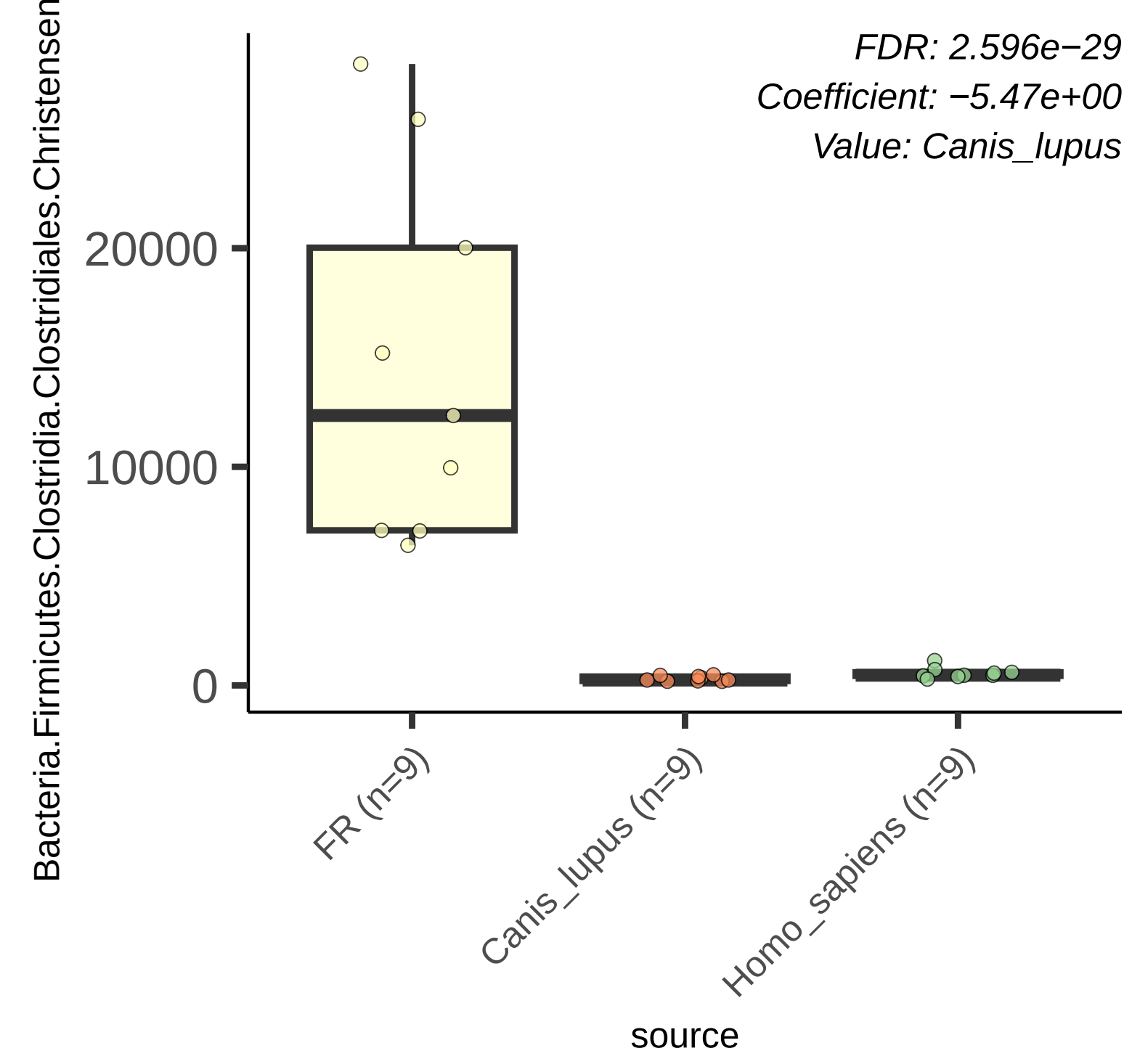
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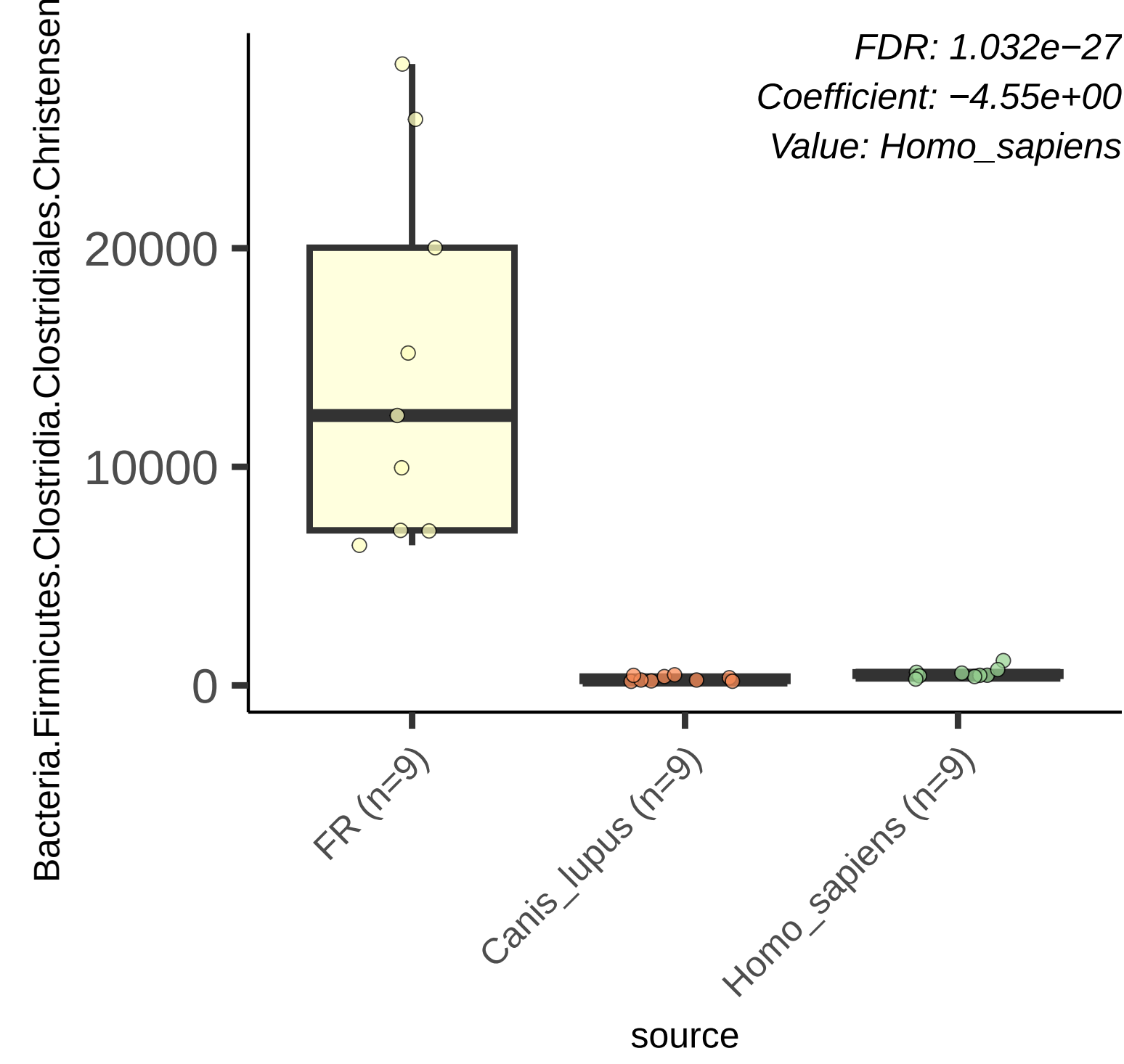
Coefficient: −5.47e+00
Value: Canis_lupus
Bacteria.Firmicutes.Clostridia.Clostridiales.Christensen
20000
10000
0
FR (n=9)
Canis_lupus (n=9)
Homo_sapiens (n=9)
source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Coefficient: −4.55e+00
Value: Homo_sapiens
Bacteria.Firmicutes.Clostridia.Clostridiales.Christensen
20000
10000
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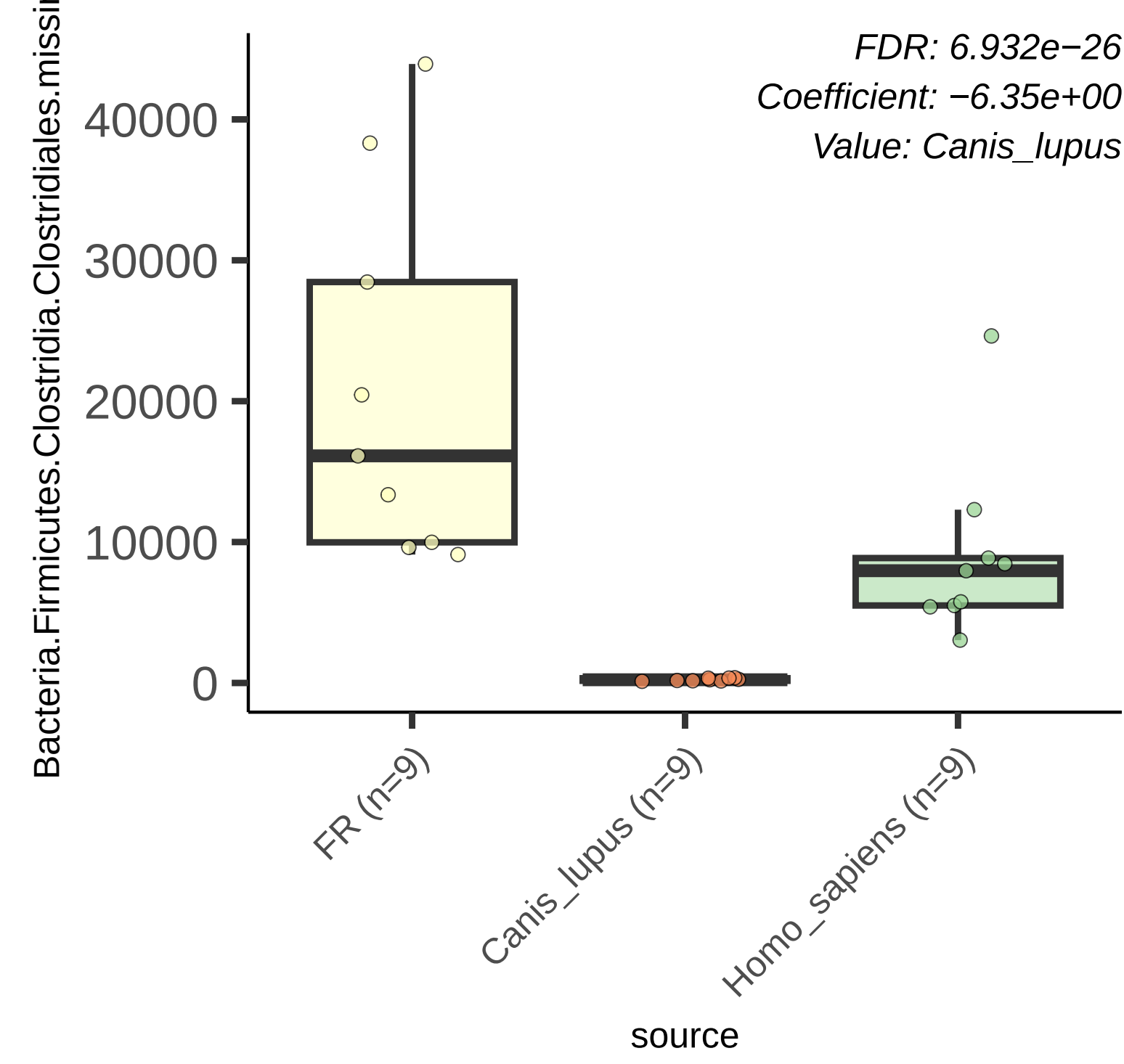
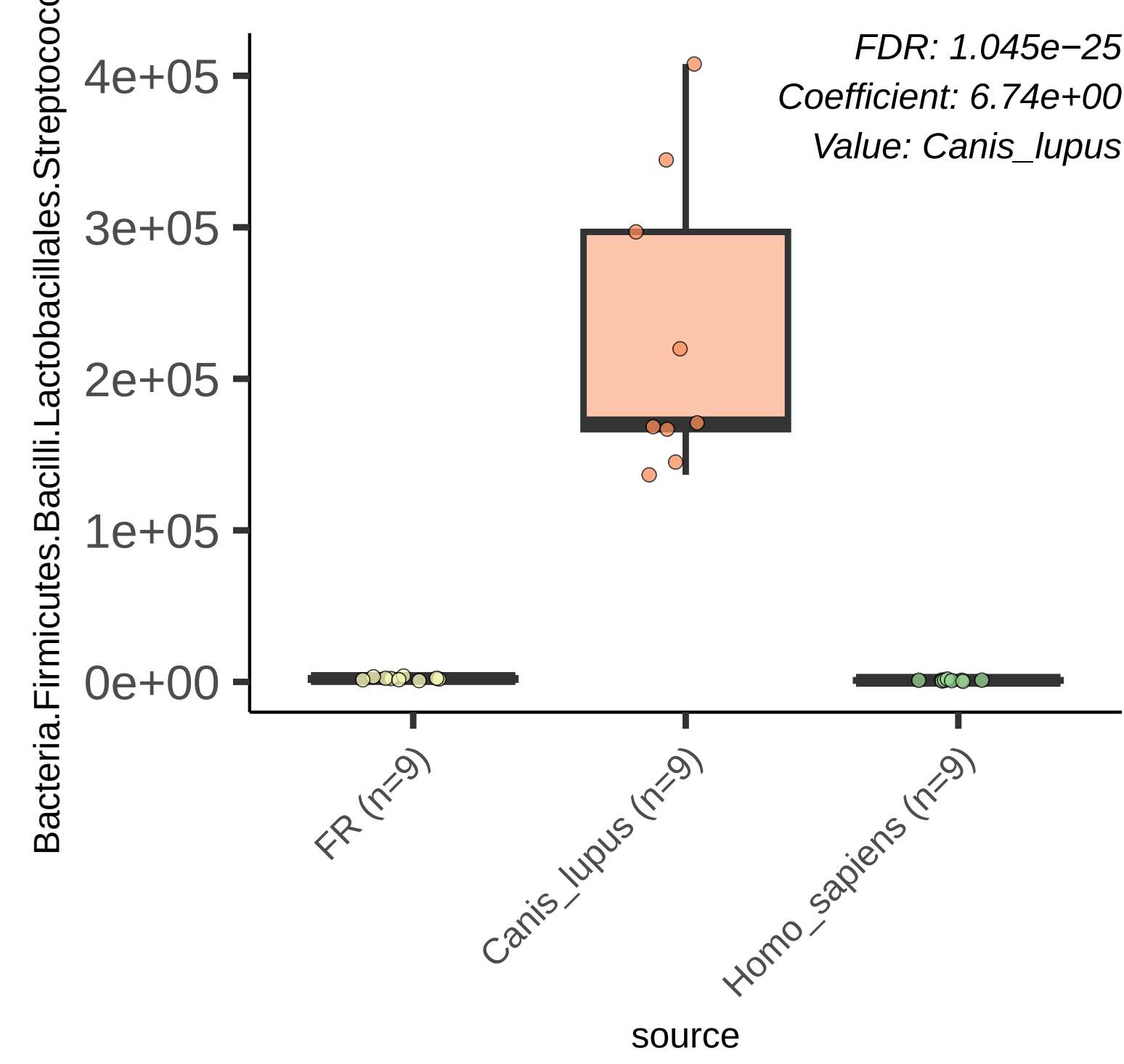
Canis_lupus (n=9)
Homo_sapiens (n=9)
source


FDR: 6.932e−26
Coefficient: −6.35e+00
Value: Canis_lupus
Bacteria.Firmicutes.Clostridia.Clostridiales.missi
40000
30000
20000
10000
0
FR (n=9)
Canis_lupus (n=9)
Homo_sapiens (n=9)
source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Coefficient: 6.74e+00
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Bacteria.Firmicutes.Bacilli.Lactobacillales.Streptococ
4e+05
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2e+05
1e+05
0e+00
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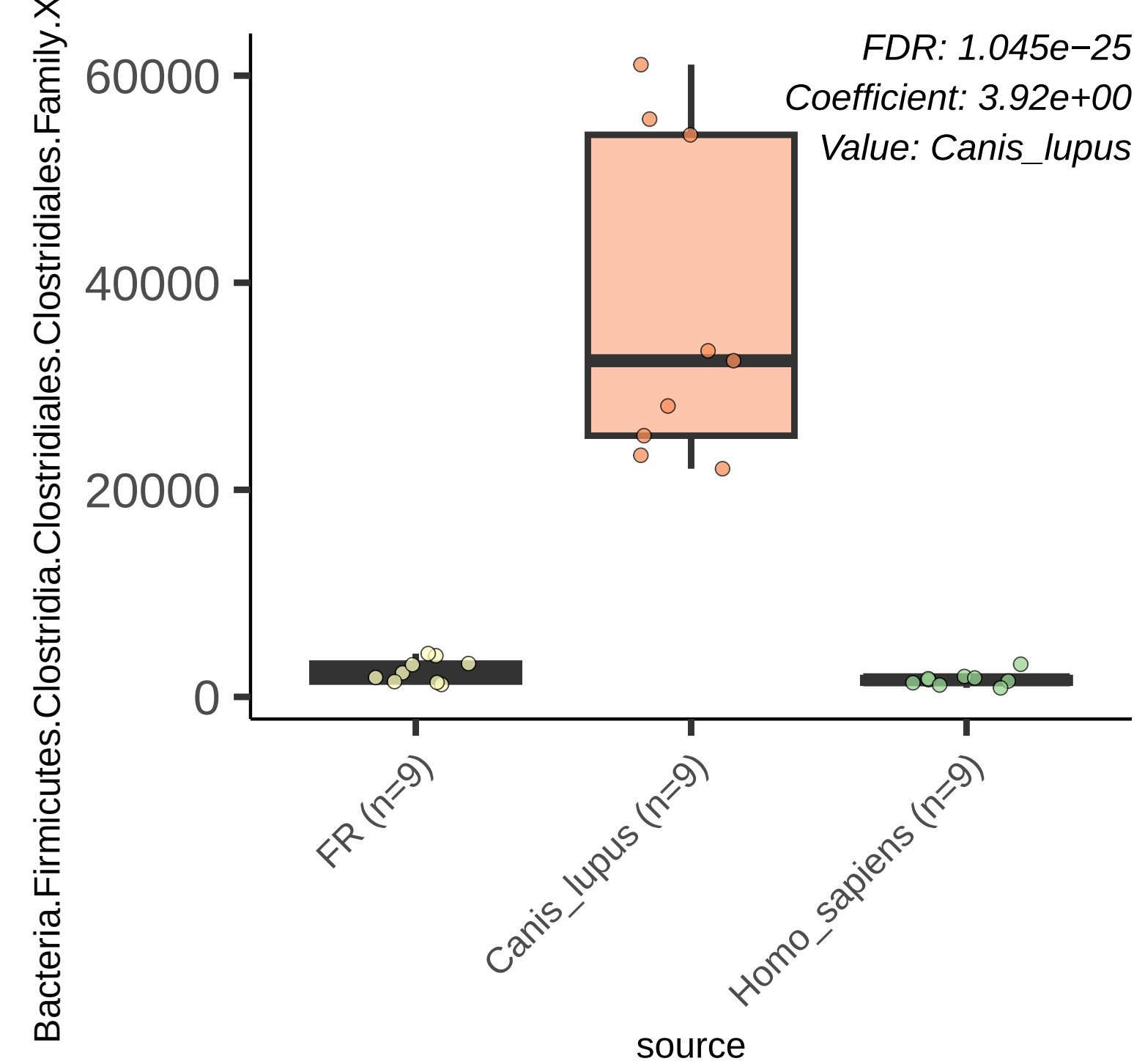
Canis_lupus (n=9)
Homo_sapiens (n=9)
source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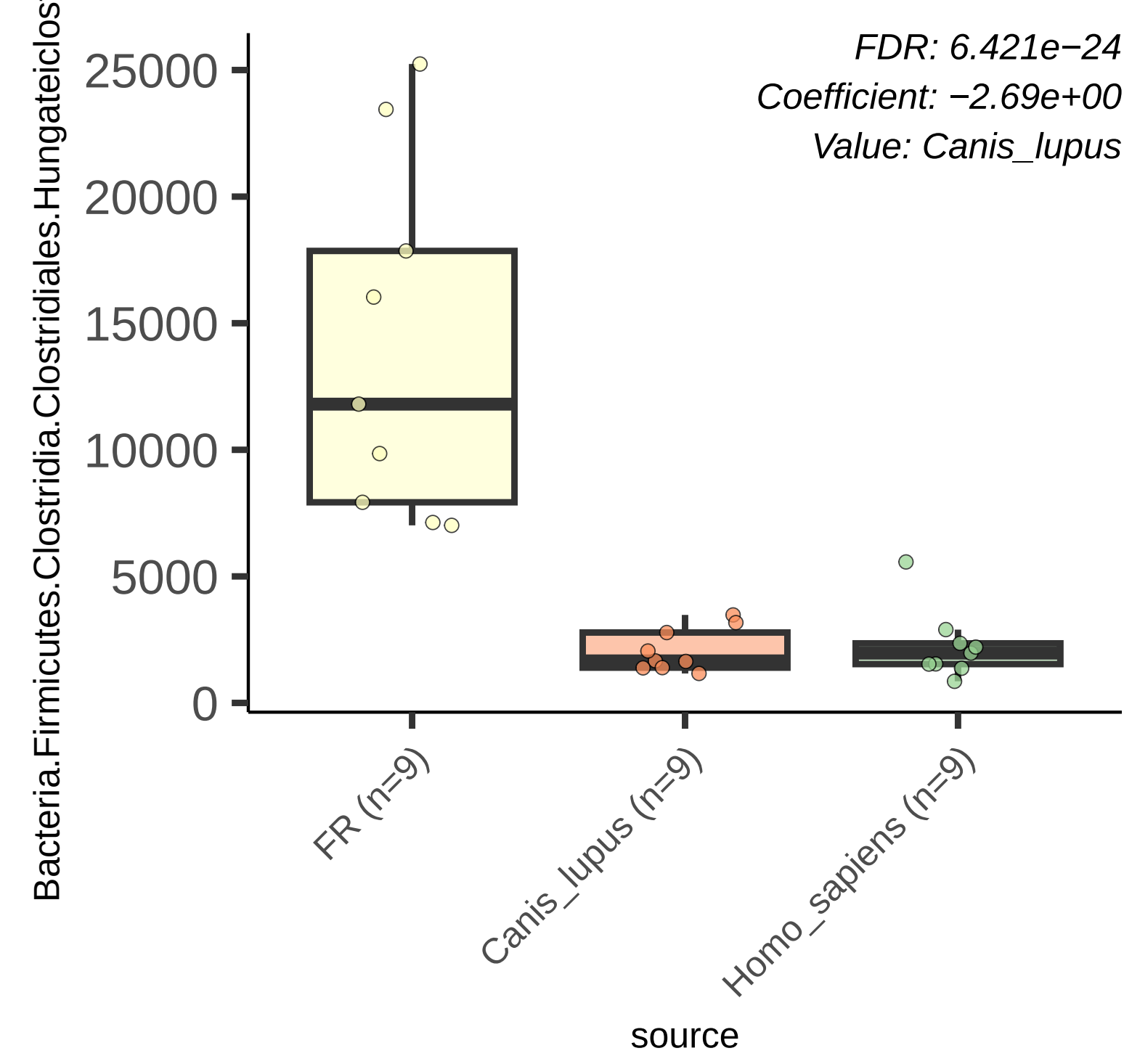
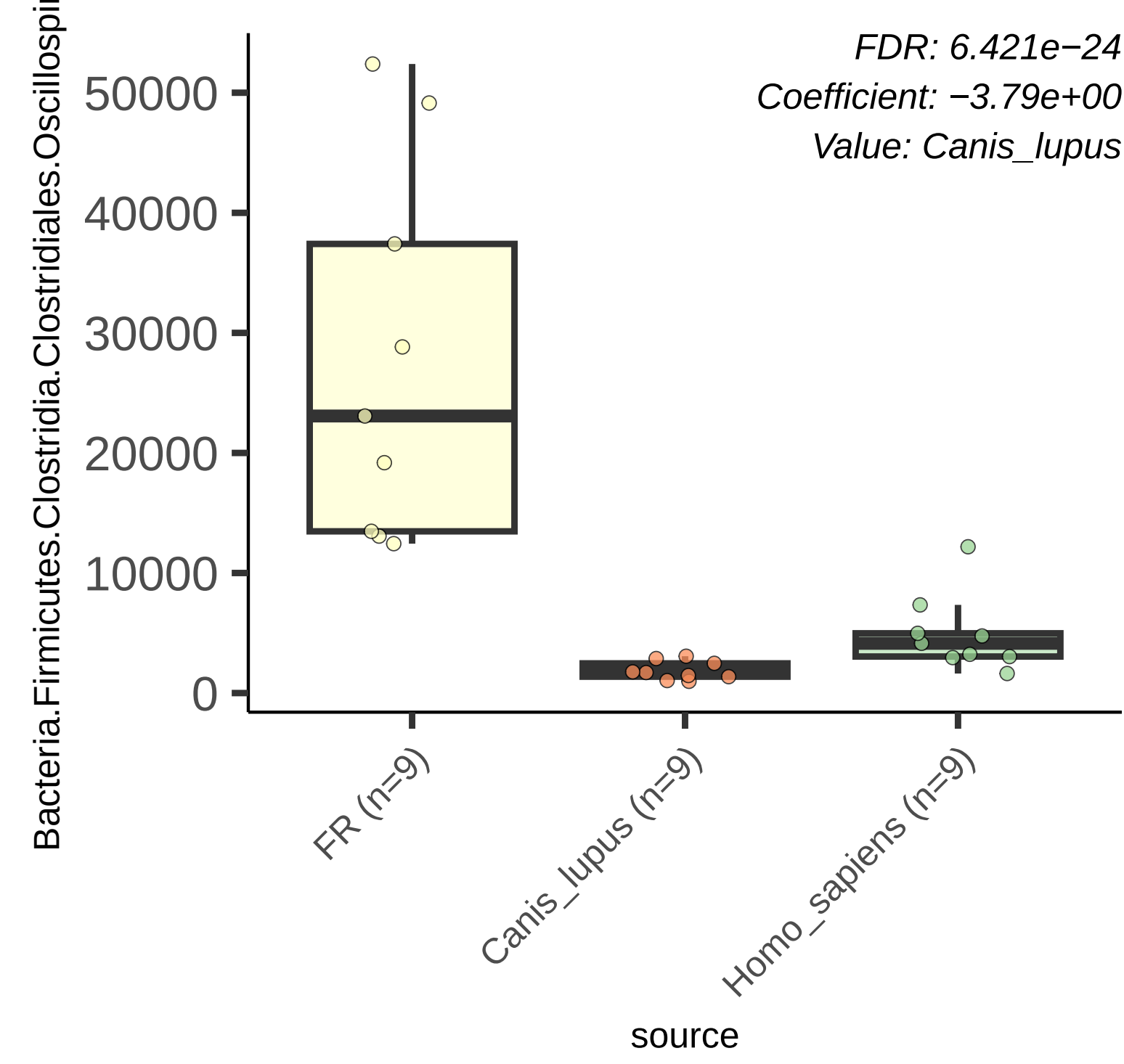
Coefficient: 3.92e+00
Value: Canis_lupus
60000
40000
20000
0
Bacteria.Firmicutes.Clostridia.Clostridiales.Clostridiales.Family.X
FR (n=9)
Canis_lupus (n=9)
Homo_sapiens (n=9)
source


FDR: 6.421e−24
Coefficient: −2.69e+00
Value: Canis_lupus
Bacteria.Firmicutes.Clostridia.Clostridiales.Hungateiclos
25000
20000
15000
10000
5000
0
FR (n=9)
Canis_lupus (n=9)
Homo_sapiens (n=9)
source


FDR: 6.421e−24
Coefficient: −3.79e+00
Value: Canis_lupus
Bacteria.Firmicutes.Clostridia.Clostridiales.Oscillospi
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40000
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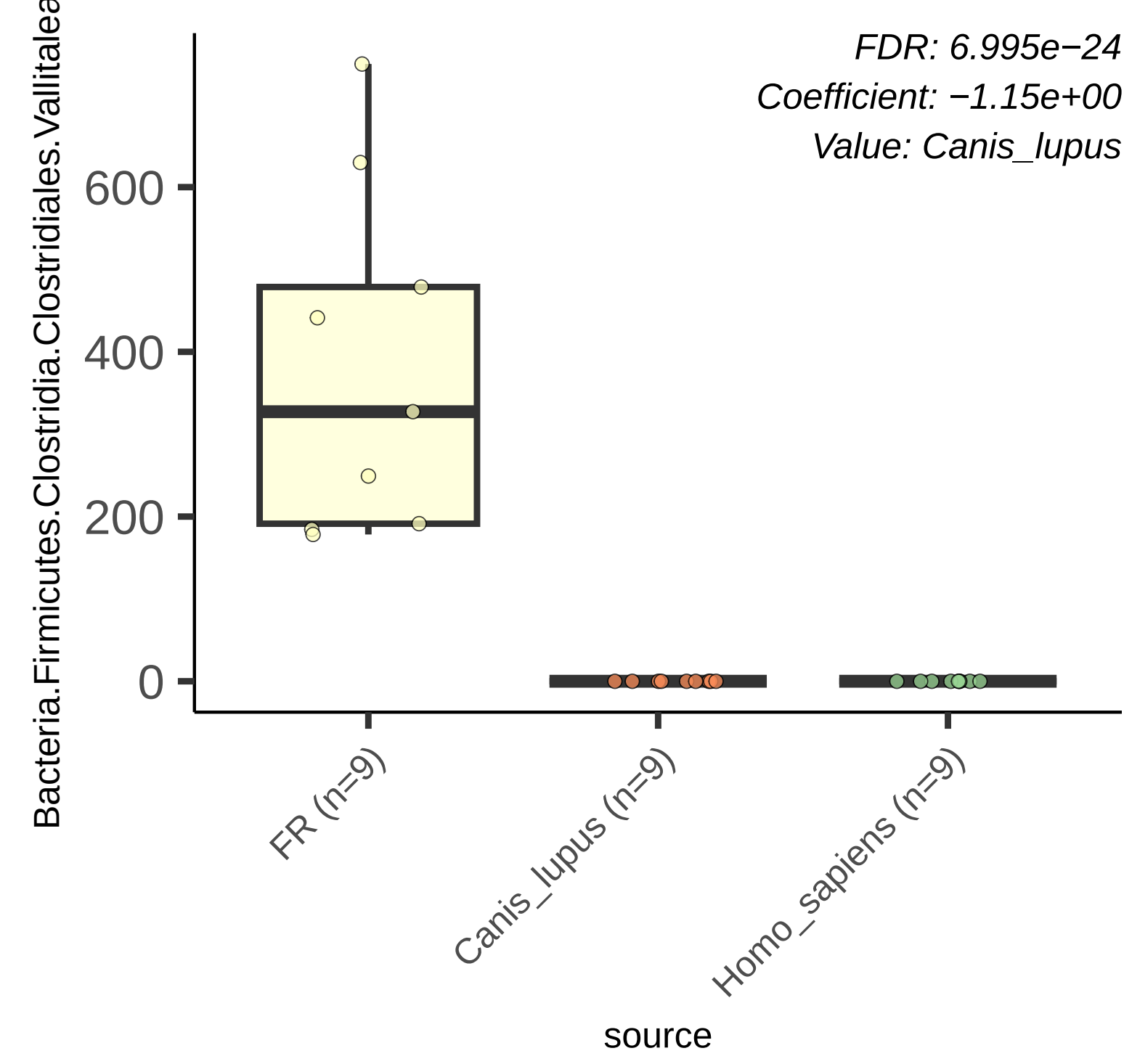
20000
10000
0
FR (n=9)
Canis_lupus (n=9)
Homo_sapiens (n=9)
source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Coefficient: −1.15e+00
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Bacteria.Firmicutes.Clostridia.Clostridiales.Vallitalea
600
400
200
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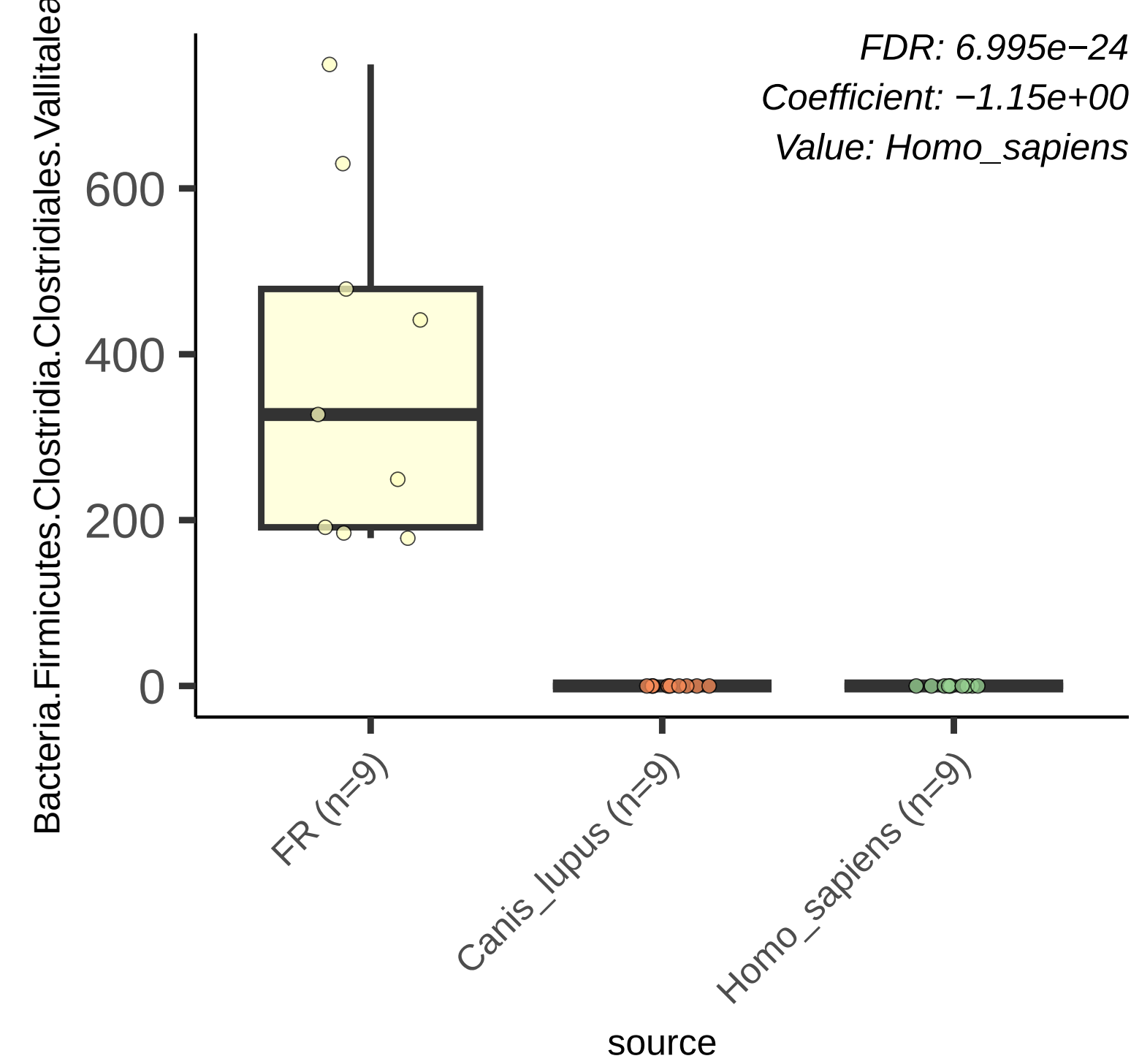
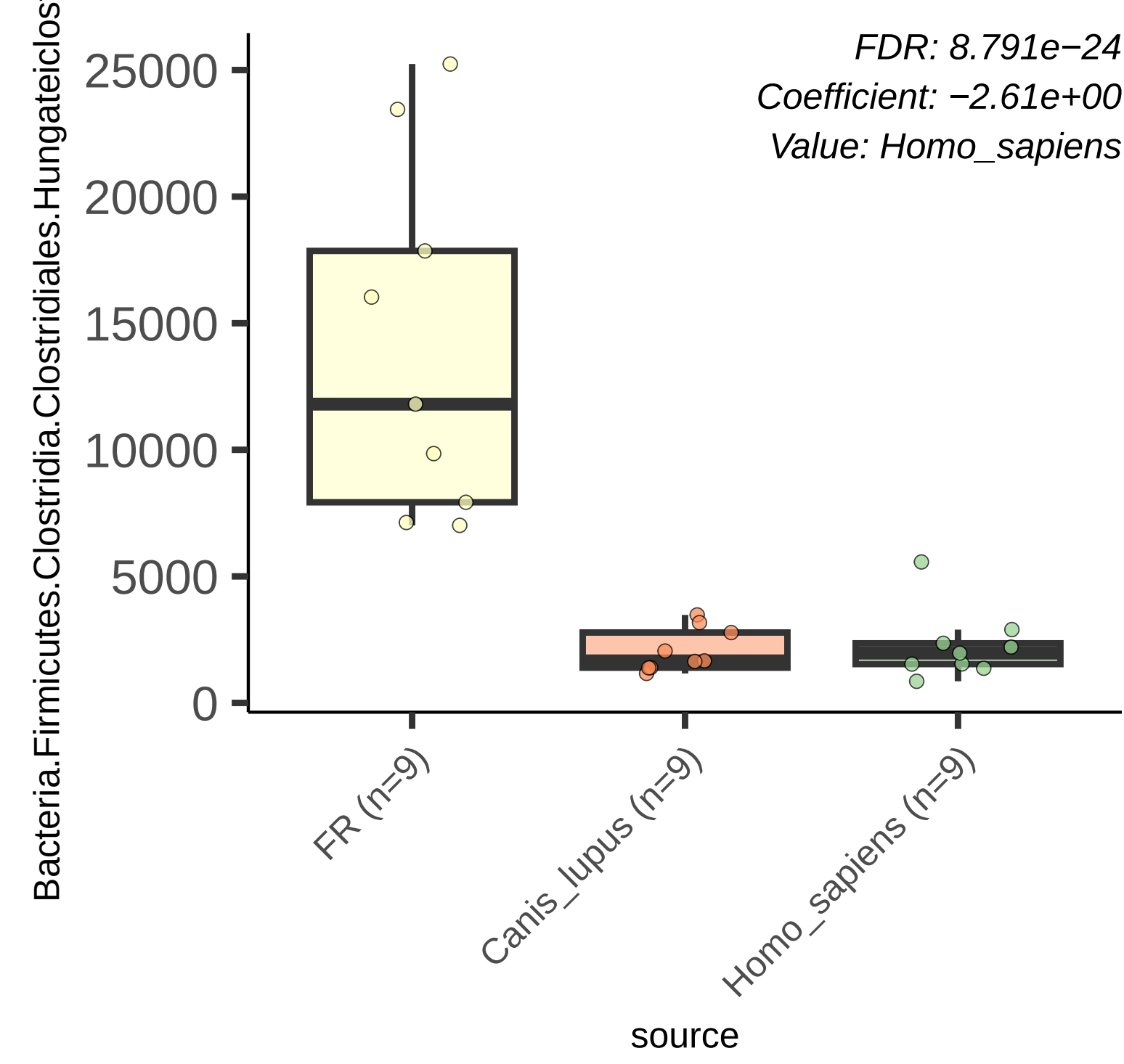
Canis_lupus (n=9)
Homo_sapiens (n=9)
source


FDR: 6.995e−24
Coefficient: −1.15e+00
Value: Homo_sapiens
Bacteria.Firmicutes.Clostridia.Clostridiales.Vallitalea
600
400
200
0
FR (n=9)
Canis_lupus (n=9)
Homo_sapiens (n=9)
source


FDR: 8.791e−24
Coefficient: −2.61e+00
Value: Homo_sapiens
Bacteria.Firmicutes.Clostridia.Clostridiales.Hungateiclos
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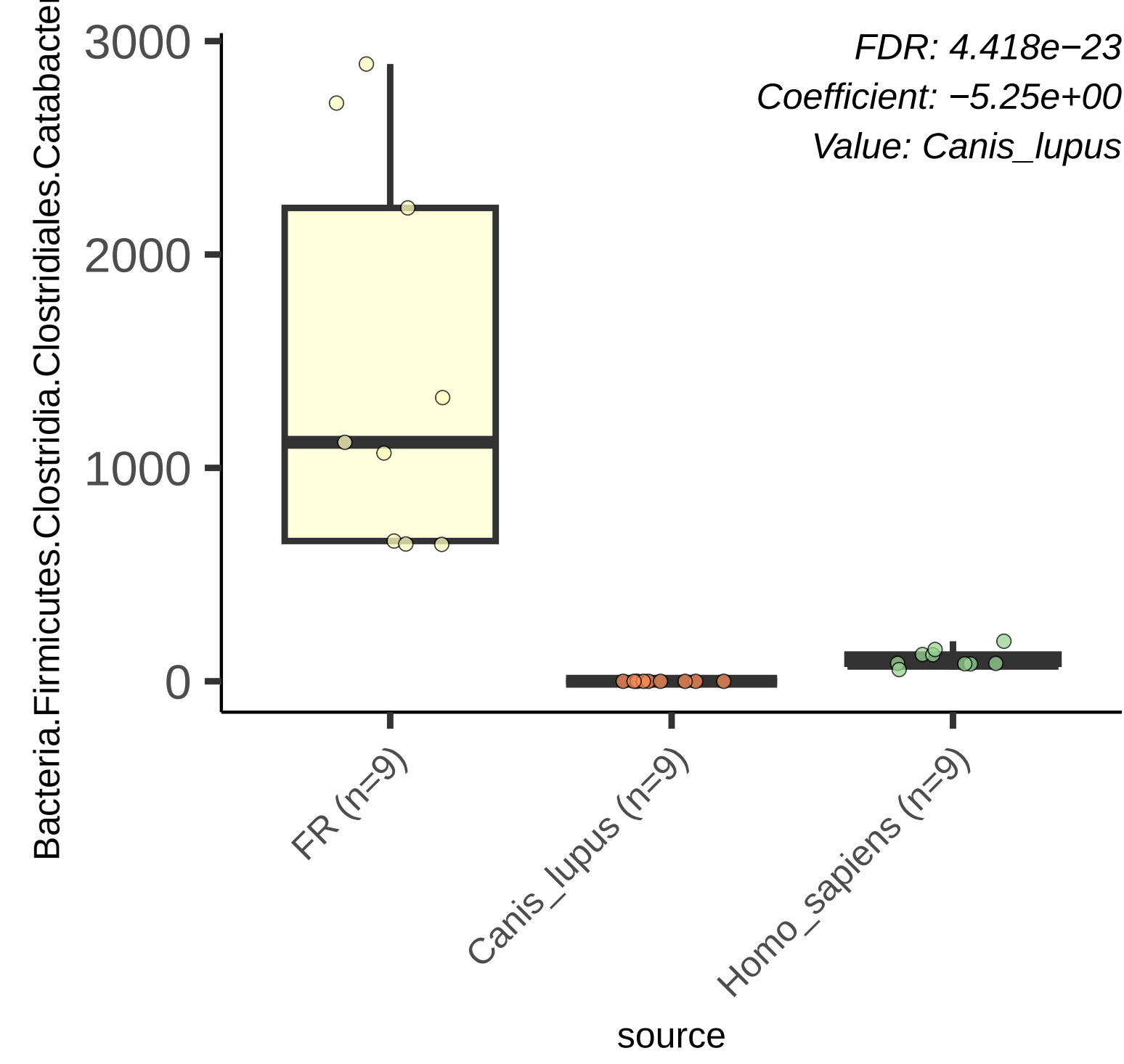
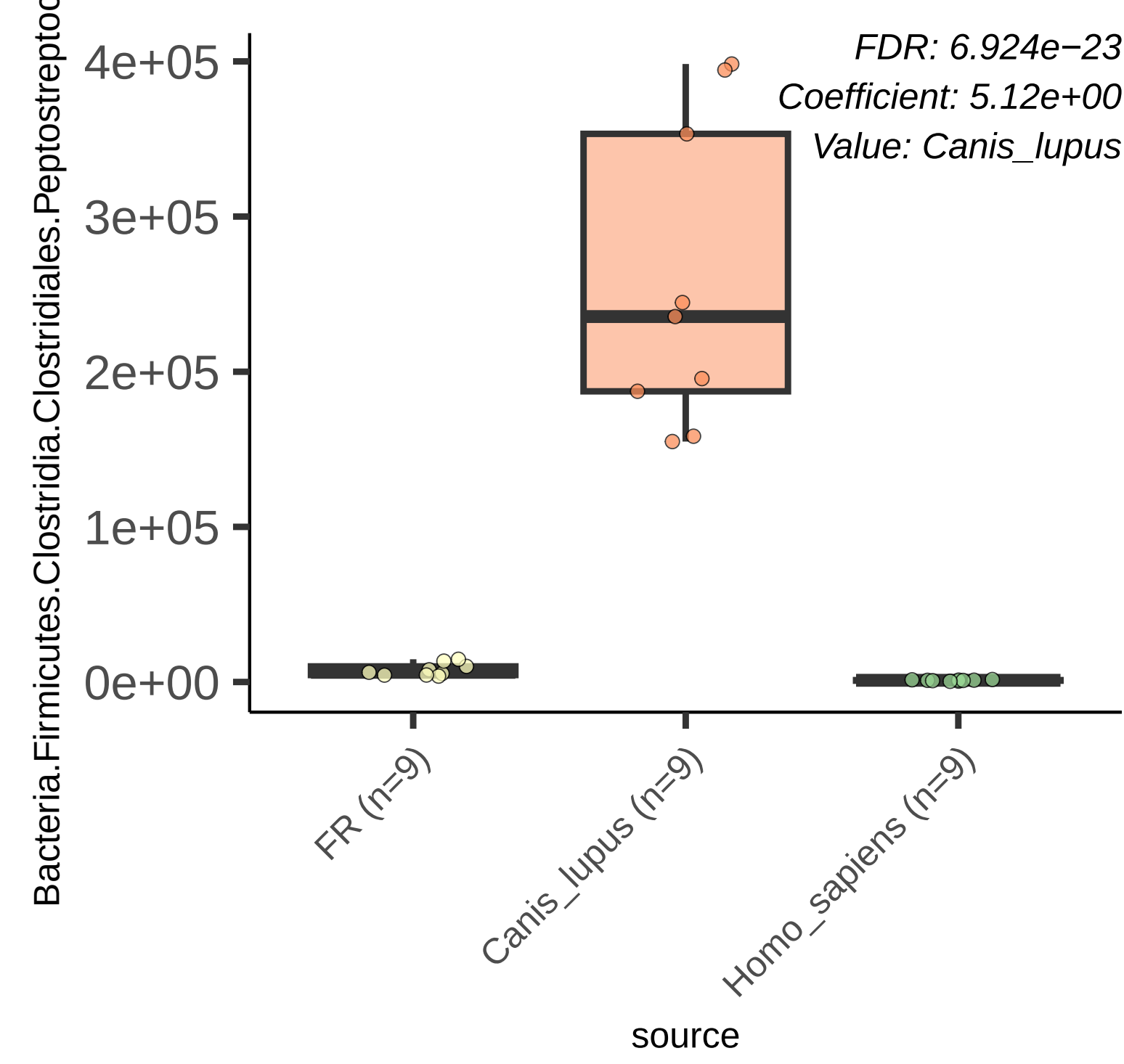
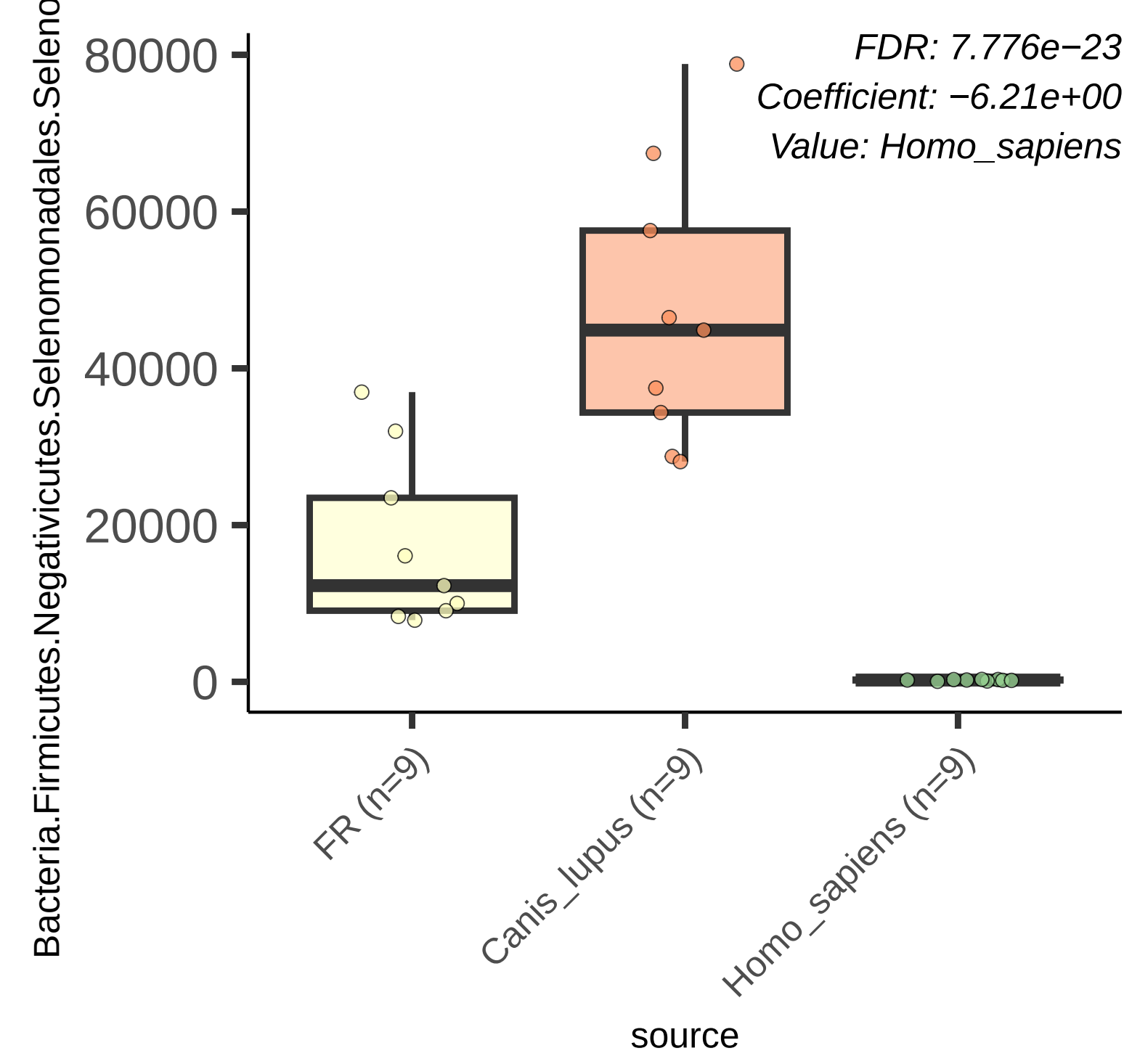
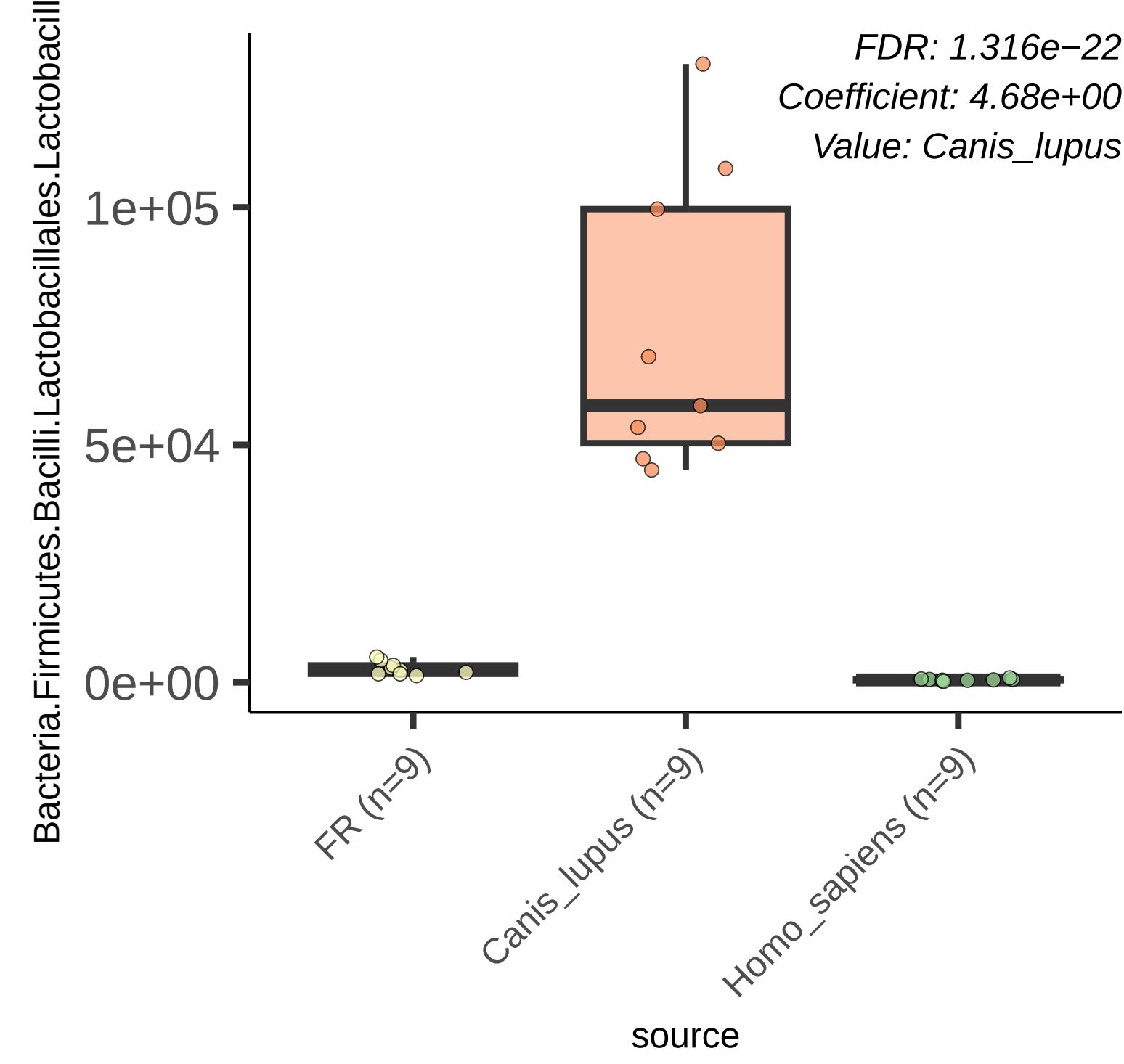
20000
15000
10000
5000
0
FR (n=9)
Canis_lupus (n=9)
Homo_sapiens (n=9)
source


FDR: 4.418e−23
Coefficient: −5.25e+00
Value: Canis_lupus
3000
2000
1000
0
Bacteria.Firmicutes.Clostridia.Clostridiales.Catabacter
FR (n=9)
Canis_lupus (n=9)
Homo_sapiens (n=9)
source


FDR: 6.924e−23
Coefficient: 5.12e+00
Value: Canis_lupus
Bacteria.Firmicutes.Clostridia.Clostridiales.Peptostreptoc
4e+05
3e+05
2e+05
1e+05
0e+00
FR (n=9)
Canis_lupus (n=9)
Homo_sapiens (n=9)
source


FDR: 7.776e−23
Coefficient: −6.21e+00
Value: Homo_sapiens
Bacteria.Firmicutes.Negativicutes.Selenomonadales.Seleno
80000
60000
40000
20000
0
FR (n=9)
Canis_lupus (n=9)
Homo_sapiens (n=9)
source


FDR: 1.316e−22
Coefficient: 4.68e+00
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Bacteria.Firmicutes.Bacilli.Lactobacillales.Lactobacill
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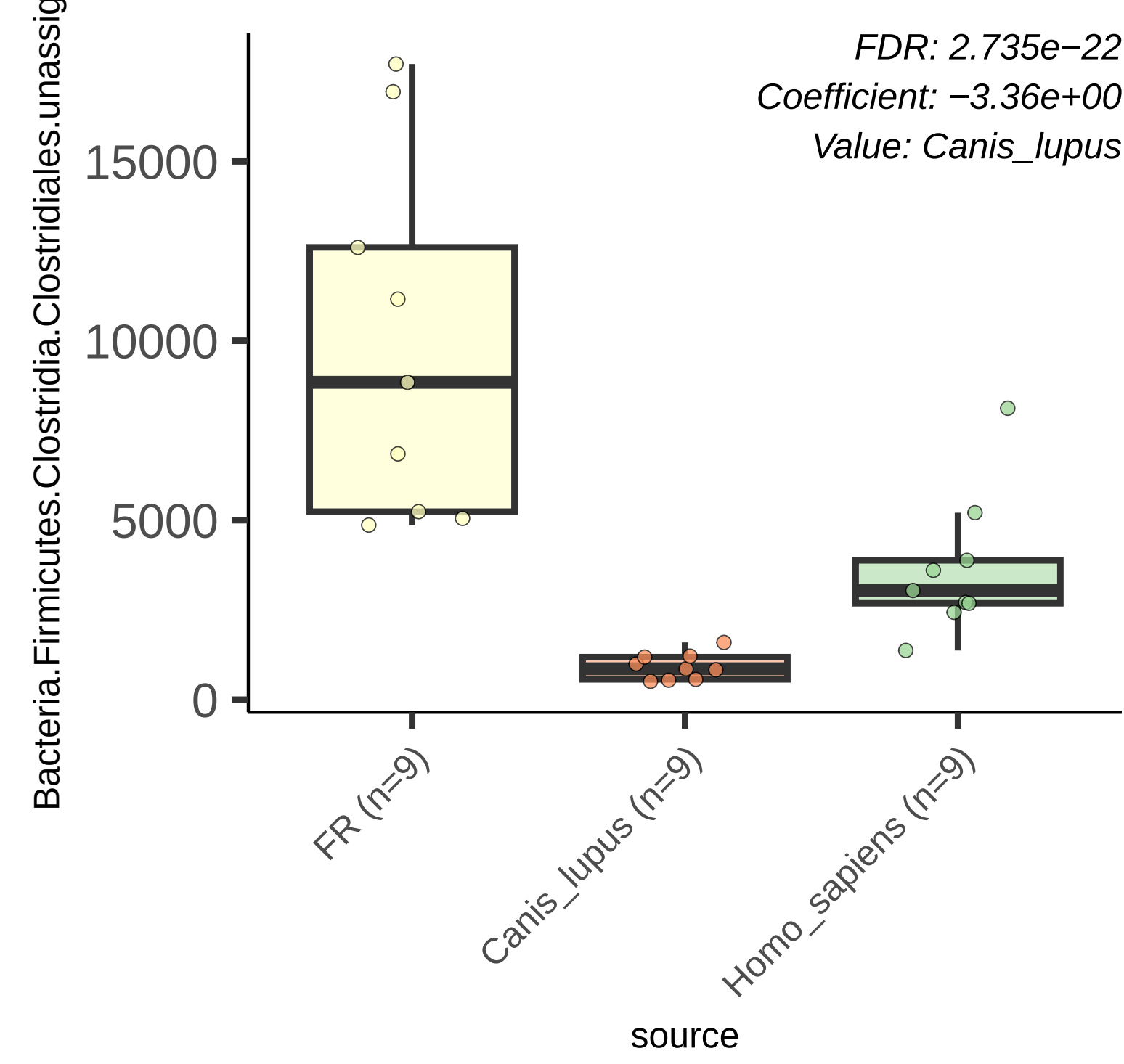
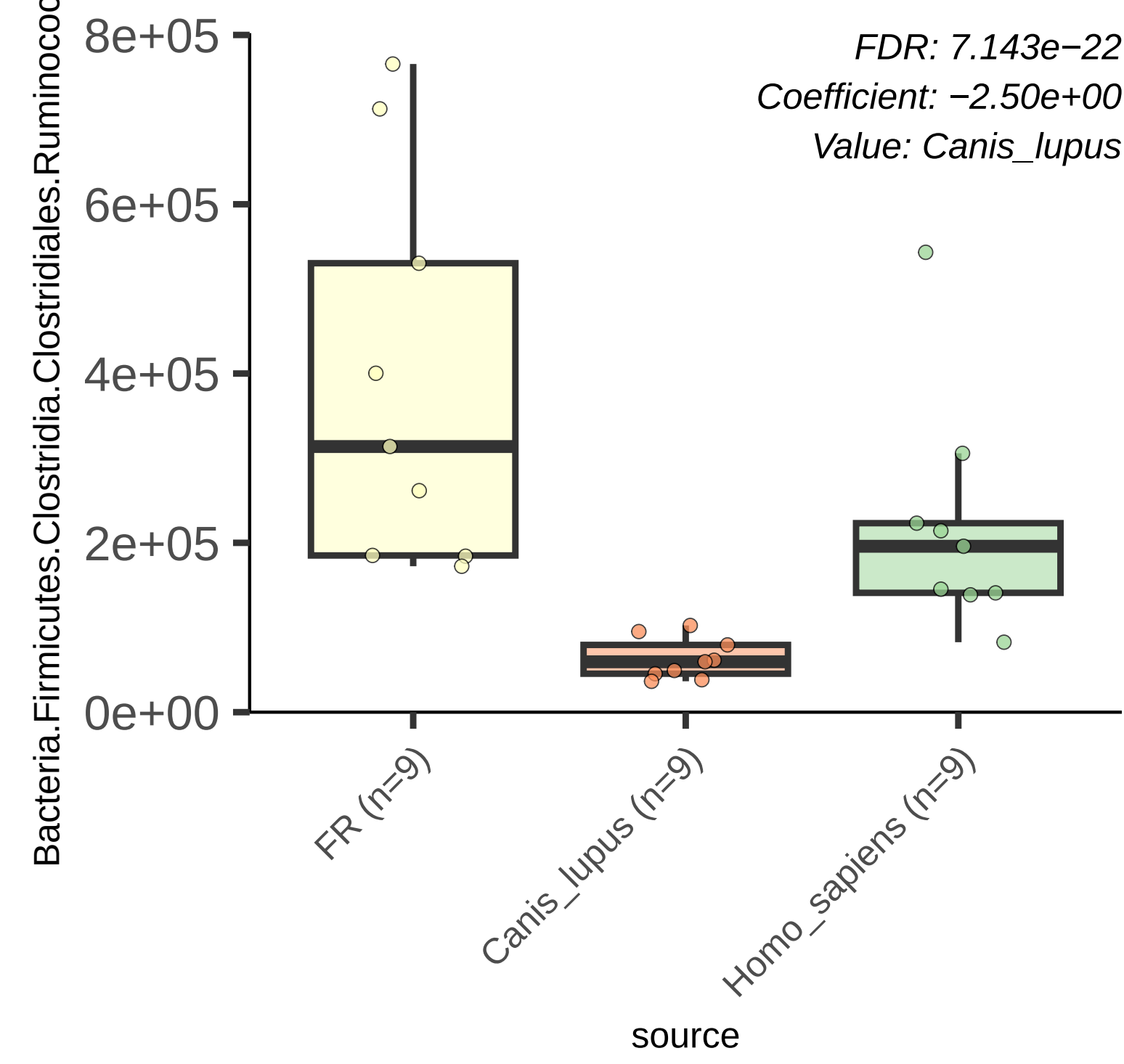
5e+04
0e+00
FR (n=9)
Canis_lupus (n=9)
Homo_sapiens (n=9)
source


FDR: 2.735e−22
Coefficient: −3.36e+00
Value: Canis_lupus
Bacteria.Firmicutes.Clostridia.Clostridiales.unassig
15000
10000
5000
0
FR (n=9)
Canis_lupus (n=9)
Homo_sapiens (n=9)
source


FDR: 7.143e−22
Coefficient: −2.50e+00
Value: Canis_lupus
Bacteria.Firmicutes.Clostridia.Clostridiales.Ruminococ
8e+05
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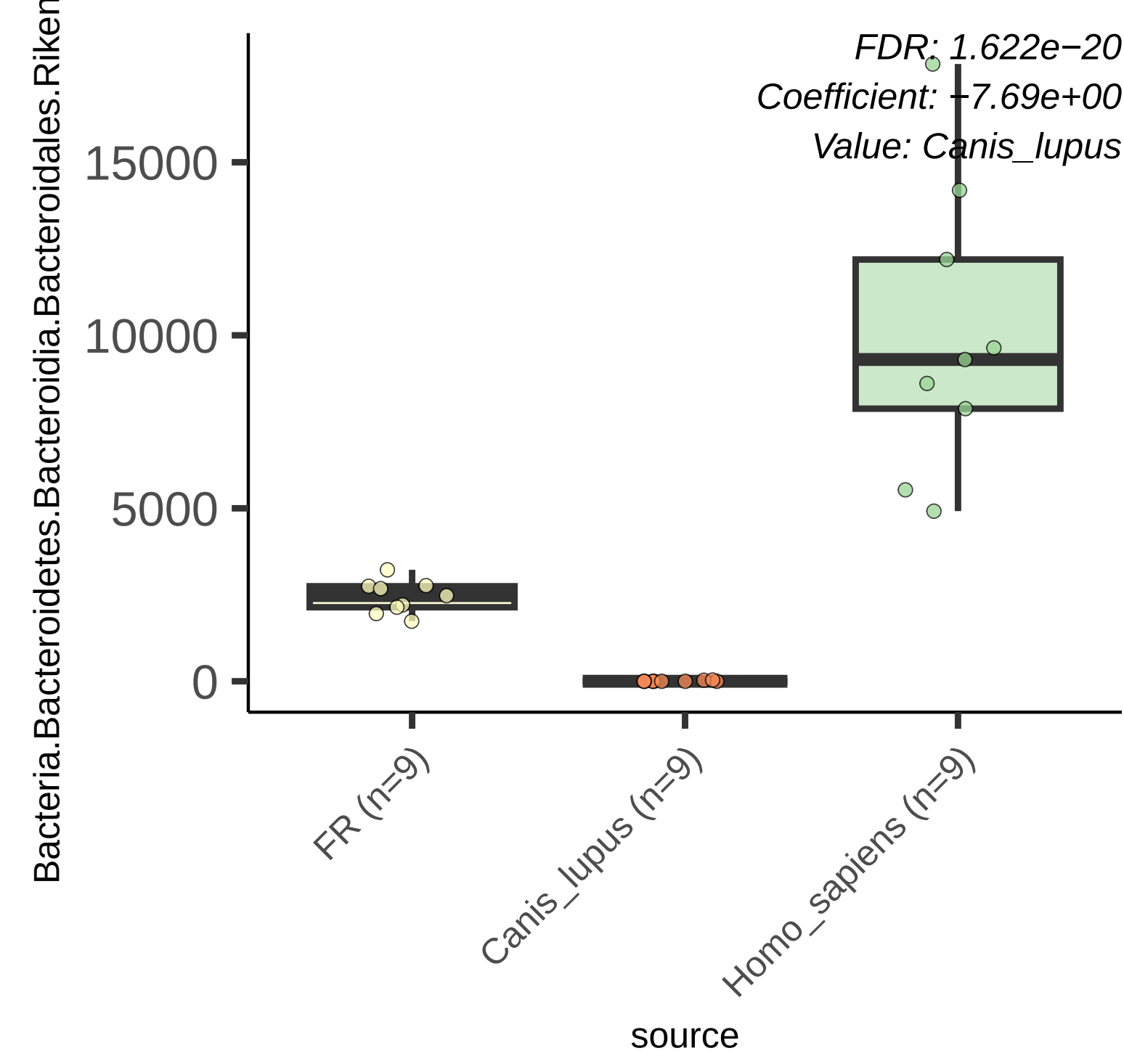
4e+05
2e+05
0e+00
FR (n=9)
Canis_lupus (n=9)
Homo_sapiens (n=9)
source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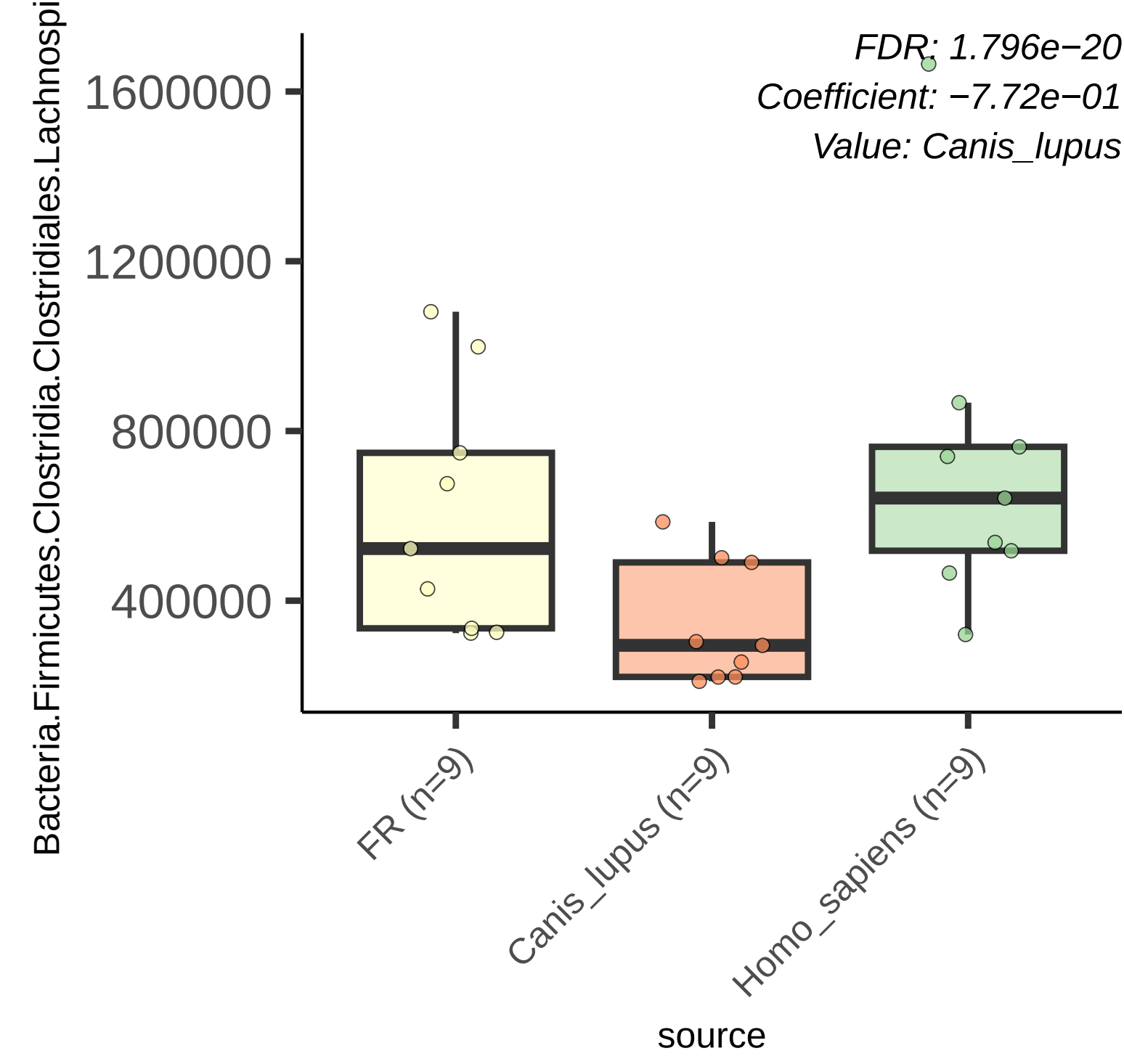
Coefficient: +7.69e+00
Value: Canis_lupus
Bacteria.Bacteroidetes.Bacteroidia.Bacteroidales.Riken
15000
10000
5000
0
FR (n=9)
Canis_lupus (n=9)
Homo_sapiens (n=9)
source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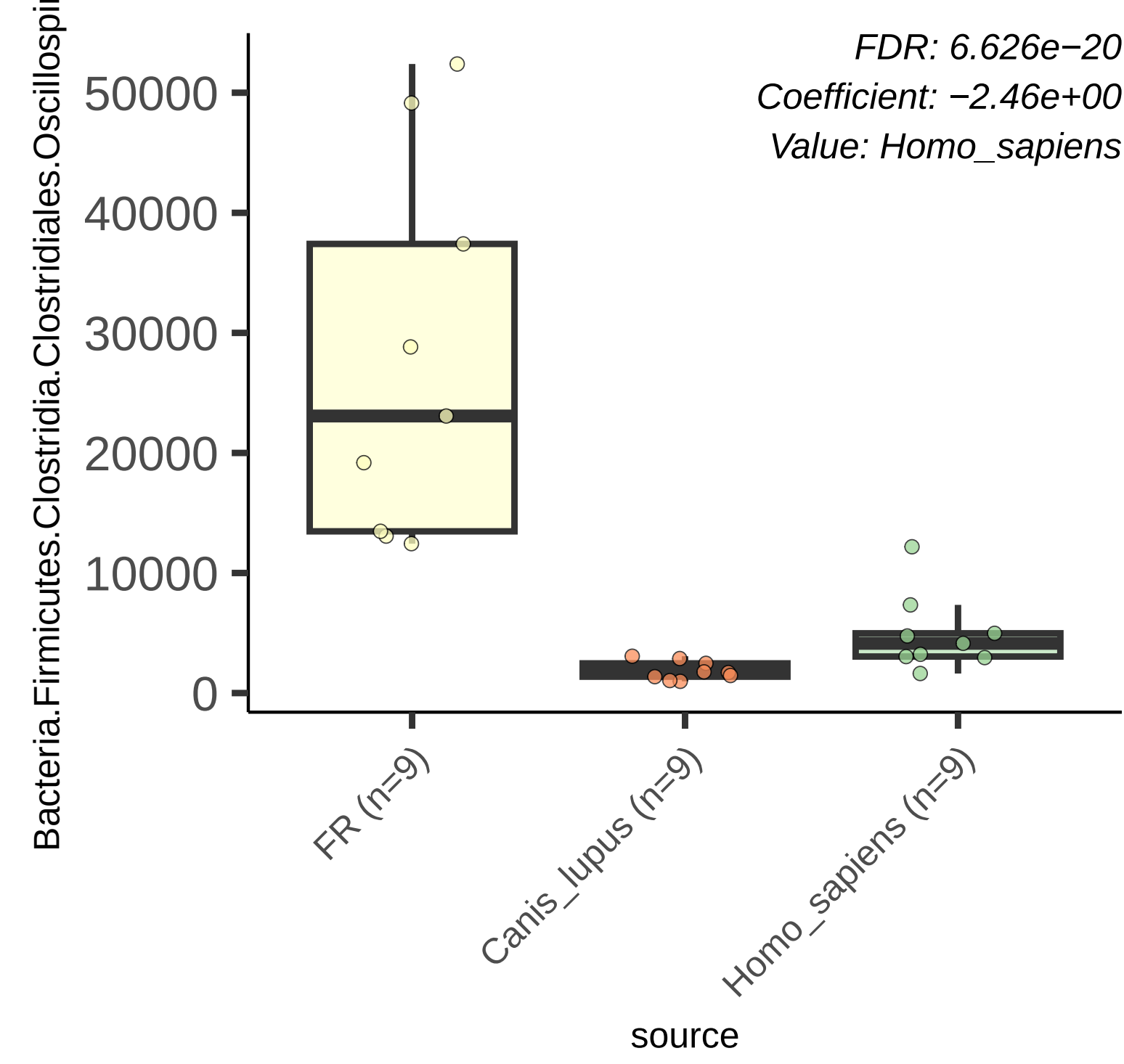
Coefficient: −7.72e−01
Value: Canis_lupus
1600000
1200000
800000
400000
Bacteria.Firmicutes.Clostridia.Clostridiales.Lachnospi
FR (n=9)
Canis_lupus (n=9)
Homo_sapiens (n=9)
source


FDR: 6.626e−20
Coefficient: −2.46e+00
Value: Homo_sapiens
Bacteria.Firmicutes.Clostridia.Clostridiales.Oscillospi
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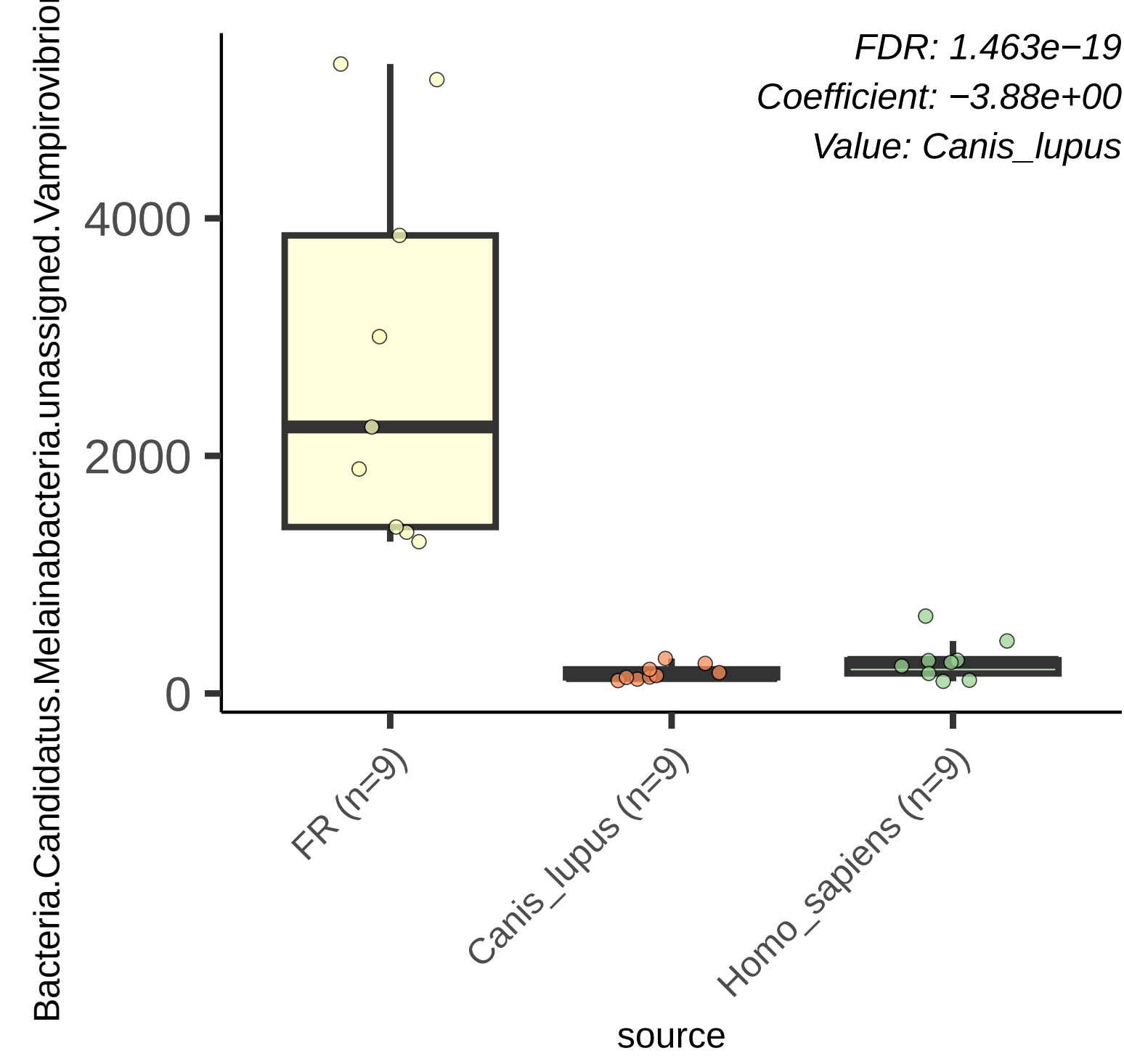
40000
30000
20000
10000
0
FR (n=9)
Canis_lupus (n=9)
Homo_sapiens (n=9)
source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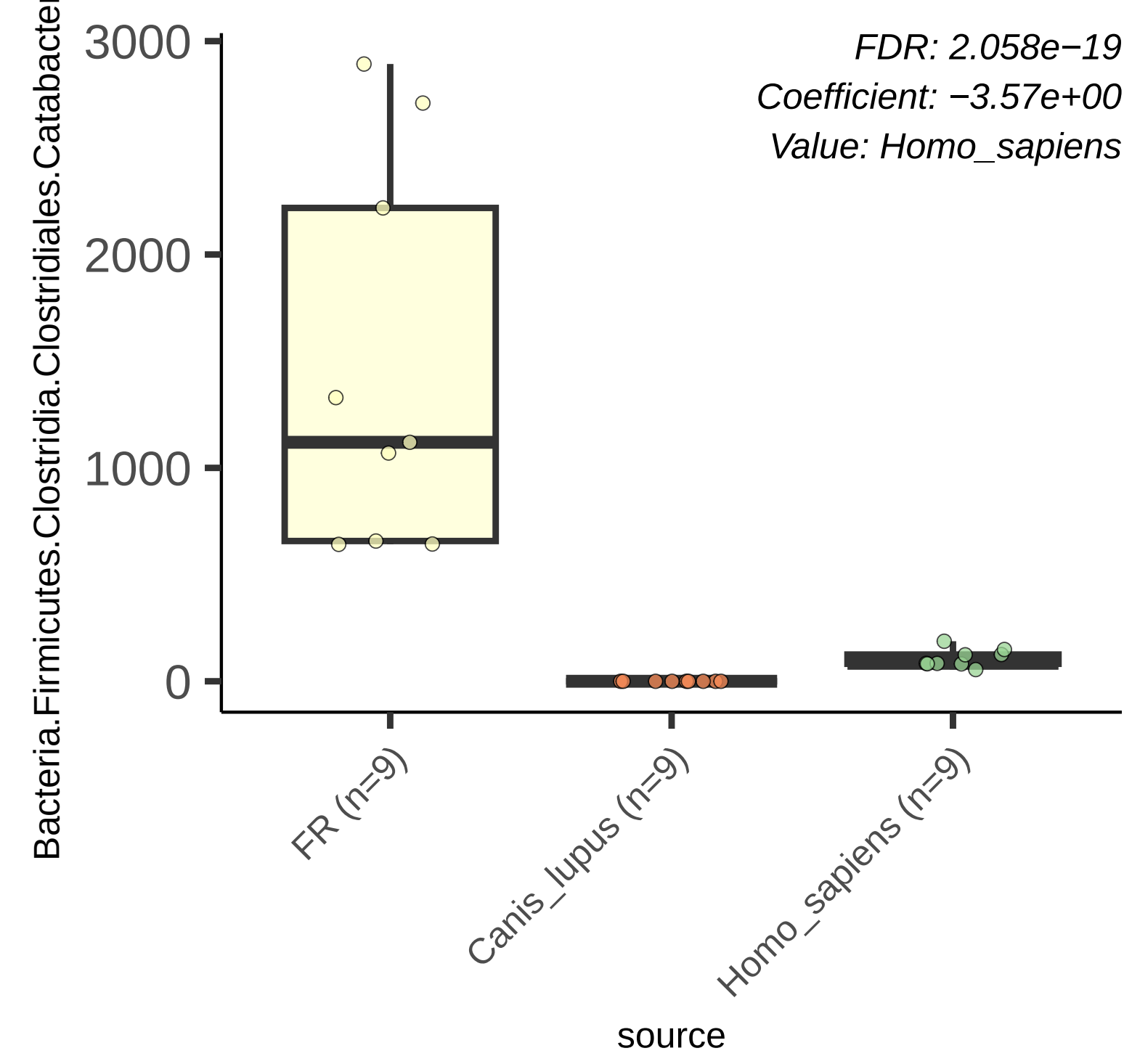
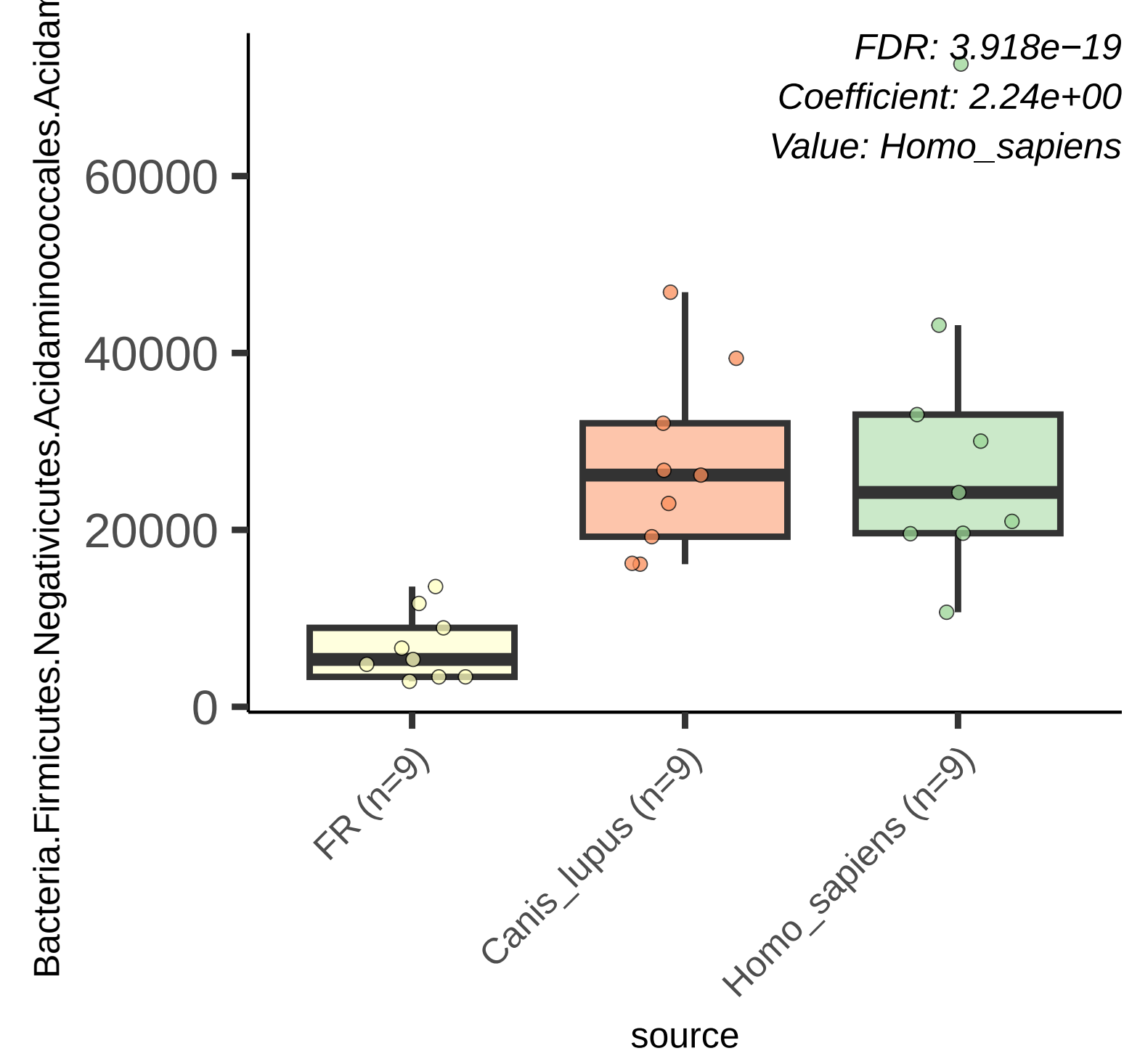
Coefficient: −3.88e+00
Value: Canis_lupus
Bacteria.Candidatus.Melainabacteria.unassigned.Vampirovibrio
4000
2000
0
FR (n=9)
Canis_lupus (n=9)
Homo_sapiens (n=9)
source


FDR: 2.058e−19
Coefficient: −3.57e+00
Value: Homo_sapiens
Bacteria.Firmicutes.Clostridia.Clostridiales.Catabacter
3000
2000
1000
0
FR (n=9)
Canis_lupus (n=9)
Homo_sapiens (n=9)
source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Coefficient: 2.24e+00
Value: Homo_sapiens
Bacteria.Firmicutes.Negativicutes.Acidaminococcales.Acidam
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40000
20000
0
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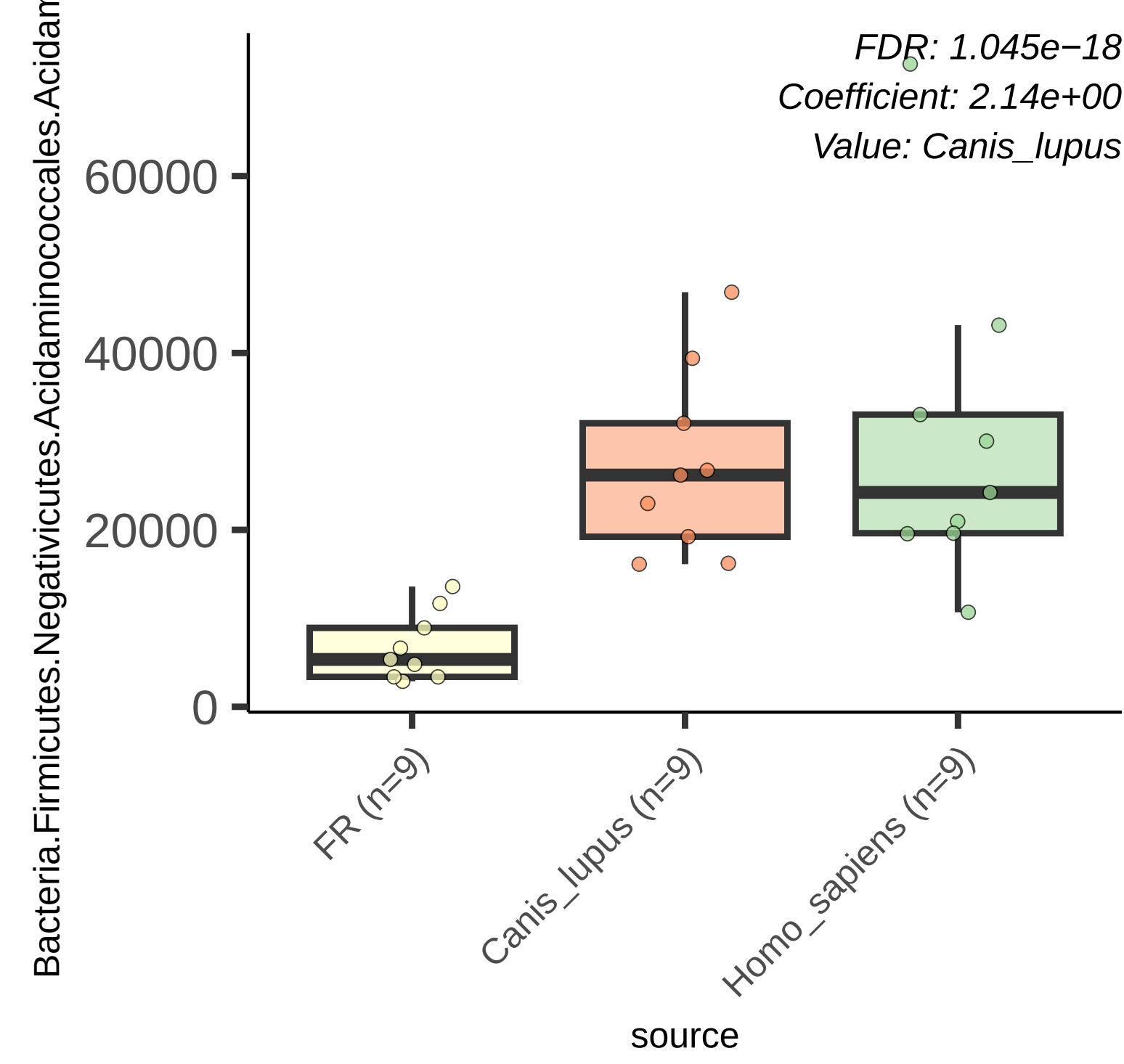
Canis_lupus (n=9)
Homo_sapiens (n=9)
source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Coefficient: 2.14e+00
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Bacteria.Firmicutes.Negativicutes.Acidaminococcales.Acidam
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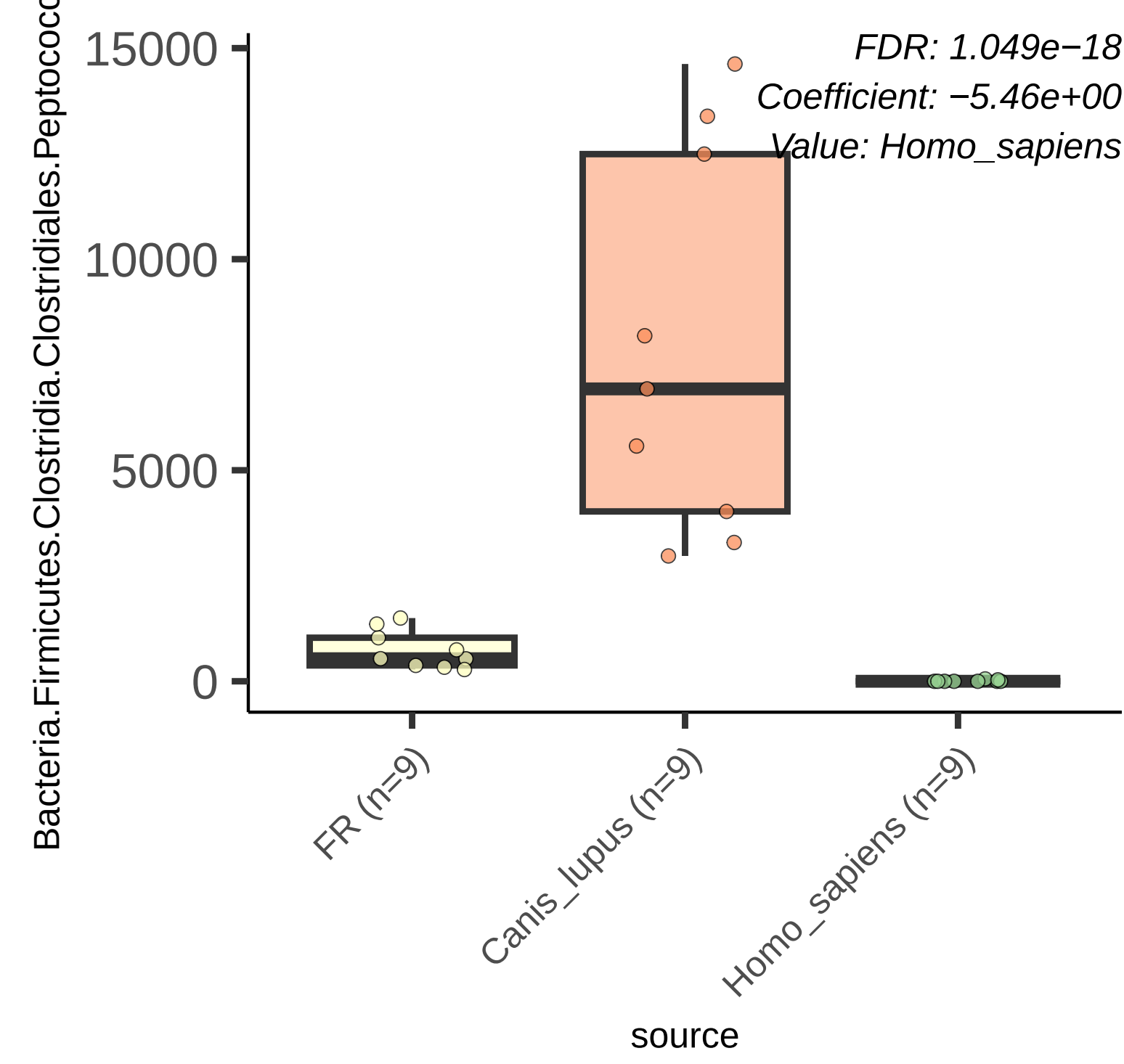
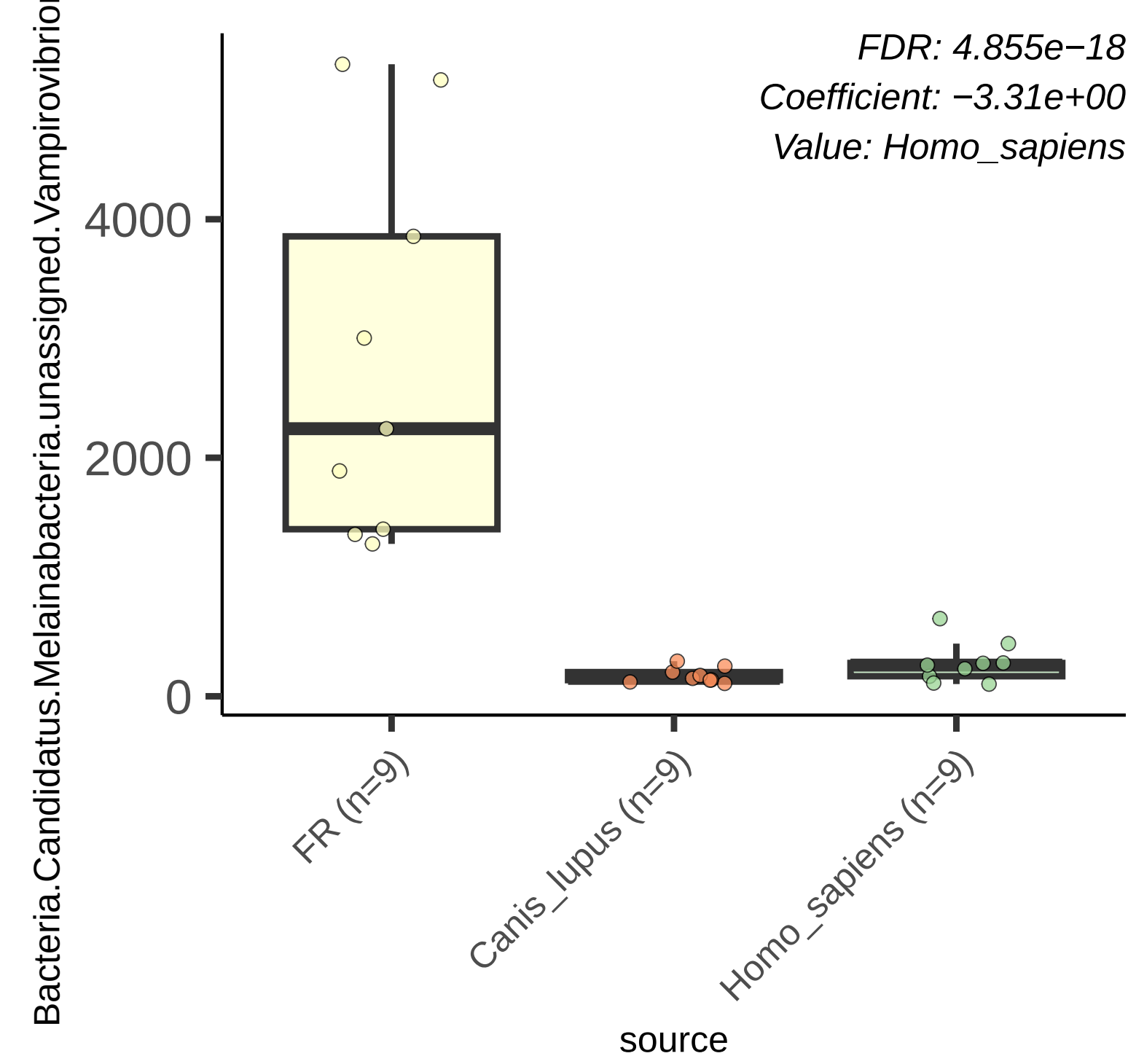
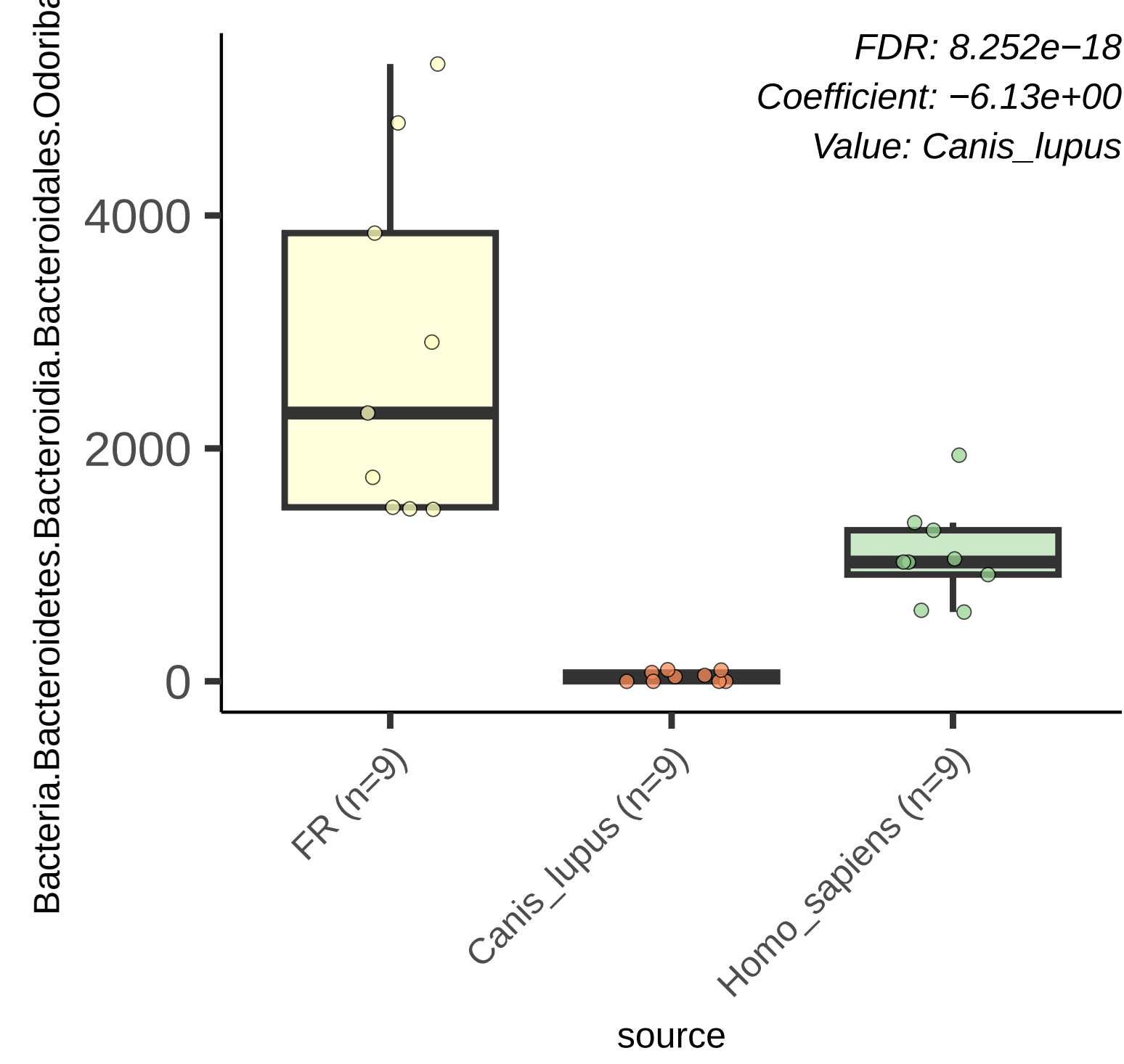
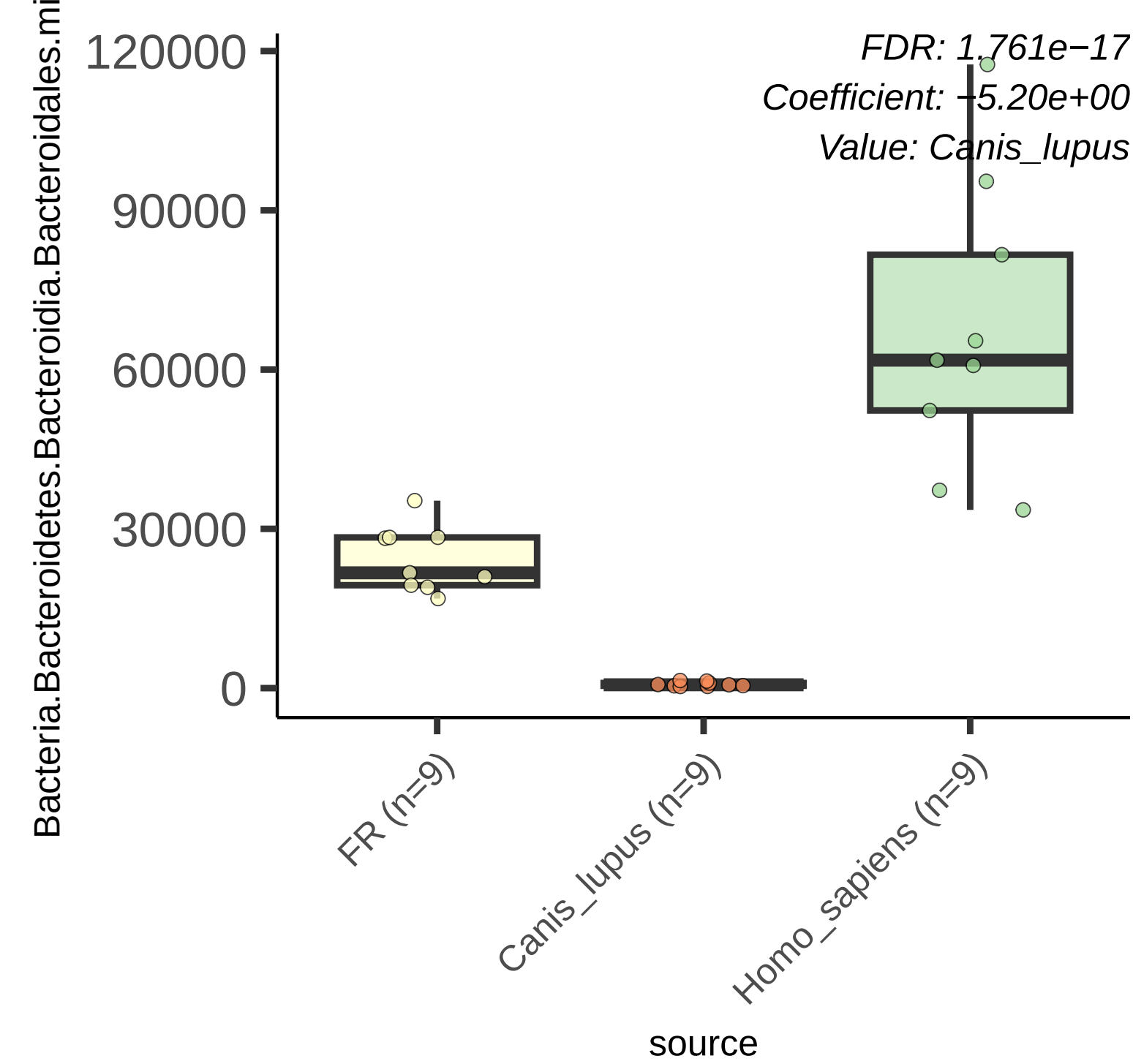
40000
20000
0
FR (n=9)
Canis_lupus (n=9)
Homo_sapiens (n=9)
source


FDR: 1.049e−18
Coefficient: −5.46e+00
Value: Homo_sapiens
15000
10000
5000
0
Bacteria.Firmicutes.Clostridia.Clostridiales.Peptococ
FR (n=9)
Canis_lupus (n=9)
Homo_sapiens (n=9)
source


FDR: 4.855e−18
Coefficient: −3.31e+00
Value: Homo_sapiens
Bacteria.Candidatus.Melainabacteria.unassigned.Vampirovibrio
4000
2000
0
FR (n=9)
Canis_lupus (n=9)
Homo_sapiens (n=9)
source


FDR: 8.252e−18
Coefficient: −6.13e+00
Value: Canis_lupus
Bacteria.Bacteroidetes.Bacteroidia.Bacteroidales.Odoriba
4000
2000
0
FR (n=9)
Canis_lupus (n=9)
Homo_sapiens (n=9)
source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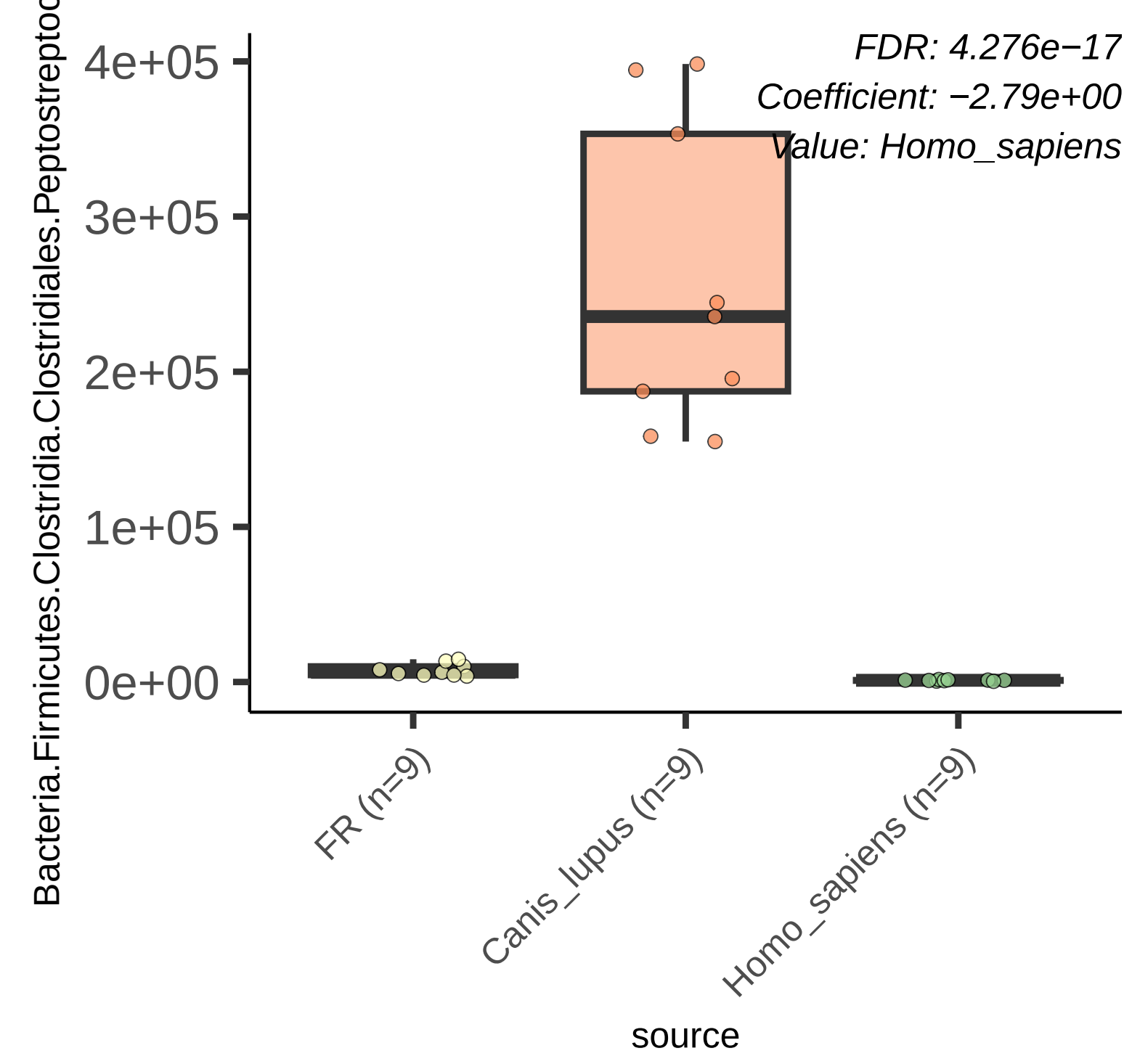
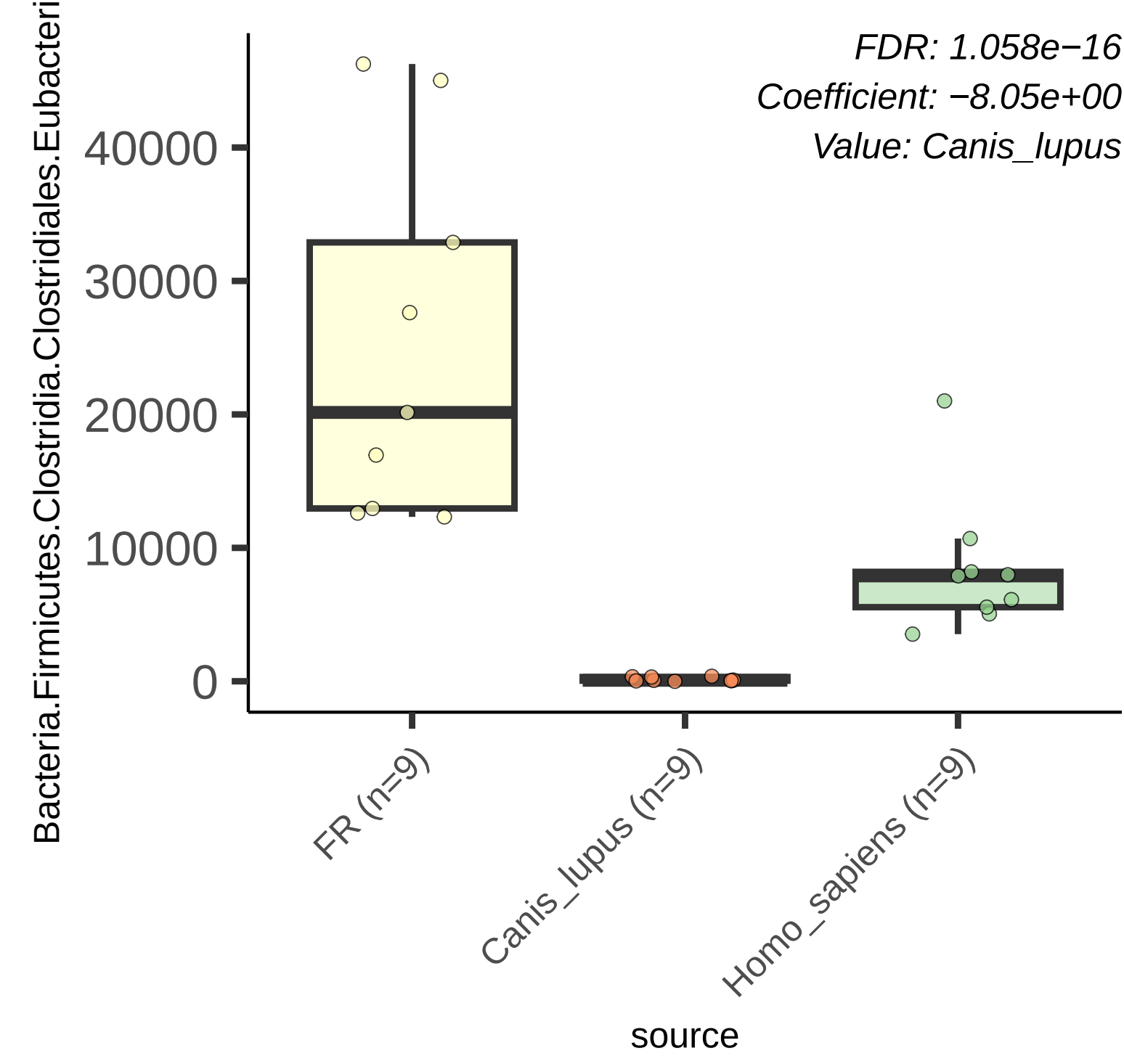
Coefficient: −5.20e+00
Value: Canis_lupus
120000
90000
60000
30000
0
Bacteria.Bacteroidetes.Bacteroidia.Bacteroidales.mi
FR (n=9)
Canis_lupus (n=9)
Homo_sapiens (n=9)
source


FDR: 4.276e−17
Coefficient: −2.79e+00
Value: Homo_sapiens
Bacteria.Firmicutes.Clostridia.Clostridiales.Peptostreptoc
4e+05
3e+05
2e+05
1e+05
0e+00
FR (n=9)
Canis_lupus (n=9)
Homo_sapiens (n=9)
source


FDR: 1.058e−16
Coefficient: −8.05e+00
Value: Canis_lupus
Bacteria.Firmicutes.Clostridia.Clostridiales.Eubacteri
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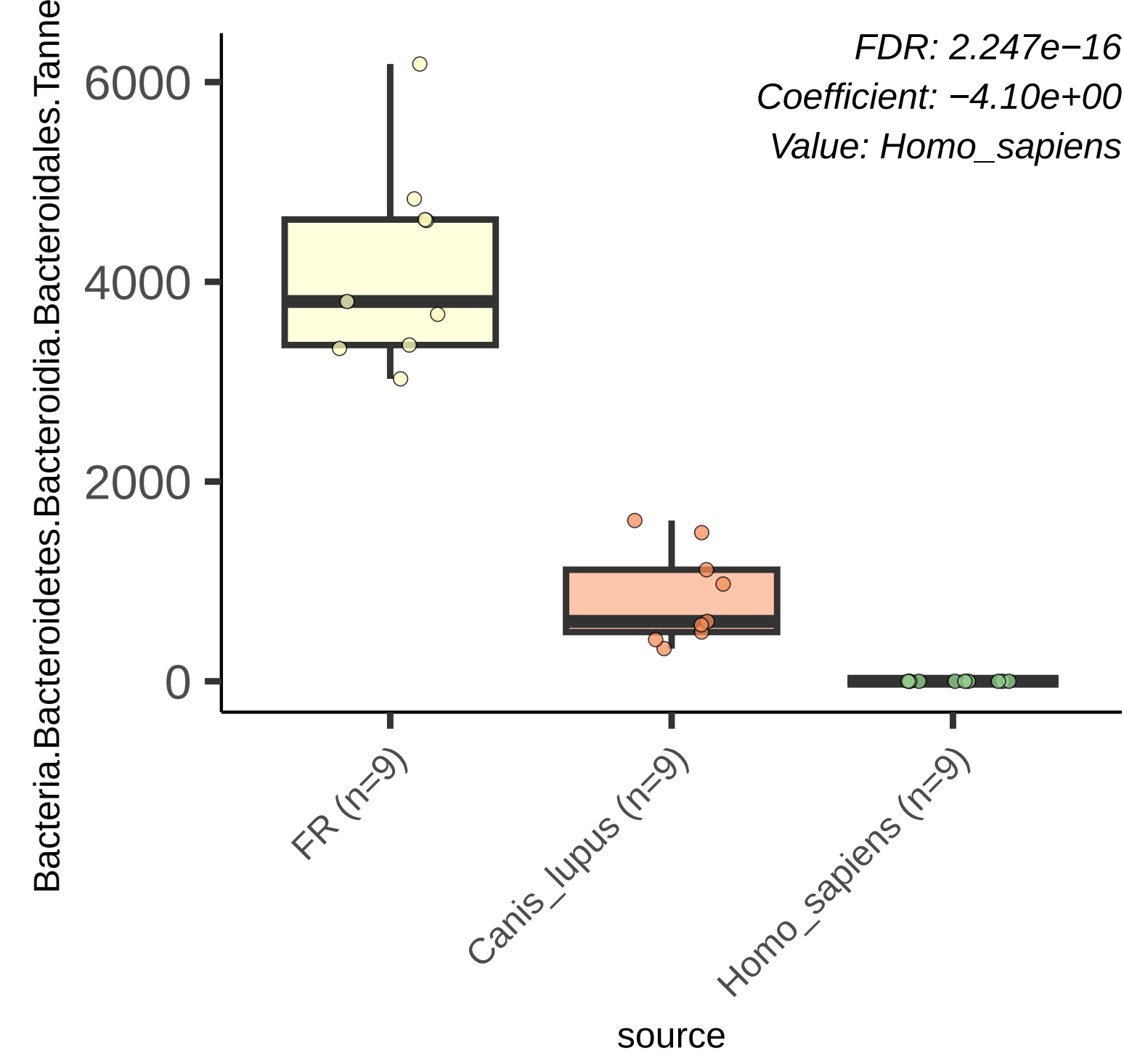
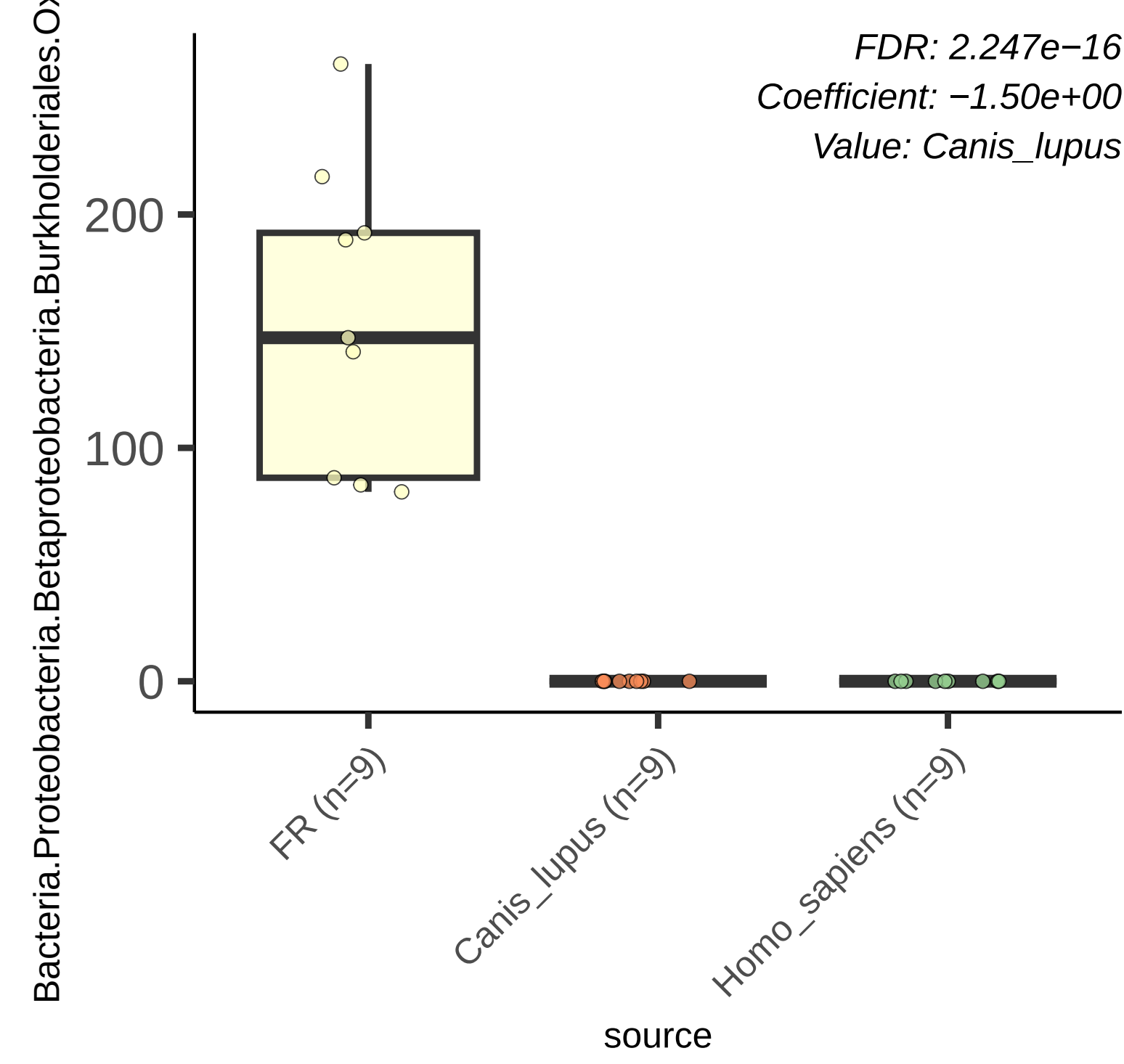
30000
20000
10000
0
FR (n=9)
Canis_lupus (n=9)
Homo_sapiens (n=9)
source


FDR: 2.247e−16
Coefficient: −4.10e+00
Value: Homo_sapiens
Bacteria.Bacteroidetes.Bacteroidia.Bacteroidales.Tanne
6000
4000
2000
0
FR (n=9)
Canis_lupus (n=9)
Homo_sapiens (n=9)
source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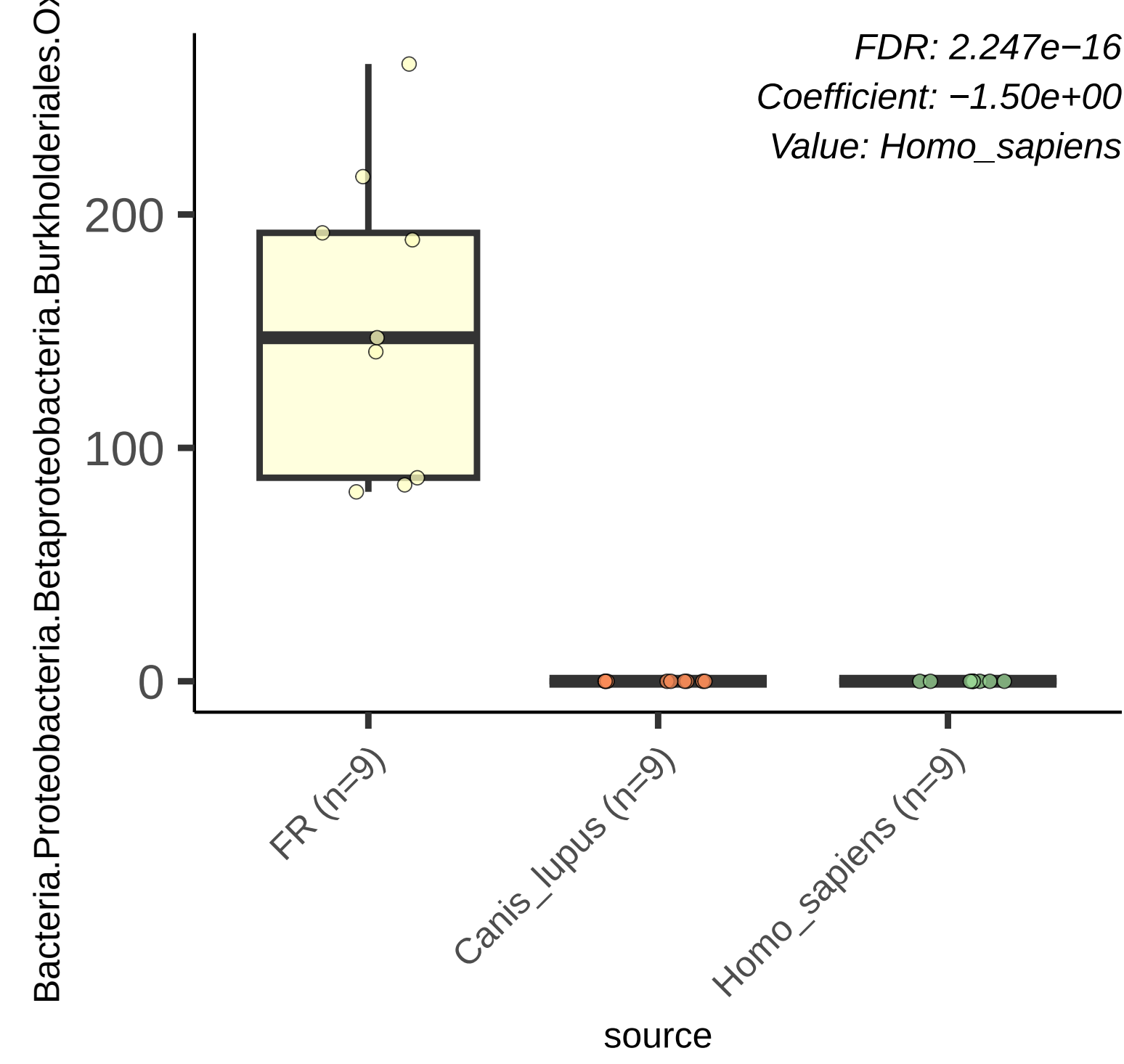
Canis_lupus (n=9)
Homo_sapiens (n=9)
source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Bacteria.Proteobacteria.Betaproteobacteria.Burkholderiales.Ox
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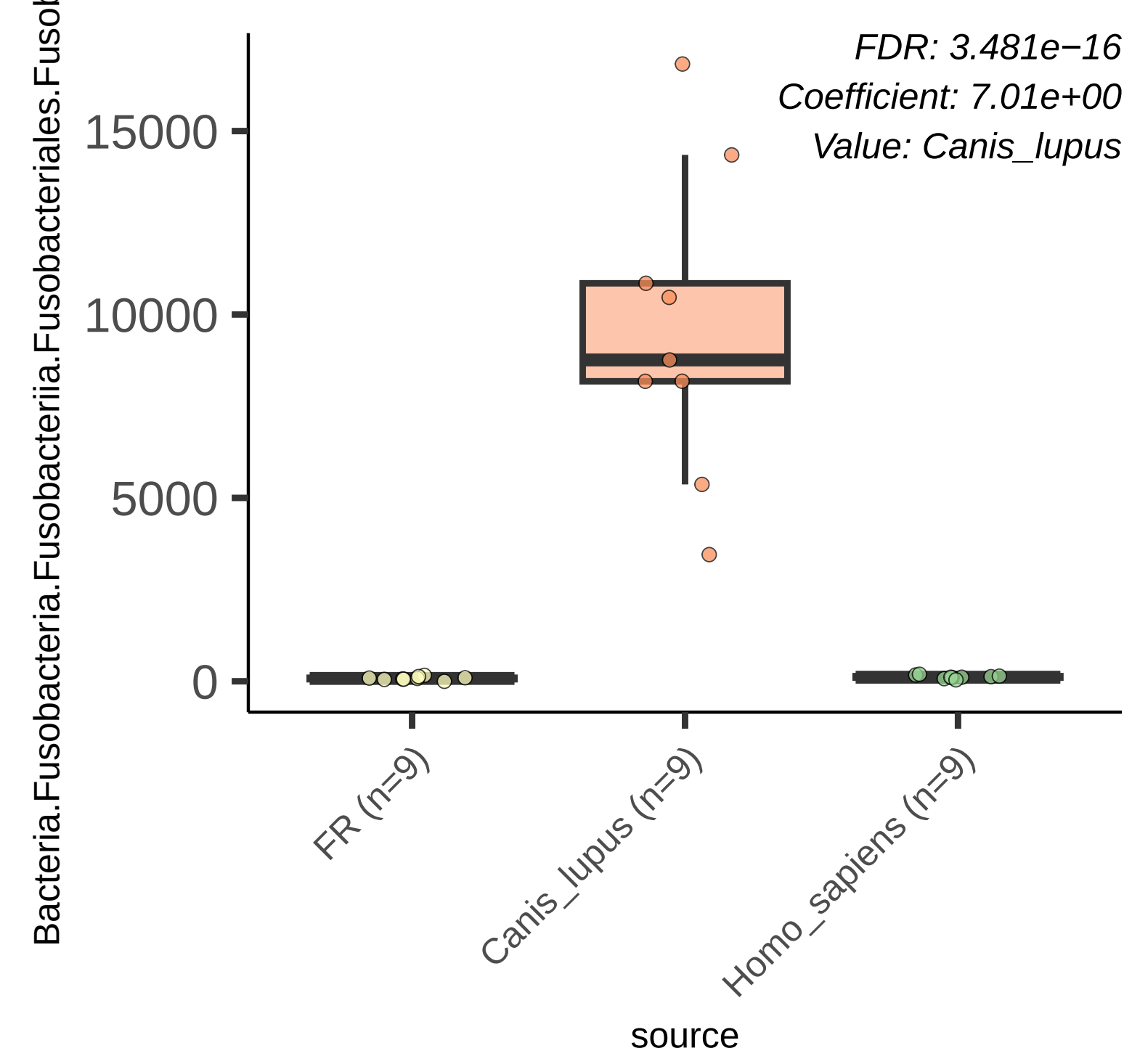
Canis_lupus (n=9)
Homo_sapiens (n=9)
source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15000
10000
5000
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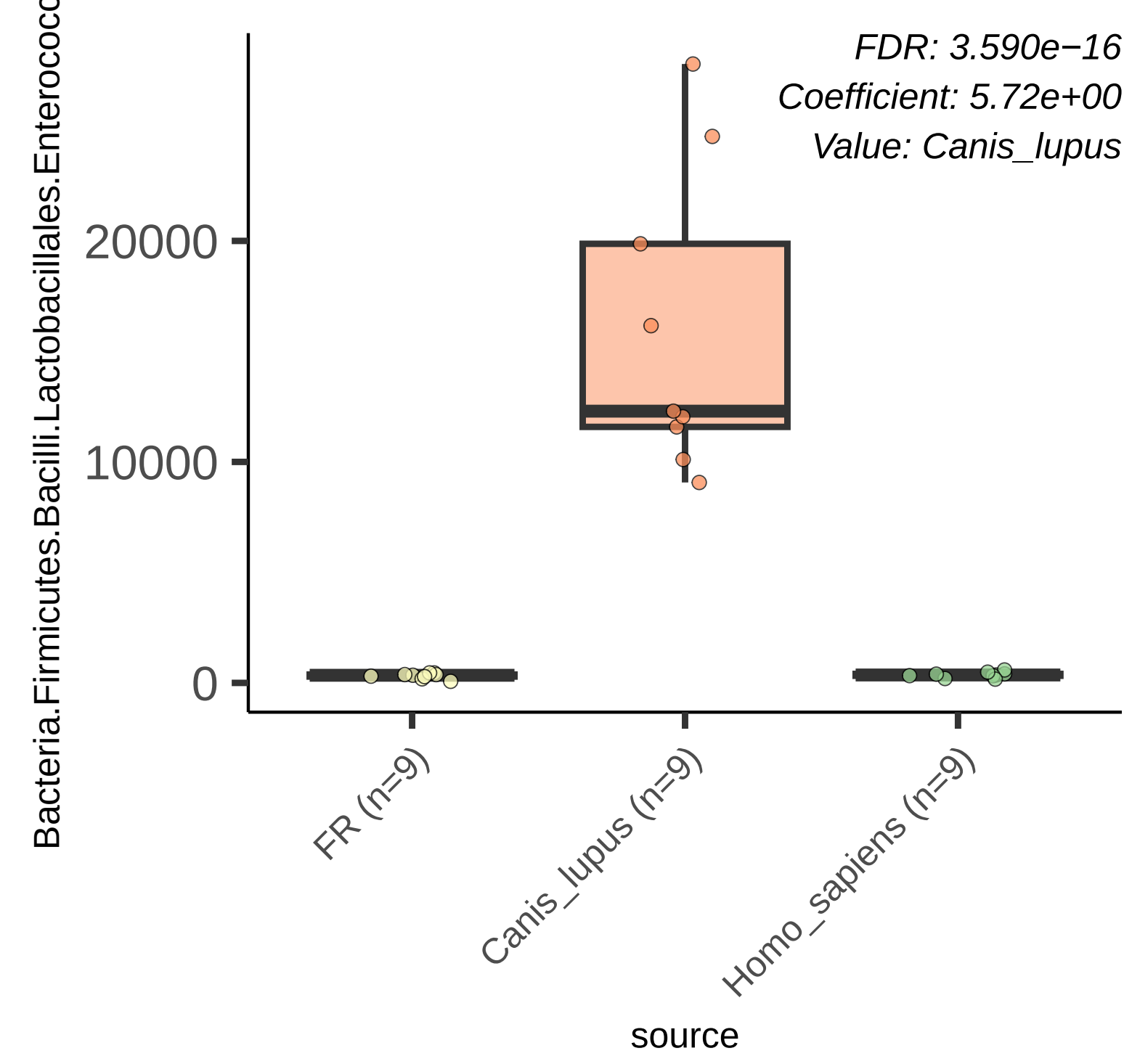
FR (n=9)
Canis_lupus (n=9)
Homo_sapiens (n=9)
source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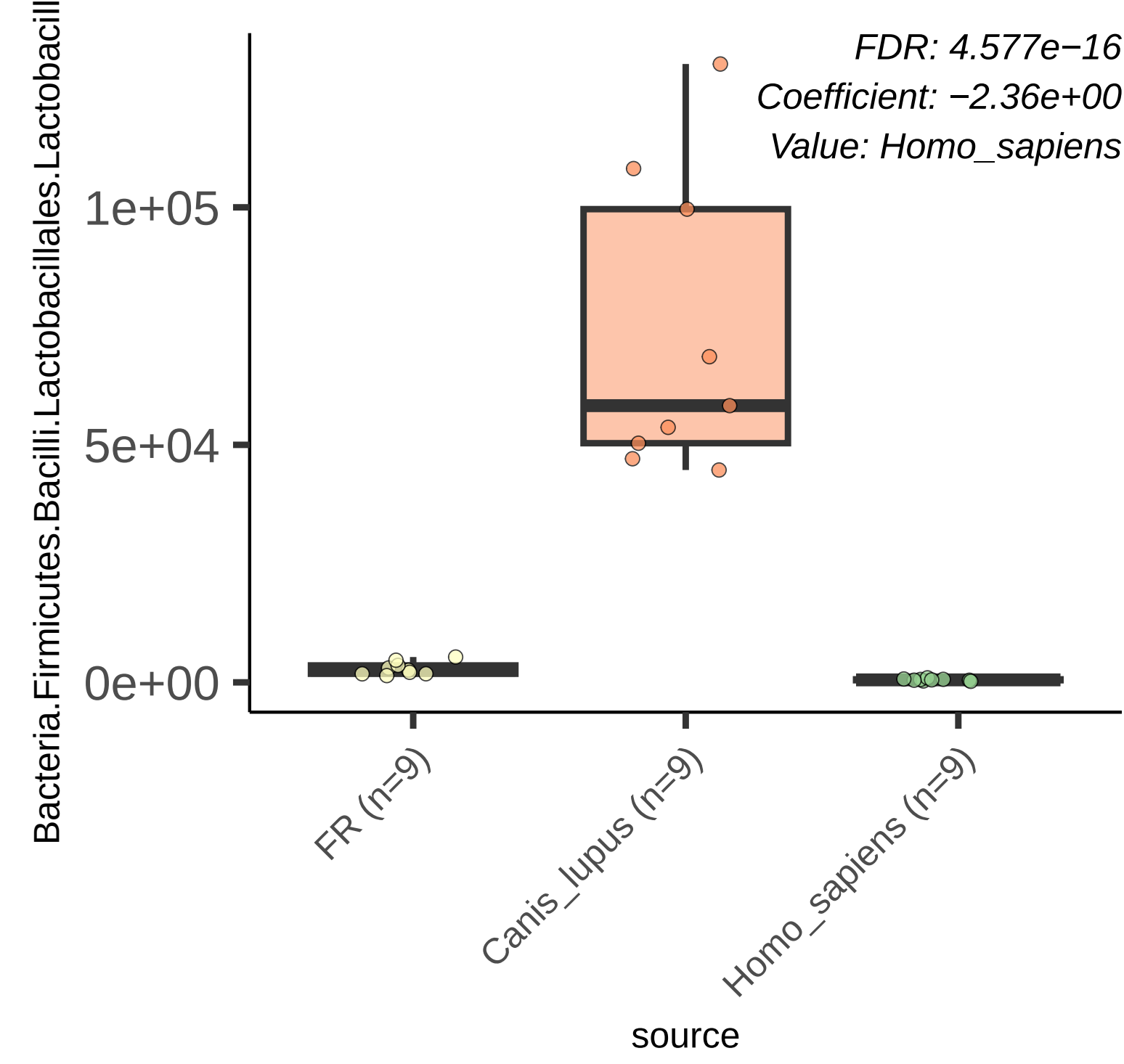
Canis_lupus (n=9)
Homo_sapiens (n=9)
source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5e+04
0e+00
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Homo_sapiens (n=9)
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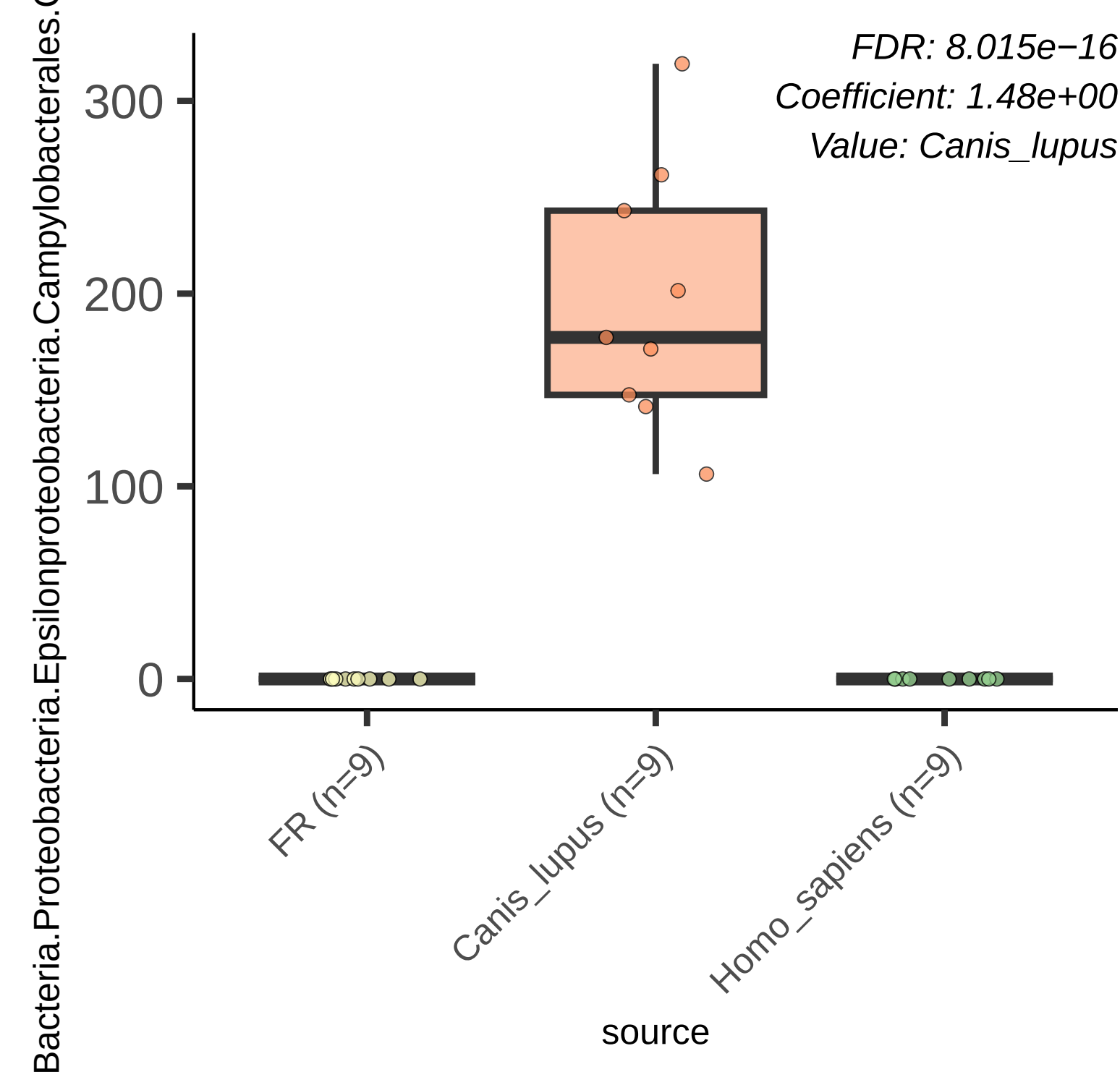
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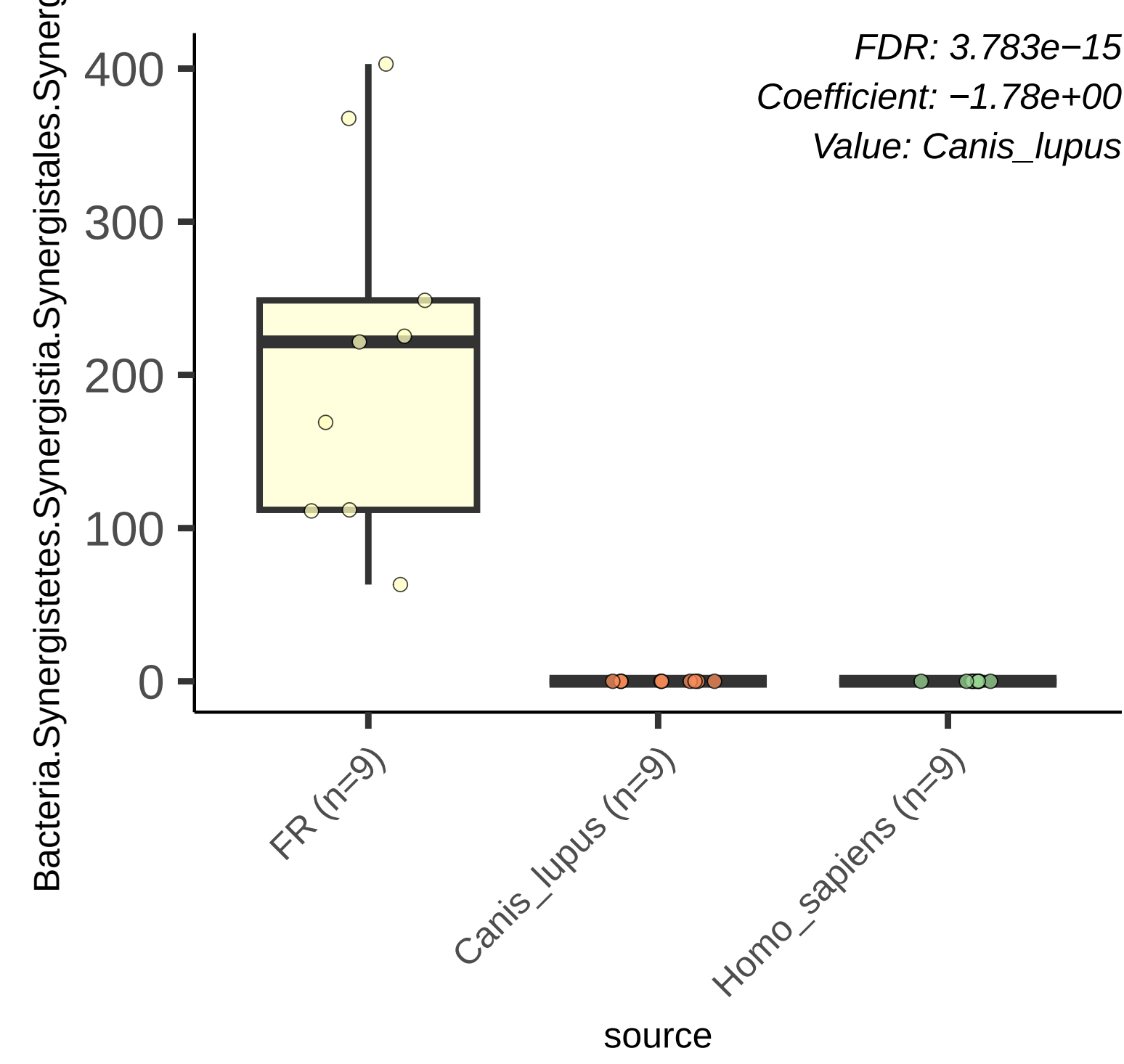
Canis_lupus (n=9)
Homo_sapiens (n=9)
source
Bacteria.Proteobacteria.Epsilonproteobacteria.Campylobacterales.C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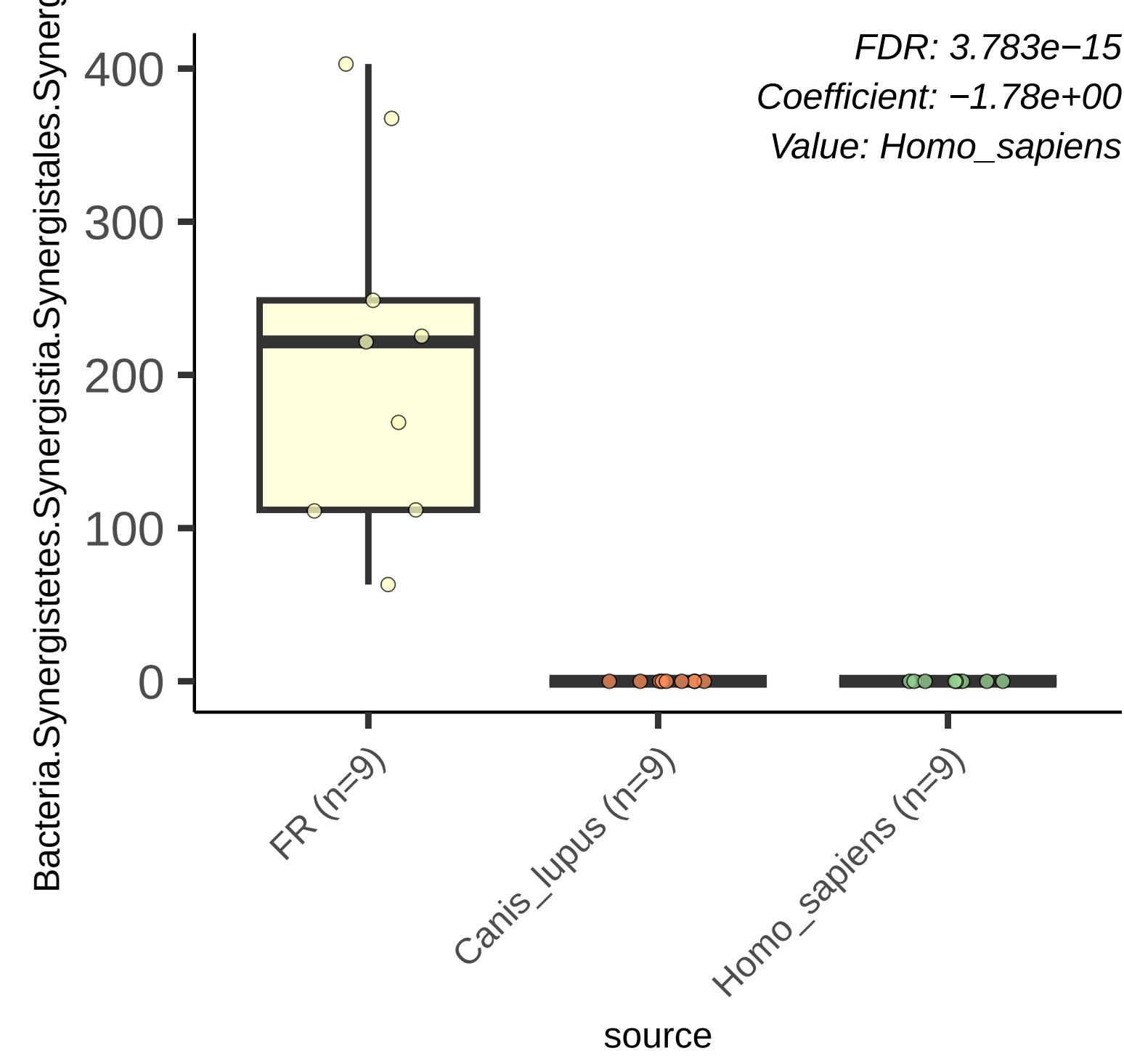
Canis_lupus (n=9)
Homo_sapiens (n=9)
source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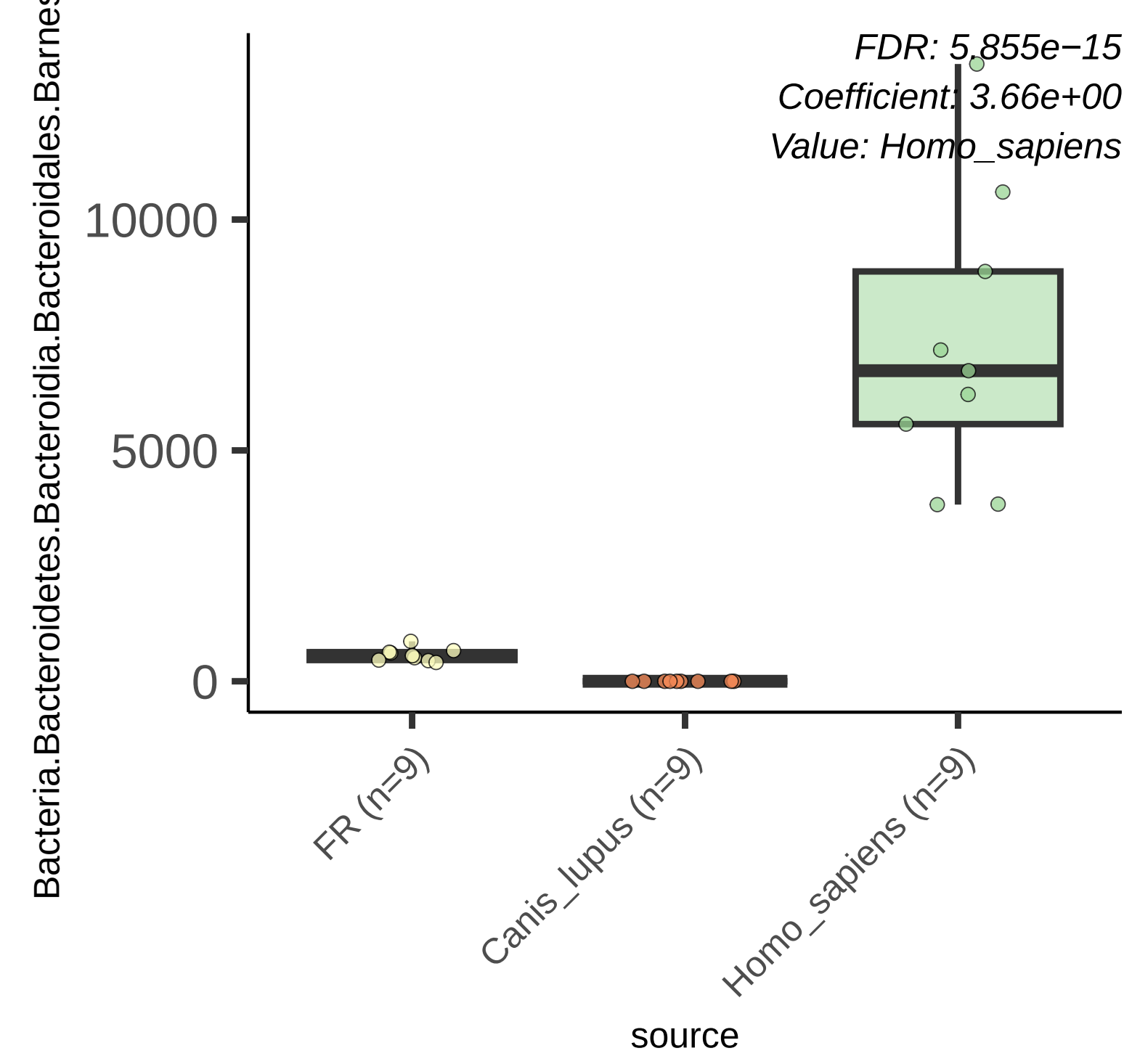
Homo_sapiens (n=9)
source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5000
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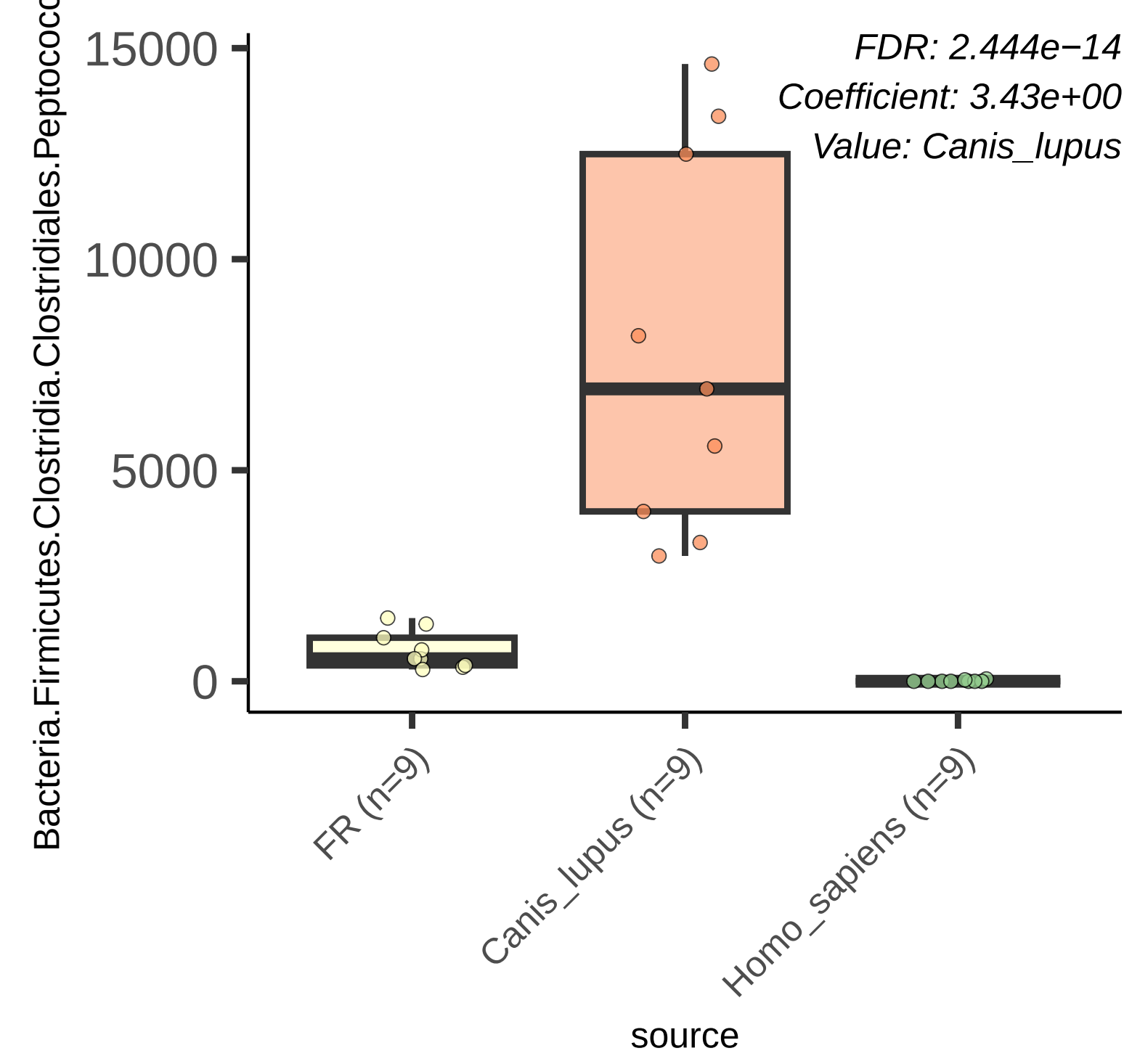
FR (n=9)
Canis_lupus (n=9)
Homo_sapiens (n=9)
source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10000
5000
0
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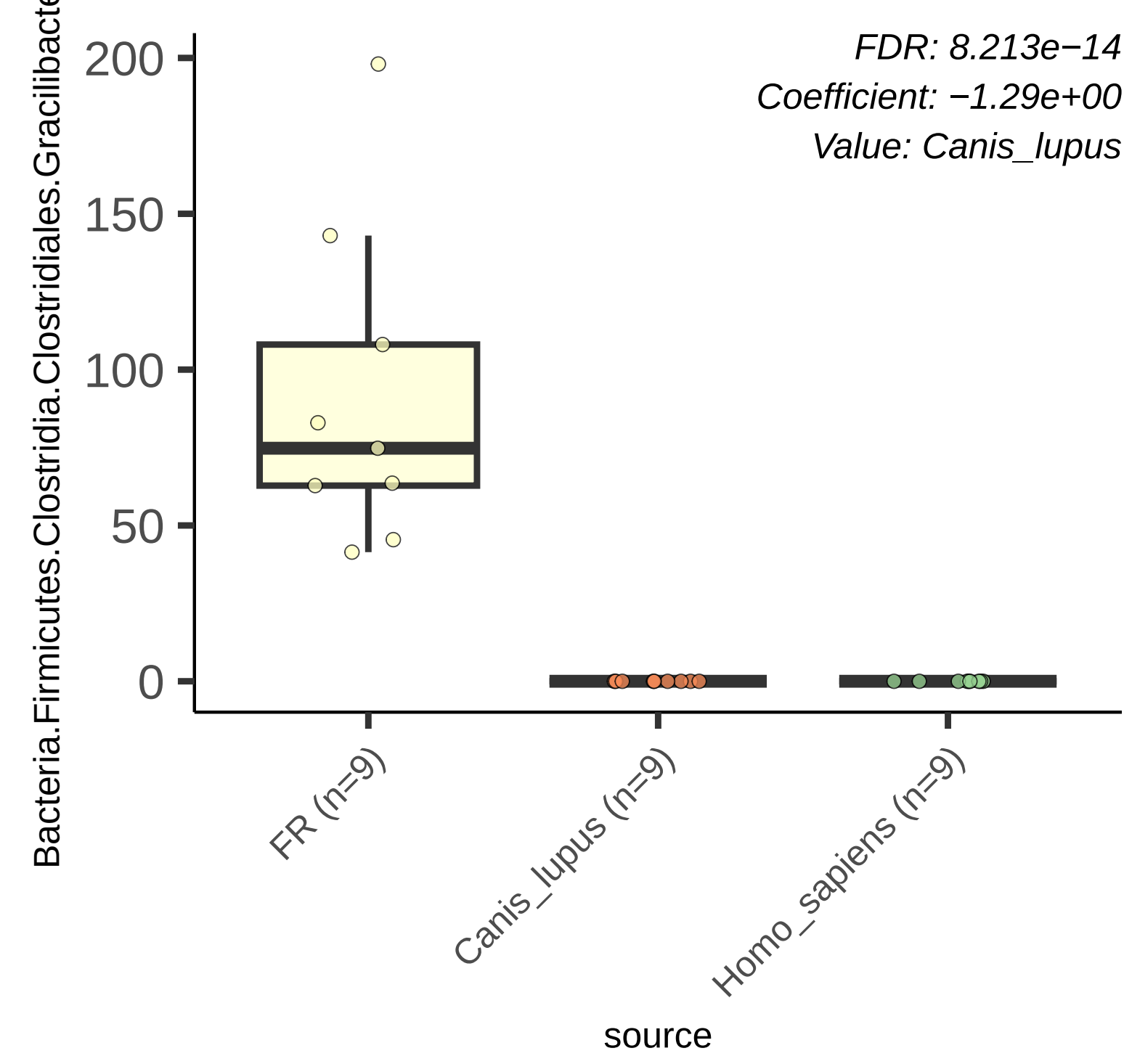
Canis_lupus (n=9)
Homo_sapiens (n=9)
source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Coefficient: −1.29e+00
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Bacteria.Firmicutes.Clostridia.Clostridiales.Gracilibacte
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150
100
50
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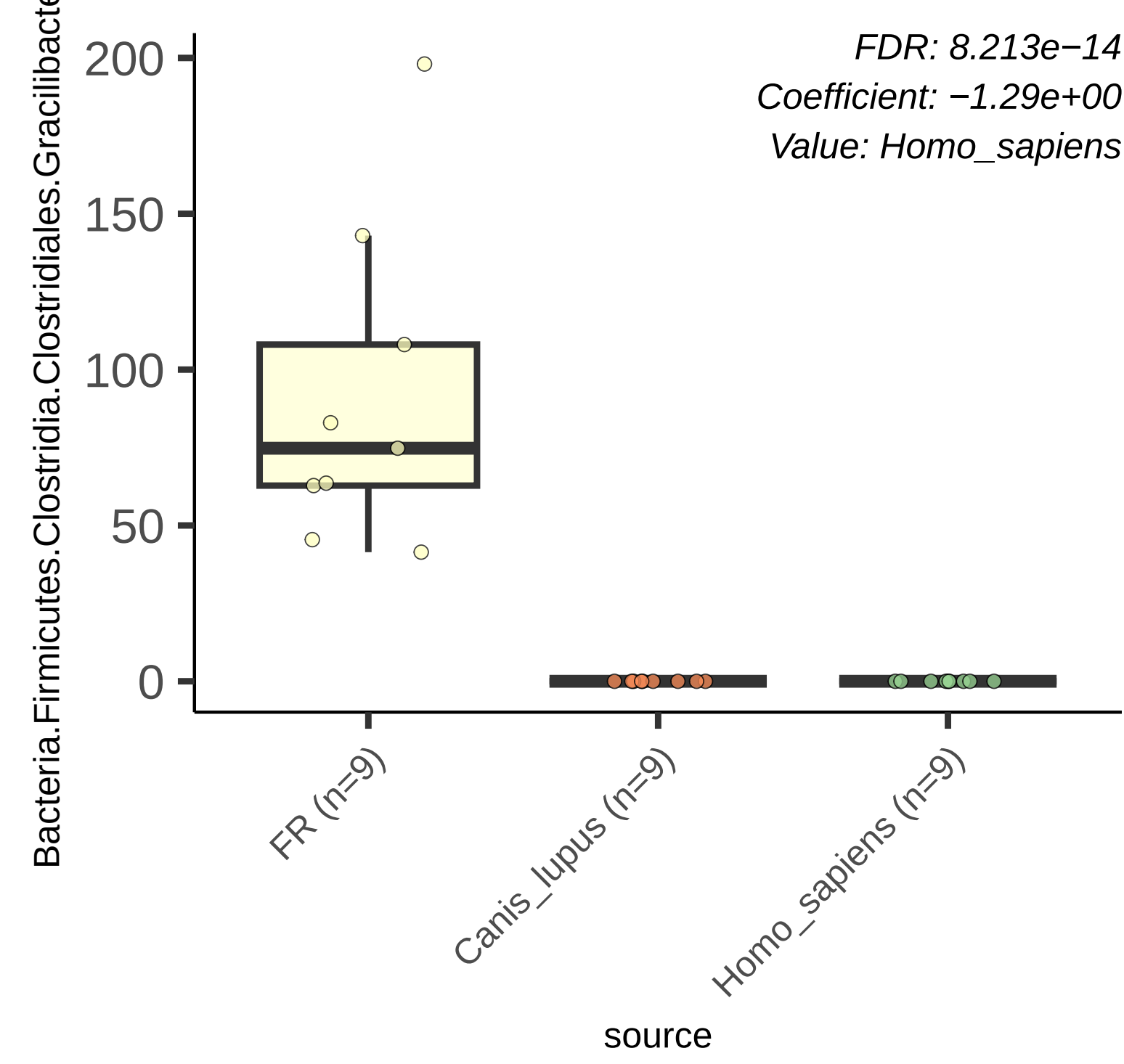
Canis_lupus (n=9)
Homo_sapiens (n=9)
source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Coefficient: −1.29e+00
Value: Homo_sapiens
Bacteria.Firmicutes.Clostridia.Clostridiales.Gracilibacte
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150
100
50
0
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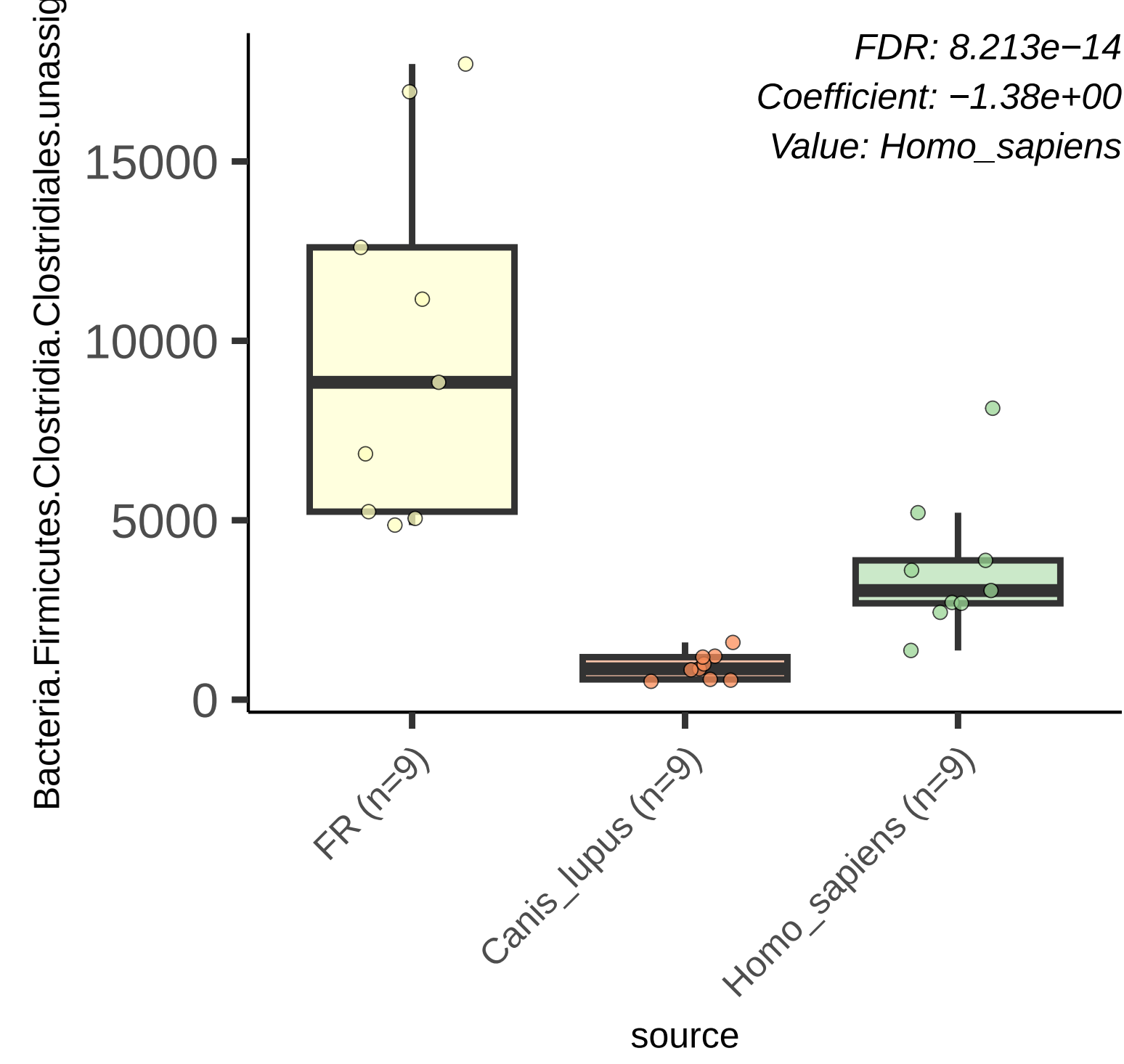
Canis_lupus (n=9)
Homo_sapiens (n=9)
source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Coefficient: −1.38e+00
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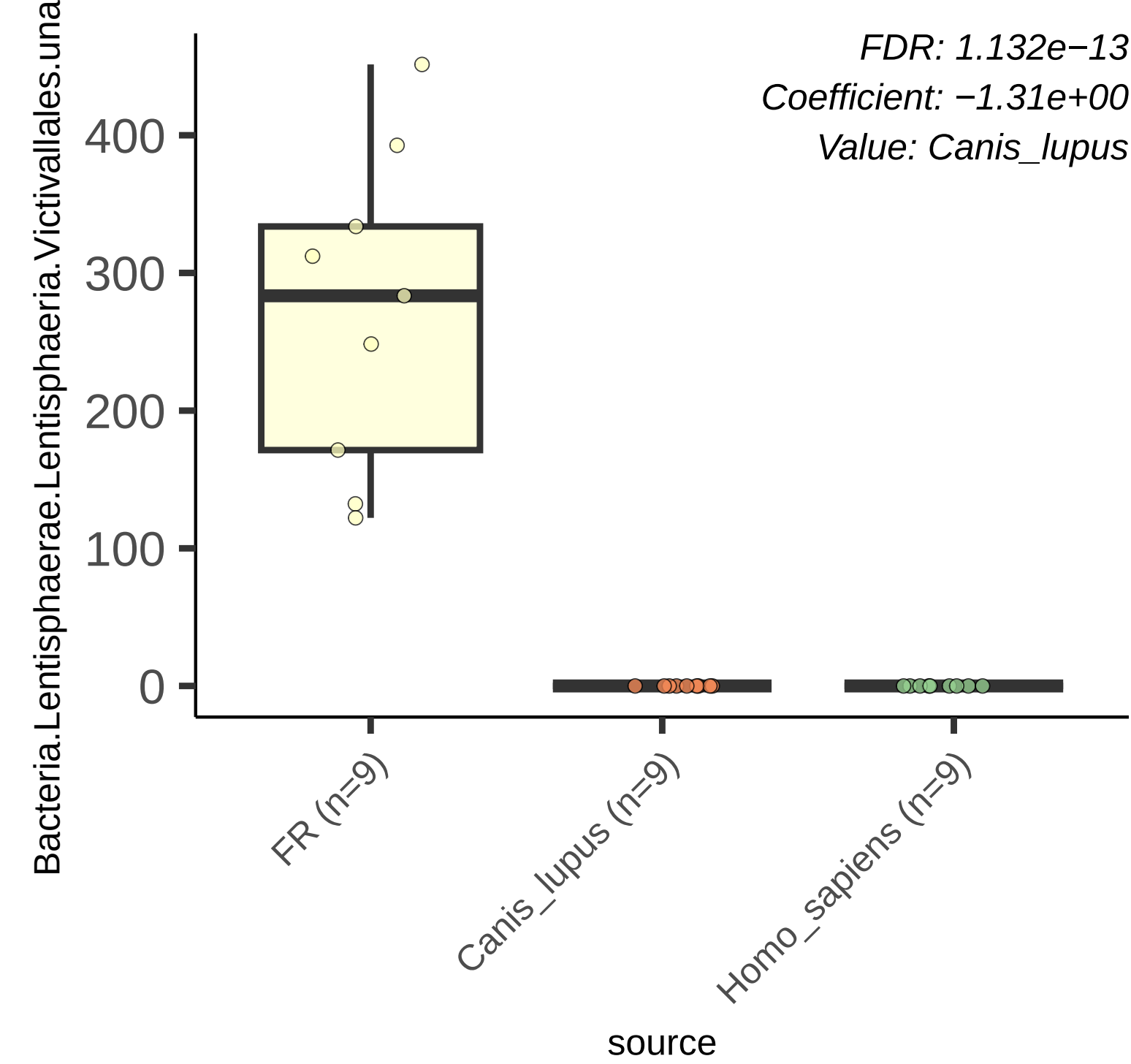
Bacteria.Firmicutes.Clostridia.Clostridiales.unassig
15000
10000
5000
0
FR (n=9)
Canis_lupus (n=9)
Homo_sapiens (n=9)
source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300
200
100
0
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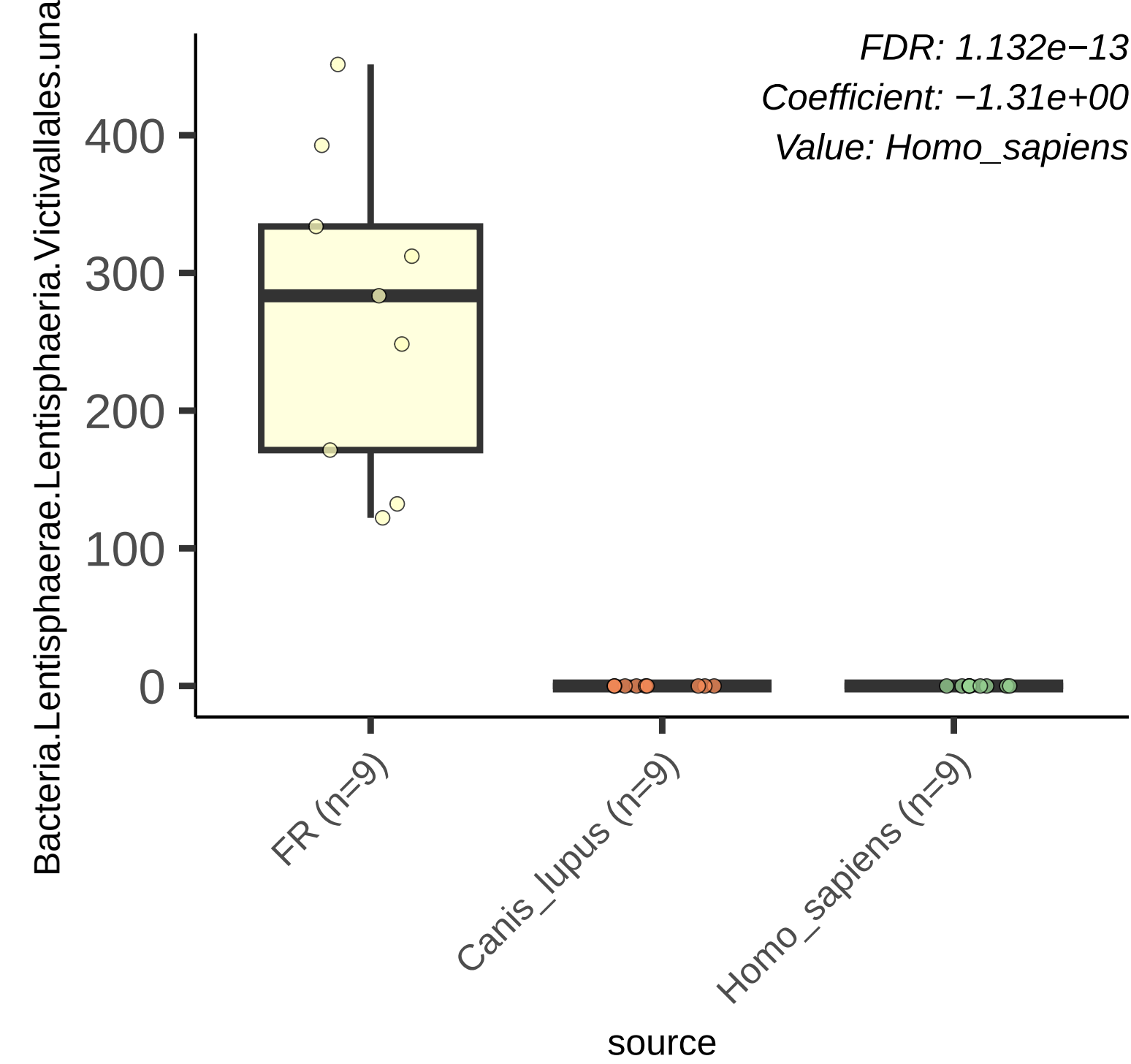
Canis_lupus (n=9)
Homo_sapiens (n=9)
source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Coefficient: −1.31e+00
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Bacteria.Lentisphaerae.Lentisphaeria.Victivallales.una
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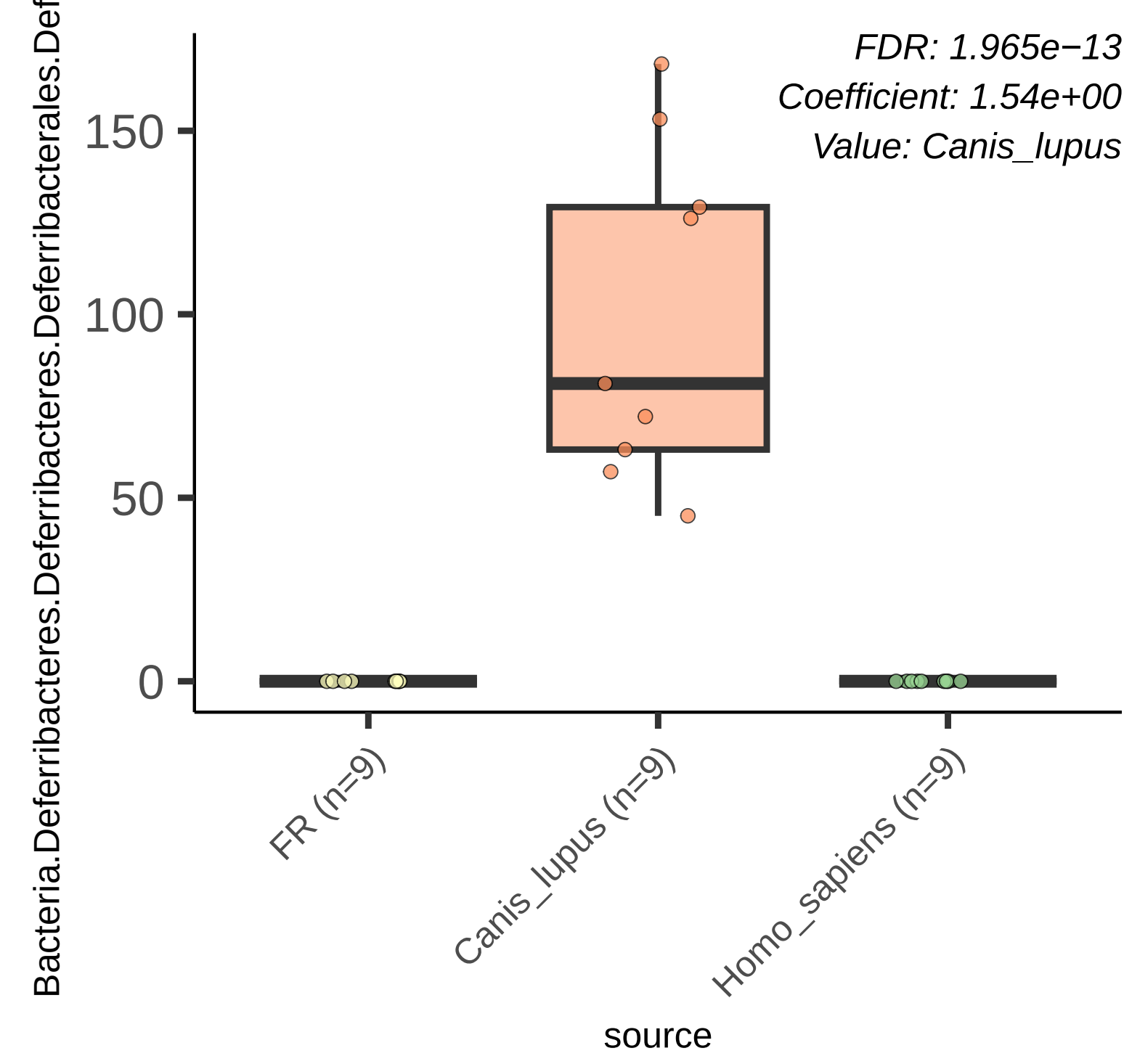
300
200
100
0
FR (n=9)
Canis_lupus (n=9)
Homo_sapiens (n=9)
source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Bacteria.Deferribacteres.Deferribacteres.Deferribacterales.Def
150
100
50
0
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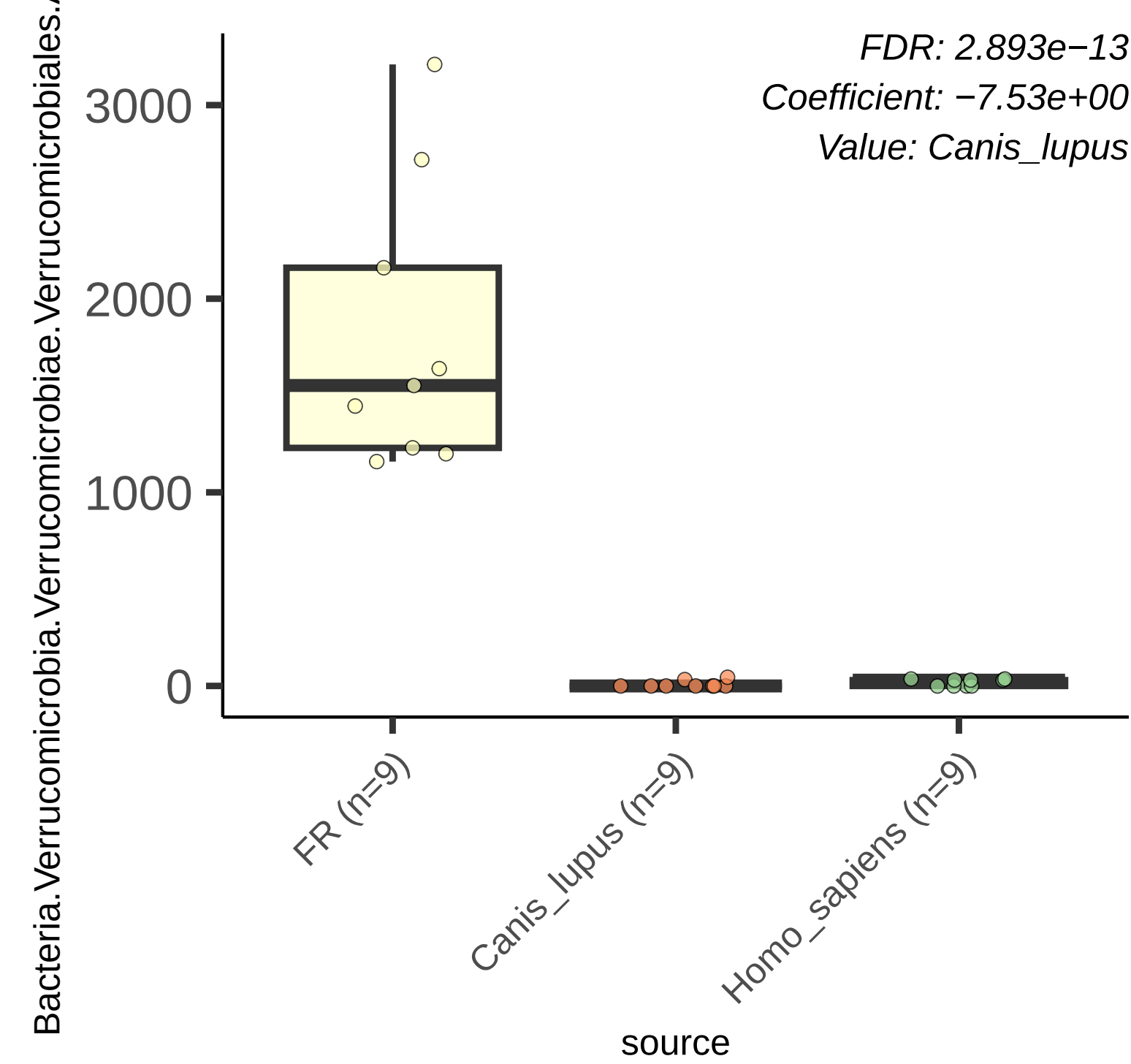
Canis_lupus (n=9)
Homo_sapiens (n=9)
source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Coefficient: −7.53e+00
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2000
1000
0
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FR (n=9)
Canis_lupus (n=9)
Homo_sapiens (n=9)
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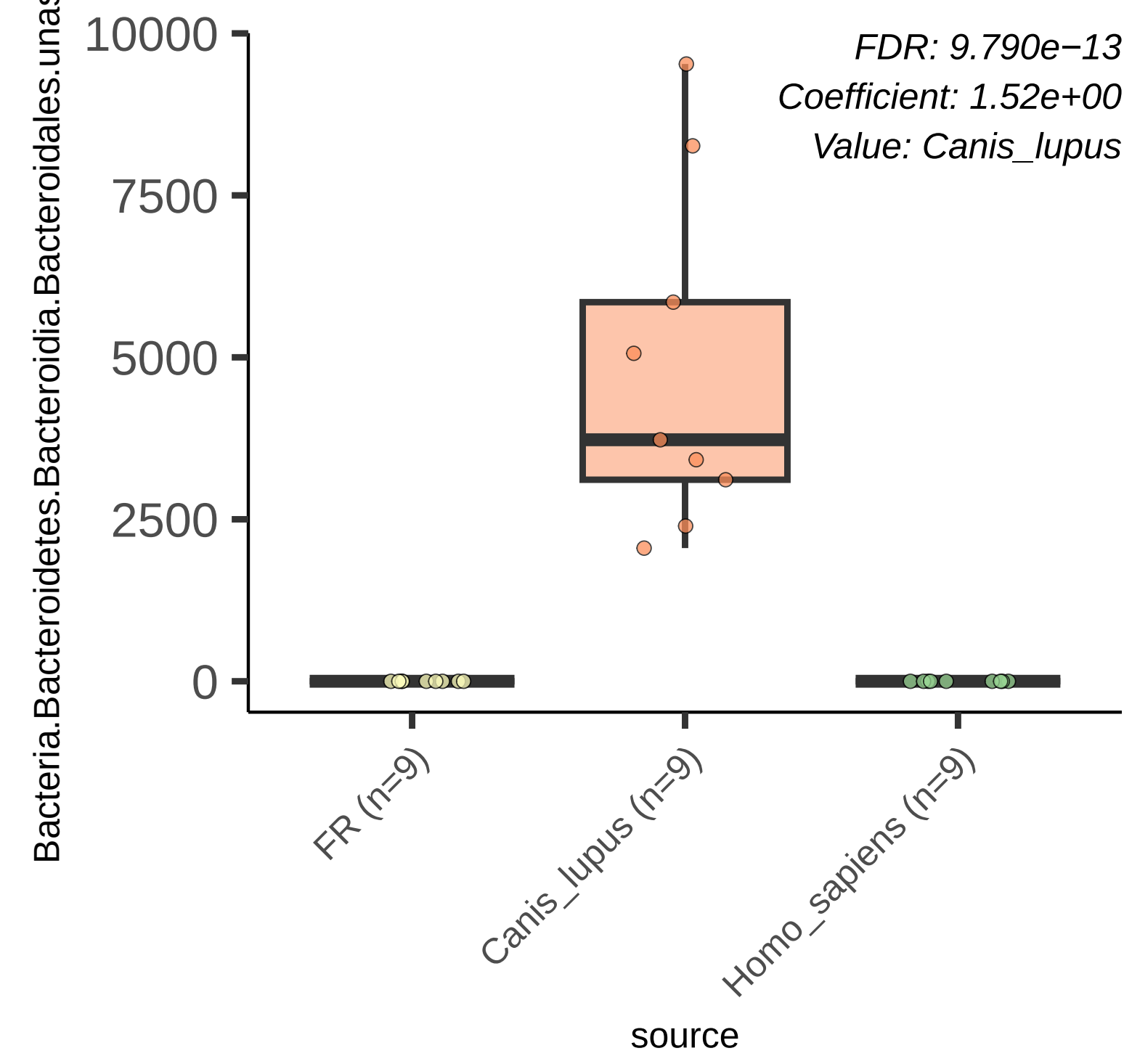
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Coefficient: 1.52e+00
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7500
5000
2500
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Canis_lupus (n=9)
Homo_sapiens (n=9)
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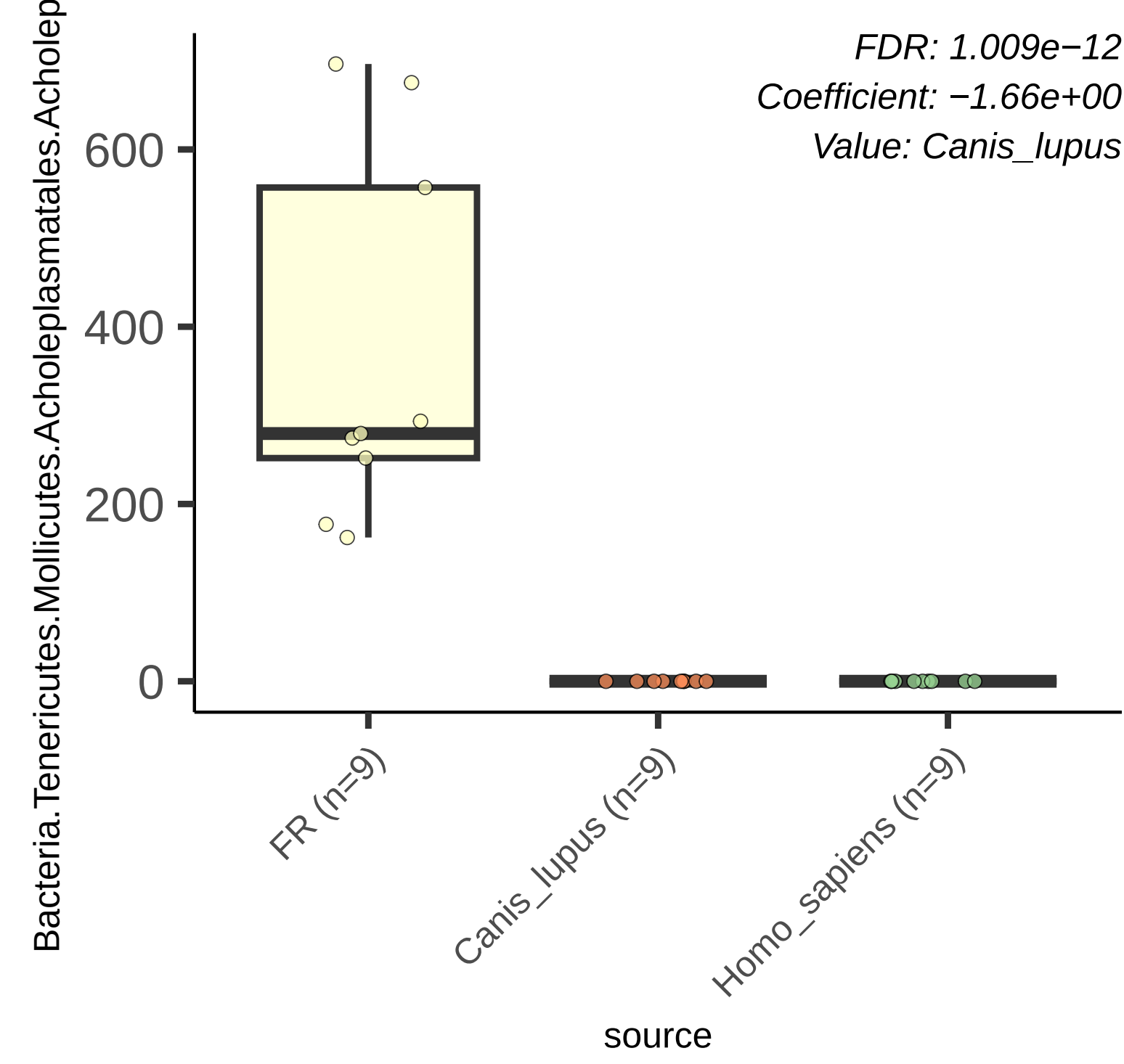
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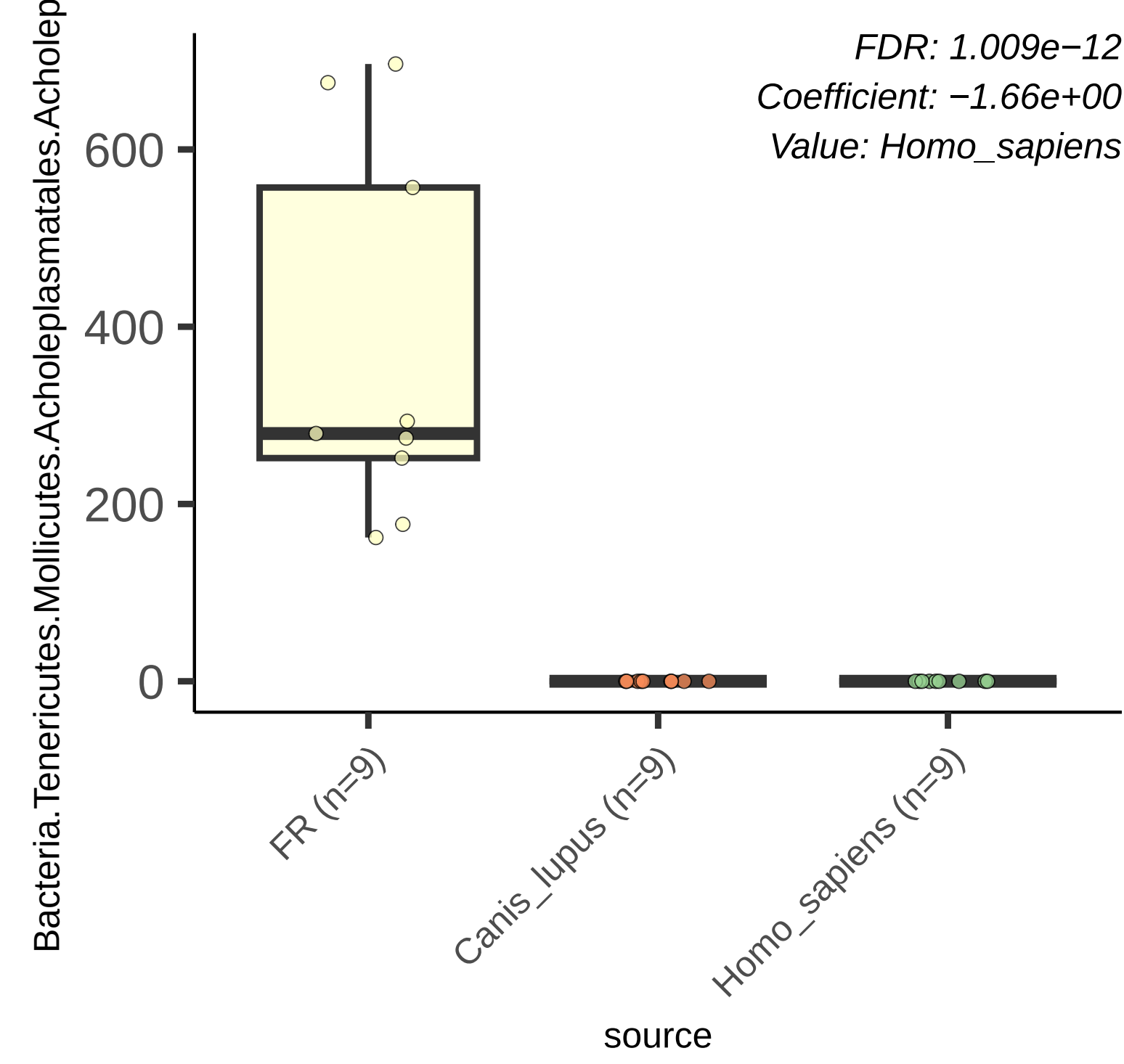
Canis_lupus (n=9)
Homo_sapiens (n=9)
source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Bacteria.Tenericutes.Mollicutes.Acholeplasmatales.Acholep
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200
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Canis_lupus (n=9)
Homo_sapiens (n=9)
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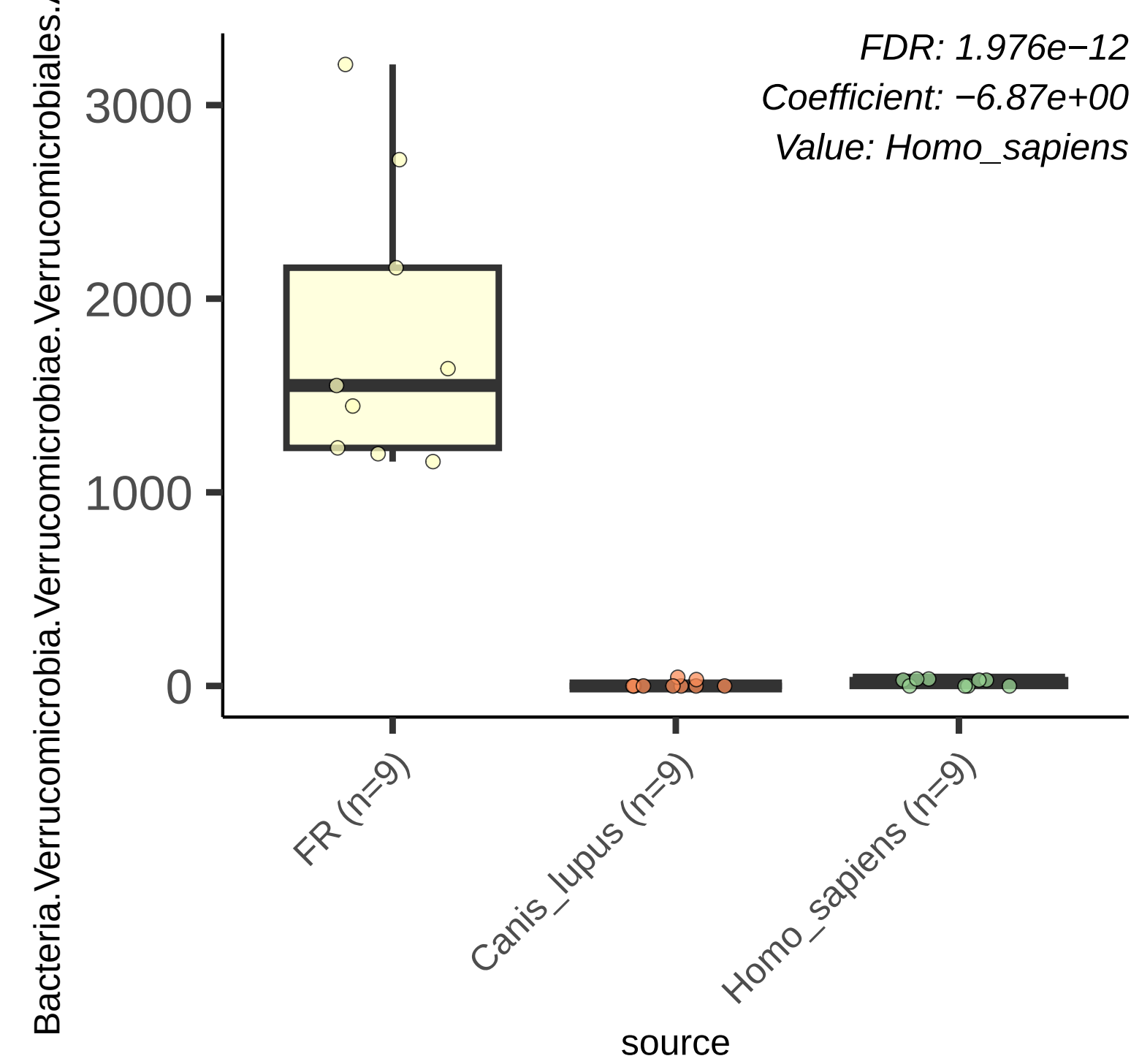
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2000
1000
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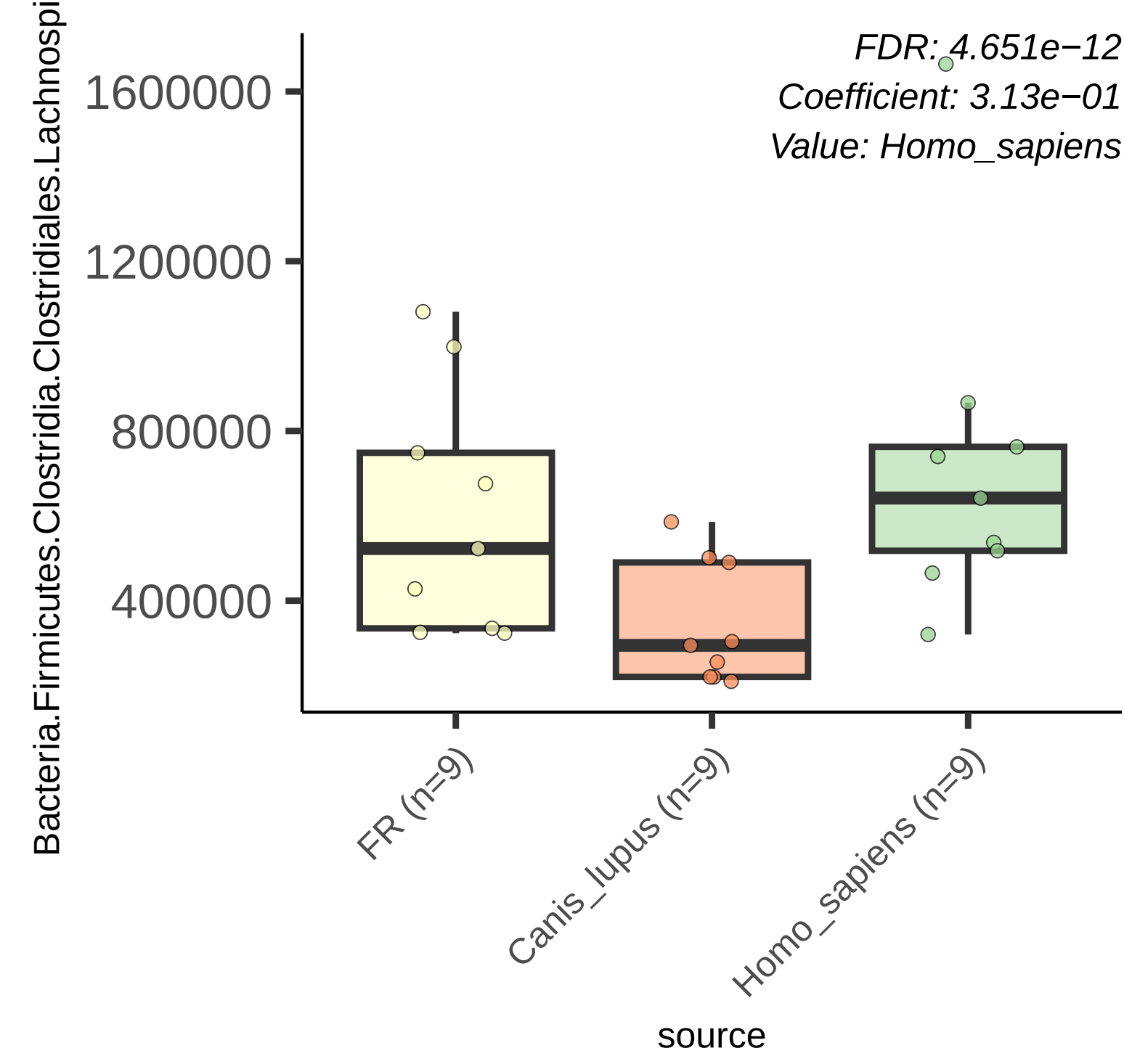
Canis_lupus (n=9)
Homo_sapiens (n=9)
source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800000
400000
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Canis_lupus (n=9)
Homo_sapiens (n=9)
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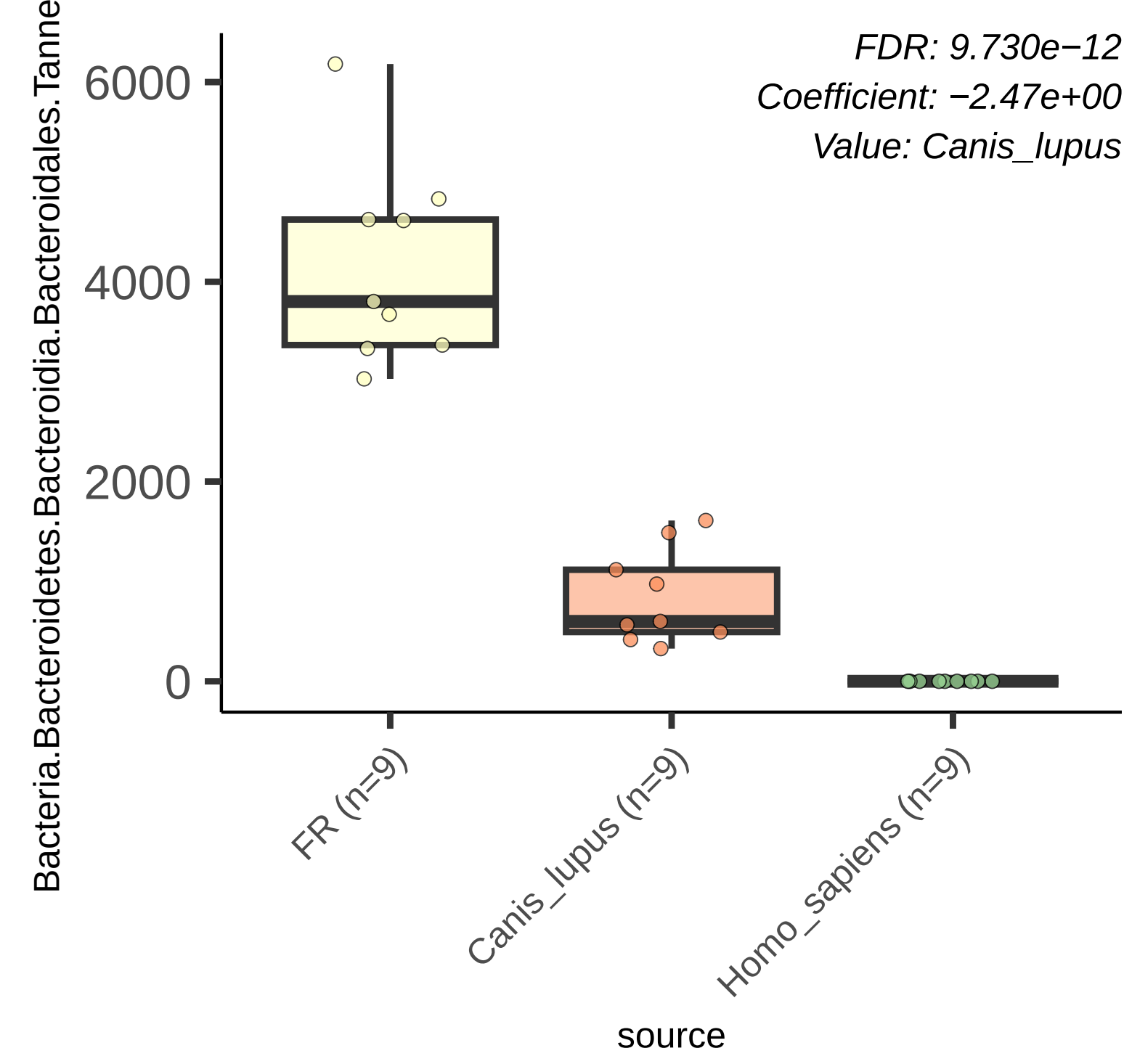
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Coefficient: −2.47e+00
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2000
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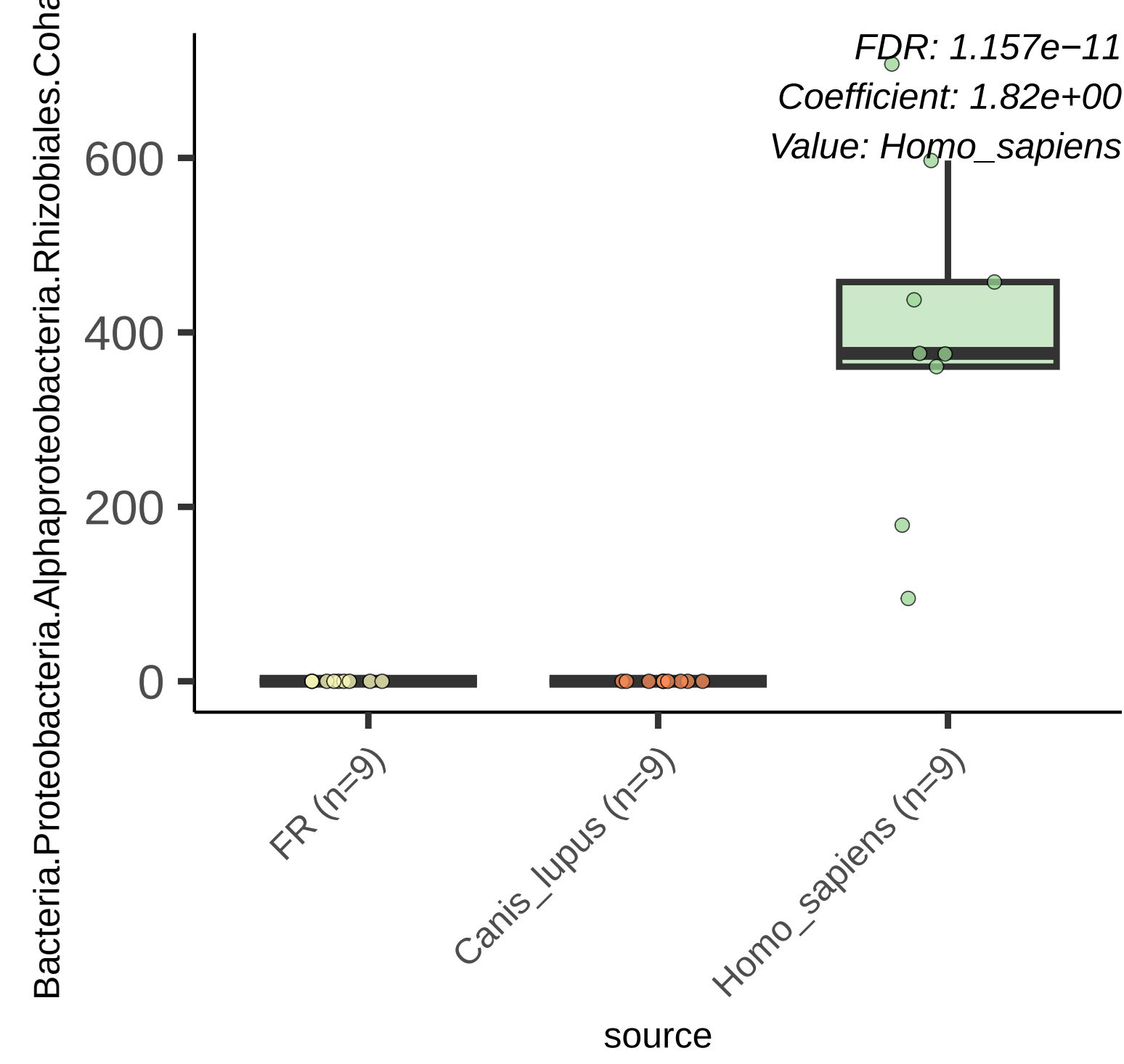
Homo_sapiens (n=9)
source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400
200
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Canis_lupus (n=9)
Homo_sapiens (n=9)
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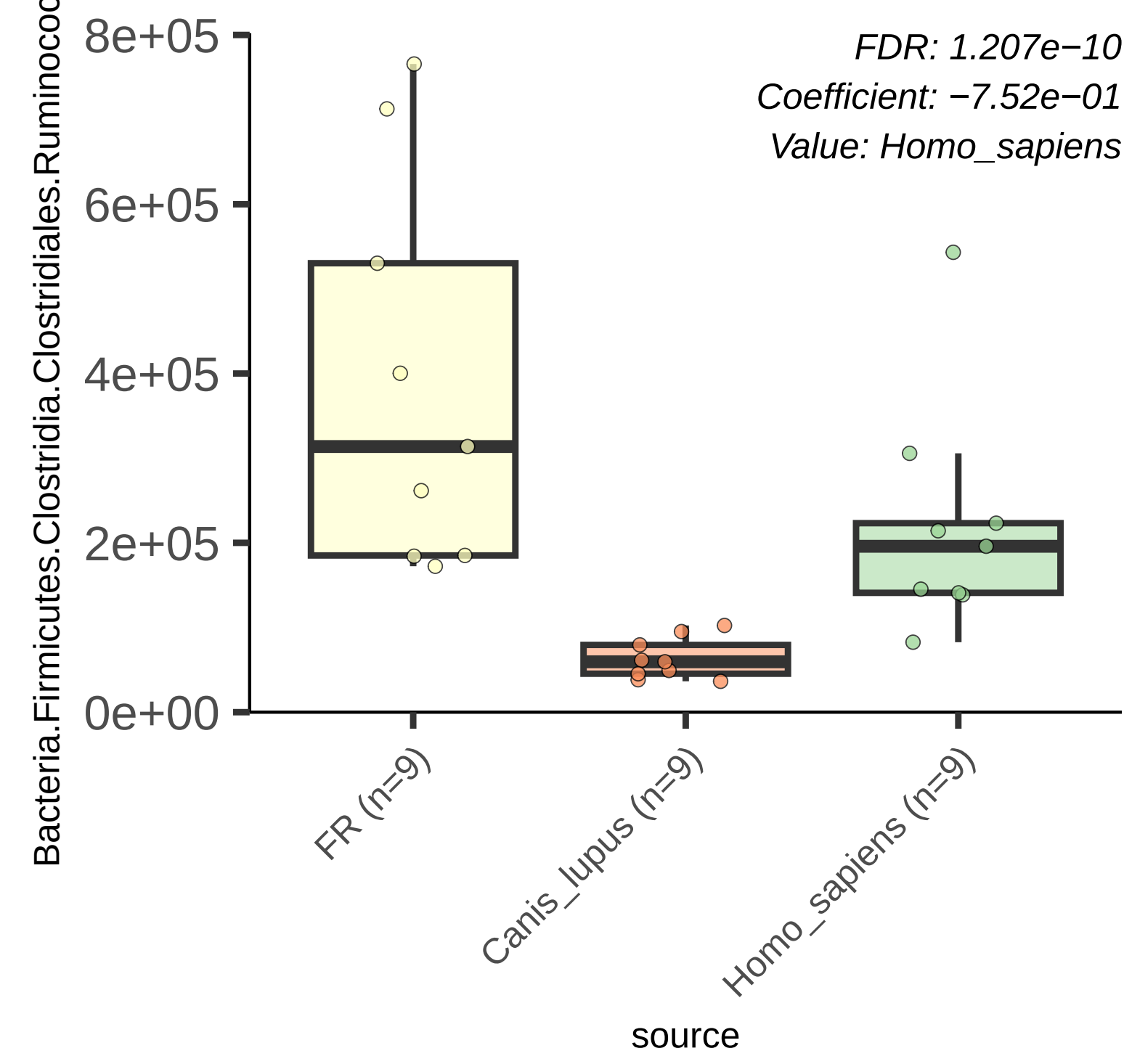
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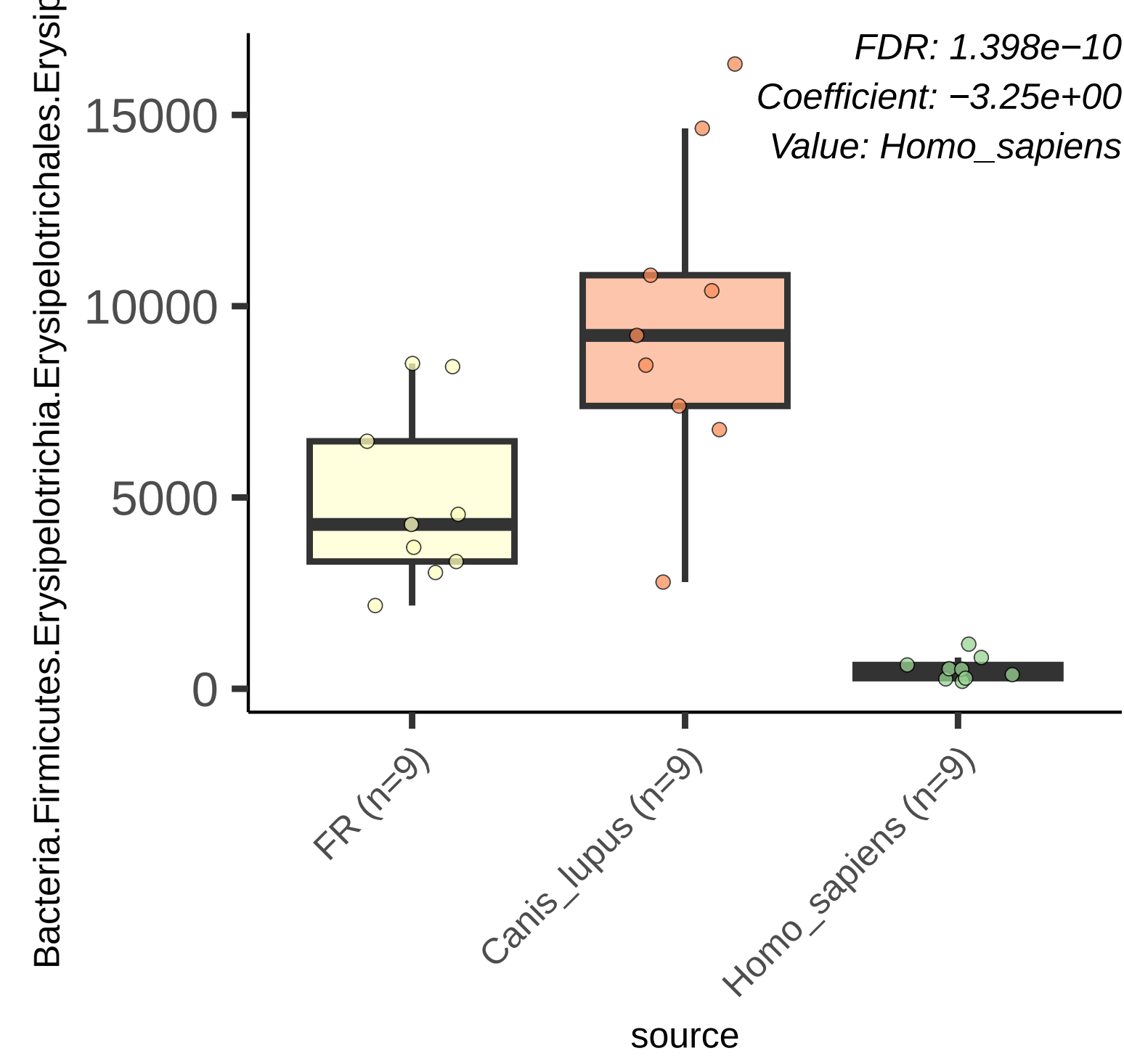
8e+05
6e+05
4e+05
2e+05
0e+00
FR (n=9)
Canis_lupus (n=9)
Homo_sapiens (n=9)
source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Coefficient: −3.25e+00
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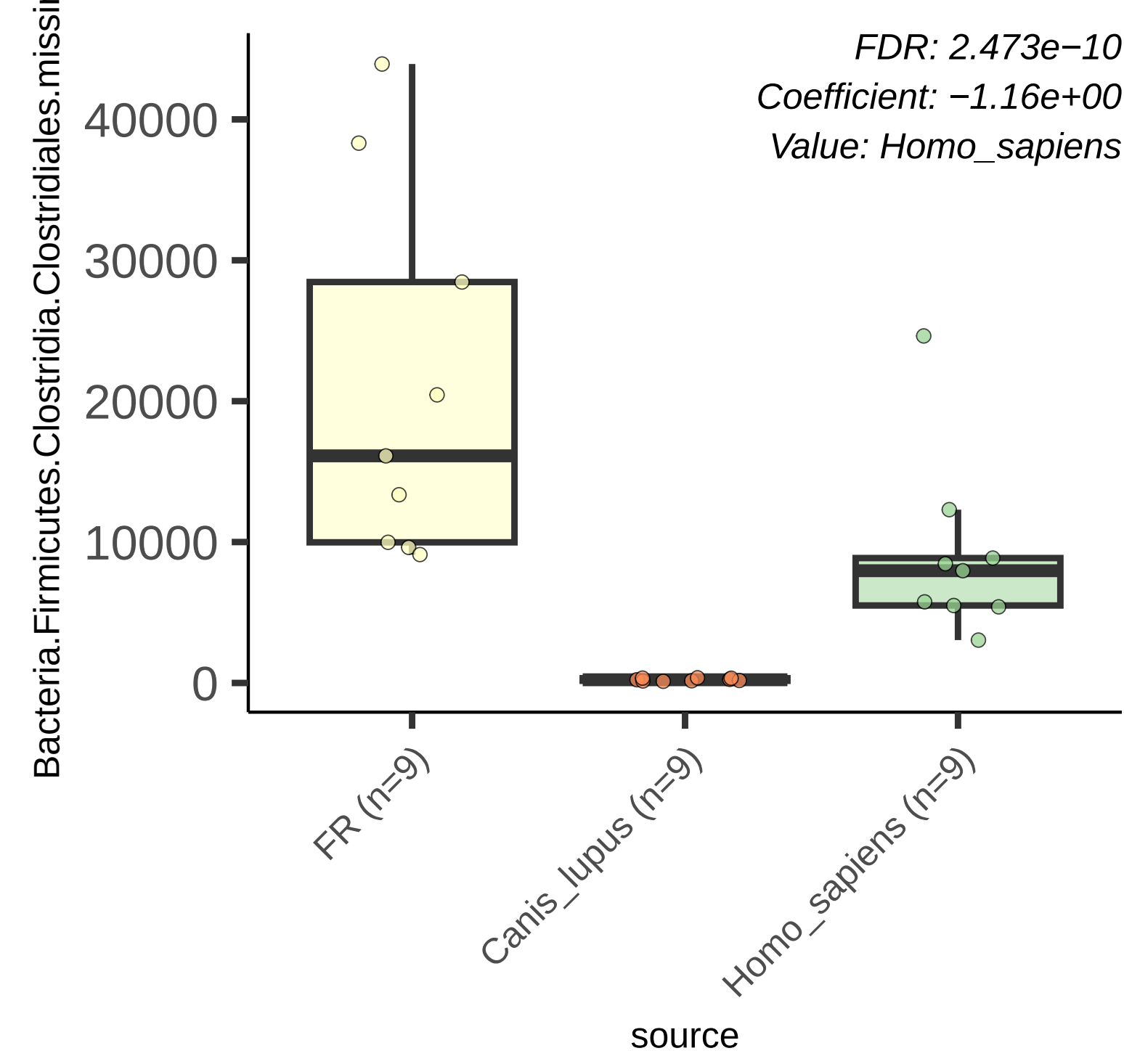
15000
10000
5000
0
Bacteria.Firmicutes.Erysipelotrichia.Erysipelotrichales.Erysip
FR (n=9)
Canis_lupus (n=9)
Homo_sapiens (n=9)
source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Bacteria.Firmicutes.Clostridia.Clostridiales.missi
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30000
20000
10000
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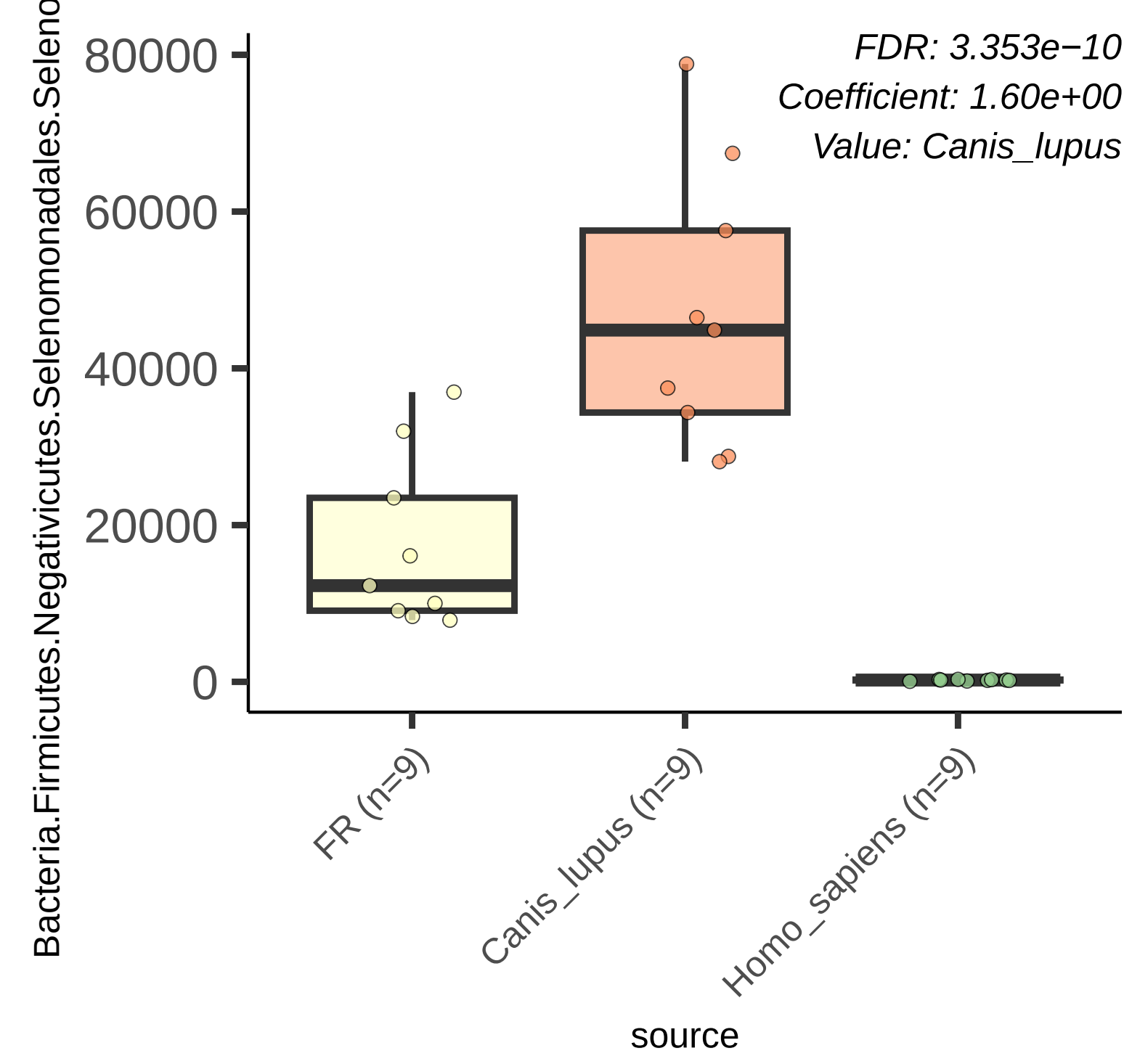
Canis_lupus (n=9)
Homo_sapiens (n=9)
source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40000
20000
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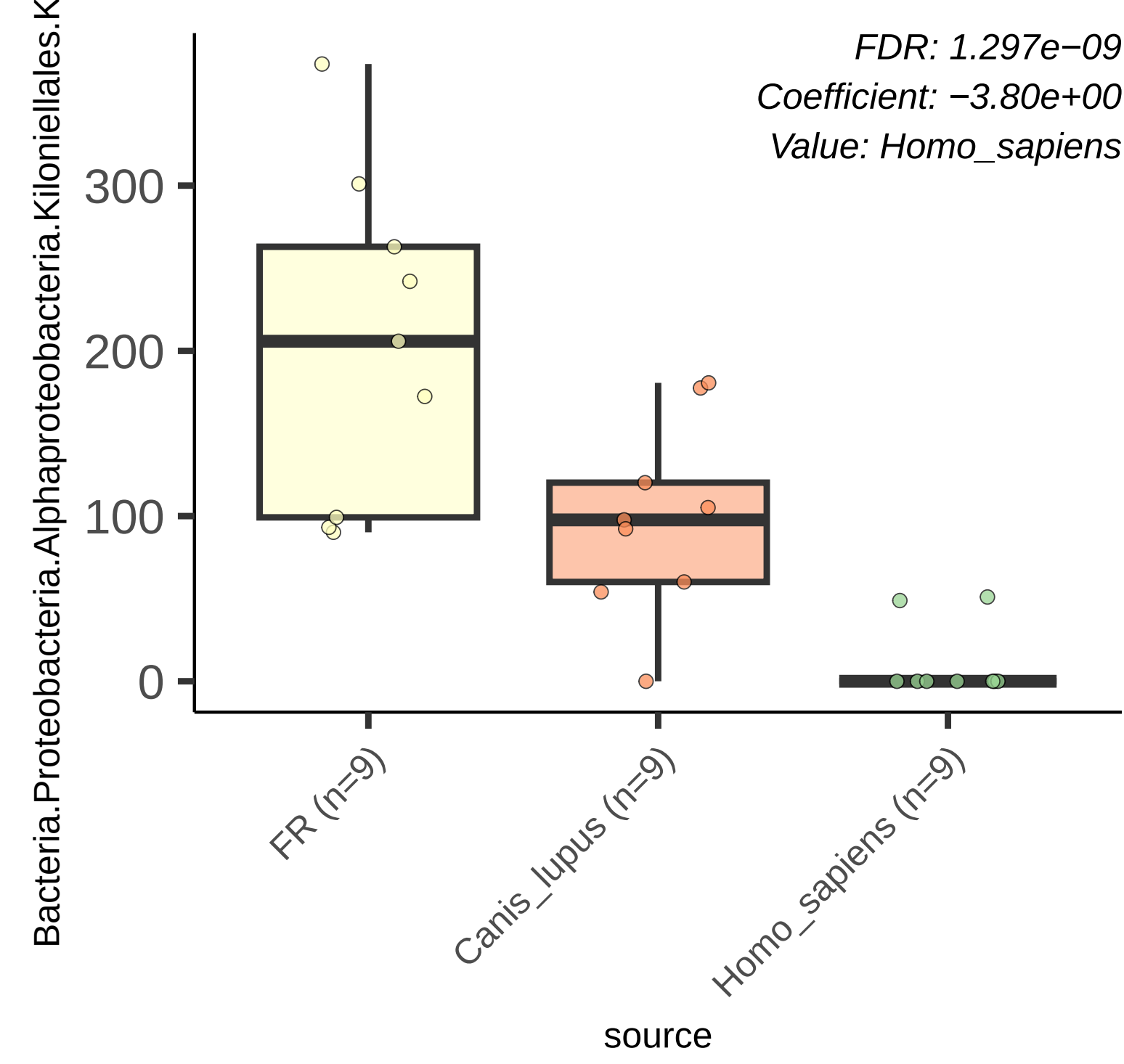
Canis_lupus (n=9)
Homo_sapiens (n=9)
source


FDR: 1.297e−09
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Bacteria.Proteobacteria.Alphaproteobacteria.Kiloniellales.K
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200
100
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Canis_lupus (n=9)
Homo_sapiens (n=9)
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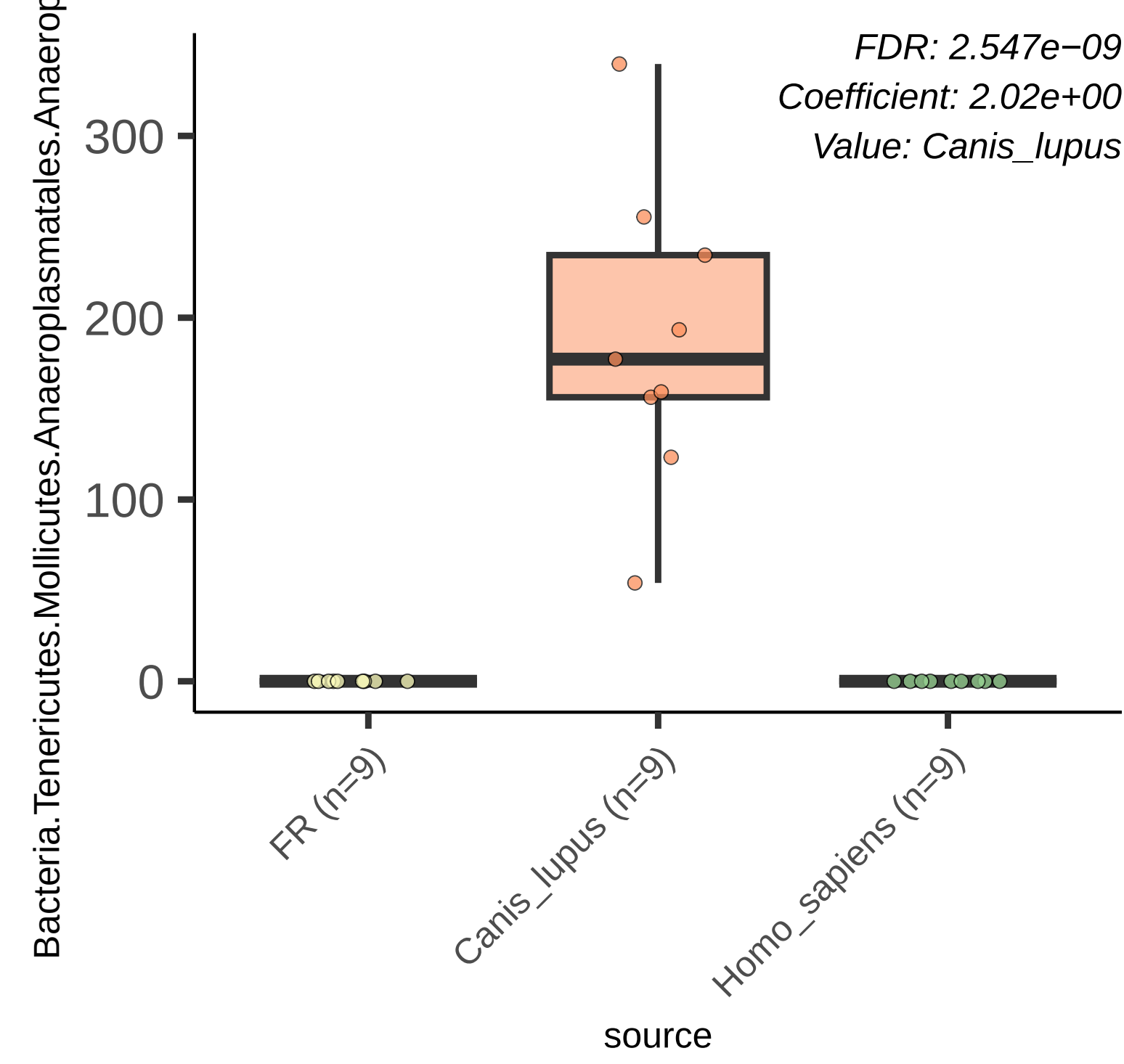
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Coefficient: 2.02e+00
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100
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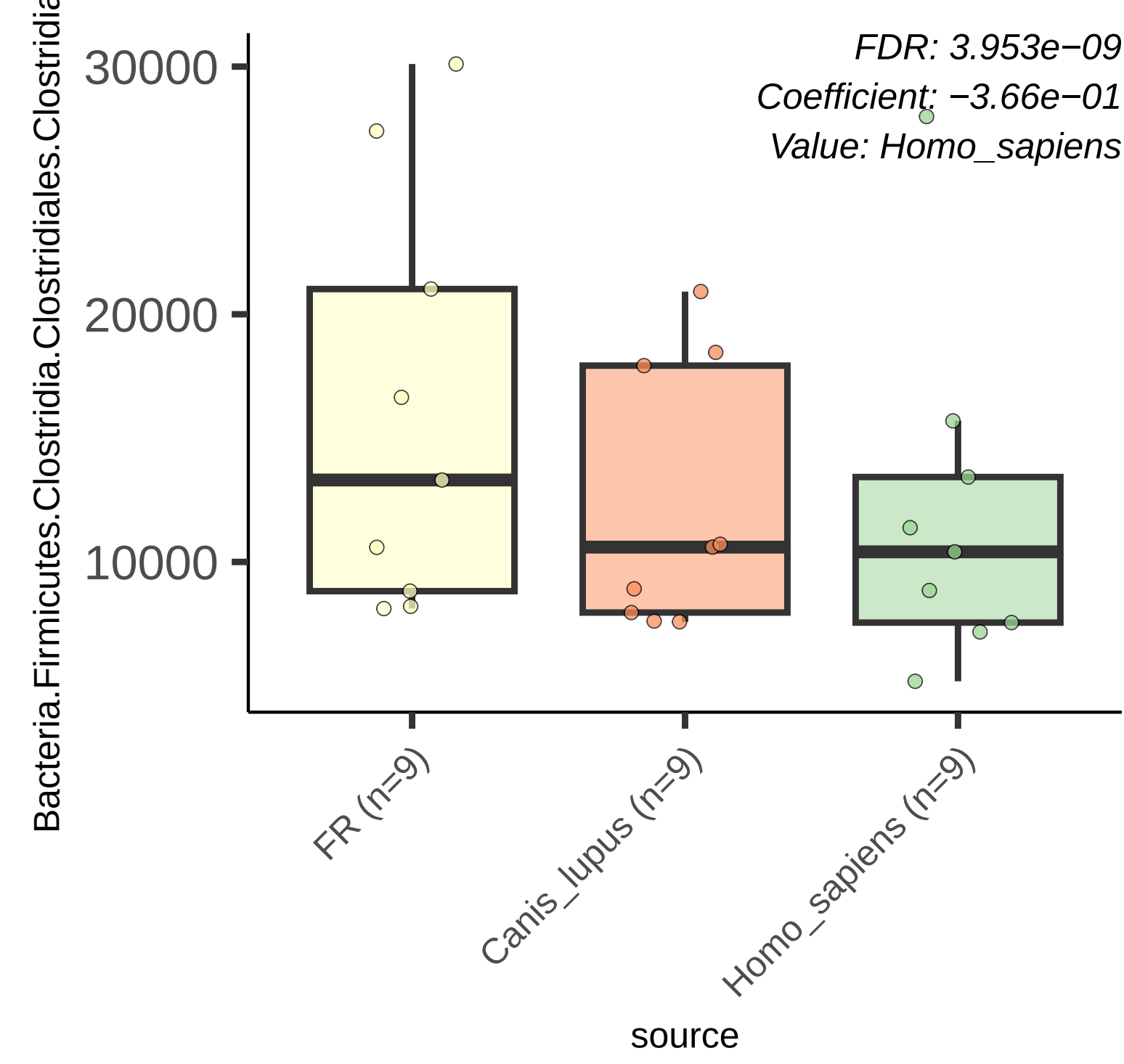
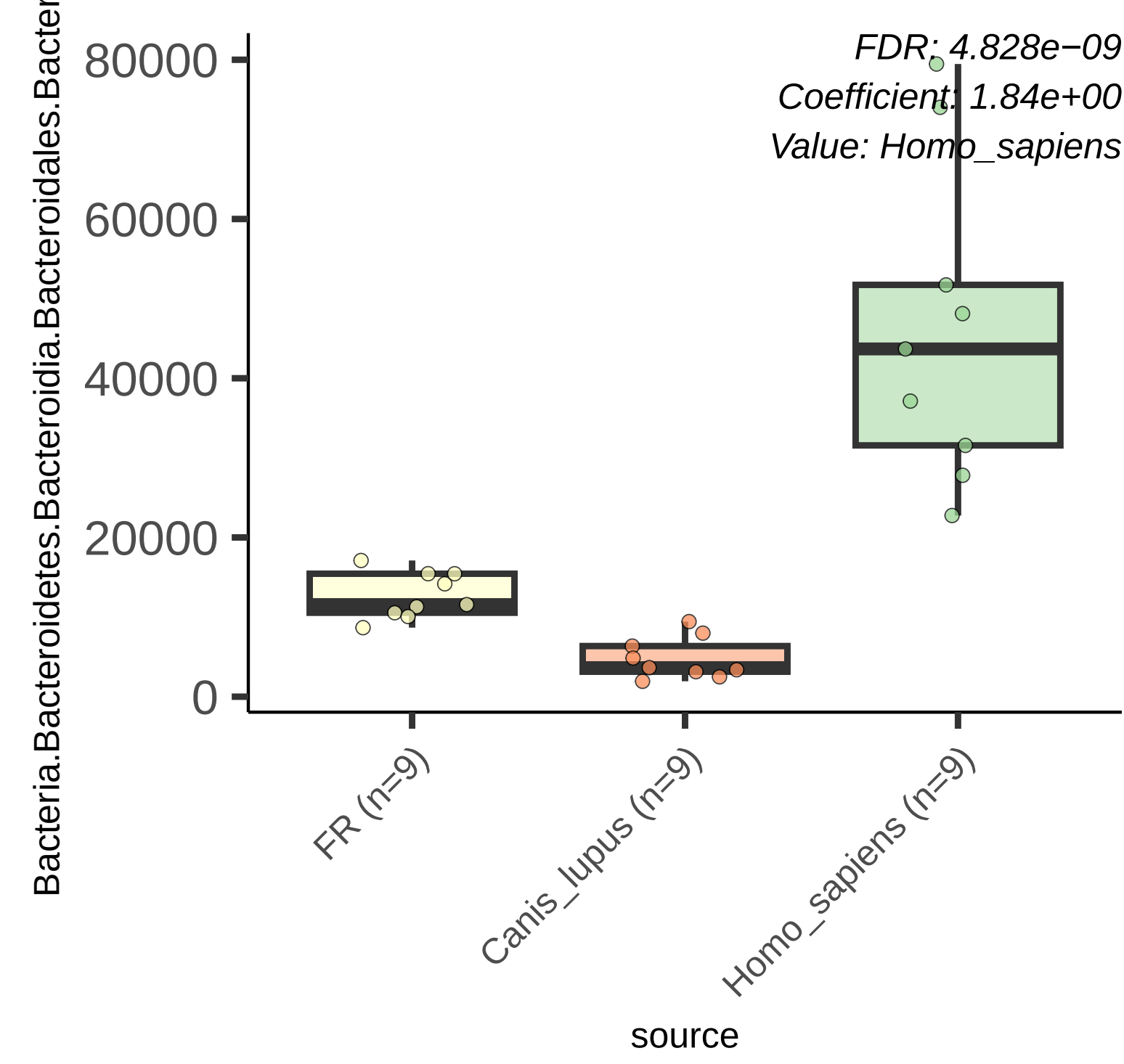
Canis_lupus (n=9)
Homo_sapiens (n=9)
source


FDR: 3.953e−09
Coefficient: −3.66e−01
Value: Homo_sapiens
Bacteria.Firmicutes.Clostridia.Clostridiales.Clostridia
30000
20000
10000
FR (n=9)
Canis_lupus (n=9)
Homo_sapiens (n=9)
source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Coefficient: 1.84e+00
Value: Homo_sapiens
Bacteria.Bacteroidetes.Bacteroidia.Bacteroidales.Bacter
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40000
20000
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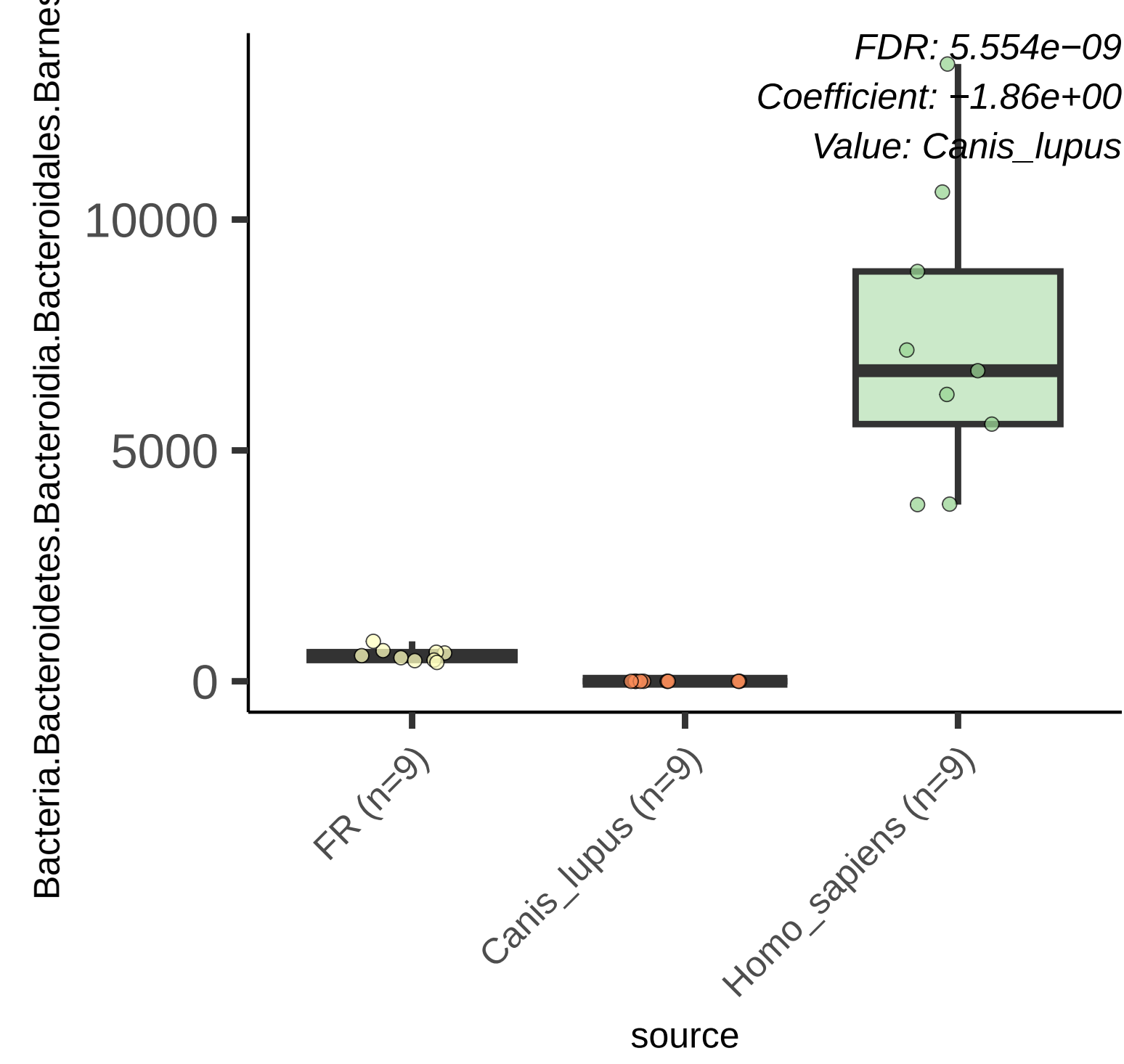
FR (n=9)
Canis_lupus (n=9)
Homo_sapiens (n=9)
source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Coefficient: −1.86e+00
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10000
5000
0
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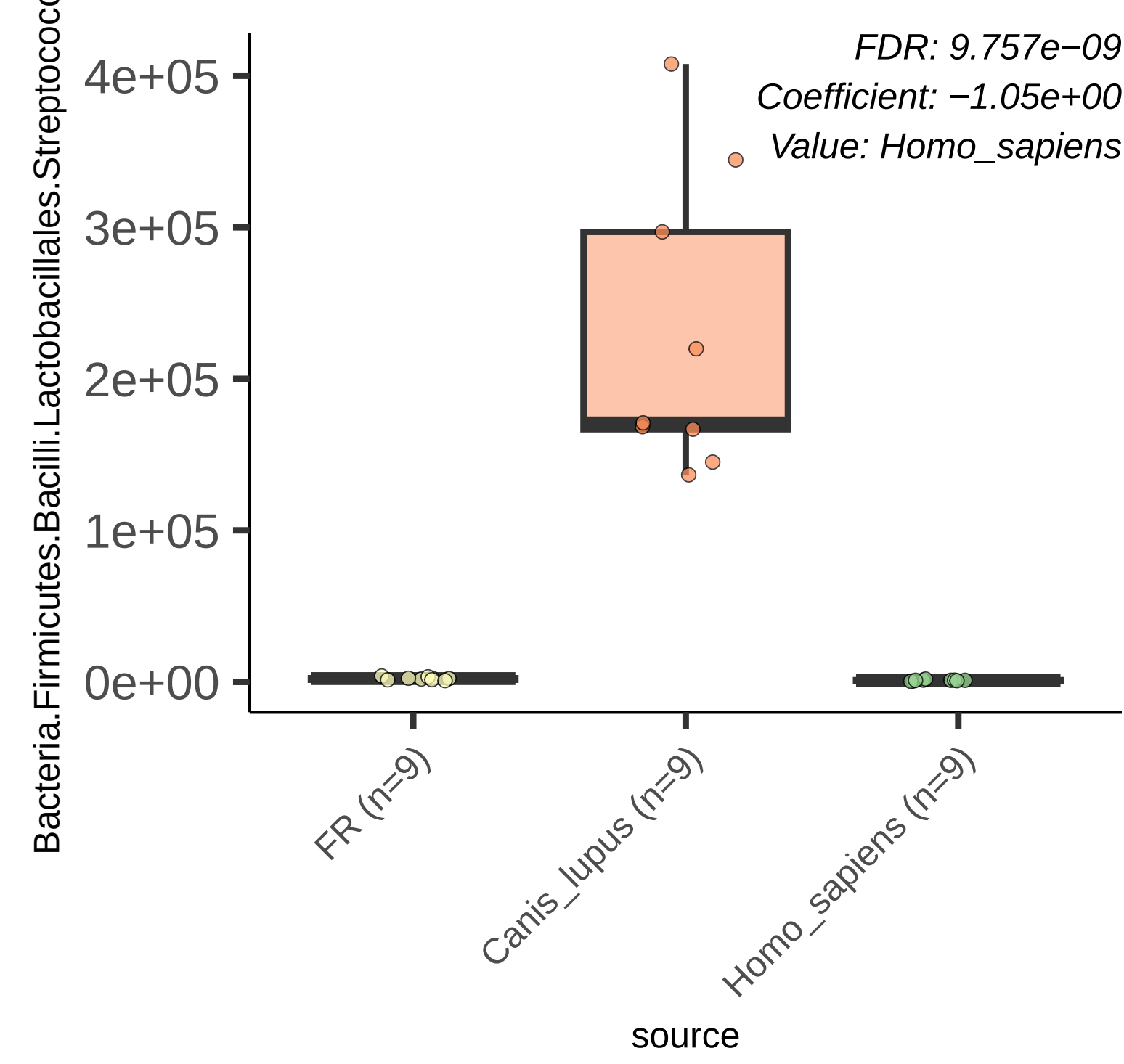
Canis_lupus (n=9)
Homo_sapiens (n=9)
source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4e+05
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1e+05
0e+00
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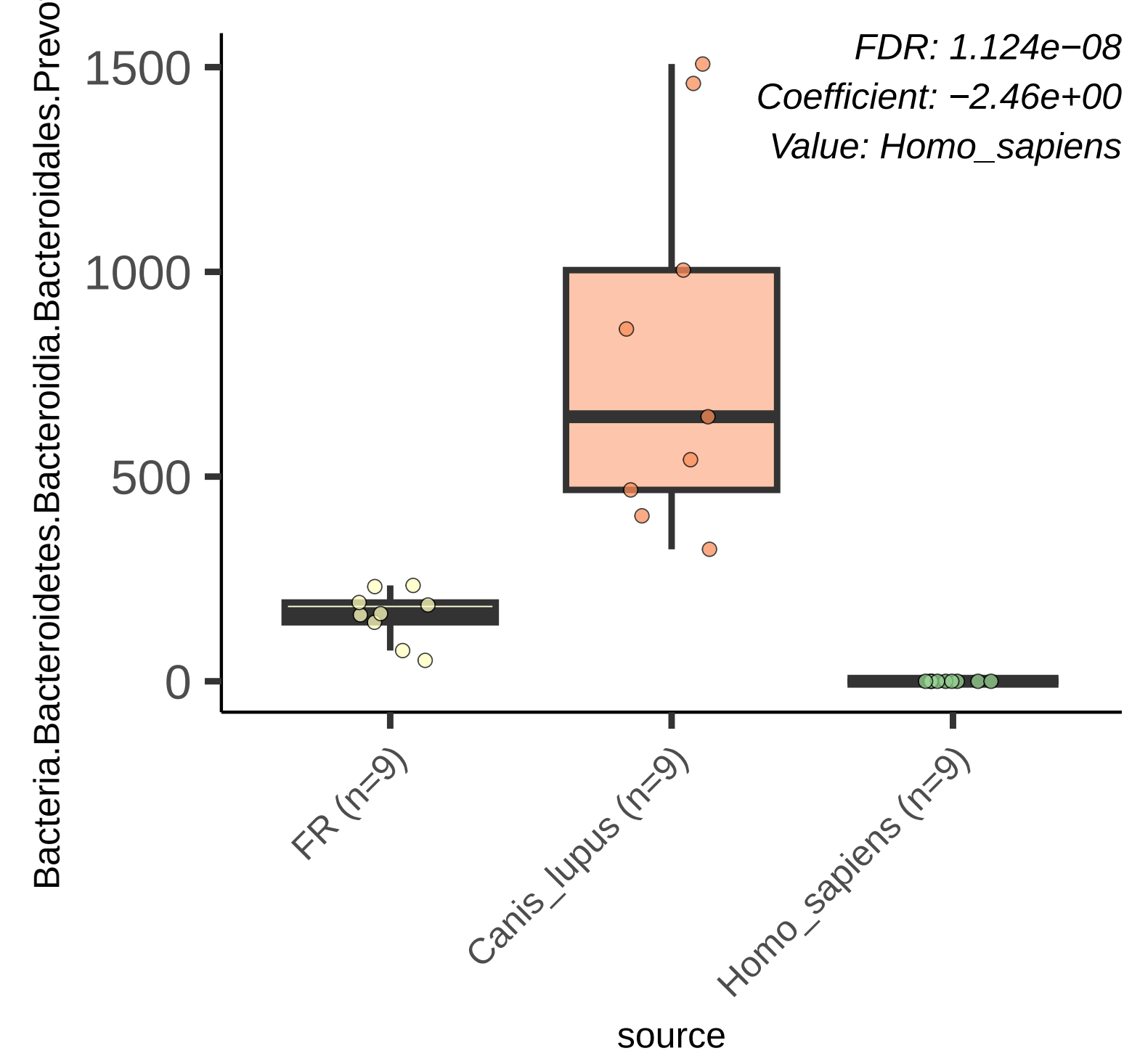
Canis_lupus (n=9)
Homo_sapiens (n=9)
source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Coefficient: −2.46e+00
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Bacteria.Bacteroidetes.Bacteroidia.Bacteroidales.Prevo
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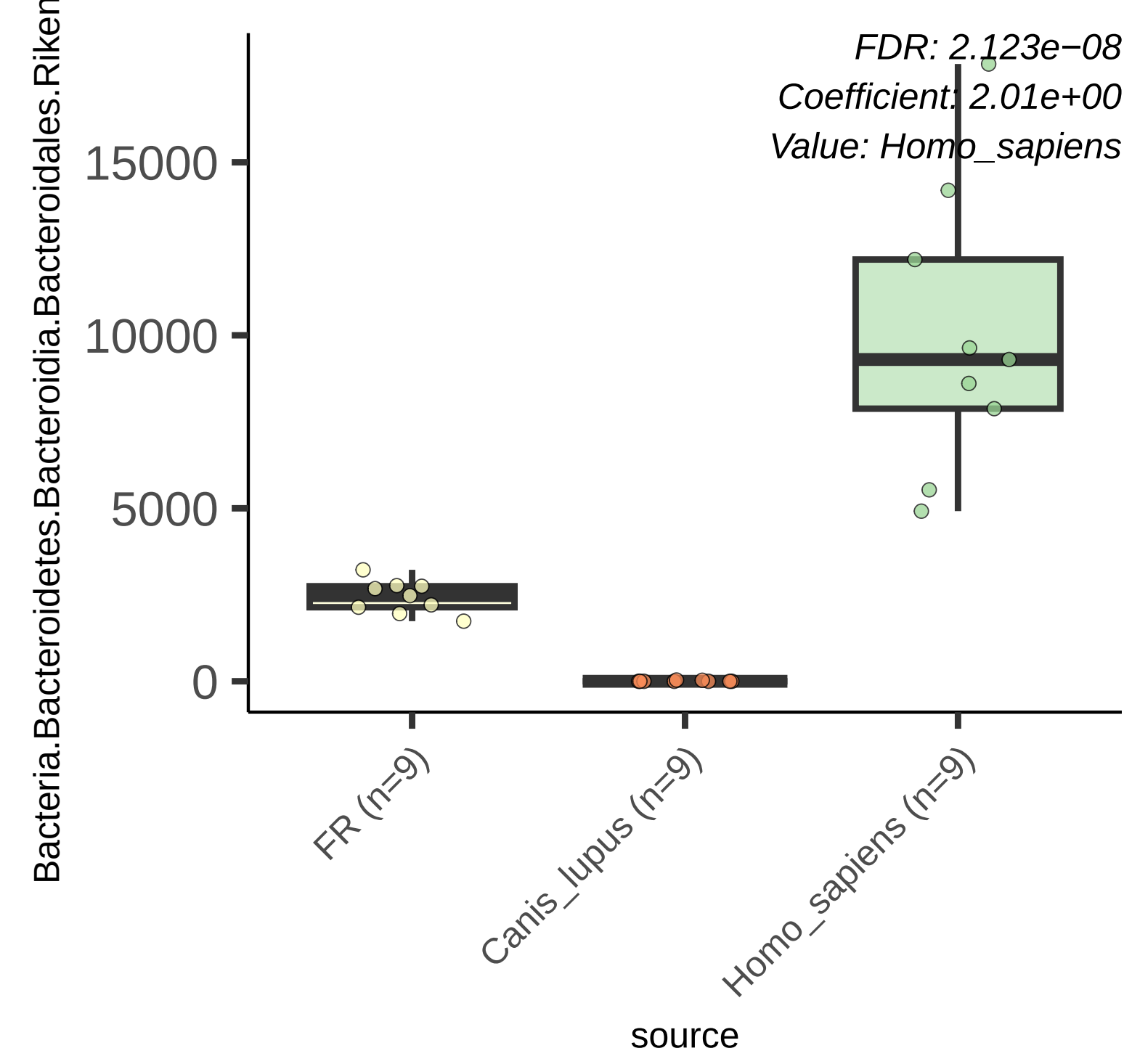
Canis_lupus (n=9)
Homo_sapiens (n=9)
source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Bacteria.Bacteroidetes.Bacteroidia.Bacteroidales.Riken
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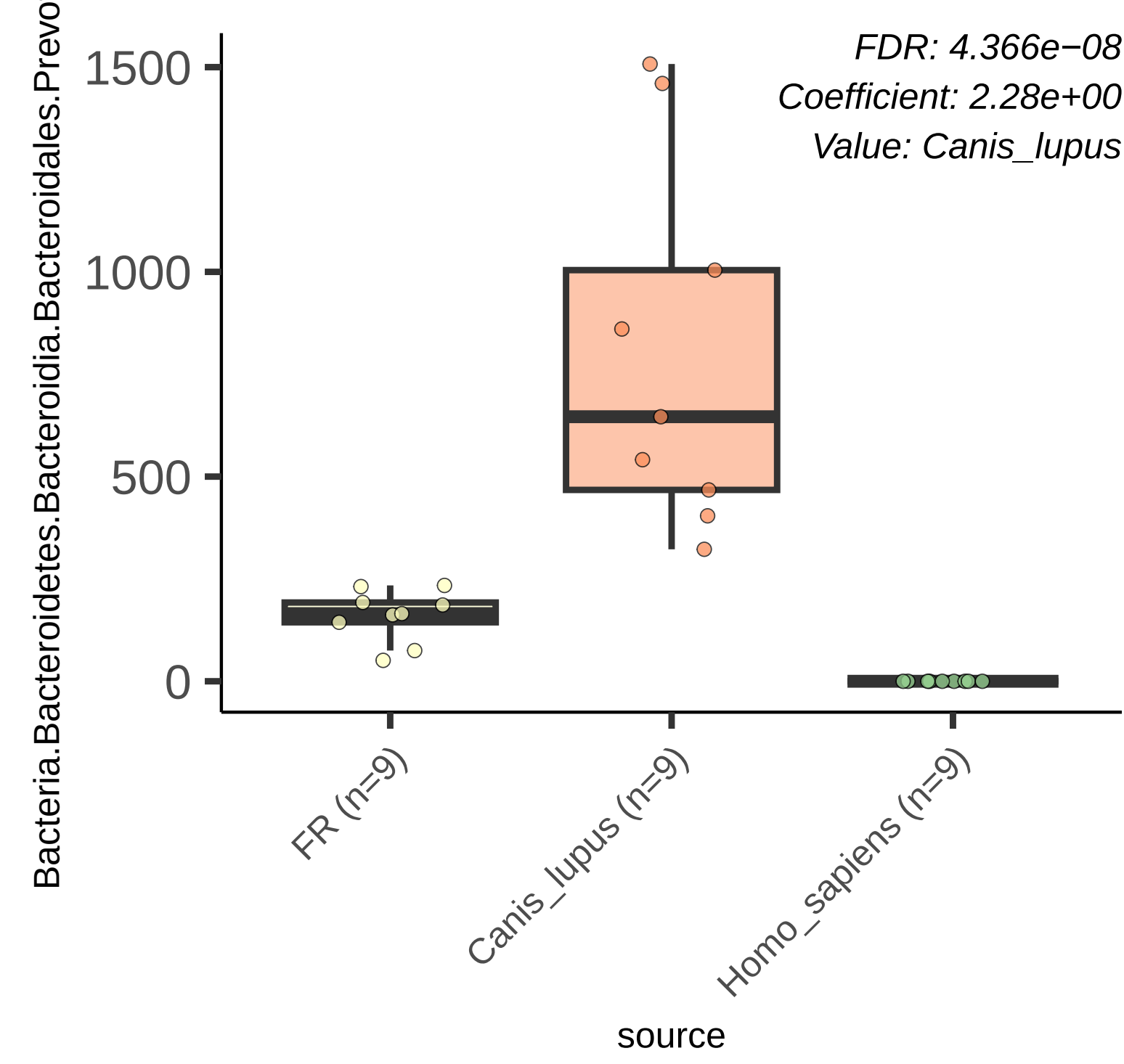
Canis_lupus (n=9)
Homo_sapiens (n=9)
source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Bacteria.Bacteroidetes.Bacteroidia.Bacteroidales.Prevo
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1000
500
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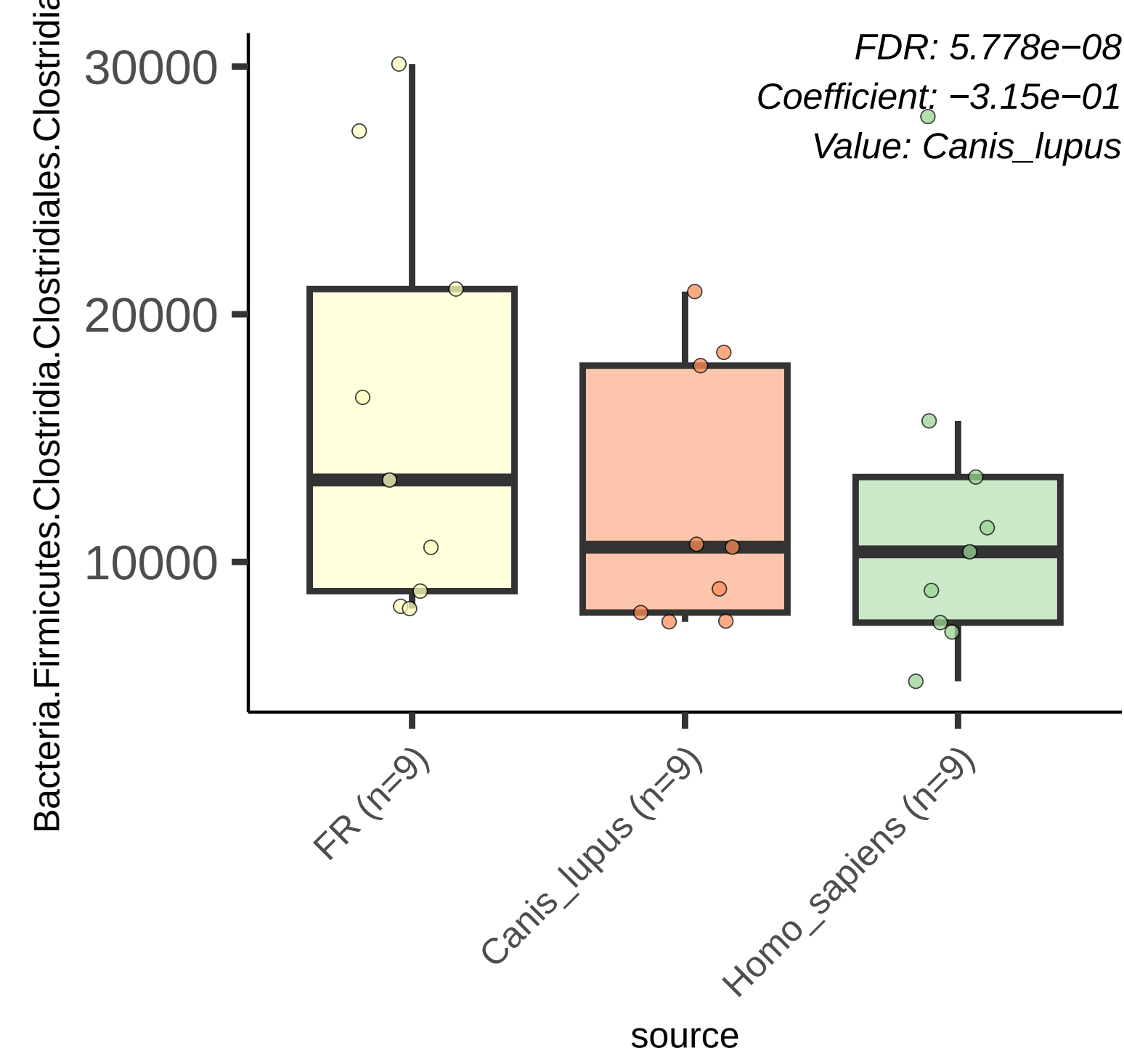
Canis_lupus (n=9)
Homo_sapiens (n=9)
source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Coefficient: −3.15e−01
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30000
20000
10000
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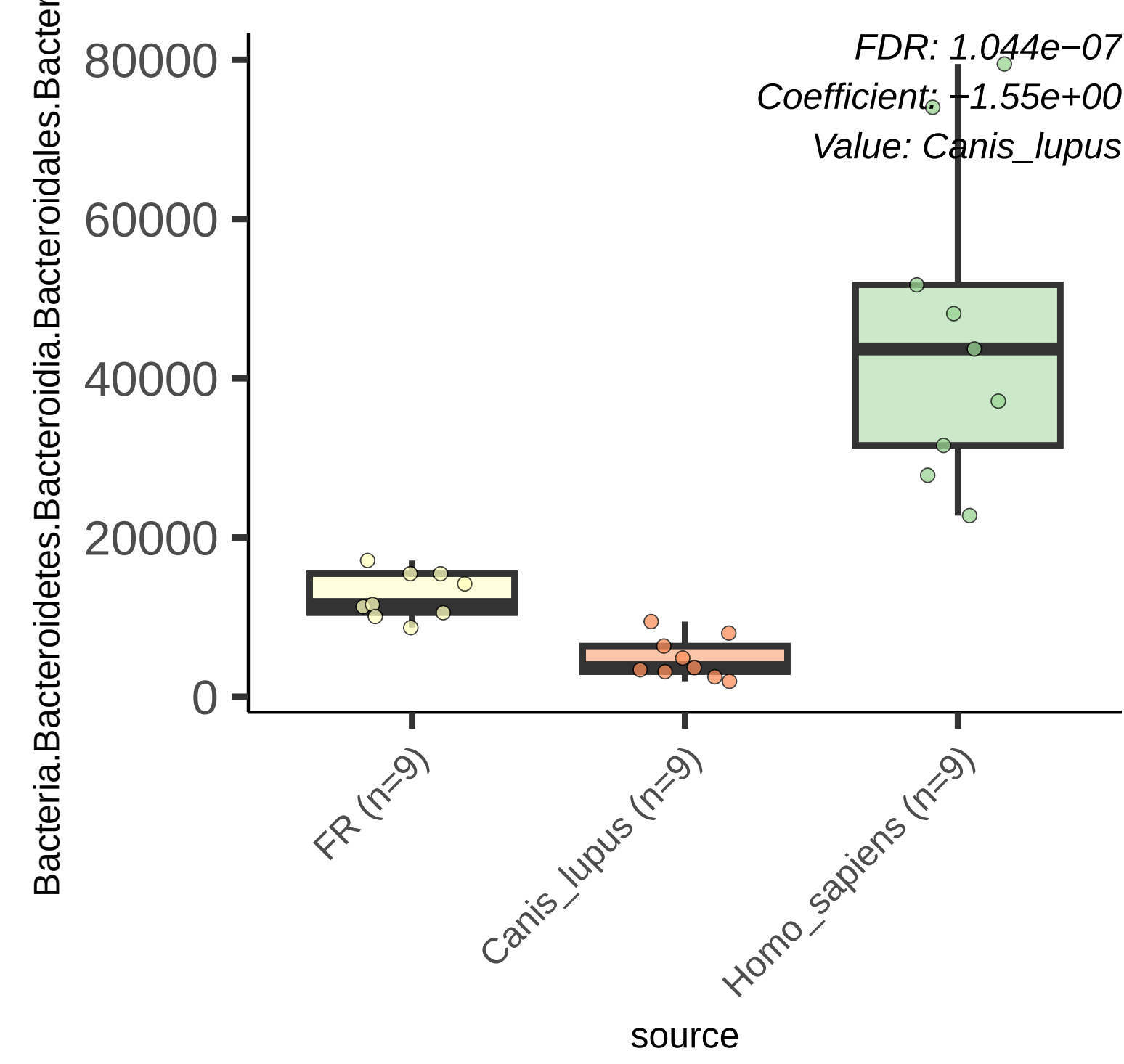
FR (n=9)
Canis_lupus (n=9)
Homo_sapiens (n=9)
source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60000
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20000
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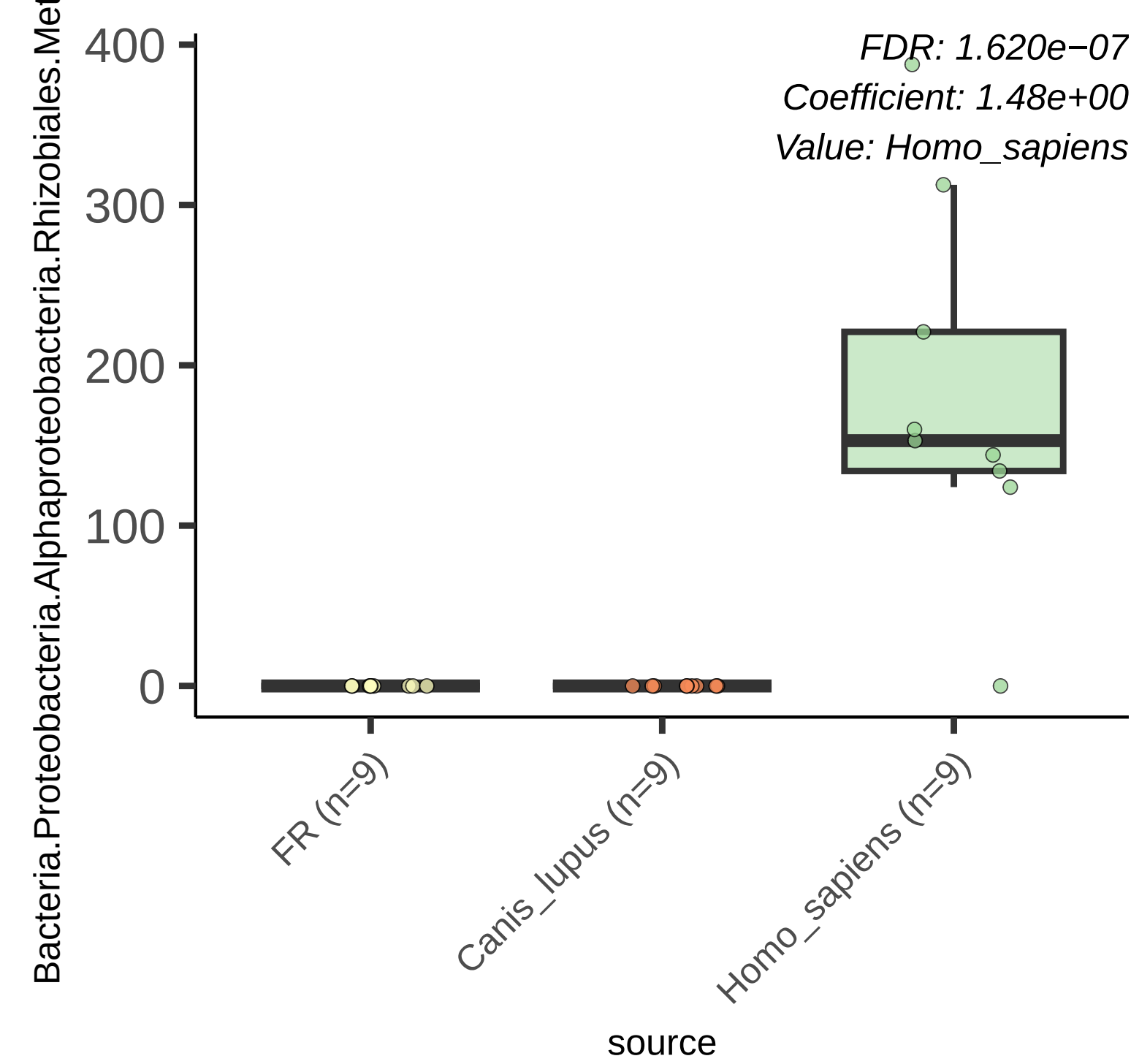
FR (n=9)
Canis_lupus (n=9)
Homo_sapiens (n=9)
source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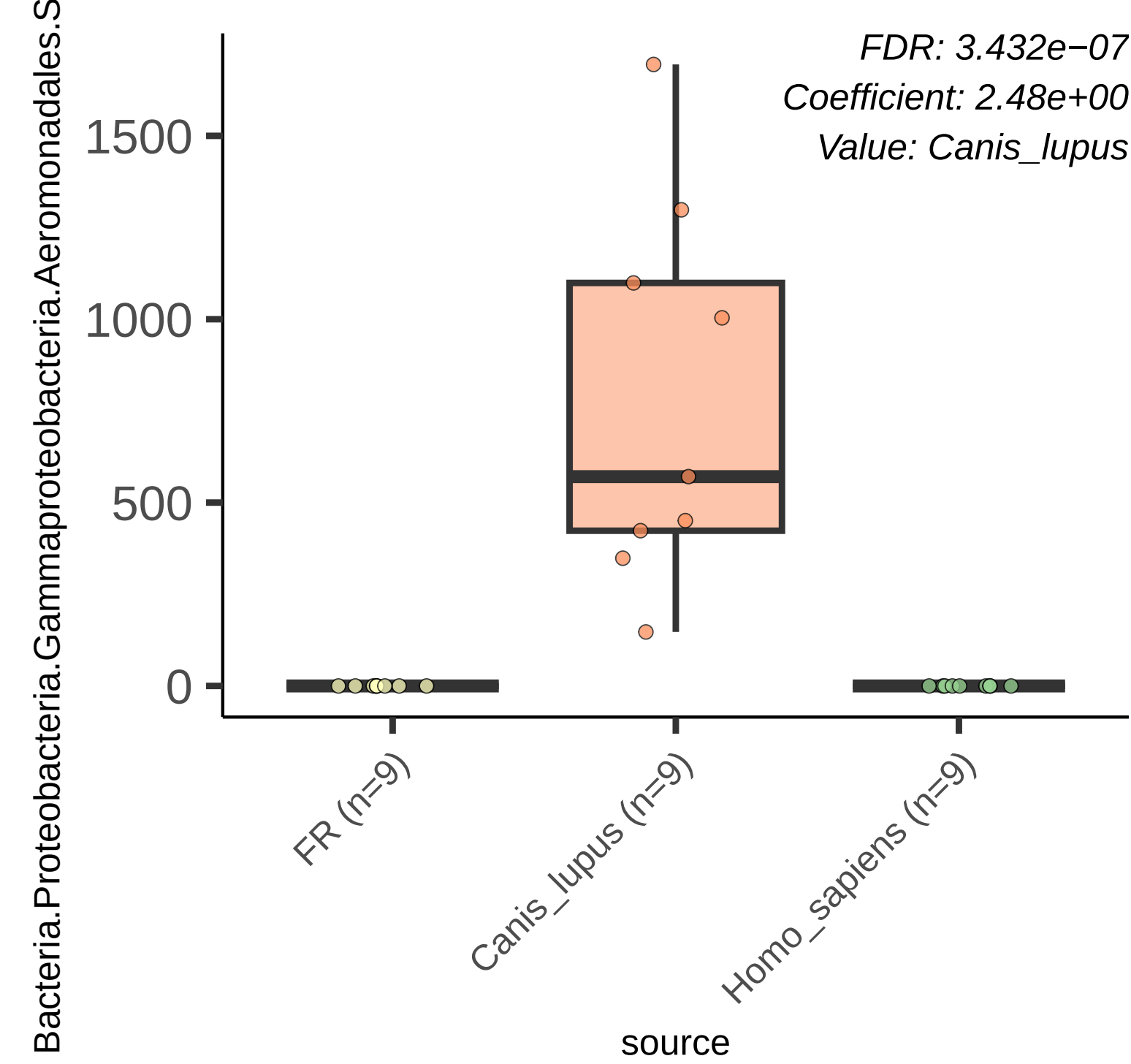
Coefficient: 1.48e+00
Value: Homo_sapiens
Bacteria.Proteobacteria.Alphaproteobacteria.Rhizobiales.Met
400
300
200
100
0
FR (n=9)
Canis_lupus (n=9)
Homo_sapiens (n=9)
source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Coefficient: 2.48e+00
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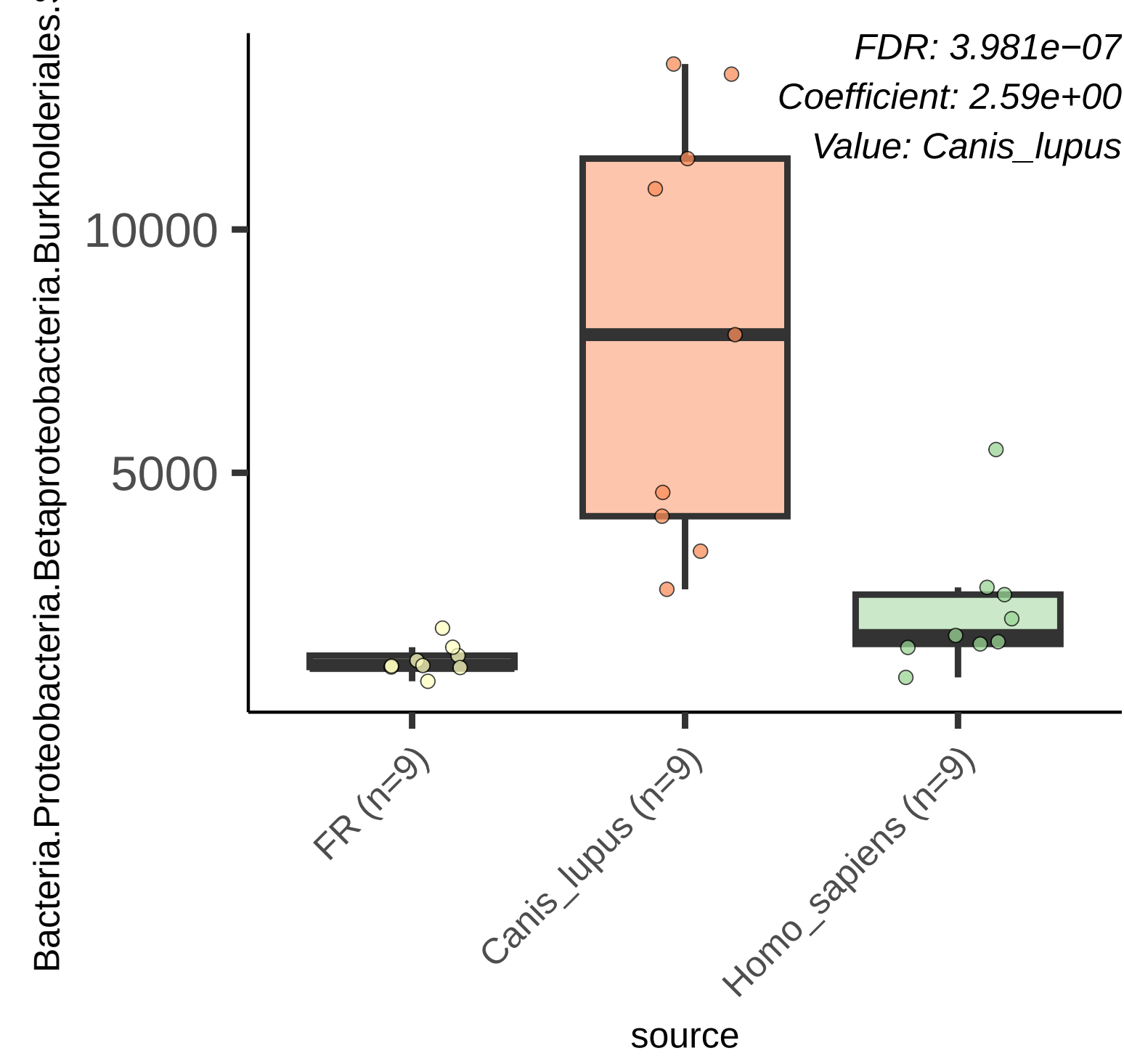
Bacteria.Proteobacteria.Gammaproteobacteria.Aeromonadales.S
1500
1000
500
0
FR (n=9)
Canis_lupus (n=9)
Homo_sapiens (n=9)
source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Coefficient: 2.59e+00
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Bacteria.Proteobacteria.Betaproteobacteria.Burkholderiales.
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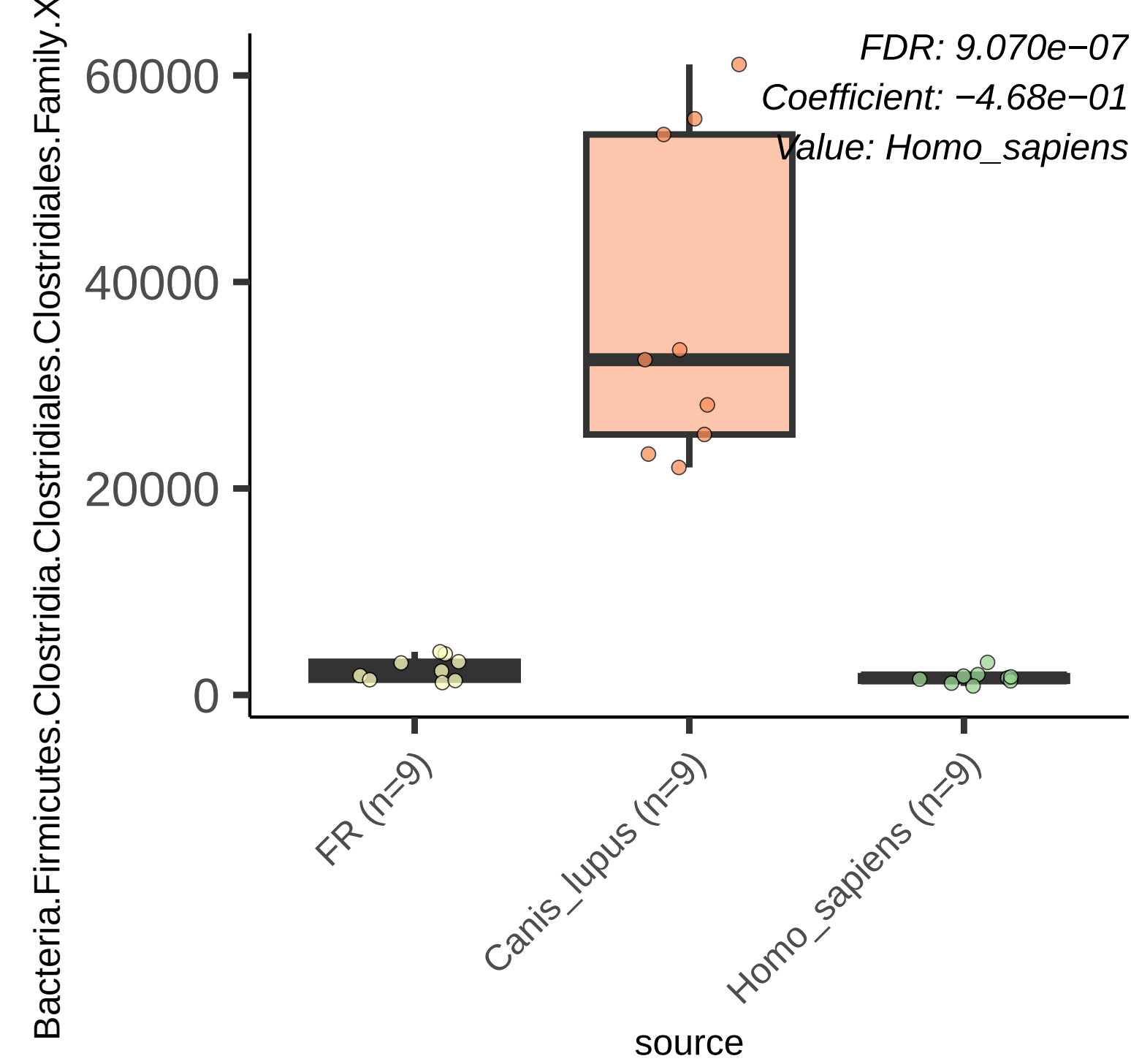
Canis_lupus (n=9)
Homo_sapiens (n=9)
source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Coefficient: −4.68e−01
Value: Homo_sapiens
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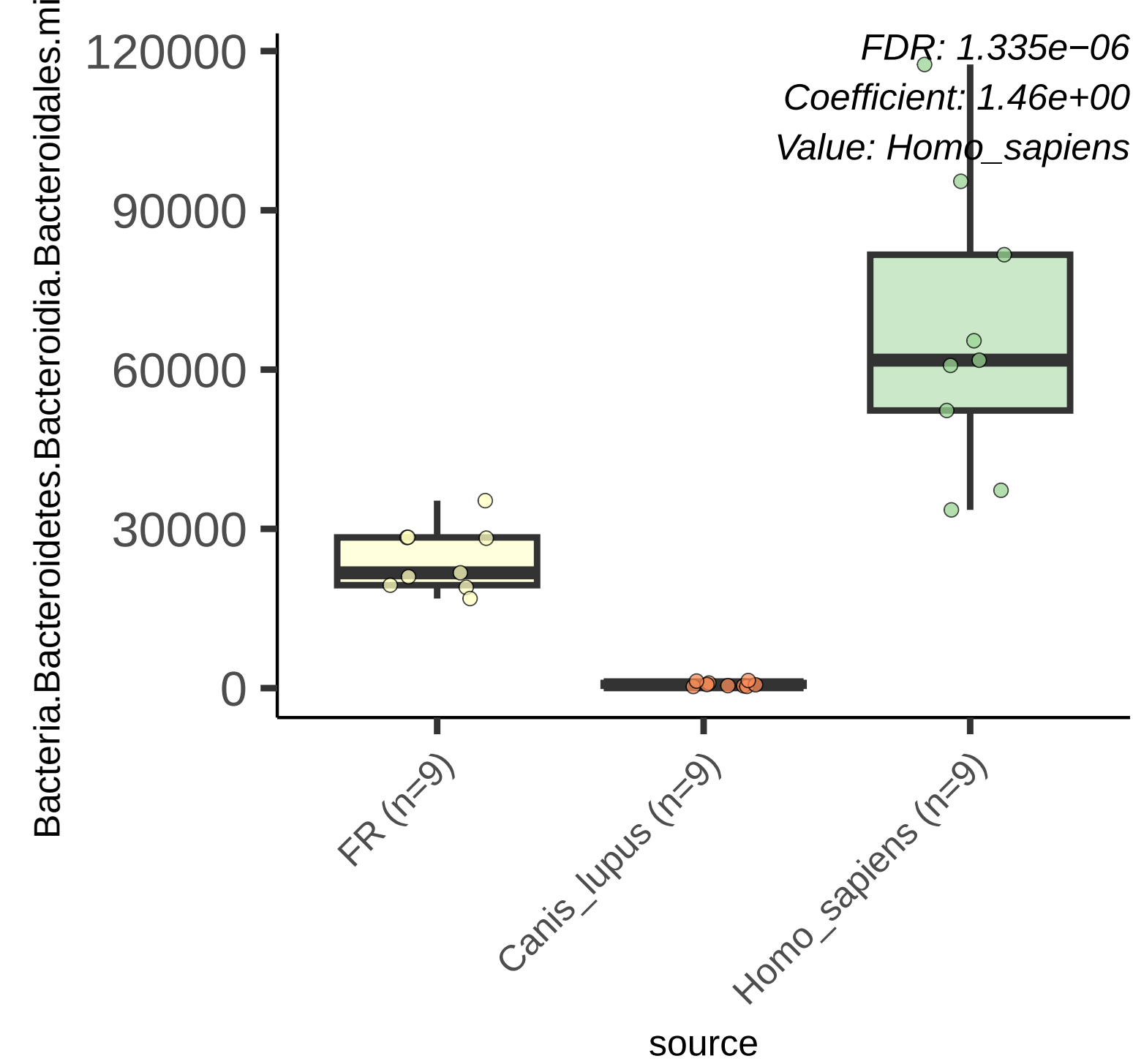
40000
20000
0
Bacteria.Firmicutes.Clostridia.Clostridiales.Clostridiales.Family.X
FR (n=9)
Canis_lupus (n=9)
Homo_sapiens (n=9)
source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Coefficient: 1.46e+00
Value: Homo_sapiens
Bacteria.Bacteroidetes.Bacteroidia.Bacteroidales.mi
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90000
60000
30000
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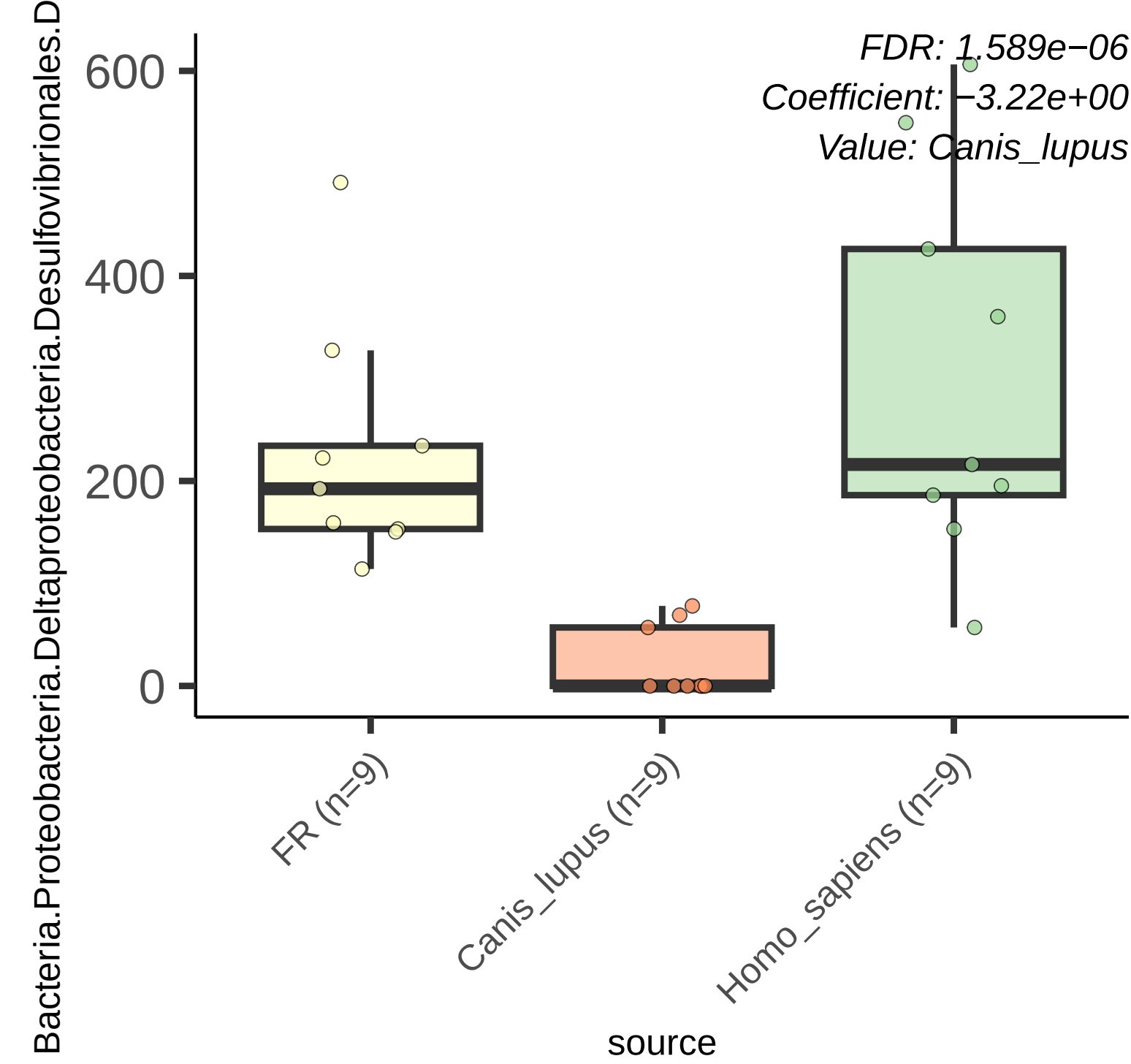
FR (n=9)
Canis_lupus (n=9)
Homo_sapiens (n=9)
source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Coefficient: −3.22e+00
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600
400
200
0
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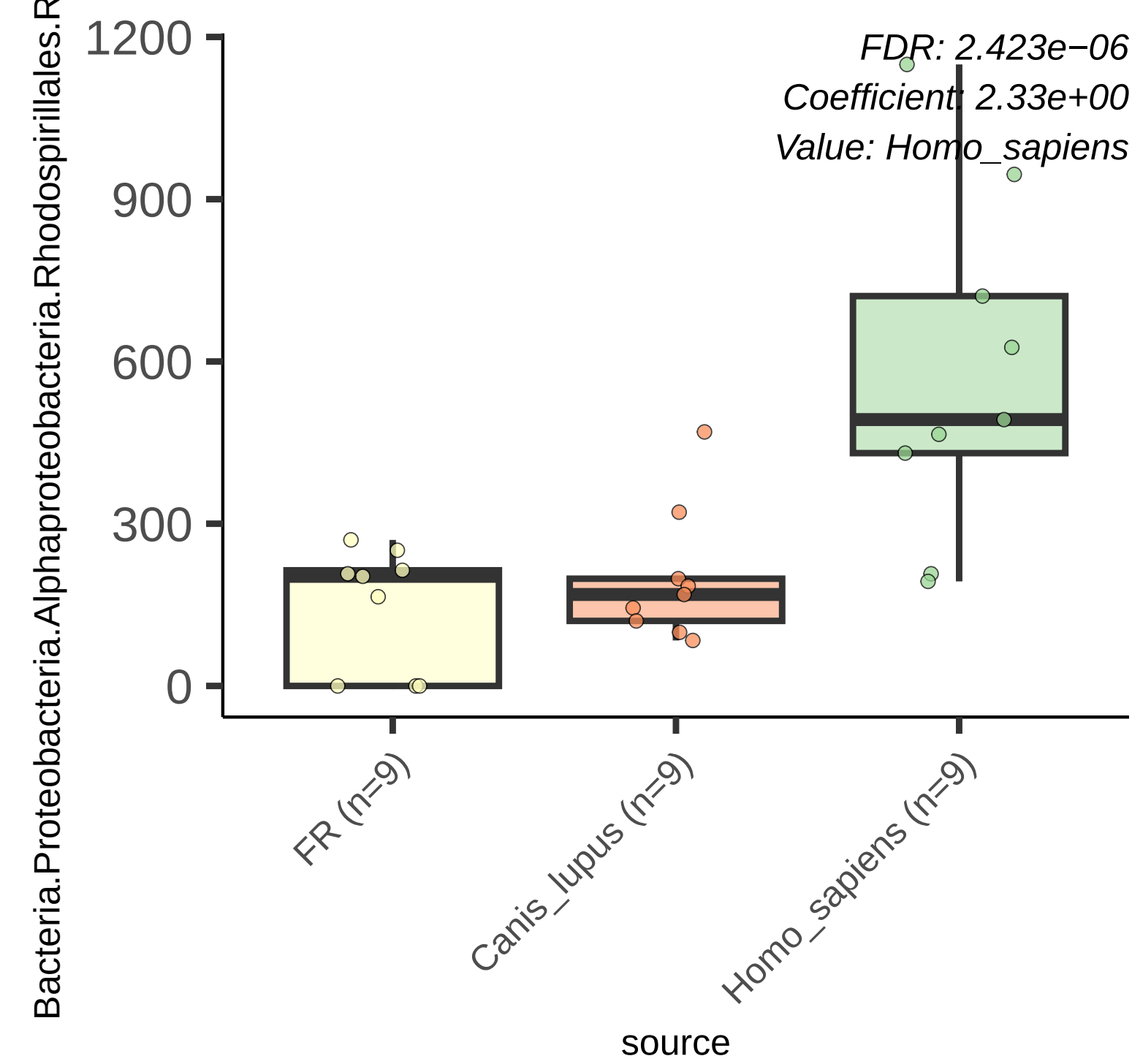
FR (n=9)
Canis_lupus (n=9)
Homo_sapiens (n=9)
source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Coefficient: 2.33e+00
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1200
900
600
300
0
Bacteria.Proteobacteria.Alphaproteobacteria.Rhodospirillales.R
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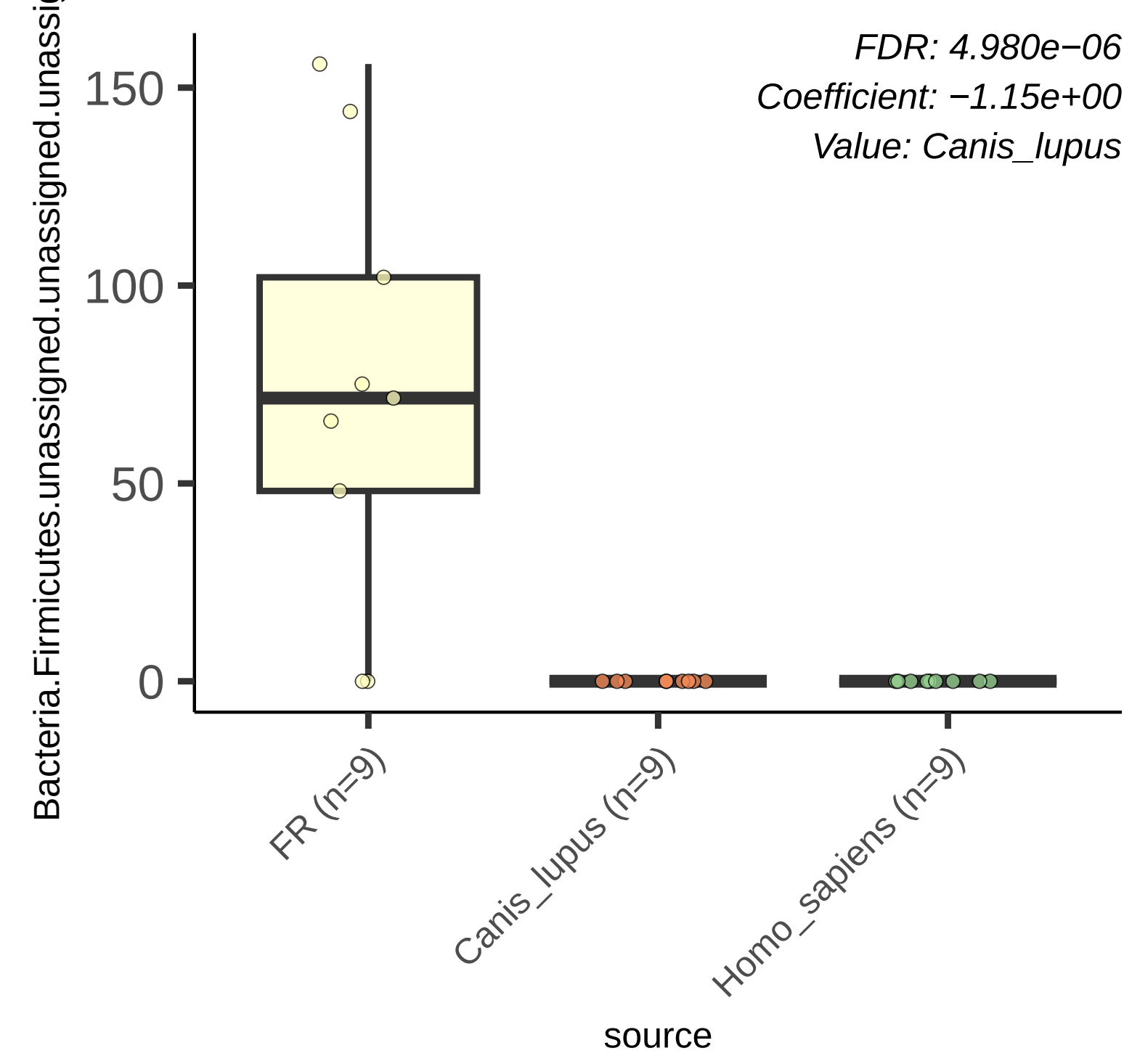
Canis_lupus (n=9)
Homo_sapiens (n=9)
source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Coefficient: −1.15e+00
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Bacteria.Firmicutes.unassigned.unassigned.unassig
150
100
50
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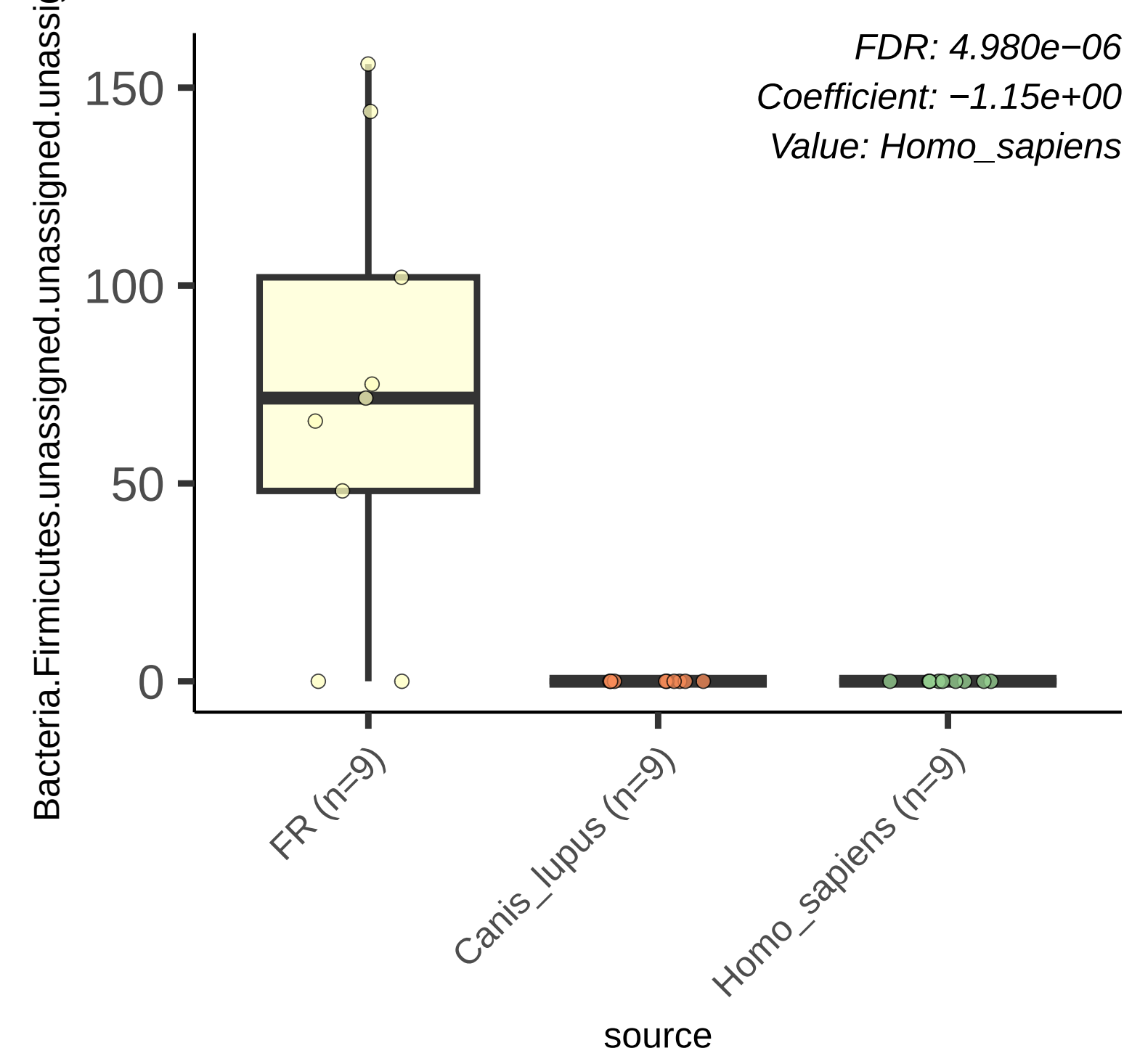
FR (n=9)
Canis_lupus (n=9)
Homo_sapiens (n=9)
source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Coefficient: −1.15e+00
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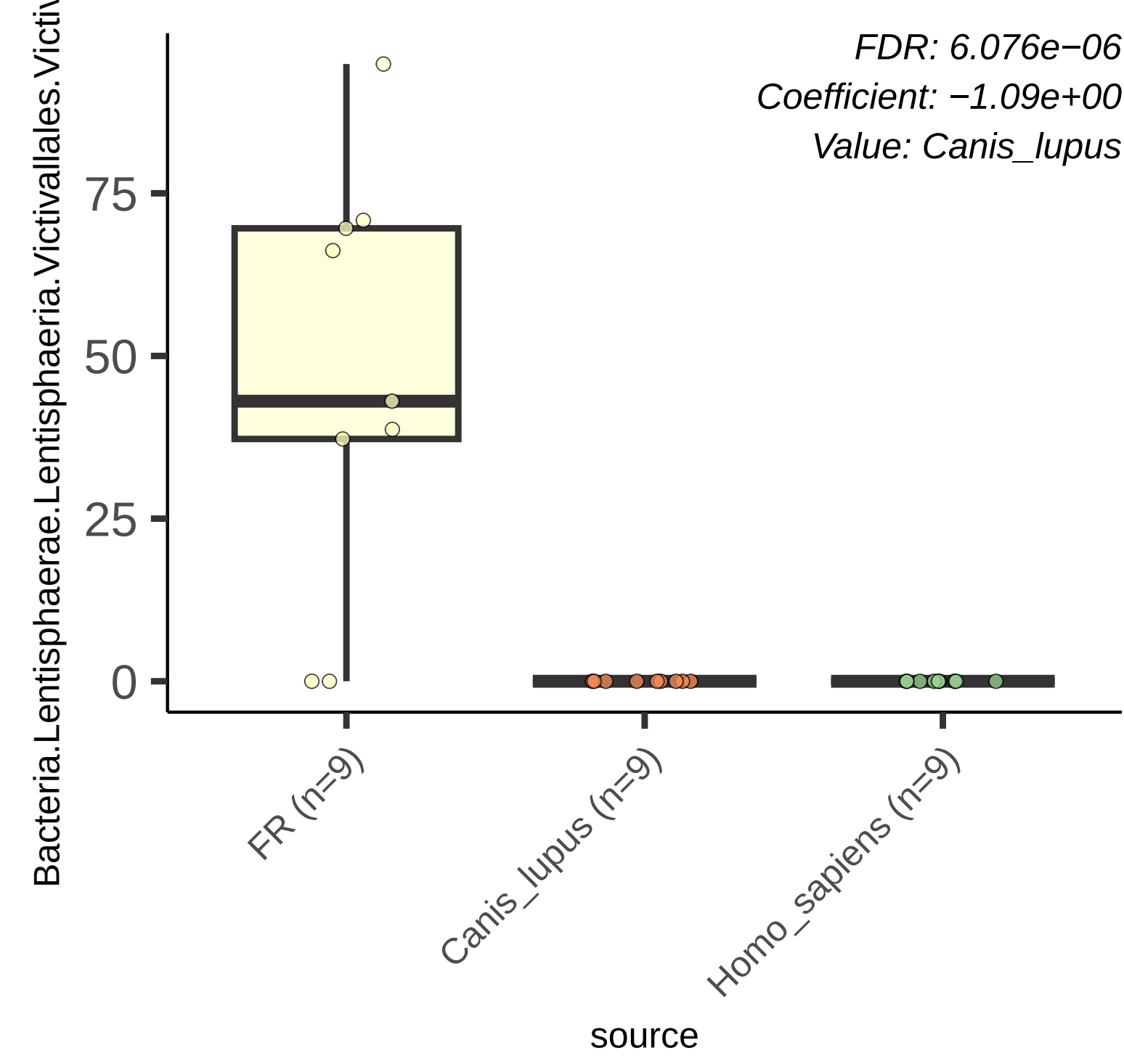
Bacteria.Firmicutes.unassigned.unassigned.unassig
150
100
50
0
FR (n=9)
Canis_lupus (n=9)
Homo_sapiens (n=9)
source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Coefficient: −1.09e+00
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Bacteria.Lentisphaerae.Lentisphaeria.Victivallales.Victi
75
50
25
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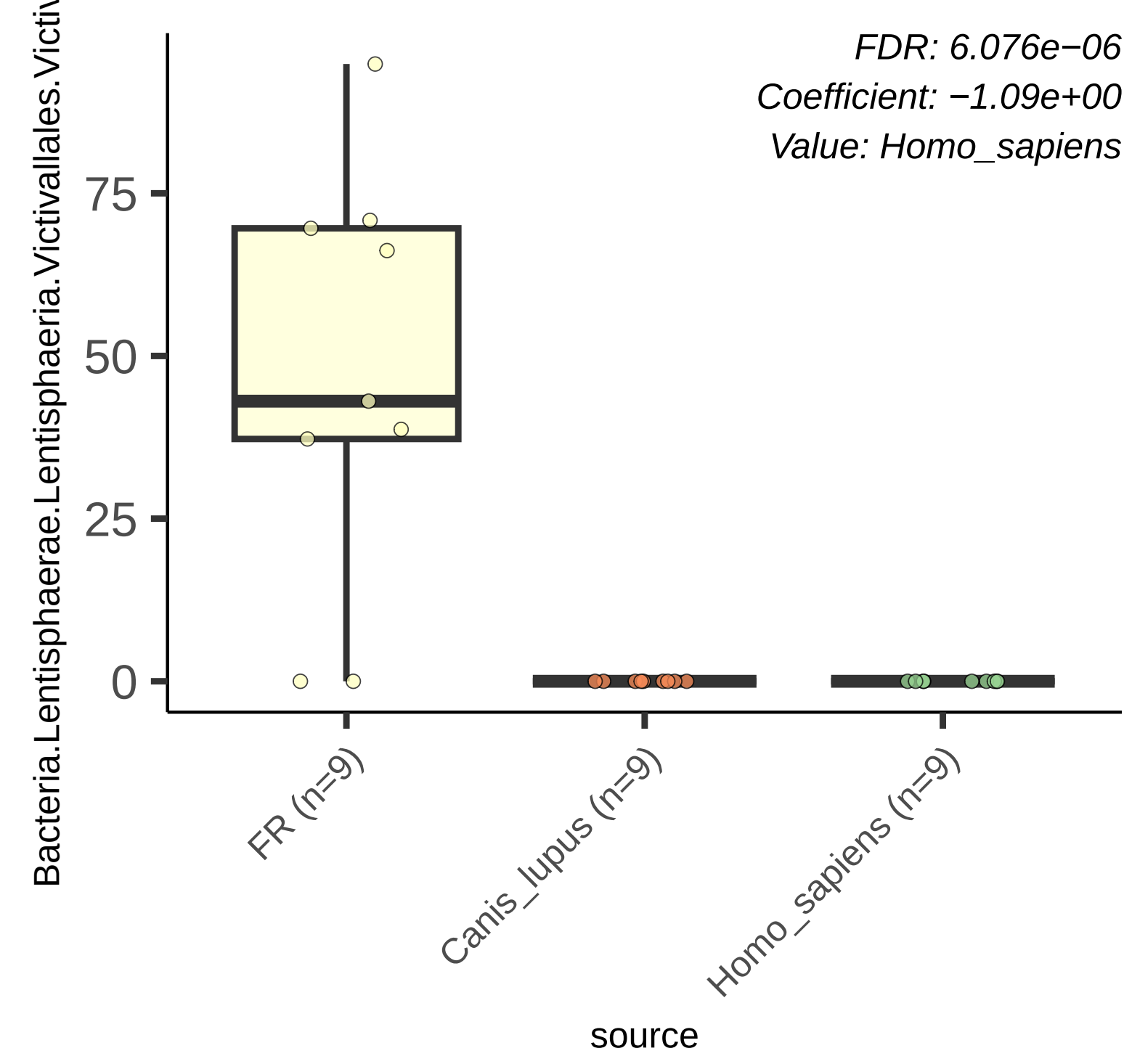
FR (n=9)
Canis_lupus (n=9)
Homo_sapiens (n=9)
source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Coefficient: −1.09e+00
Value: Homo_sapiens
Bacteria.Lentisphaerae.Lentisphaeria.Victivallales.Victiv
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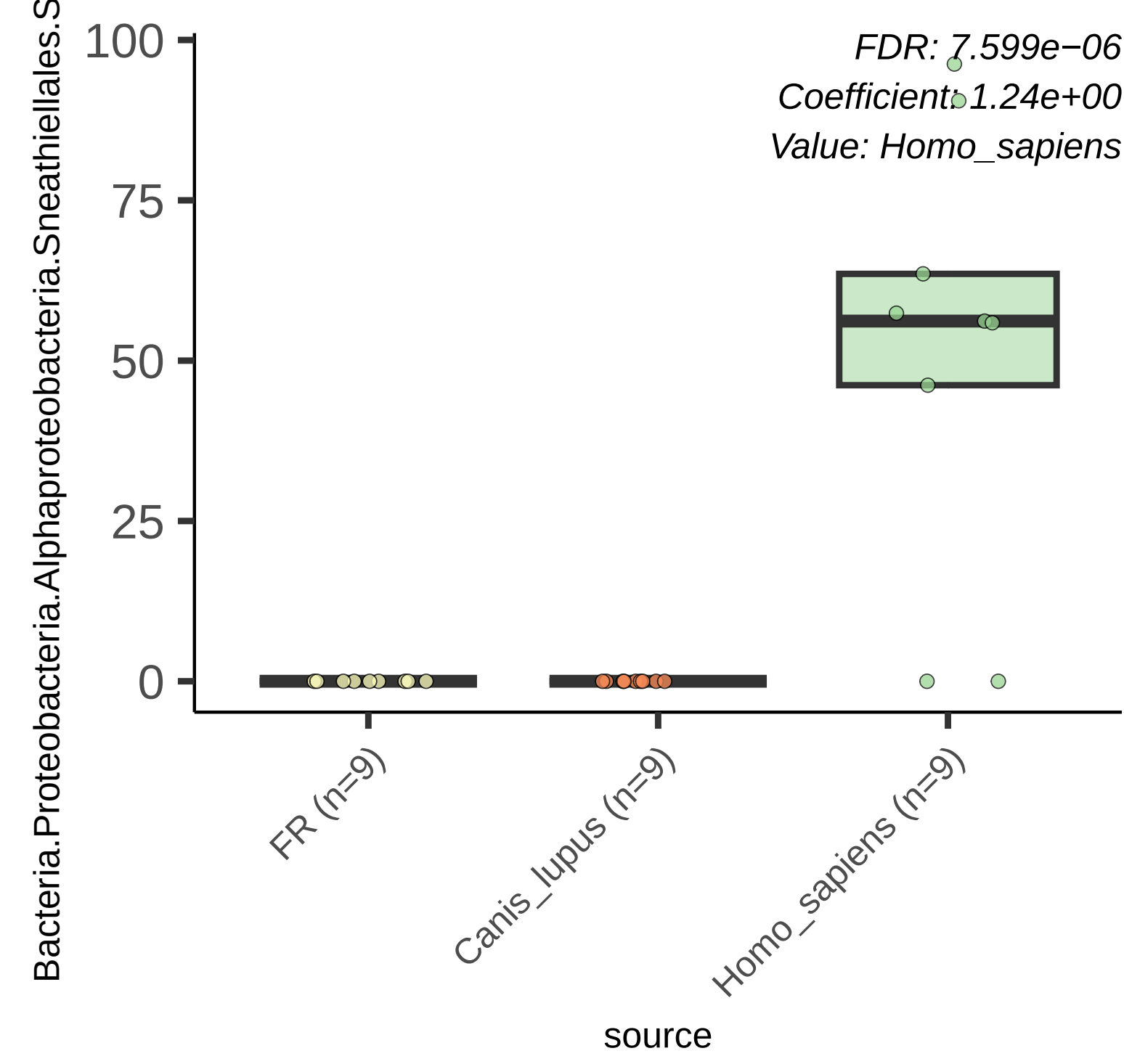
50
25
0
FR (n=9)
Canis_lupus (n=9)
Homo_sapiens (n=9)
source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Coefficient: 1.24e+00
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75
50
25
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Bacteria.Proteobacteria.Alphaproteobacteria.Sneathiellales.S
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Canis_lupus (n=9)
Homo_sapiens (n=9)
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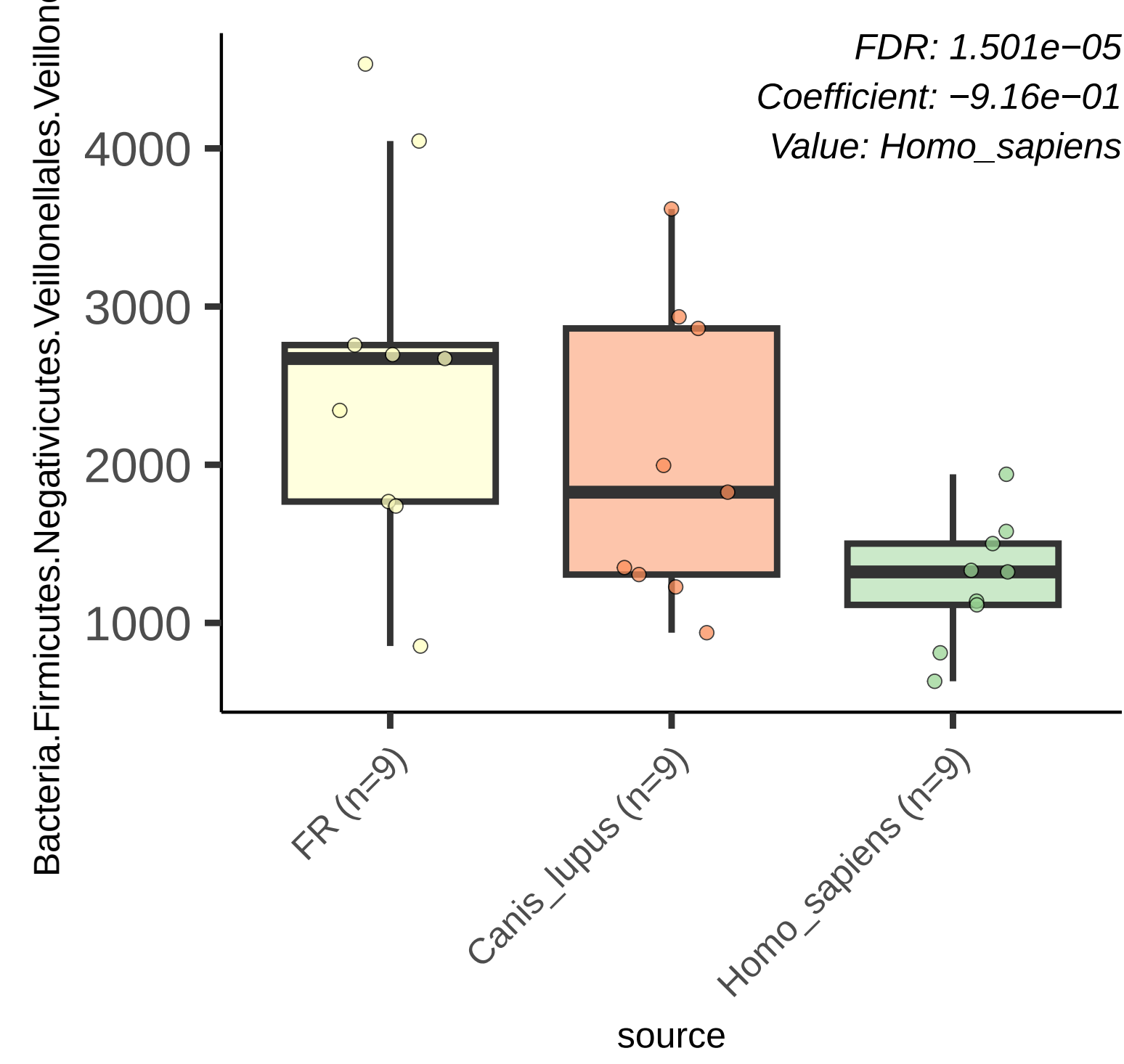
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Coefficient: −9.16e−01
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Bacteria.Firmicutes.Negativicutes.Veillonellales.Veillonellales
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3000
2000
1000
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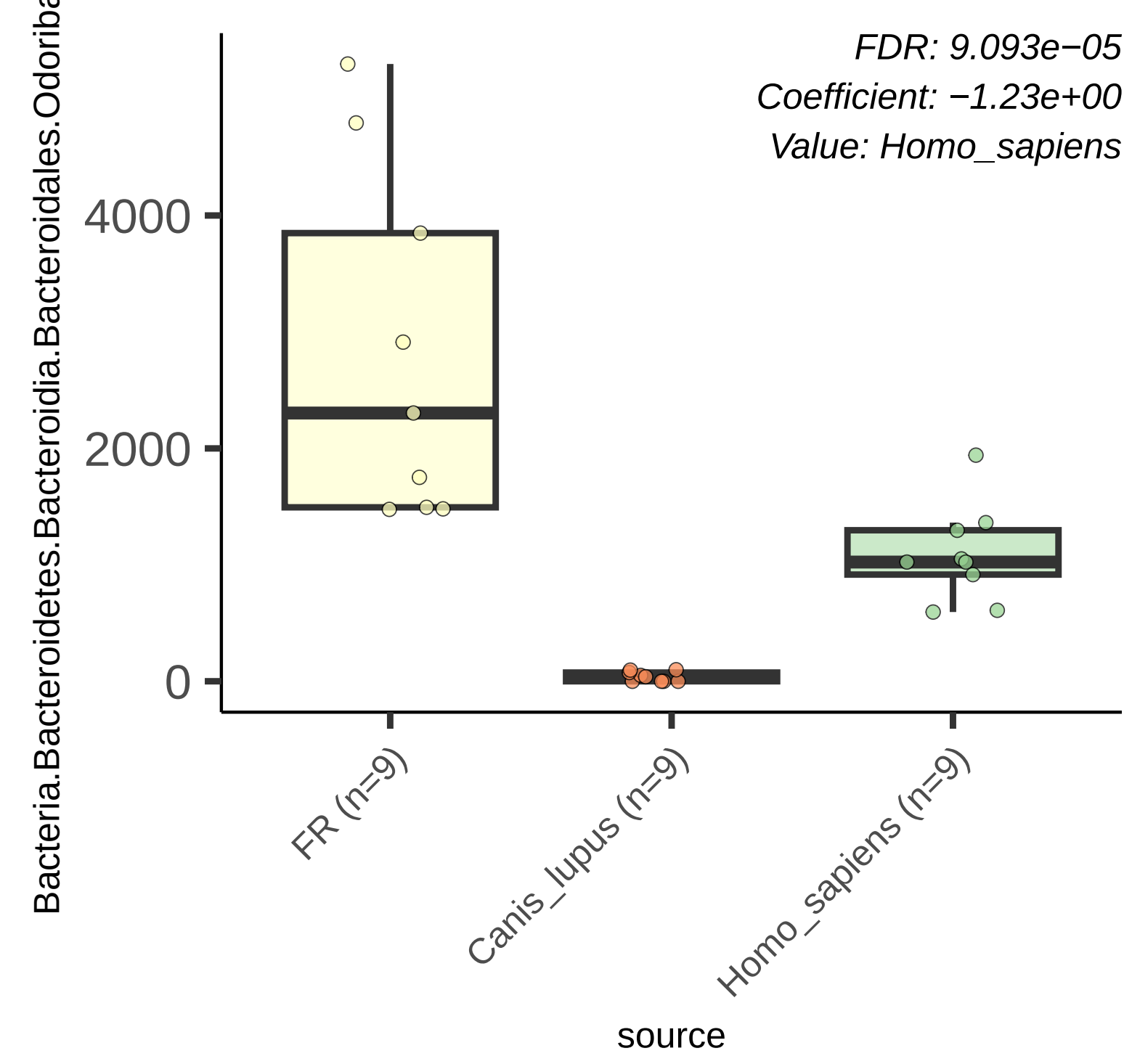
Canis_lupus (n=9)
Homo_sapiens (n=9)
source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Coefficient: −1.23e+00
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Bacteria.Bacteroidetes.Bacteroidia.Bacteroidales.Odoriba
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2000
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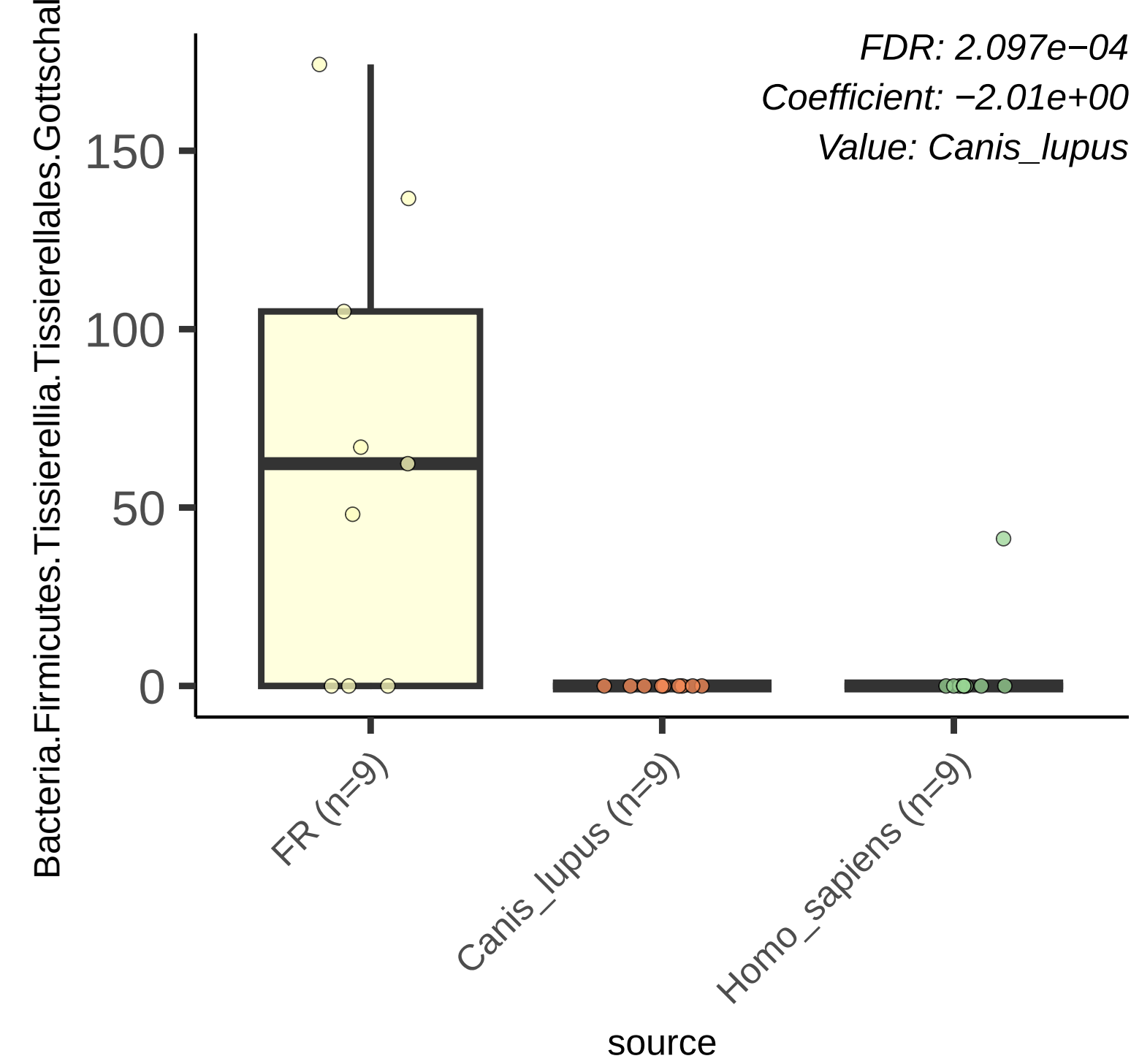
Canis_lupus (n=9)
Homo_sapiens (n=9)
source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Coefficient: −2.01e+00
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Bacteria.Firmicutes.Tissierellia.Tissierellales.Gottschal
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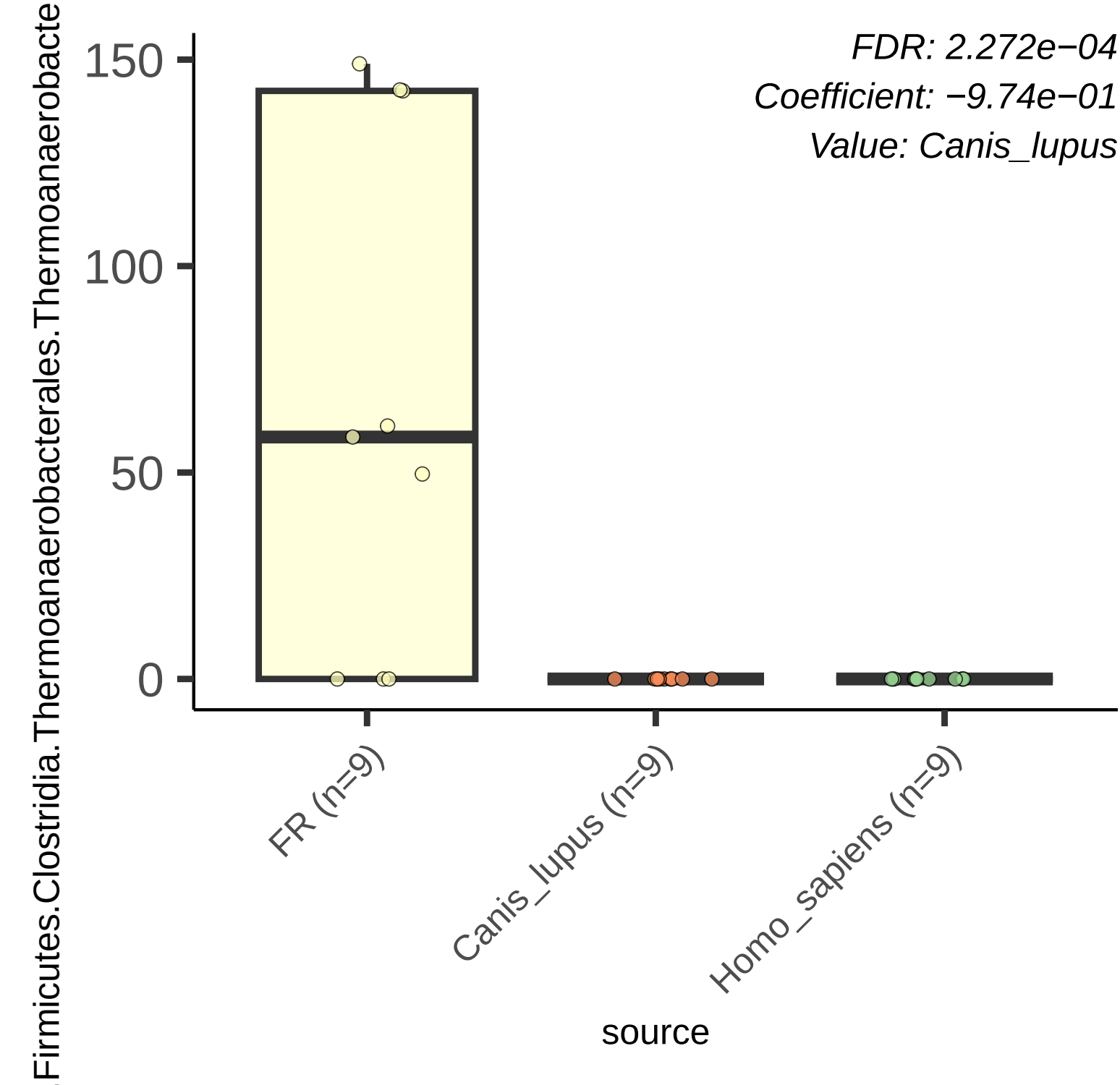
Canis_lupus (n=9)
Homo_sapiens (n=9)
source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Coefficient: −9.74e−01
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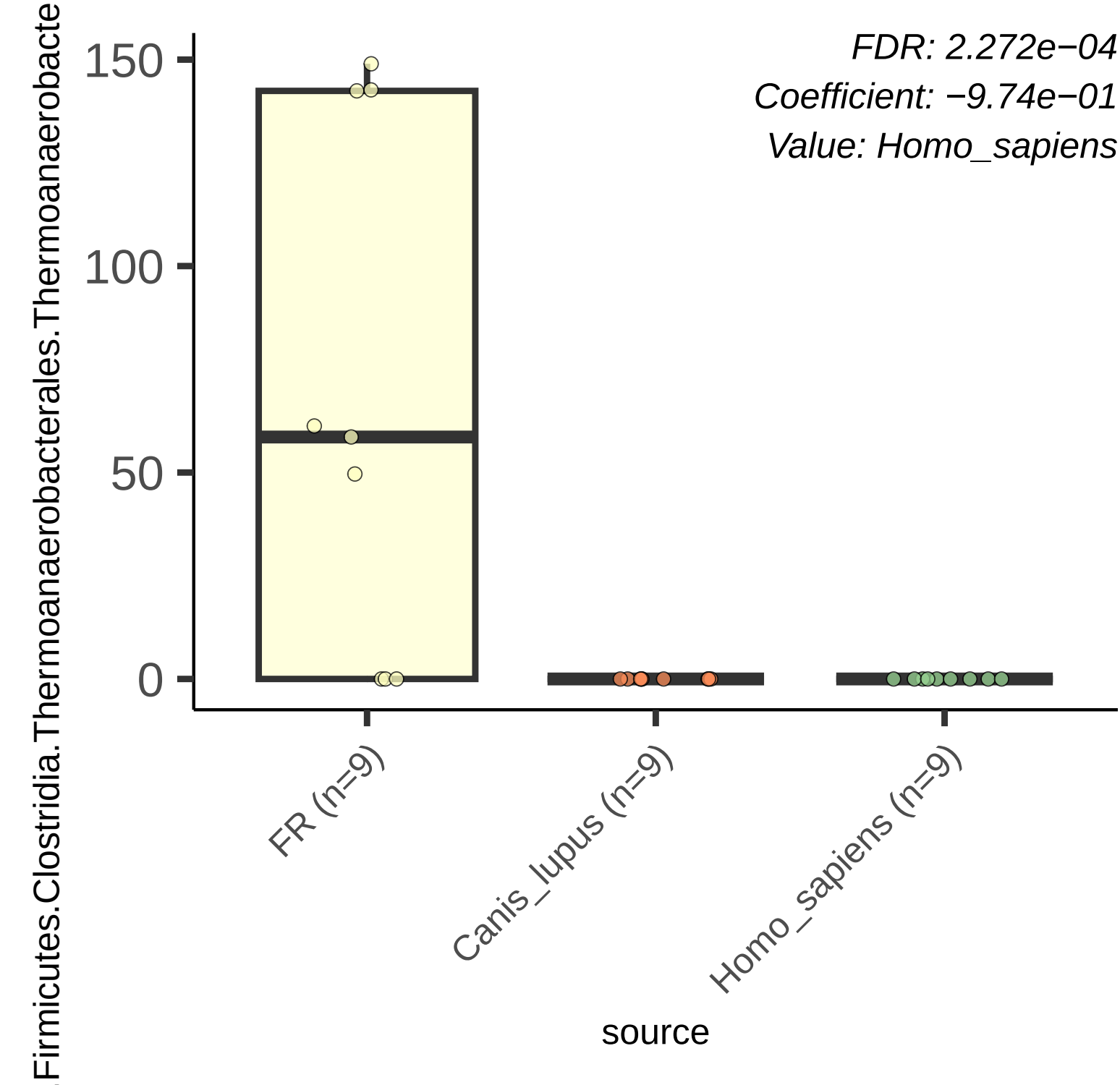
Canis_lupus (n=9)
Homo_sapiens (n=9)
source
Firmicutes.Clostridia.Thermoanaerobacterales.Thermoanaerobacte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Coefficient: −9.74e−01
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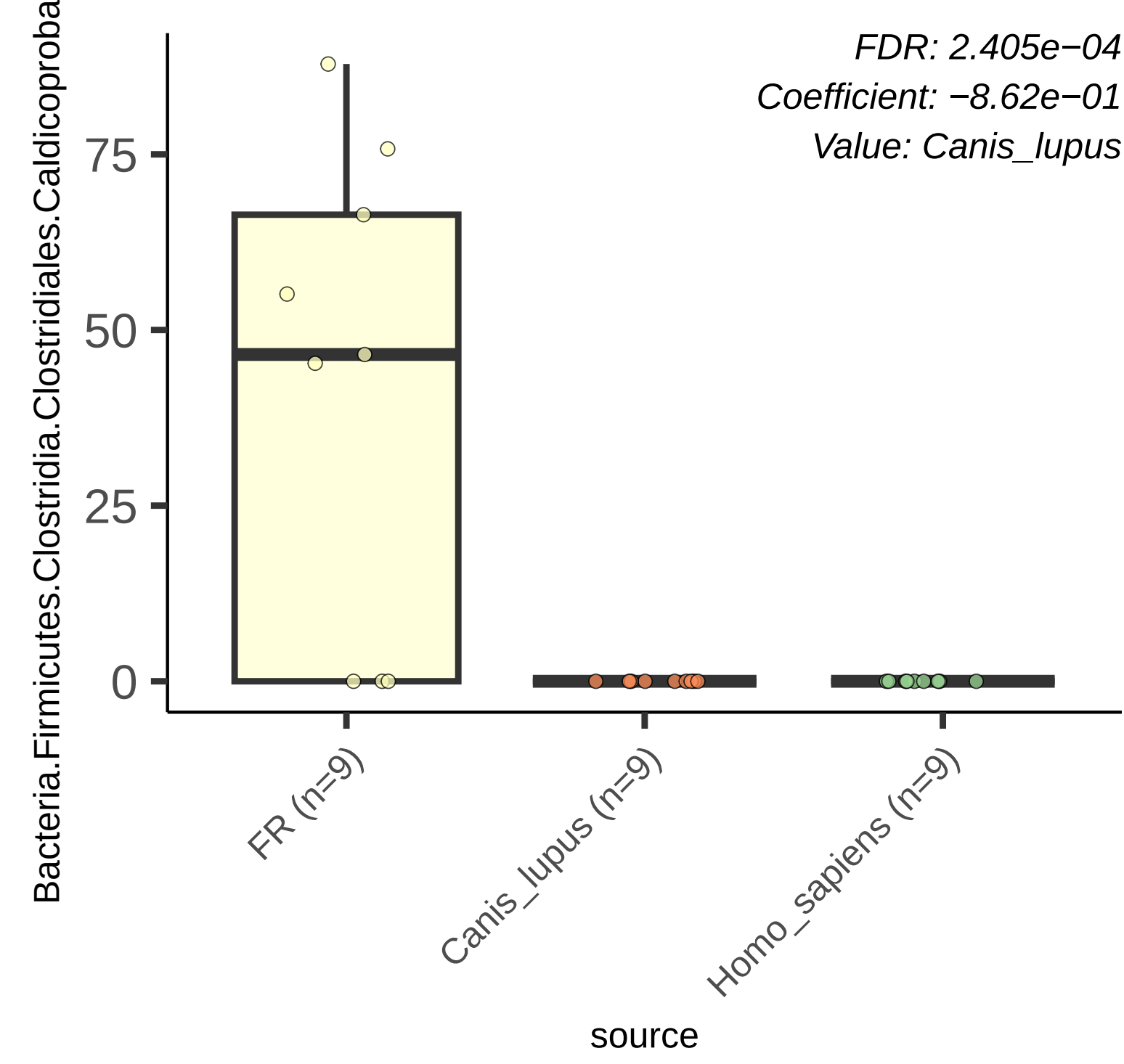
0
FR (n=9)
Canis_lupus (n=9)
Homo_sapiens (n=9)
source
Firmicutes.Clostridia.Thermoanaerobacterales.Thermoanaerobacte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Coefficient: −8.62e−01
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Bacteria.Firmicutes.Clostridia.Clostridiales.Caldicoprobacteraceae.Caldicoprobacter
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25
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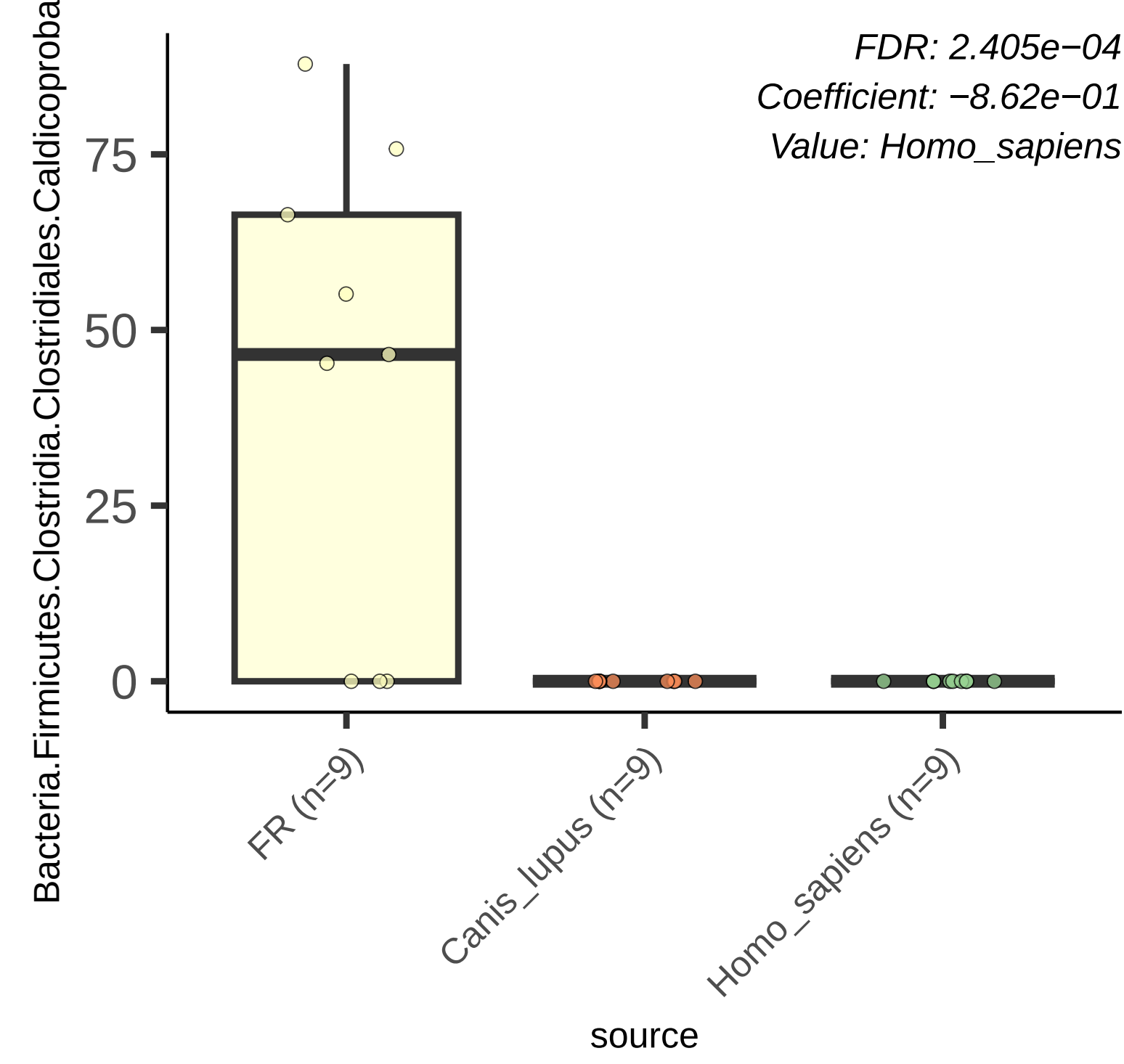
Canis_lupus (n=9)
Homo_sapiens (n=9)
source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Coefficient: −8.62e−01
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Bacteria.Firmicutes.Clostridia.Clostridiales.Caldicoprobacteraceae.Caldicoprobacter
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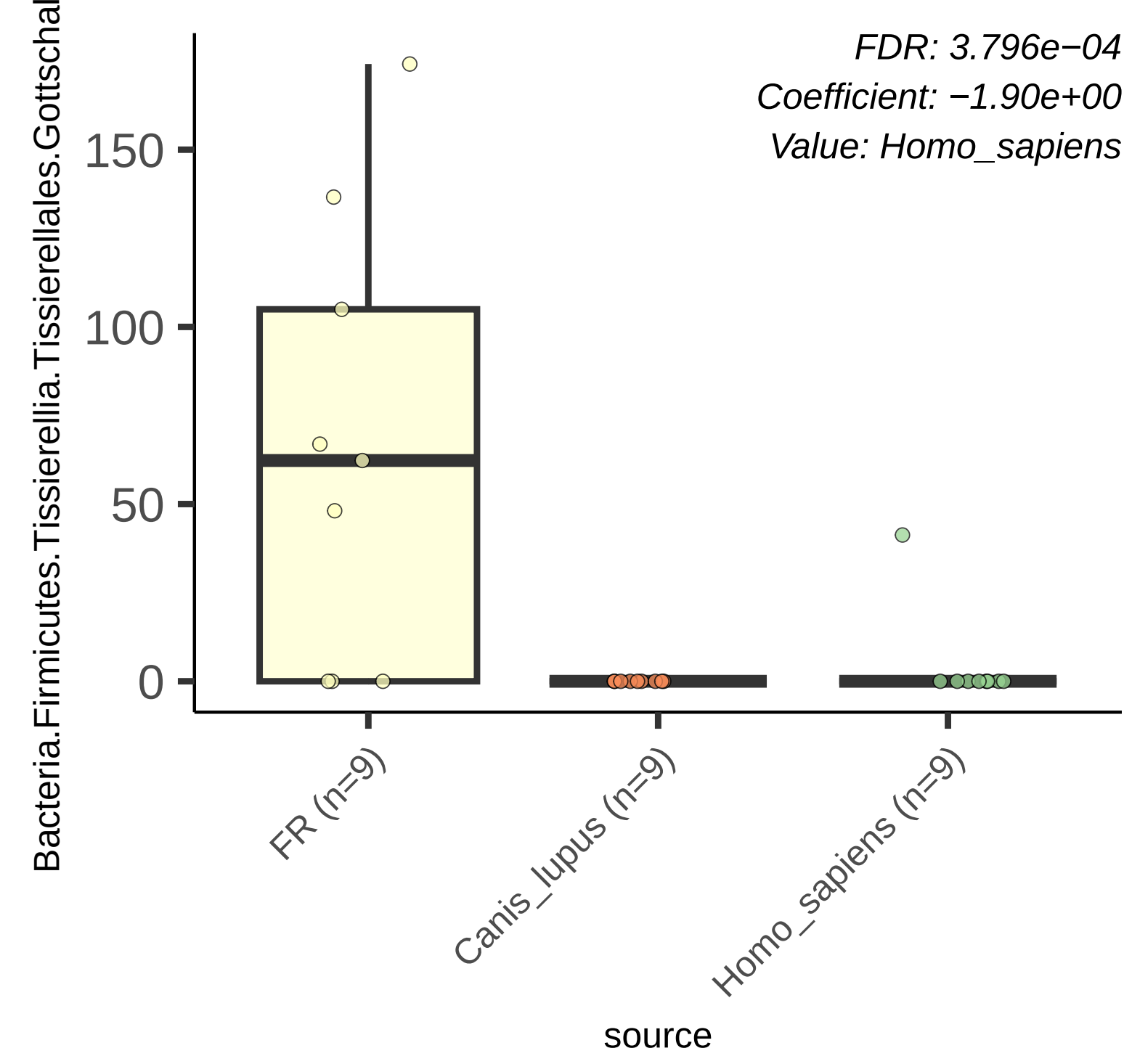
50
25
0
FR (n=9)
Canis_lupus (n=9)
Homo_sapiens (n=9)
source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Coefficient: −1.90e+00
Value: Homo_sapiens
Bacteria.Firmicutes.Tissierellia.Tissierellales.Gottschal
150
100
50
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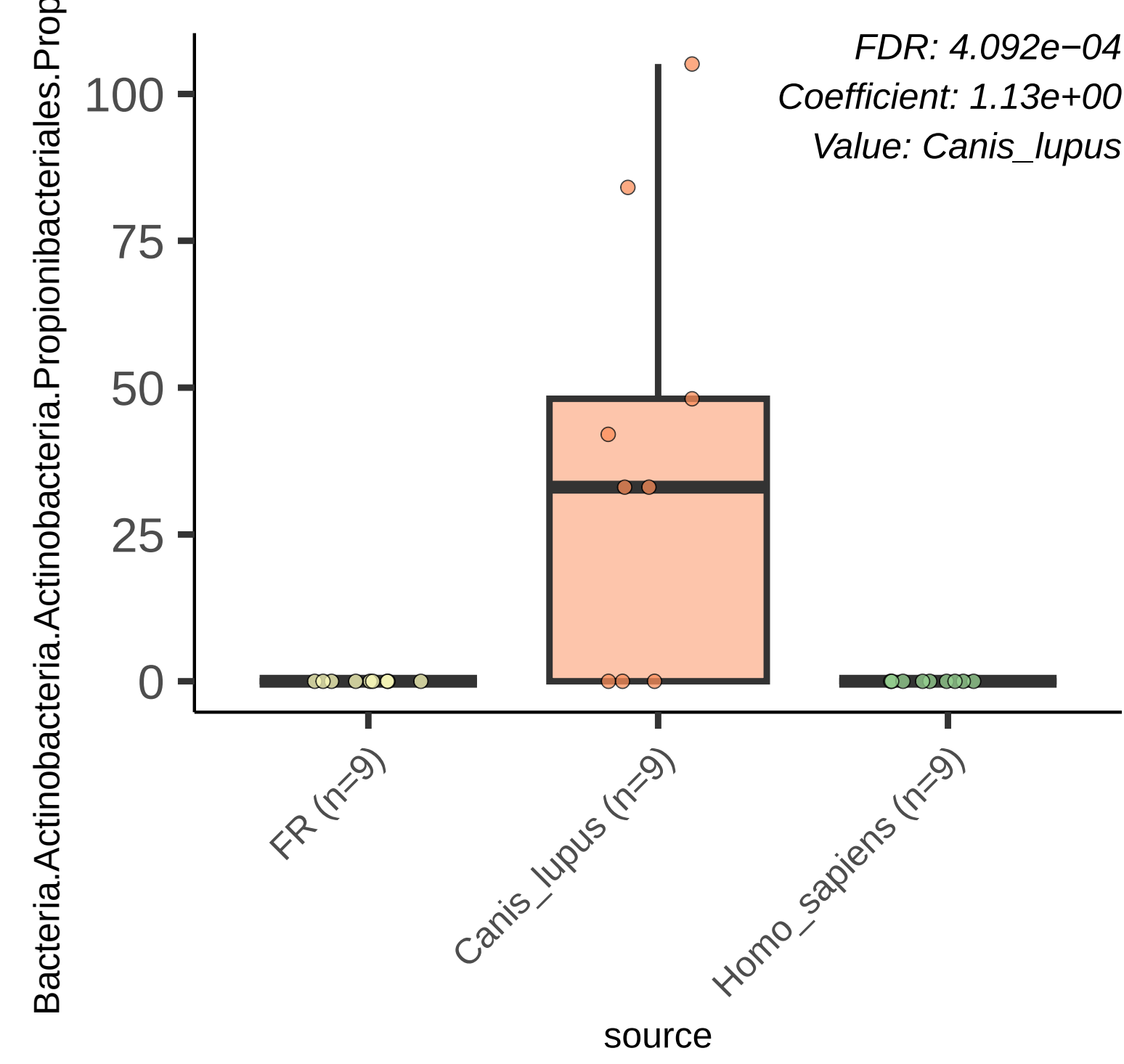
FR (n=9)
Canis_lupus (n=9)
Homo_sapiens (n=9)
source


FDR: 4.092e−04
Coefficient: 1.13e+00
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Bacteria.Actinobacteria.Actinobacteria.Propionibacteriales.Prop
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50
25
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Canis_lupus (n=9)
Homo_sapiens (n=9)
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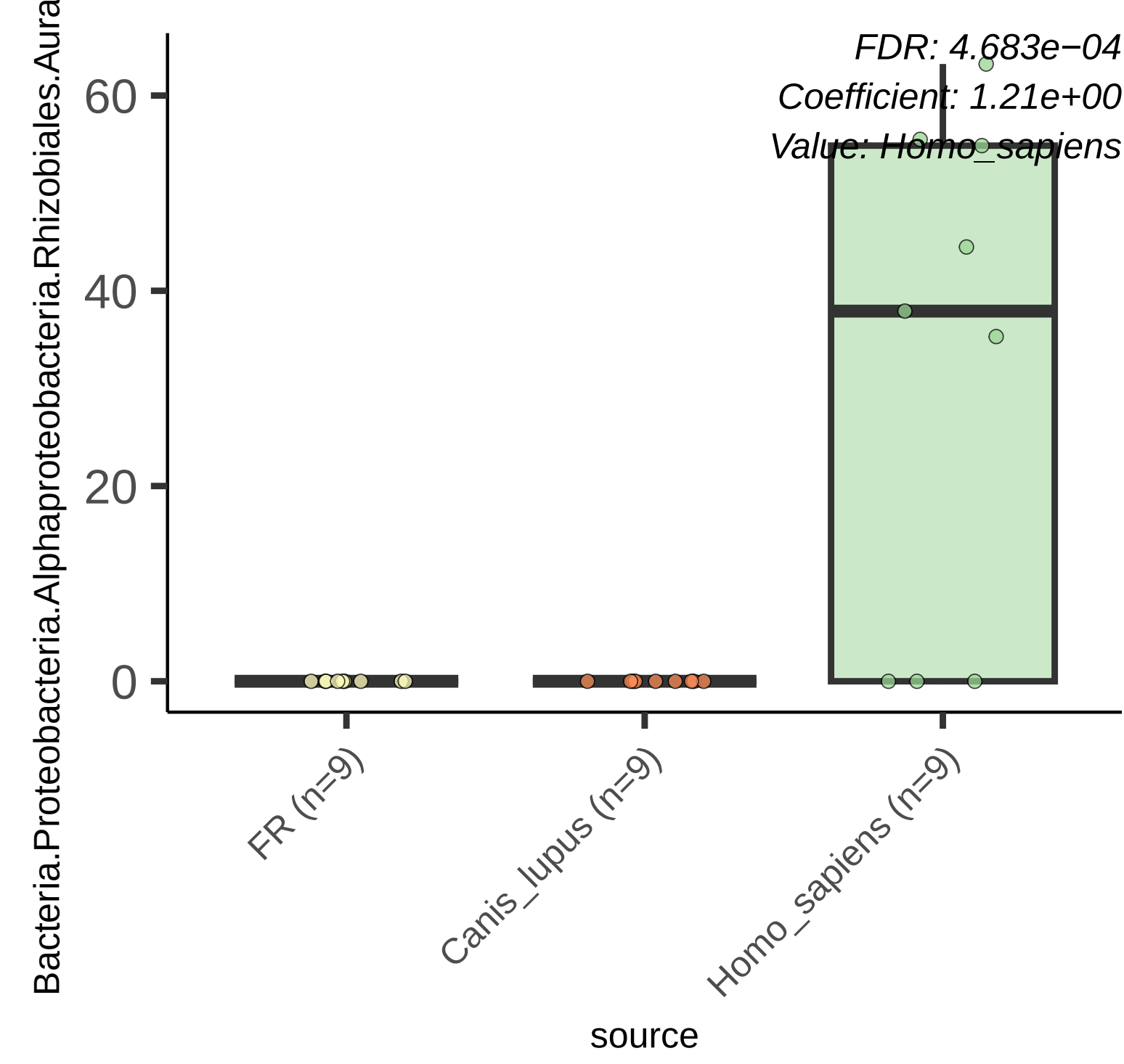
FDR: 4.683e−04
Coefficient: 1.21e+00
Value: Homo_sapiens
Bacteria.Proteobacteria.Alphaproteobacteria.Rhizobiales.Aura
60
40
20
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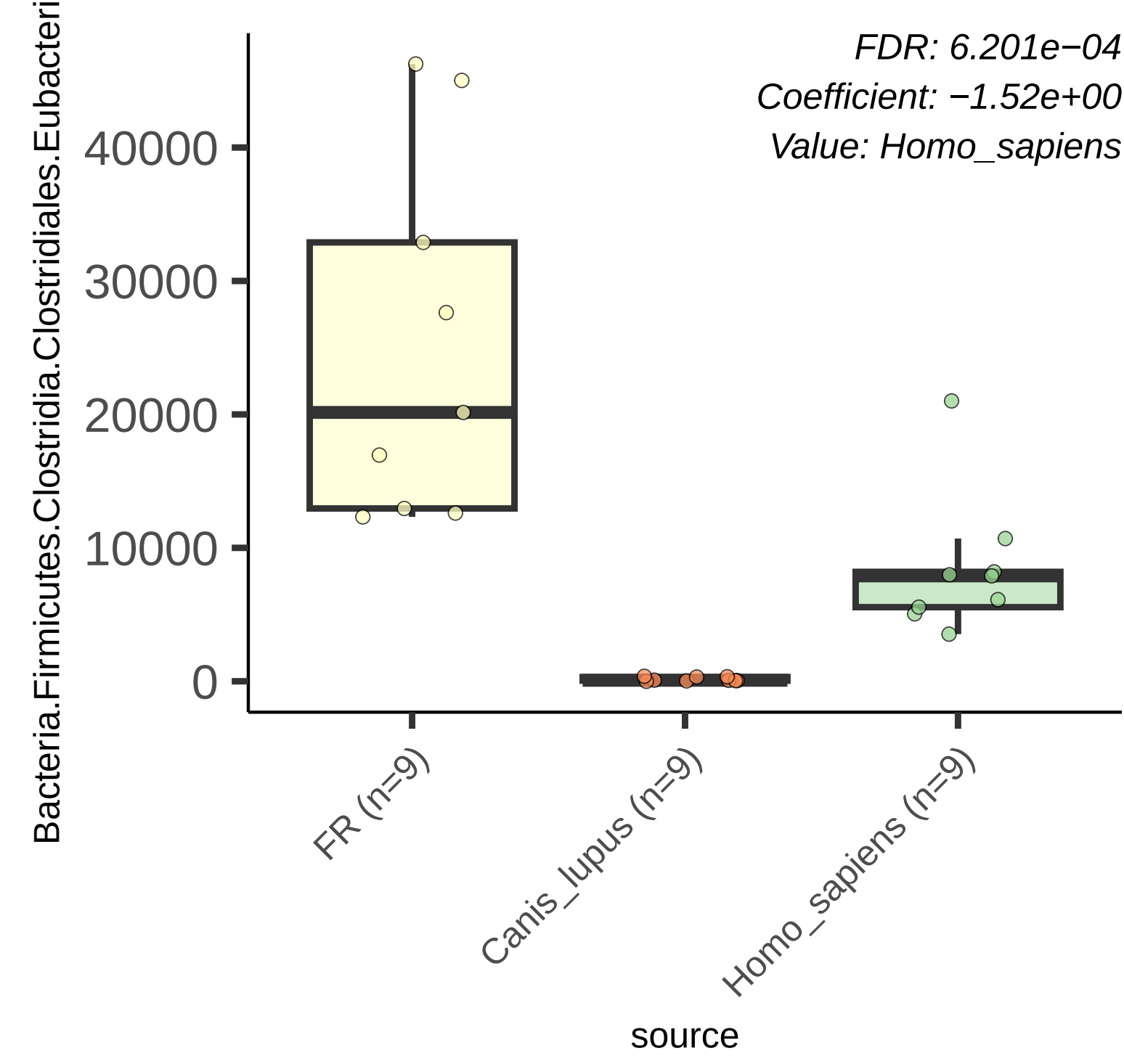
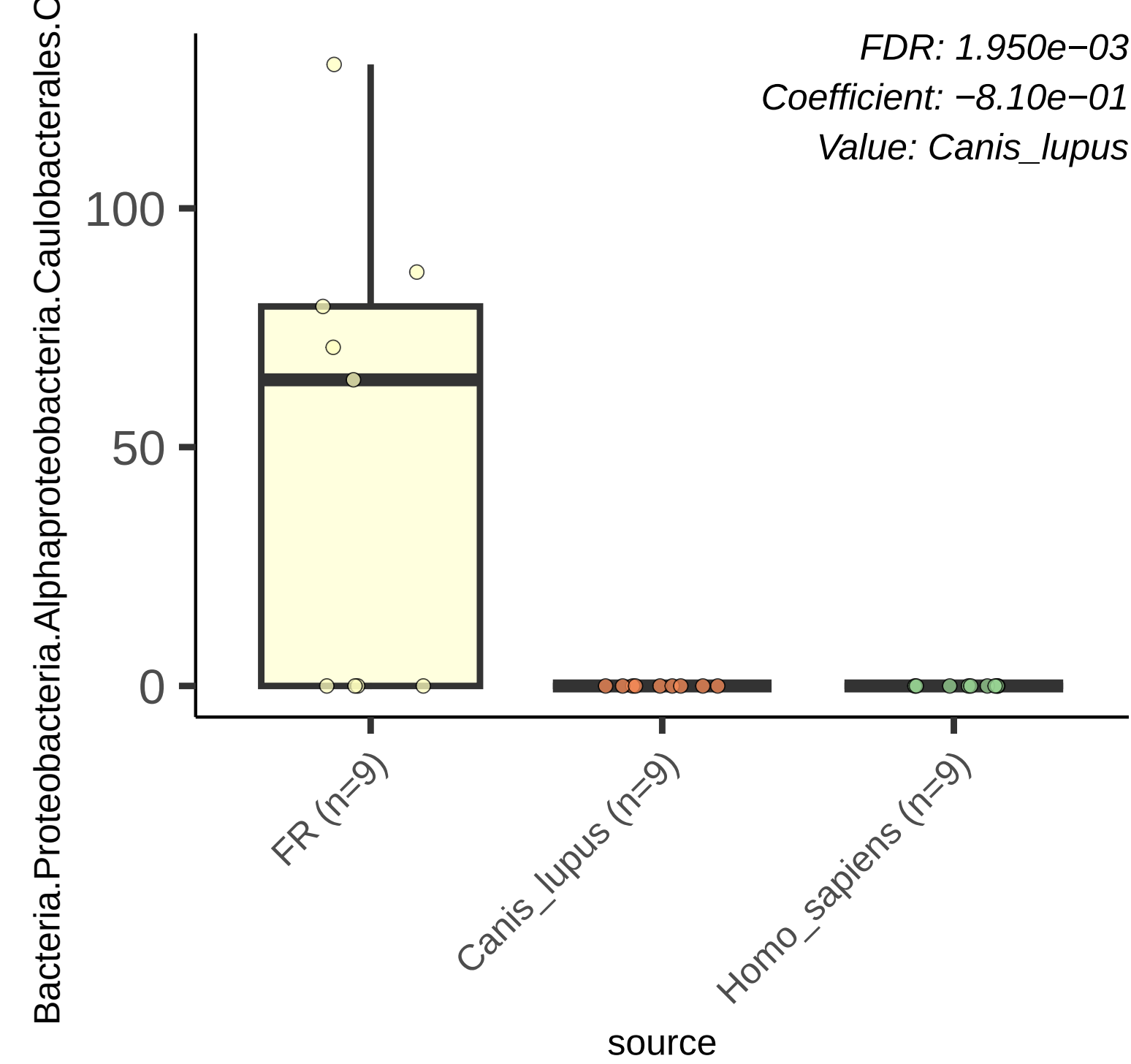
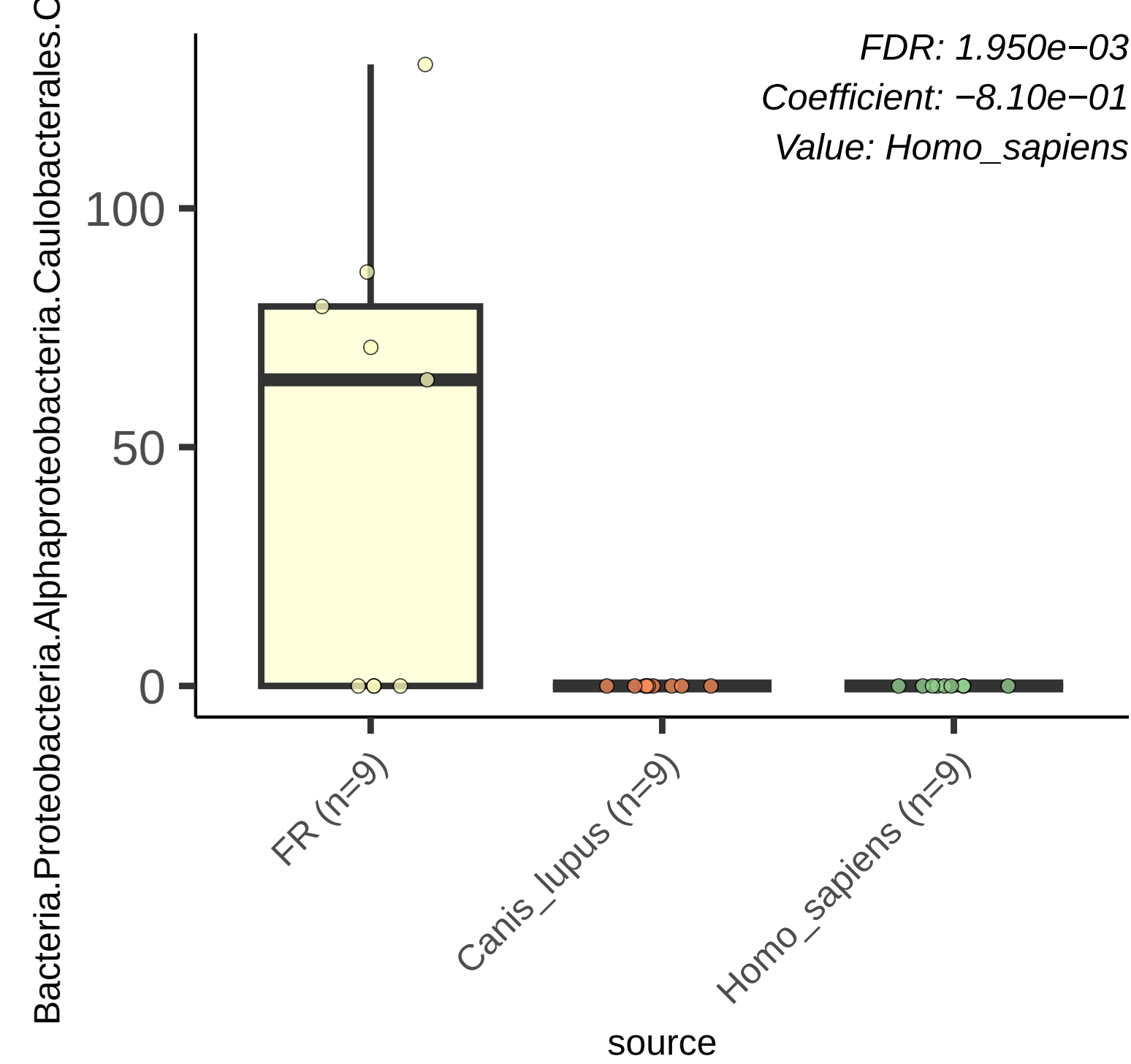
FR (n=9)
Canis_lupus (n=9)
Homo_sapiens (n=9)
source


FDR: 6.201e−04
Coefficient: −1.52e+00
Value: Homo_sapiens
Bacteria.Firmicutes.Clostridia.Clostridiales.Eubacteri
40000
30000
20000
10000
0
FR (n=9)
Canis_lupus (n=9)
Homo_sapiens (n=9)
source


FDR: 1.950e−03
Coefficient: −8.10e−01
Value: Canis_lupus
Bacteria.Proteobacteria.Alphaproteobacteria.Caulobacterales.C
100
50
0
FR (n=9)
Canis_lupus (n=9)
Homo_sapiens (n=9)
source


FDR: 1.950e−03
Coefficient: −8.10e−01
Value: Homo_sapiens
Bacteria.Proteobacteria.Alphaproteobacteria.Caulobacterales.C
100
50
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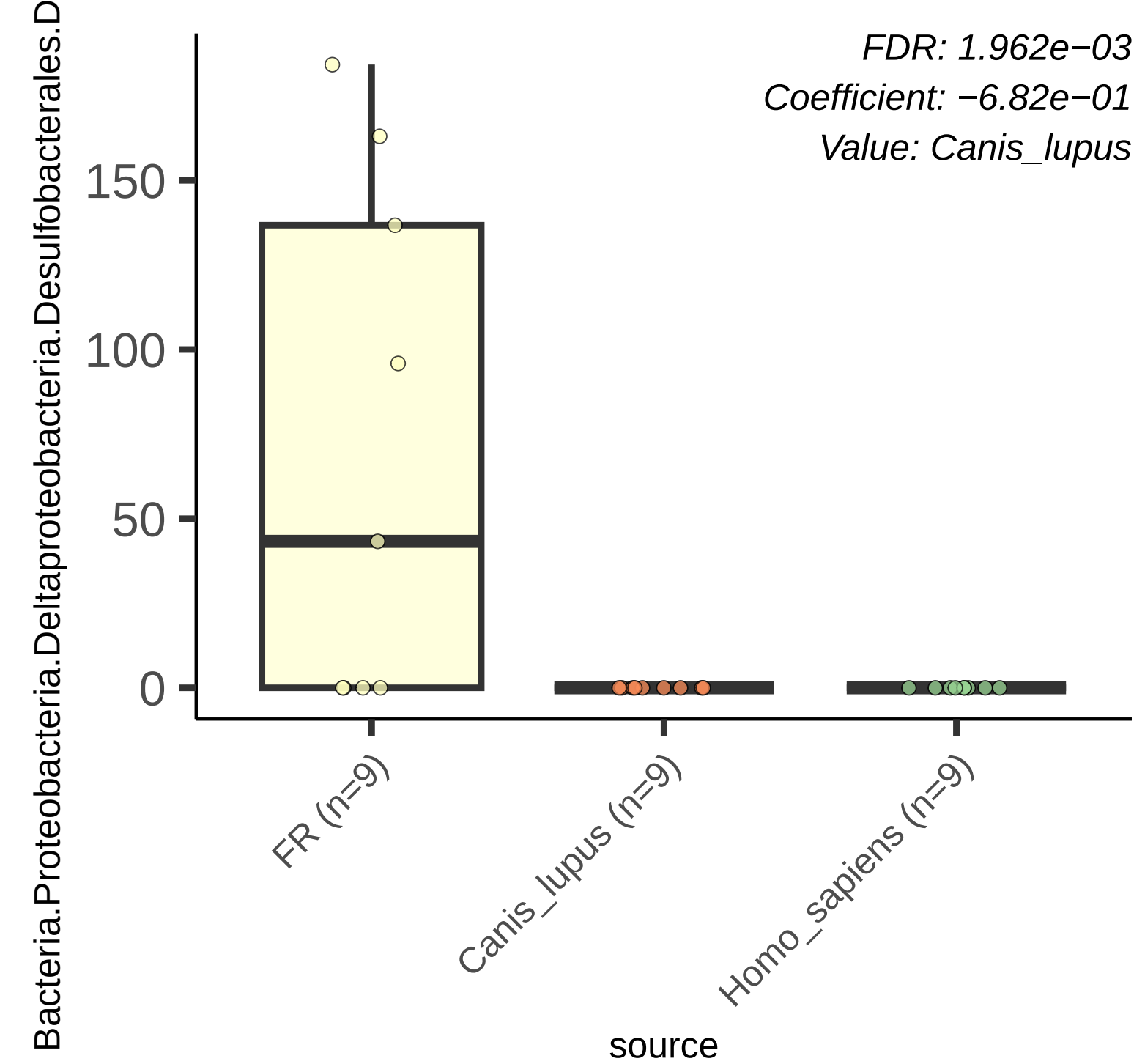
FR (n=9)
Canis_lupus (n=9)
Homo_sapiens (n=9)
source


FDR: 1.962e−03
Coefficient: −6.82e−01
Value: Canis_lupus
Bacteria.Proteobacteria.Deltaproteobacteria.Desulfobacterales.D
150
100
50
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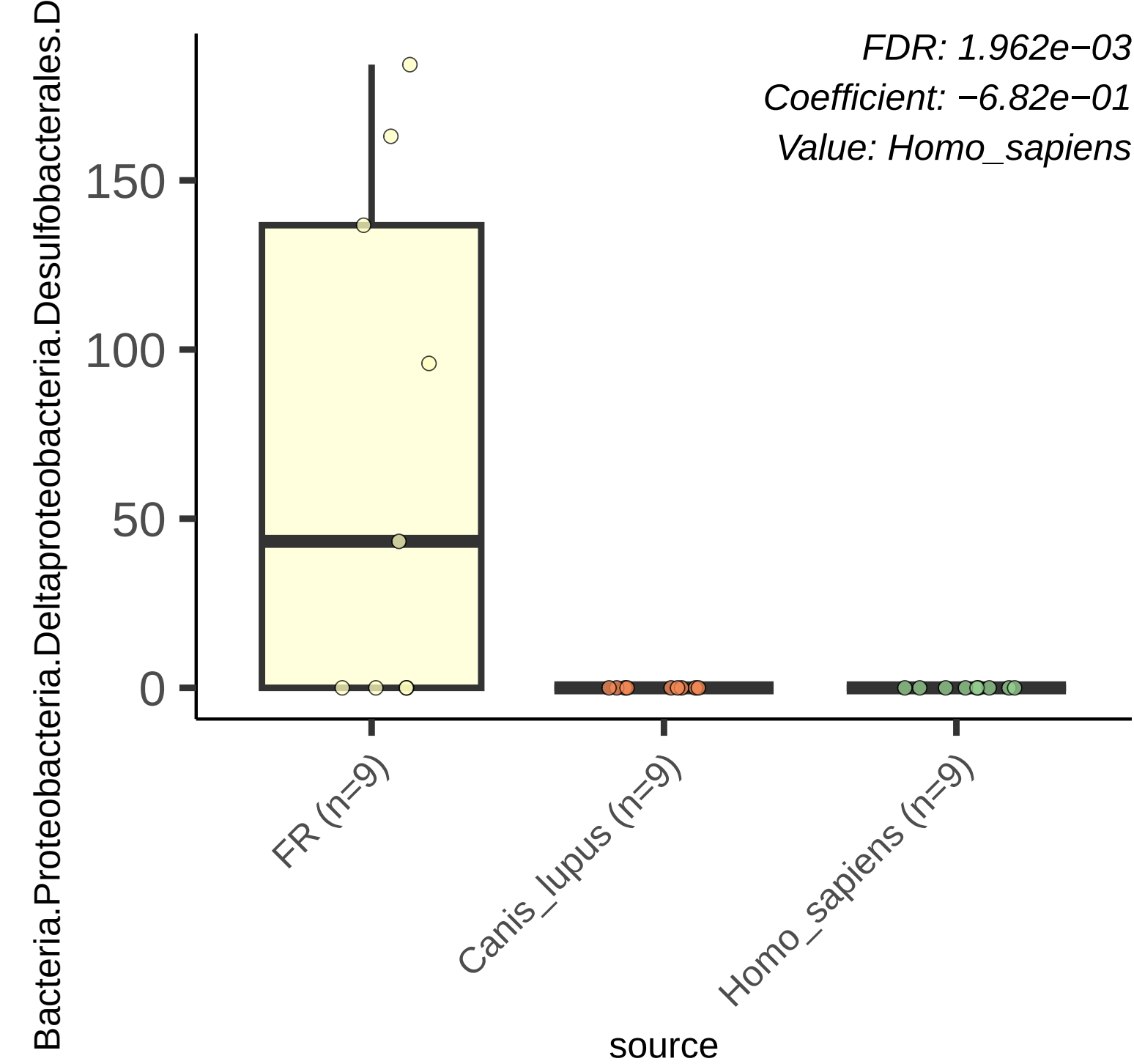
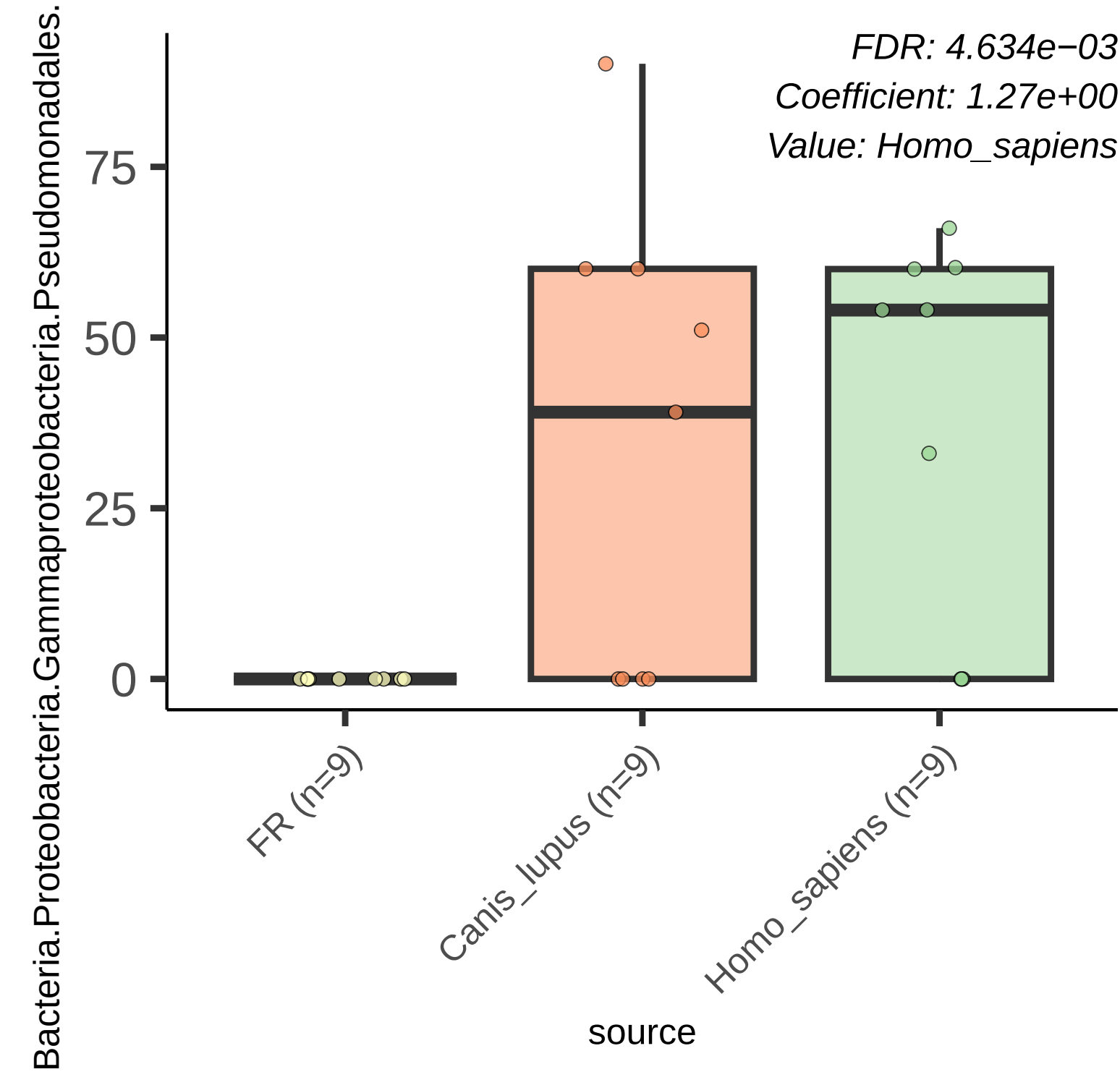
FR (n=9)
Canis_lupus (n=9)
Homo_sapiens (n=9)
source


FDR: 1.962e−03
Coefficient: −6.82e−01
Value: Homo_sapiens
Bacteria.Proteobacteria.Deltaproteobacteria.Desulfobacterales.D
150
100
50
0
FR (n=9)
Canis_lupus (n=9)
Homo_sapiens (n=9)
source


FDR: 4.634e−03
Coefficient: 1.27e+00
Value: Homo_sapiens
Bacteria.Proteobacteria.Gammaproteobacteria.Pseudomonadales.
75
50
25
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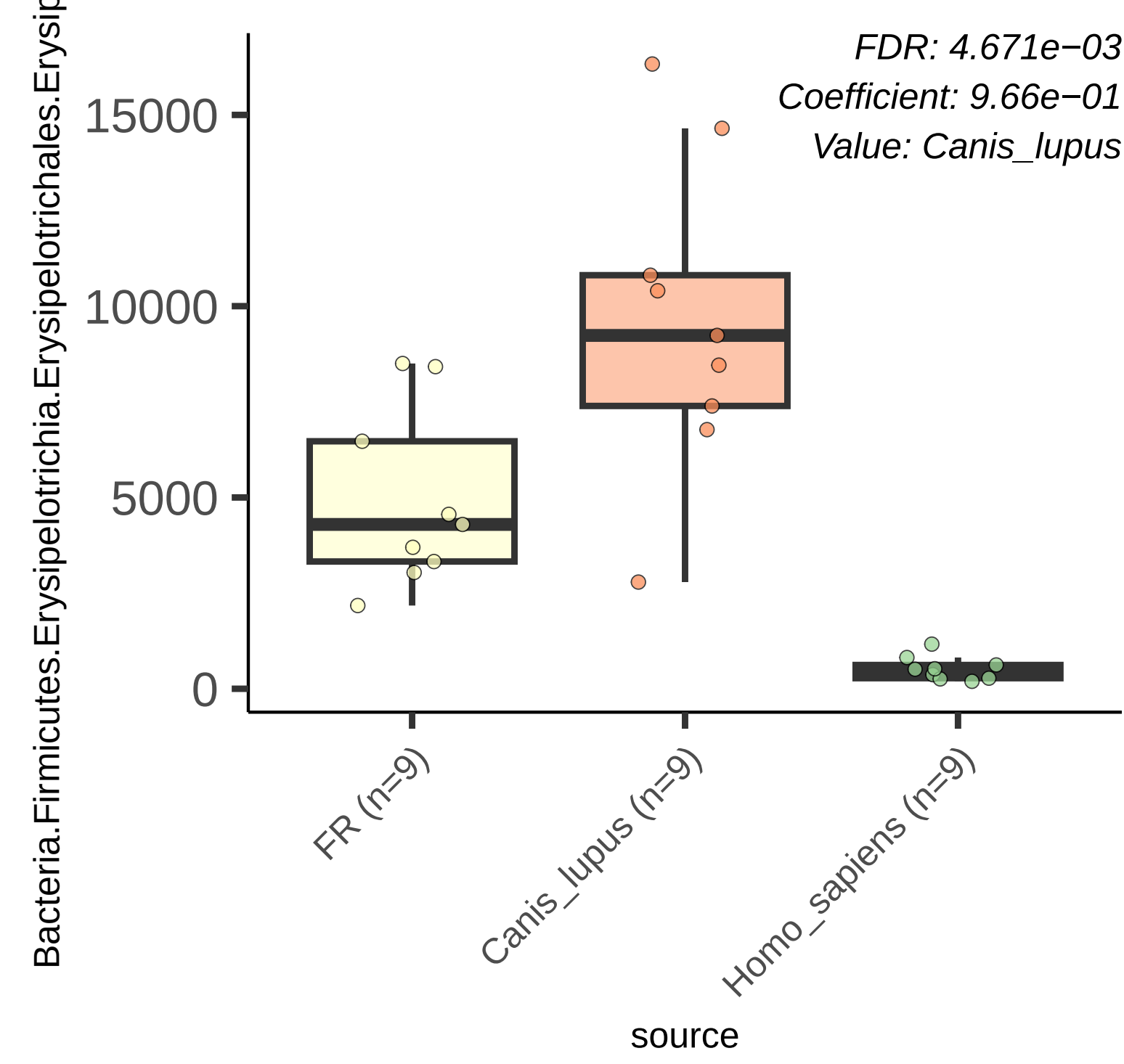
Canis_lupus (n=9)
Homo_sapiens (n=9)
source


FDR: 4.671e−03
Coefficient: 9.66e−01
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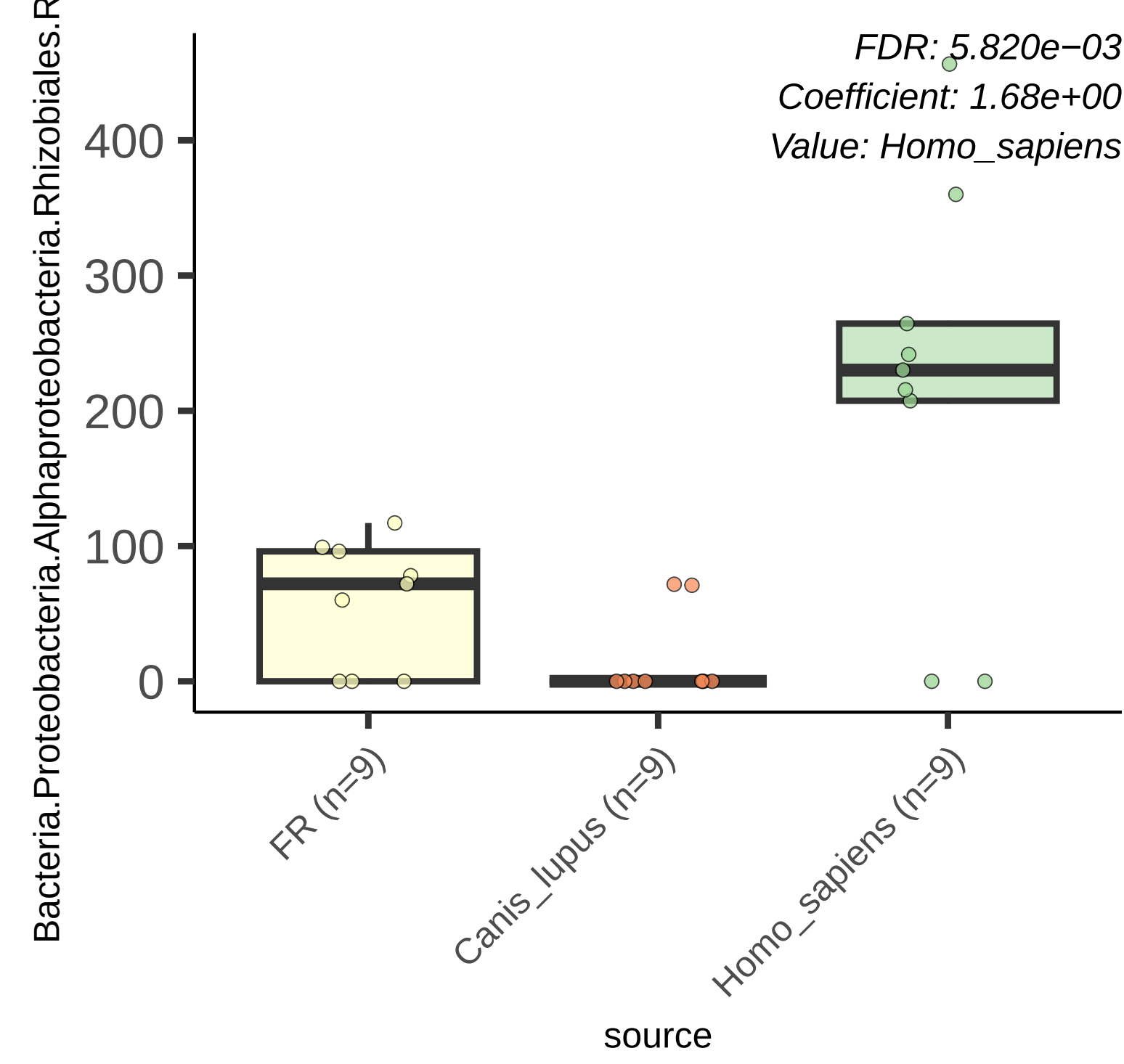
15000
10000
5000
0
Bacteria.Firmicutes.Erysipelotrichia.Erysipelotrichales.Erysip
FR (n=9)
Canis_lupus (n=9)
Homo_sapiens (n=9)
source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Coefficient: 1.68e+00
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Bacteria.Proteobacteria.Alphaproteobacteria.Rhizobiales.R
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300
200
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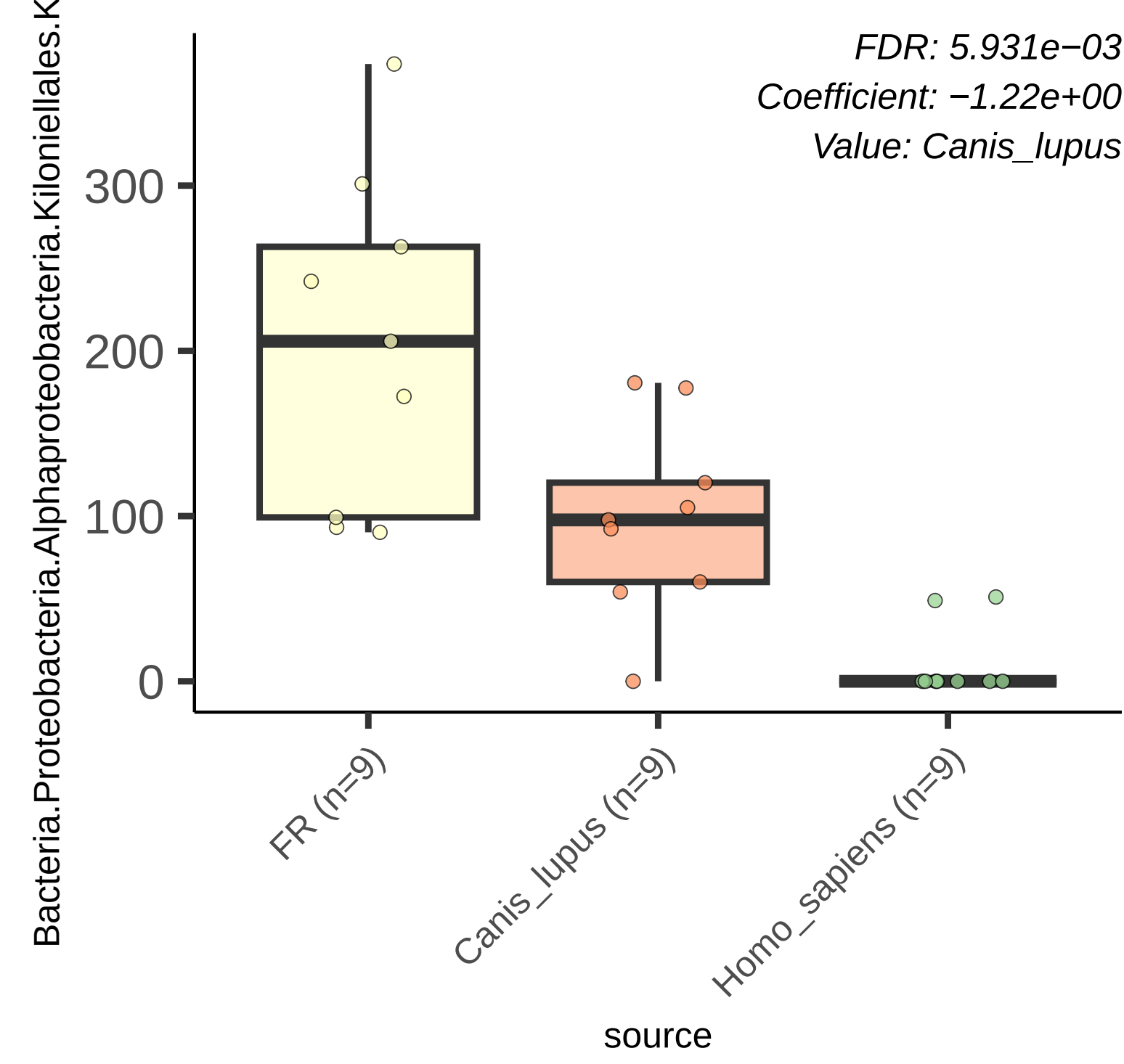
Canis_lupus (n=9)
Homo_sapiens (n=9)
source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Coefficient: −1.22e+00
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Bacteria.Proteobacteria.Alphaproteobacteria.Kiloniellales.K
300
200
100
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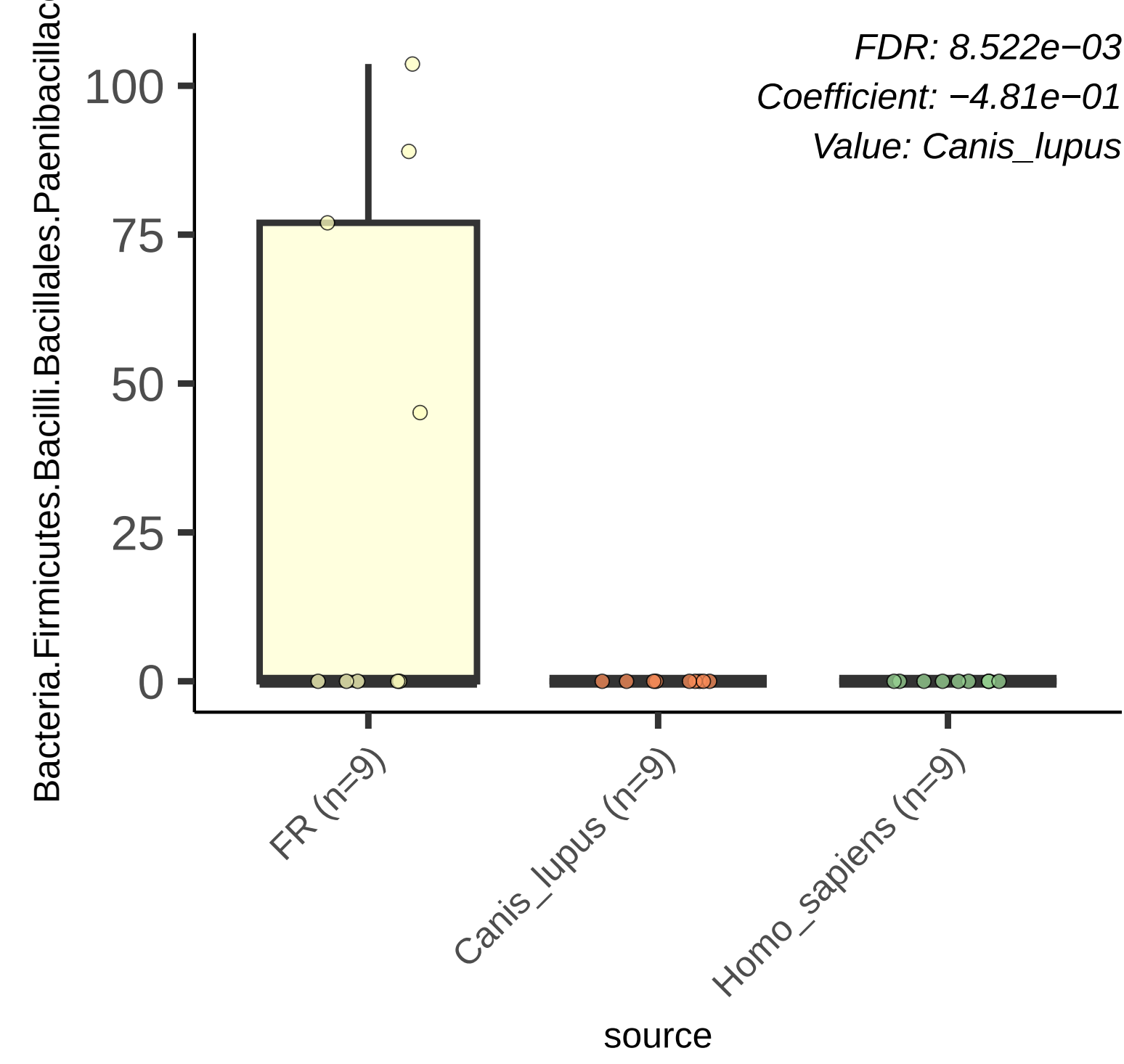
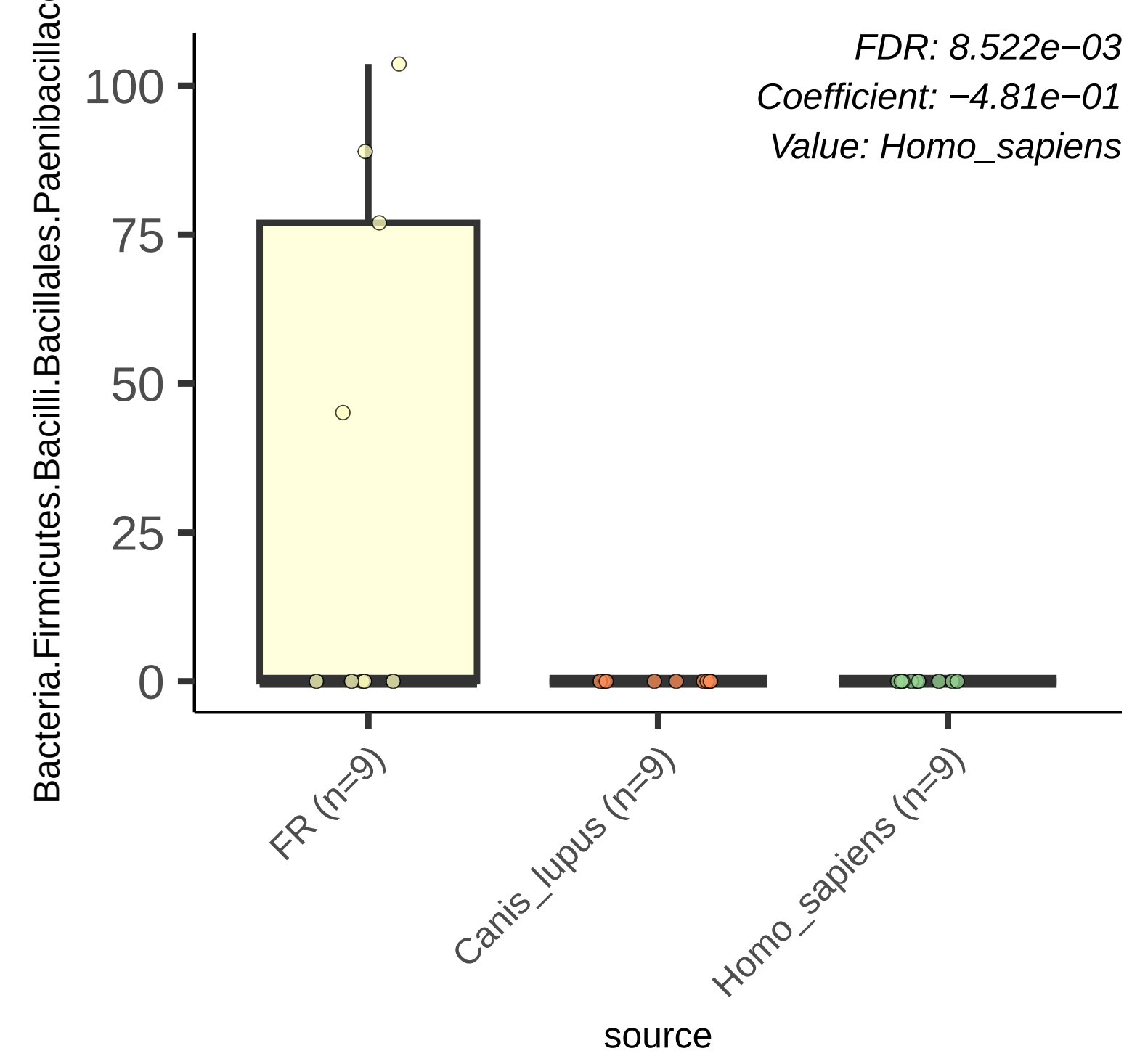
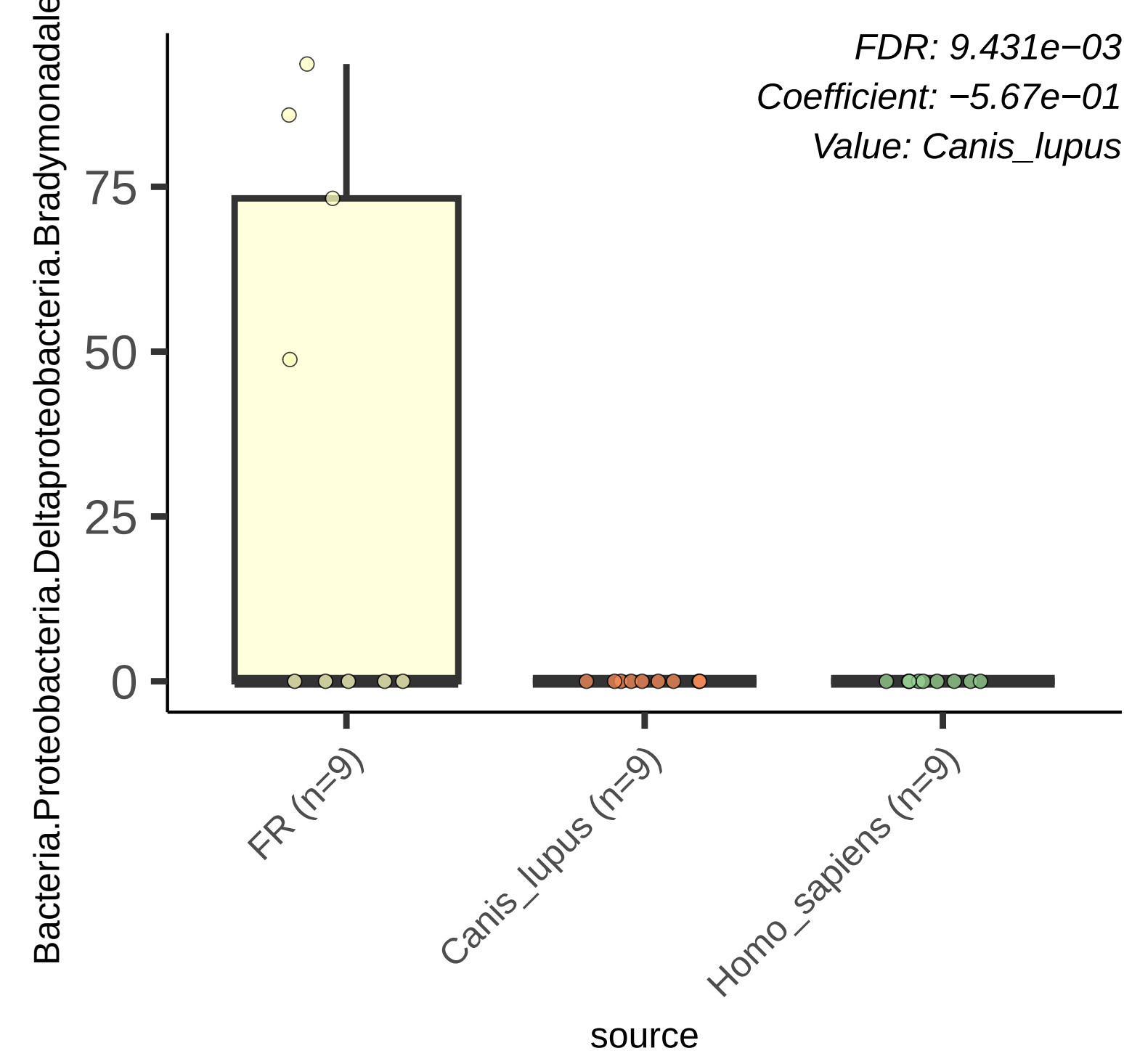
Canis_lupus (n=9)
Homo_sapiens (n=9)
source


FDR: 8.522e−03
Coefficient: −4.81e−01
Value: Canis_lupus
Bacteria.Firmicutes.Bacilli.Bacillales.Paenibacillac
100
75
50
25
0
FR (n=9)
Canis_lupus (n=9)
Homo_sapiens (n=9)
source


FDR: 8.522e−03
Coefficient: −4.81e−01
Value: Homo_sapiens
Bacteria.Firmicutes.Bacilli.Bacillales.Paenibacillac
100
75
50
25
0
FR (n=9)
Canis_lupus (n=9)
Homo_sapiens (n=9)
source


FDR: 9.431e−03
Coefficient: −5.67e−01
Value: Canis_lupus
Bacteria.Proteobacteria.Deltaproteobacteria.Bradymonadale
75
50
25
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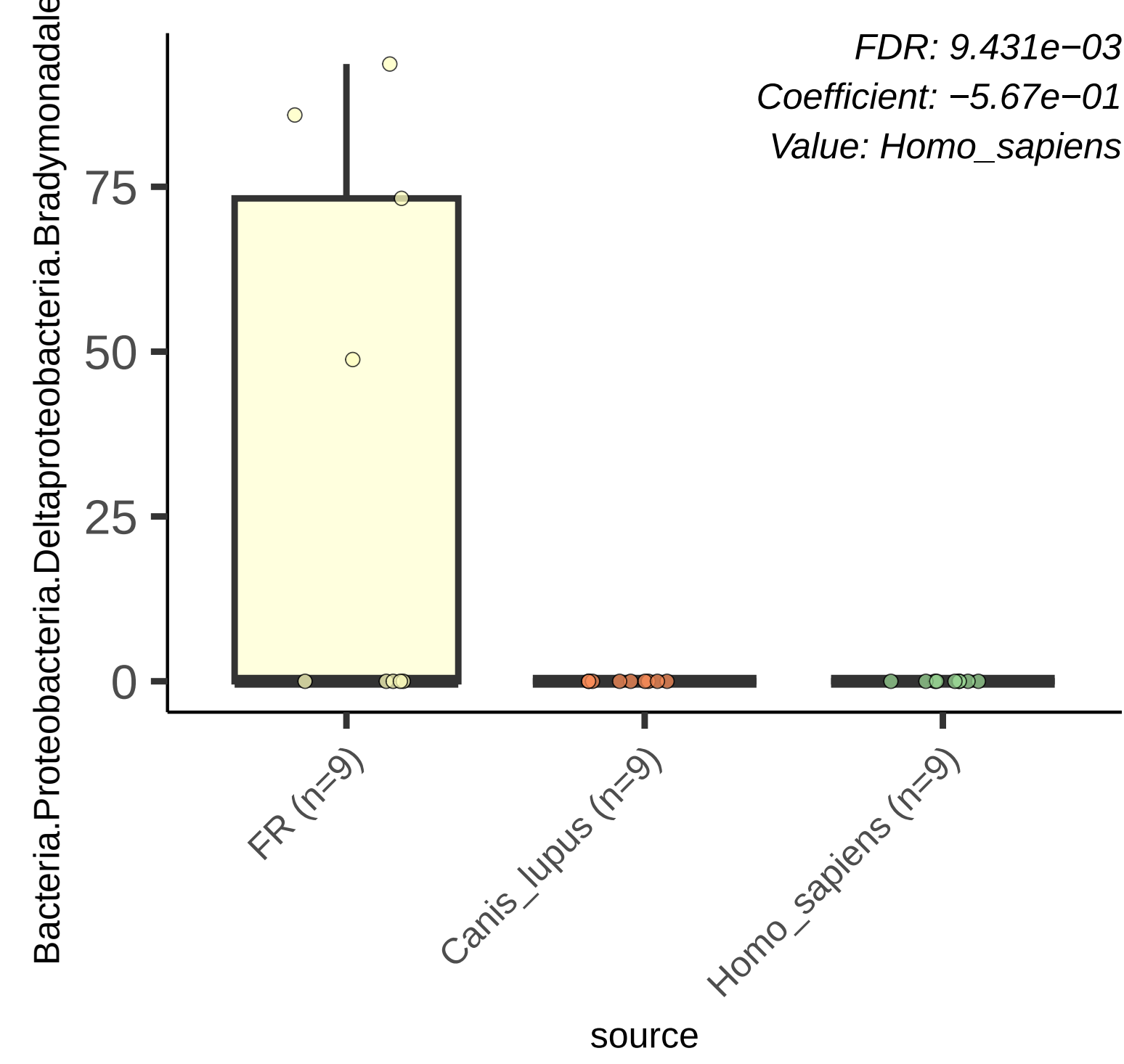
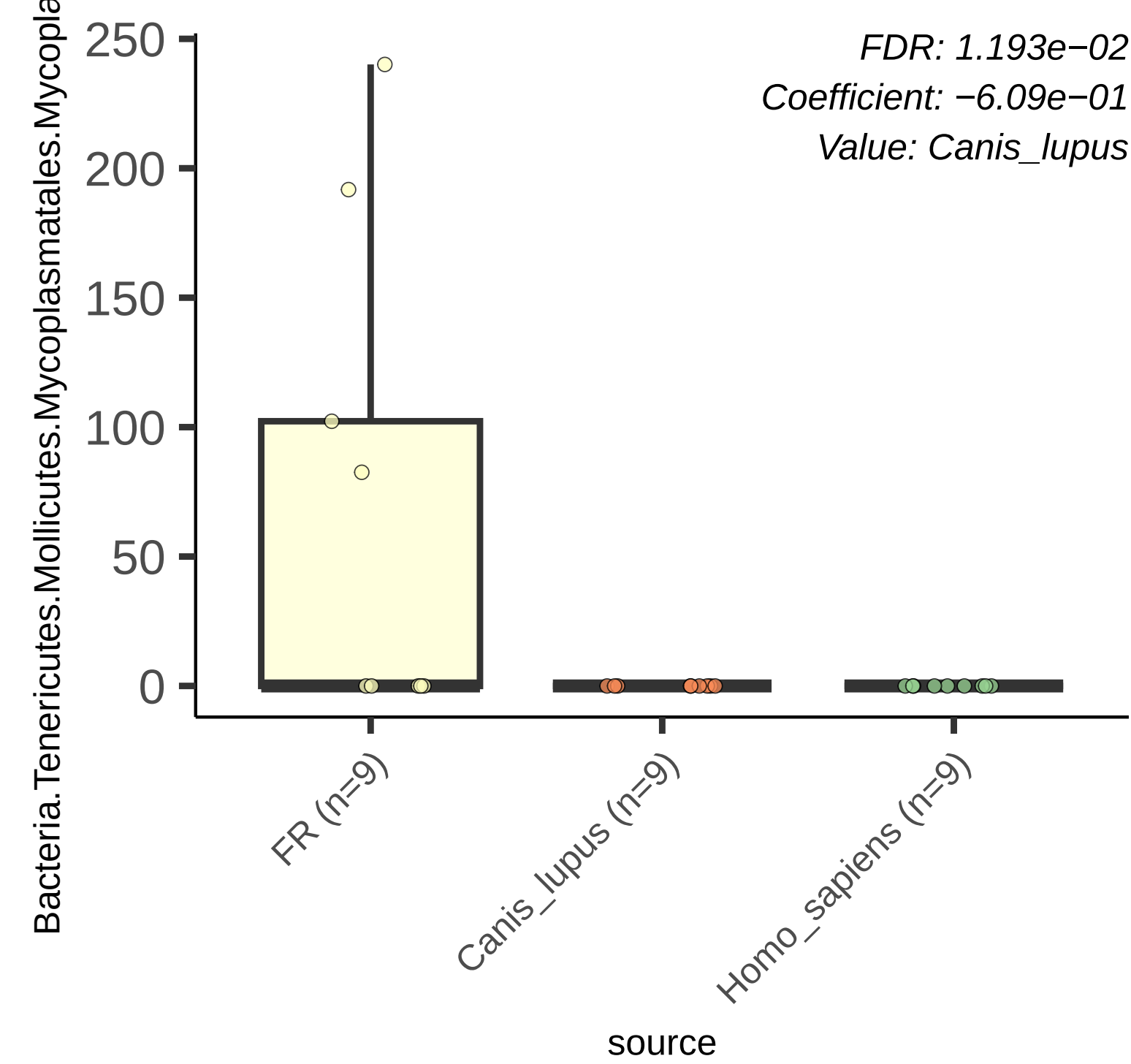
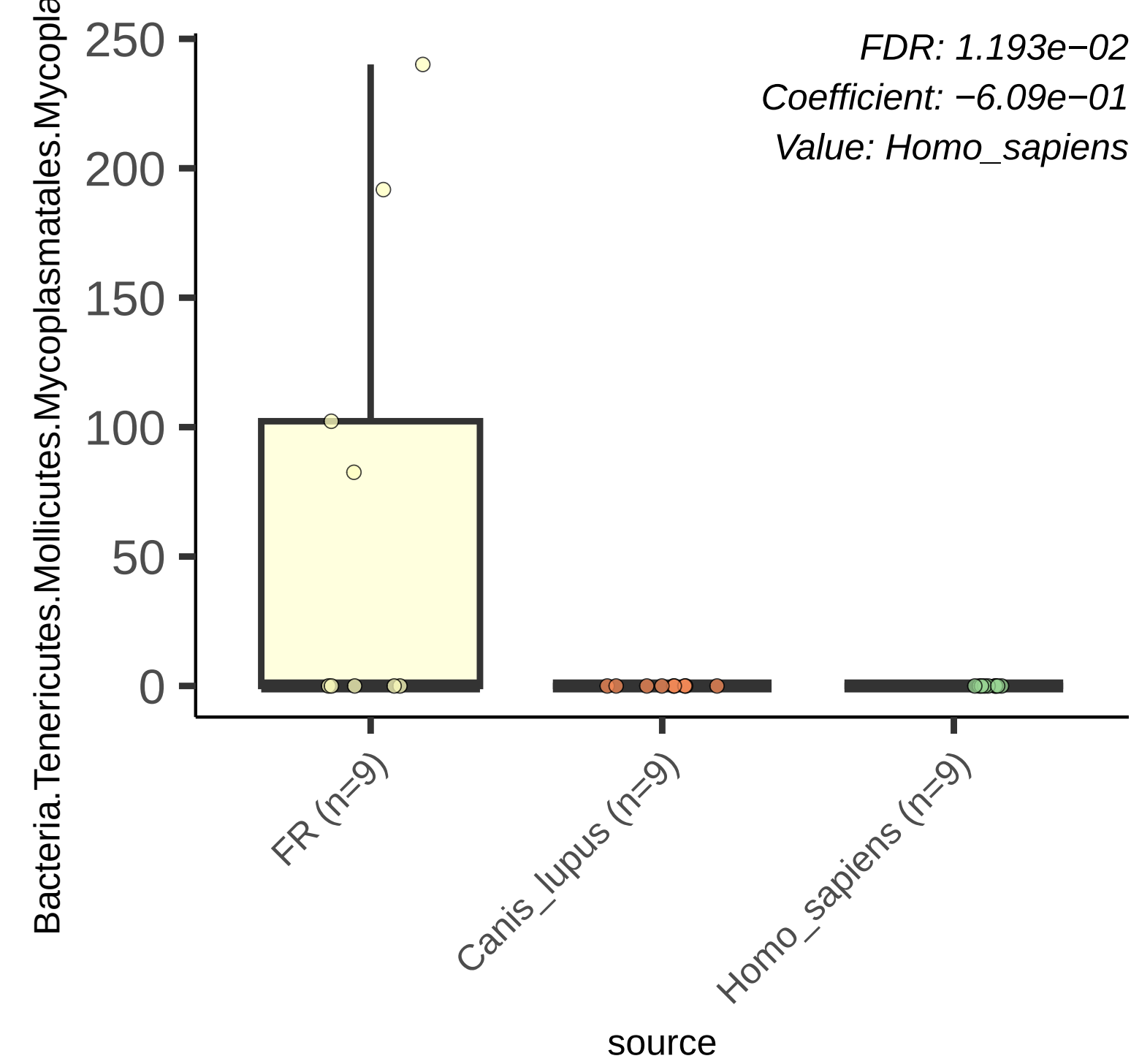
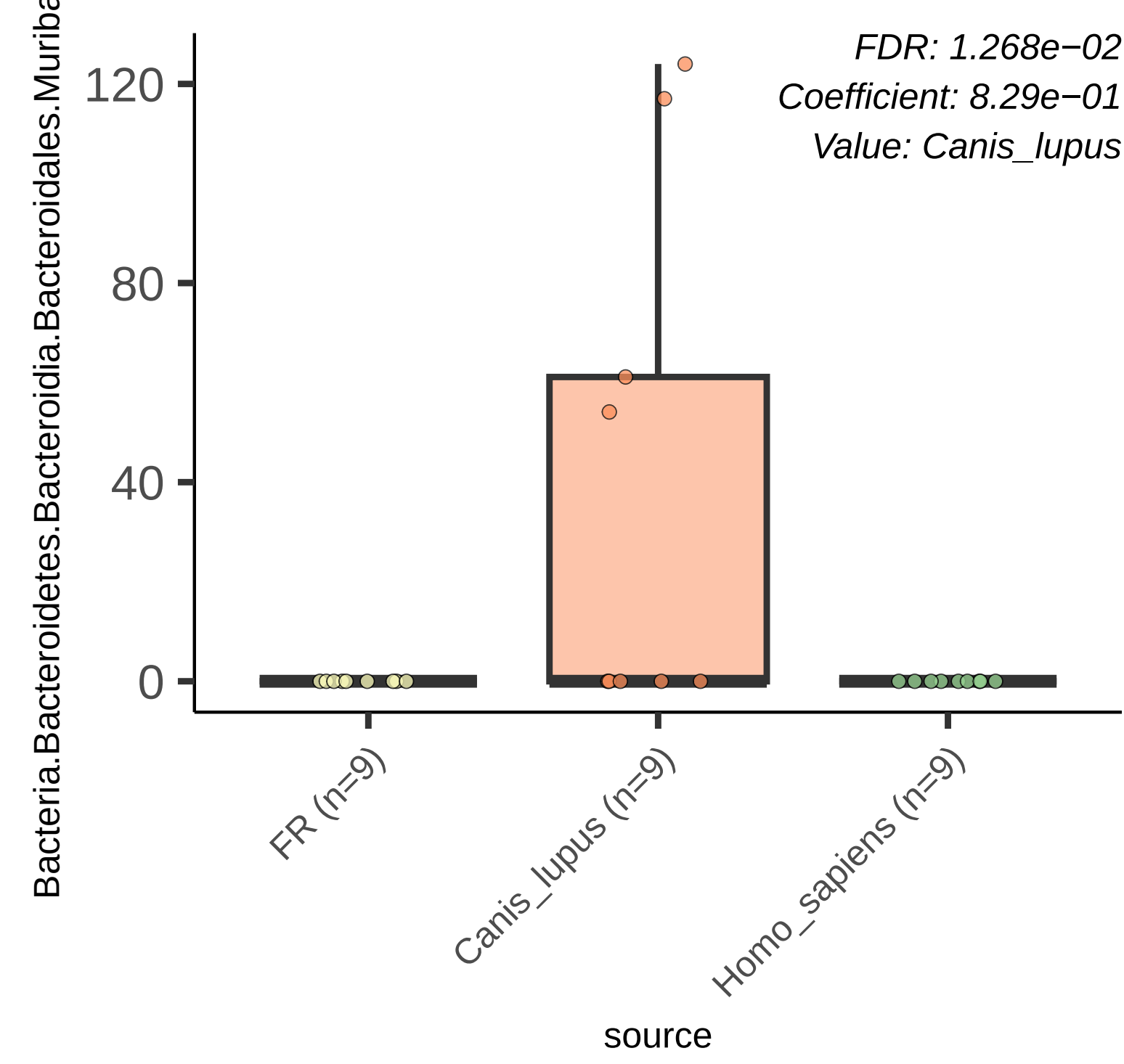
FR (n=9)
Canis_lupus (n=9)
Homo_sapiens (n=9)
source


FDR: 9.431e−03
Coefficient: −5.67e−01
Value: Homo_sapiens
Bacteria.Proteobacteria.Deltaproteobacteria.Bradymonadale
75
50
25
0
FR (n=9)
Canis_lupus (n=9)
Homo_sapiens (n=9)
source


FDR: 1.193e−02
Coefficient: −6.09e−01
Value: Canis_lupus
Bacteria.Tenericutes.Mollicutes.Mycoplasmatales.Mycopla
250
200
150
100
50
0
FR (n=9)
Canis_lupus (n=9)
Homo_sapiens (n=9)
source


FDR: 1.193e−02
Coefficient: −6.09e−01
Value: Homo_sapiens
Bacteria.Tenericutes.Mollicutes.Mycoplasmatales.Mycopla
250
200
150
100
50
0
FR (n=9)
Canis_lupus (n=9)
Homo_sapiens (n=9)
source


FDR: 1.268e−02
Coefficient: 8.29e−01
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Bacteria.Bacteroidetes.Bacteroidia.Bacteroidales.Muriba
120
80
40
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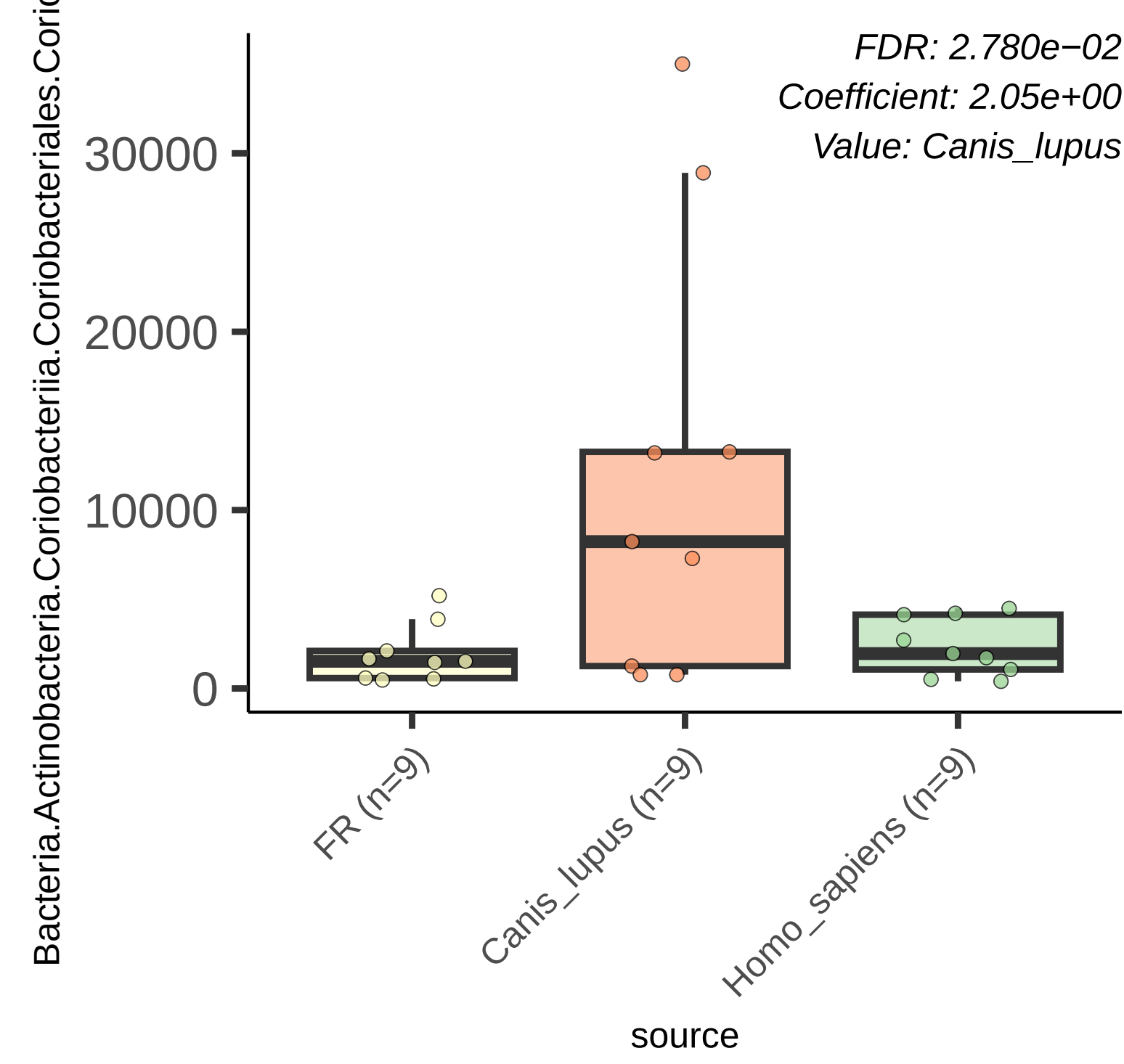
Canis_lupus (n=9)
Homo_sapiens (n=9)
source


FDR: 2.780e−02
Coefficient: 2.05e+00
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Bacteria.Actinobacteria.Coriobacteriia.Coriobacteriales.Corio
30000
20000
10000
0
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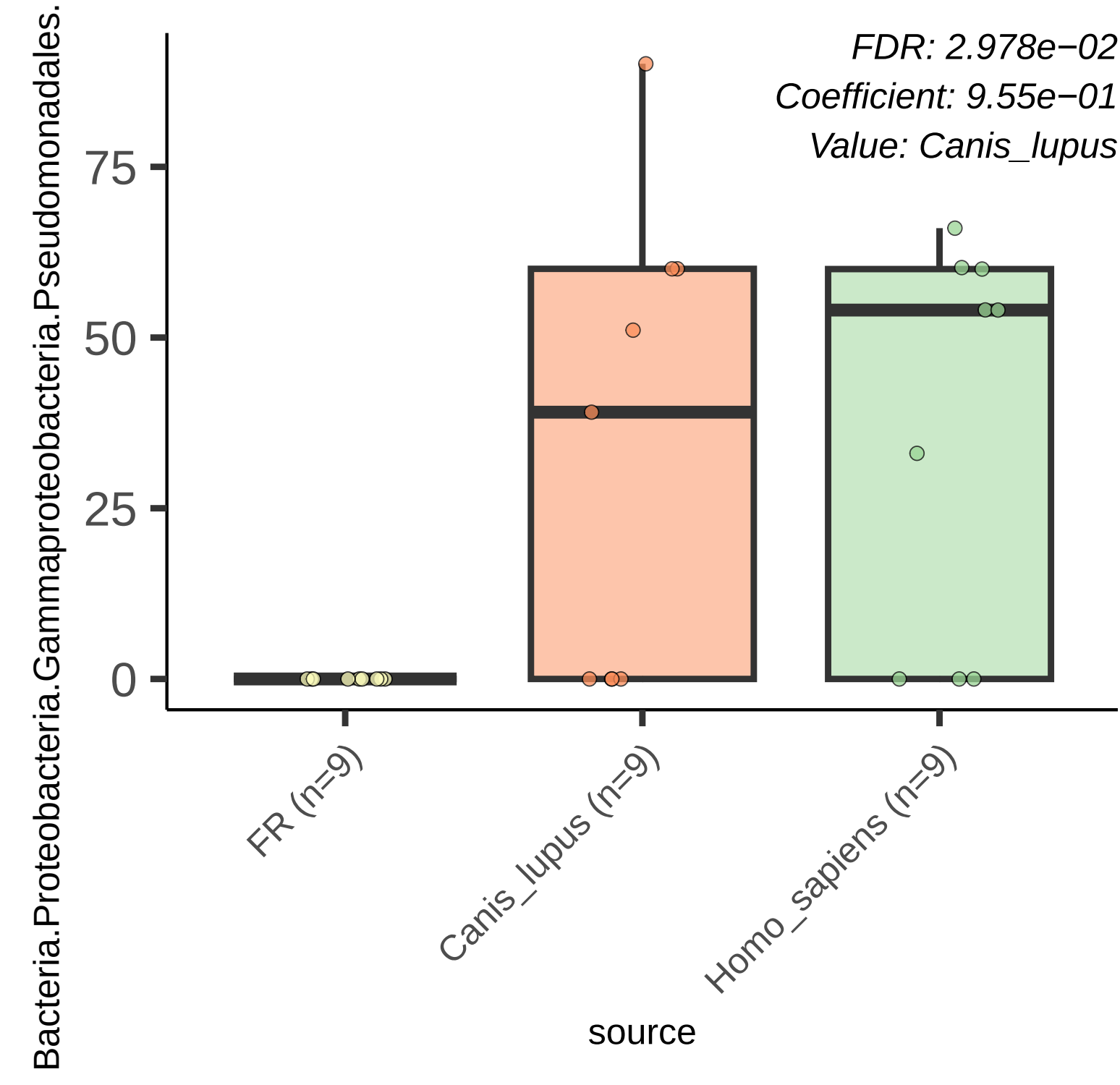
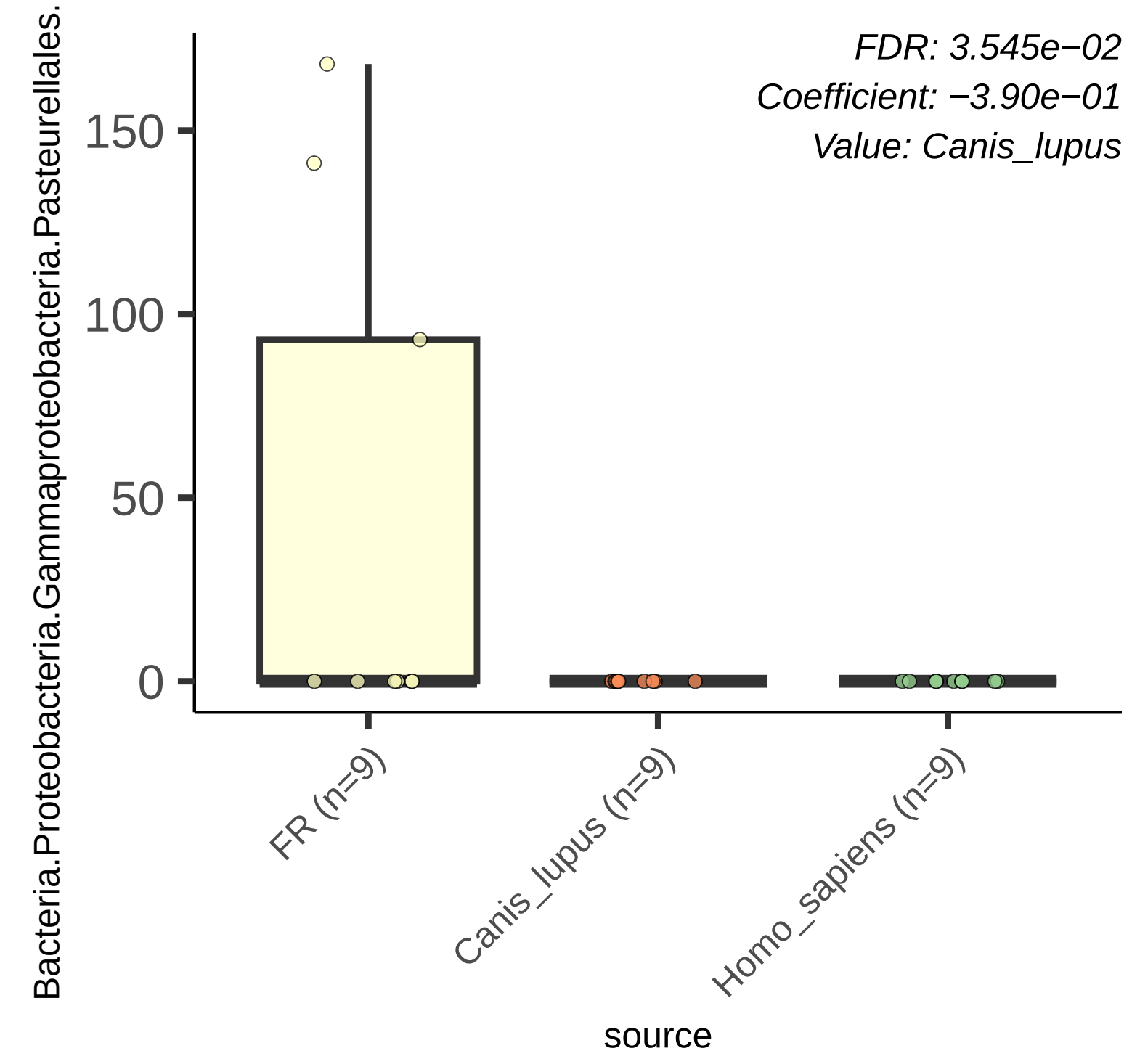
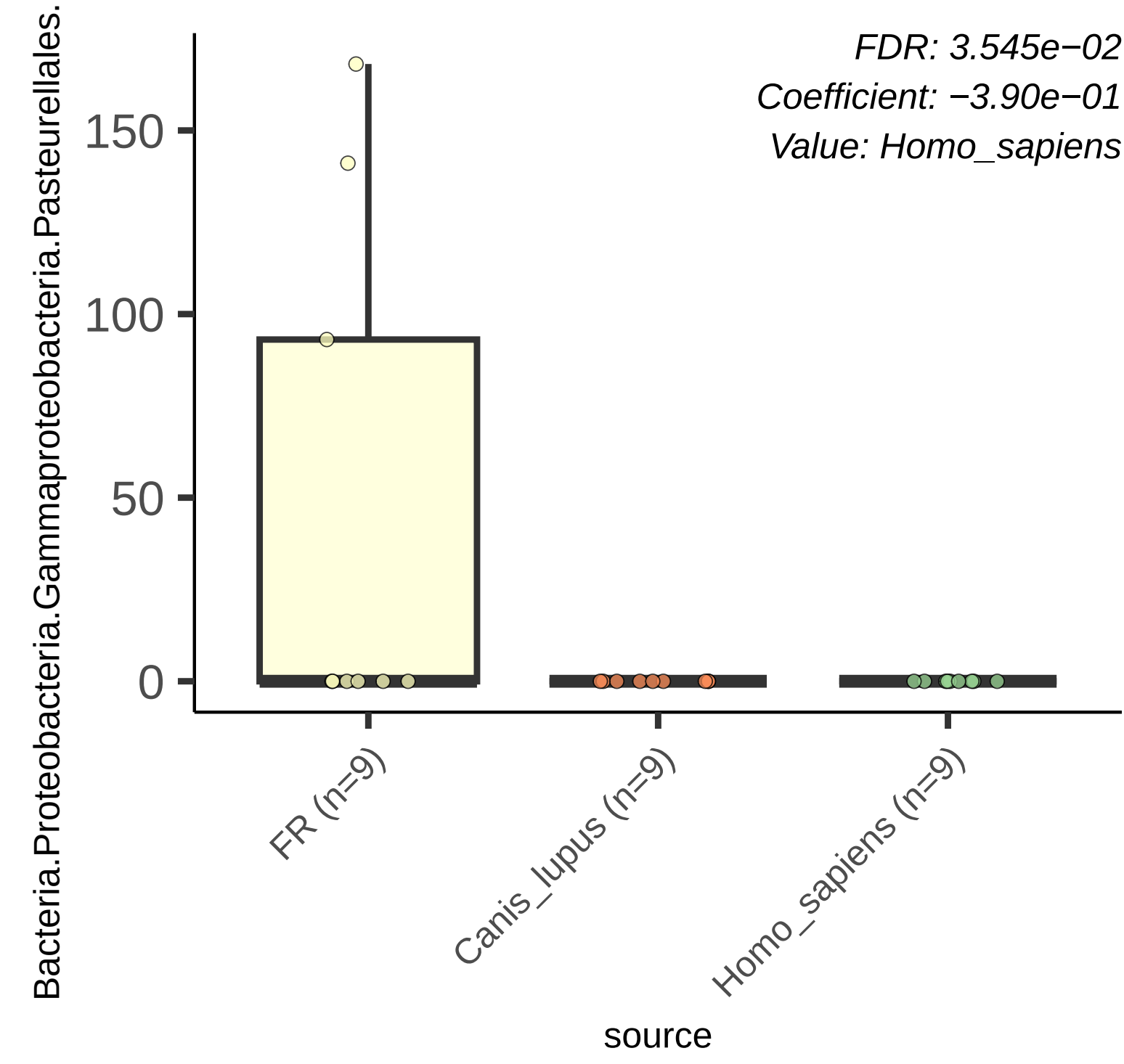
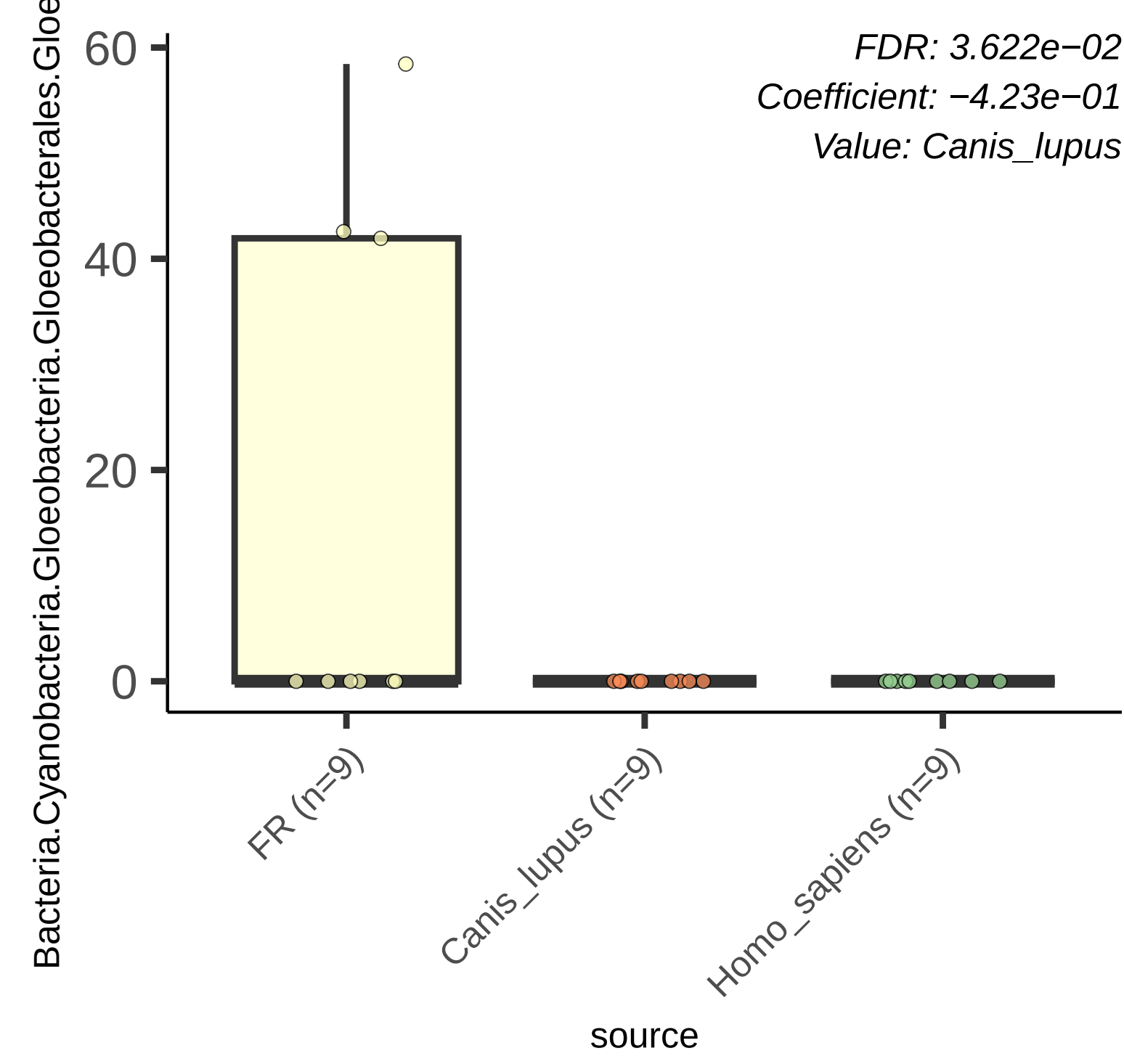
Canis_lupus (n=9)
Homo_sapiens (n=9)
source


FDR: 2.978e−02
Coefficient: 9.55e−01
Value: Canis_lupus
Bacteria.Proteobacteria.Gammaproteobacteria.Pseudomonadales.
75
50
25
0
FR (n=9)
Canis_lupus (n=9)
Homo_sapiens (n=9)
source


FDR: 3.545e−02
Coefficient: −3.90e−01
Value: Canis_lupus
Bacteria.Proteobacteria.Gammaproteobacteria.Pasteurellales.
150
100
50
0
FR (n=9)
Canis_lupus (n=9)
Homo_sapiens (n=9)
source


FDR: 3.545e−02
Coefficient: −3.90e−01
Value: Homo_sapiens
Bacteria.Proteobacteria.Gammaproteobacteria.Pasteurellales.
150
100
50
0
FR (n=9)
Canis_lupus (n=9)
Homo_sapiens (n=9)
source


FDR: 3.622e−02
Coefficient: −4.23e−01
Value: Canis_lupus
Bacteria.Cyanobacteria.Gloeobacteria.Gloeobacterales.Gloe
60
40
20
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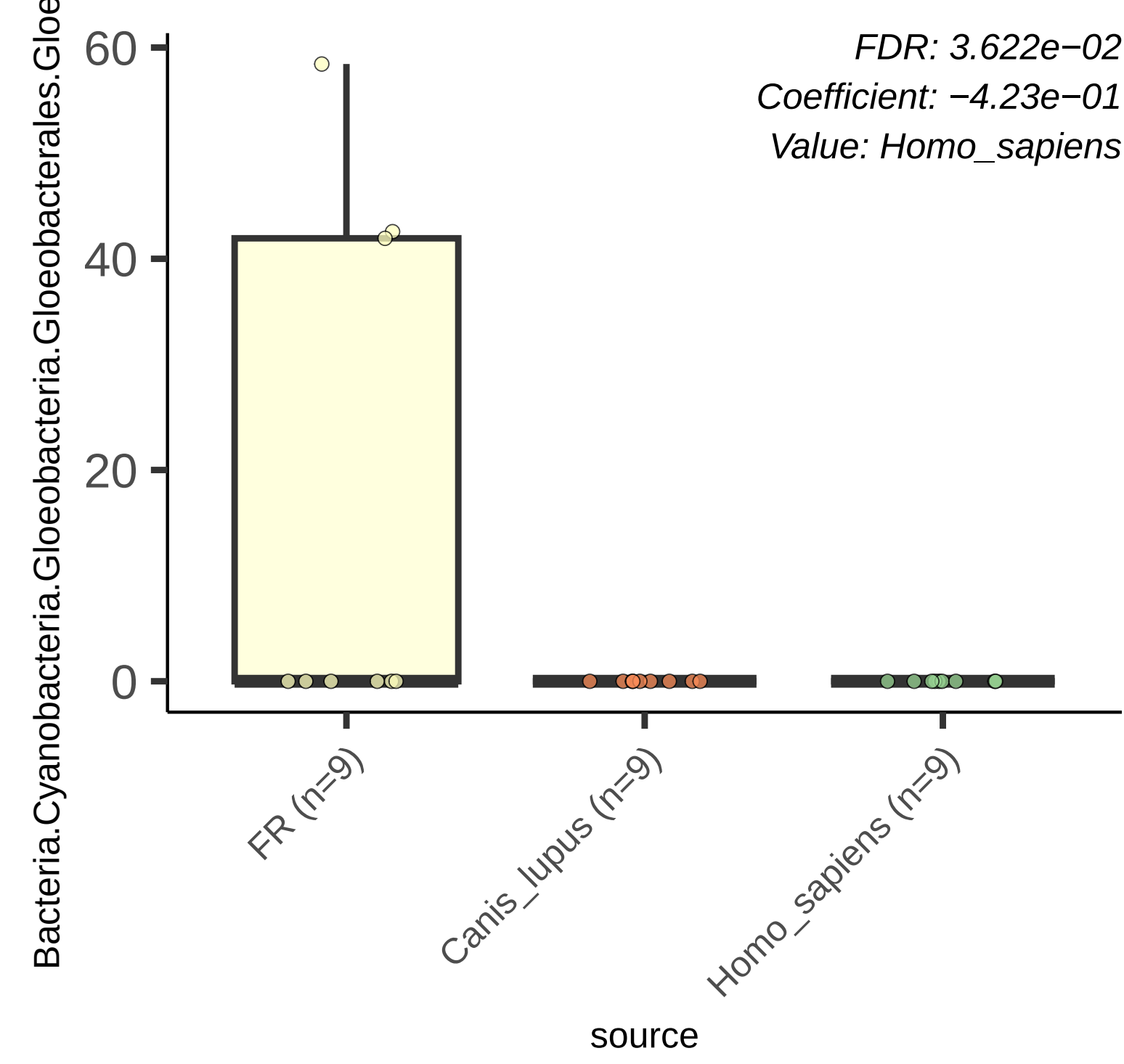
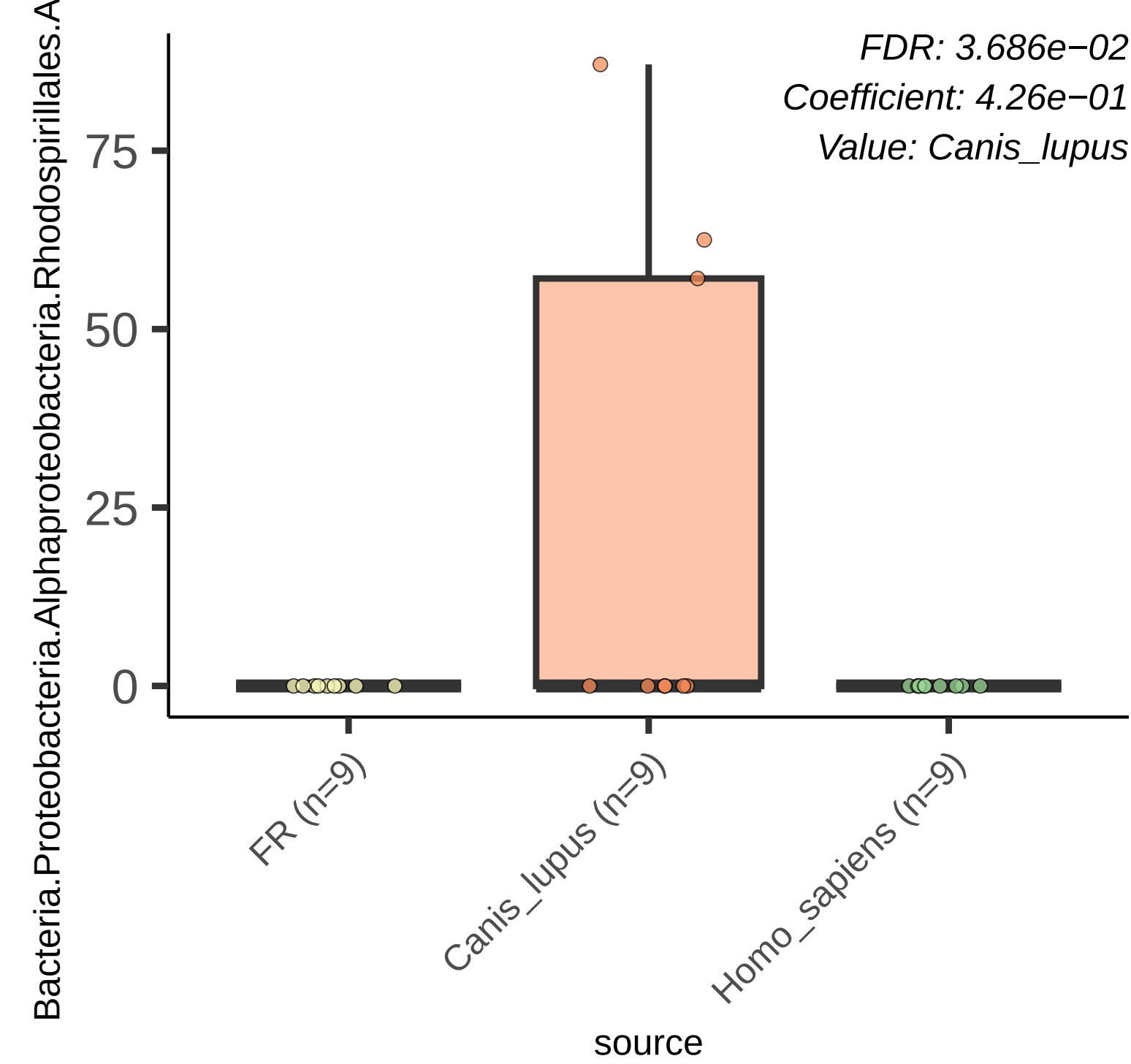
FR (n=9)
Canis_lupus (n=9)
Homo_sapiens (n=9)
source


FDR: 3.622e−02
Coefficient: −4.23e−01
Value: Homo_sapiens
Bacteria.Cyanobacteria.Gloeobacteria.Gloeobacterales.Gloe
60
40
20
0
FR (n=9)
Canis_lupus (n=9)
Homo_sapiens (n=9)
source


FDR: 3.686e−02
Coefficient: 4.26e−01
Value: Canis_lupus
Bacteria.Proteobacteria.Alphaproteobacteria.Rhodospirillales.A
75
50
25
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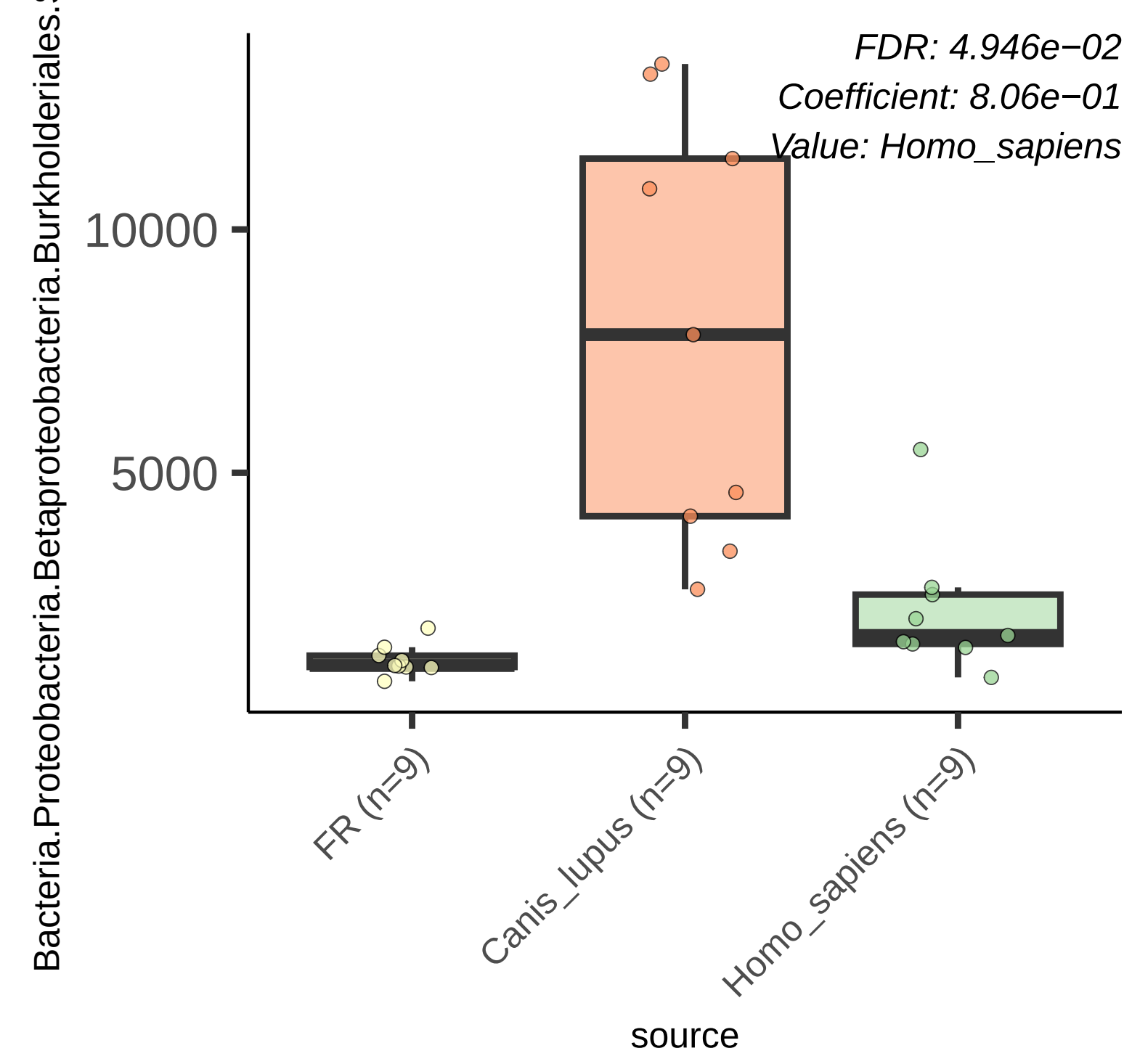
FR (n=9)
Canis_lupus (n=9)
Homo_sapiens (n=9)
source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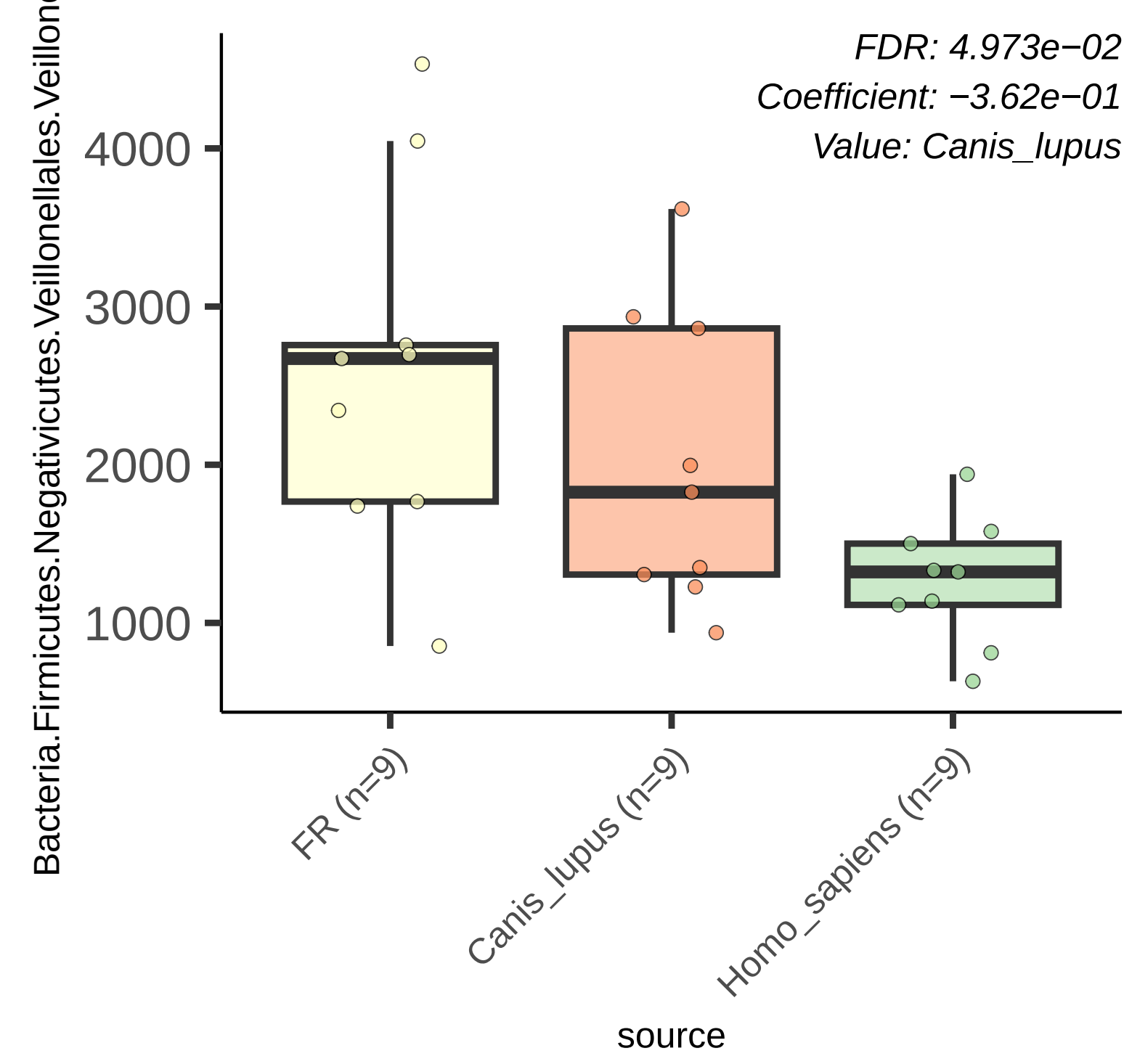
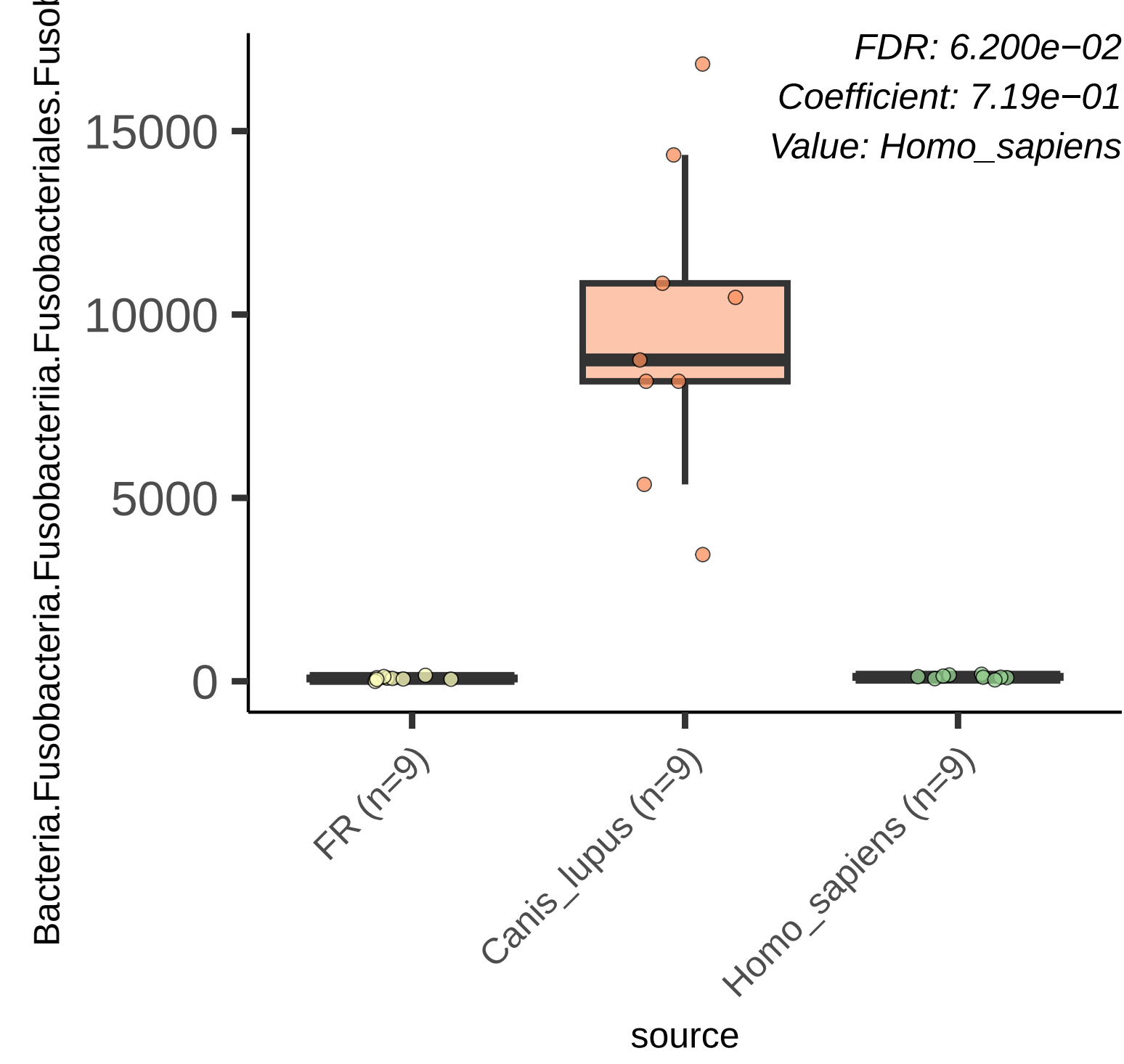
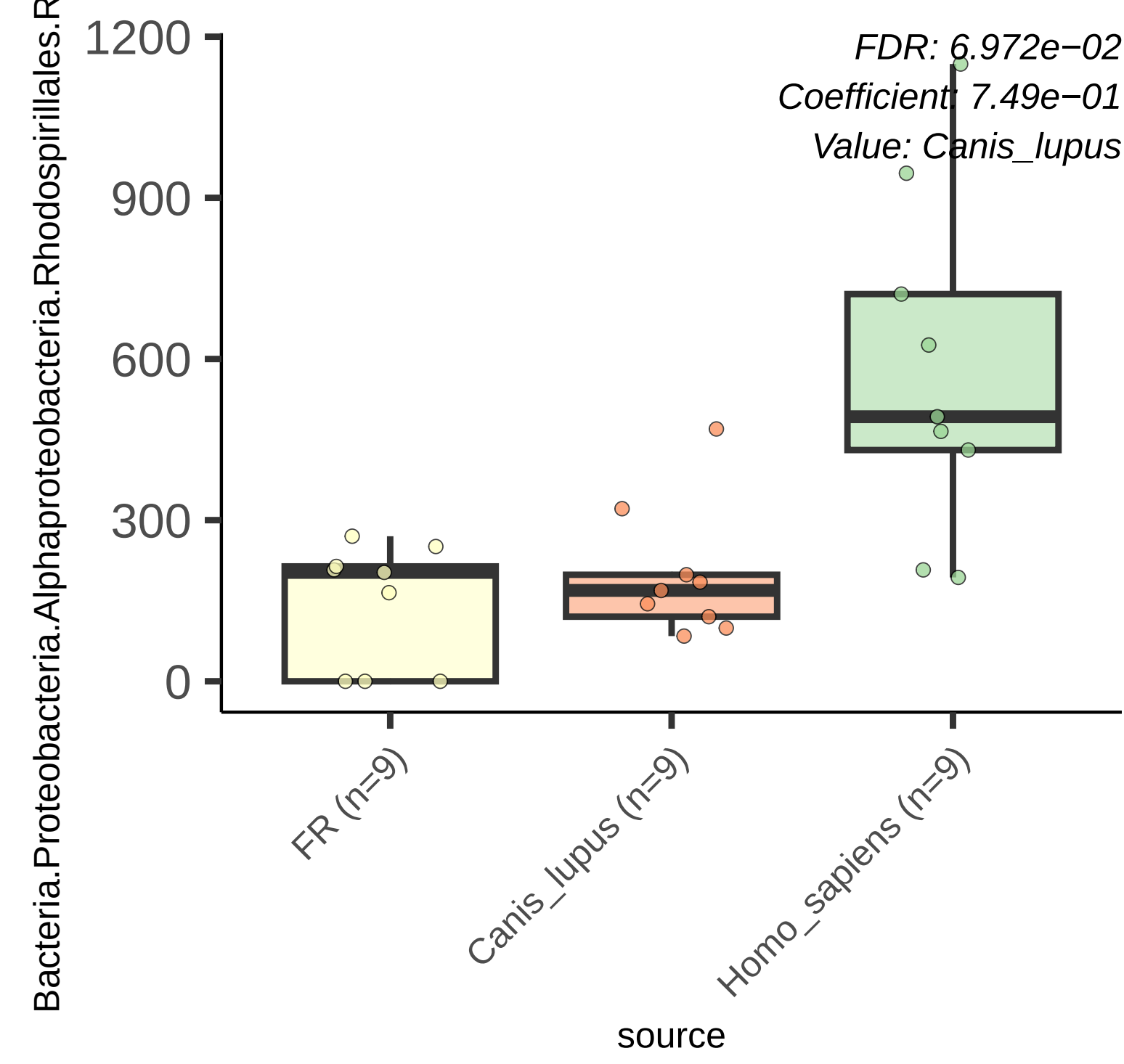
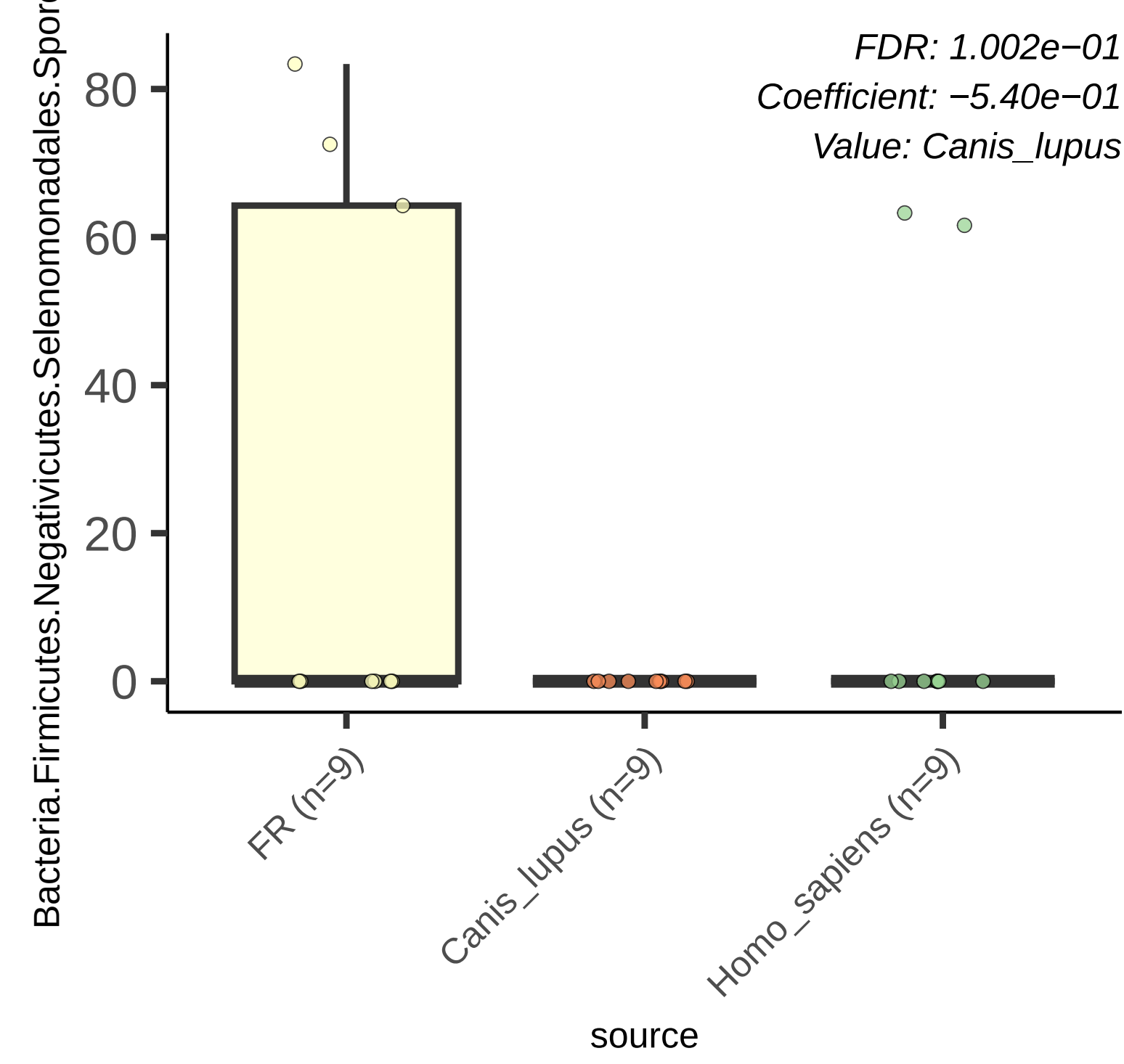
Coefficient: 8.06e−01
Value: Homo_sapiens
10000
5000
Bacteria.Proteobacteria.Betaproteobacteria.Burkholderiales.
FR (n=9)
Canis_lupus (n=9)
Homo_sapiens (n=9)
source


FDR: 4.973e−02
Coefficient: −3.62e−01
Value: Canis_lupus
Bacteria.Firmicutes.Negativicutes.Veillonellales.Veillone
4000
3000
2000
1000
FR (n=9)
Canis_lupus (n=9)
Homo_sapiens (n=9)
source


FDR: 6.200e−02
Coefficient: 7.19e−01
Value: Homo_sapiens
15000
10000
5000
0
Bacteria.Fusobacteria.Fusobacteriia.Fusobacteriales.Fusob
FR (n=9)
Canis_lupus (n=9)
Homo_sapiens (n=9)
source


FDR: 6.972e−02
Coefficient: 7.49e−01
Value: Canis_lupus
1200
900
600
300
0
Bacteria.Proteobacteria.Alphaproteobacteria.Rhodospirillales.R
FR (n=9)
Canis_lupus (n=9)
Homo_sapiens (n=9)
source


FDR: 1.002e−01
Coefficient: −5.40e−01
Value: Canis_lupus
Bacteria.Firmicutes.Negativicutes.Selenomonadales.Spor
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60
40
20
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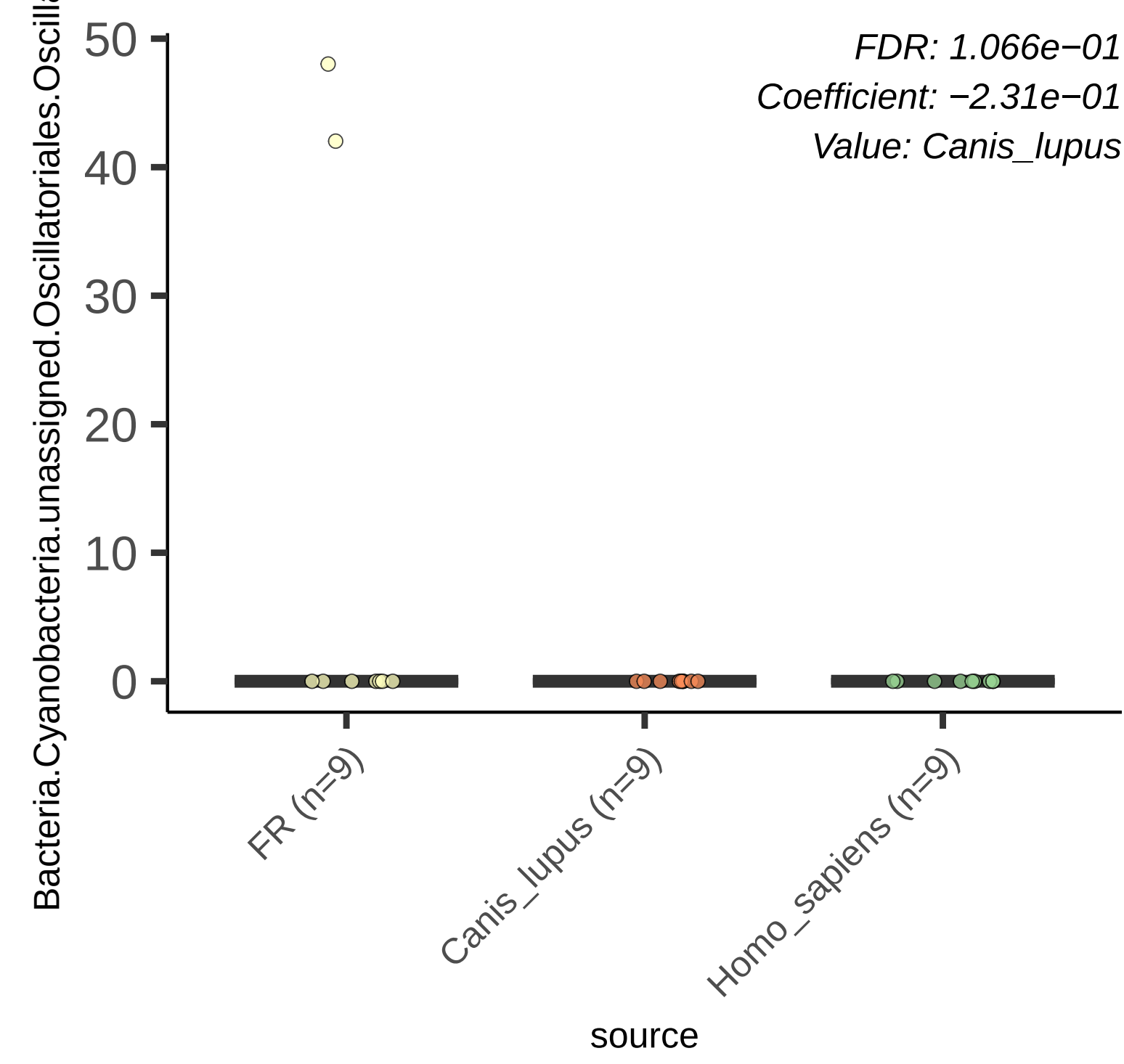
FR (n=9)
Canis_lupus (n=9)
Homo_sapiens (n=9)
source


FDR: 1.066e−01
Coefficient: −2.31e−01
Value: Canis_lupus
Bacteria.Cyanobacteria.unassigned.Oscillatoriales.Oscilla
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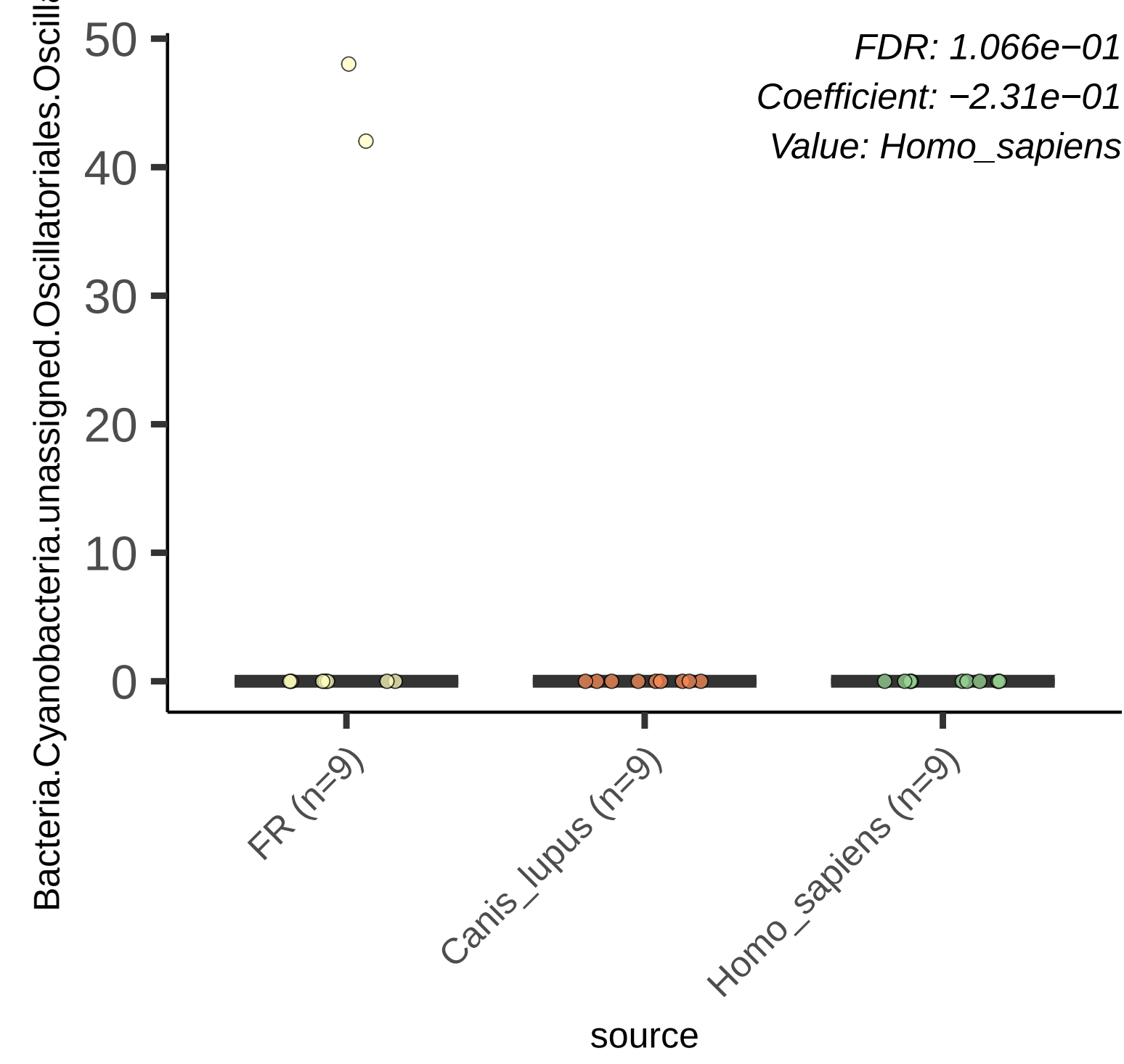
40
30
20
10
0
FR (n=9)
Canis_lupus (n=9)
Homo_sapiens (n=9)
source


FDR: 1.066e−01
Coefficient: −2.31e−01
Value: Homo_sapiens
Bacteria.Cyanobacteria.unassigned.Oscillatoriales.Oscilla
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30
20
10
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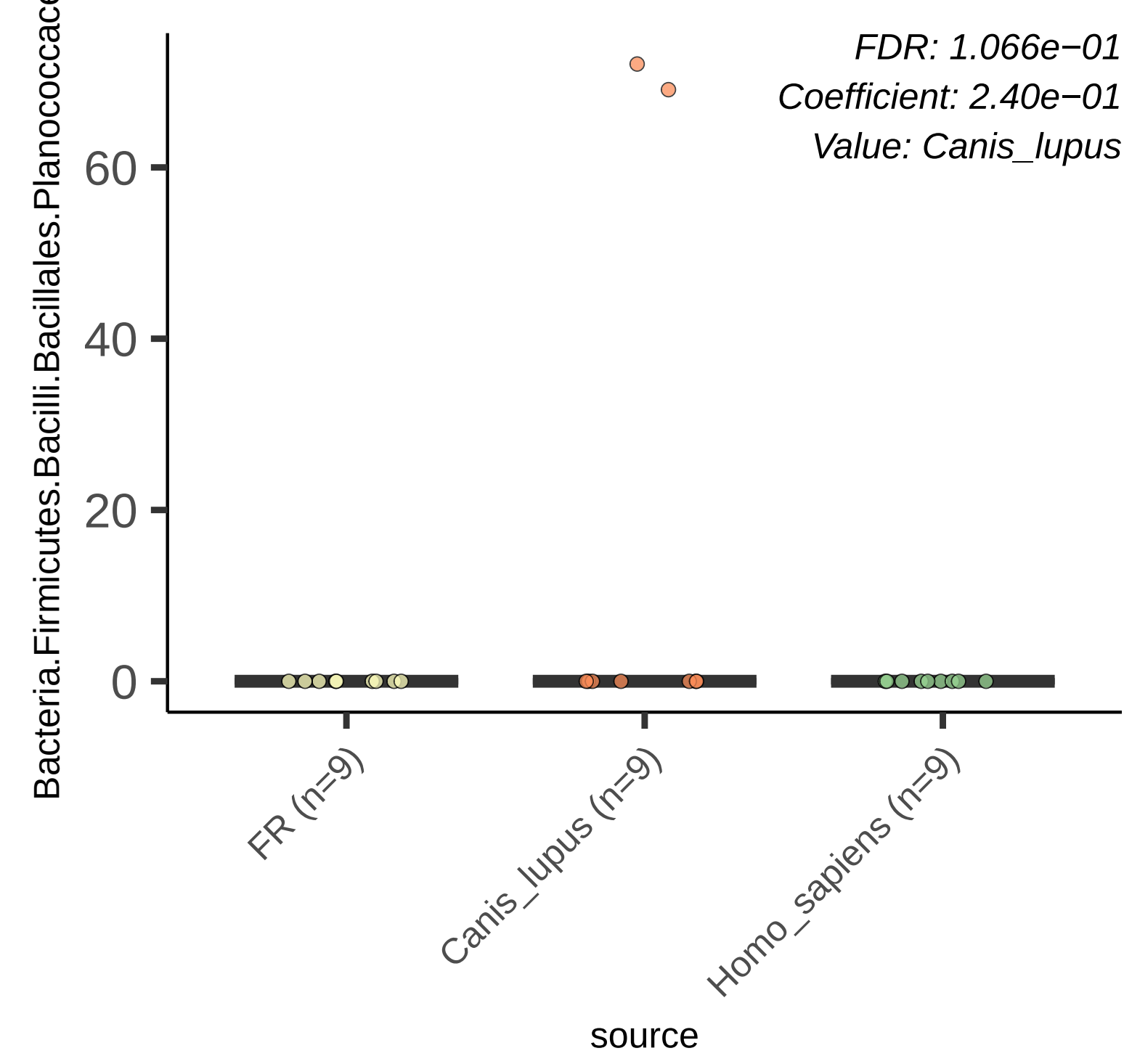
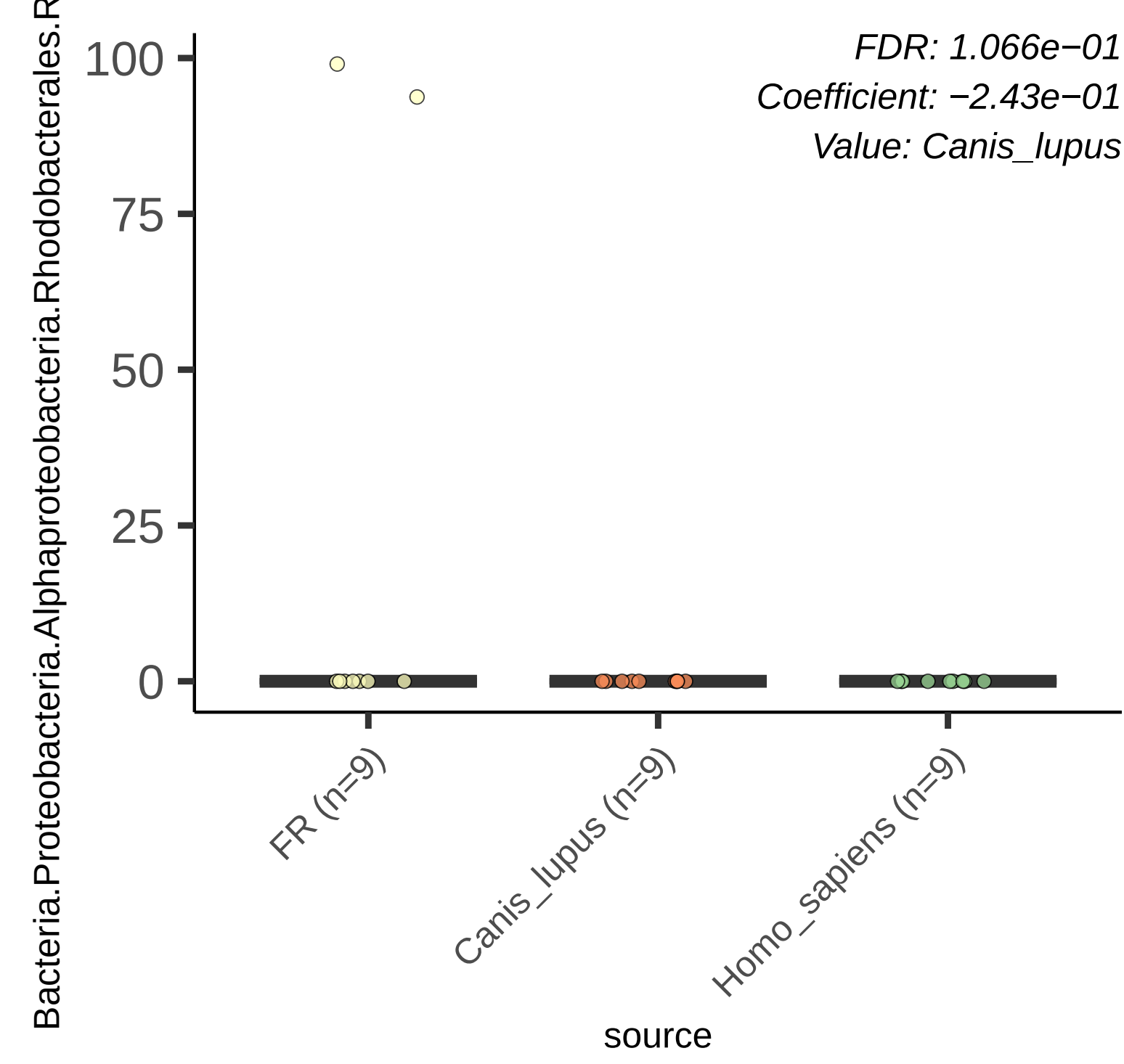
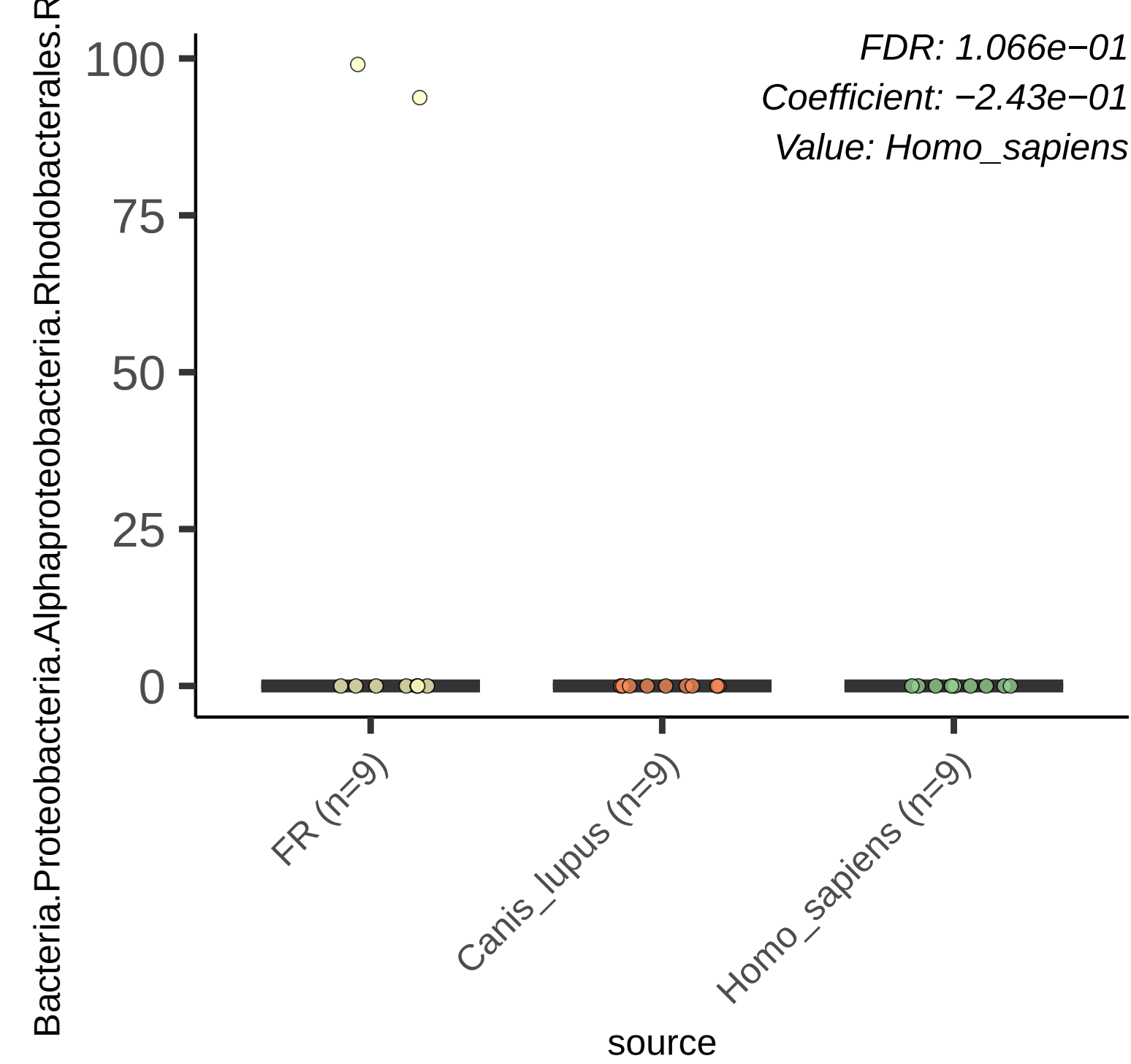
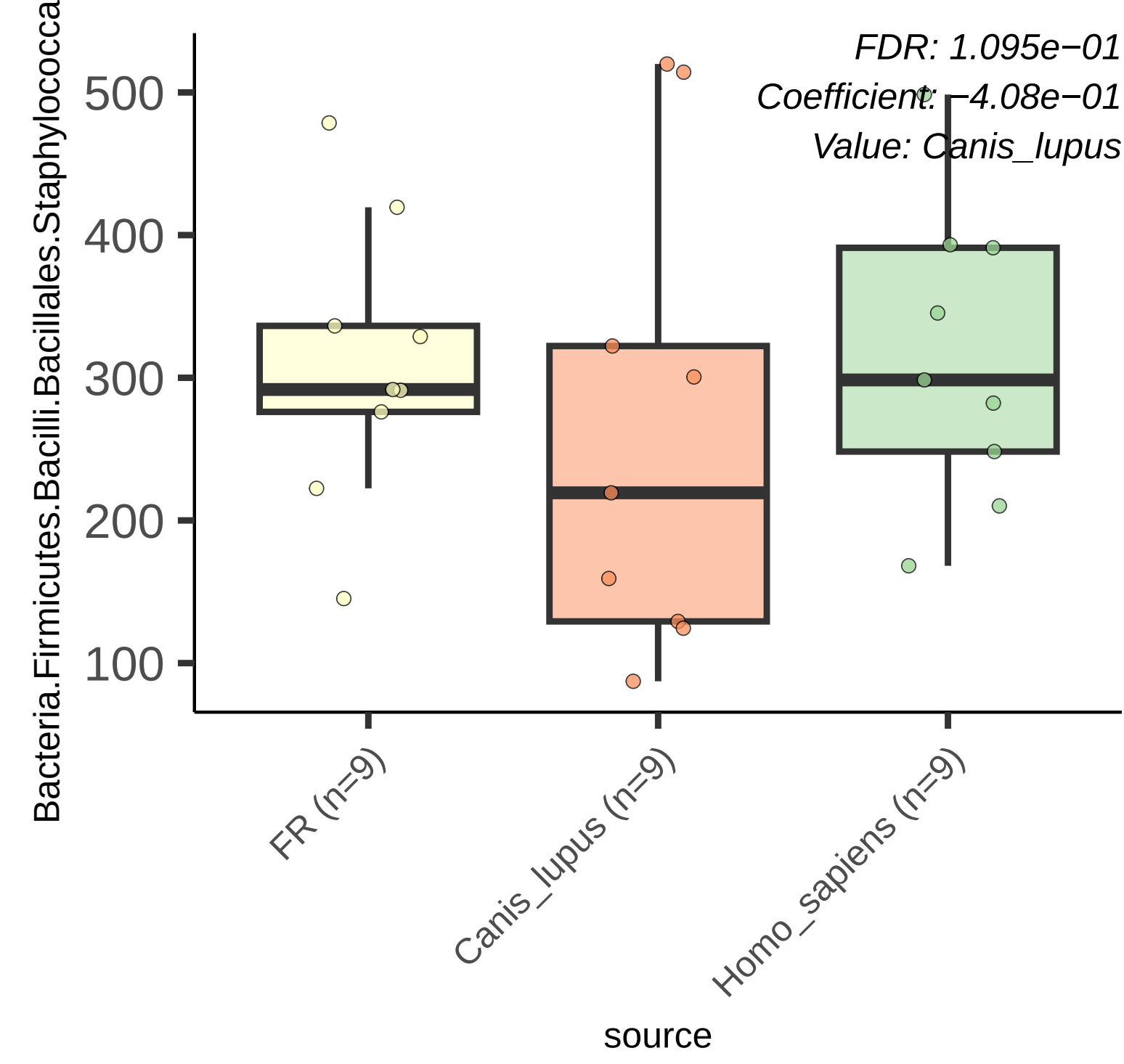
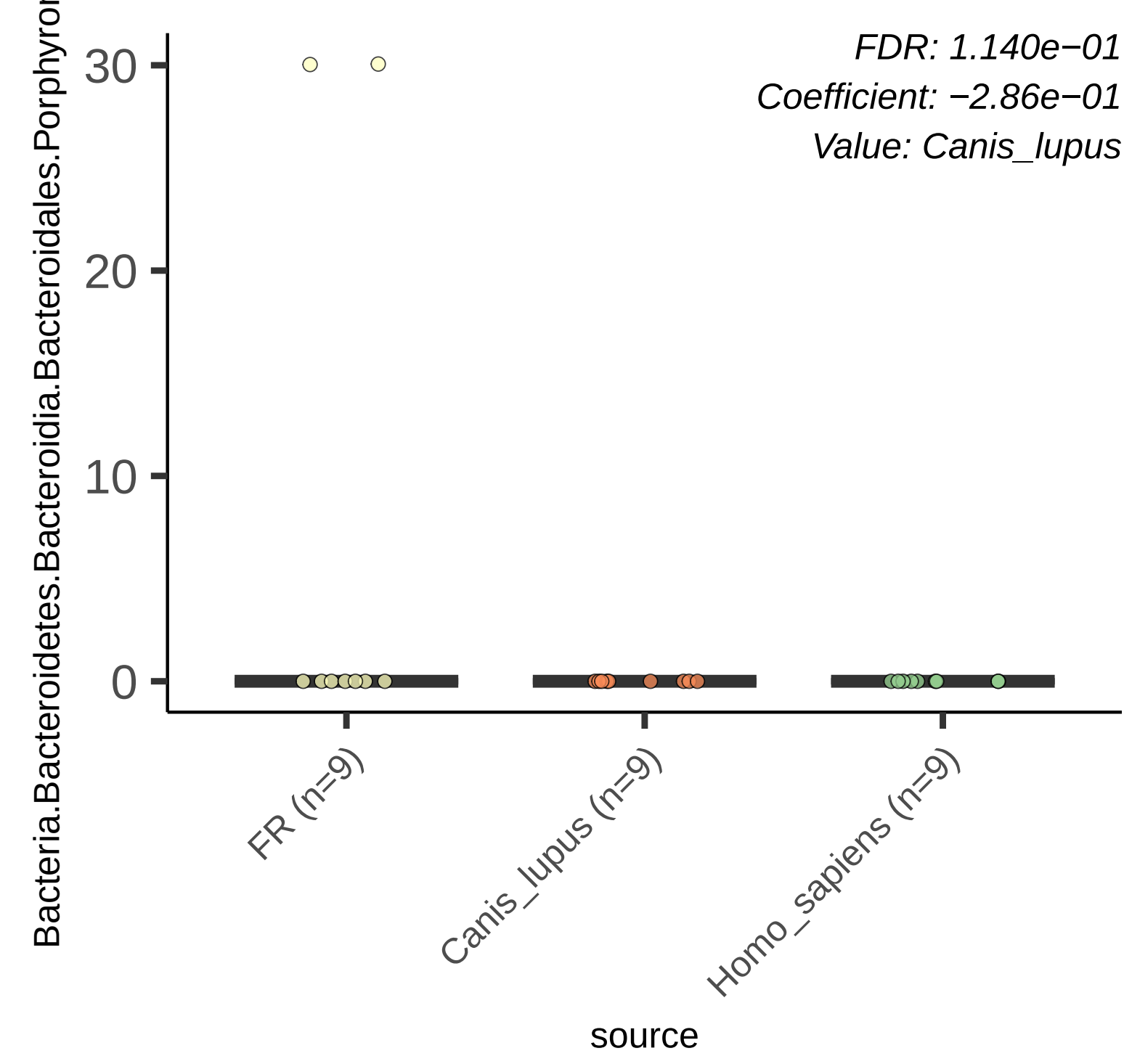
FR (n=9)
Canis_lupus (n=9)
Homo_sapiens (n=9)
source

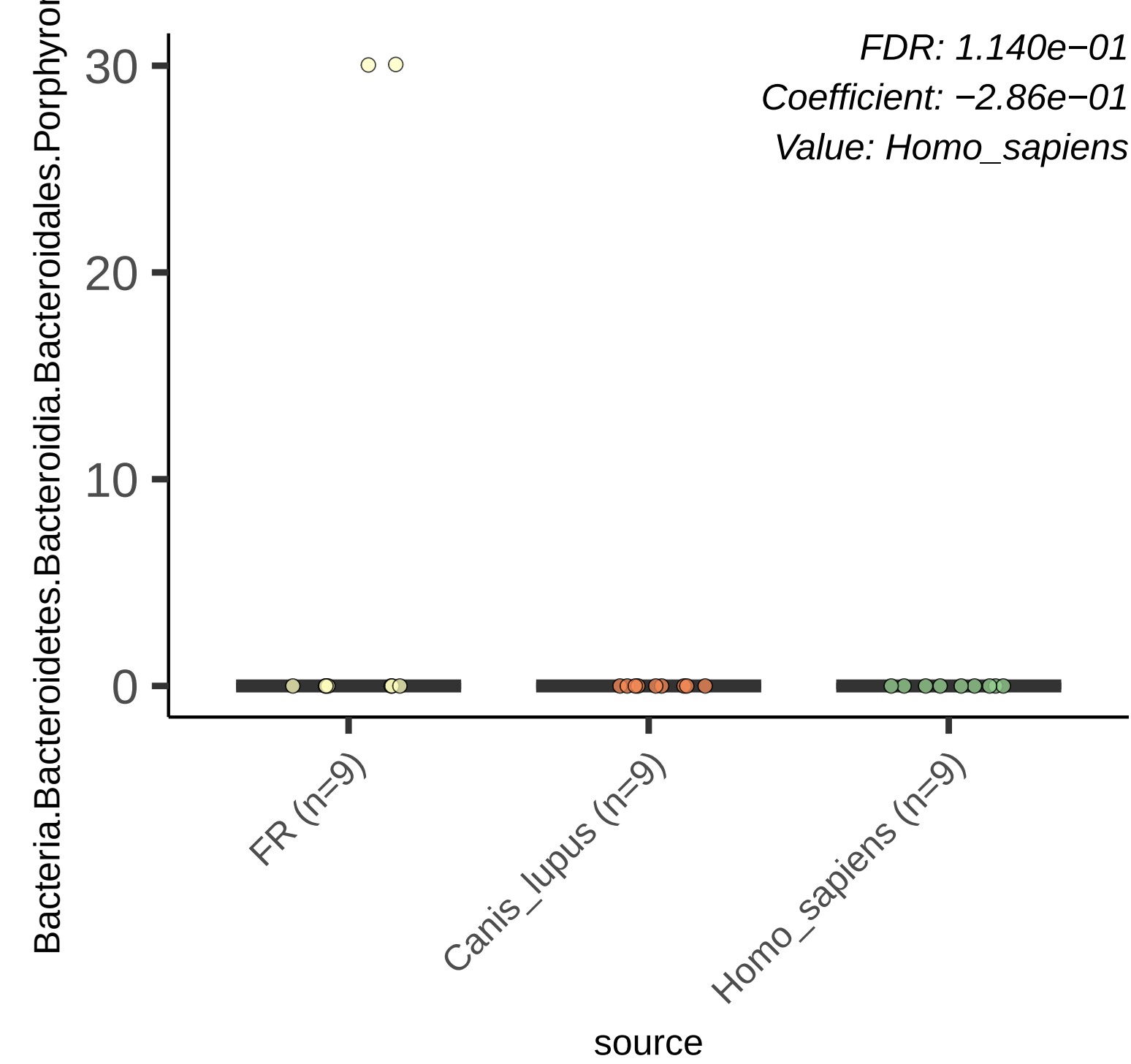

FDR: 1.066e−01
Coefficient: 2.40e−01
Value: Canis_lupus
Bacteria.Firmicutes.Bacilli.Bacillales.Planococcac
60
40
20
0
FR (n=9)
Canis_lupus (n=9)
Homo_sapiens (n=9)
source

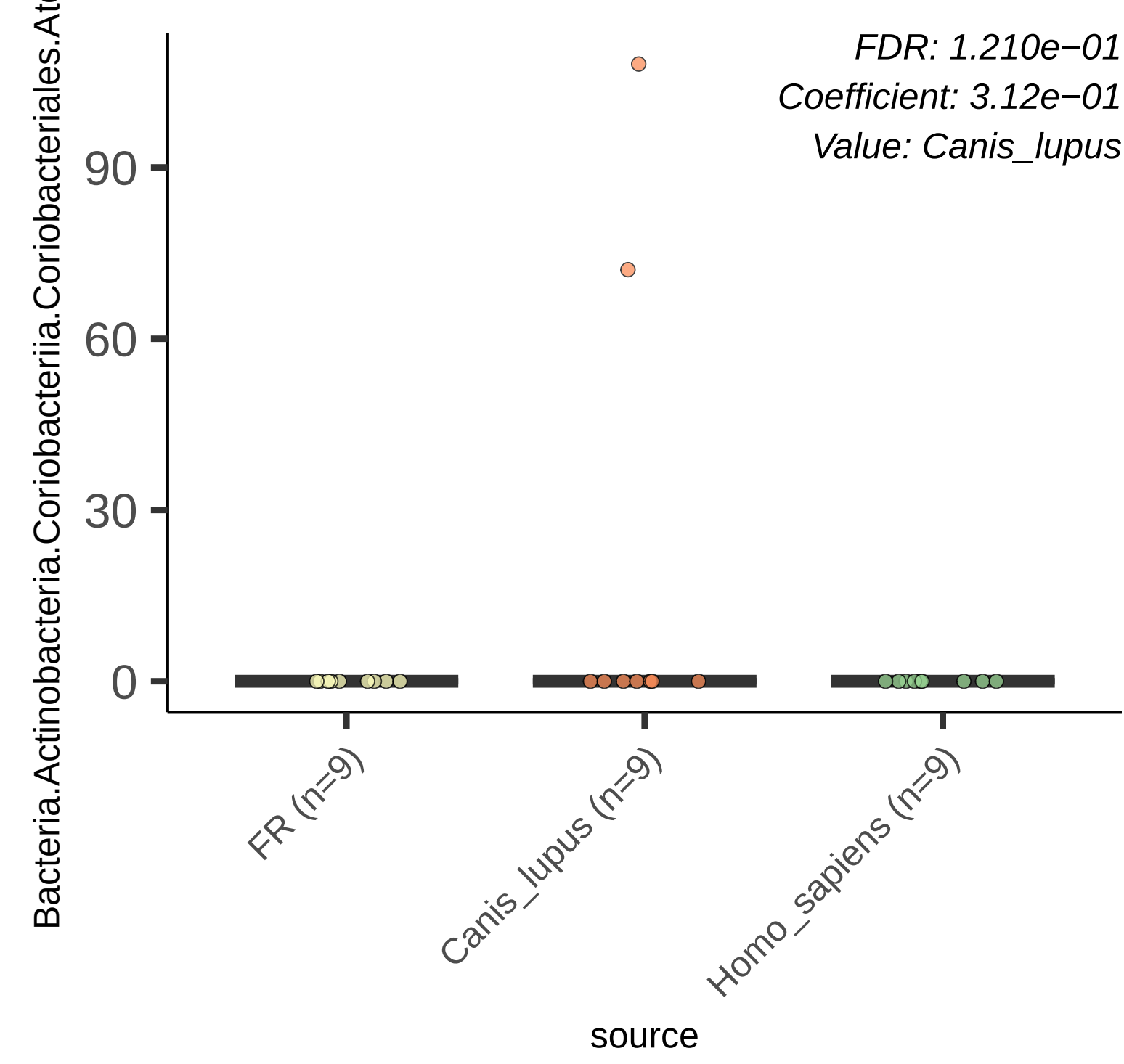

FDR: 1.066e−01
Coefficient: −2.43e−01
Value: Canis_lupus
100
75
50
25
0
Bacteria.Proteobacteria.Alphaproteobacteria.Rhodobacterales.R
FR (n=9)
Canis_lupus (n=9)
Homo_sapiens (n=9)
source

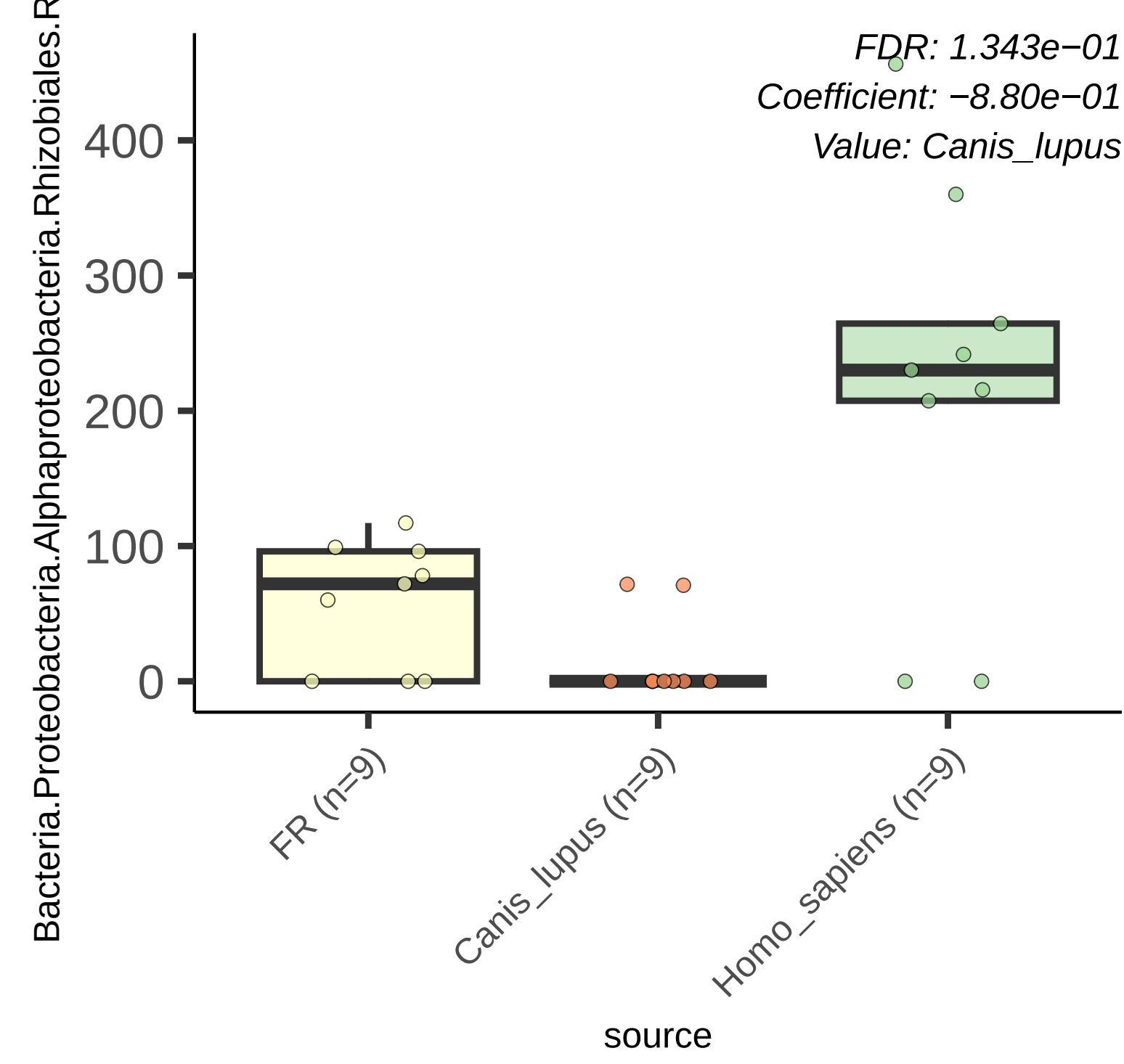

FDR: 1.066e−01
Coefficient: −2.43e−01
Value: Homo_sapiens
Bacteria.Proteobacteria.Alphaproteobacteria.Rhodobacterales.R
100
75
50
25
0
FR (n=9)
Canis_lupus (n=9)
Homo_sapiens (n=9)
source

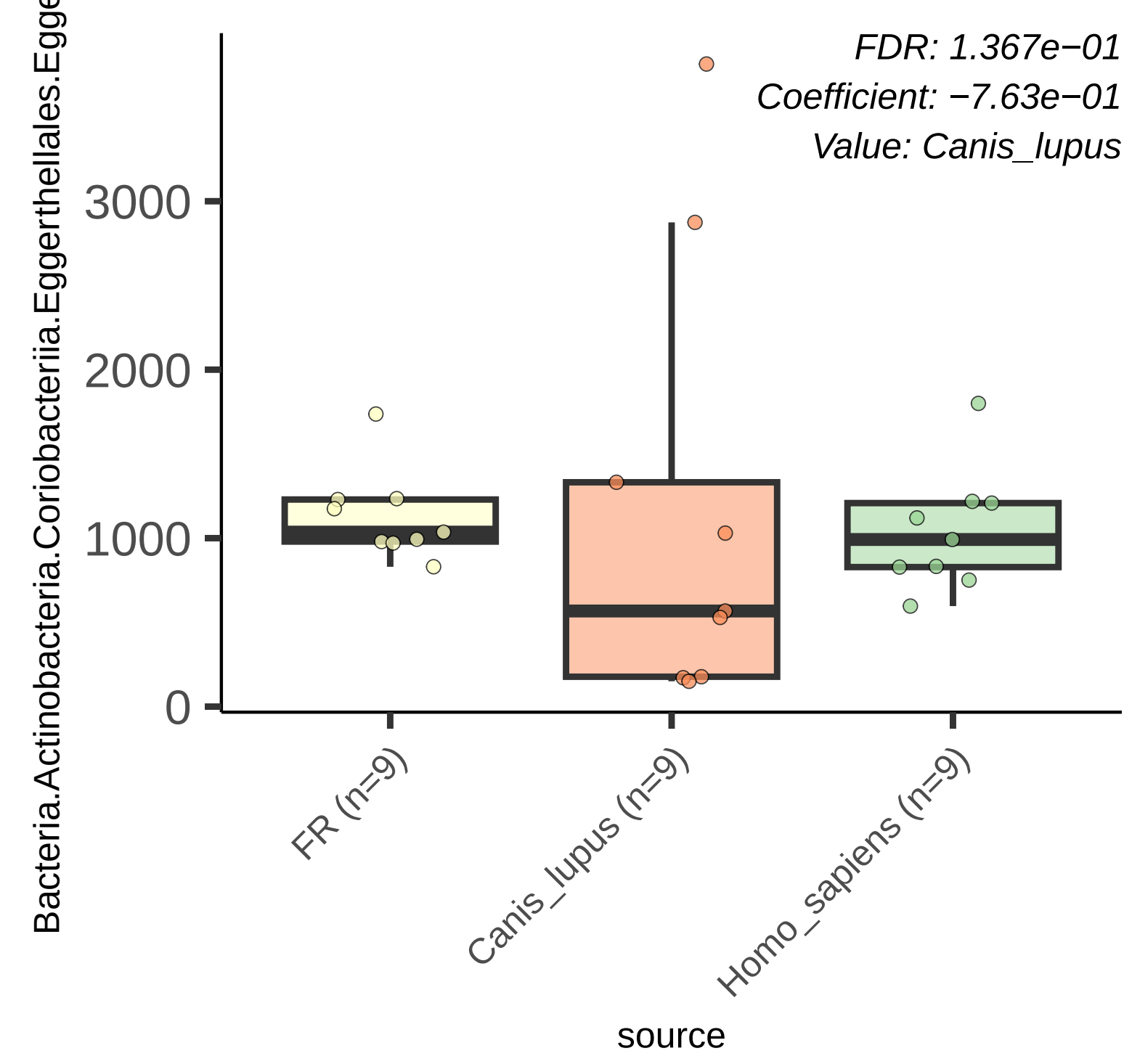

FDR: 1.095e−01
Coefficient: −4.08e−01
Value: Canis_lupus
Bacteria.Firmicutes.Bacilli.Bacillales.Staphylococca
500
400
300
200
100
FR (n=9)
Canis_lupus (n=9)
Homo_sapiens (n=9)
source

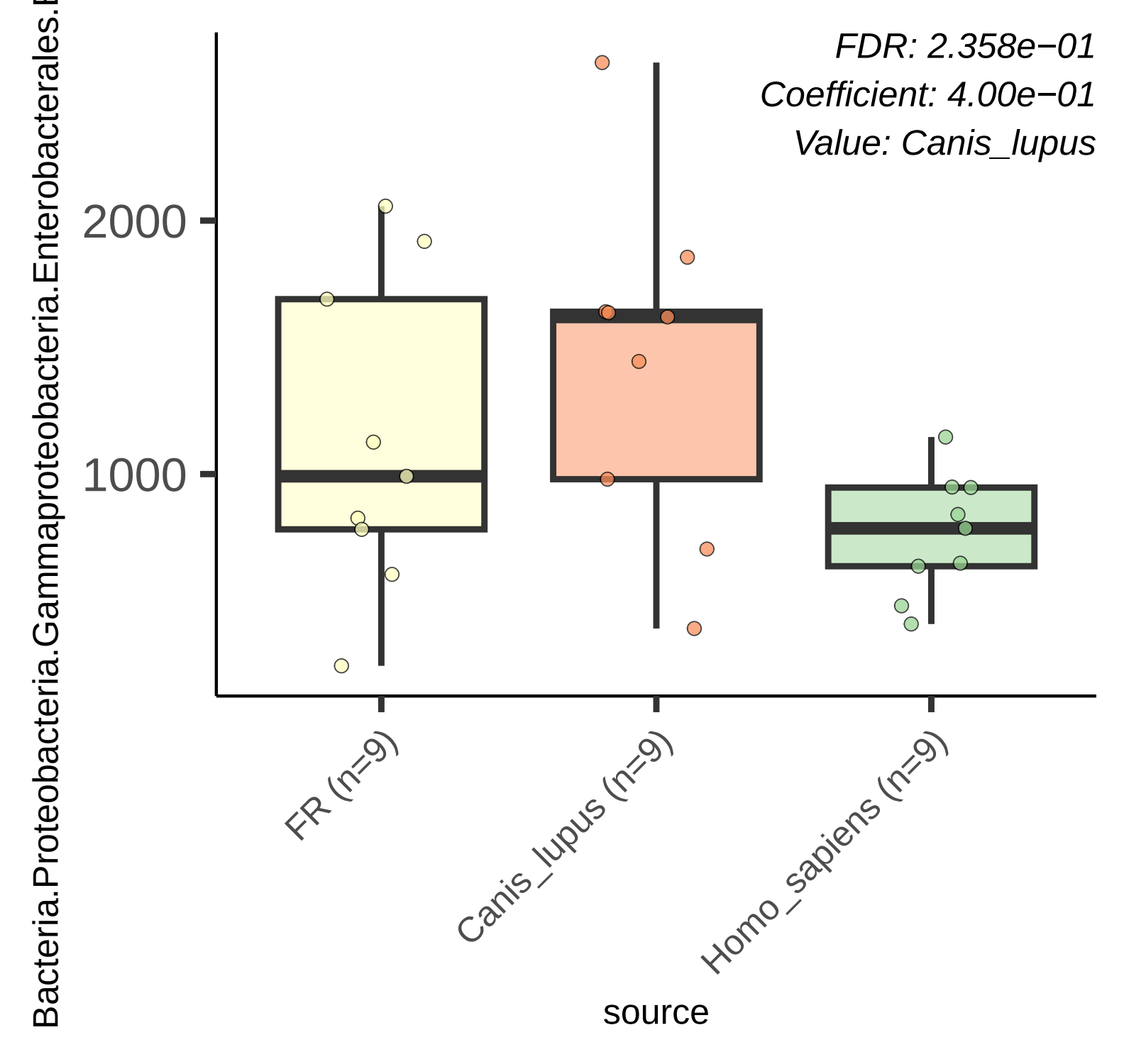

FDR: 1.140e−01
Coefficient: −2.86e−01
Value: Canis_lupus
Bacteria.Bacteroidetes.Bacteroidia.Bacteroidales.Porphyro
30
20
10
0
FR (n=9)
Canis_lupus (n=9)
Homo_sapiens (n=9)
source

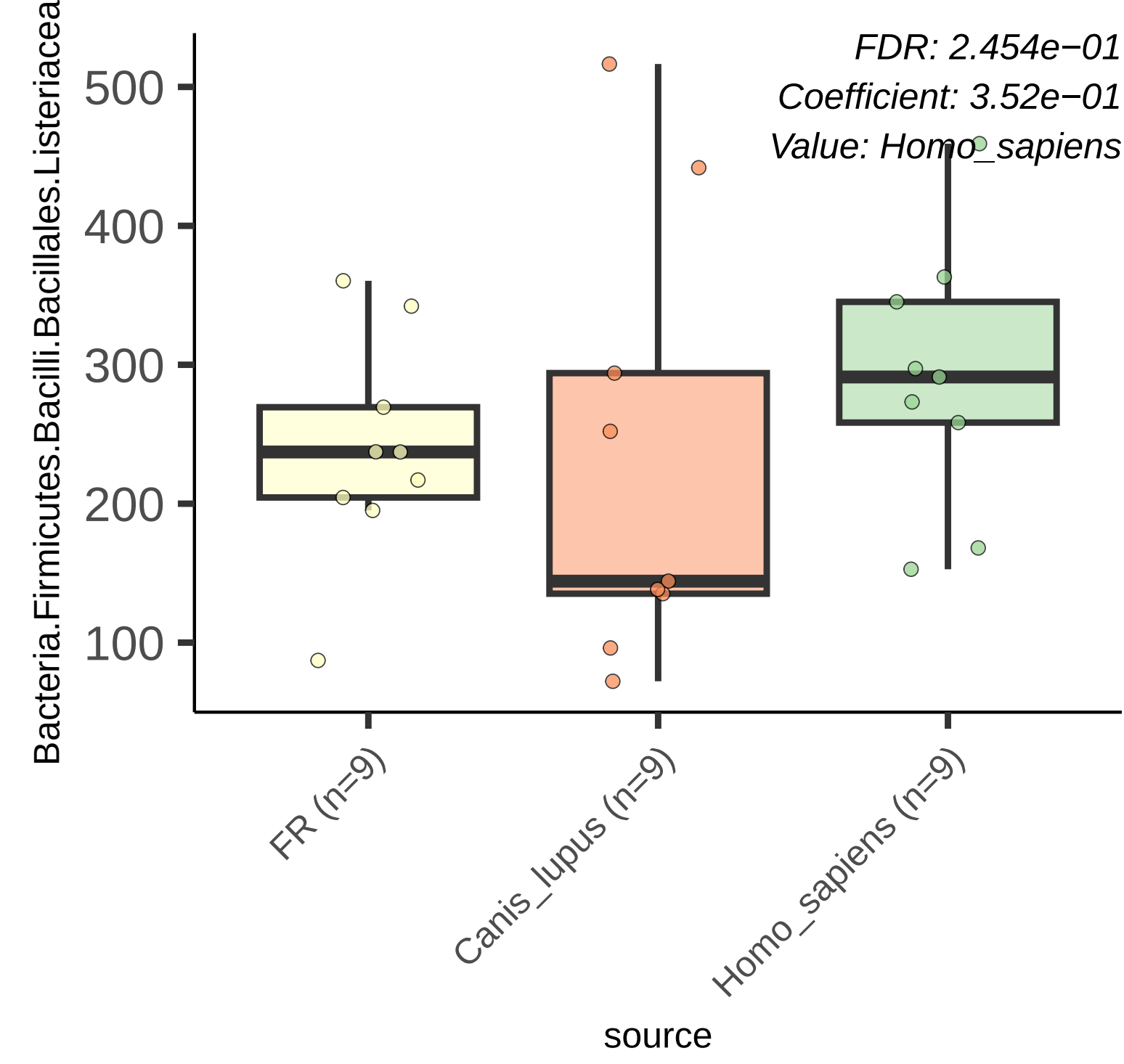

FDR: 1.140e−01
Coefficient: −2.86e−01
Value: Homo_sapiens
Bacteria.Bacteroidetes.Bacteroidia.Bacteroidales.Porphyro
30
20
10
0
FR (n=9)
Canis_lupus (n=9)
Homo_sapiens (n=9)
source


FDR: 1.210e−01
Coefficient: 3.12e−01
Value: Canis_lupus
Bacteria.Actinobacteria.Coriobacteriia.Coriobacteriales.Atopobiaceae.Atopobium
90
60
30
0
FR (n=9)
Canis_lupus (n=9)
Homo_sapiens (n=9)
source


FDR: 1.343e−01
Coefficient: −8.80e−01
Value: Canis_lupus
Bacteria.Proteobacteria.Alphaproteobacteria.Rhizobiales.R
400
300
200
100
0
FR (n=9)
Canis_lupus (n=9)
Homo_sapiens (n=9)
source


FDR: 1.367e−01
Coefficient: −7.63e−01
Value: Canis_lupus
Bacteria.Actinobacteria.Coriobacteriia.Eggerthellales.Egge
3000
2000
1000
0
FR (n=9)
Canis_lupus (n=9)
Homo_sapiens (n=9)
source


FDR: 2.358e−01
Coefficient: 4.00e−01
Value: Canis_lupus
2000
1000
Bacteria.Proteobacteria.Gammaproteobacteria.Enterobacterales.E
FR (n=9)
Canis_lupus (n=9)
Homo_sapiens (n=9)
source


FDR: 2.454e−01
Coefficient: 3.52e−01
Value: Homo_sapiens
Bacteria.Firmicutes.Bacilli.Bacillales.Listeriacea
500
400
300
200
100
FR (n=9)
Canis_lupus (n=9)
Homo_sapiens (n=9)
source