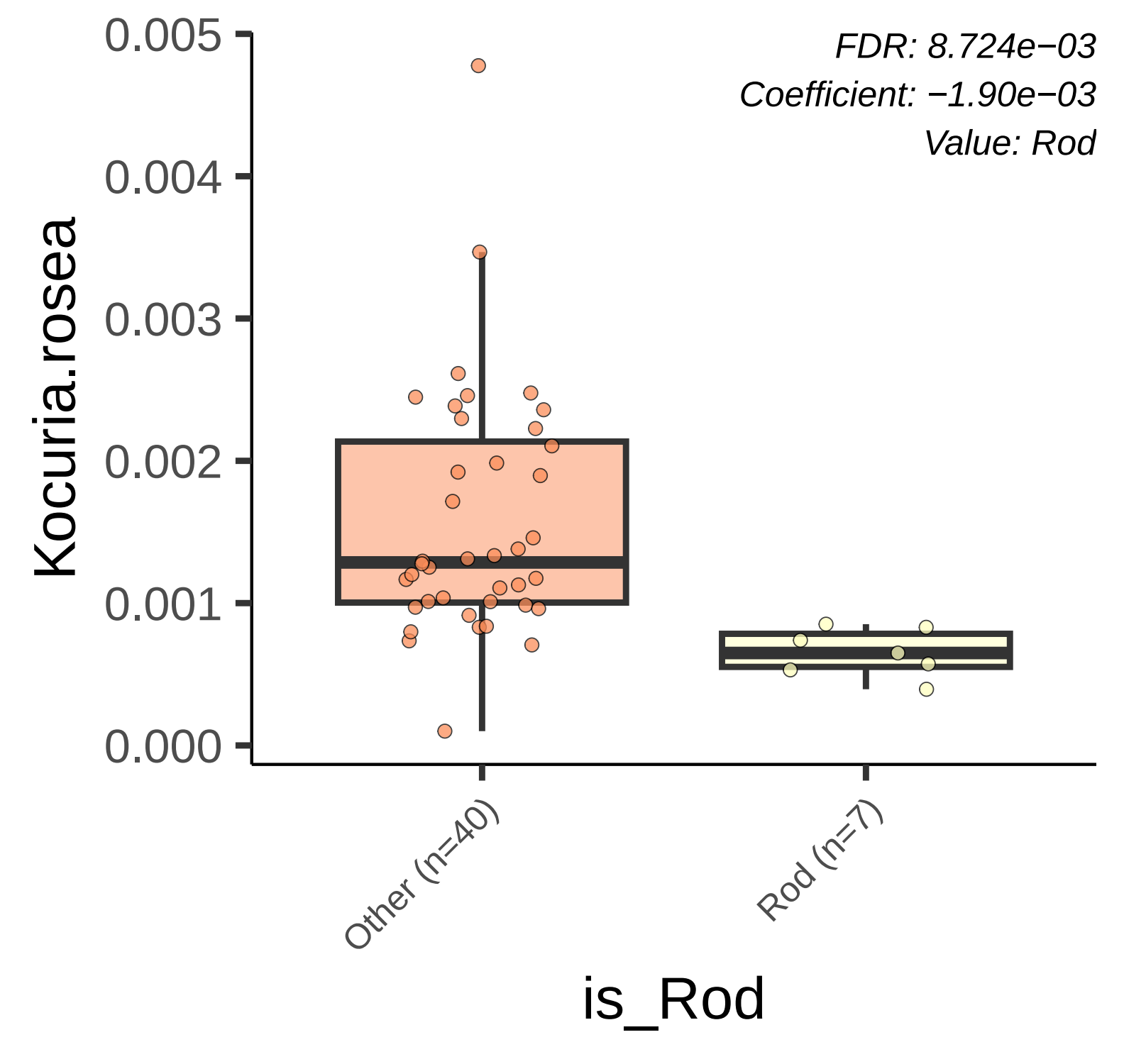

FDR: 7.754e−03
Coefficient: −8.38e−03
Value: Rod
Paracoccus.sp..AK26
0.02
0.01
0.00
Other (n=40)
Rod (n=7)
is_Rod

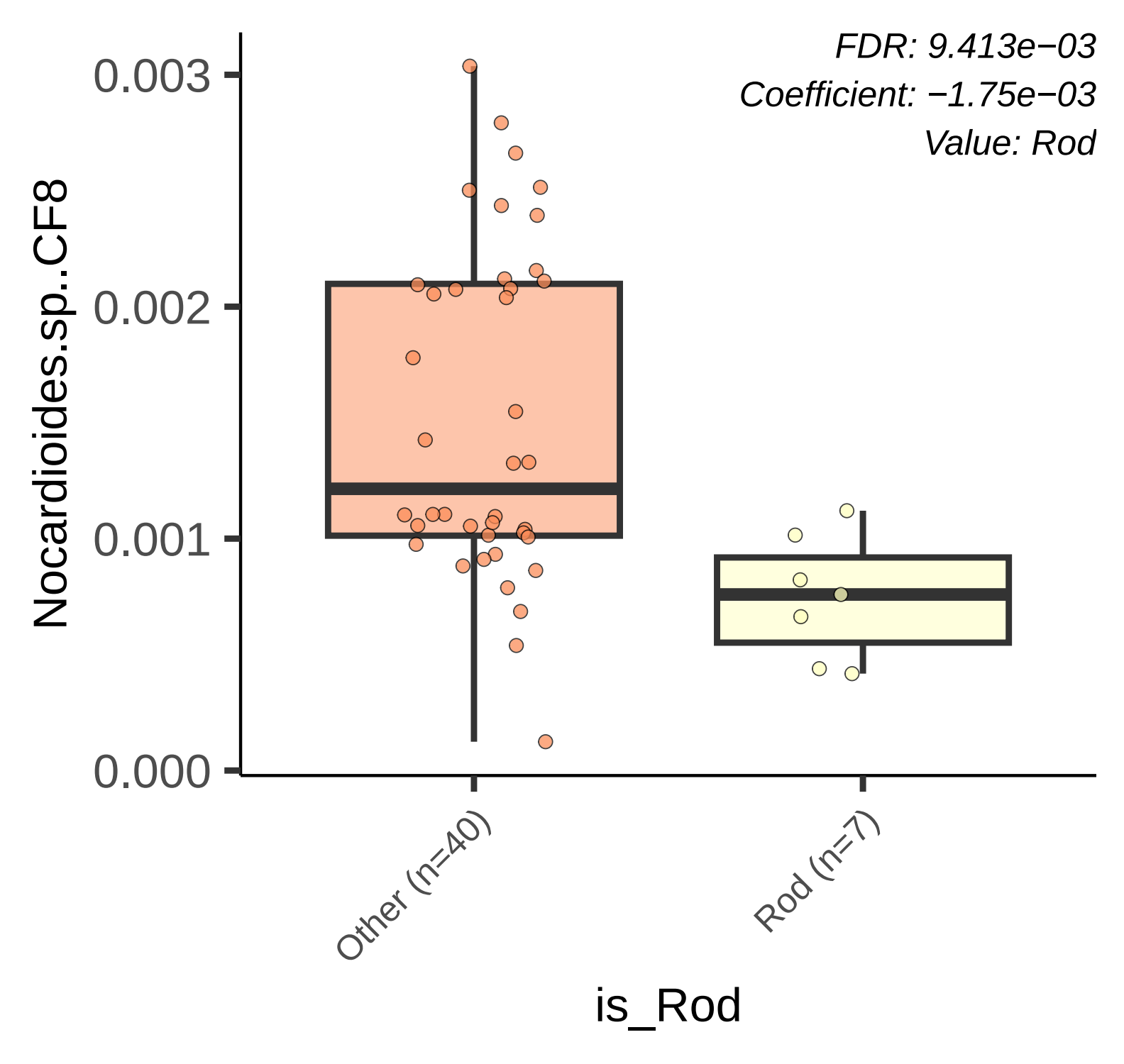

FDR: 7.754e−03
Coefficient: −2.72e−03
Value: Rod
Paracoccus.contaminans
0.006
0.004
0.002
0.000
Other (n=40)
Rod (n=7)
is_Rod

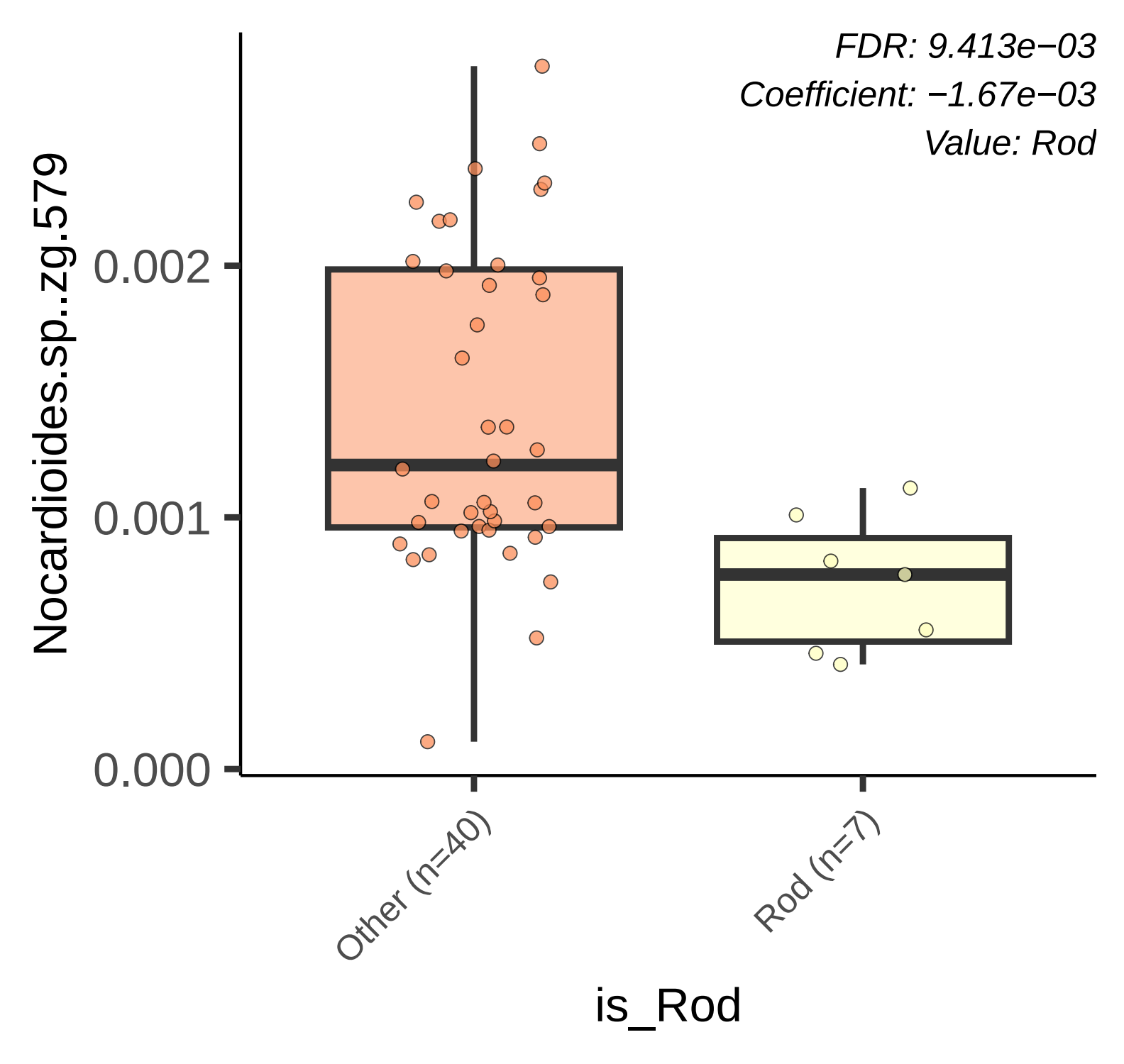

FDR: 7.796e−03
Coefficient: −1.95e−03
Value: Rod
Micropruina.glycogenica
0.003
0.002
0.001
0.000
Other (n=40)
Rod (n=7)
is_Rod

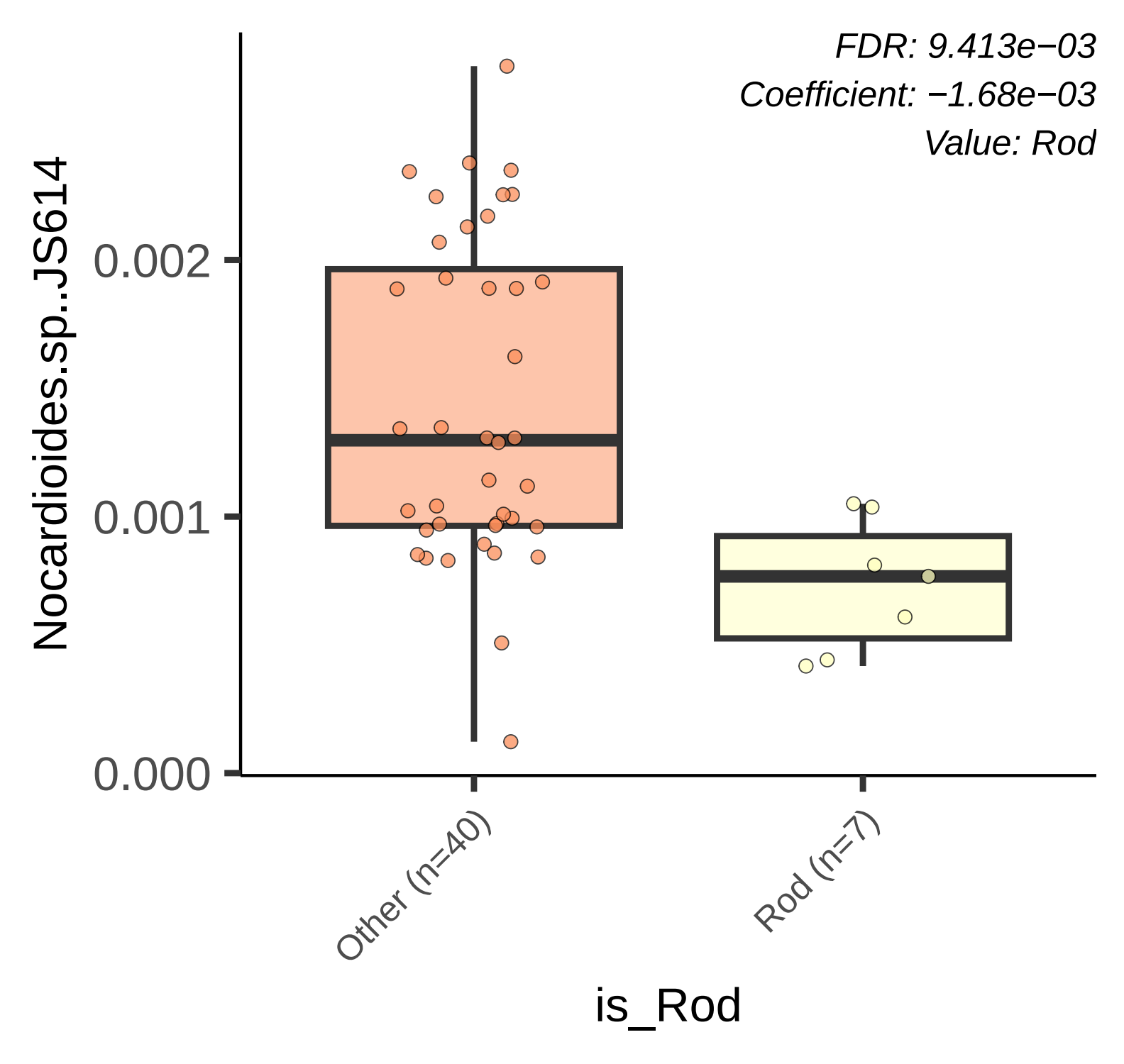

FDR: 7.796e−03
Coefficient: −1.34e−03
Value: Rod
Stenotrophomonas.acidaminiphila
0.004
0.003
0.002
0.001
Other (n=40)
Rod (n=7)
is_Rod

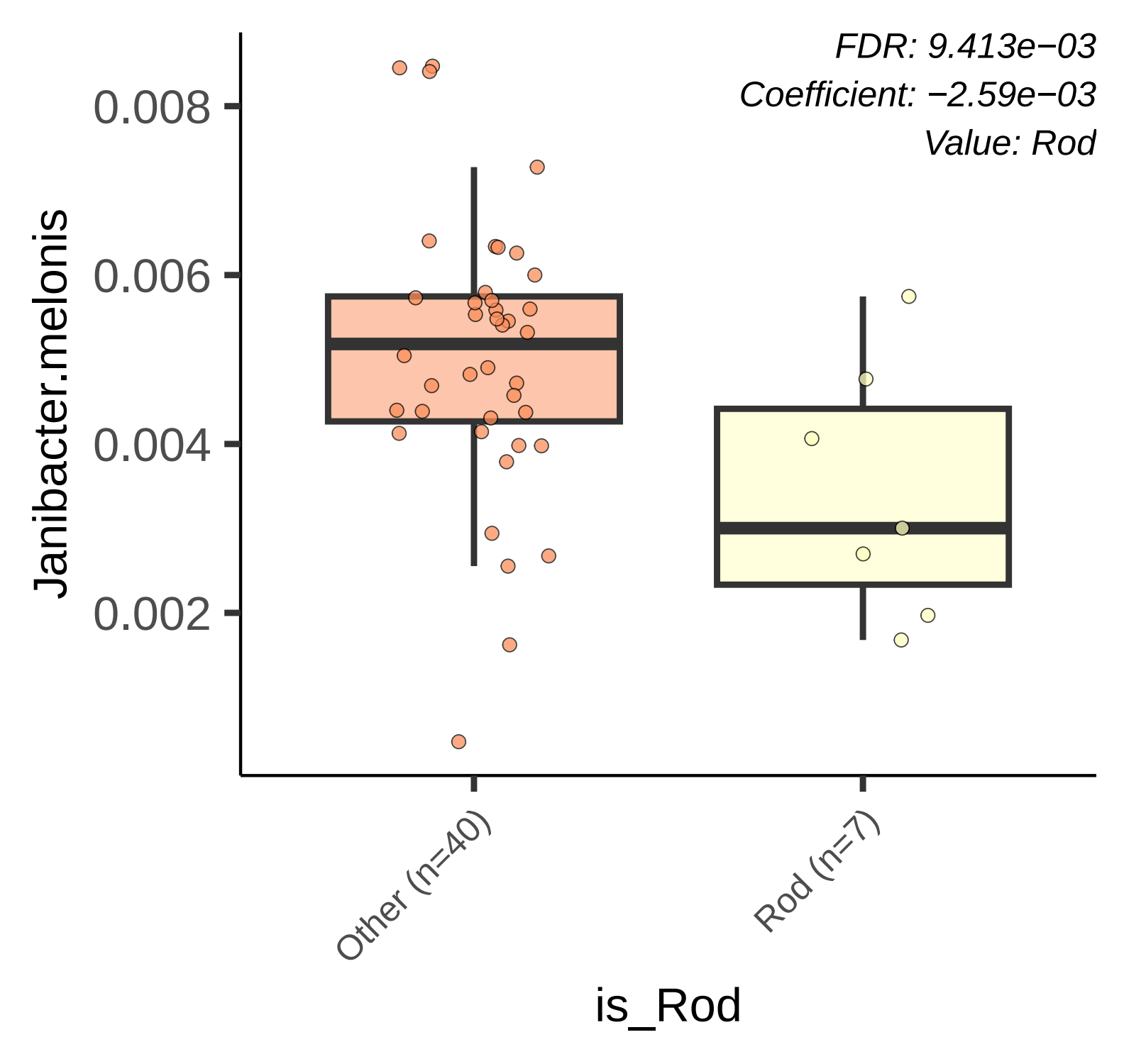

FDR: 8.472e−03
Coefficient: −1.88e−03
Value: Rod
Kocuria.turfanensis
0.003
0.002
0.001
0.000
Other (n=40)
Rod (n=7)
is_Rod

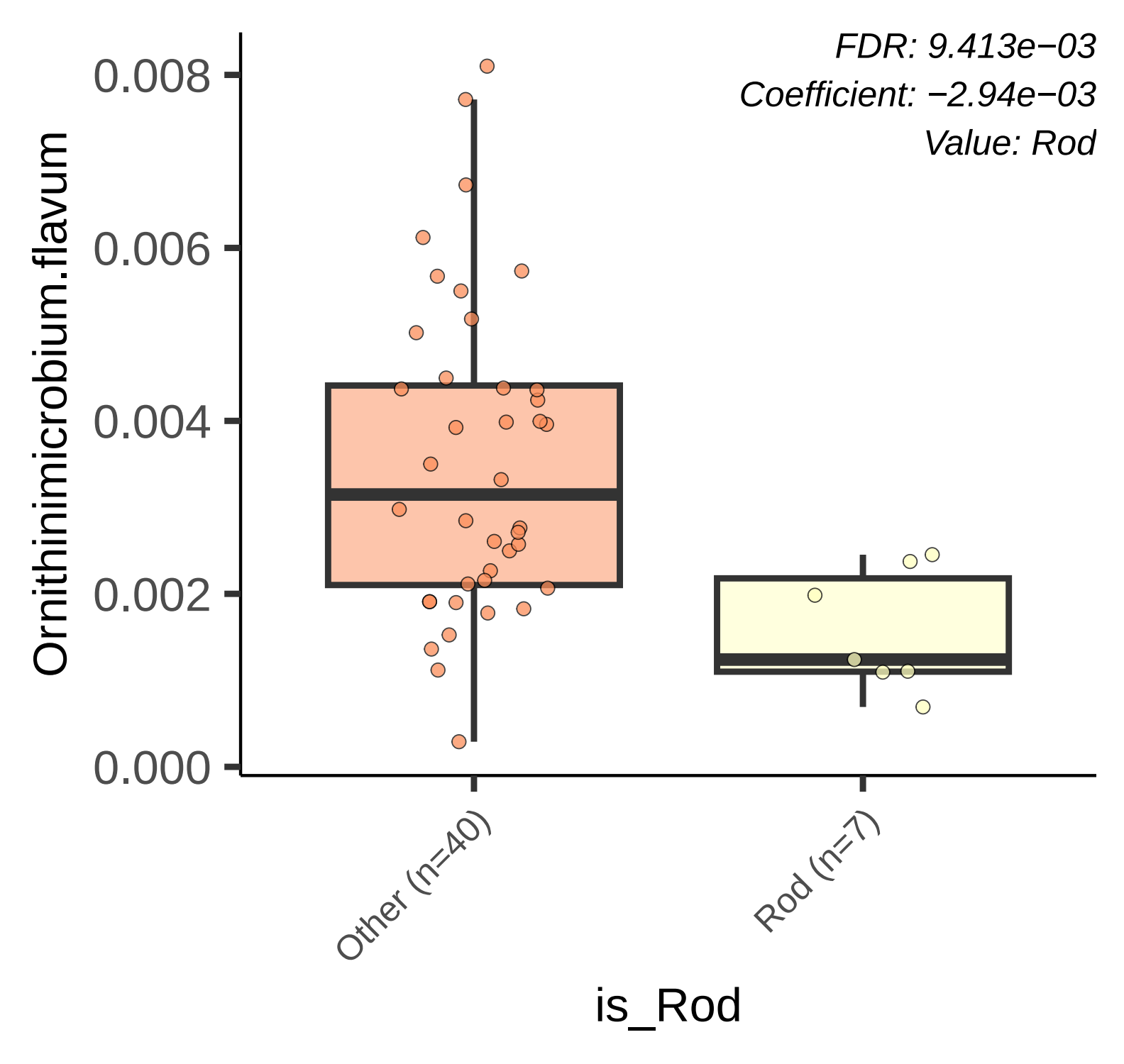

FDR: 8.724e−03
Coefficient: −1.90e−03
Value: Rod
Kocuria.rosea
0.005
0.004
0.003
0.002
0.001
0.000
Other (n=40)
Rod (n=7)
is_Rod

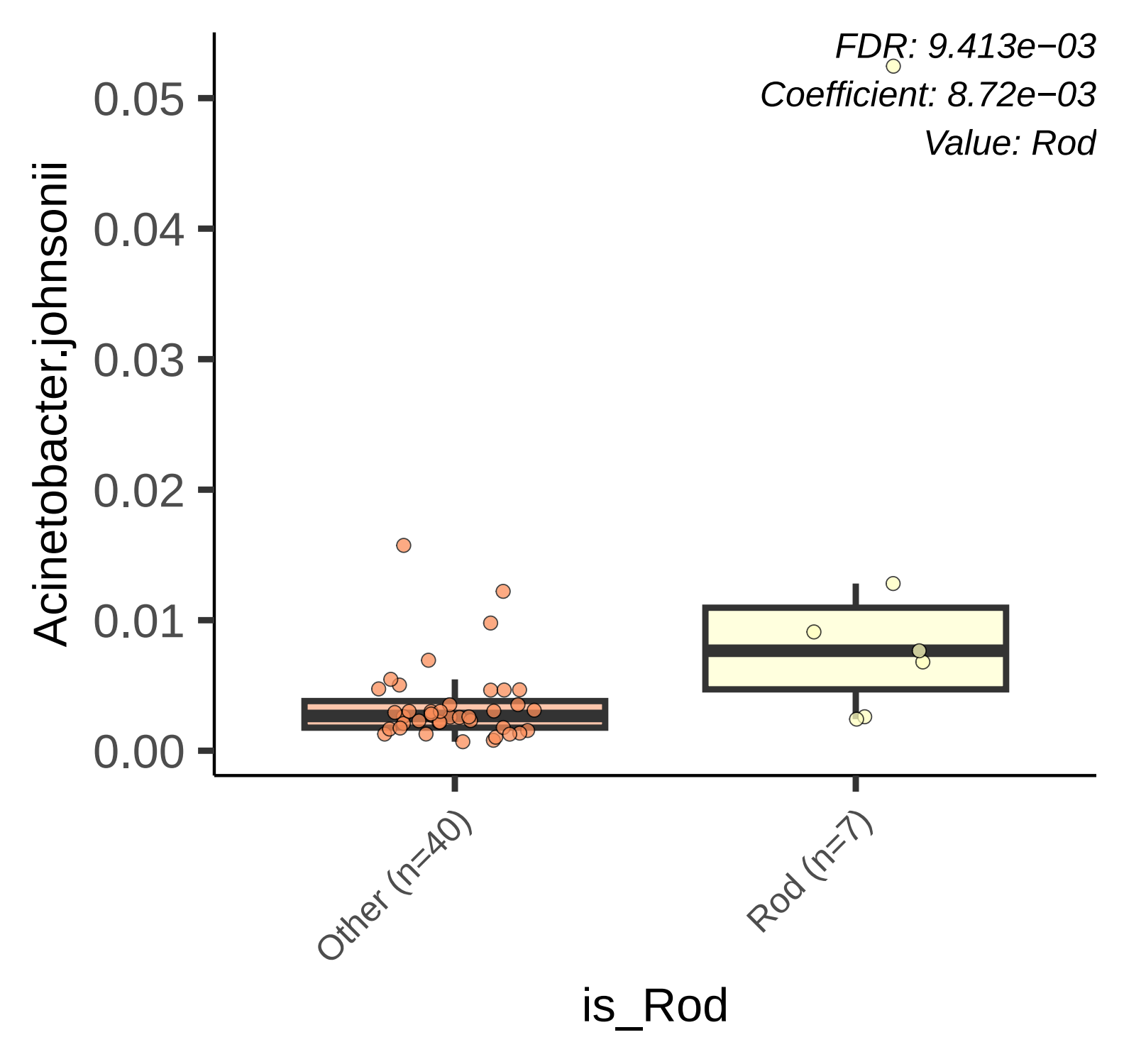

FDR: 9.413e−03
Coefficient: −1.75e−03
Value: Rod
Nocardioides.sp..CF8
0.003
0.002
0.001
0.000
Other (n=40)
Rod (n=7)
is_Rod

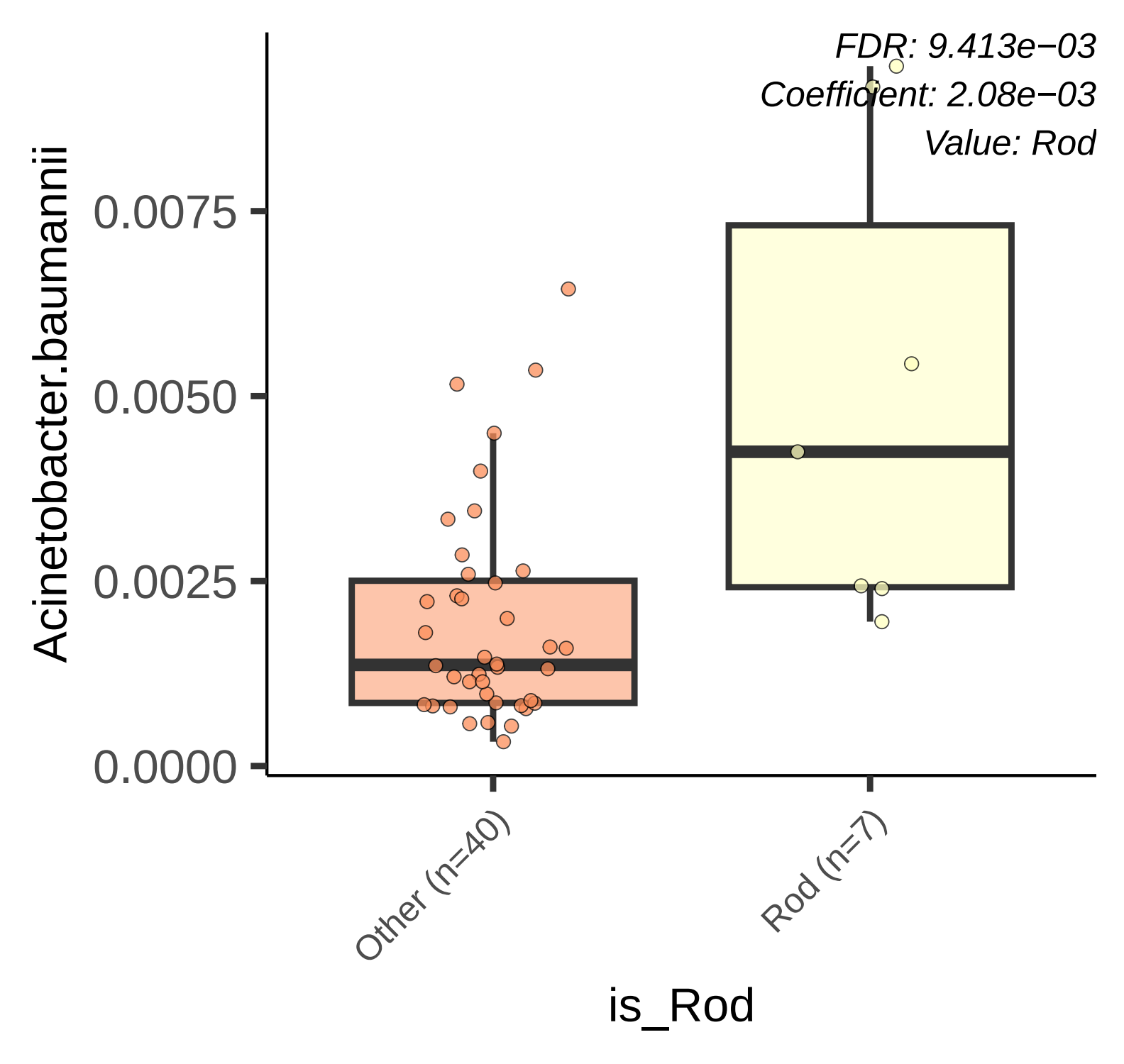

FDR: 9.413e−03
Coefficient: −1.67e−03
Value: Rod
Nocardioides.sp..zg.579
0.002
0.001
0.000
Other (n=40)
Rod (n=7)
is_Rod

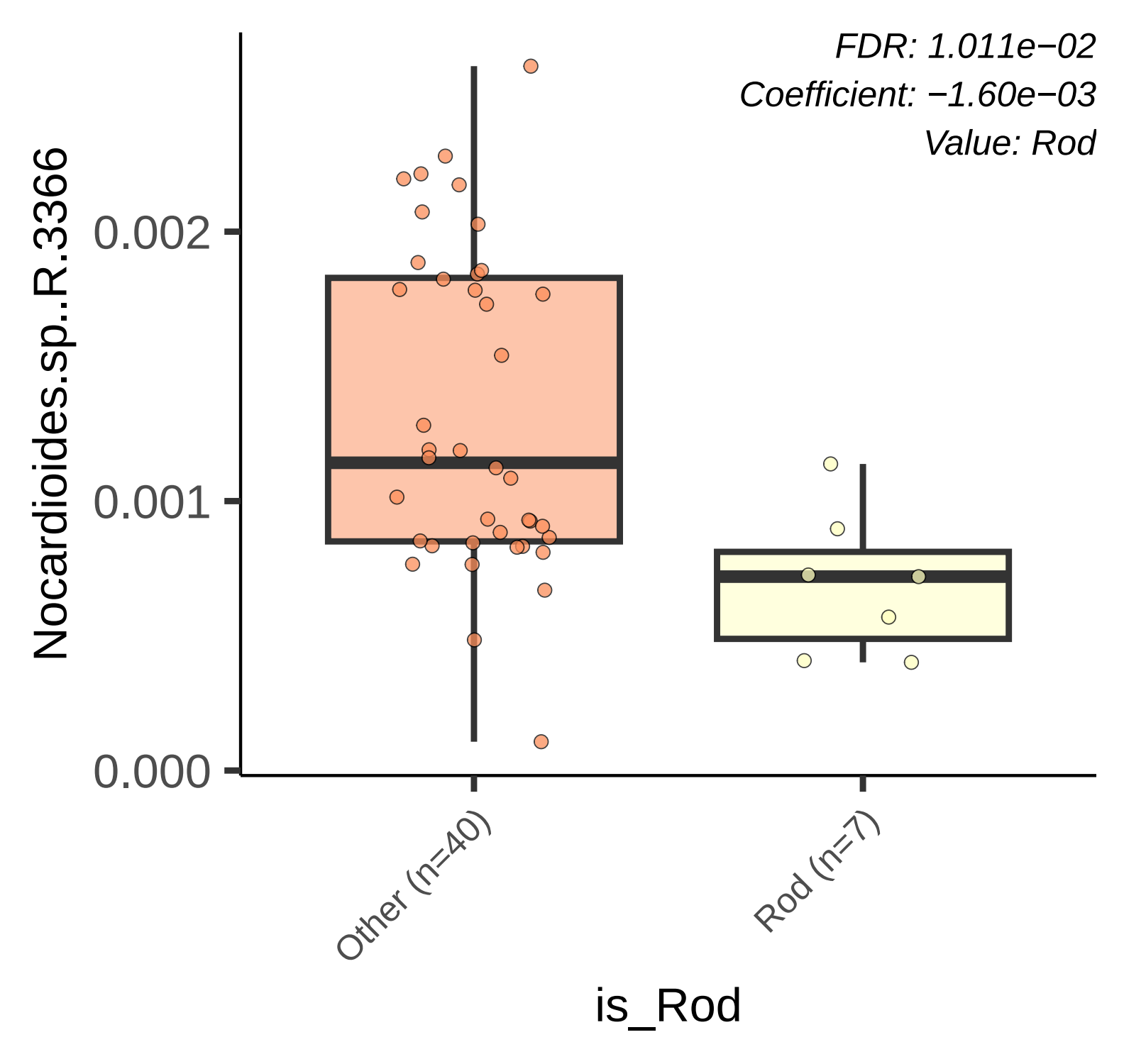

FDR: 9.413e−03
Coefficient: −1.68e−03
Value: Rod
Nocardioides.sp..JS614
0.002
0.001
0.000
Other (n=40)
Rod (n=7)
is_Rod

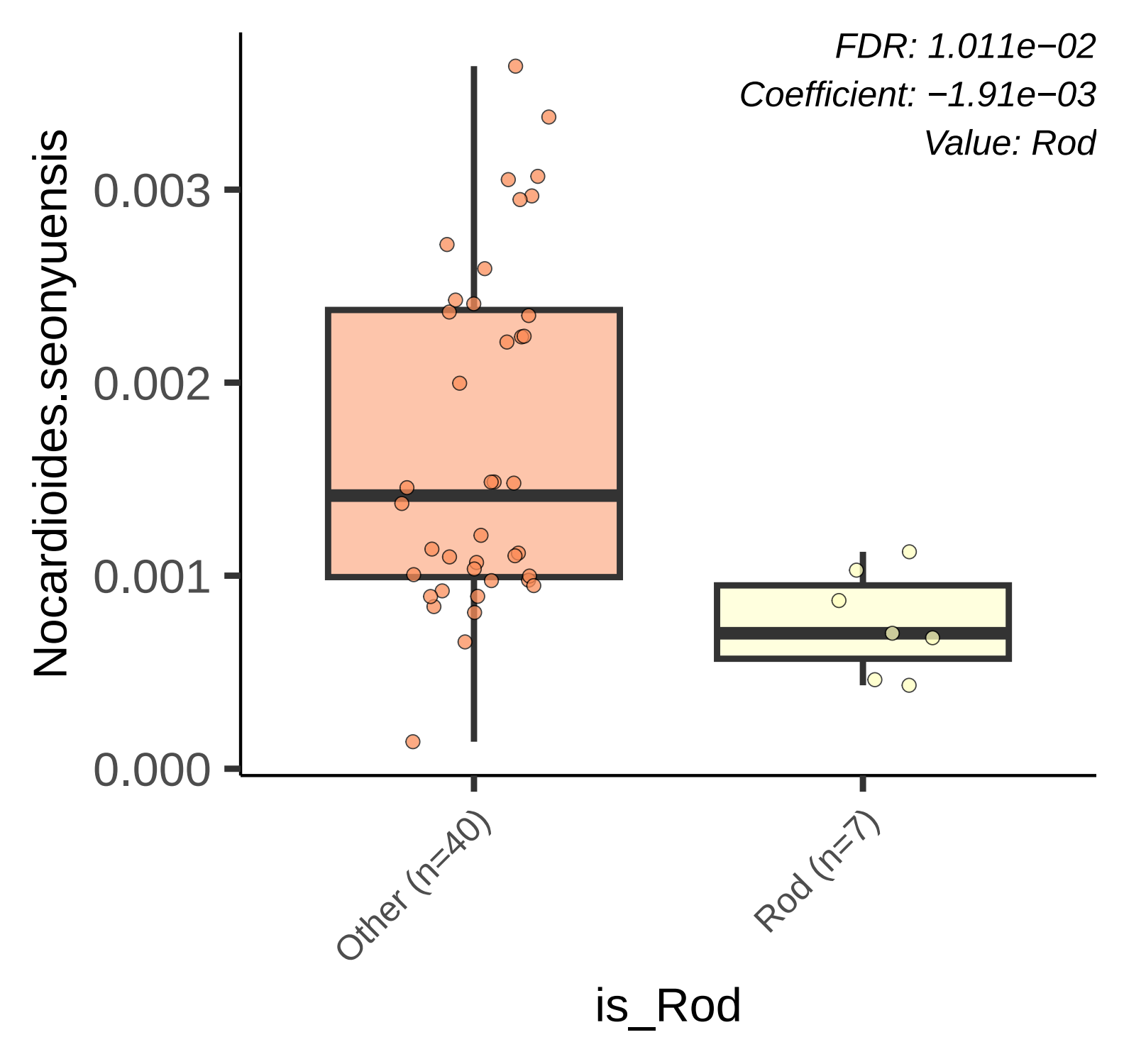

FDR: 9.413e−03
Coefficient: −2.59e−03
Value: Rod
0.008
0.006
0.004
0.002
Janibacter.melonis
Other (n=40)
Rod (n=7)
is_Rod

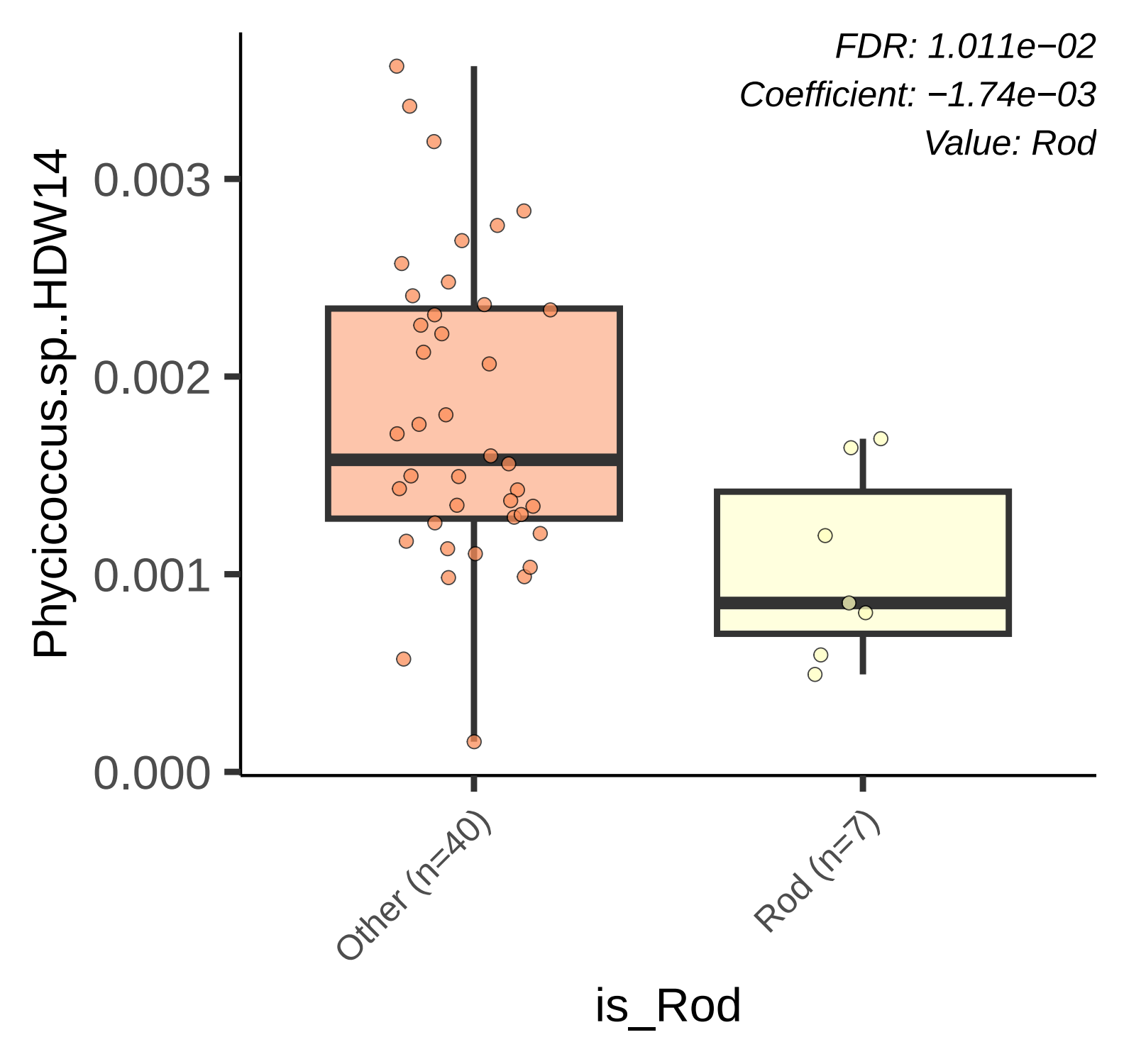

FDR: 9.413e−03
Coefficient: −2.94e−03
Value: Rod
Ornithinimicrobium.flavum
0.008
0.006
0.004
0.002
0.000
Other (n=40)
Rod (n=7)
is_Rod

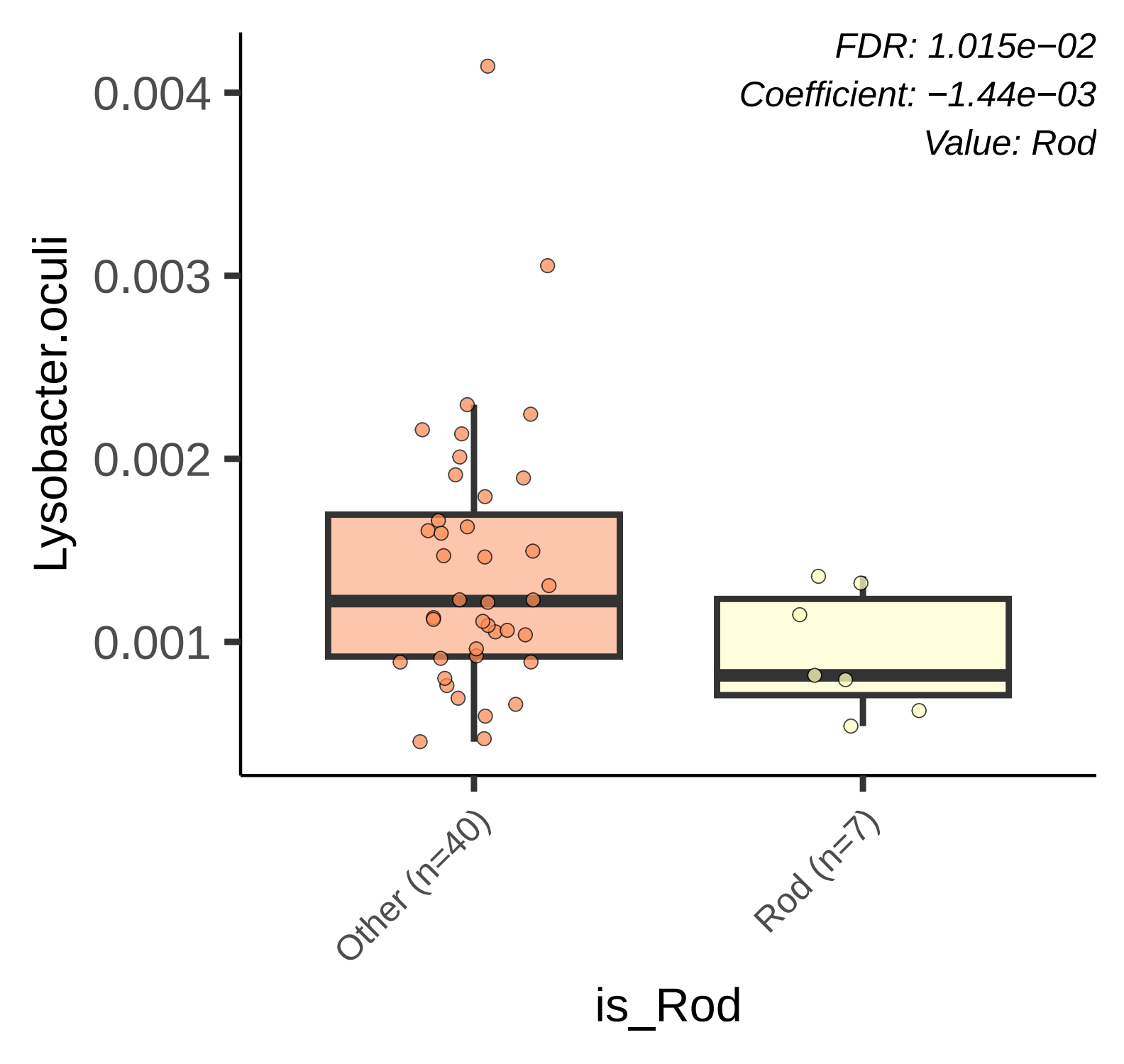

FDR: 9.413e−03
Coefficient: 8.72e−03
Value: Rod
Acinetobacter.johnsonii
0.05
0.04
0.03
0.02
0.01
0.00
Other (n=40)
Rod (n=7)
is_Rod

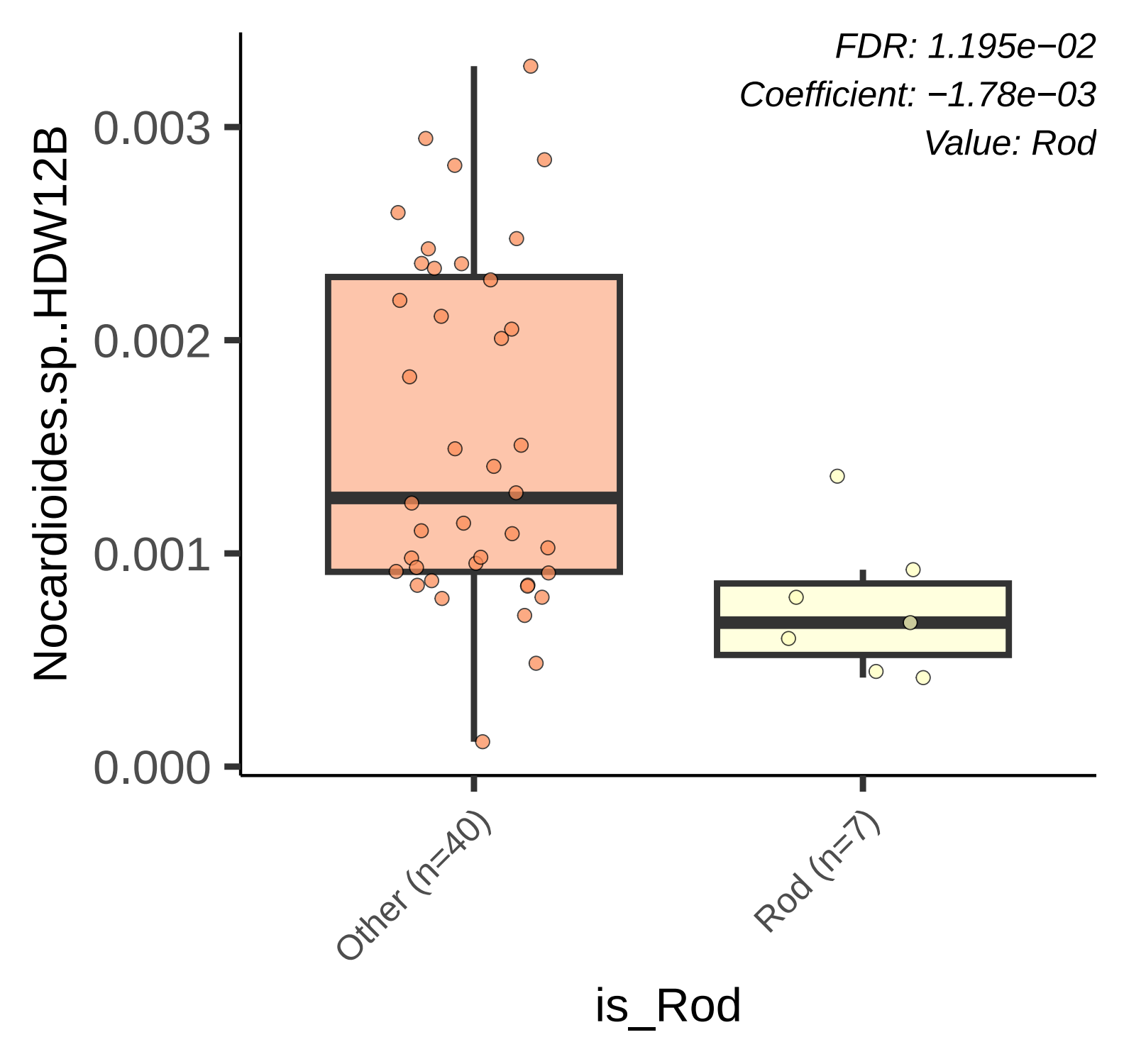

FDR: 9.413e−03
Coefficient: 2.08e−03
Value: Rod
Acinetobacter.baumannii
0.0075
0.0050
0.0025
0.0000
Other (n=40)
Rod (n=7)
is_Rod

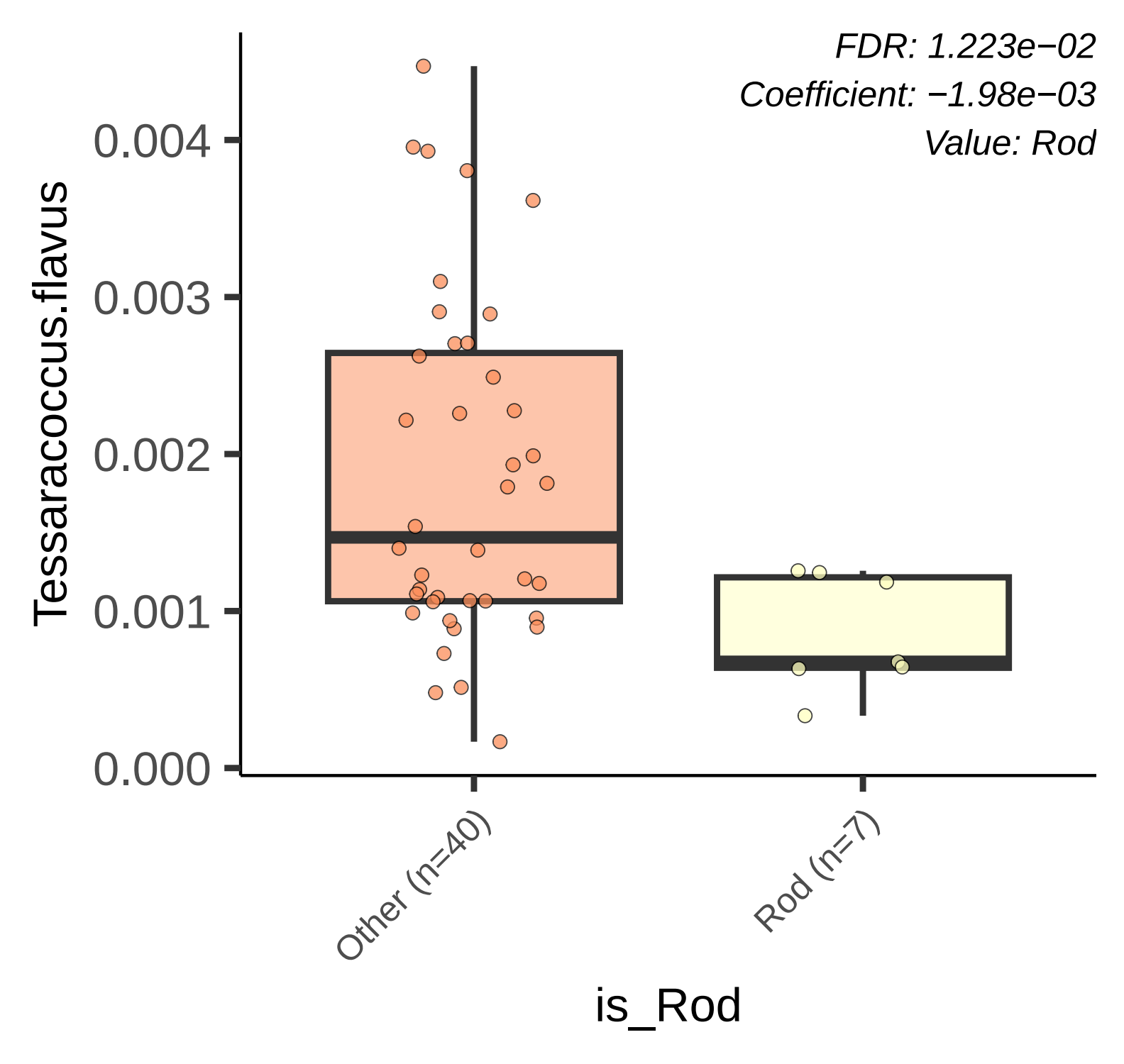

FDR: 1.011e−02
Coefficient: −1.60e−03
Value: Rod
Nocardioides.sp..R.3366
0.002
0.001
0.000
Other (n=40)
Rod (n=7)
is_Rod

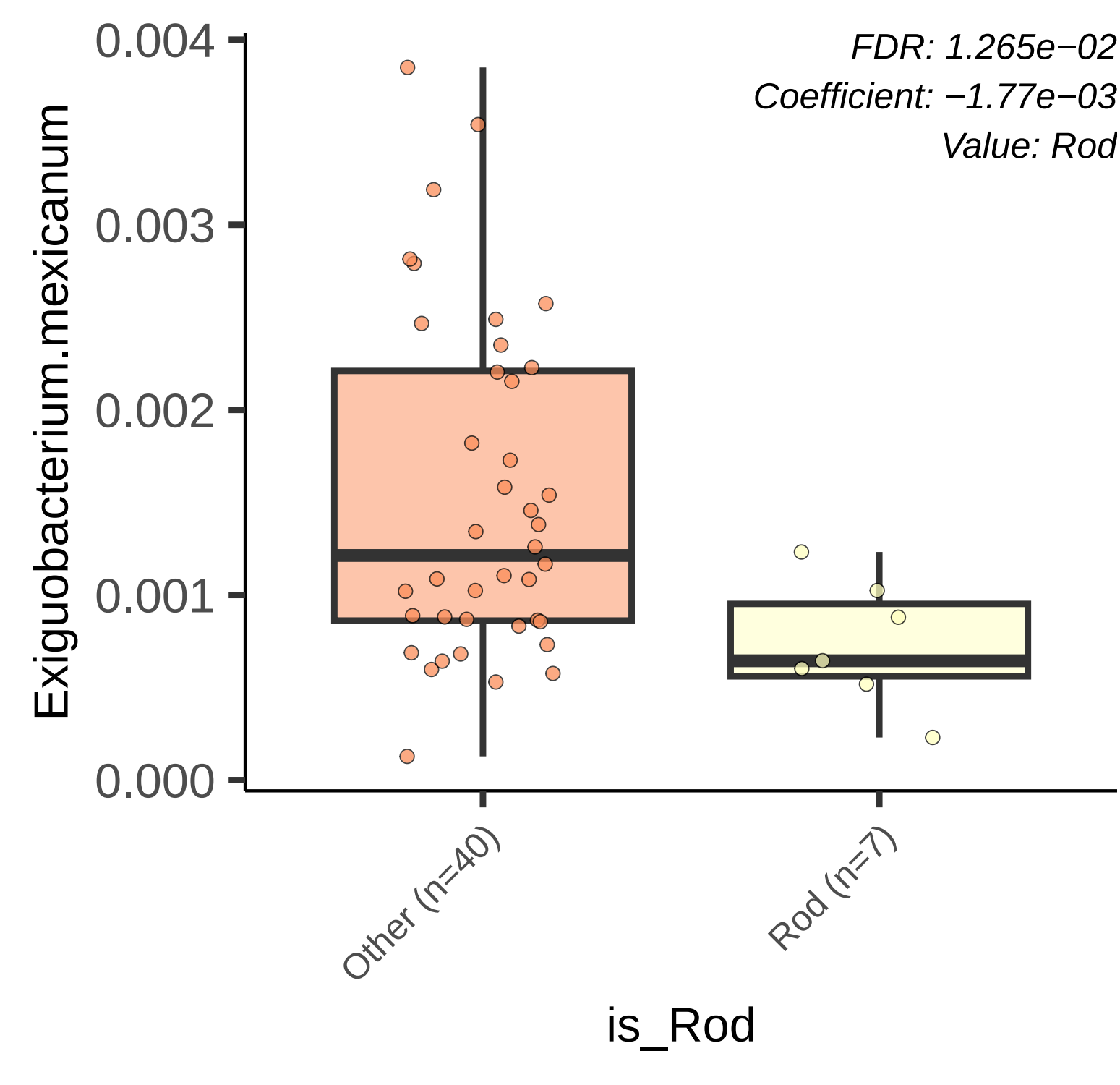

FDR: 1.011e−02
Coefficient: −1.91e−03
Value: Rod
Nocardioides.seonyuensis
0.003
0.002
0.001
0.000
Other (n=40)
Rod (n=7)
is_Rod

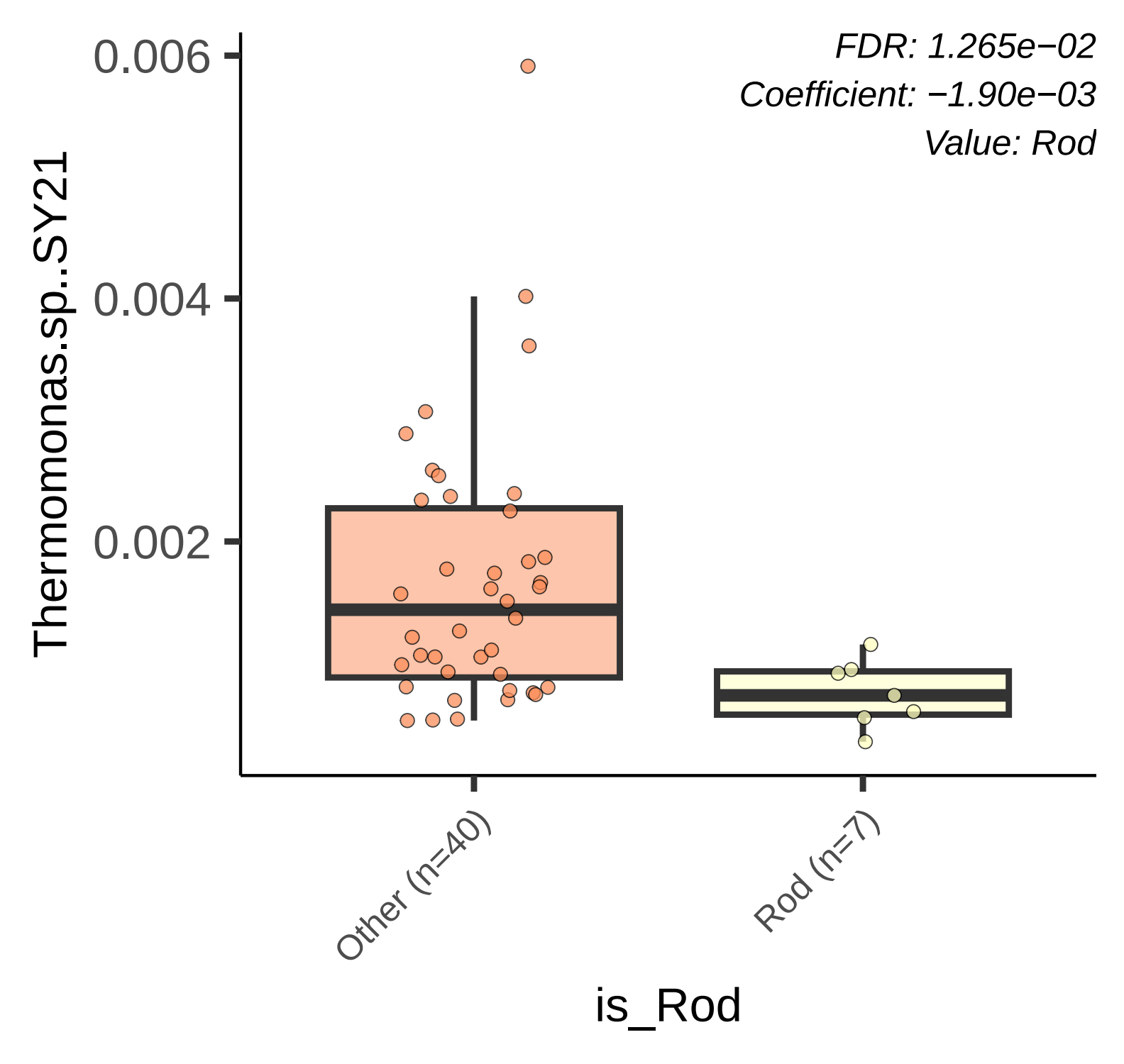

FDR: 1.011e−02
Coefficient: −1.74e−03
Value: Rod
Phycicoccus.sp..HDW14
0.003
0.002
0.001
0.000
Other (n=40)
Rod (n=7)
is_Rod

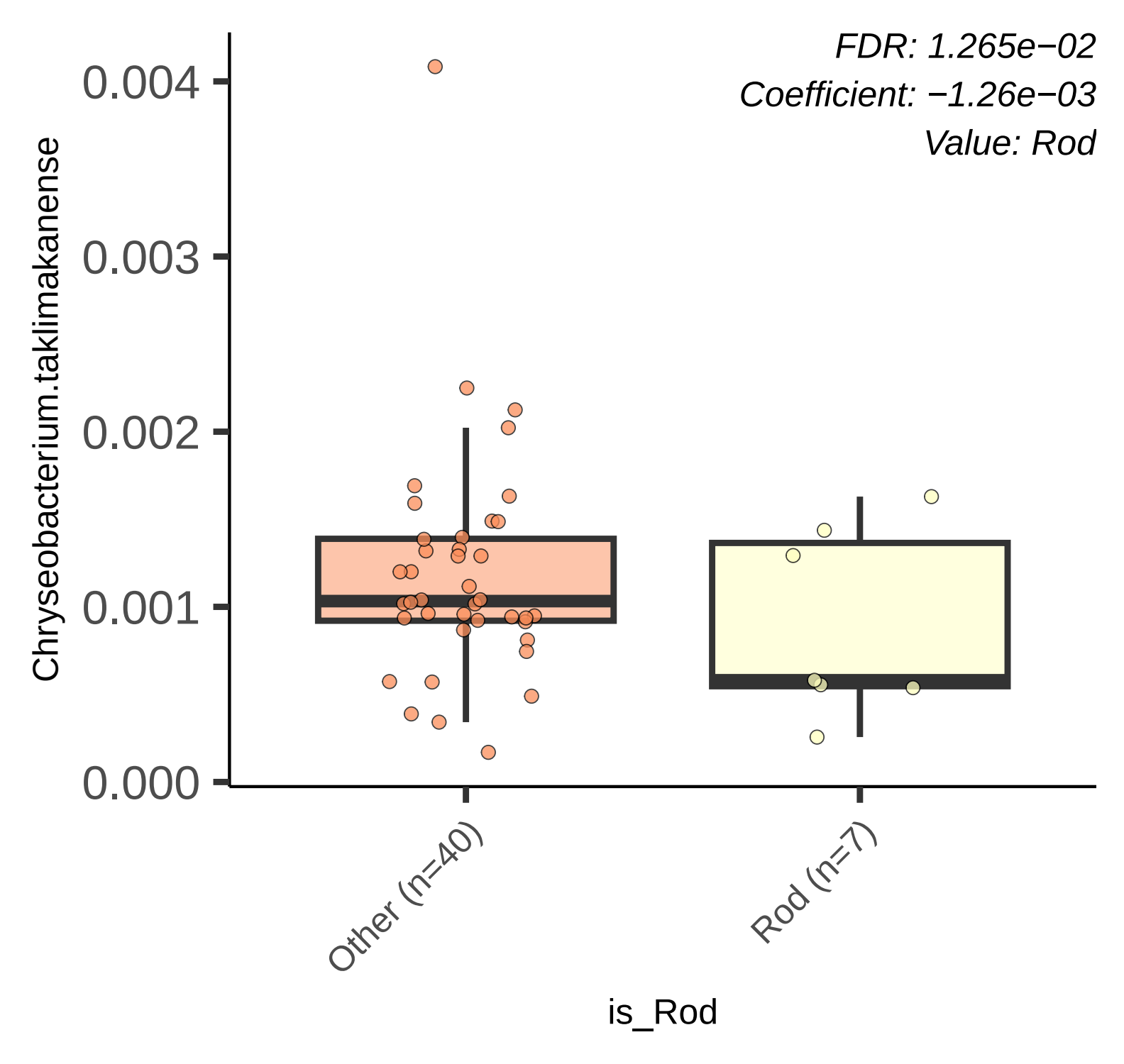

FDR: 1.015e−02
Coefficient: −1.44e−03
Value: Rod
0.004
0.003
0.002
0.001
Lysobacter.oculi
Other (n=40)
Rod (n=7)
is_Rod

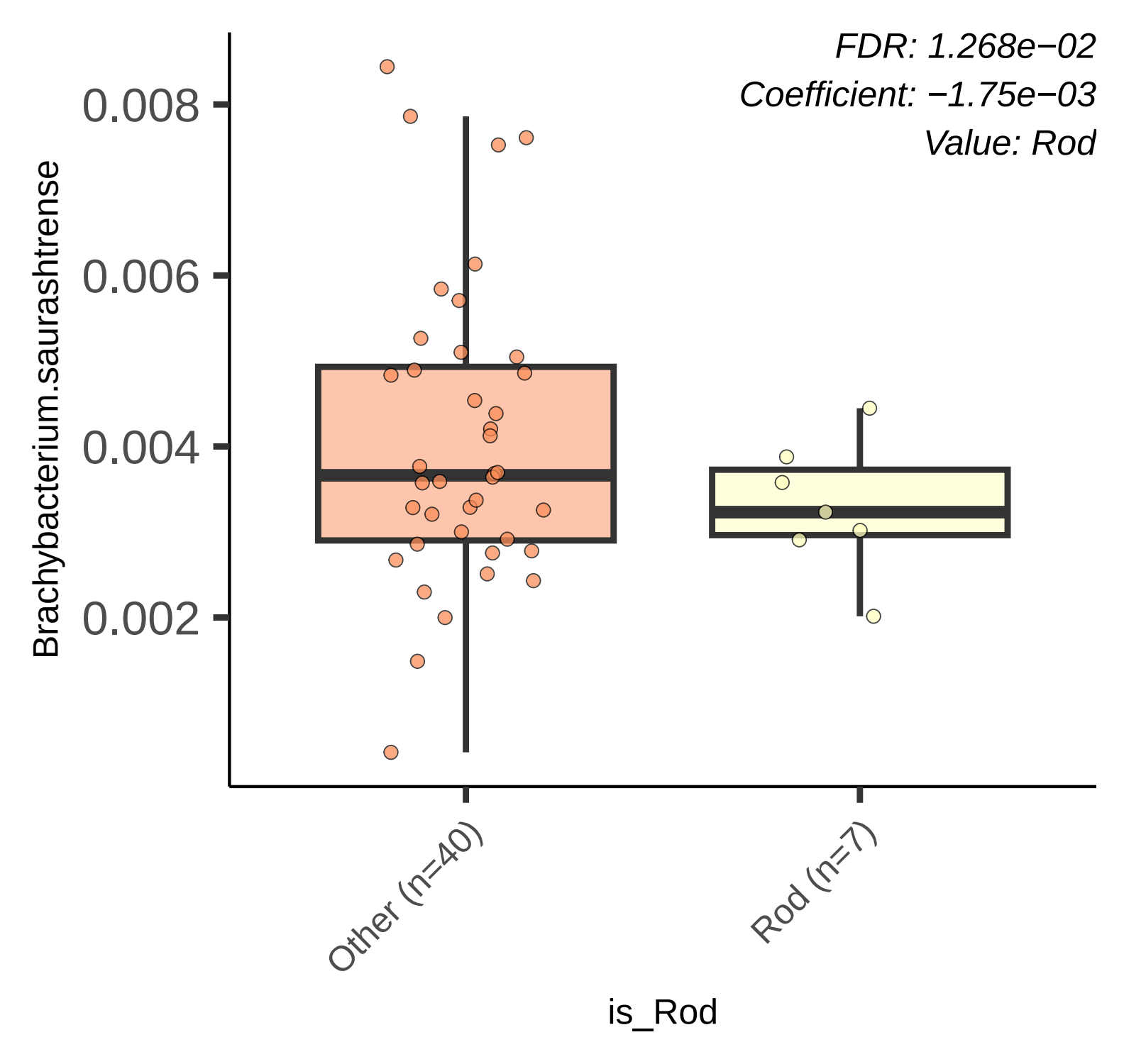

FDR: 1.195e−02
Coefficient: −1.78e−03
Value: Rod
Nocardioides.sp..HDW12B
0.003
0.002
0.001
0.000
Other (n=40)
Rod (n=7)
is_Rod

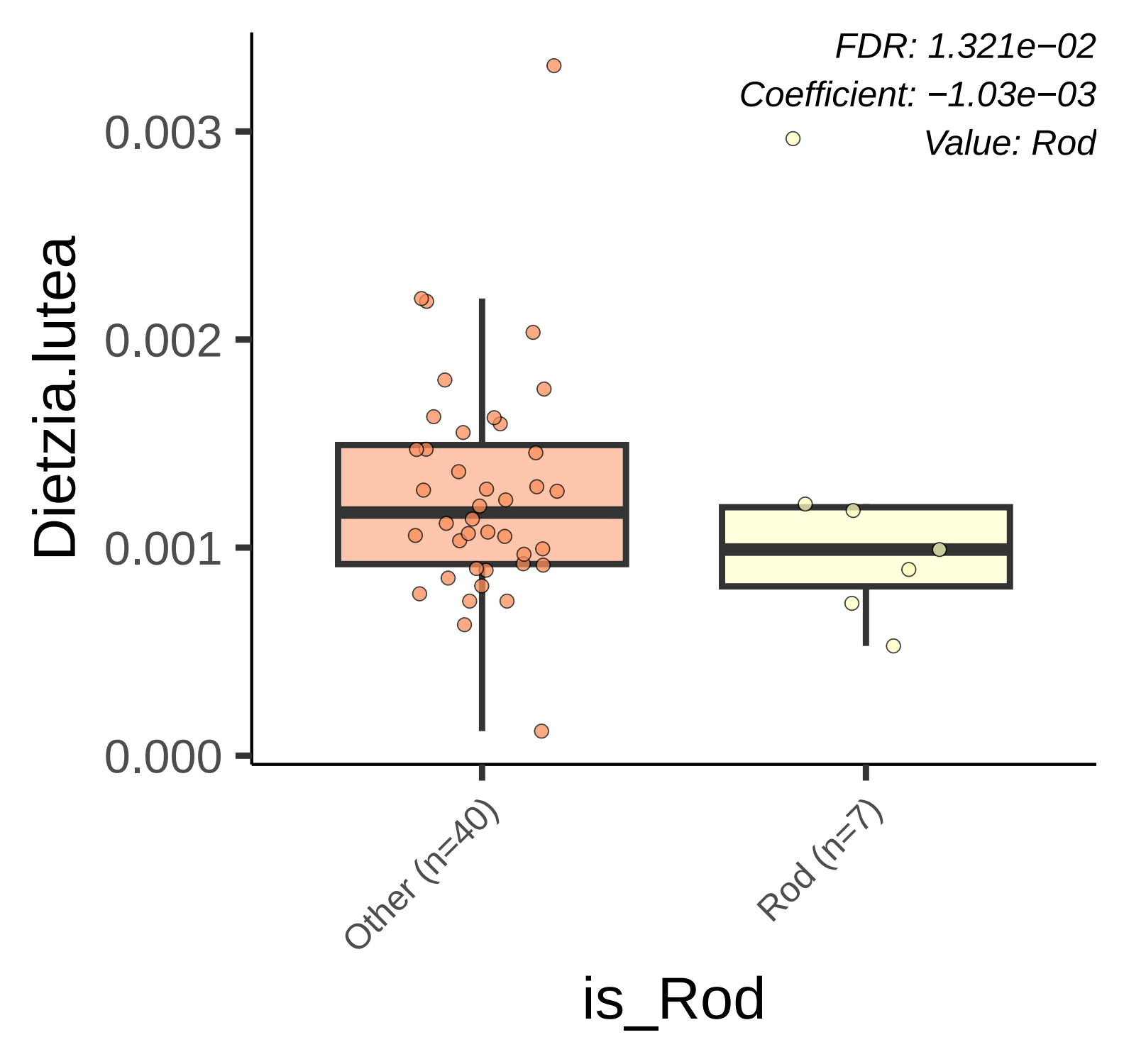

FDR: 1.223e−02
Coefficient: −1.98e−03
Value: Rod
Tessaracoccus.flavus
0.004
0.003
0.002
0.001
0.000
Other (n=40)
Rod (n=7)
is_Rod

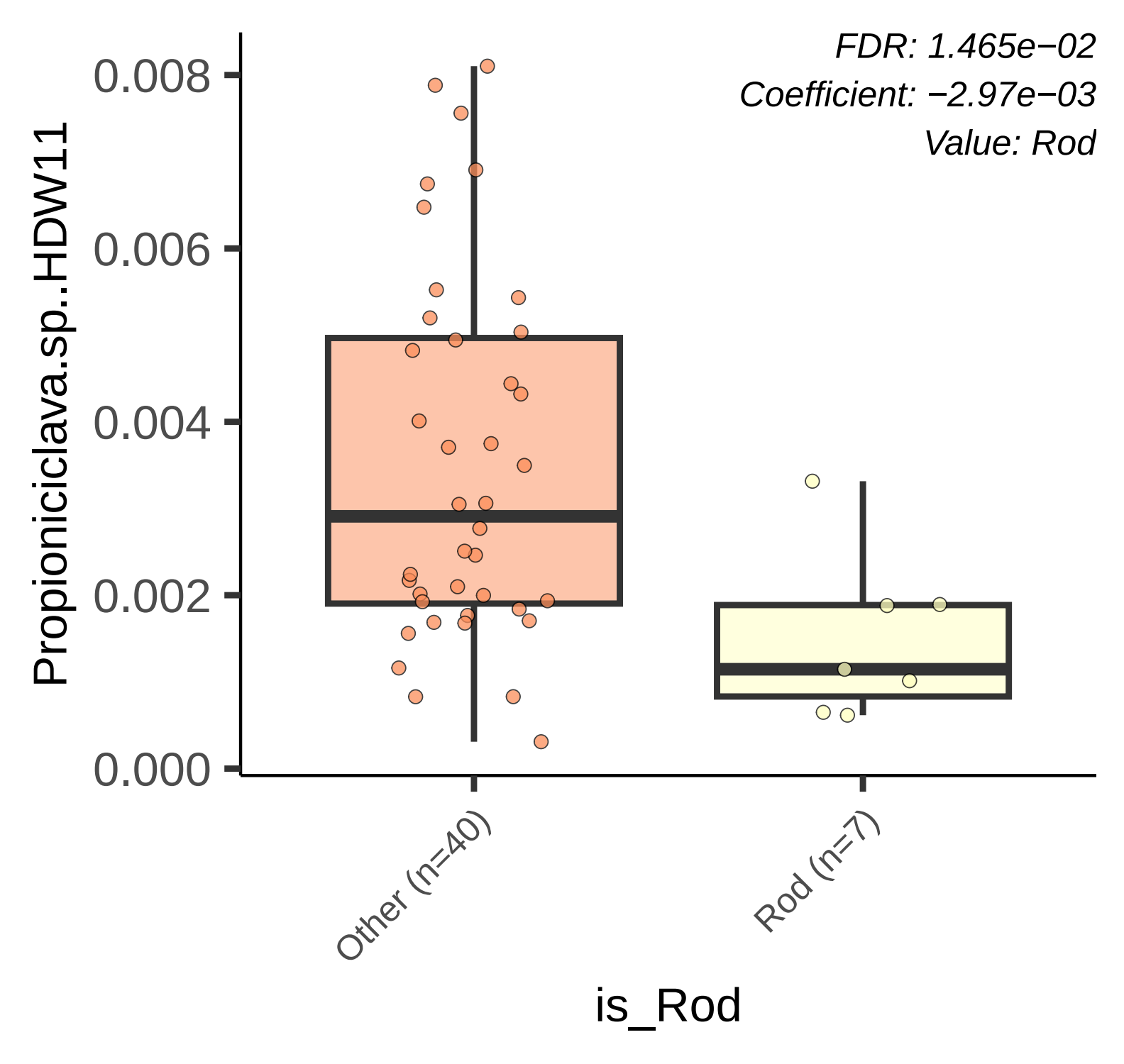

FDR: 1.265e−02
Coefficient: −1.77e−03
Value: Rod
Exiguobacterium.mexicanum
0.004
0.003
0.002
0.001
0.000
Other (n=40)
Rod (n=7)
is_Rod

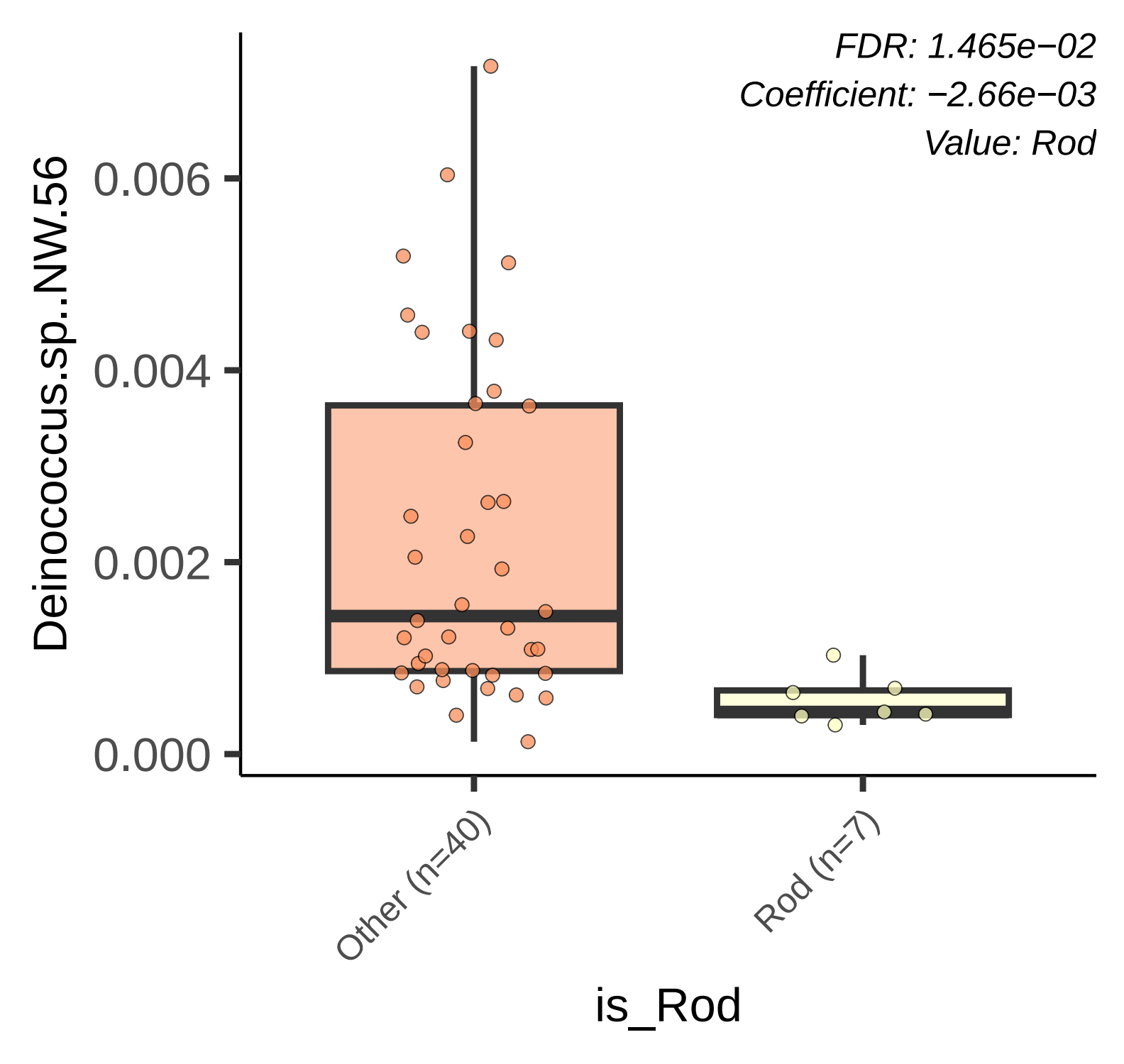

FDR: 1.265e−02
Coefficient: −1.90e−03
Value: Rod
0.006
0.004
0.002
Thermomonas.sp..SY21
Other (n=40)
Rod (n=7)
is_Rod

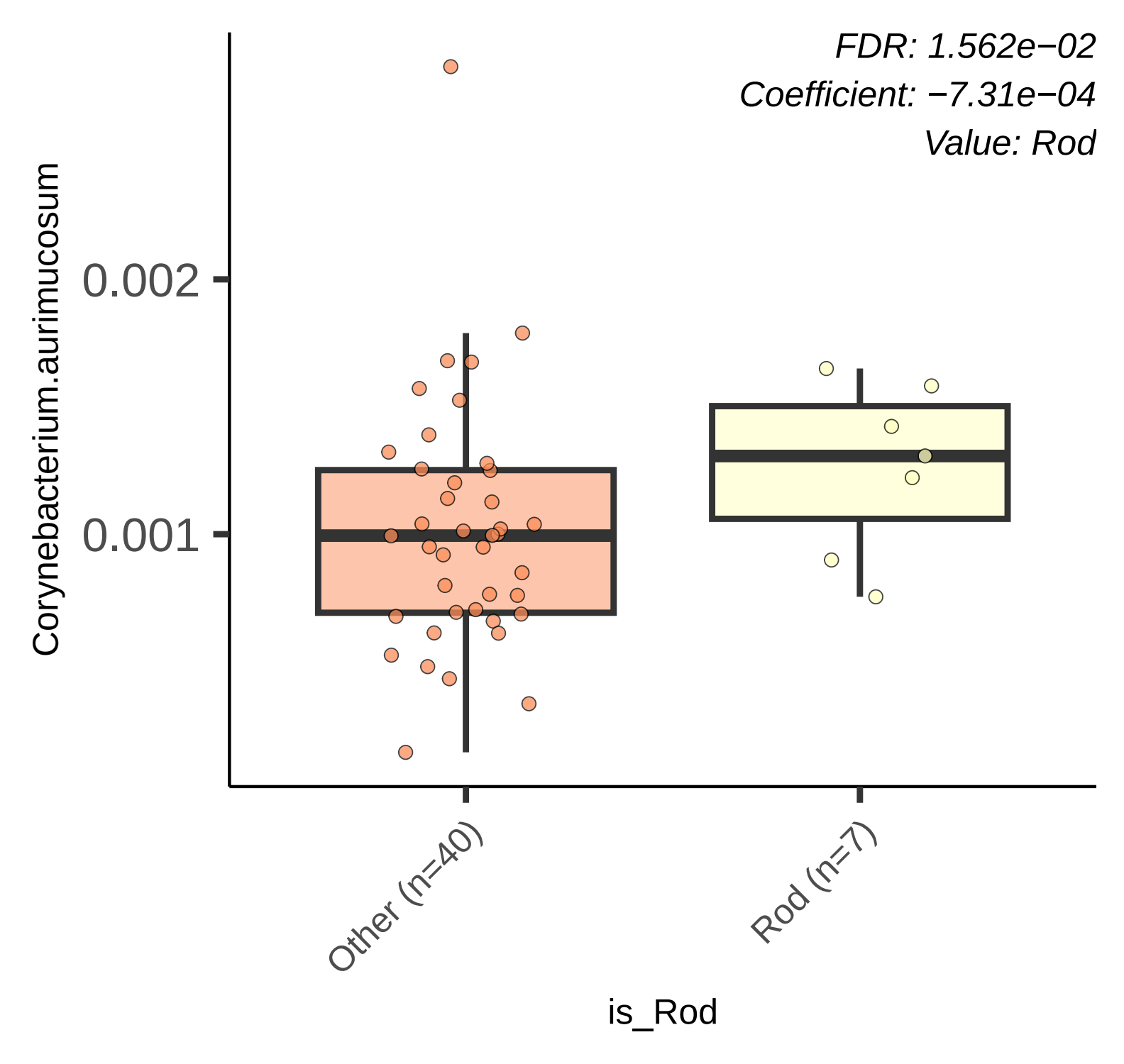

FDR: 1.265e−02
Coefficient: −1.26e−03
Value: Rod
Chryseobacterium.taklimakanense
0.004
0.003
0.002
0.001
0.000
Other (n=40)
Rod (n=7)
is_Rod

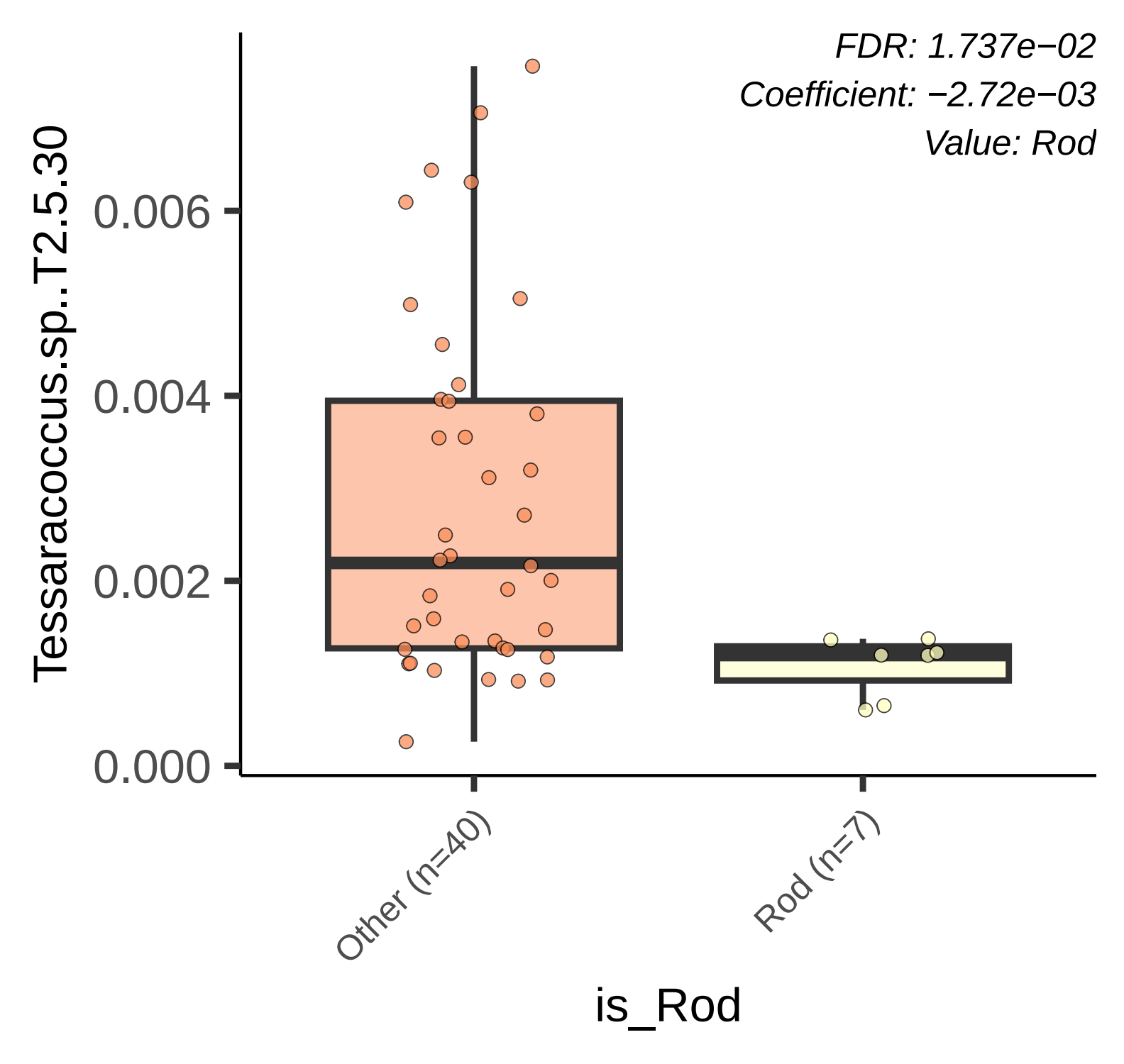

FDR: 1.268e−02
Coefficient: −1.75e−03
Value: Rod
0.008
0.006
0.004
0.002
Brachybacterium.saurashtrense
Other (n=40)
Rod (n=7)
is_Rod

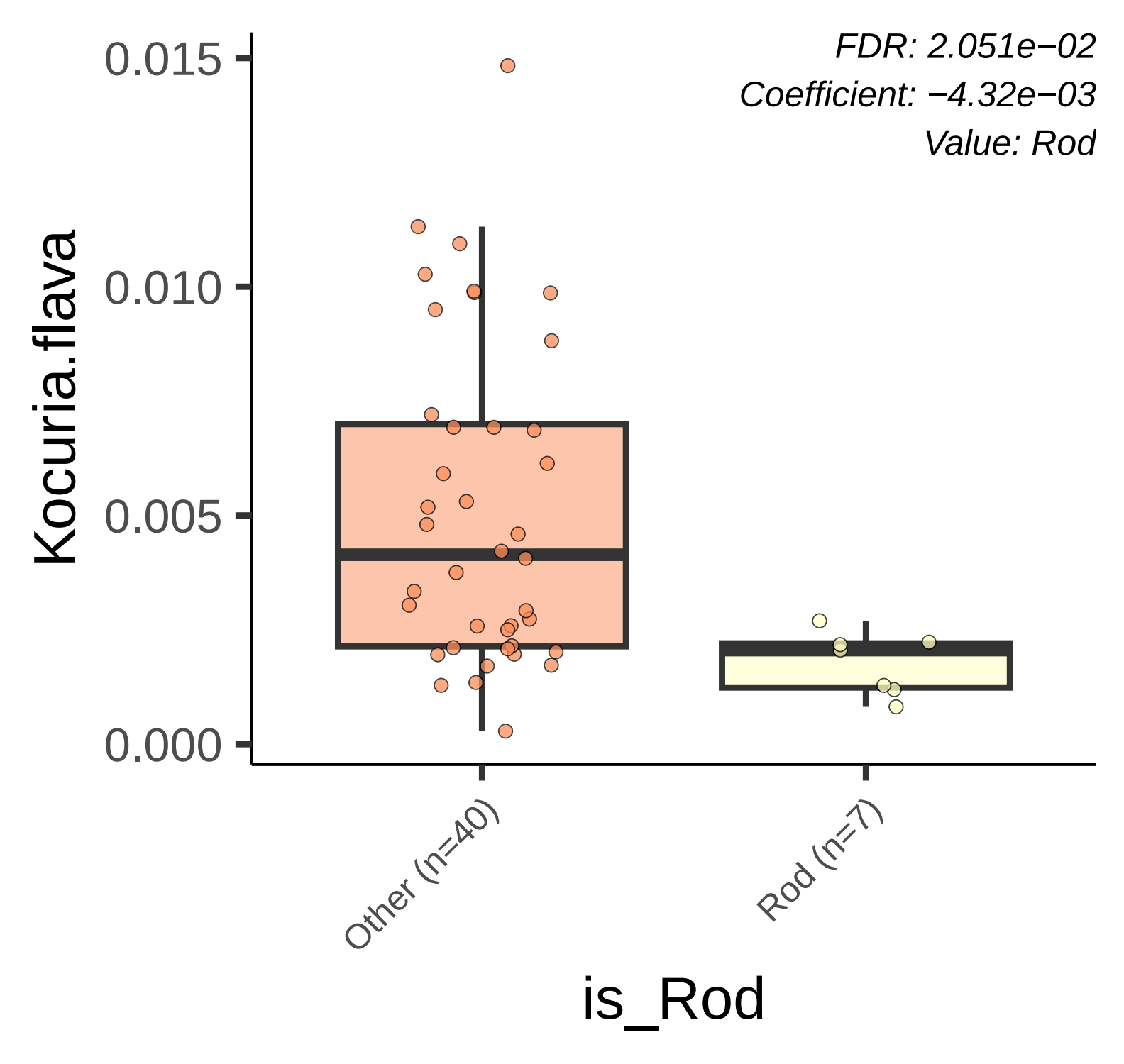

FDR: 1.321e−02
Coefficient: −1.03e−03
Value: Rod
Dietzia.lutea
0.003
0.002
0.001
0.000
Other (n=40)
Rod (n=7)
is_Rod

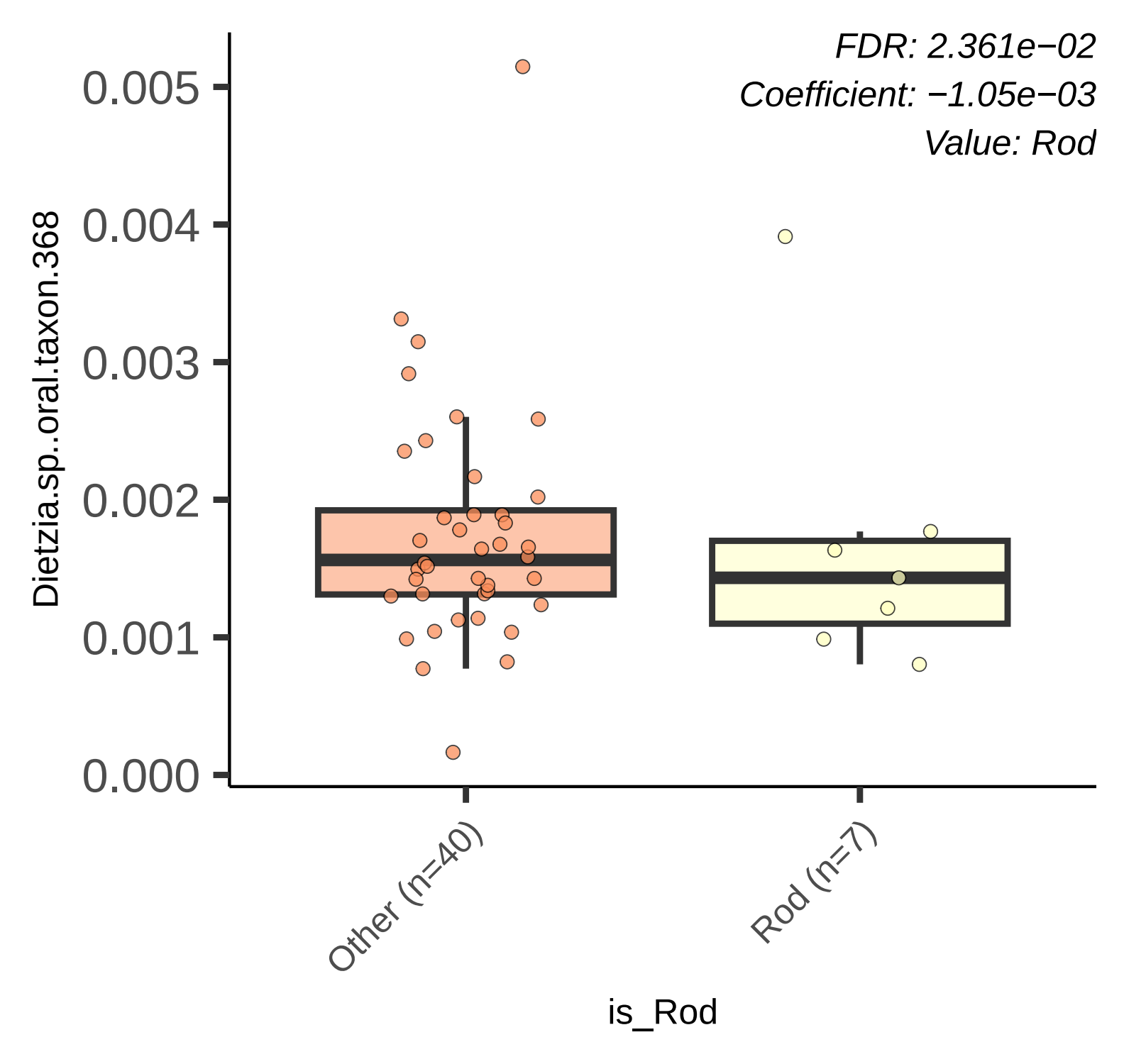

FDR: 1.465e−02
Coefficient: −2.97e−03
Value: Rod
Propioniciclava.sp..HDW11
0.008
0.006
0.004
0.002
0.000
Other (n=40)
Rod (n=7)
is_Rod

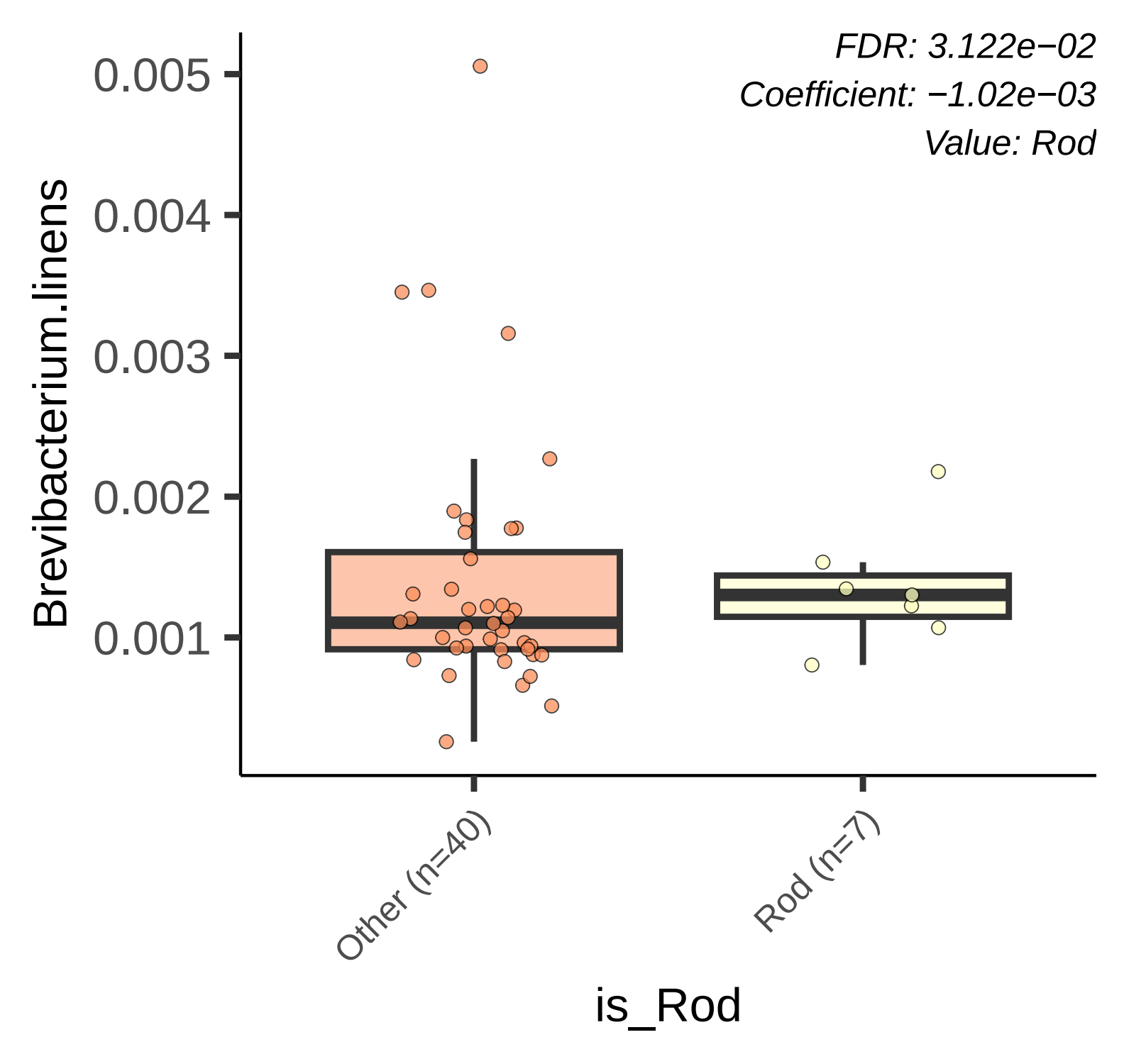

FDR: 1.465e−02
Coefficient: −2.66e−03
Value: Rod
Deinococcus.sp..NW.56
0.006
0.004
0.002
0.000
Other (n=40)
Rod (n=7)
is_Rod

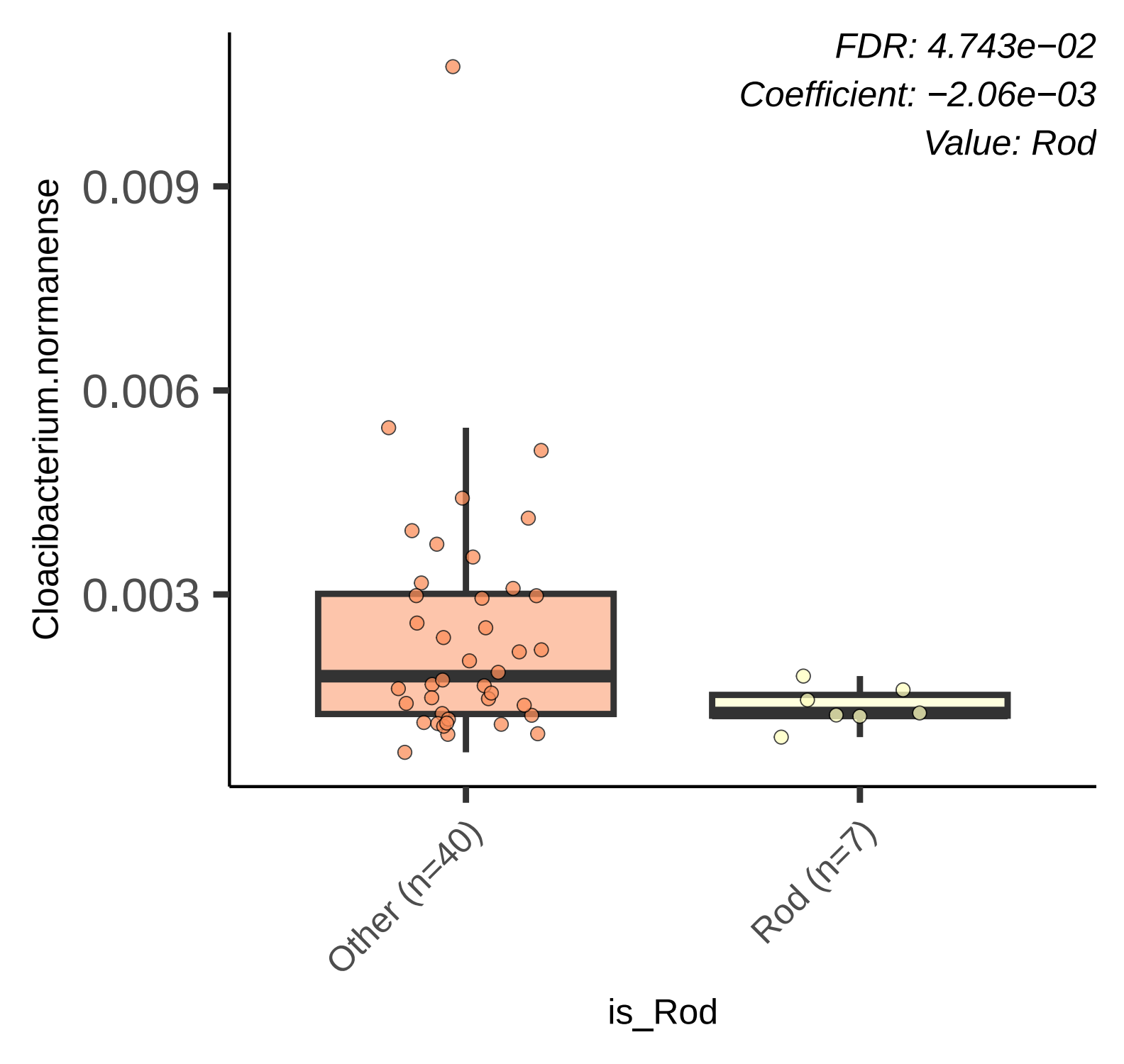

FDR: 1.562e−02
Coefficient: −7.31e−04
Value: Rod
Corynebacterium.aurimucosum
0.002
0.001
Other (n=40)
Rod (n=7)
is_Rod

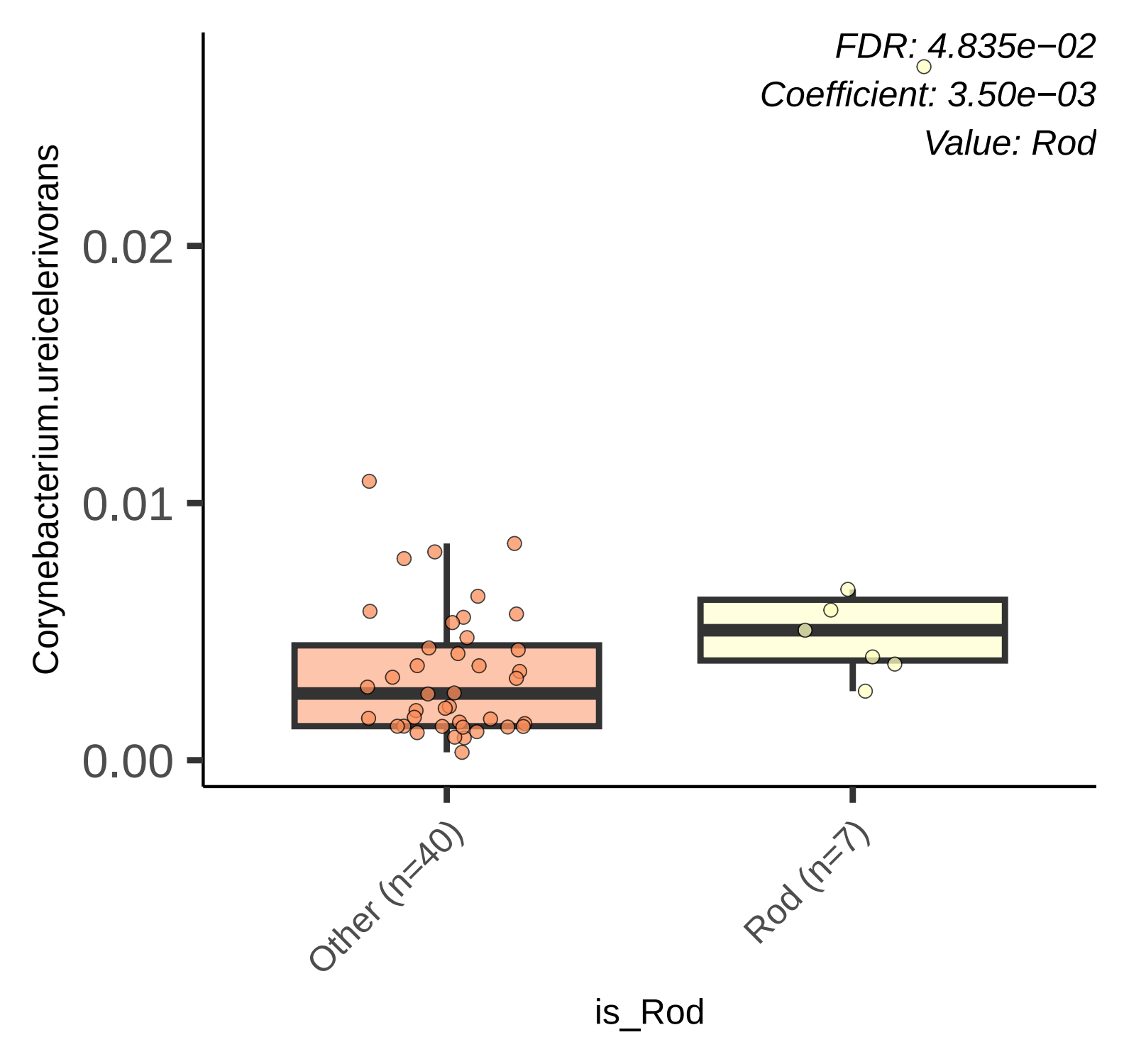

FDR: 1.737e−02
Coefficient: −2.72e−03
Value: Rod
Tessaracoccus.sp..T2.5.30
0.006
0.004
0.002
0.000
Other (n=40)
Rod (n=7)
is_Rod


FDR: 2.051e−02
Coefficient: −4.32e−03
Value: Rod
Kocuria.flava
0.015
0.010
0.005
0.000
Other (n=40)
Rod (n=7)
is_Rod


FDR: 2.361e−02
Coefficient: −1.05e−03
Value: Rod
0.005
0.004
0.003
0.002
0.001
0.000
Dietzia.sp..oral.taxon.368
Other (n=40)
Rod (n=7)
is_Rod


FDR: 3.122e−02
Coefficient: −1.02e−03
Value: Rod
Brevibacterium.linens
0.005
0.004
0.003
0.002
0.001
Other (n=40)
Rod (n=7)
is_Rod


FDR: 4.743e−02
Coefficient: −2.06e−03
Value: Rod
Cloacibacterium.normanense
0.009
0.006
0.003
Other (n=40)
Rod (n=7)
is_Rod


FDR: 4.835e−02
Coefficient: 3.50e−03
Value: Rod
Corynebacterium.ureicelerivorans
0.02
0.01
0.00
Other (n=40)
Rod (n=7)
is_Rod