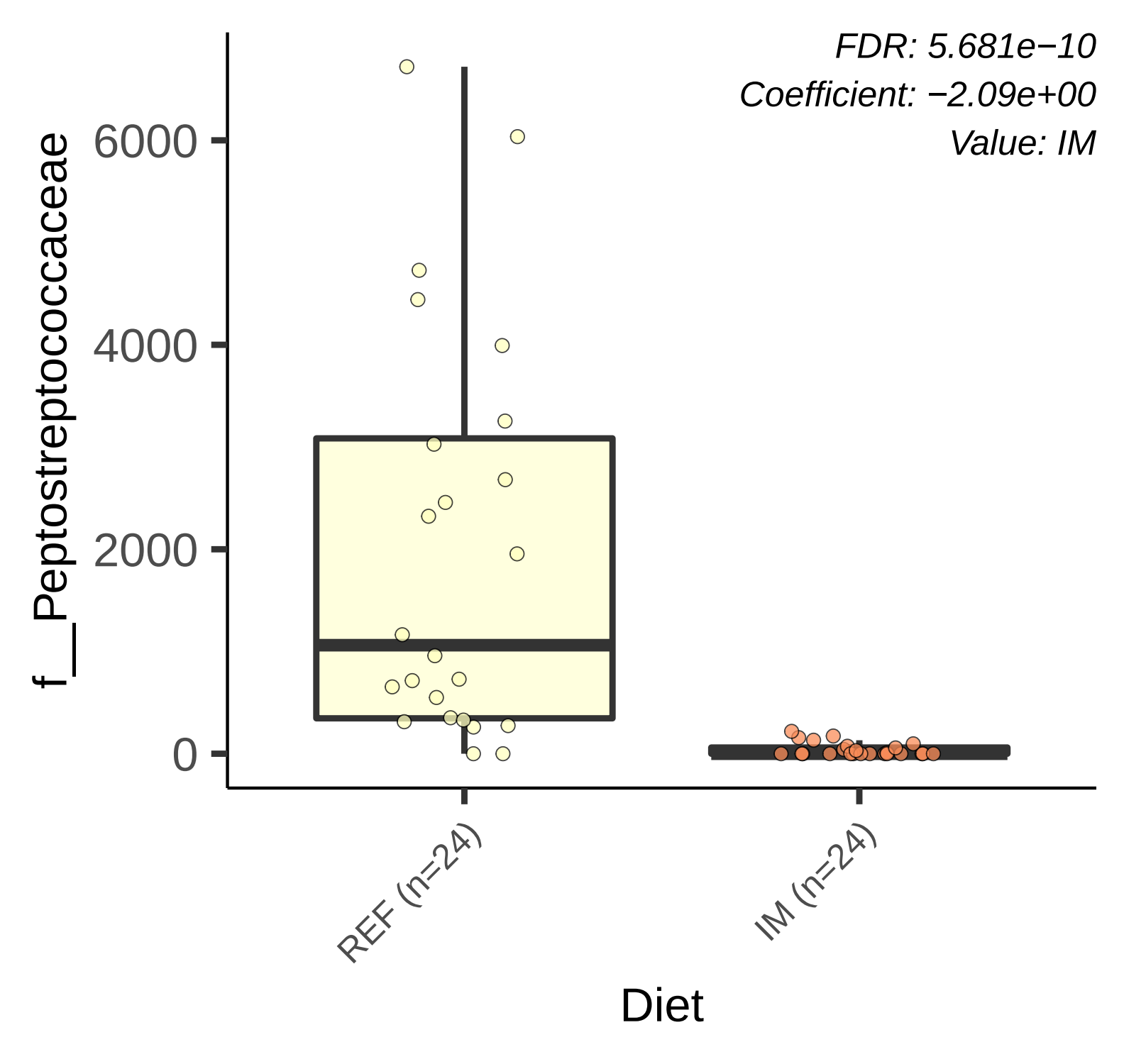
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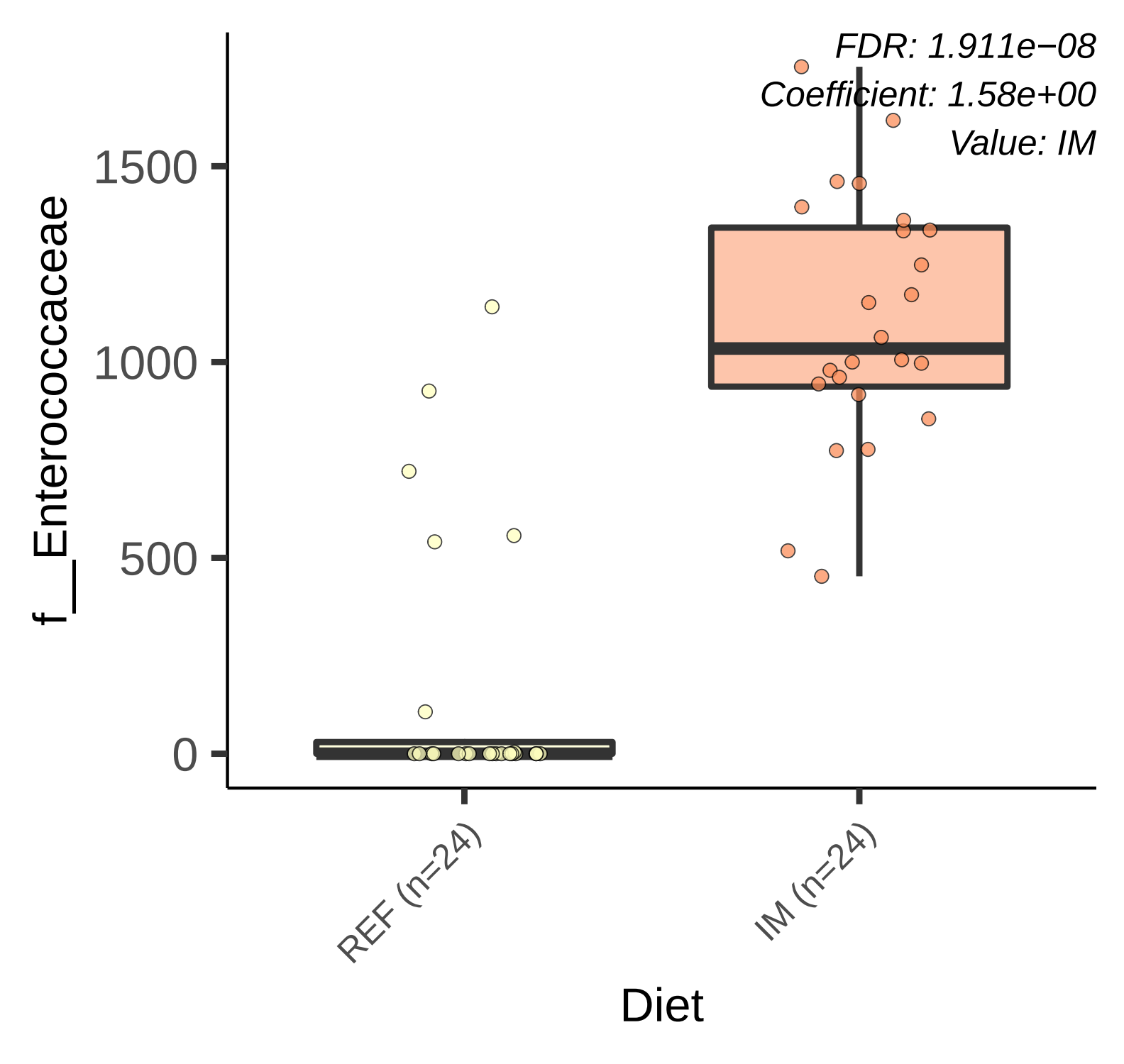
Diet


FDR: 1.911e−08
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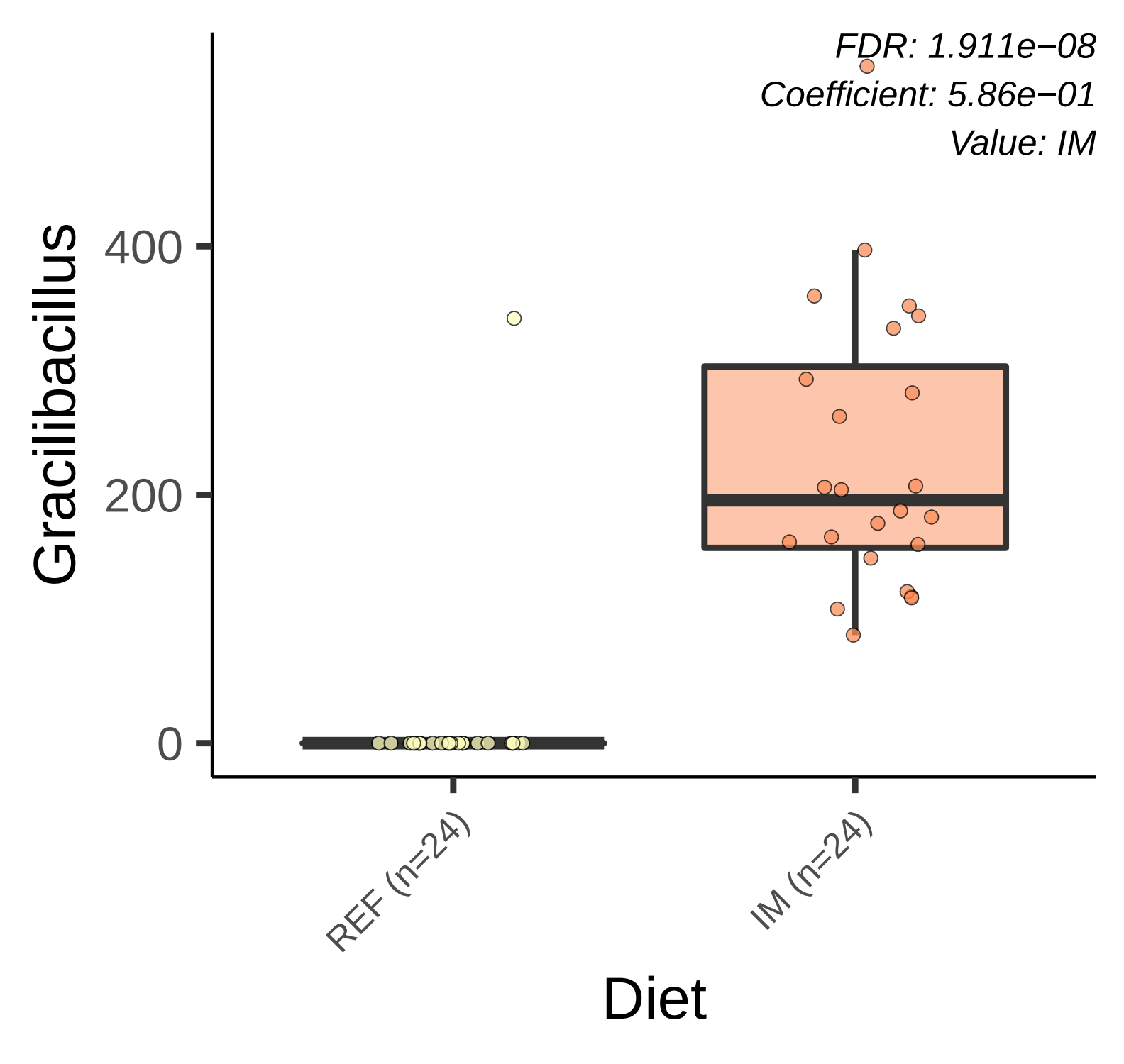
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Gracilibacillus
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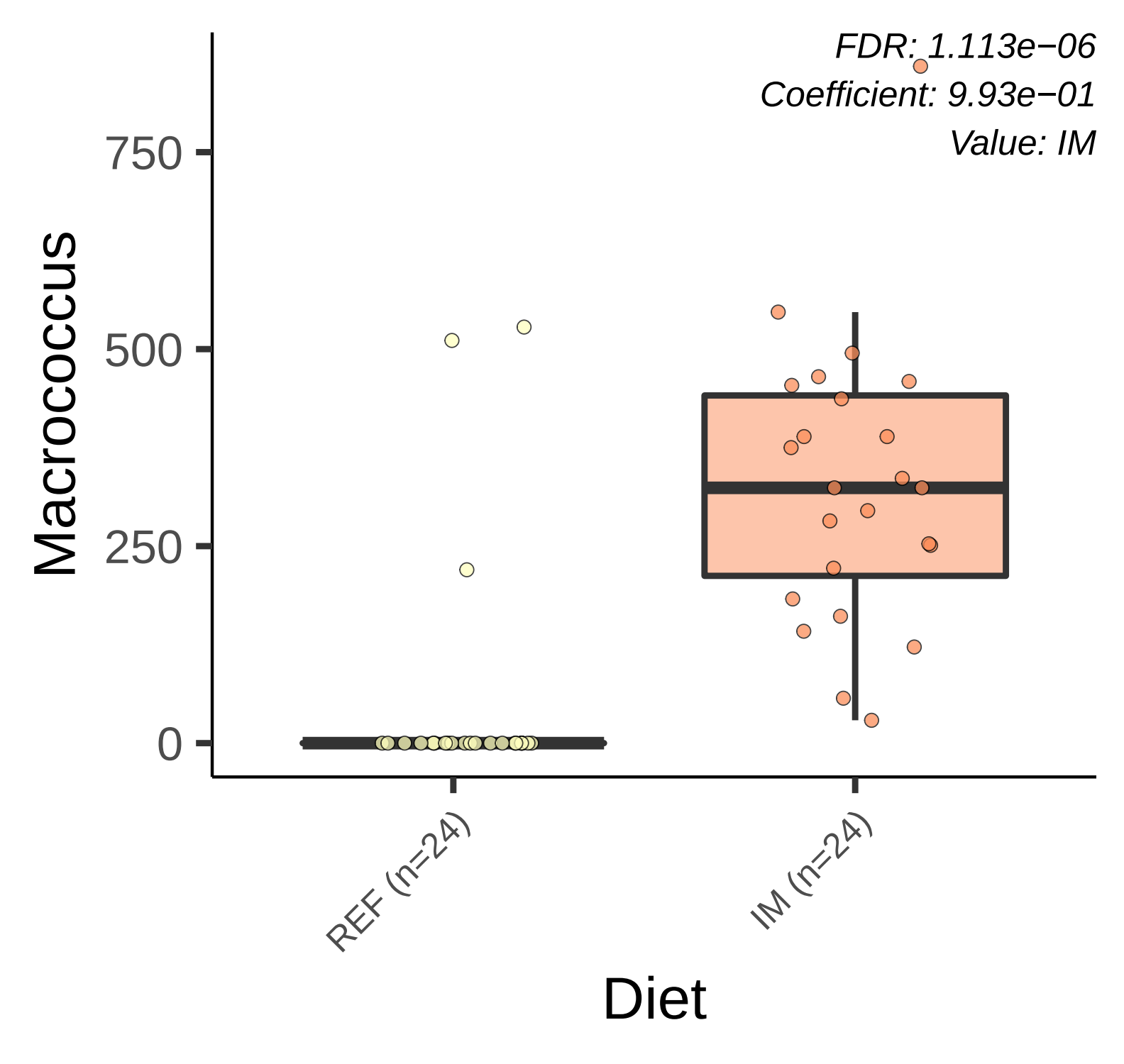
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Macrococcus
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IM (n=24)
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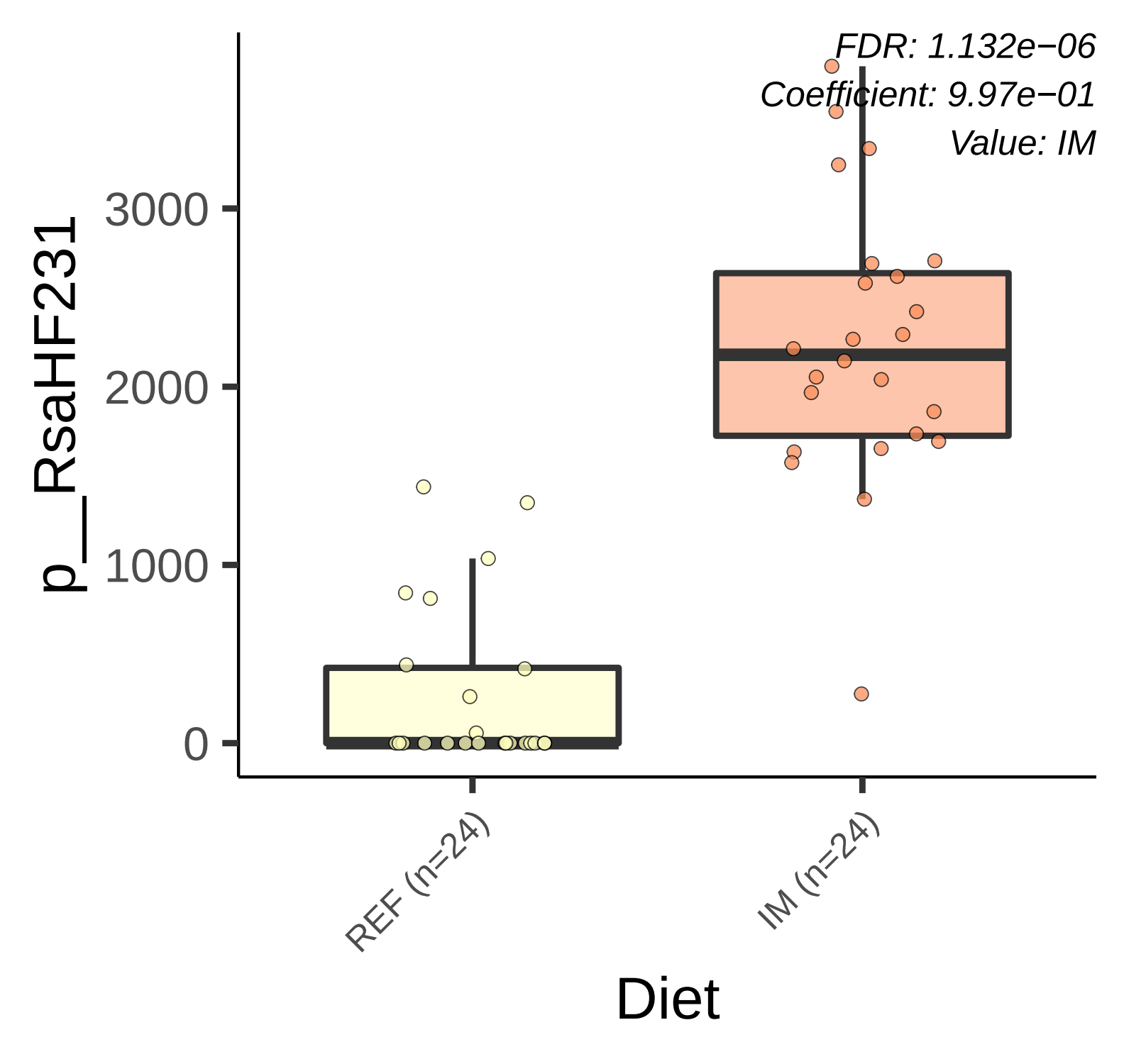
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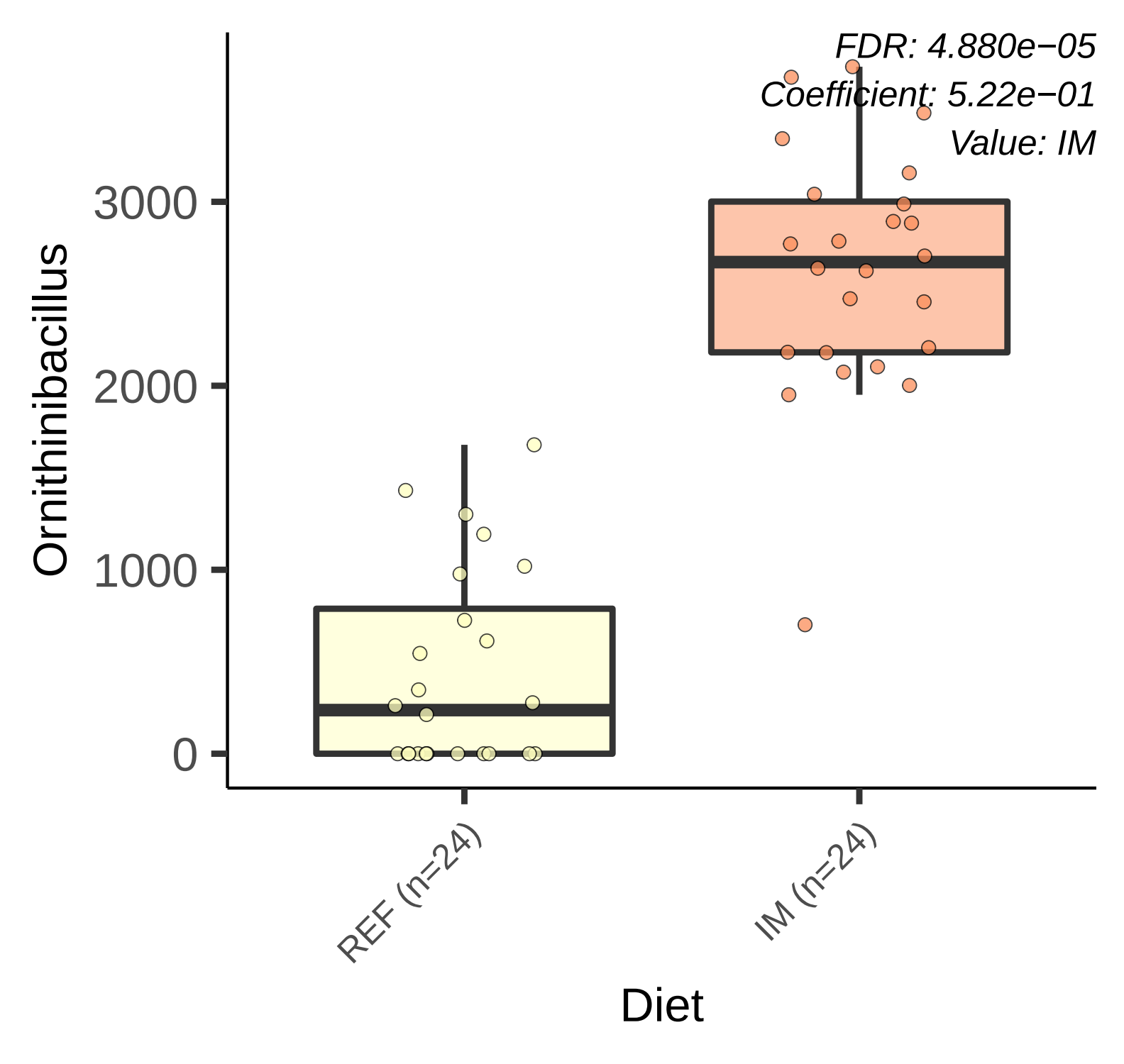
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Ornithinibacillus
3000
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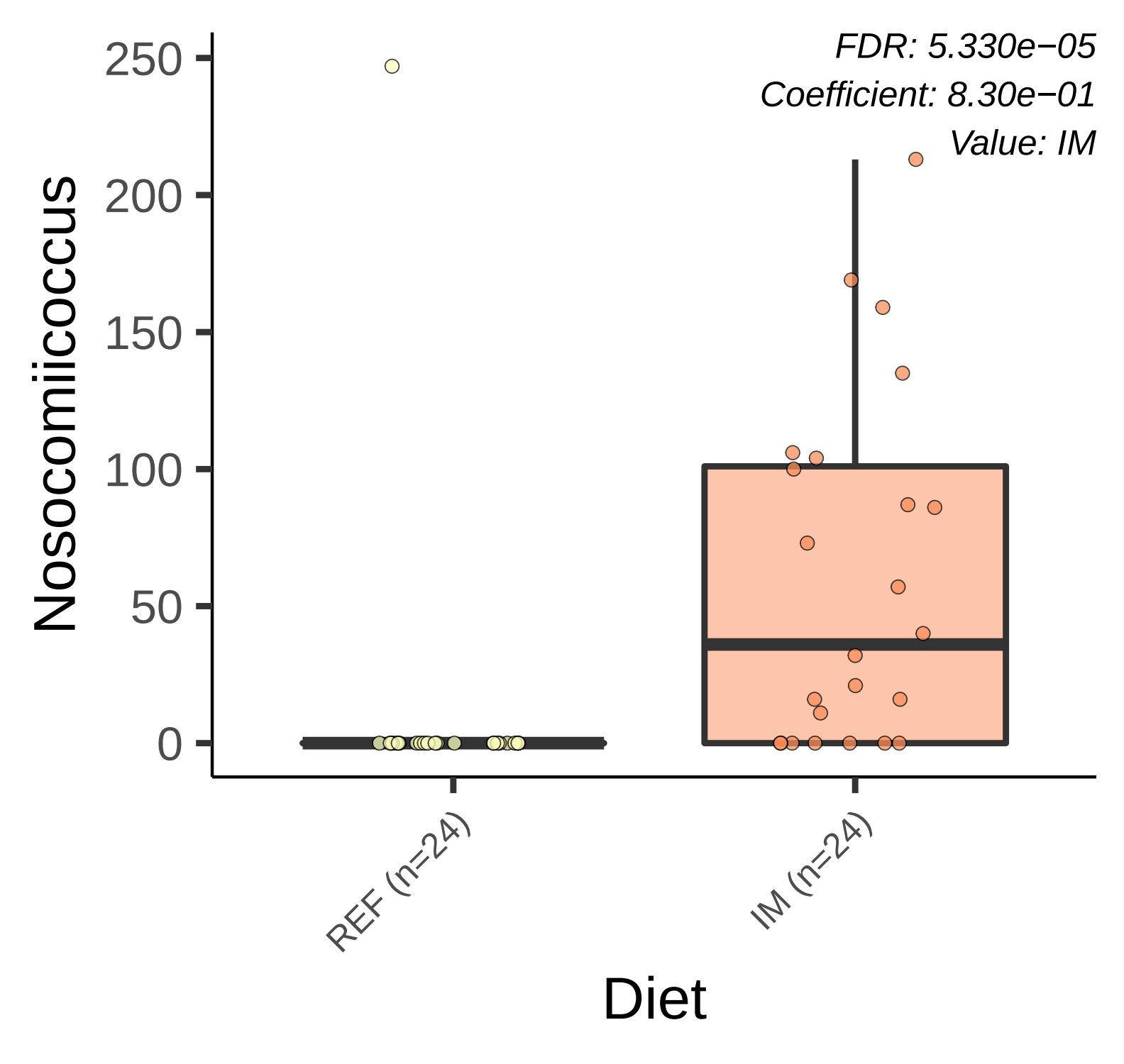
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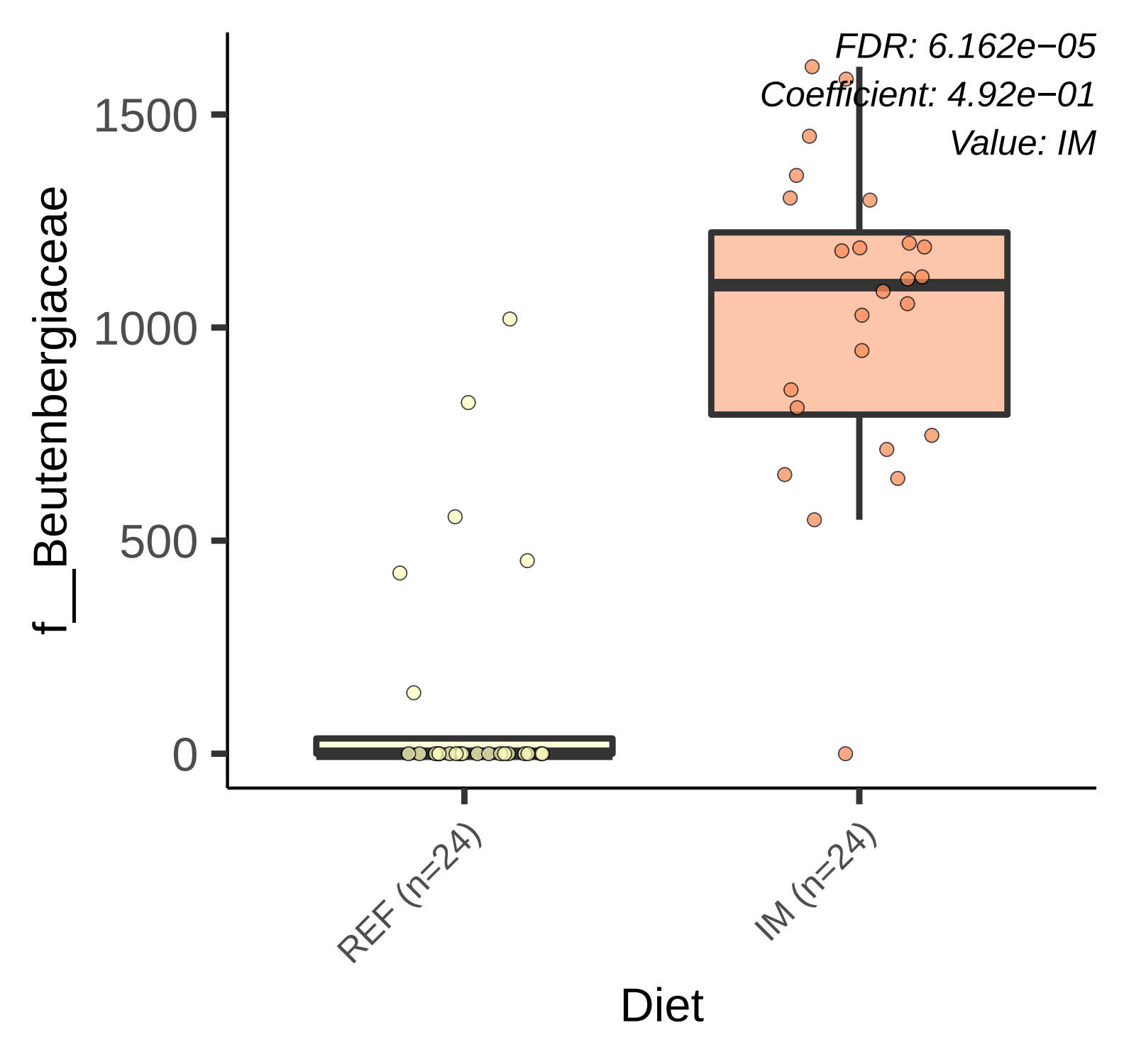
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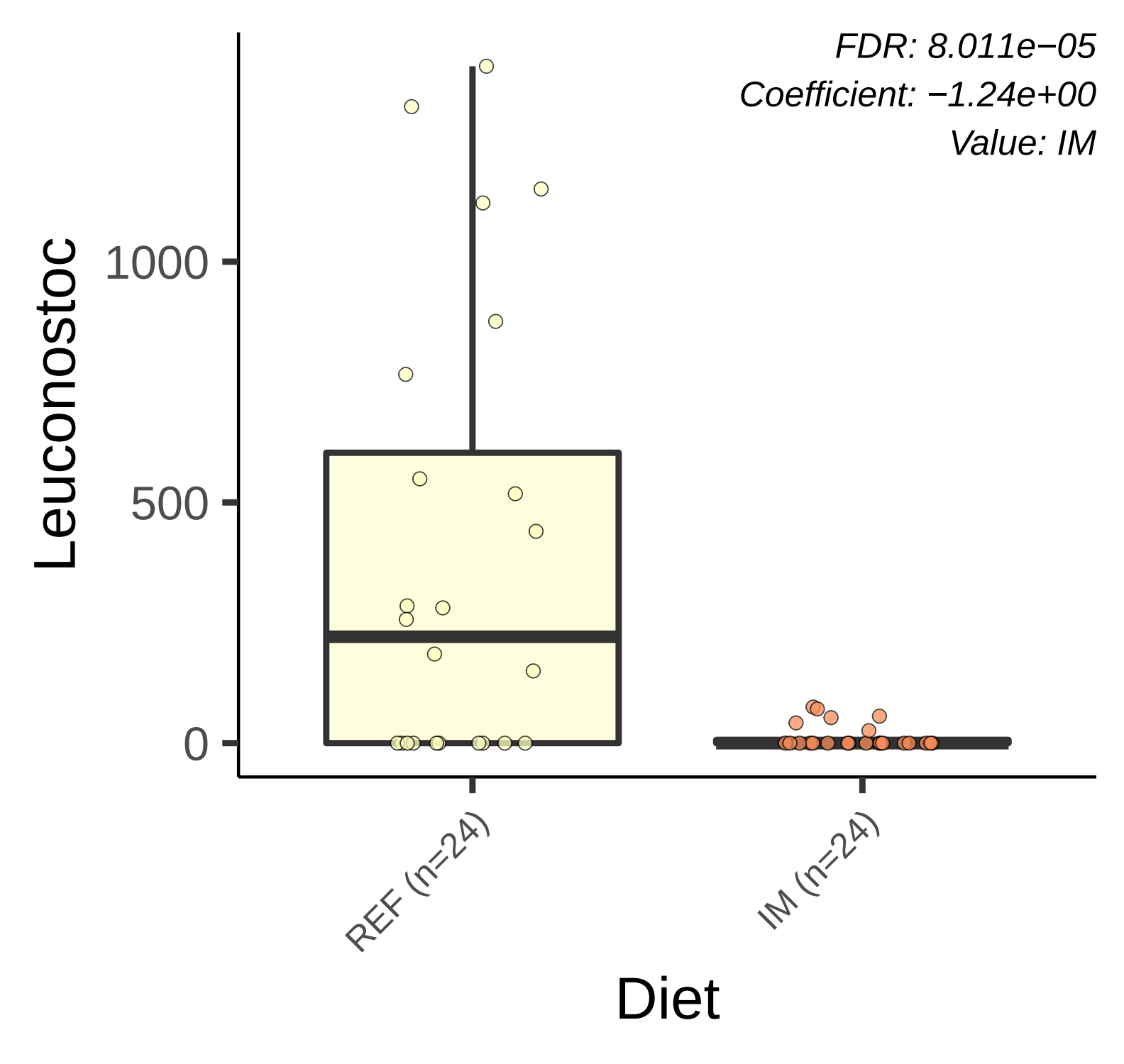
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Leuconostoc
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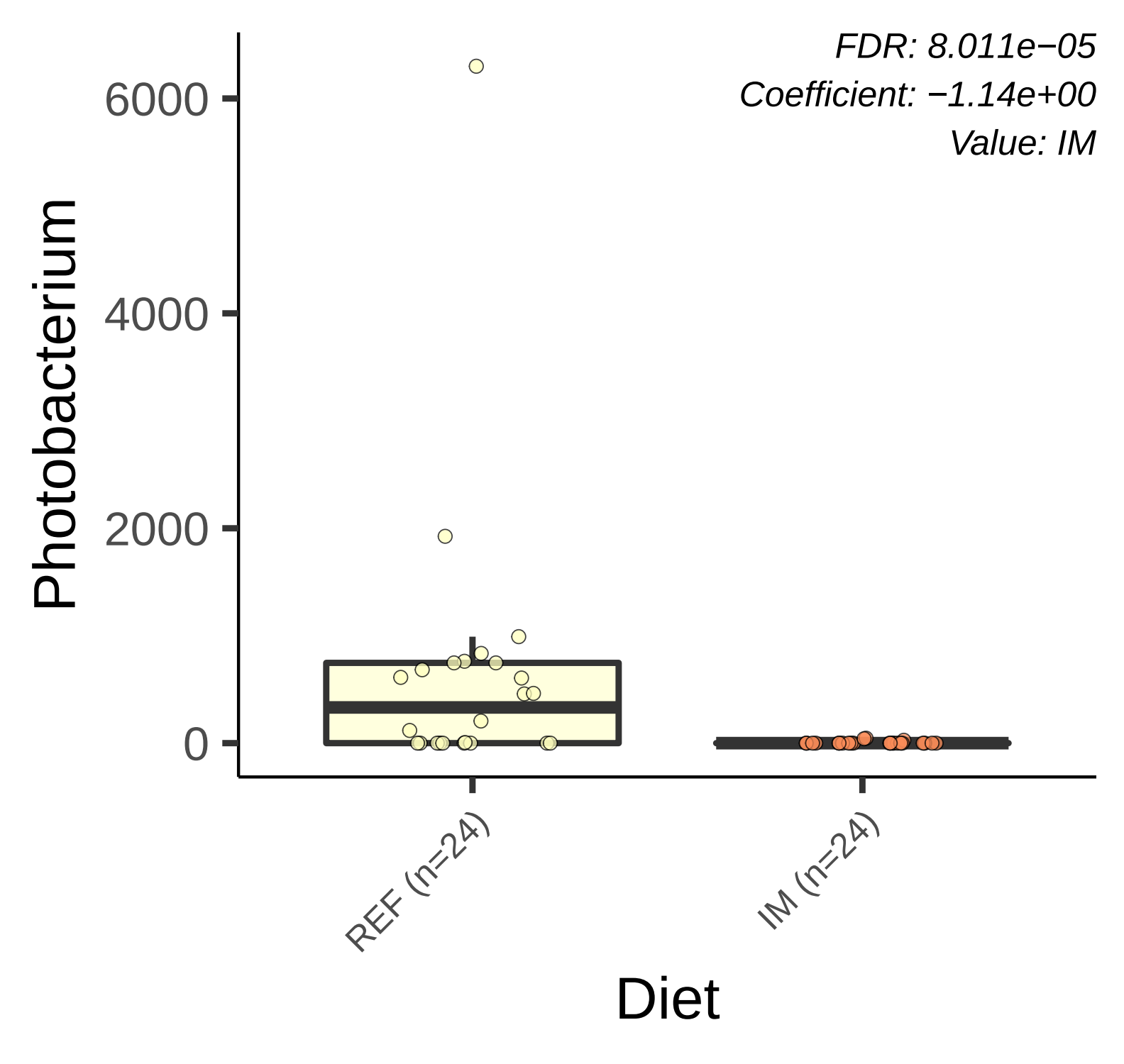
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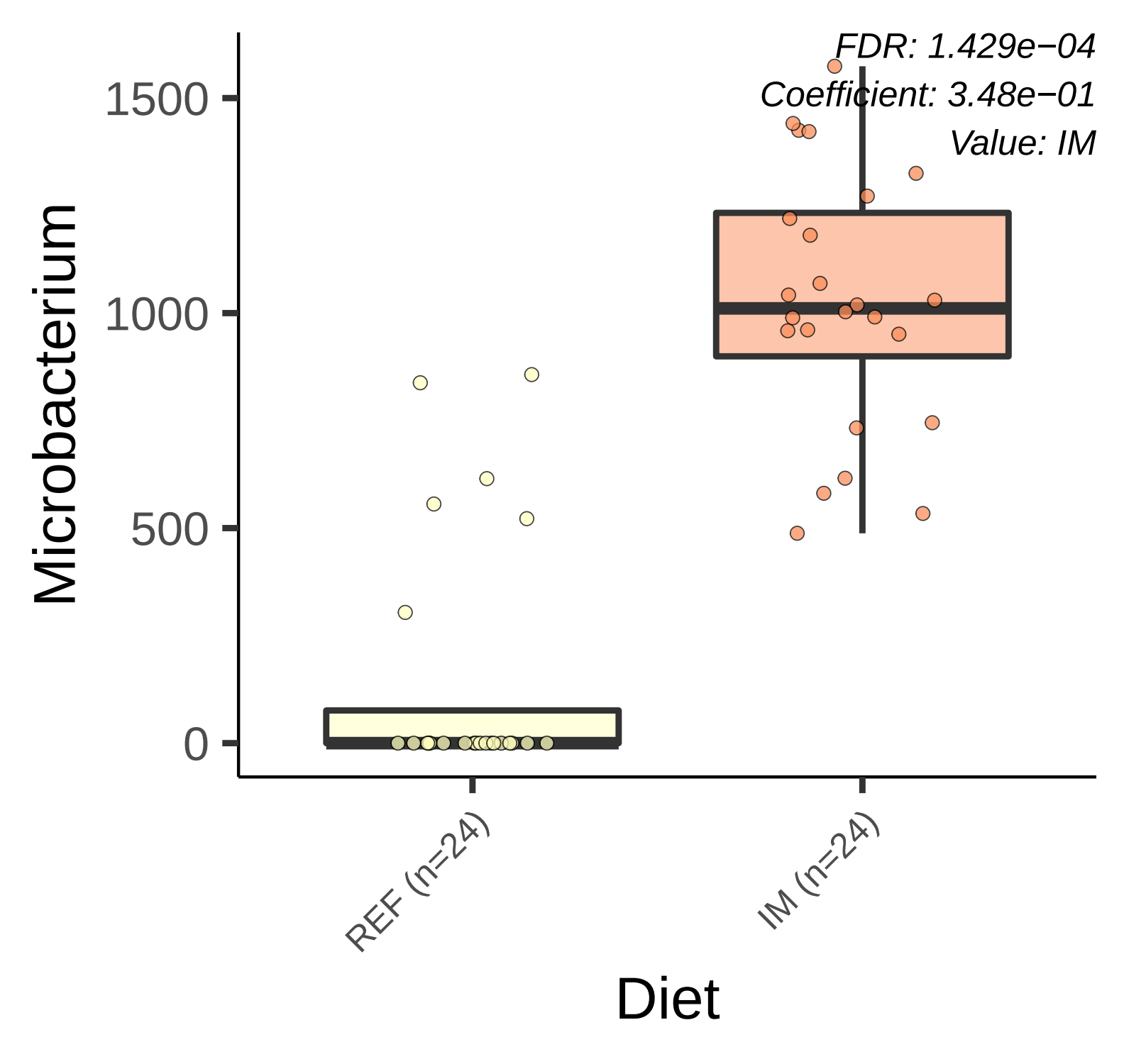
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Microbacterium
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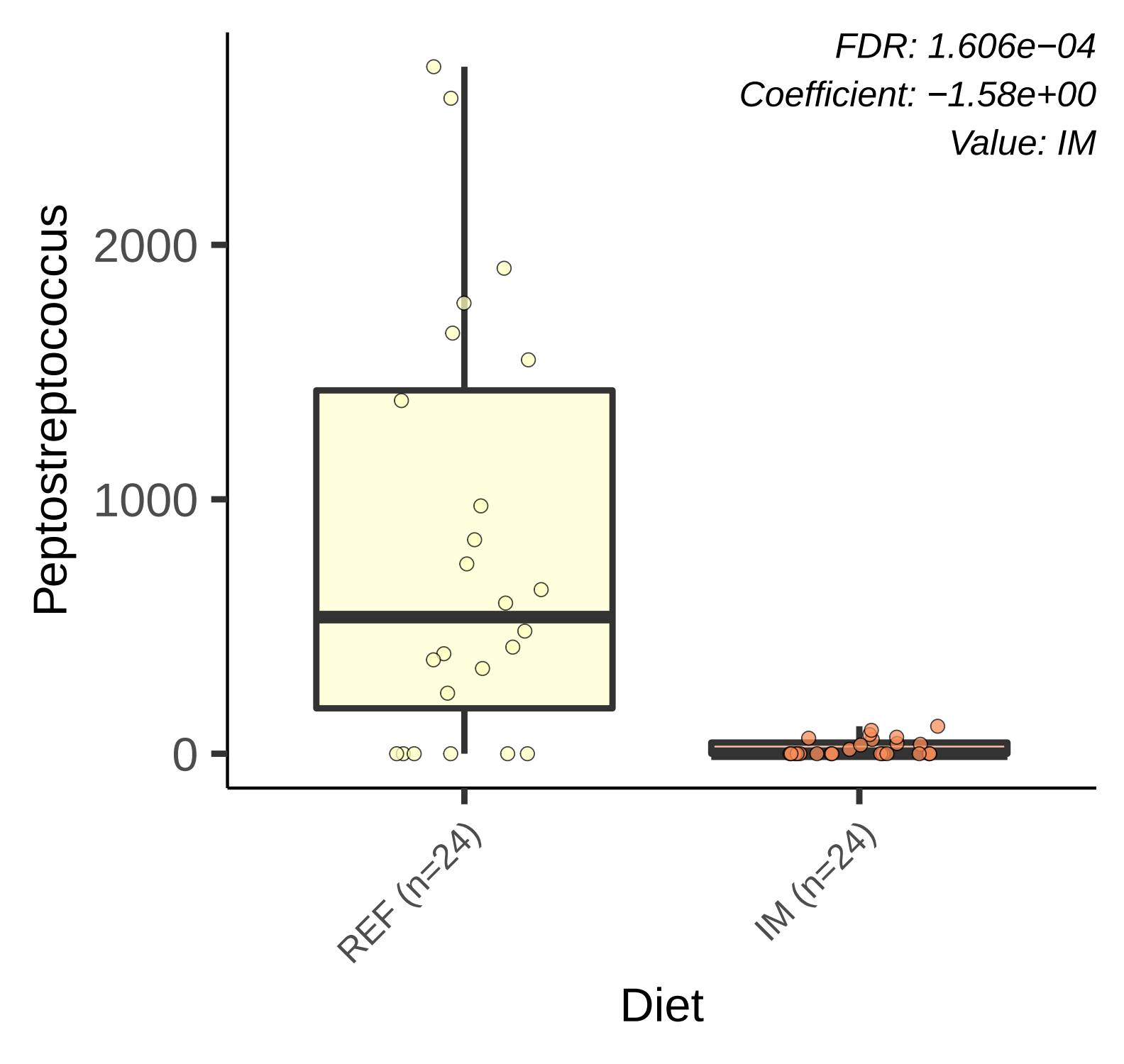
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Peptostreptococcus
2000
1000
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IM (n=24)
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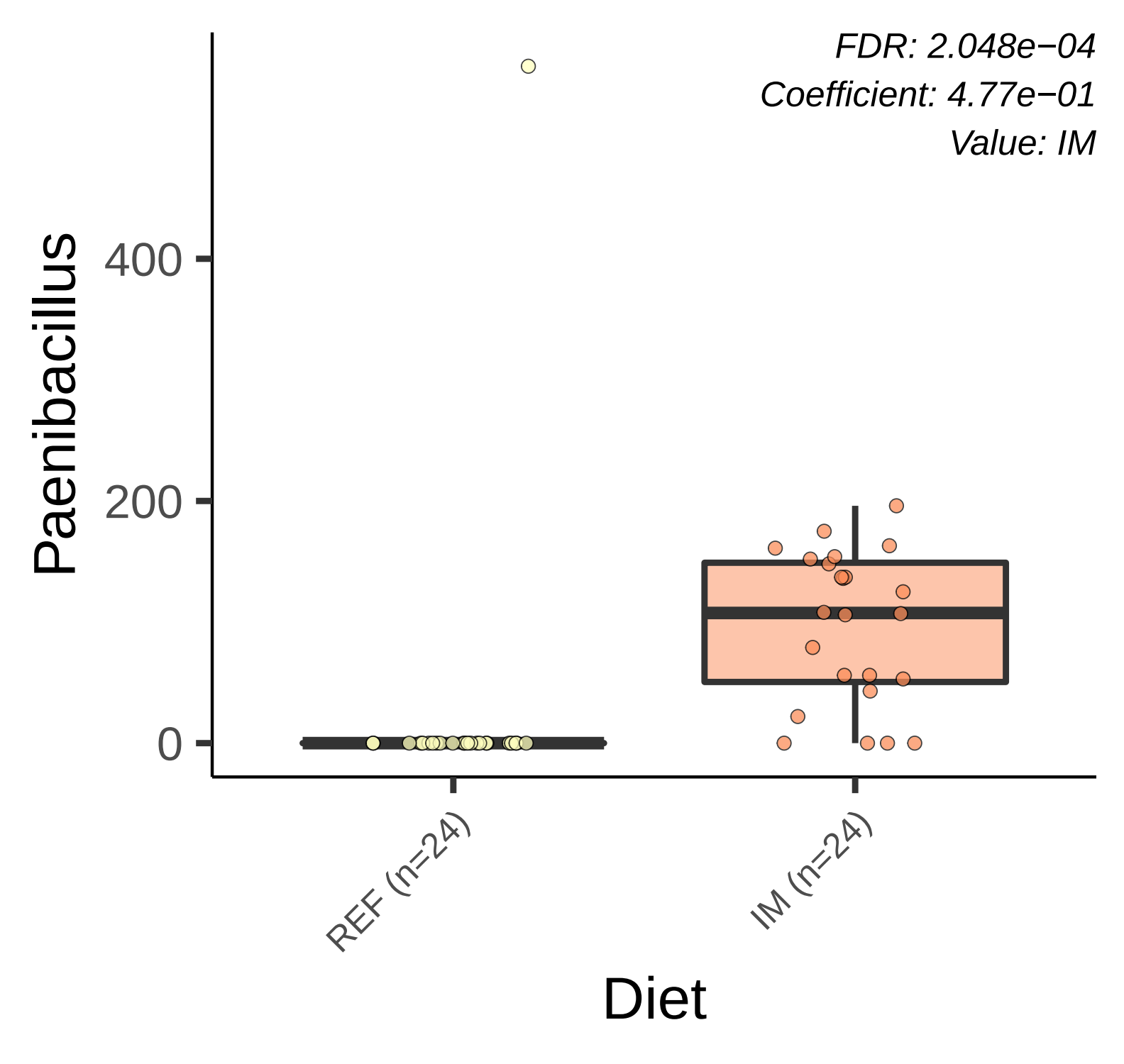
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Paenibacillus
400
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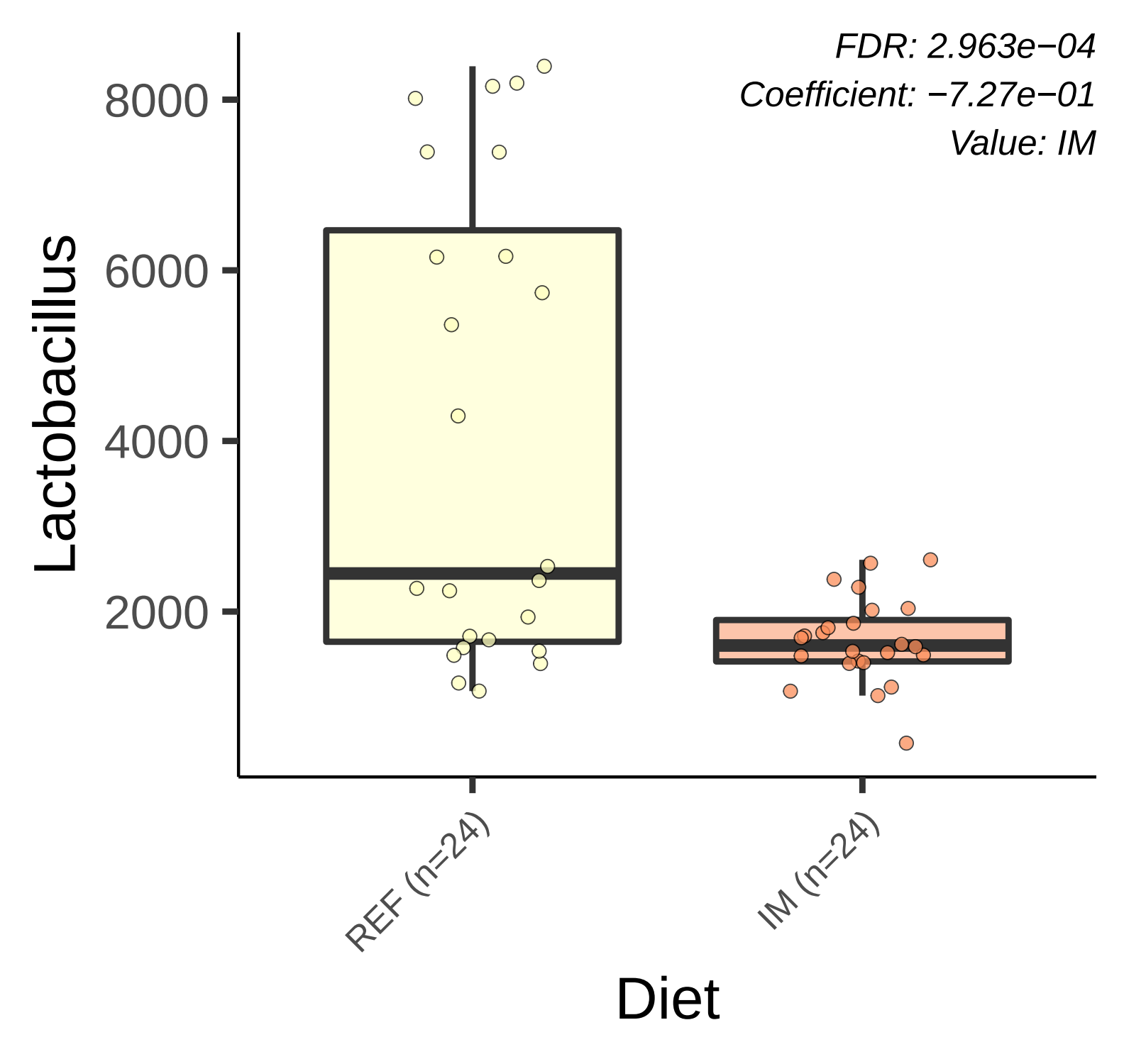
IM (n=24)
Diet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8000
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4000
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Lactobacillus
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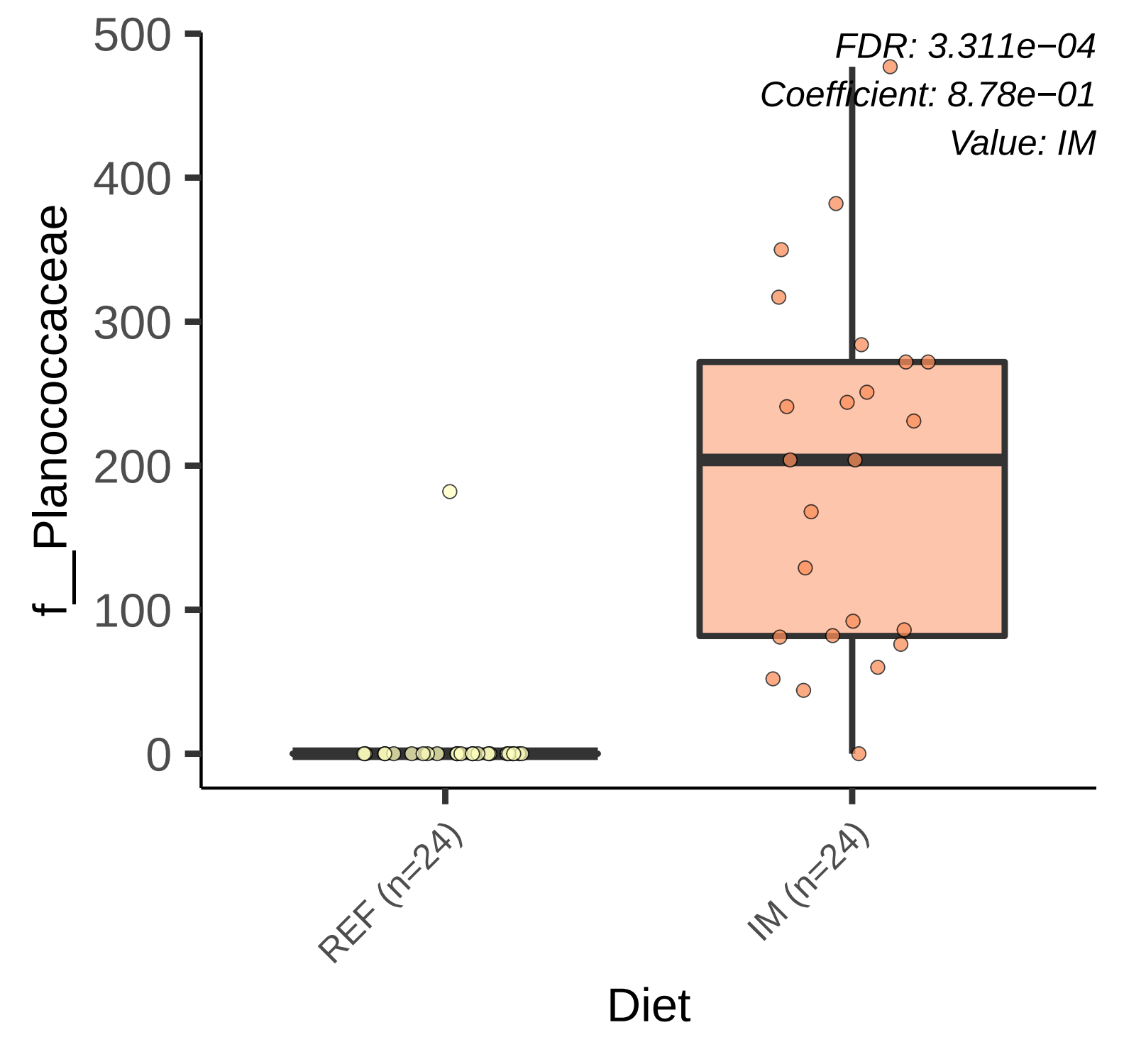
IM (n=24)
Diet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IM (n=24)
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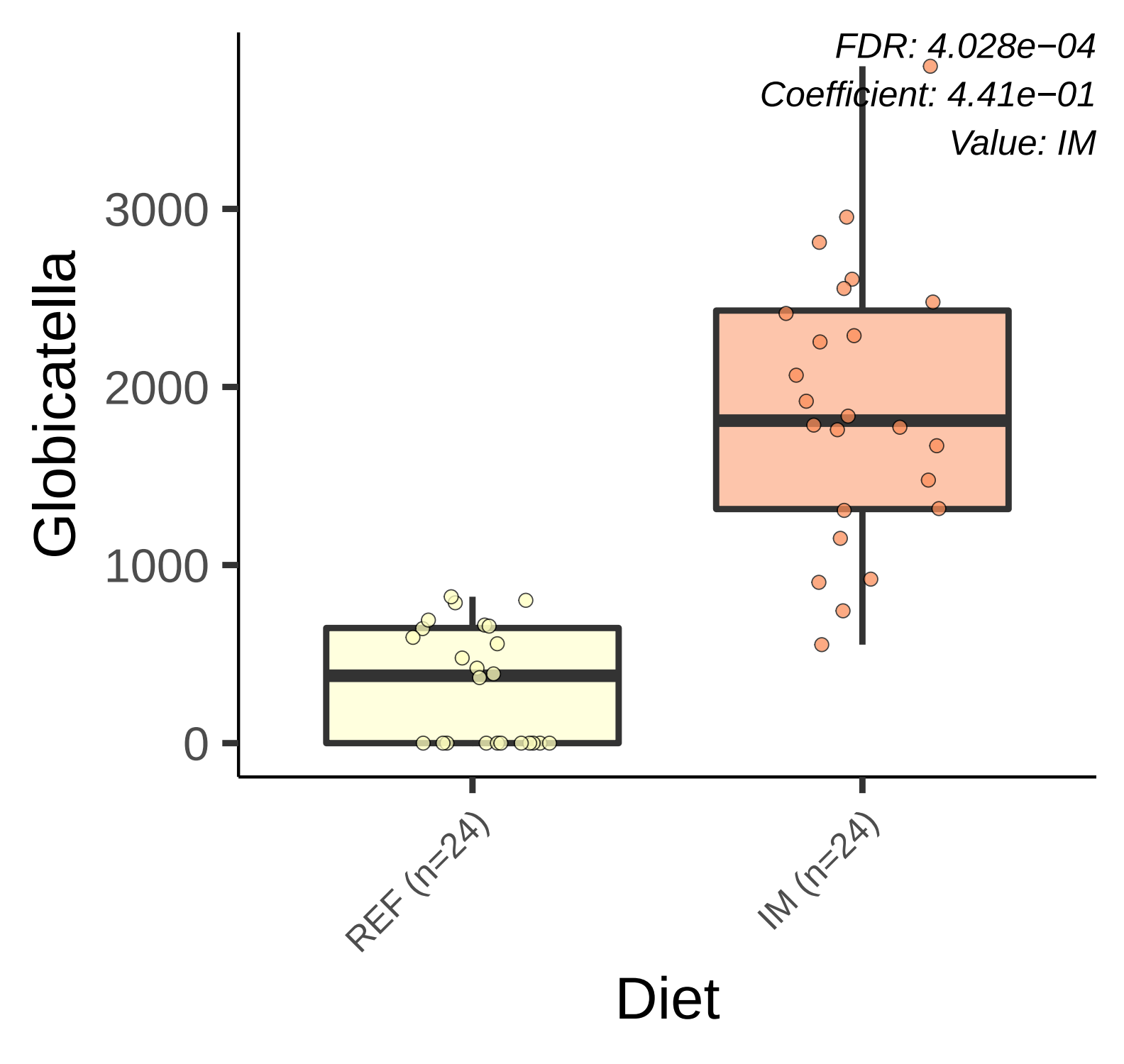
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Coefficient: 4.41e−01
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Globicatella
3000
2000
1000
0
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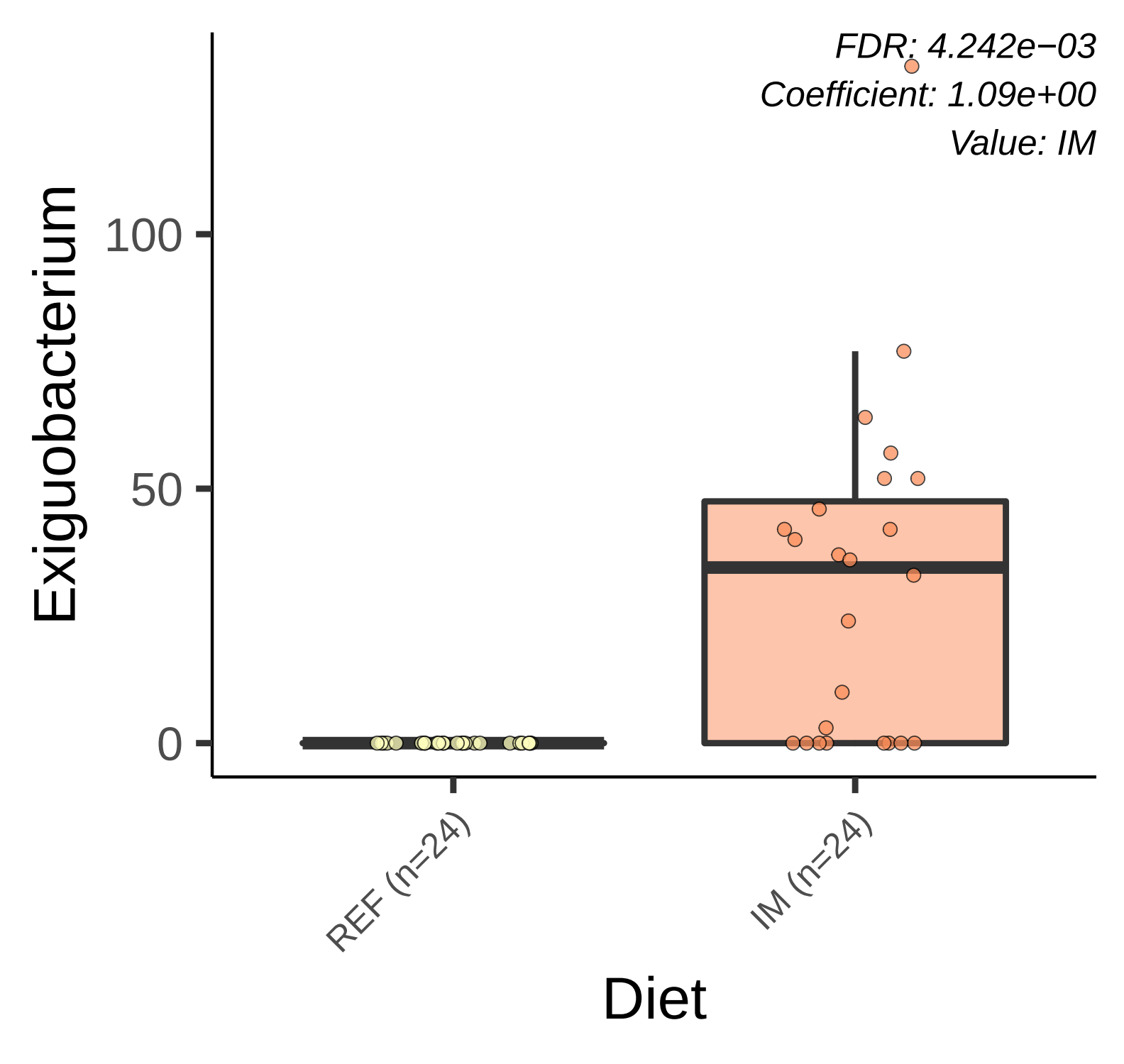
IM (n=24)
Diet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Exiguobacterium
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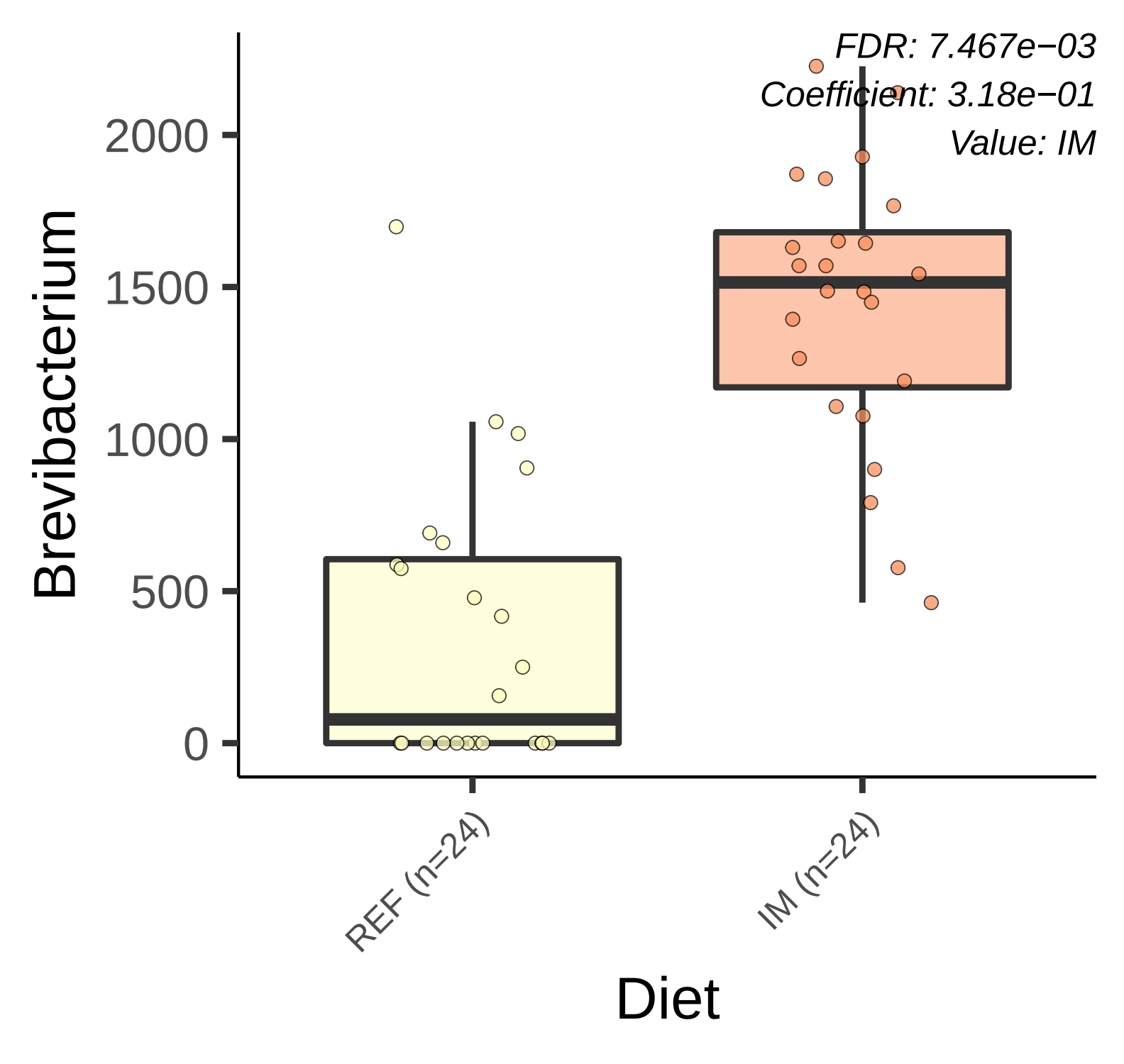
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Brevibacterium
2000
1500
1000
500
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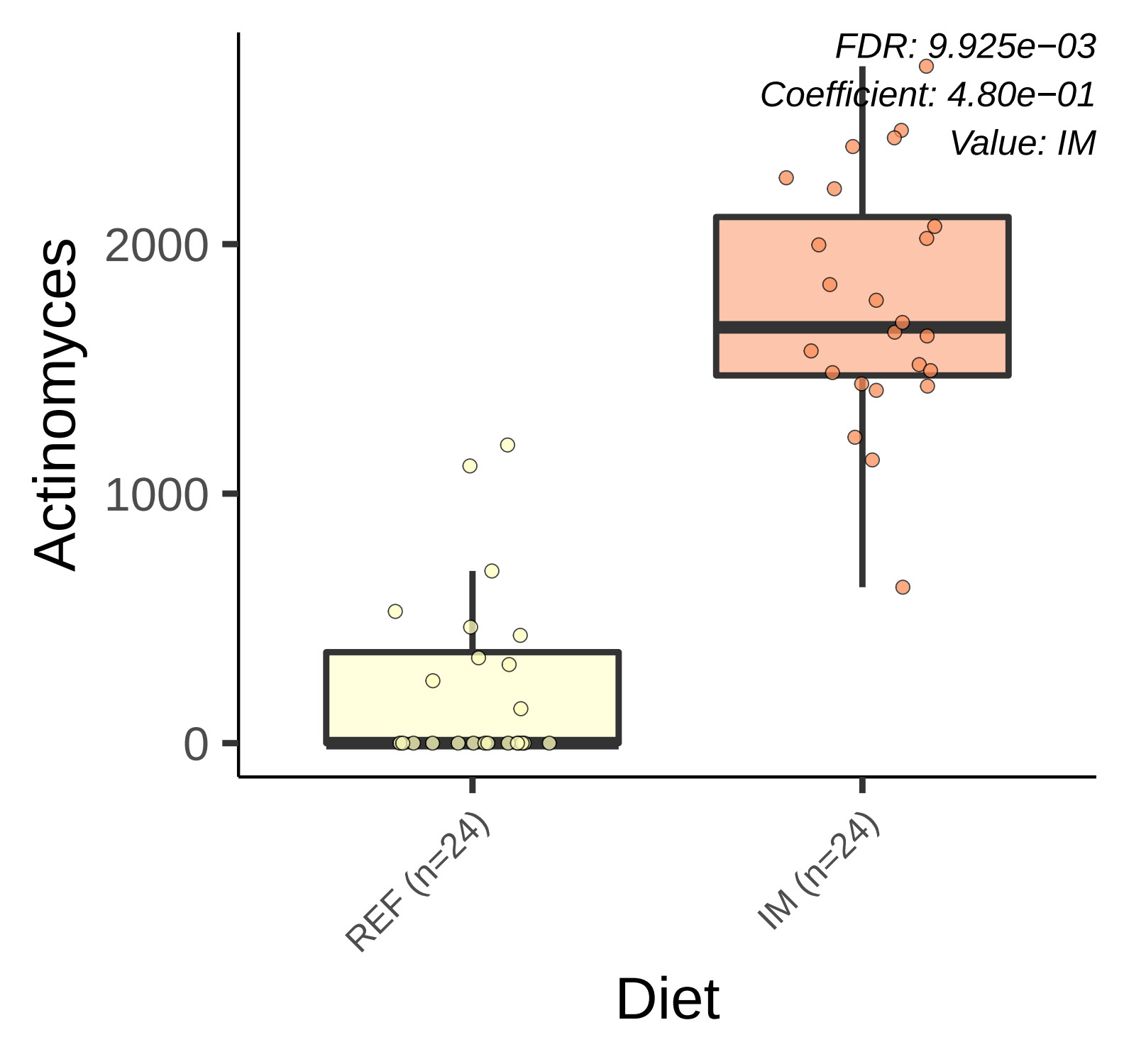
Diet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Actinomyces
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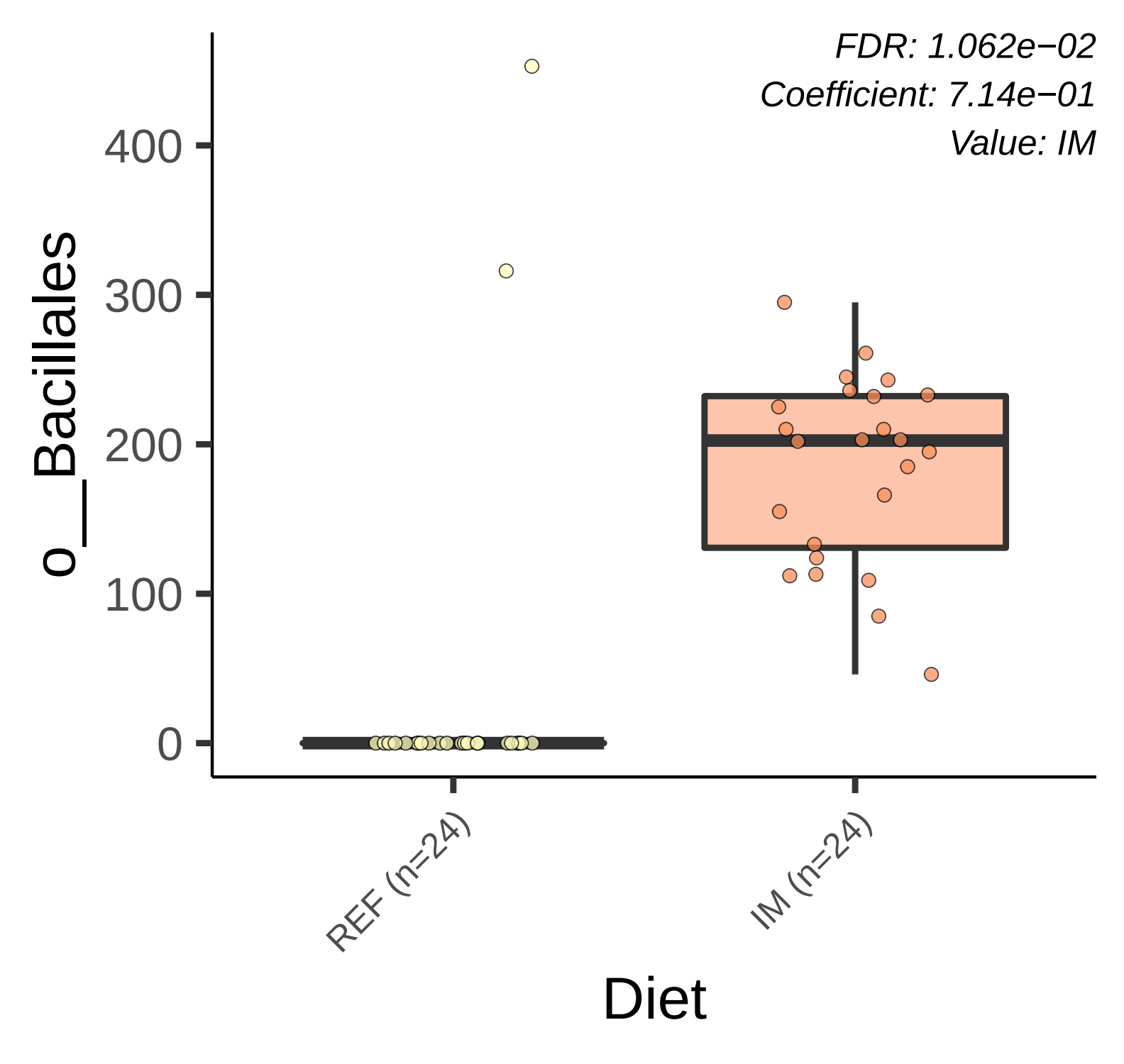
FDR: 1.062e−02
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o__Bacillales
400
300
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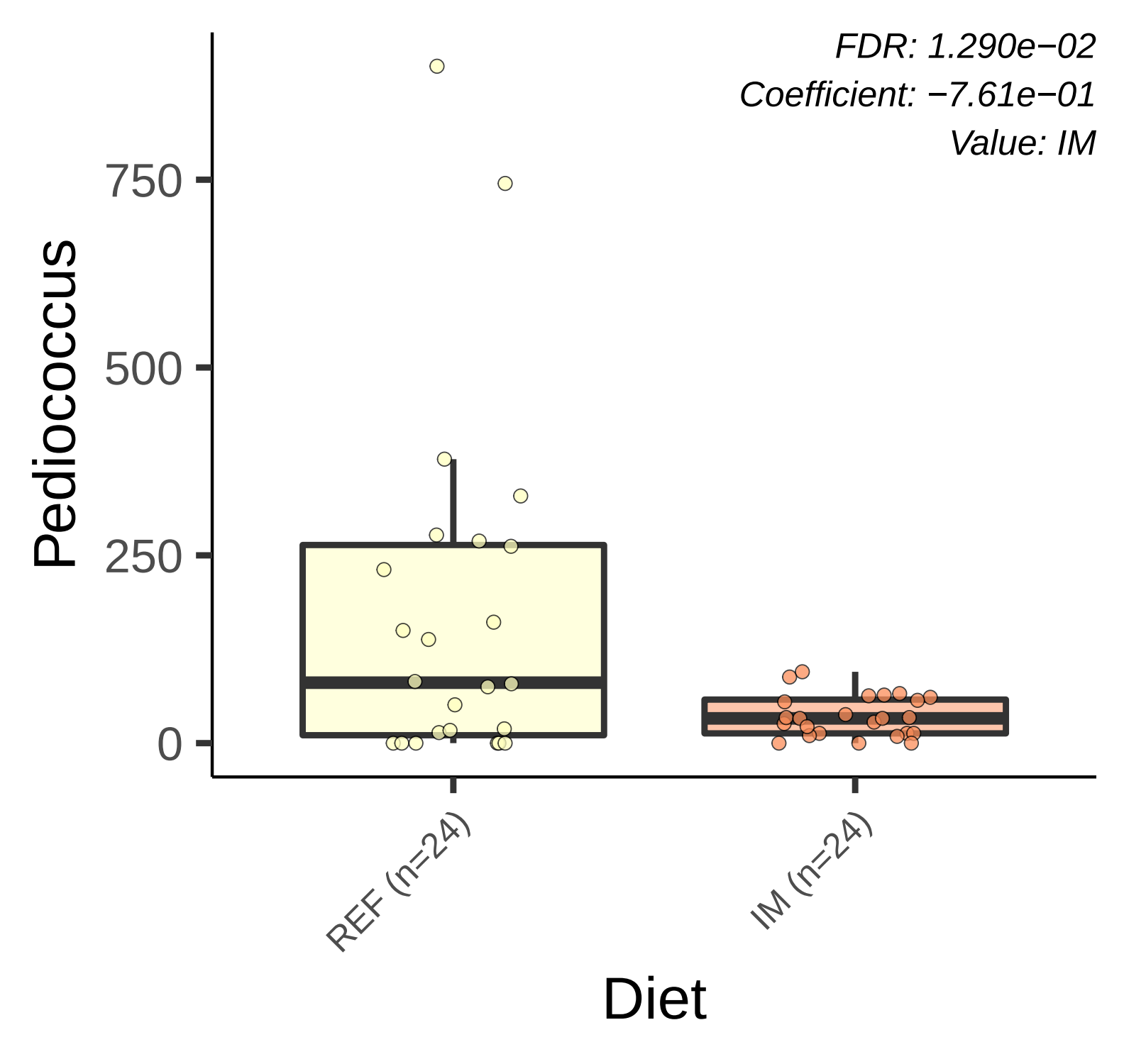
IM (n=24)
Diet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Pediococcus
750
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250
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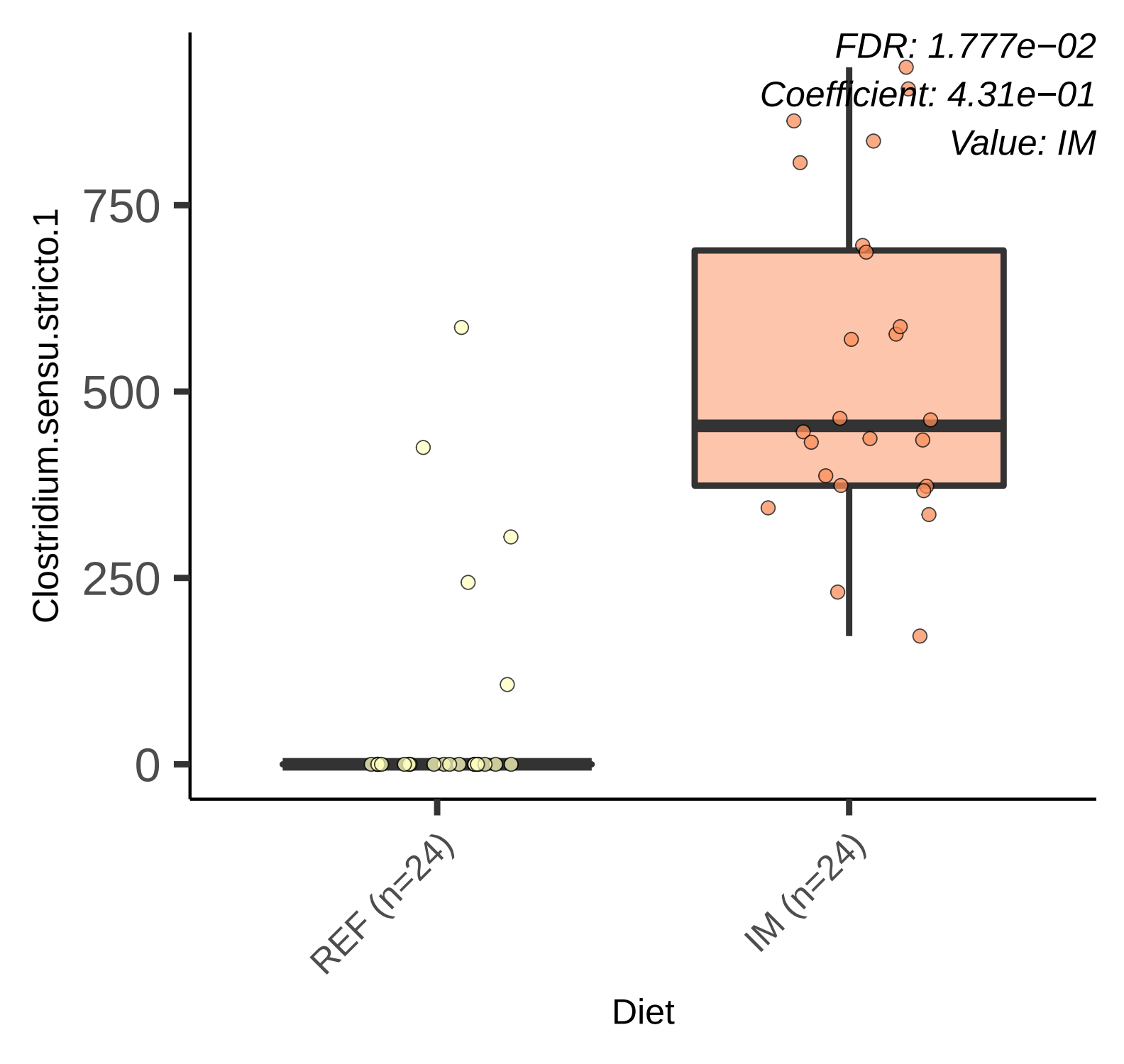
IM (n=24)
Diet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Clostridium.sensu.stricto.1
750
500
250
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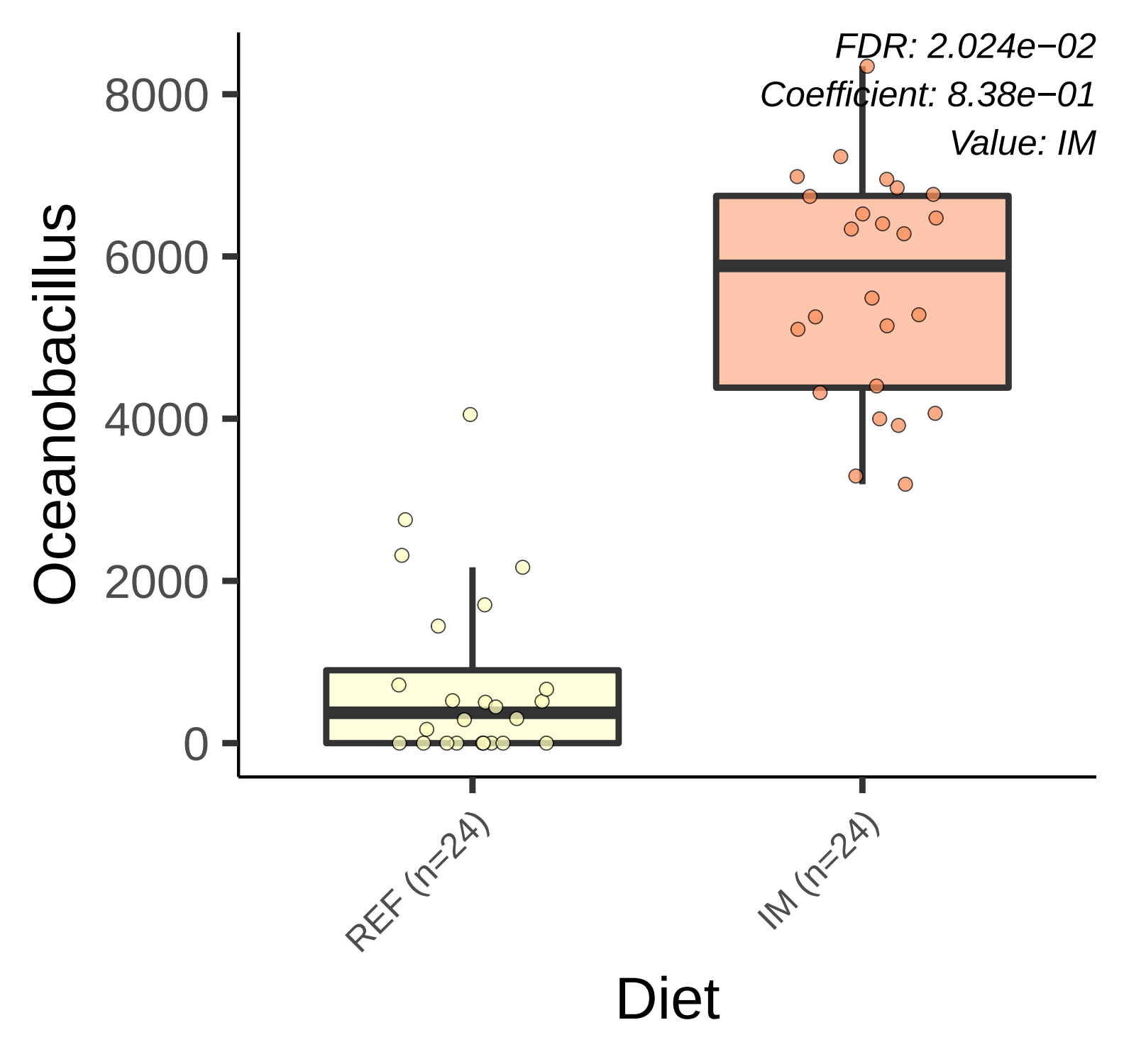
IM (n=24)
Diet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2000
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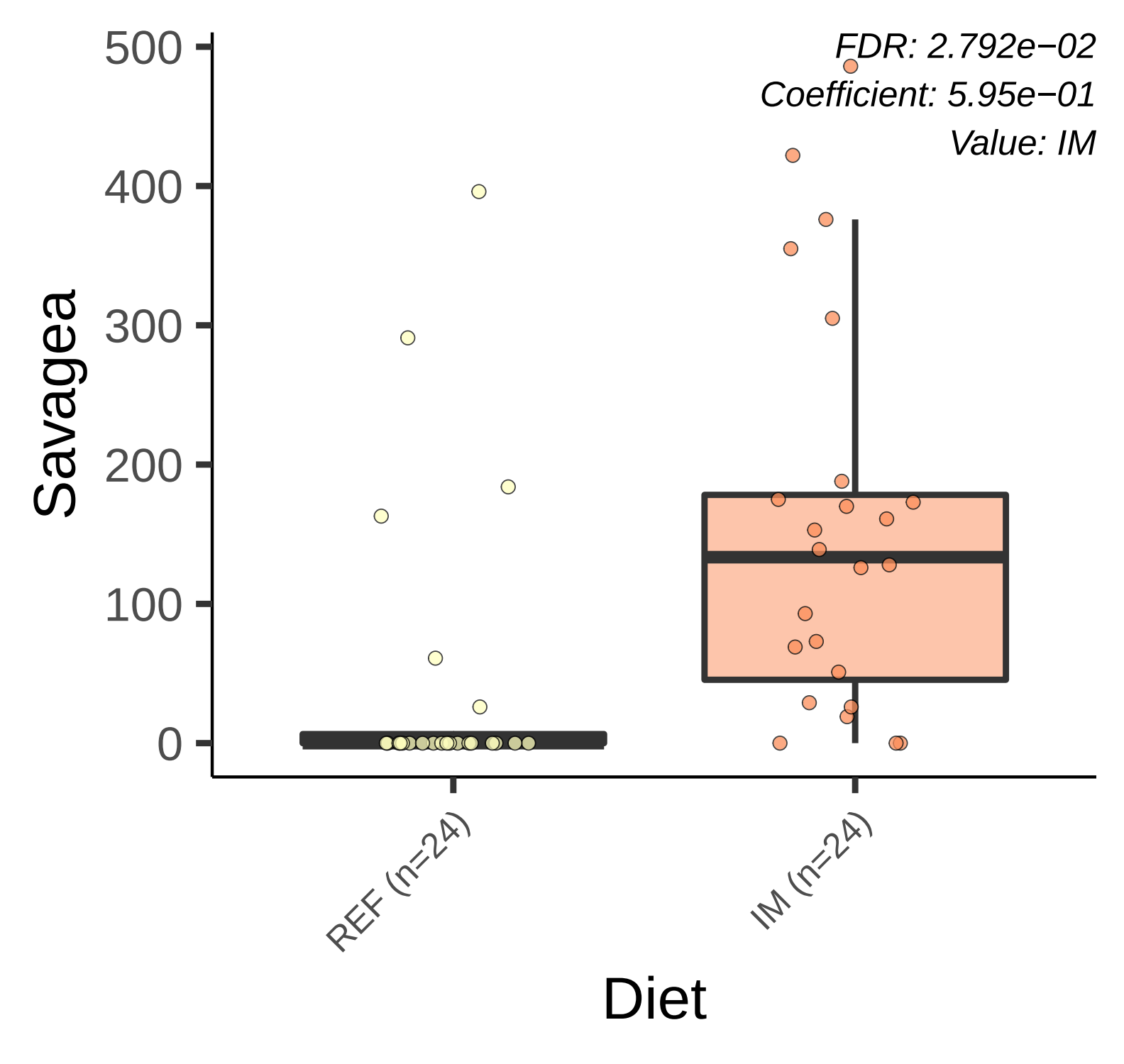
IM (n=24)
Diet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Savagea
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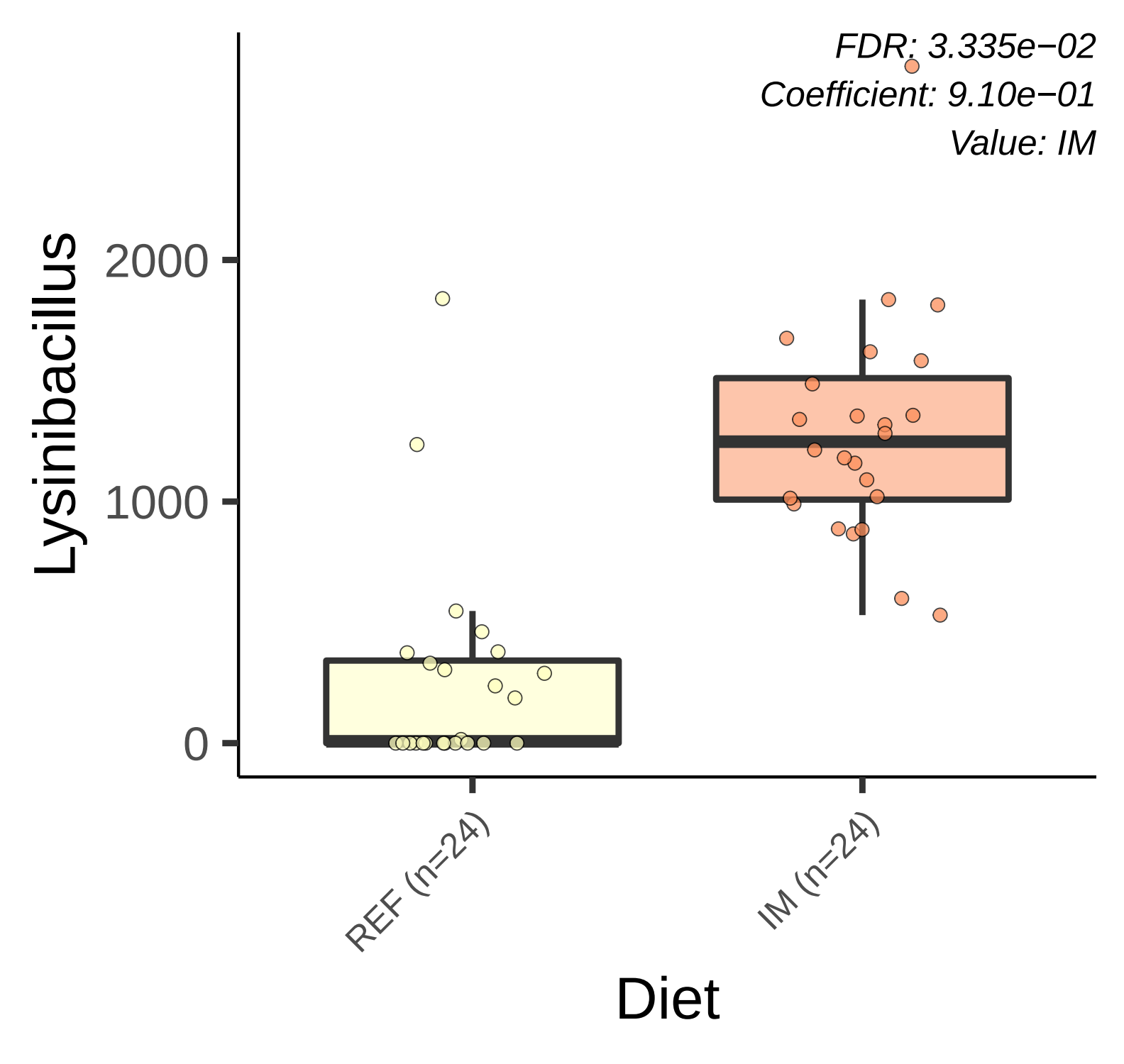
Diet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Lysinibacillus
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1000
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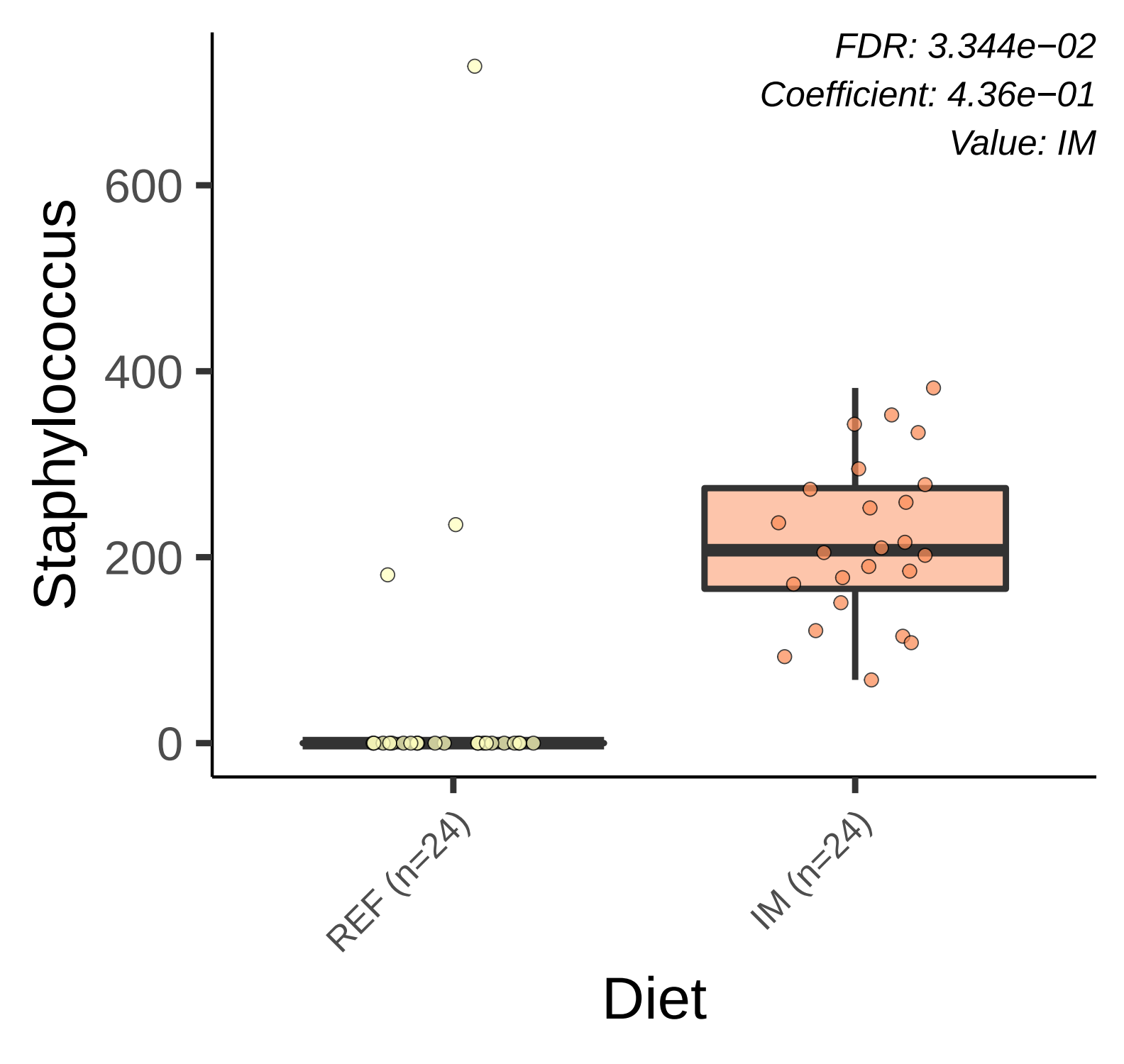
IM (n=24)
Diet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Staphylococcus
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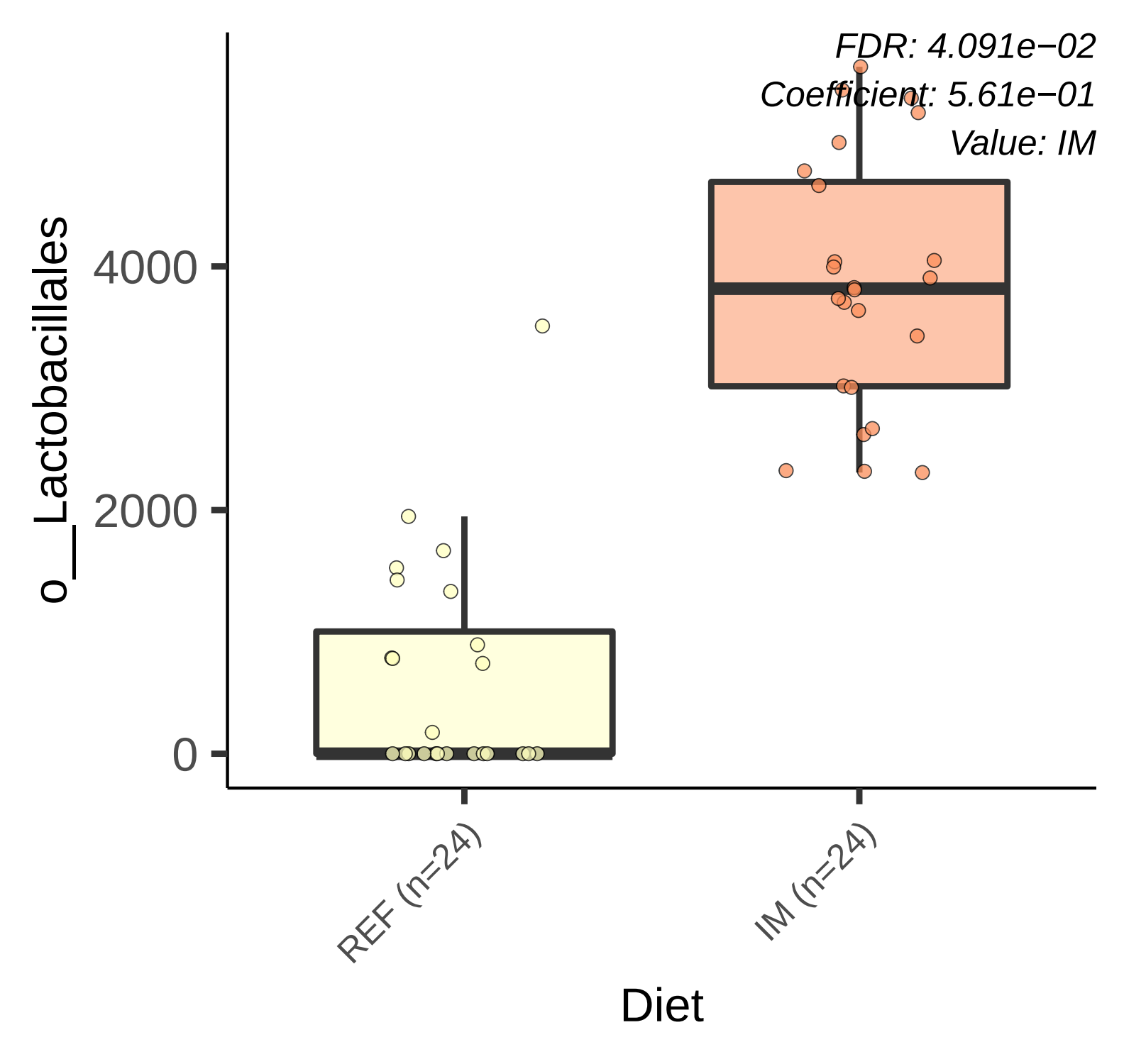
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2000
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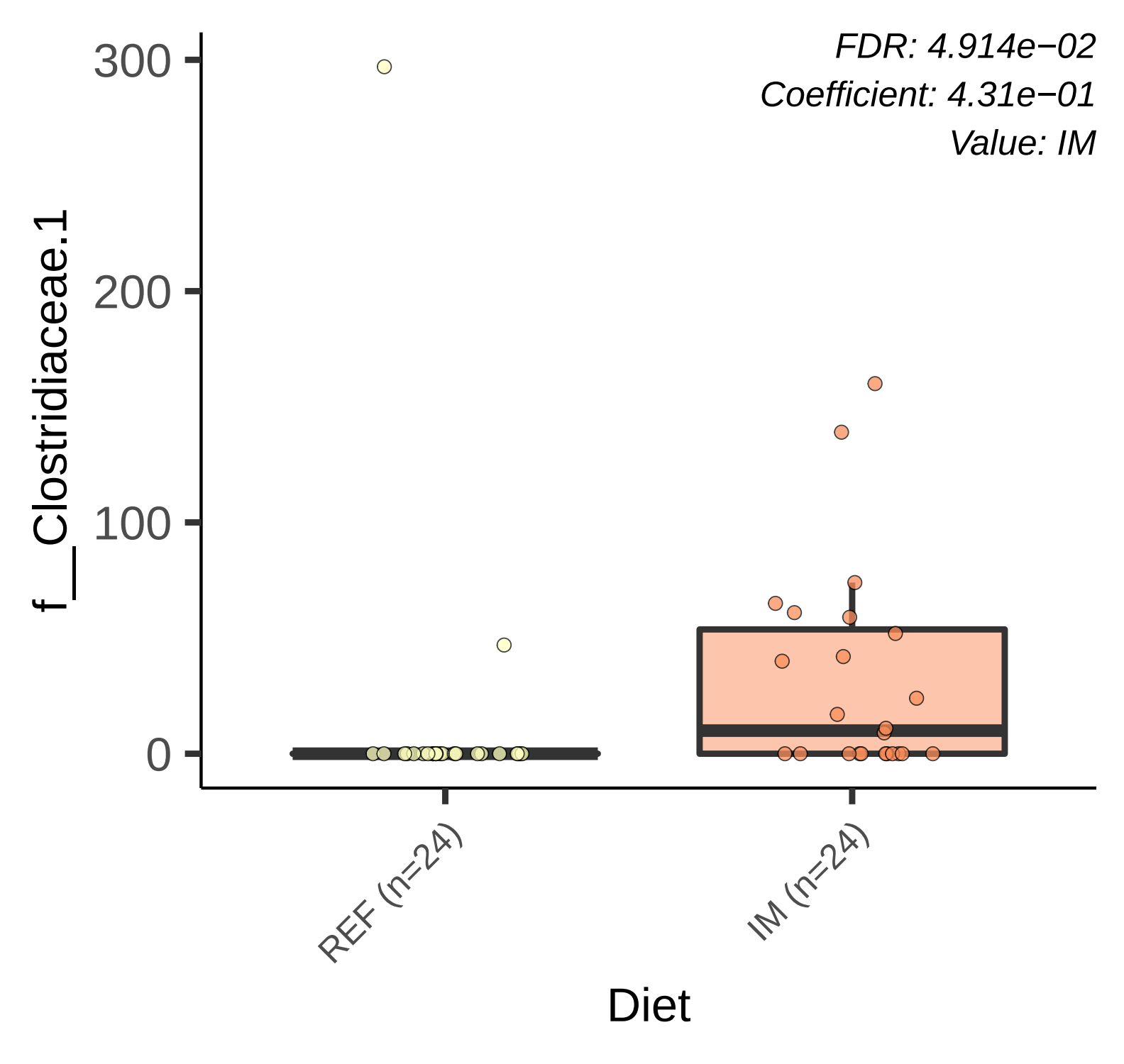
IM (n=24)
Diet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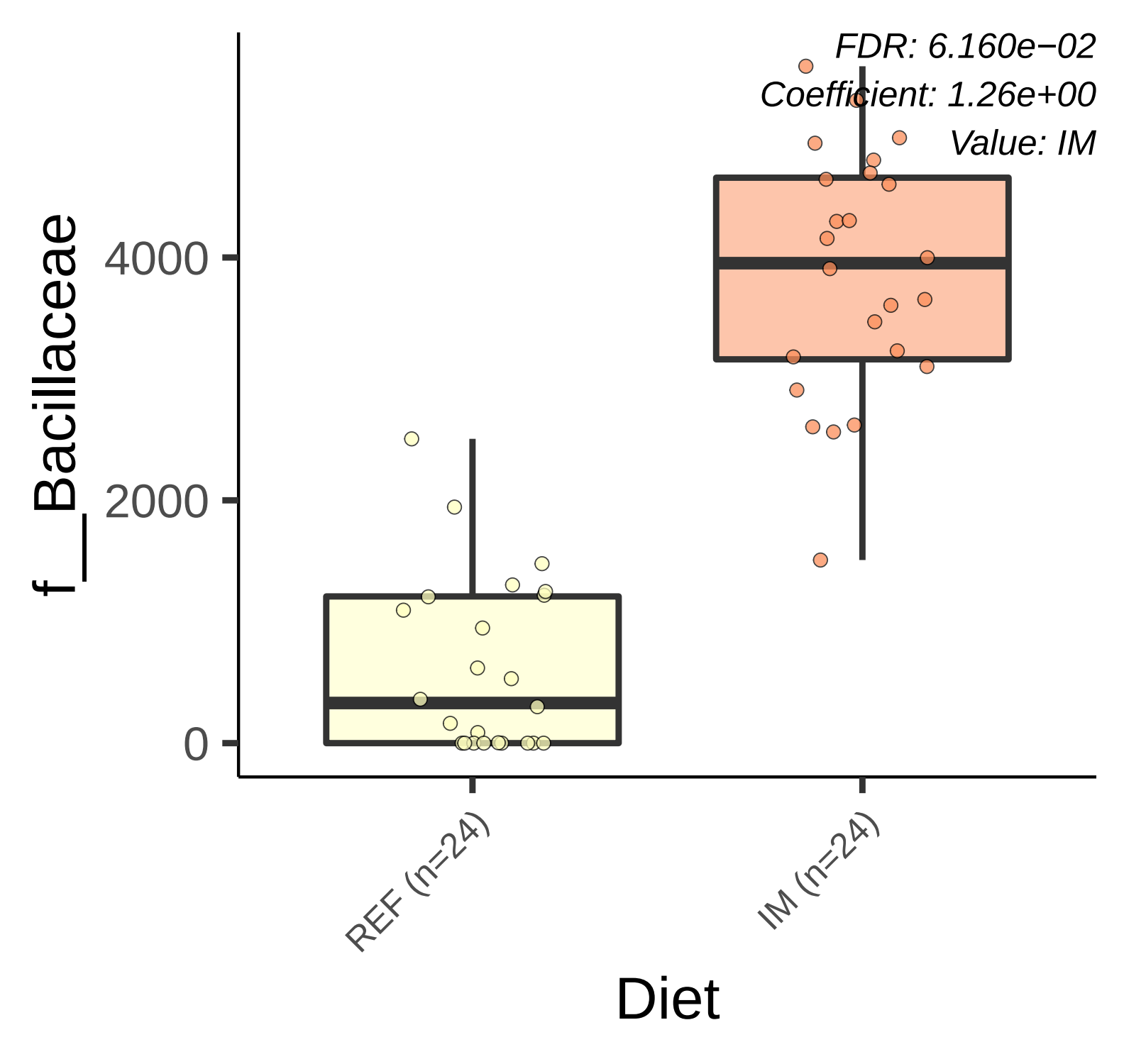
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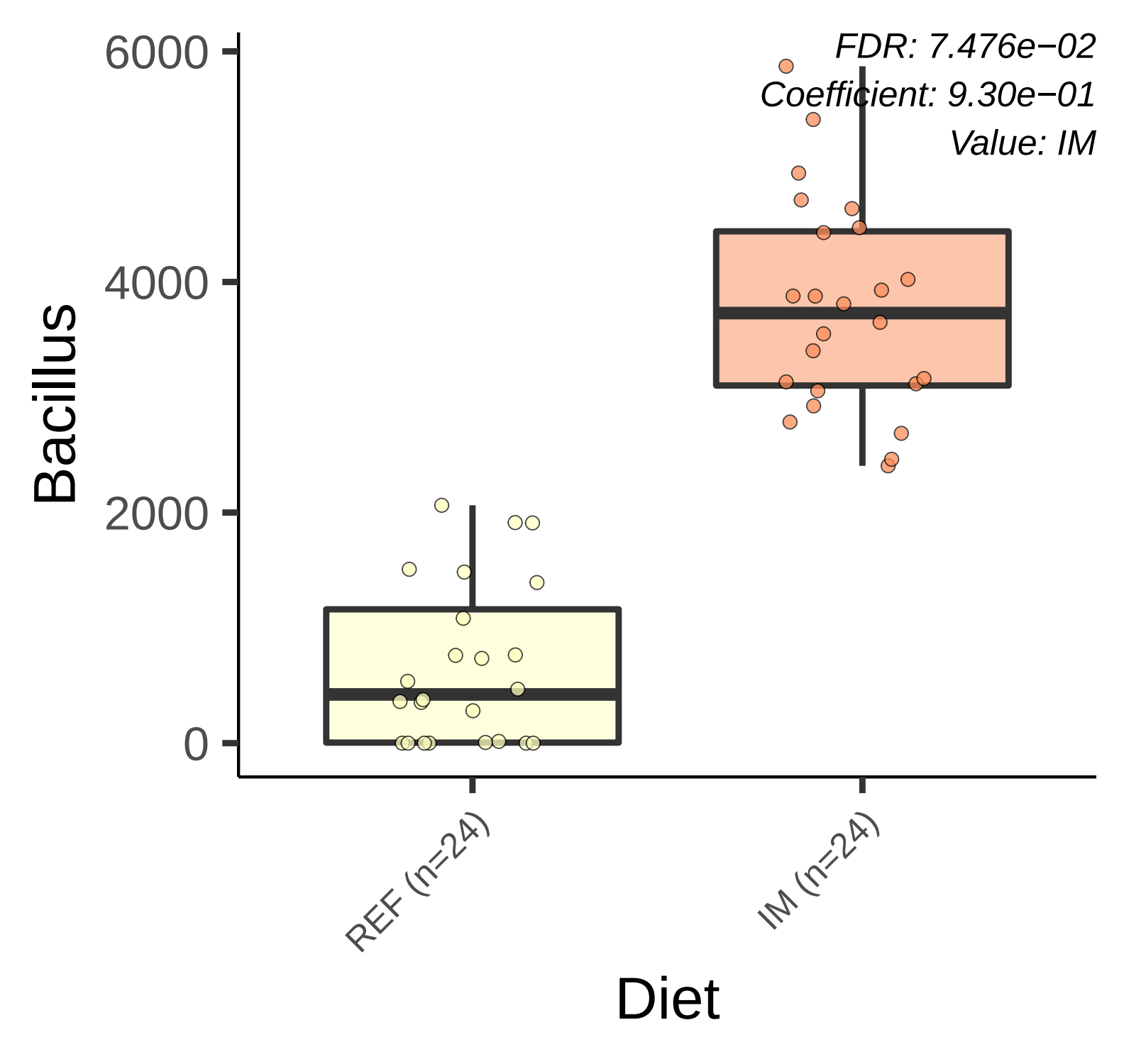
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Bacillus
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IM (n=24)
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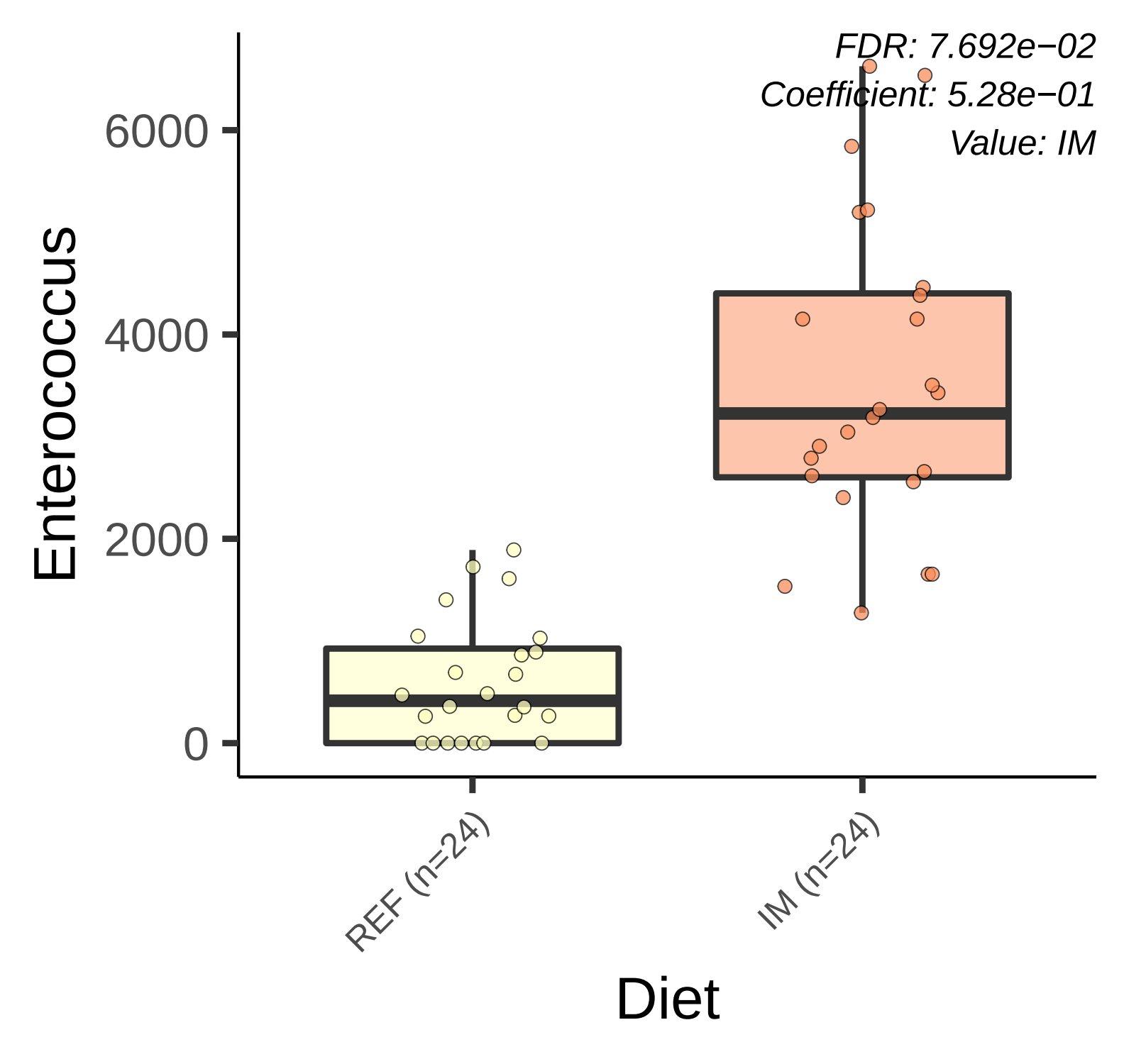
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6000
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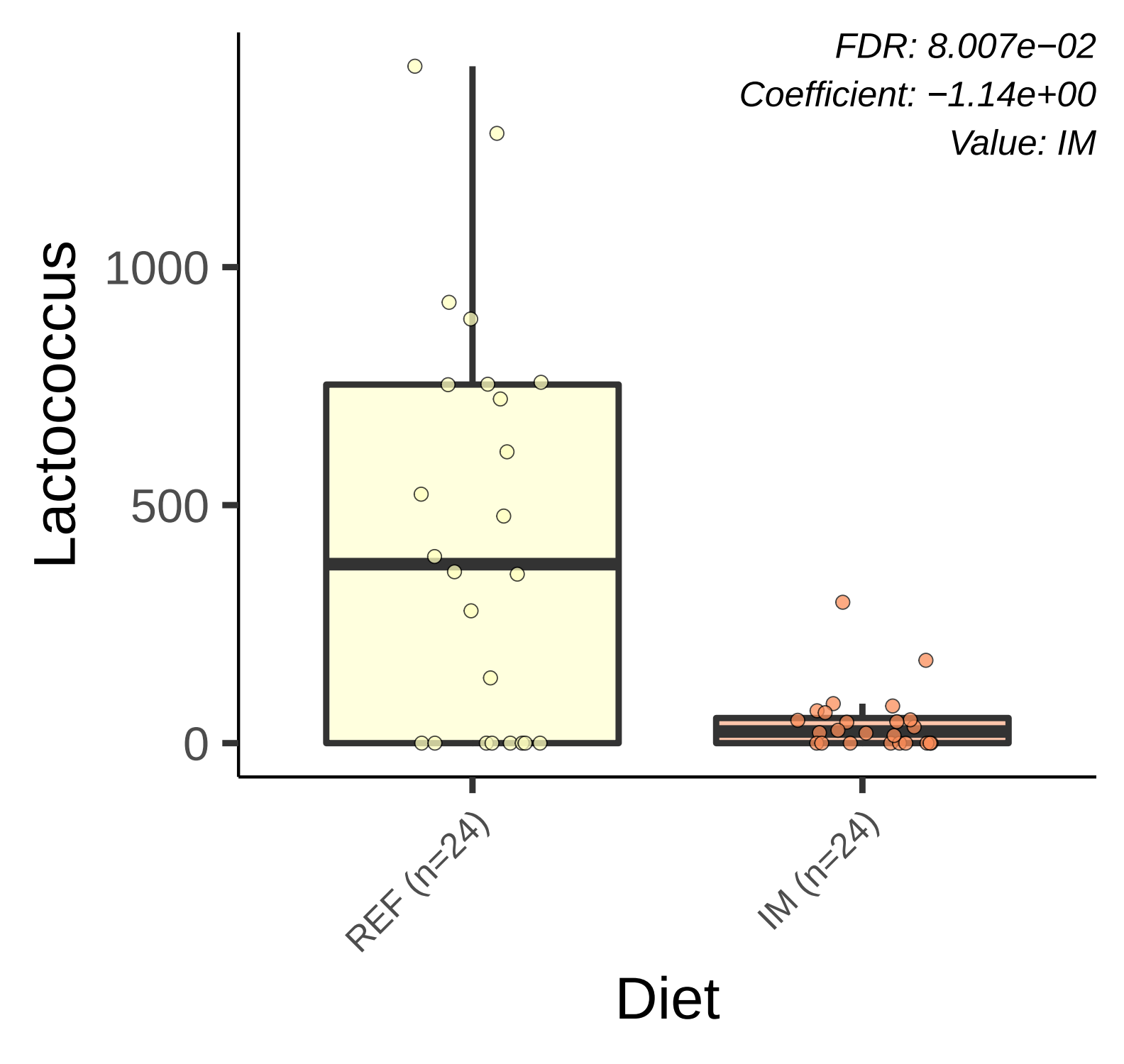
IM (n=24)
Diet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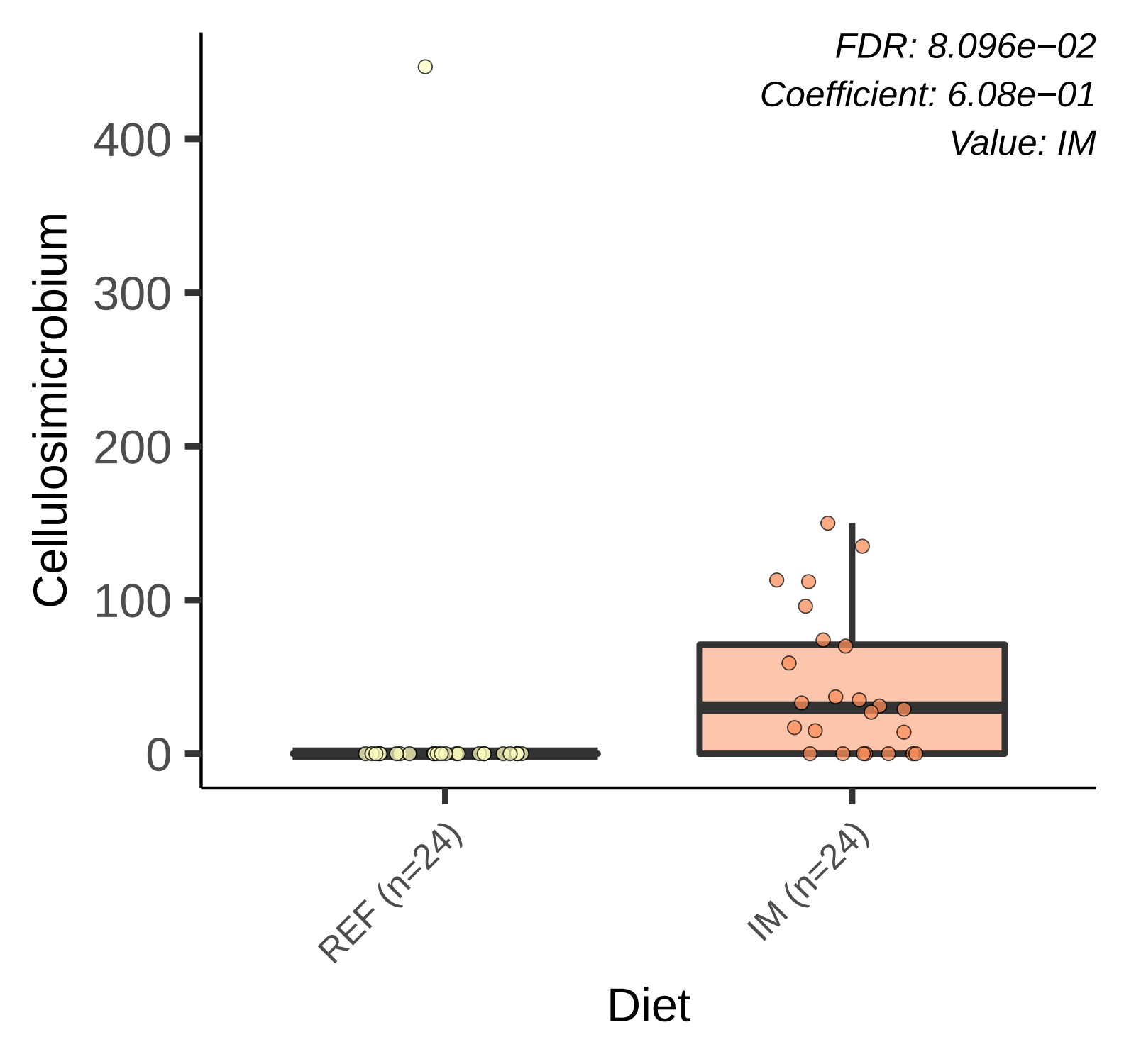
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Cellulosimicrobium
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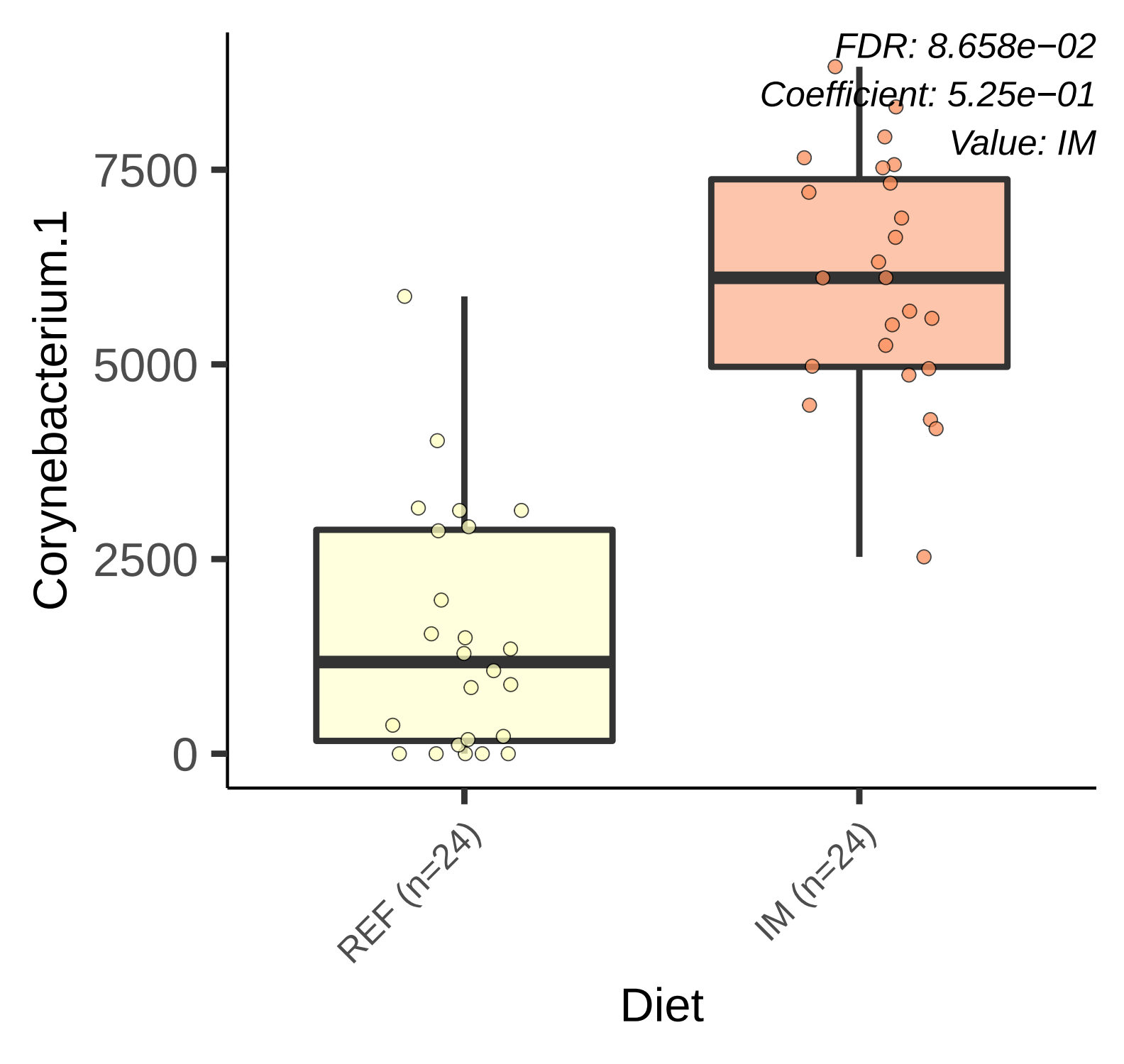
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Corynebacterium.1
7500
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2500
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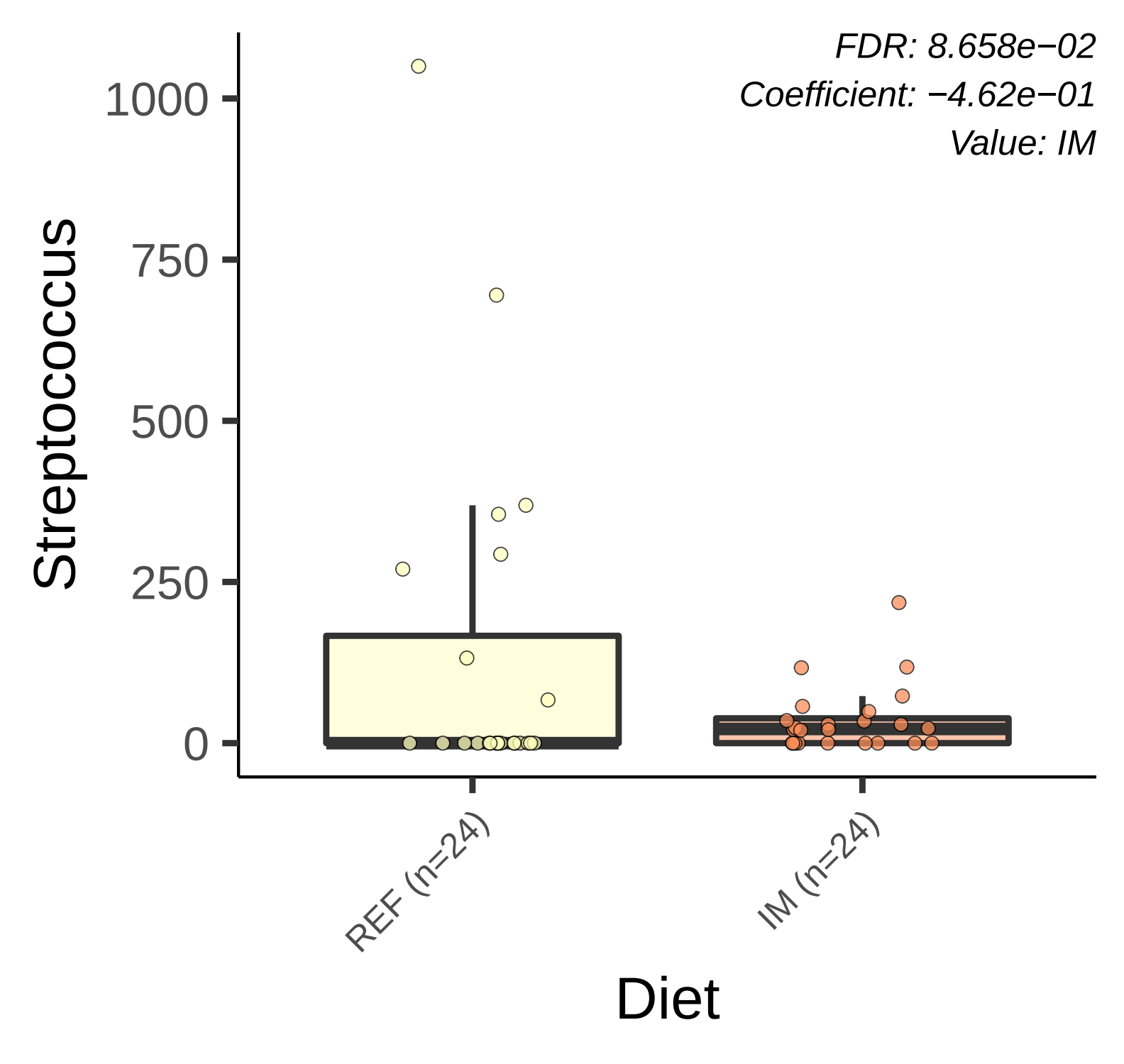
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IM (n=24)
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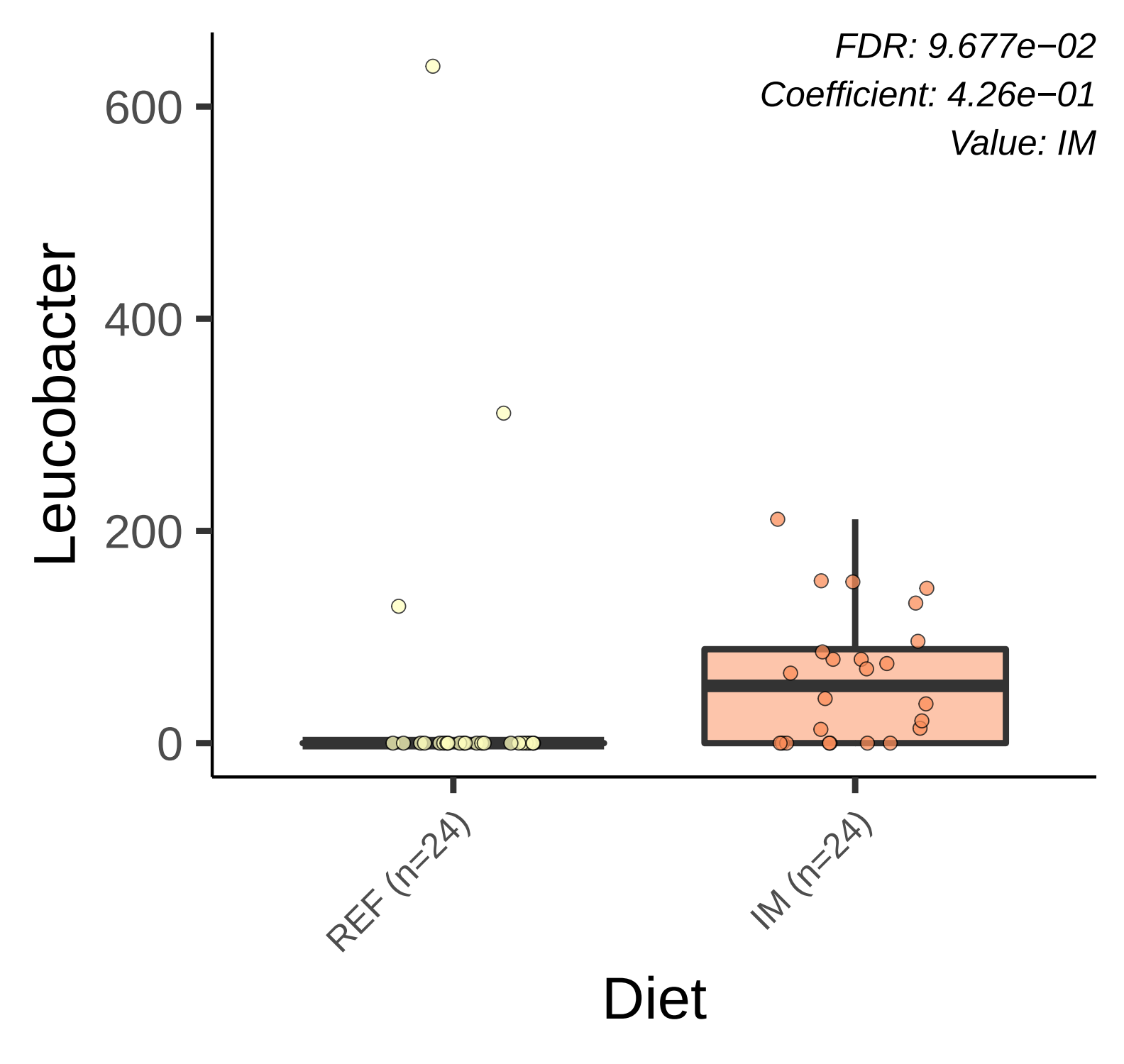
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Coefficient: 4.26e−01
Value: IM
Leucobacter
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400
200
0
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IM (n=24)
Diet