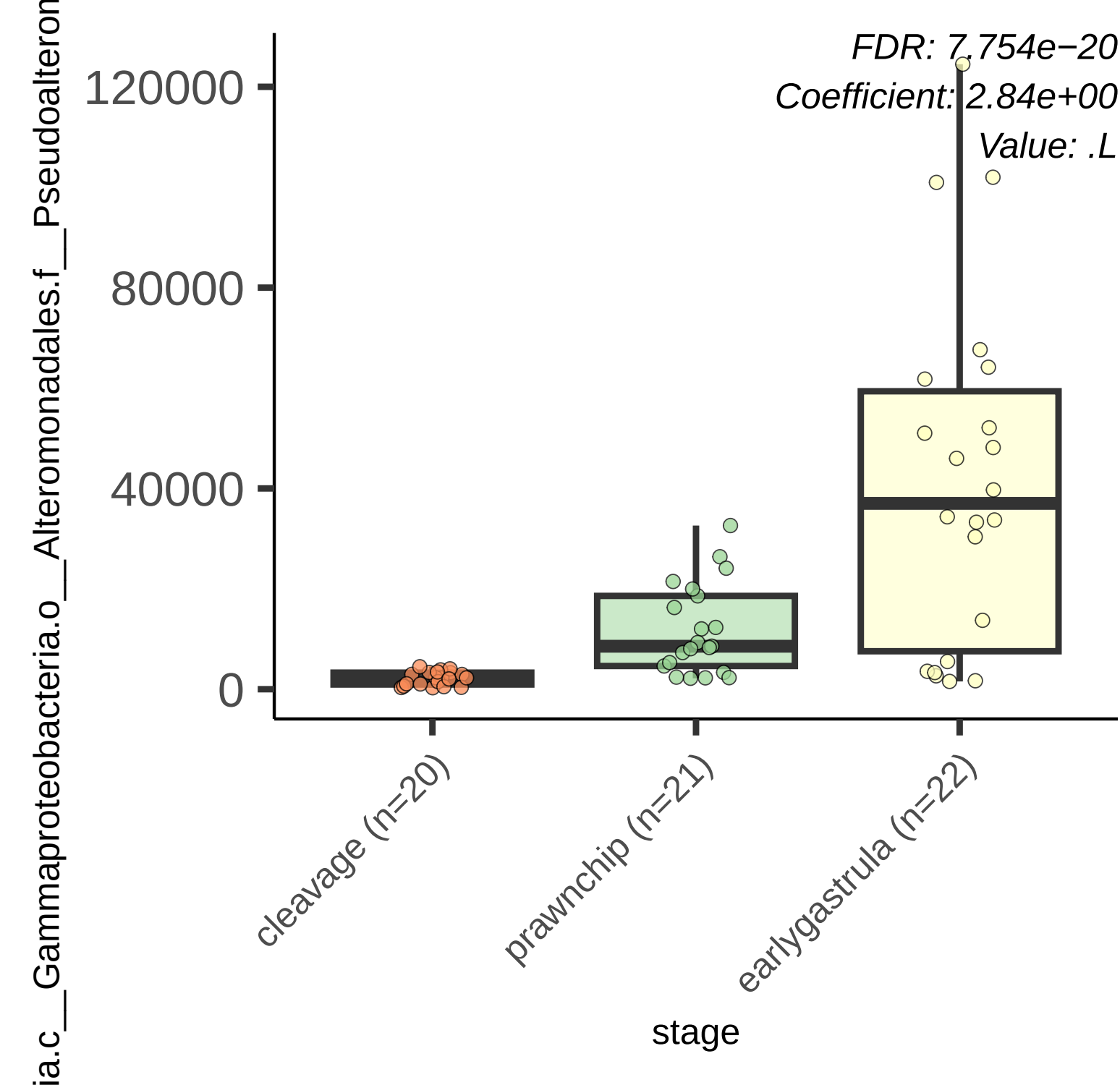
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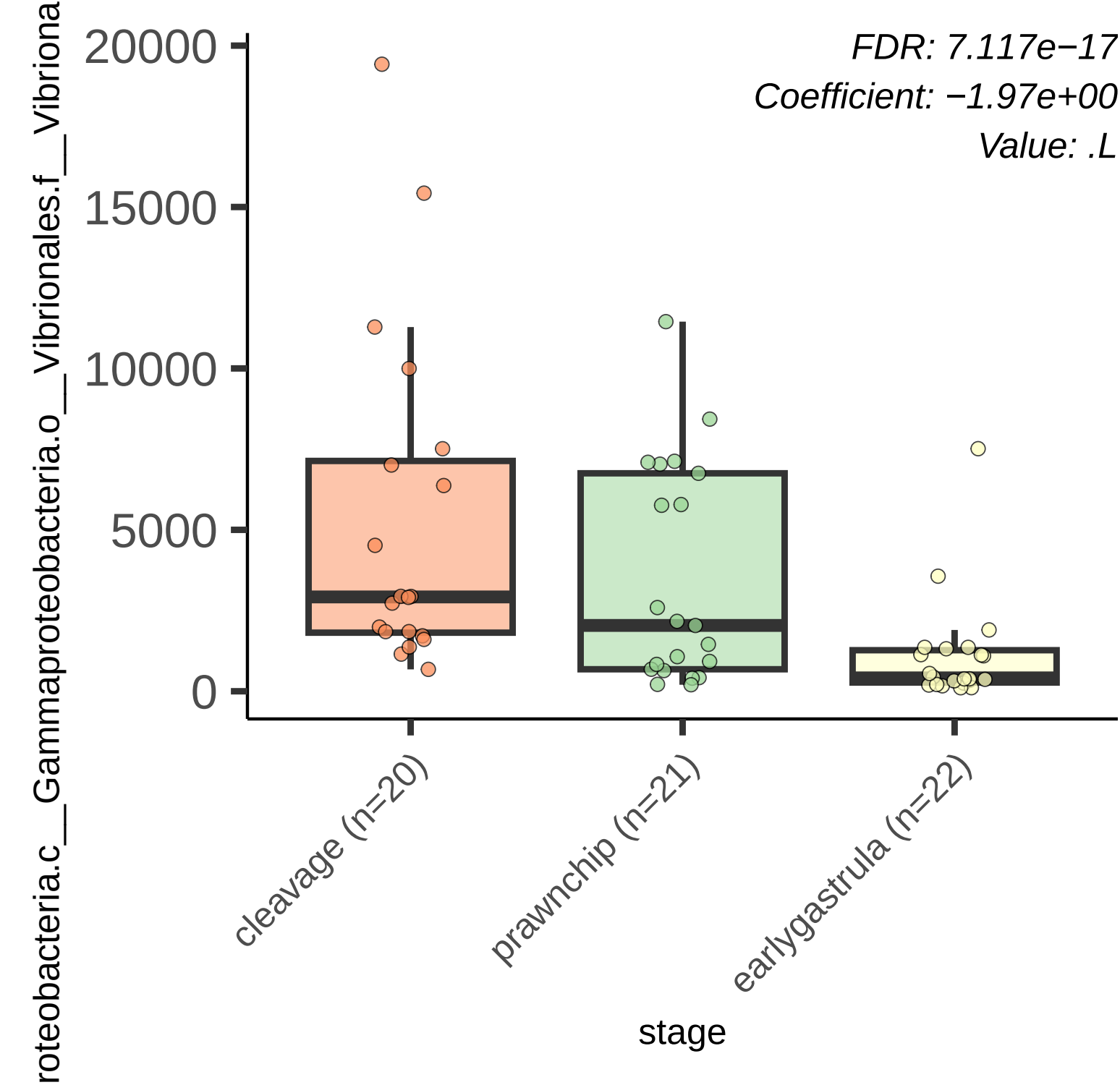
120000
80000
40000
0
cleavage (n=20)
prawnchip (n=21)
earlygastrula (n=22)
stage
ia.c__Gammaproteobacteria.o__Alteromonadales.f__Pseudoalteron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15000
10000
5000
0
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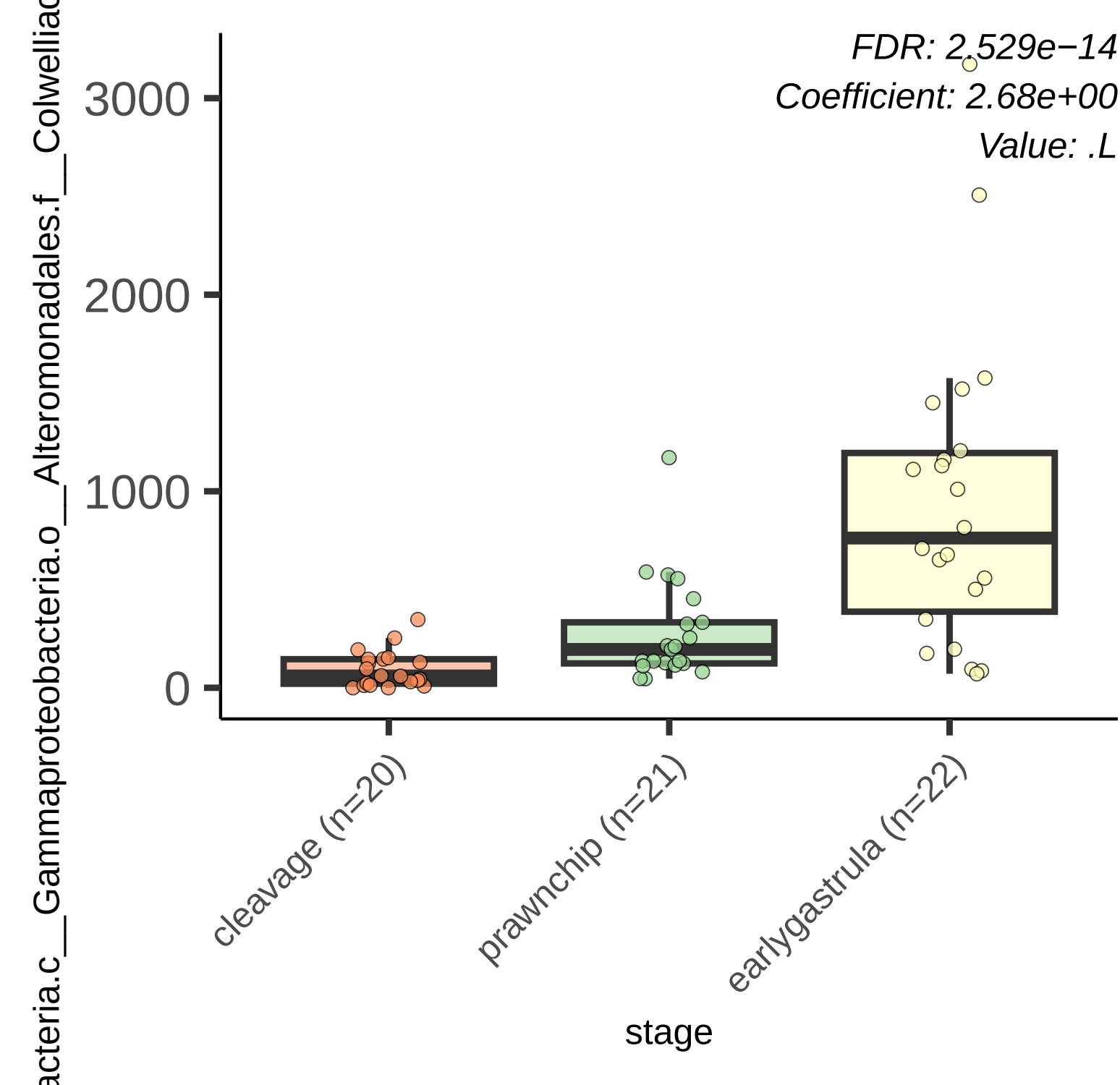
prawnchip (n=21)
earlygastrula (n=22)
stage
roteobacteria.c__Gammaproteobacteria.o__Vibrionales.f__Vibriona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2000
1000
0
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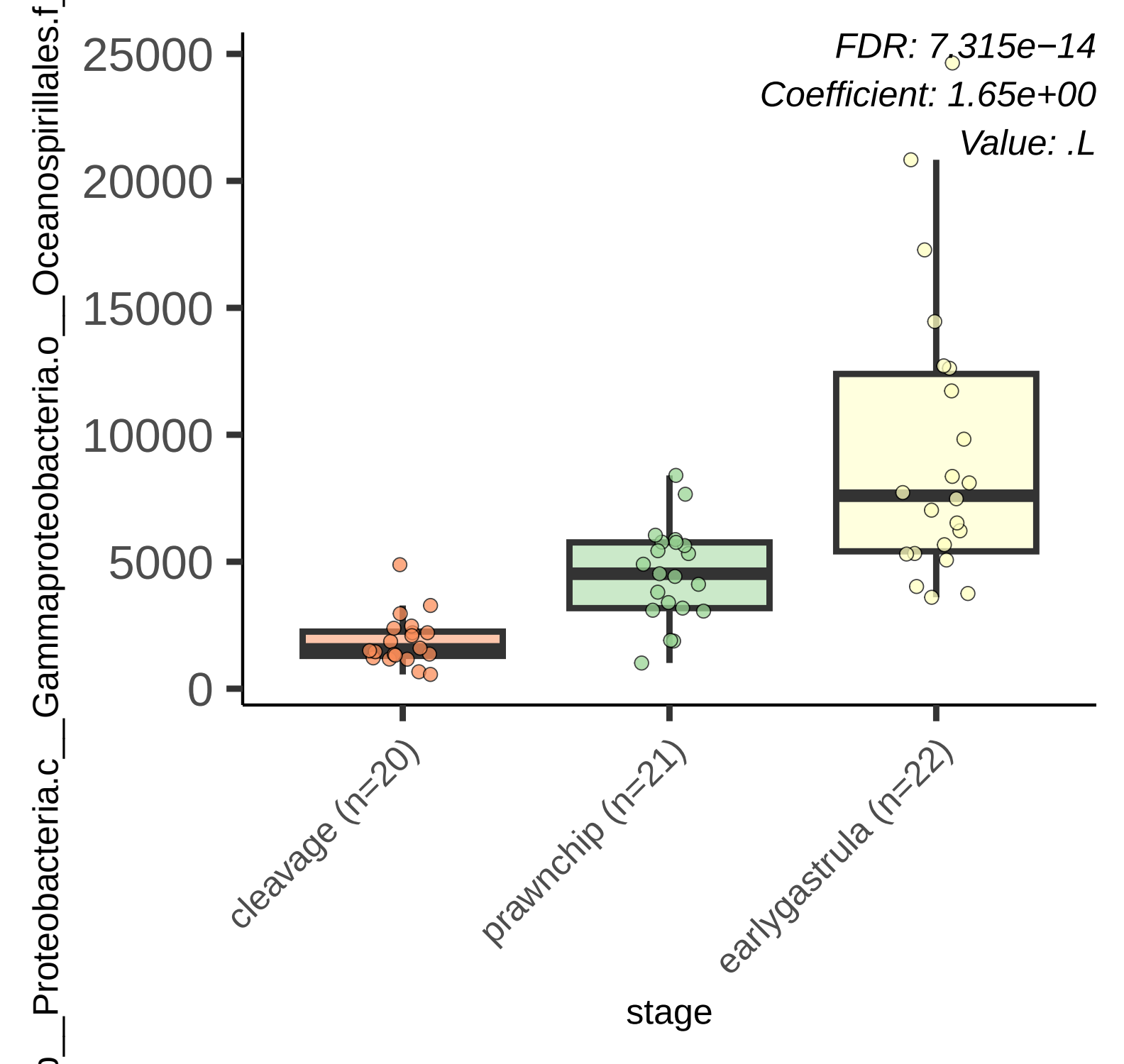
prawnchip (n=21)
earlygastrula (n=22)
stage
acteria.c__Gammaproteobacteria.o__Alteromonadales.f__Colwelliac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20000
15000
10000
5000
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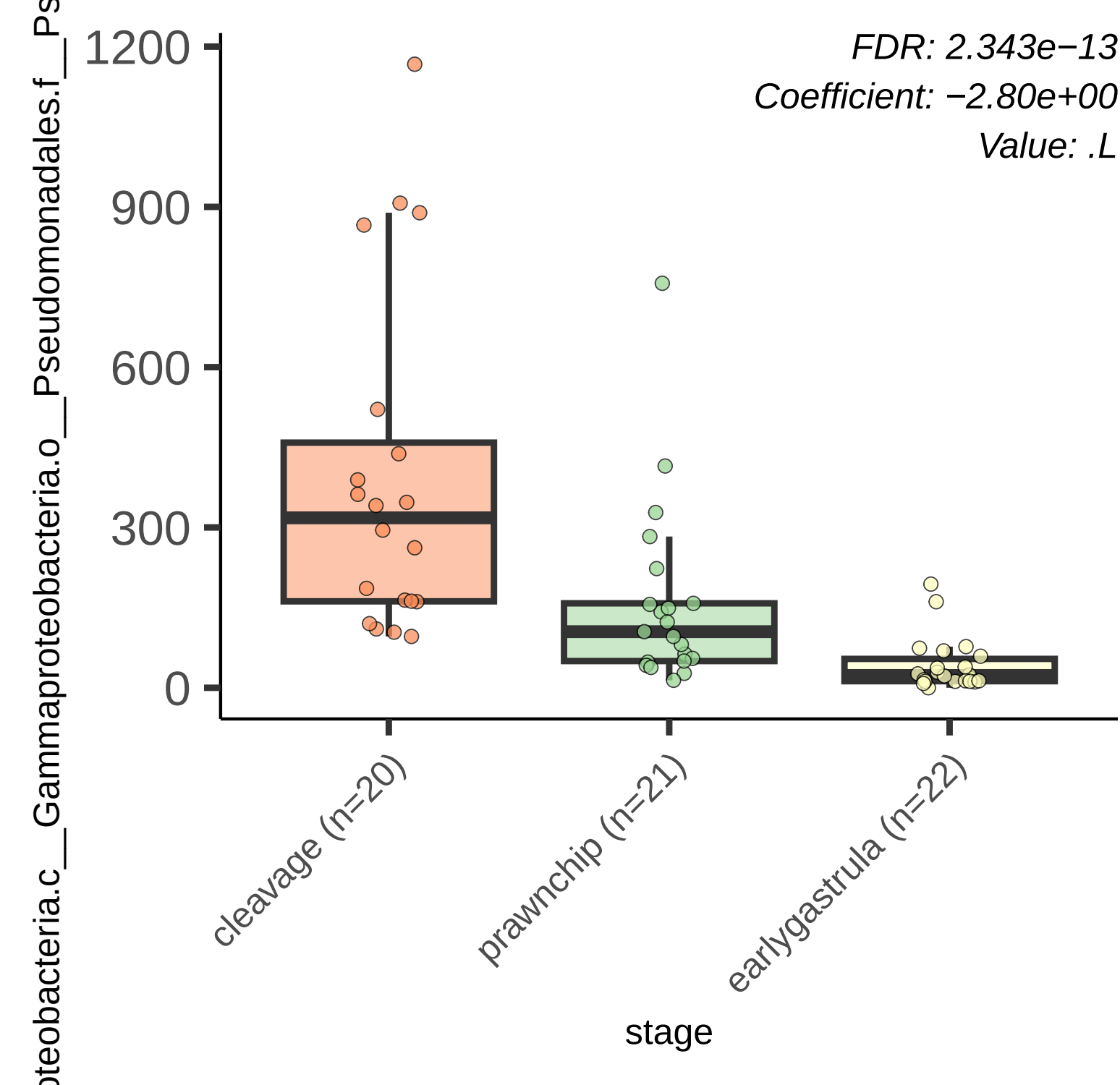
cleavage (n=20)
prawnchip (n=21)
earlygastrula (n=22)
stage
o__Proteobacteria.c__Gammaproteobacteria.o__Oceanospirillales.f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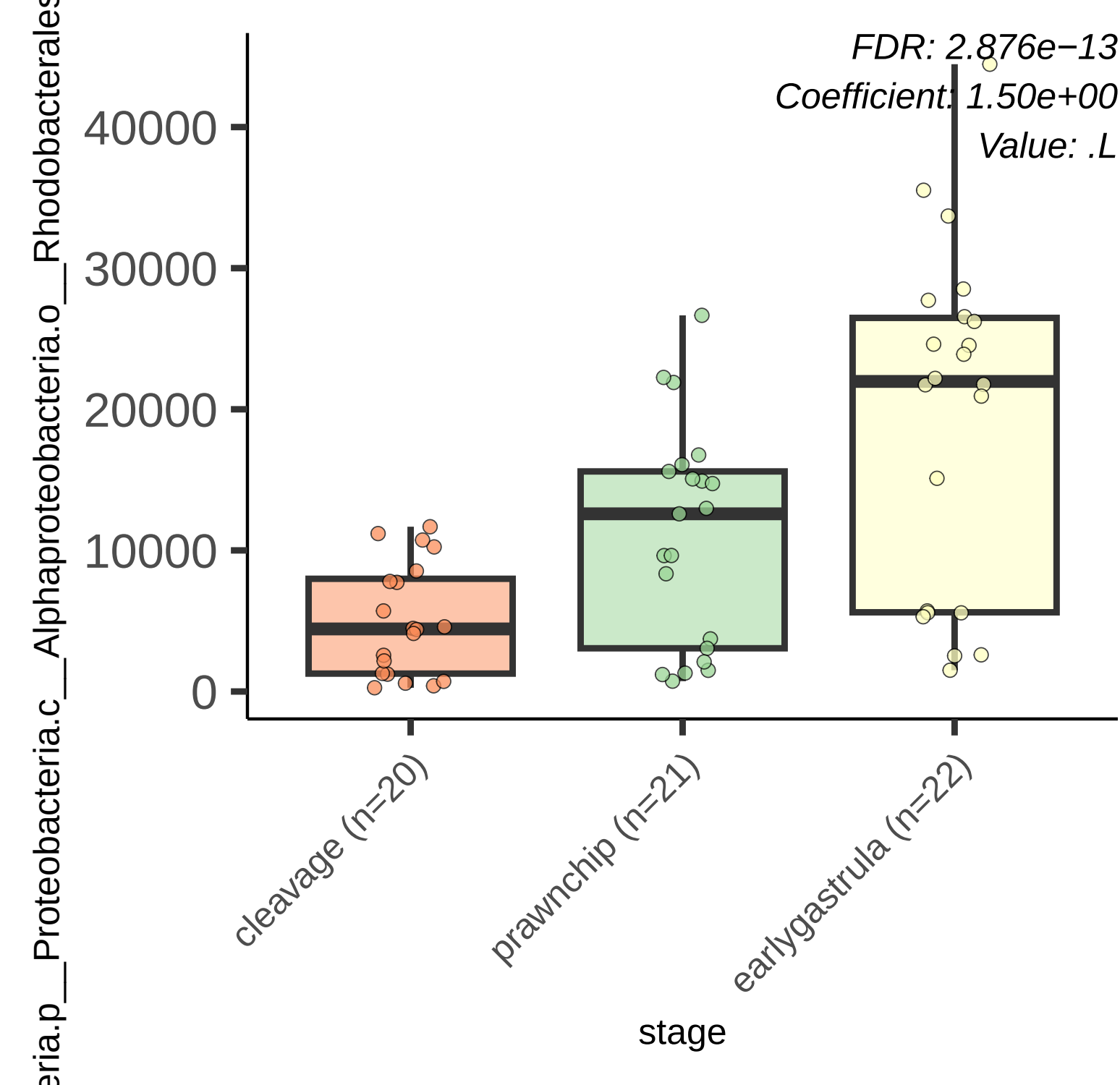
900
600
300
0
cleavage (n=20)
prawnchip (n=21)
earlygastrula (n=22)
stage
teobacteria.c__Gammaproteobacteria.o__Pseudomonadales.f__Ps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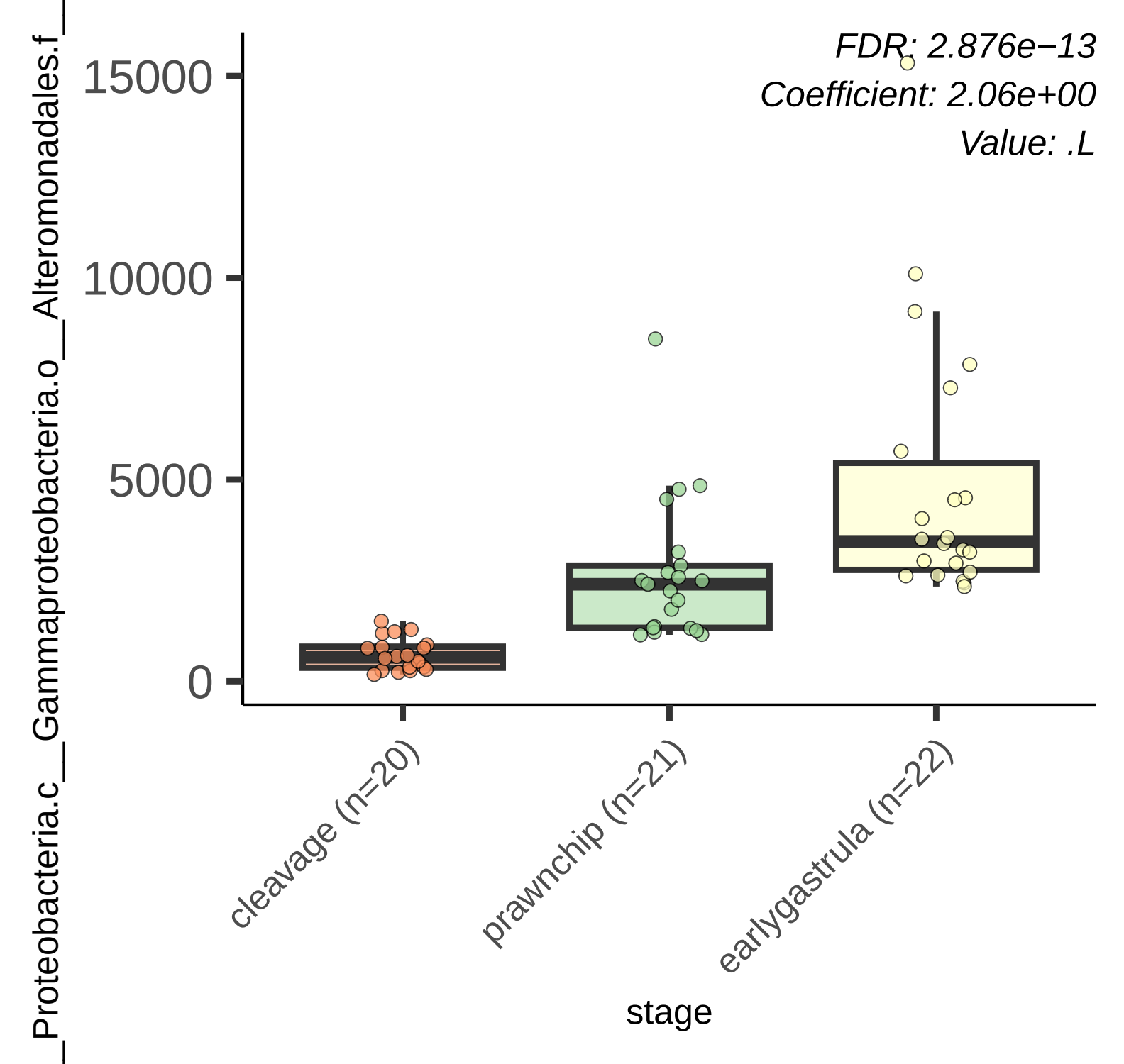
30000
20000
10000
0
cleavage (n=20)
prawnchip (n=21)
earlygastrula (n=22)
stage
eria.p__Proteobacteria.c__Alphaproteobacteria.o__Rhodobacterales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Coefficient: 2.06e+00
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10000
5000
0
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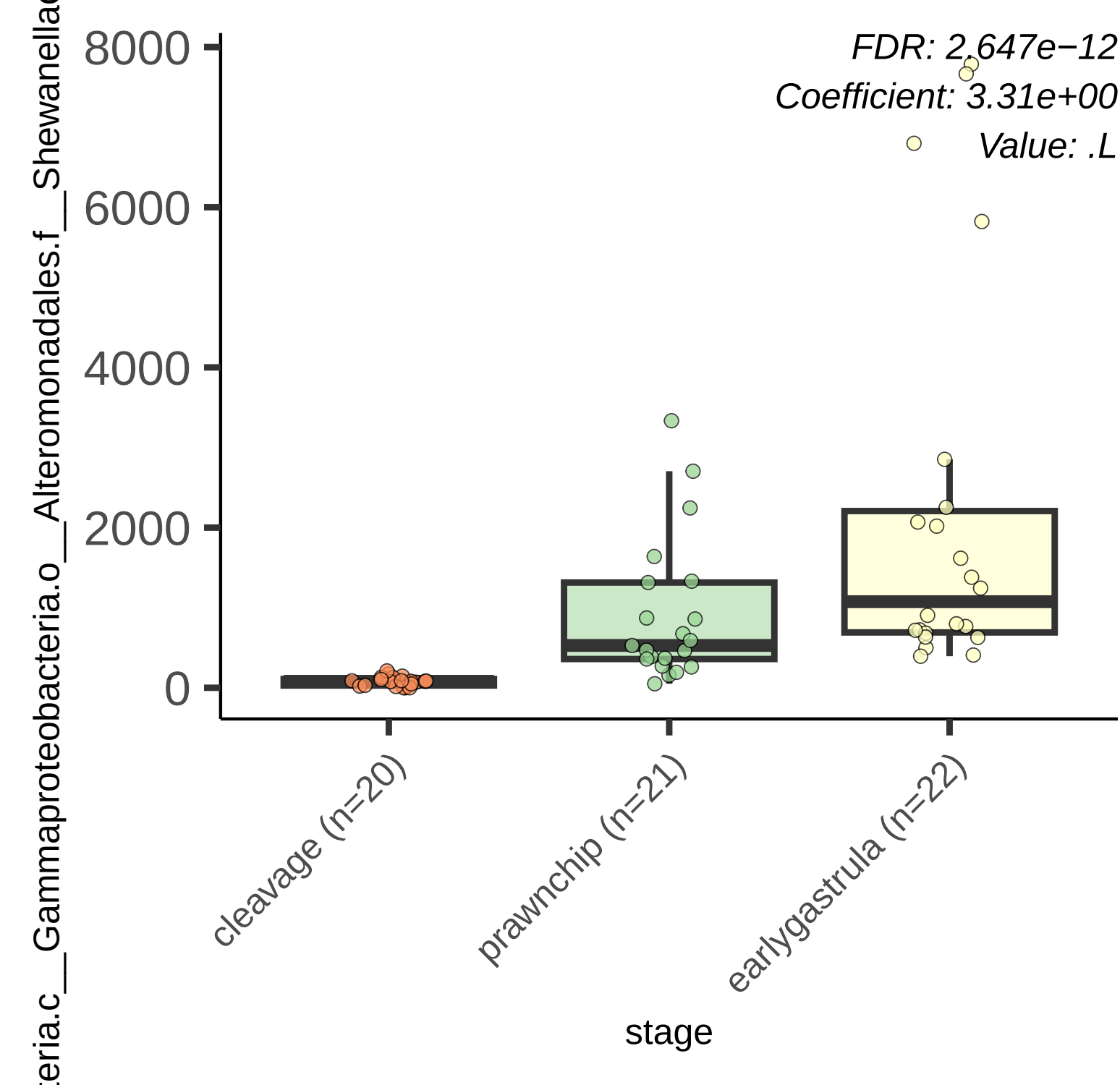
prawnchip (n=21)
earlygastrula (n=22)
stage
__Proteobacteria.c__Gammaproteobacteria.o__Alteromonadales.f__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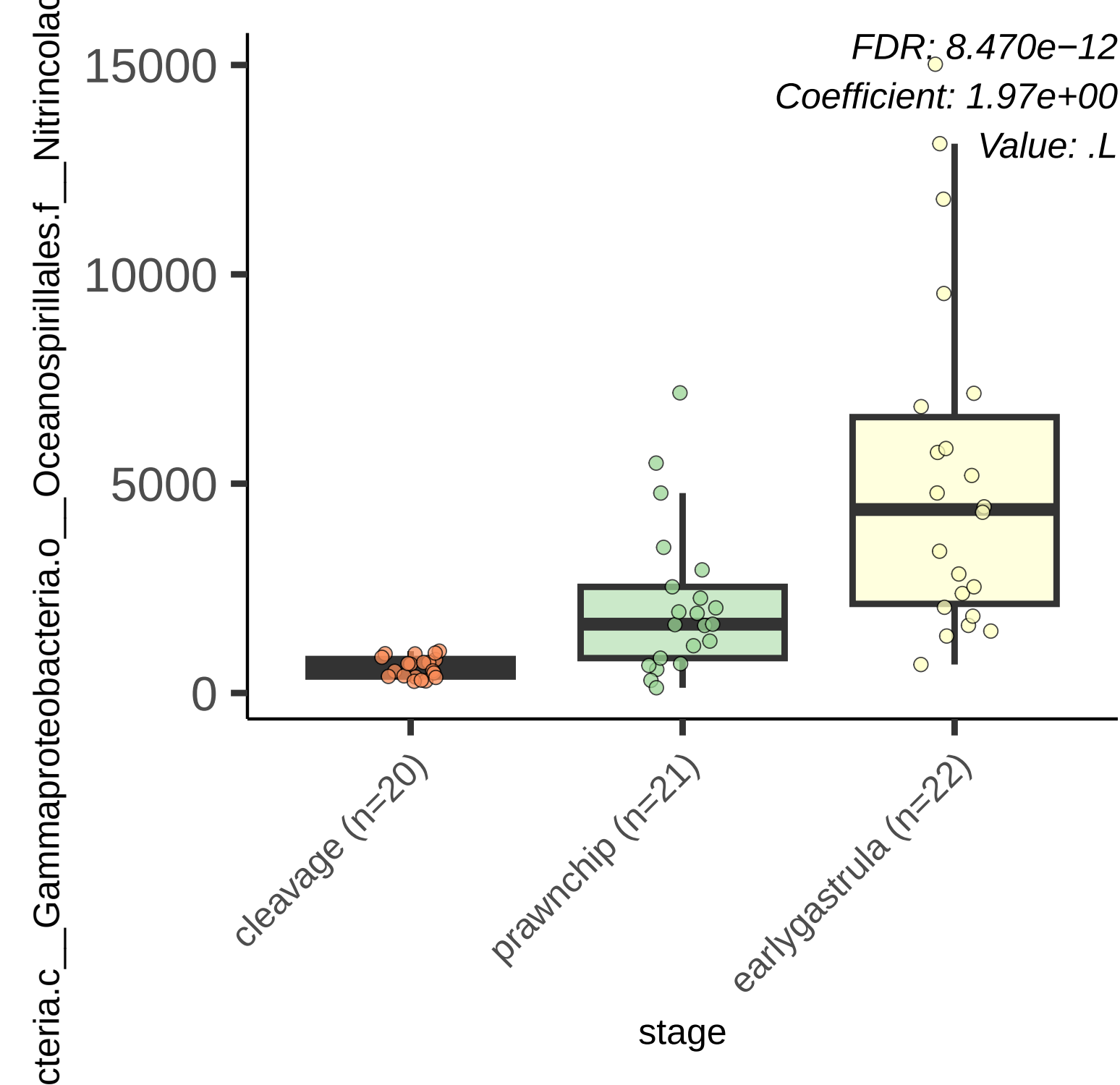
8000
6000
4000
2000
0
cleavage (n=20)
prawnchip (n=21)
earlygastrula (n=22)
stage
teria.c__Gammaproteobacteria.o__Alteromonadales.f__Shewanellac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Coefficient: 1.97e+00
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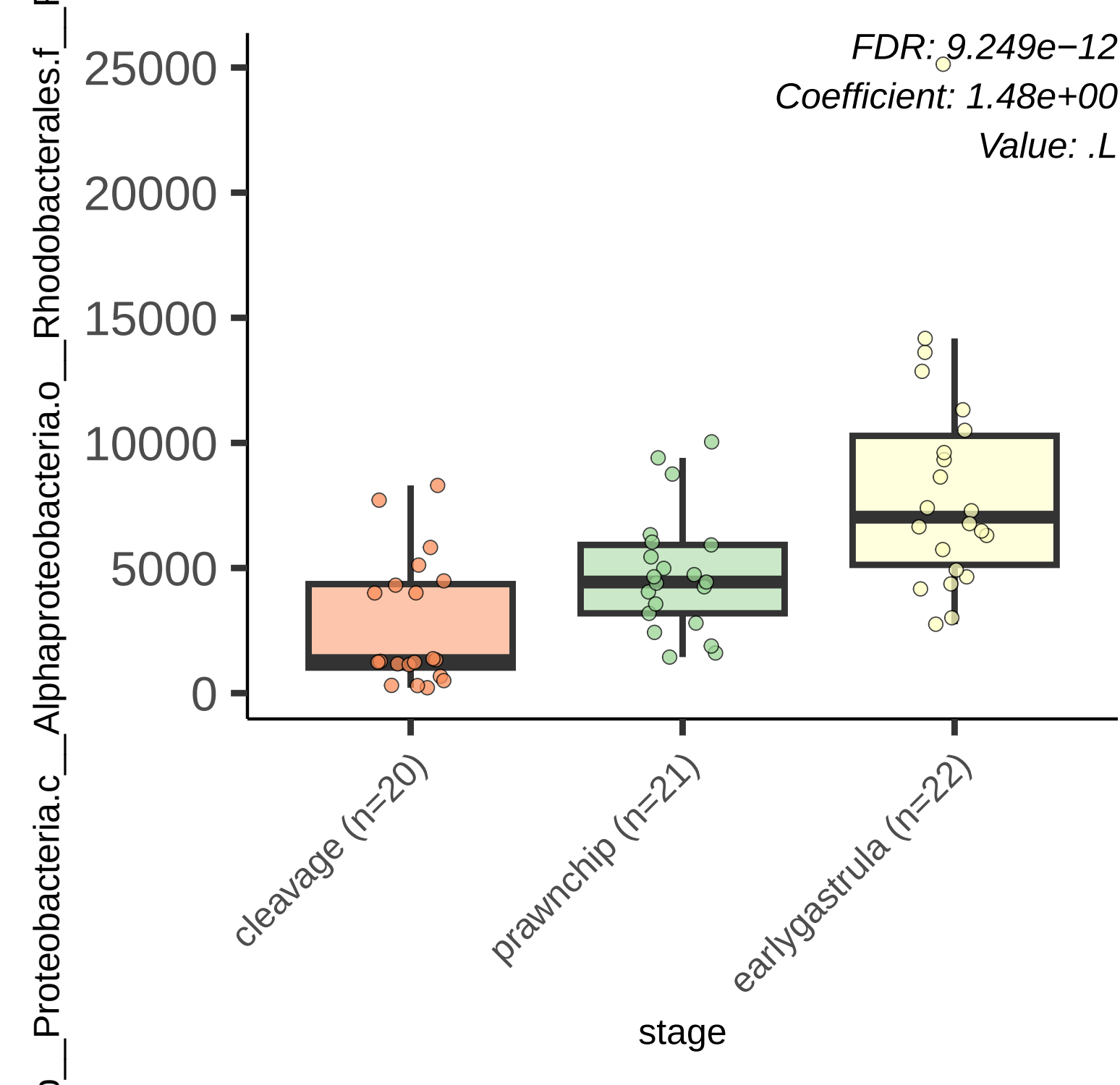
10000
5000
0
cleavage (n=20)
prawnchip (n=21)
earlygastrula (n=22)
stage
cteria.c__Gammaproteobacteria.o__Oceanospirillales.f__Nitrincolac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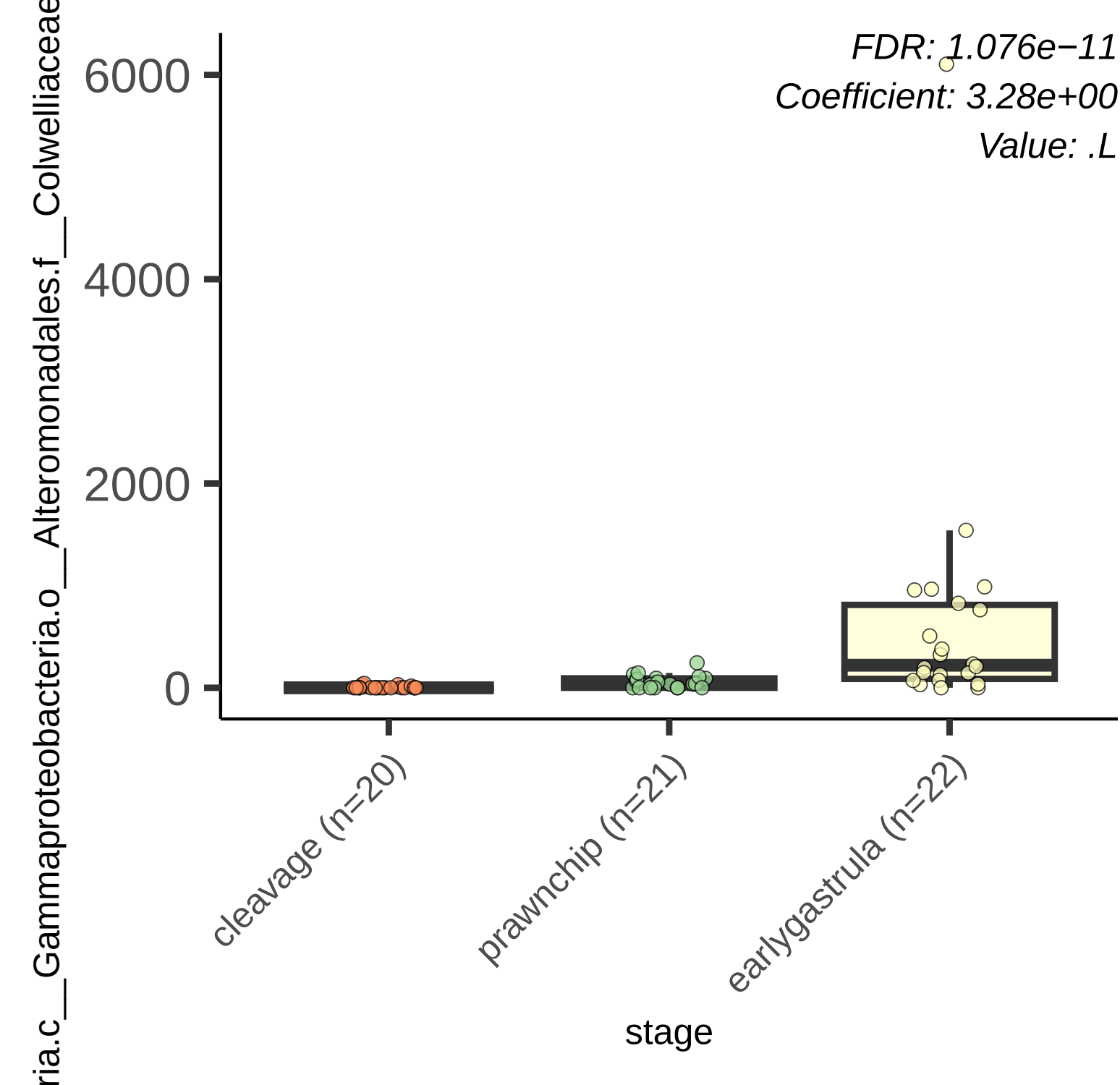
20000
15000
10000
5000
0
cleavage (n=20)
prawnchip (n=21)
earlygastrula (n=22)
stage
o__Proteobacteria.c__Alphaproteobacteria.o__Rhodobacterales.f__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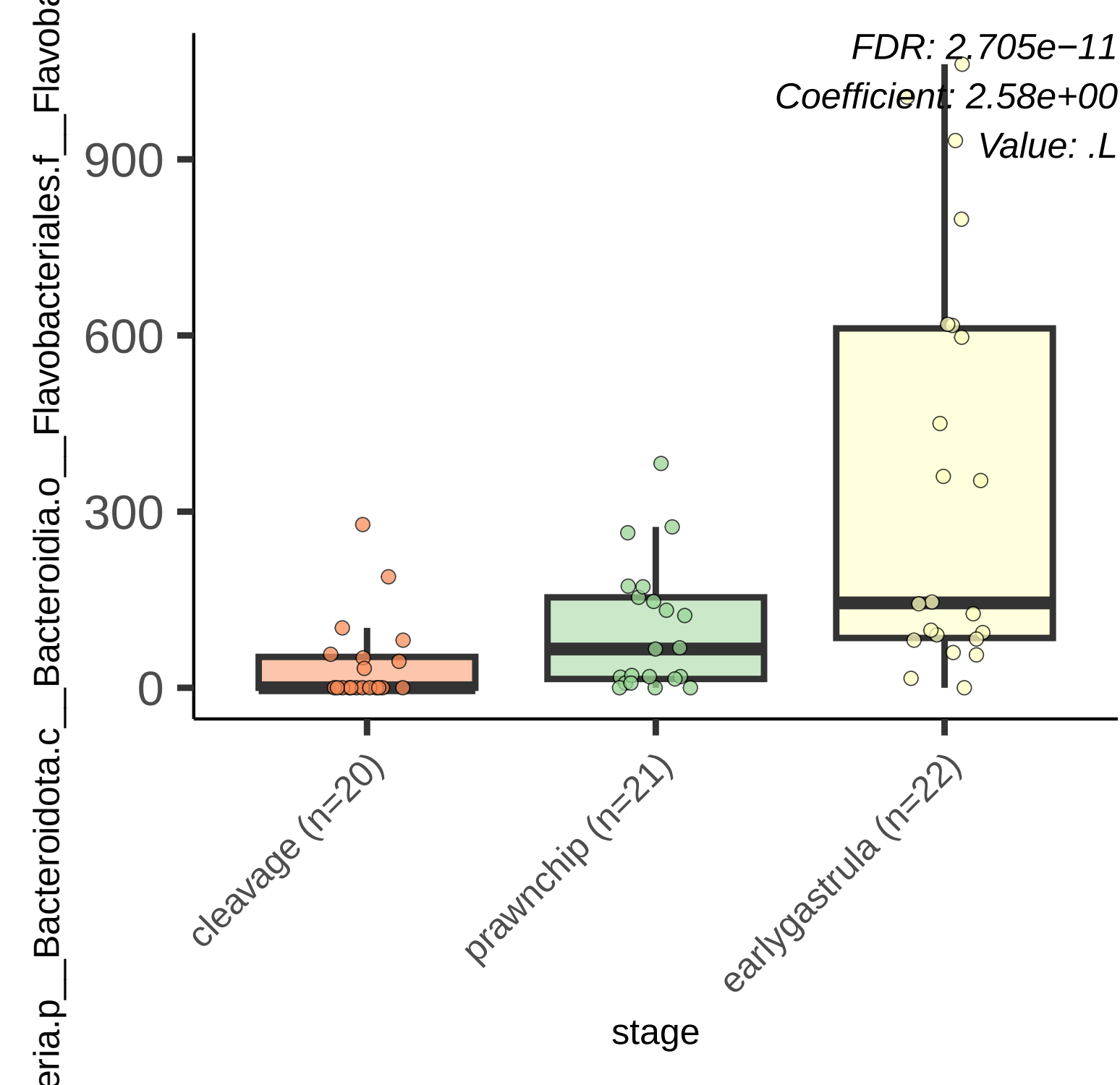
4000
2000
0
cleavage (n=20)
prawnchip (n=21)
earlygastrula (n=22)
stage
ria.c__Gammaproteobacteria.o__Alteromonadales.f__Colwelliaceae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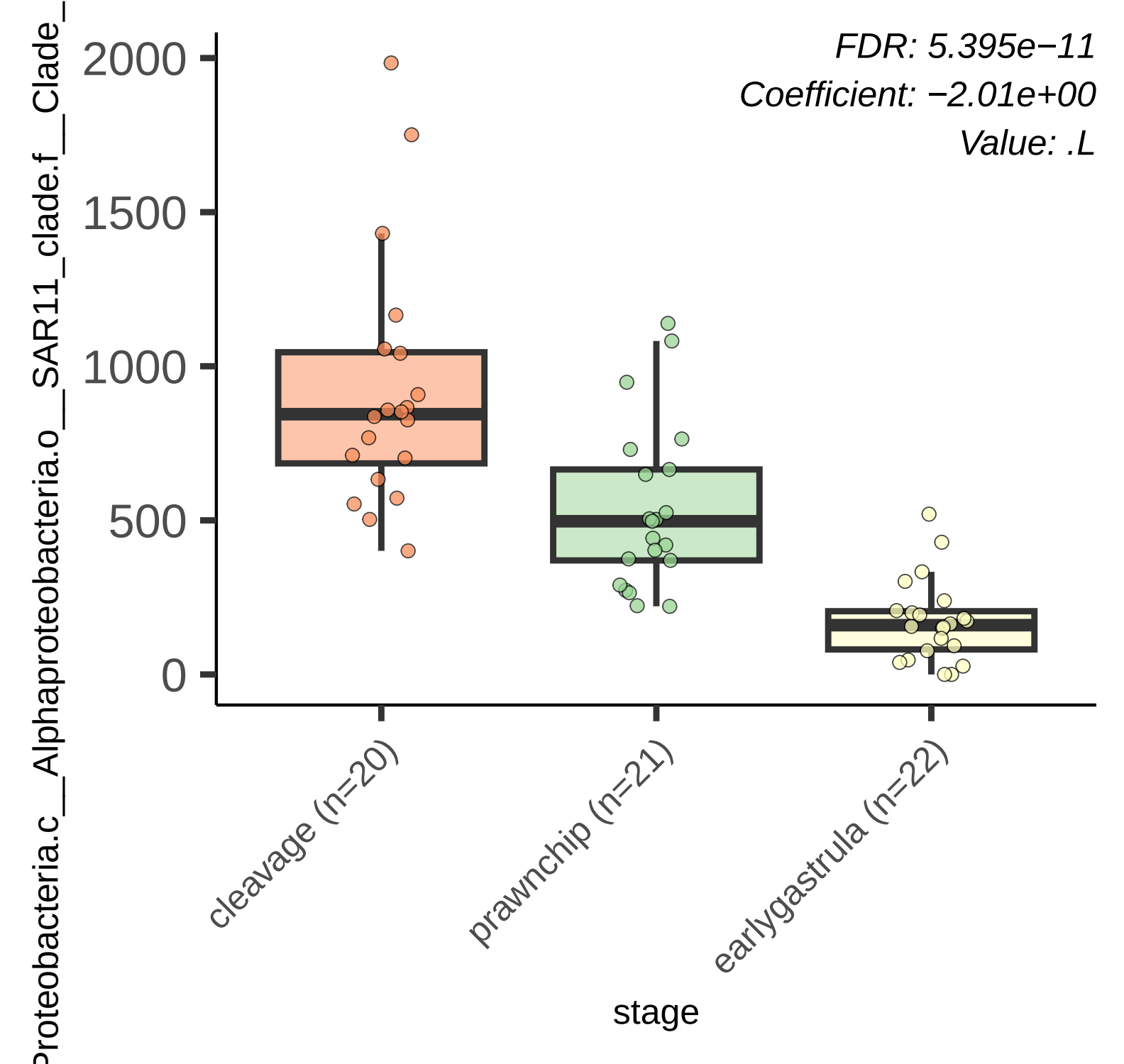
Coefficient: 2.58e+00
Value: .L
900
600
300
0
cleavage (n=20)
prawnchip (n=21)
earlygastrula (n=22)
stage
eria.p__Bacteroidota.c__Bacteroidia.o__Flavobacteriales.f__Flavoba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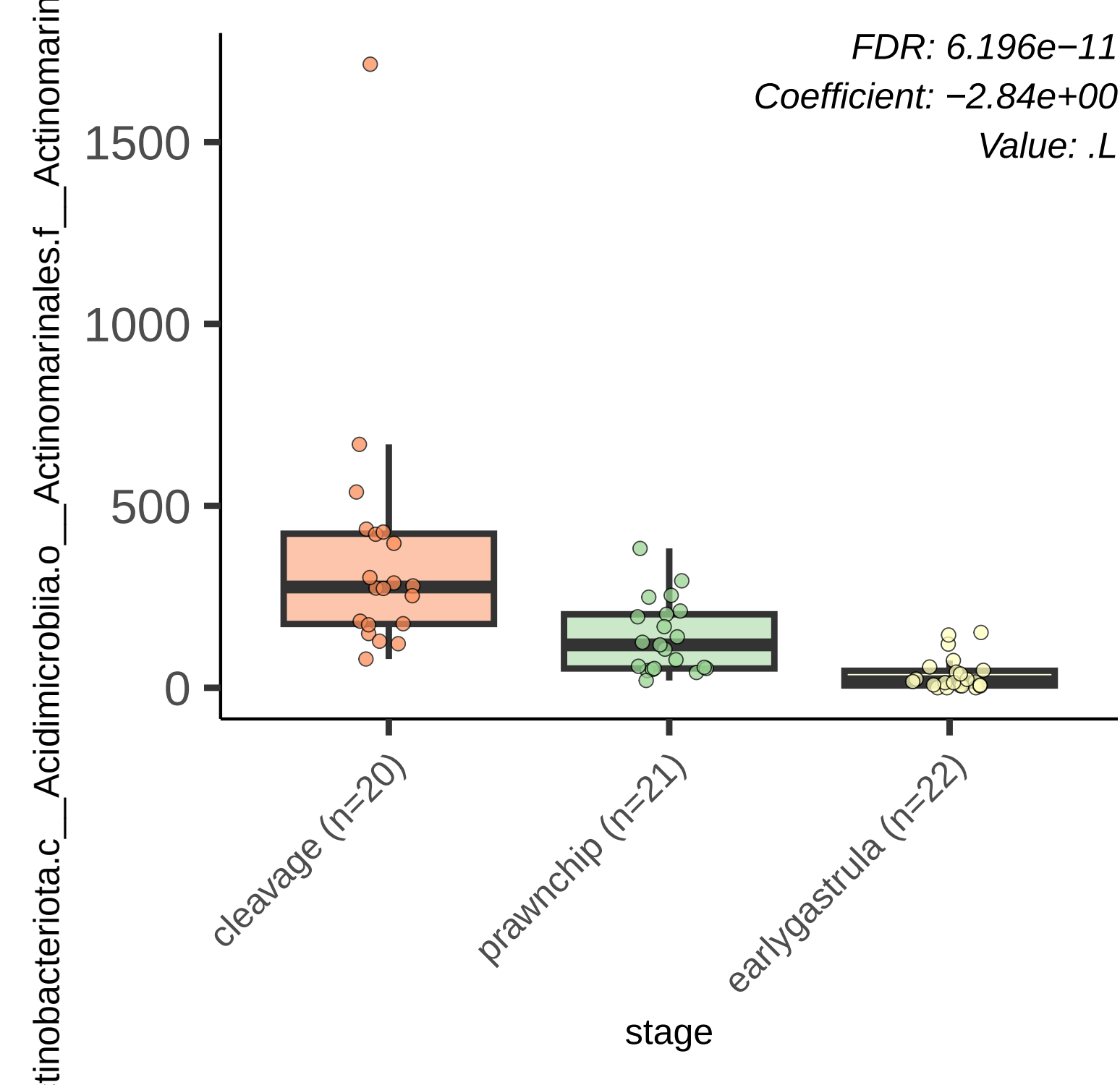
Coefficient: −2.01e+00
Value: .L
2000
1500
1000
500
0
cleavage (n=20)
prawnchip (n=21)
earlygastrula (n=22)
stage
Proteobacteria.c__Alphaproteobacteria.o__SAR11_clade.f__Clade_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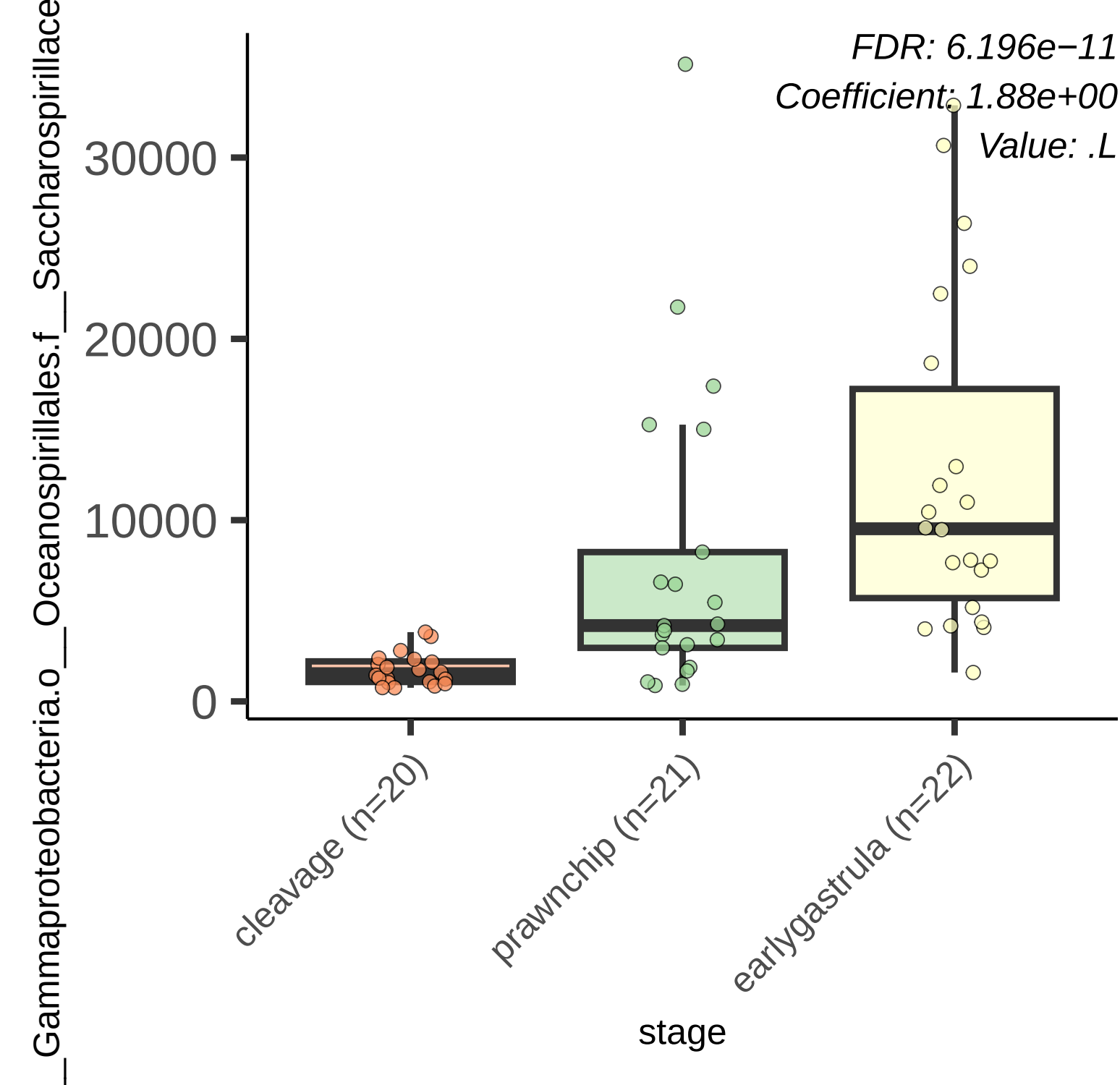
Coefficient: −2.84e+00
Value: .L
1500
1000
500
0
cleavage (n=20)
prawnchip (n=21)
earlygastrula (n=22)
stage
ctinobacteriota.c__Acidimicrobiia.o__Actinomarinales.f__Actinomarin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20000
10000
0
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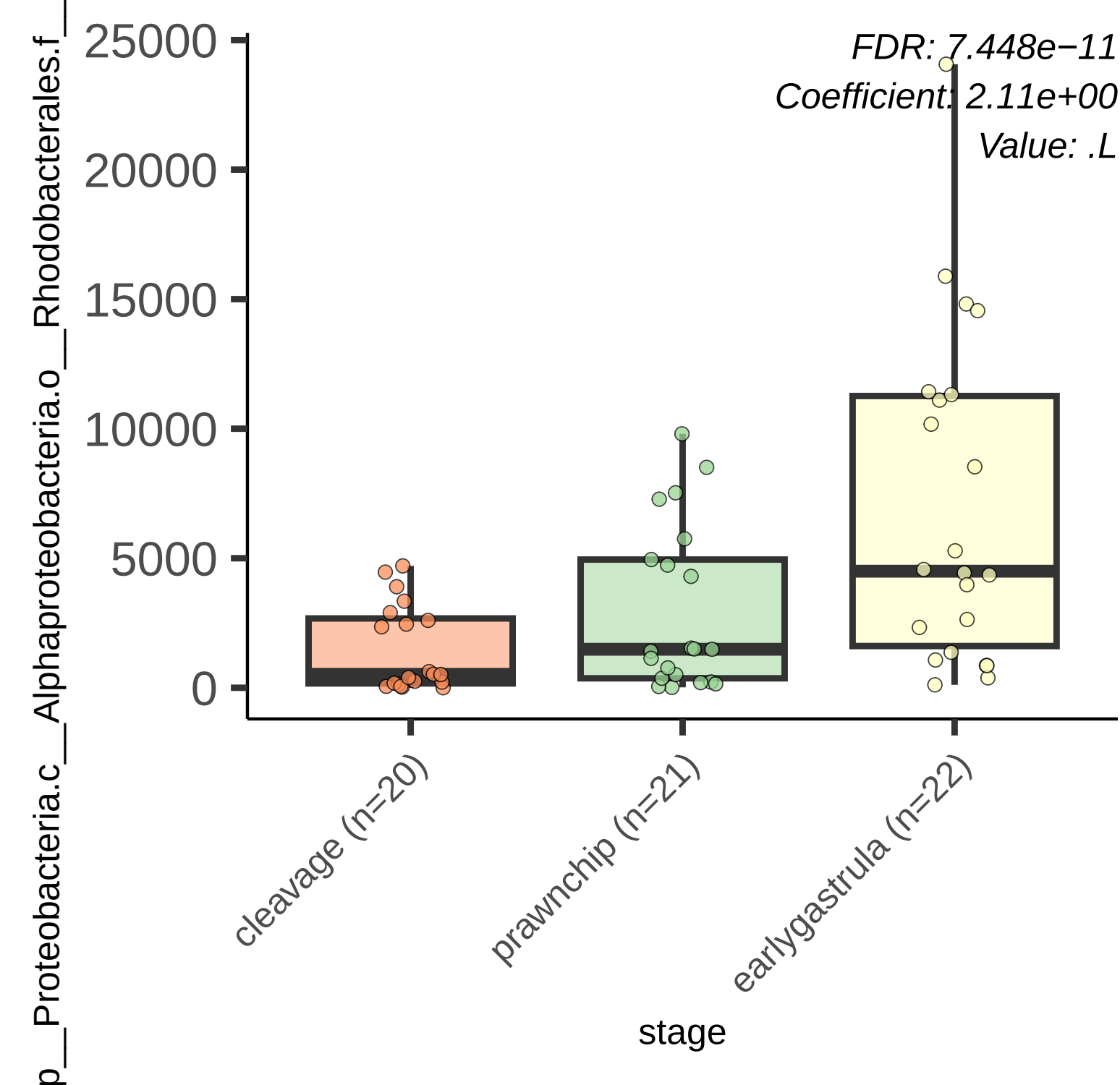
prawnchip (n=21)
earlygastrula (n=22)
stage
__Gammaproteobacteria.o__Oceanospirillales.f__Saccharospirillace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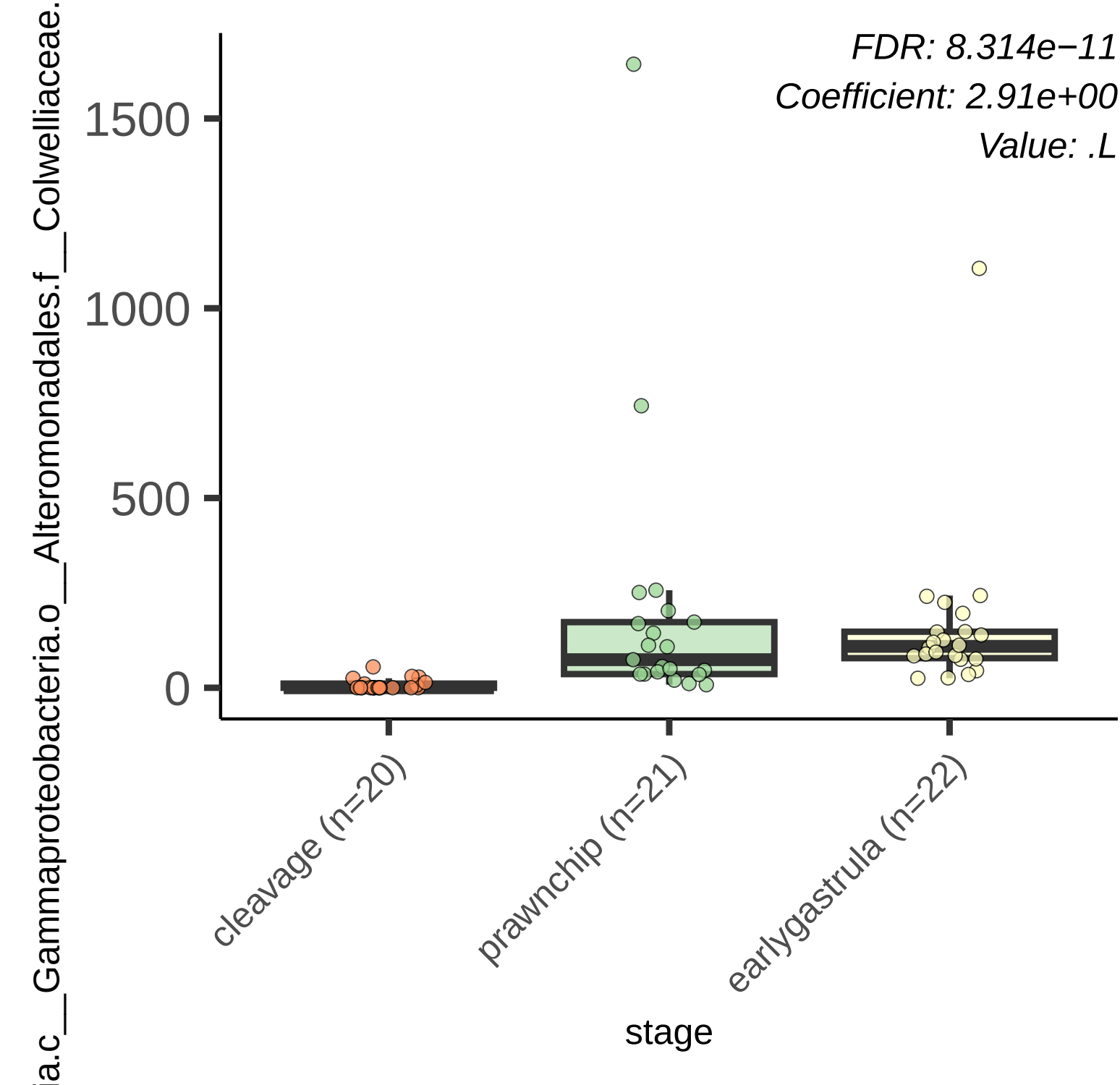
20000
15000
10000
5000
0
cleavage (n=20)
prawnchip (n=21)
earlygastrula (n=22)
stage
p__Proteobacteria.c__Alphaproteobacteria.o__Rhodobacterales.f__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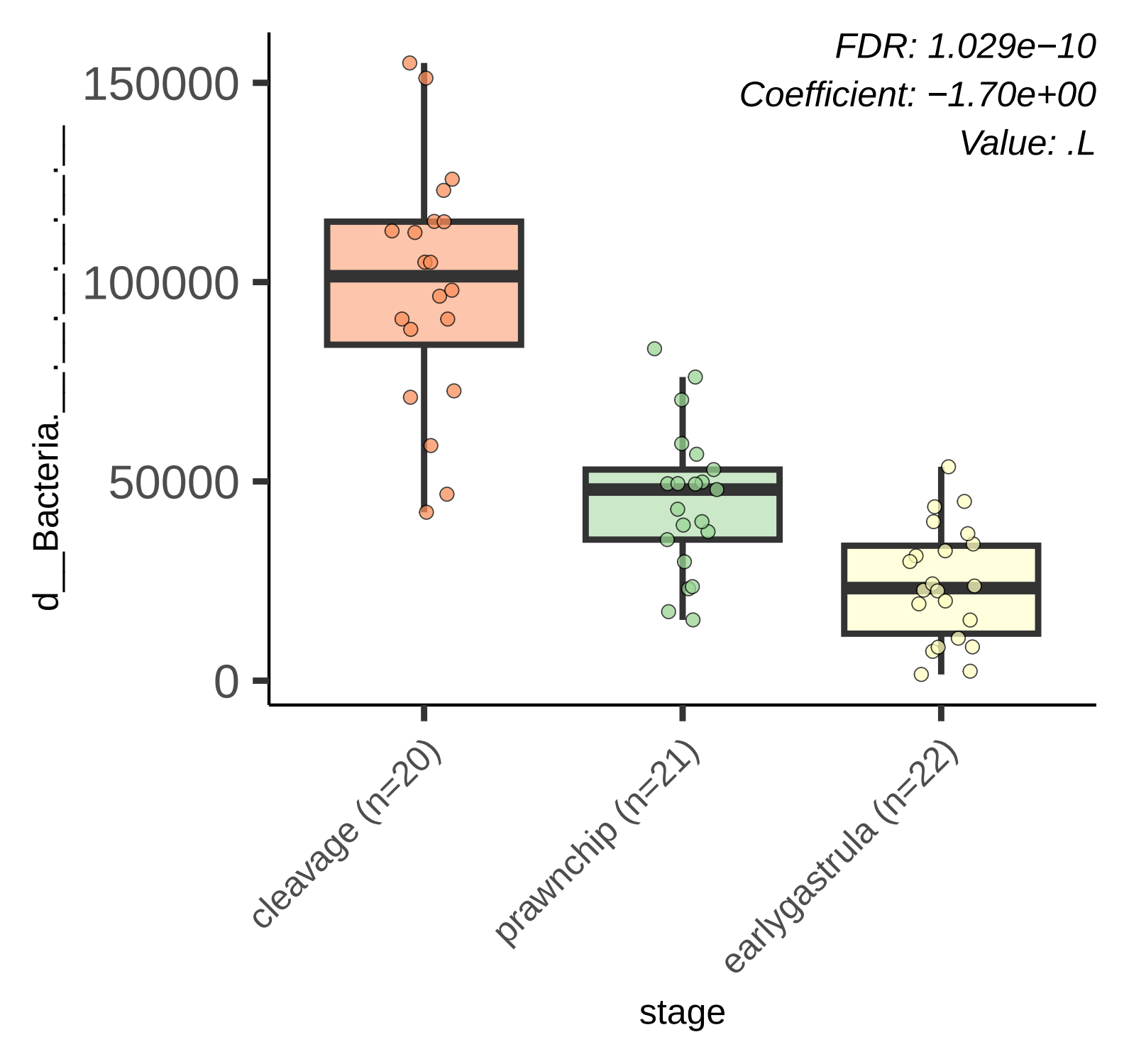
1500
1000
500
0
cleavage (n=20)
prawnchip (n=21)
earlygastrula (n=22)
stage
ia.c__Gammaproteobacteria.o__Alteromonadales.f__Colwelliaceae.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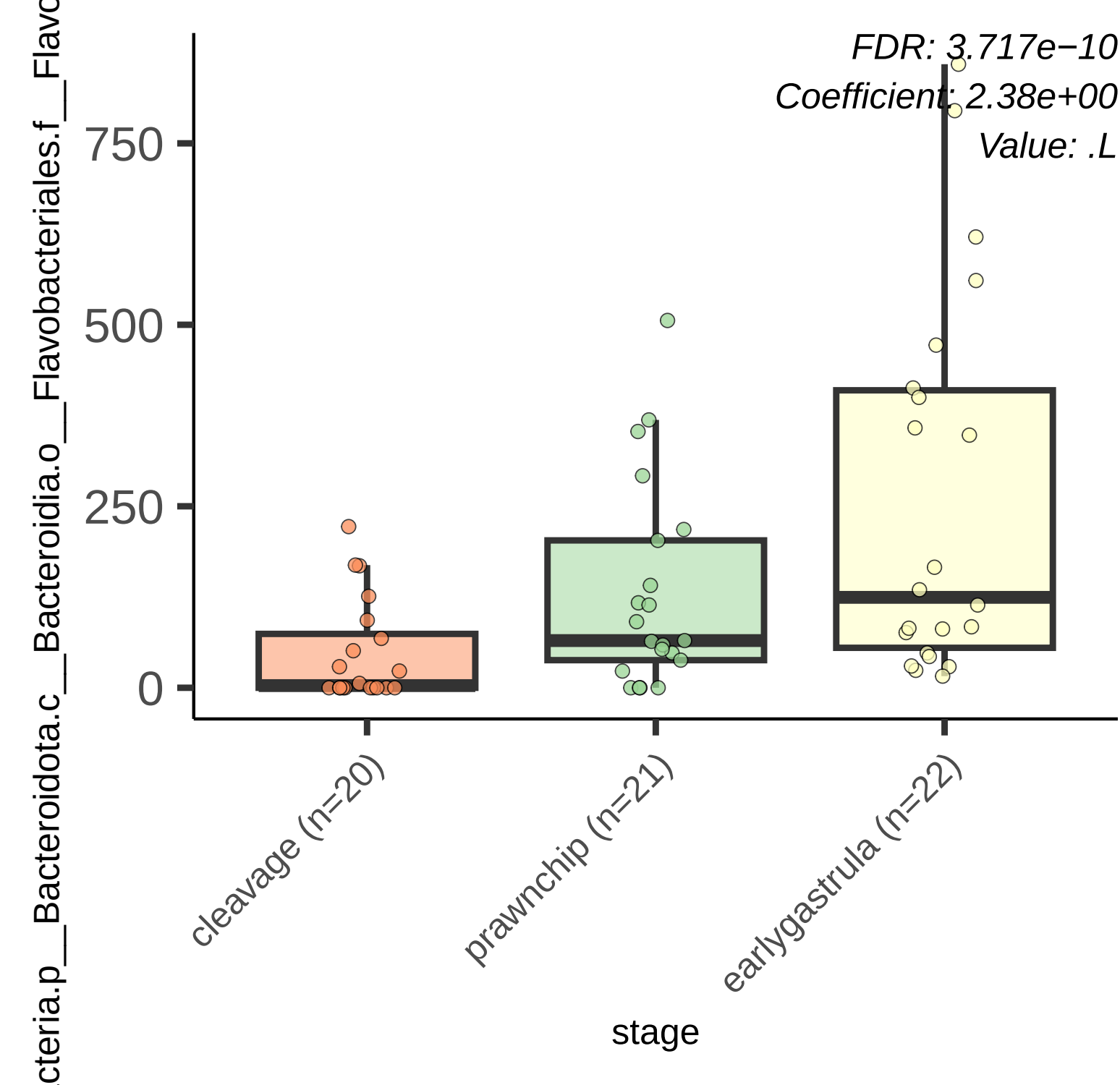
150000
100000
50000
0
d__Bacteria.__.__.__.__.__.__
cleavage (n=20)
prawnchip (n=21)
earlygastrula (n=22)
stage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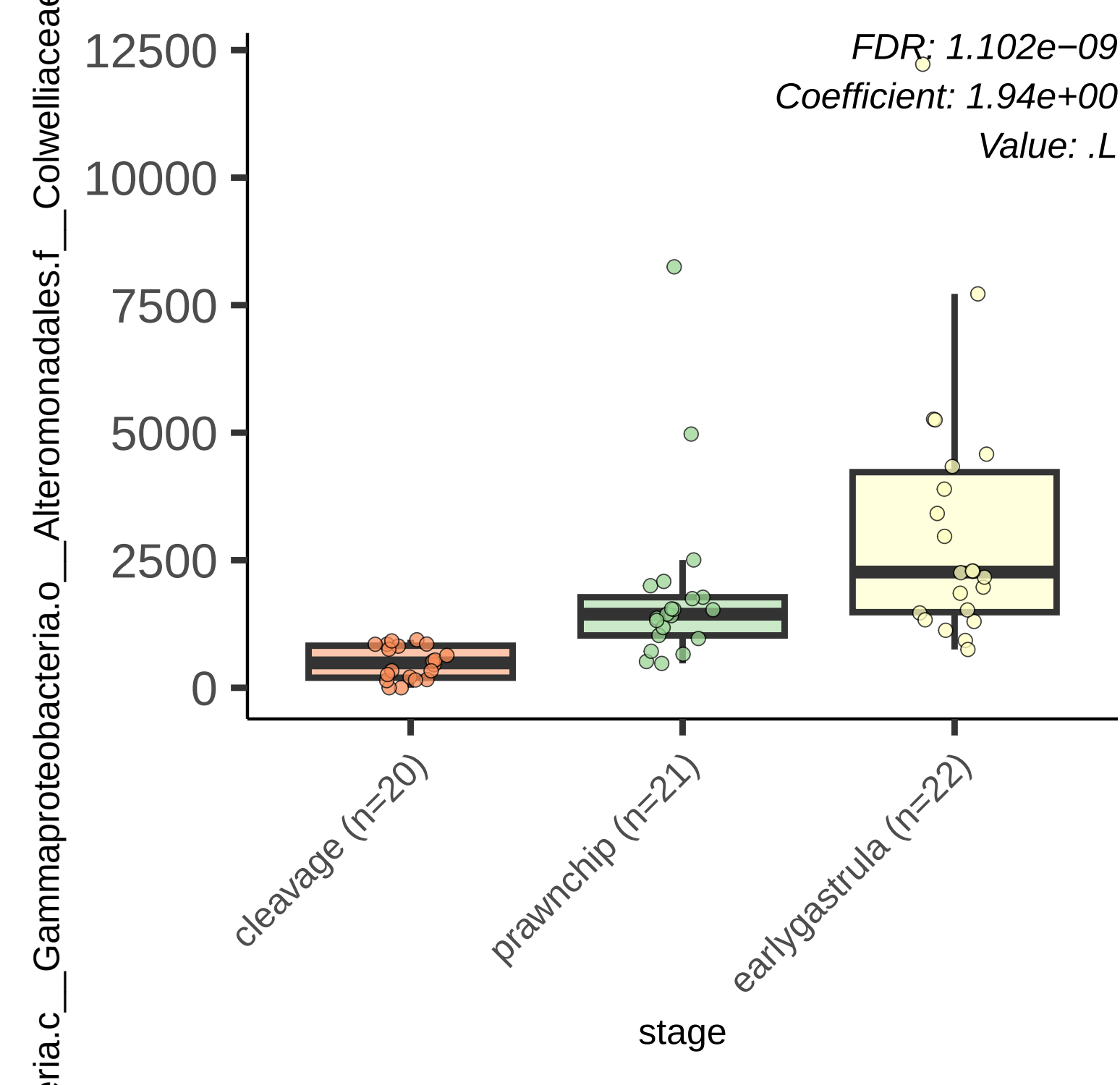
Coefficient: 2.38e+00
Value: .L
750
500
250
0
cleavage (n=20)
prawnchip (n=21)
earlygastrula (n=22)
stage
cteria.p__Bacteroidota.c__Bacteroidia.o__Flavobacteriales.f__Flavo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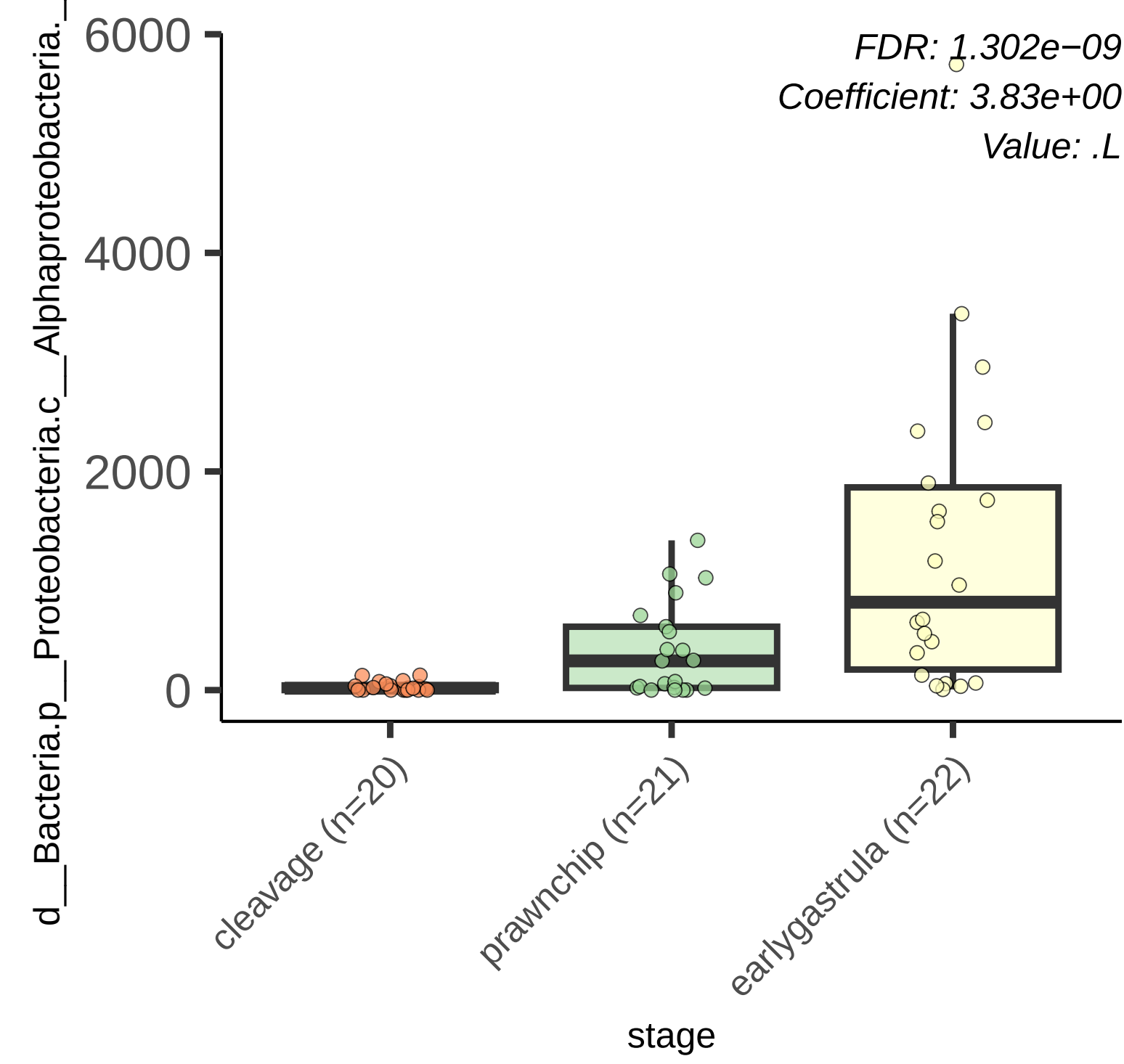
Coefficient: 1.94e+00
Value: .L
12500
10000
7500
5000
2500
0
cleavage (n=20)
prawnchip (n=21)
earlygastrula (n=22)
stage
ria.c__Gammaproteobacteria.o__Alteromonadales.f__Colwelliaceae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Coefficient: 3.83e+00
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d__Bacteria.p__Proteobacteria.c__Alphaproteobacteria.
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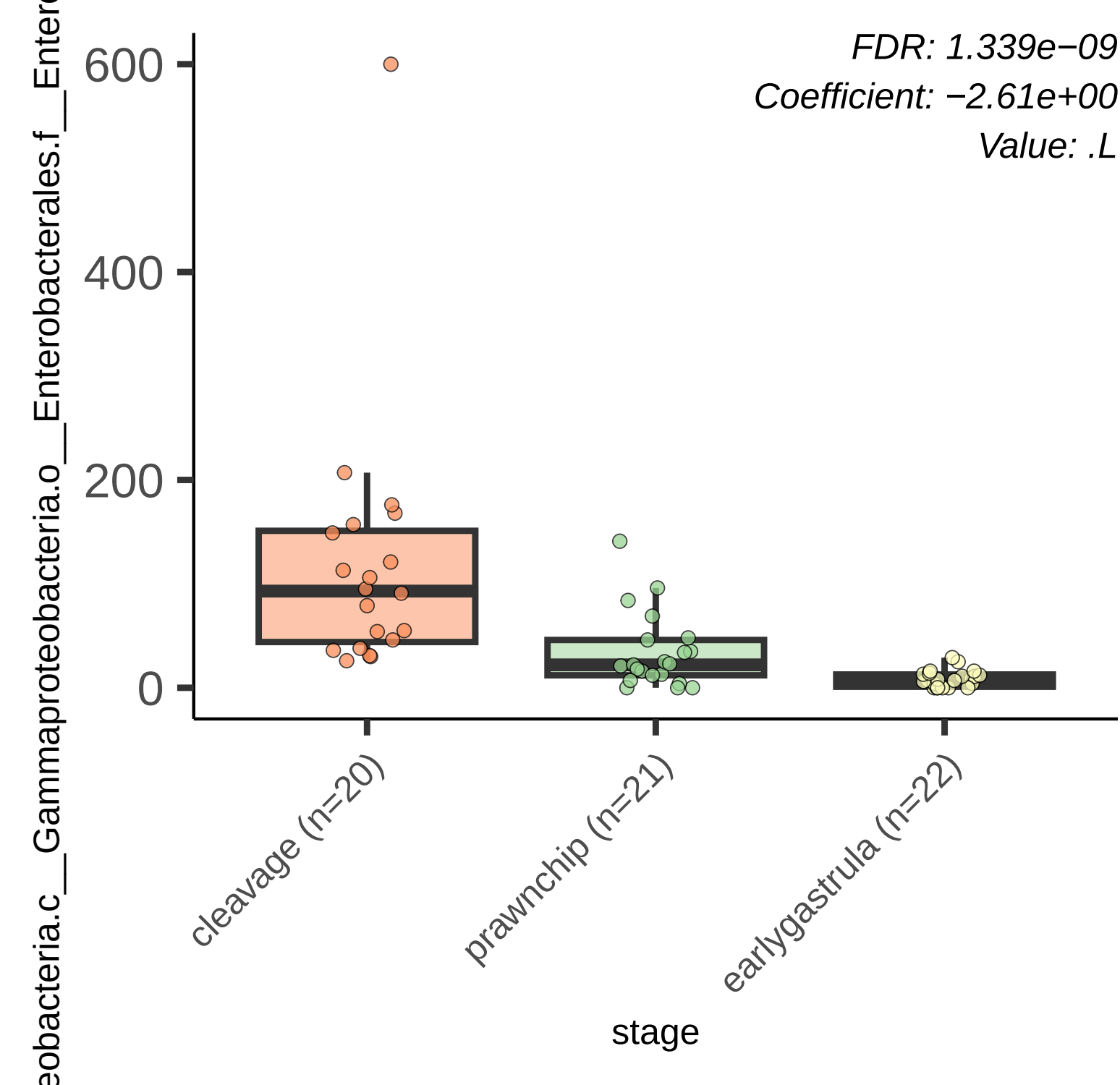
4000
2000
0
cleavage (n=20)
prawnchip (n=21)
earlygastrula (n=22)
stage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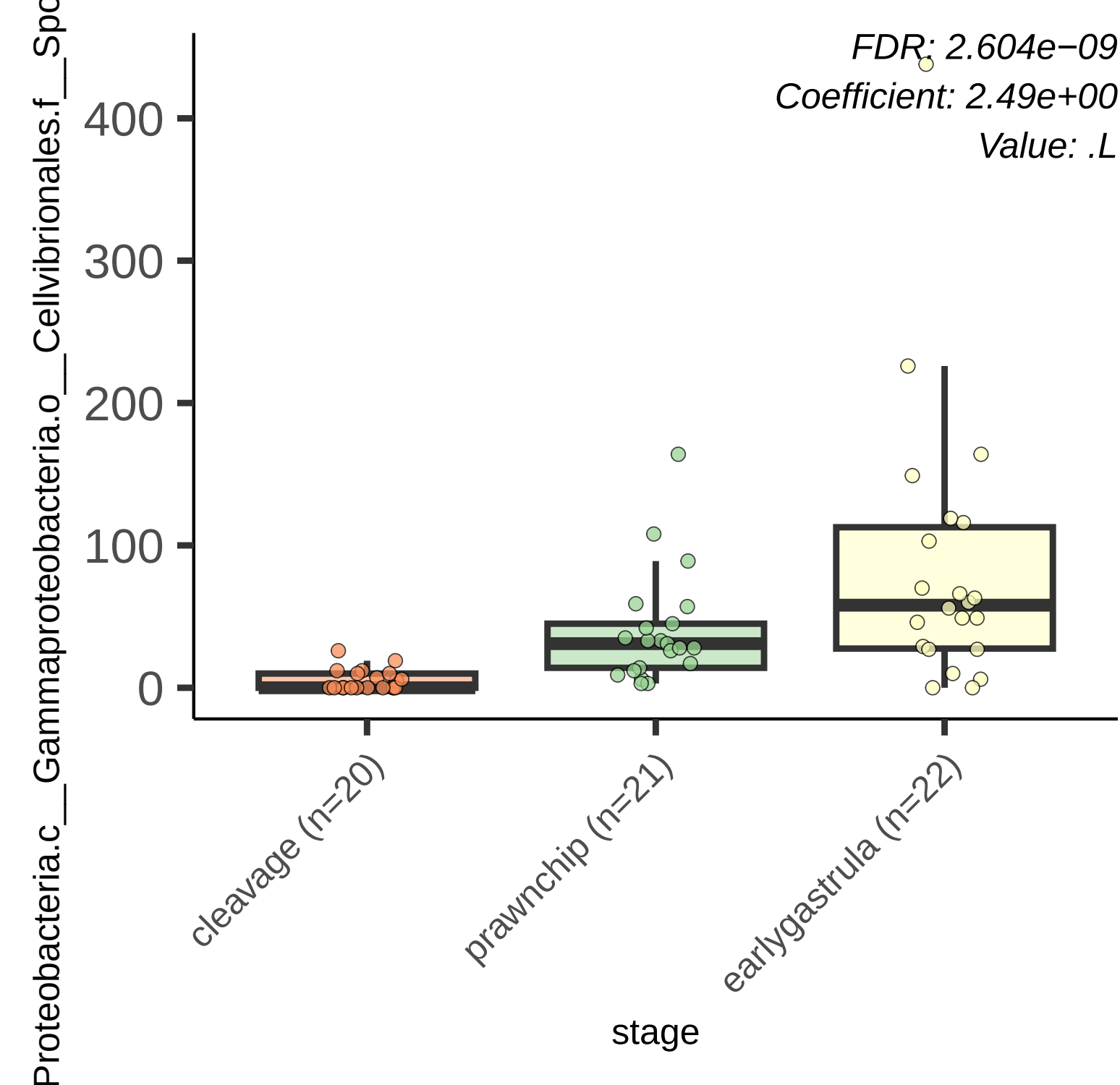
400
200
0
cleavage (n=20)
prawnchip (n=21)
earlygastrula (n=22)
stage
teobacteria.c__Gammaproteobacteria.o__Enterobacterales.f__Entero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Coefficient: 2.49e+00
Value: .L
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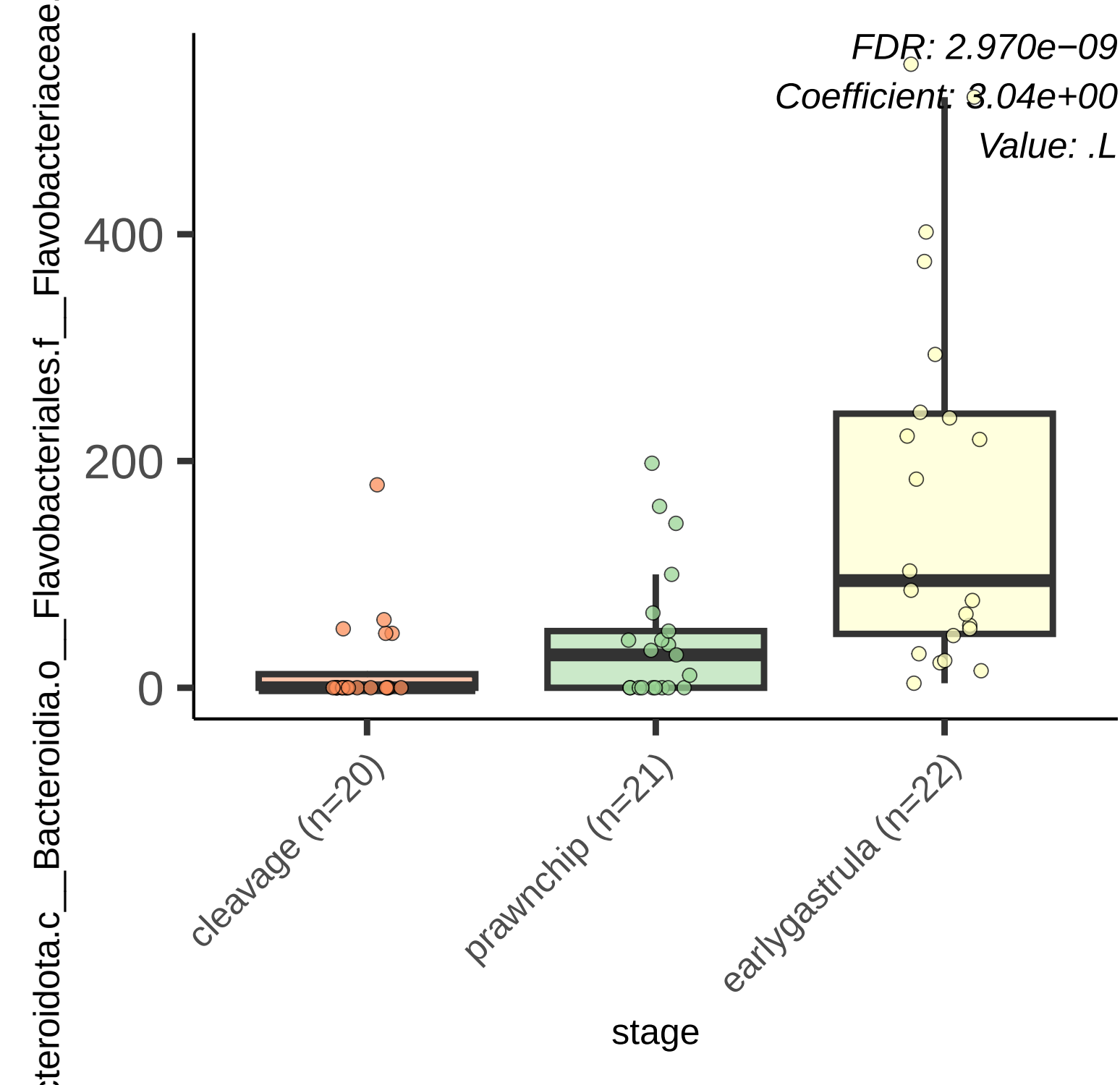
300
200
100
0
cleavage (n=20)
prawnchip (n=21)
earlygastrula (n=22)
stage
Proteobacteria.c__Gammaproteobacteria.o__Cellvibrionales.f__Spo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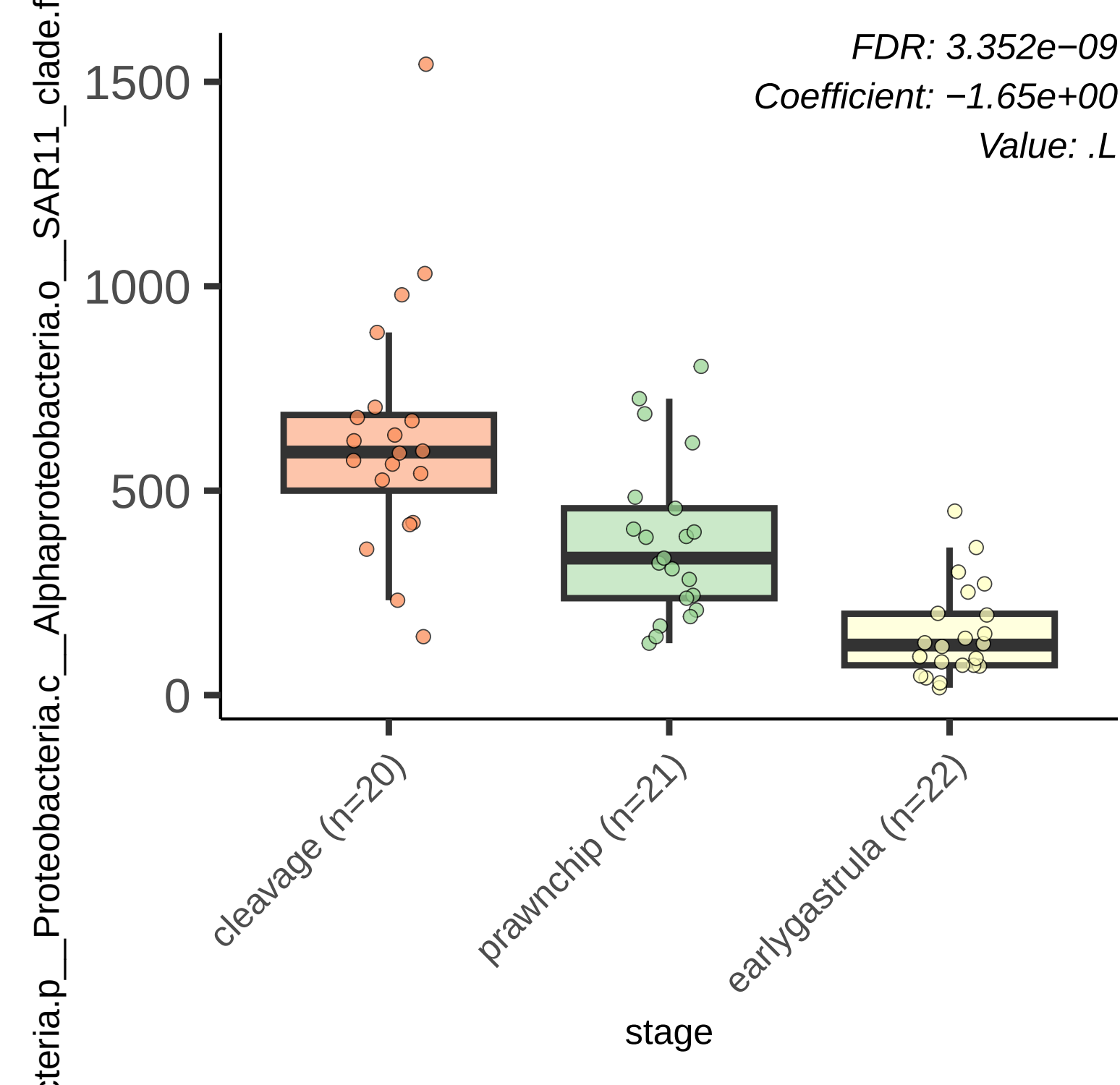
200
0
cleavage (n=20)
prawnchip (n=21)
earlygastrula (n=22)
stage
cteroidota.c__Bacteroidia.o__Flavobacteriales.f__Flavobacteriaceae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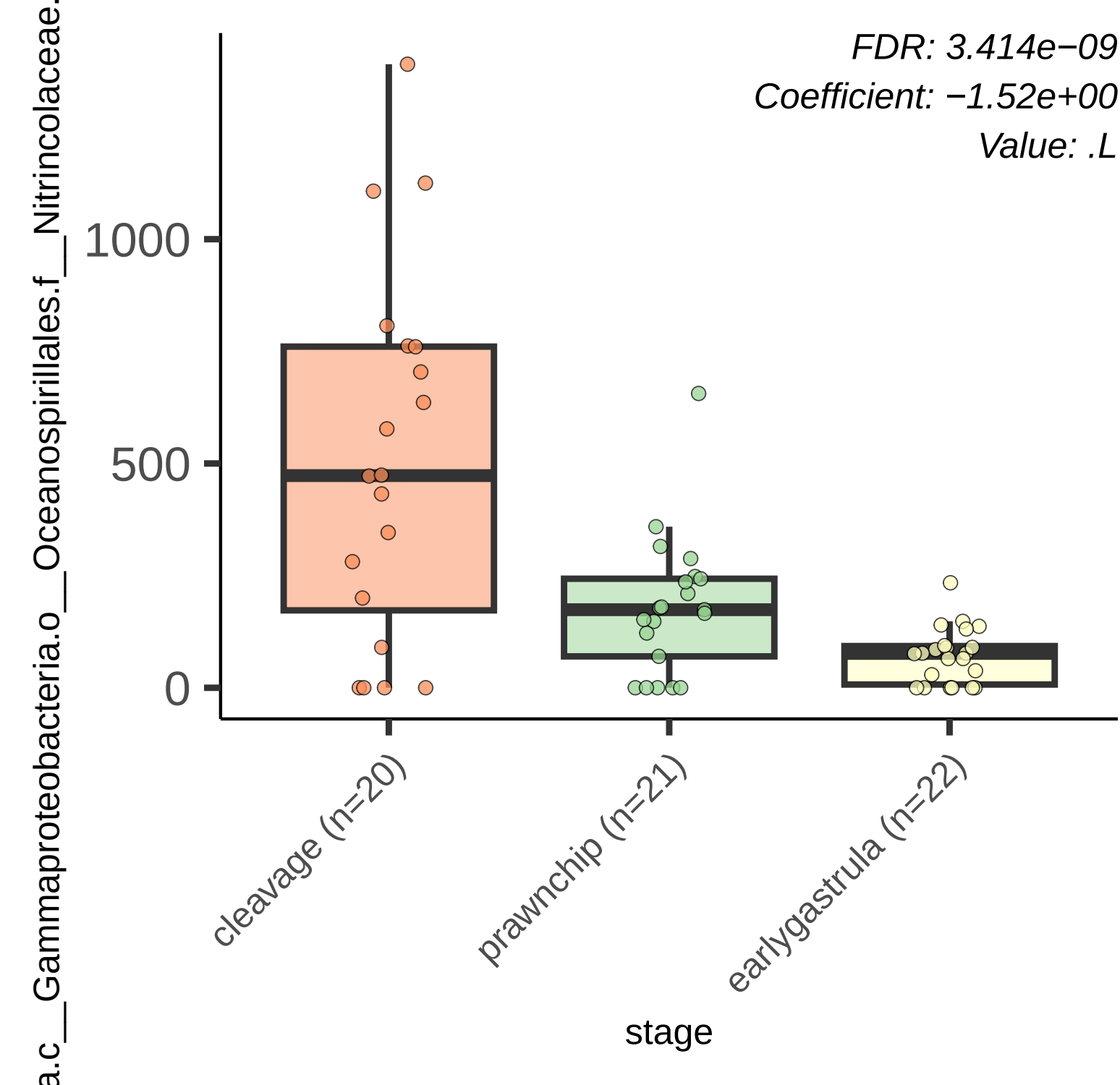
Value: .L
1500
1000
500
0
cleavage (n=20)
prawnchip (n=21)
earlygastrula (n=22)
stage
cteria.p__Proteobacteria.c__Alphaproteobacteria.o__SAR11_clade.f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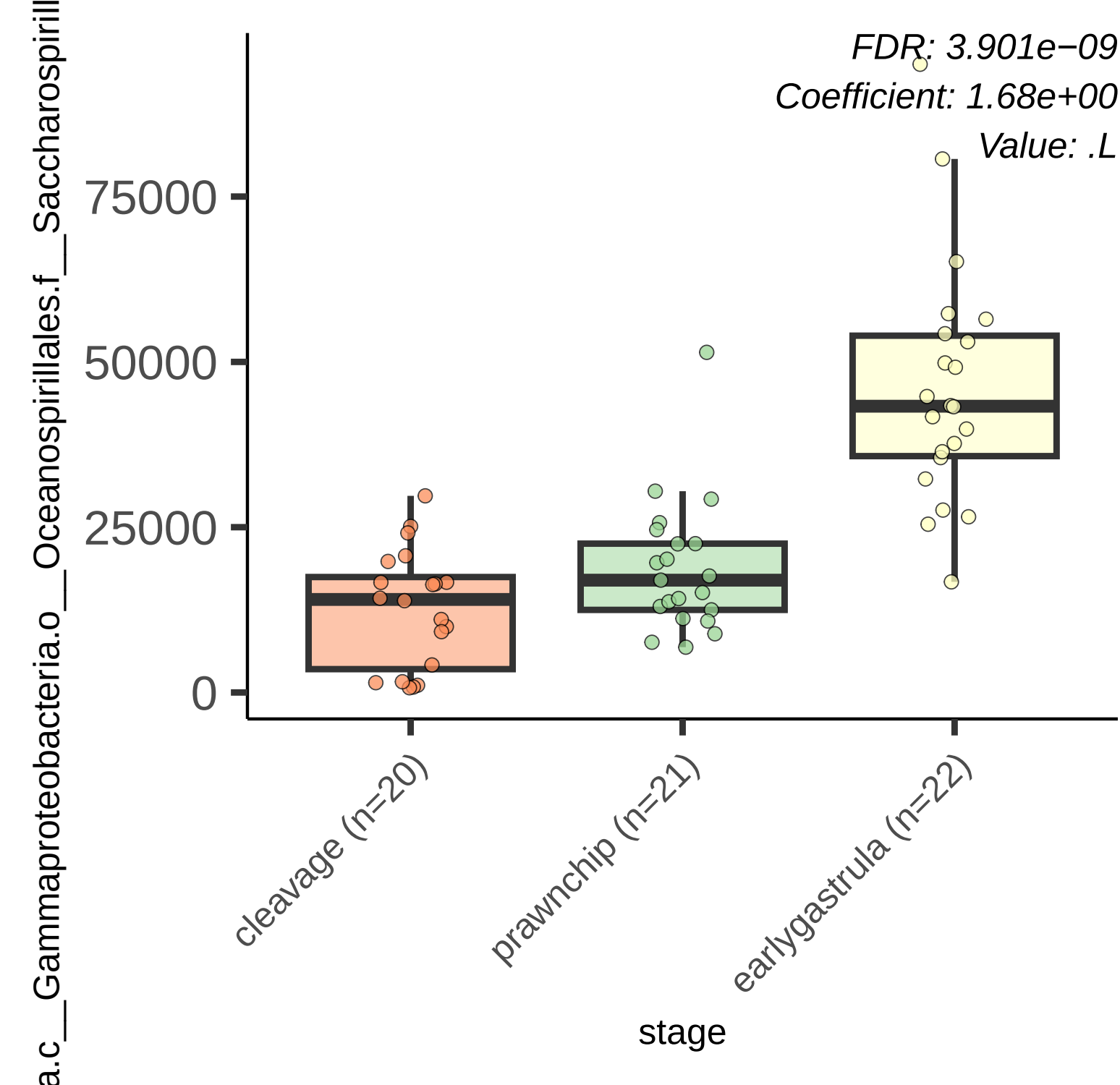
Coefficient: −1.52e+00
Value: .L
1000
500
0
cleavage (n=20)
prawnchip (n=21)
earlygastrula (n=22)
stage
a.c__Gammaproteobacteria.o__Oceanospirillales.f__Nitrincolaceae.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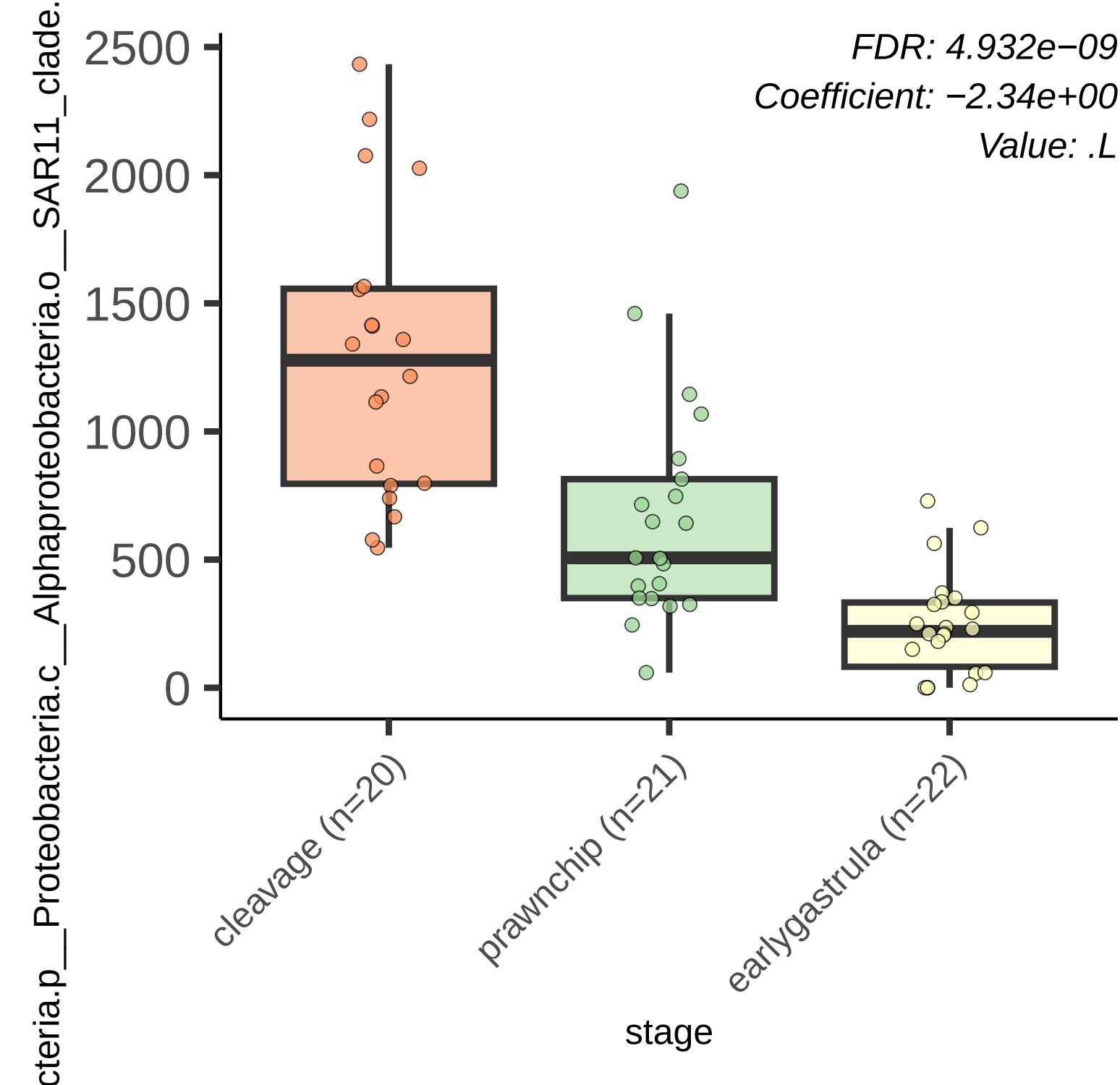
50000
25000
0
cleavage (n=20)
prawnchip (n=21)
earlygastrula (n=22)
stage
a.c__Gammaproteobacteria.o__Oceanospirillales.f__Saccharospirill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Coefficient: −2.34e+00
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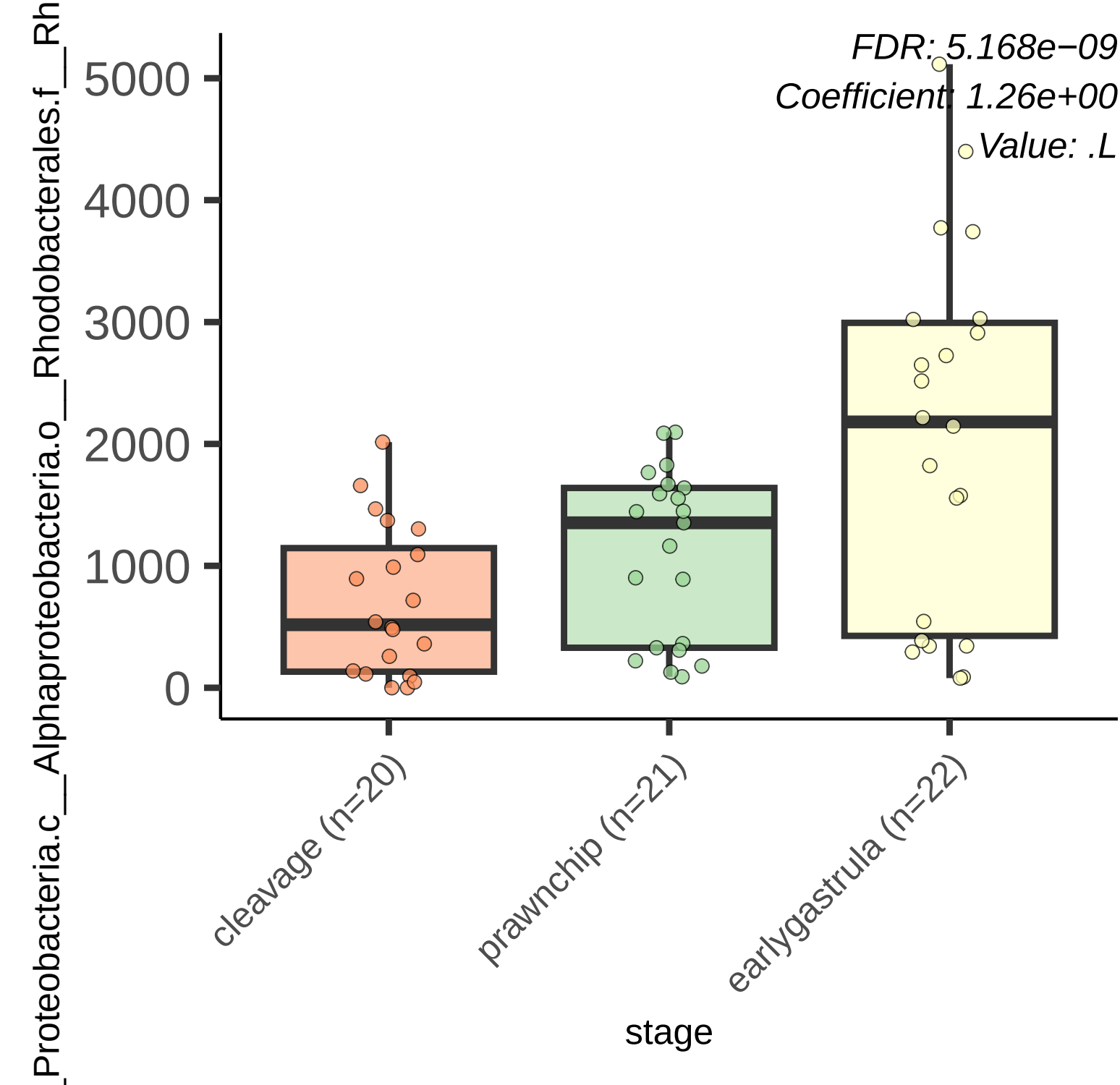
2500
2000
1500
1000
500
0
cleavage (n=20)
prawnchip (n=21)
earlygastrula (n=22)
stage
cteria.p__Proteobacteria.c__Alphaproteobacteria.o__SAR11_clade.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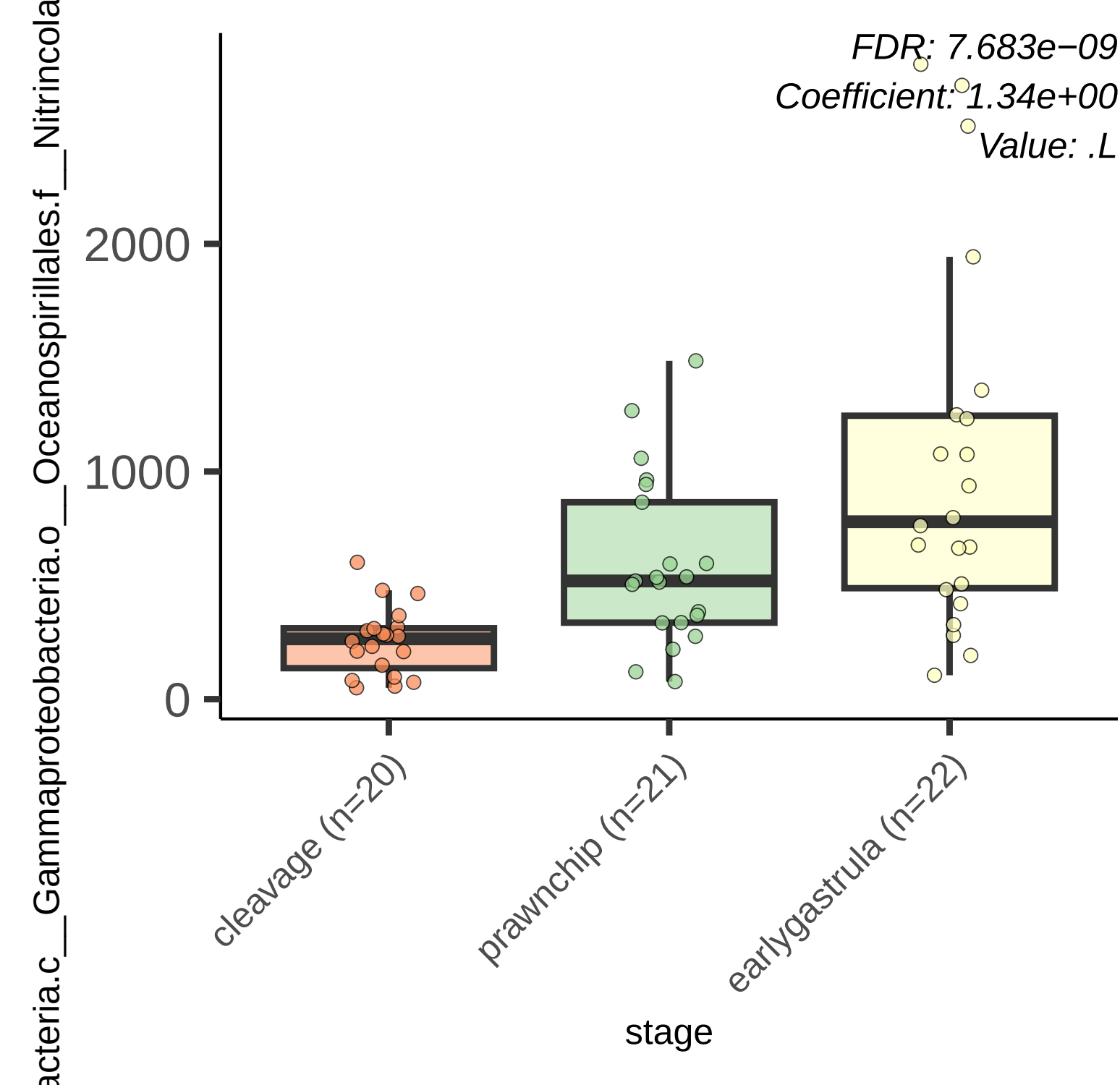
Coefficient: 1.26e+00
Value: .L
5000
4000
3000
2000
1000
0
cleavage (n=20)
prawnchip (n=21)
earlygastrula (n=22)
stage
_Proteobacteria.c__Alphaproteobacteria.o__Rhodobacterales.f__Rh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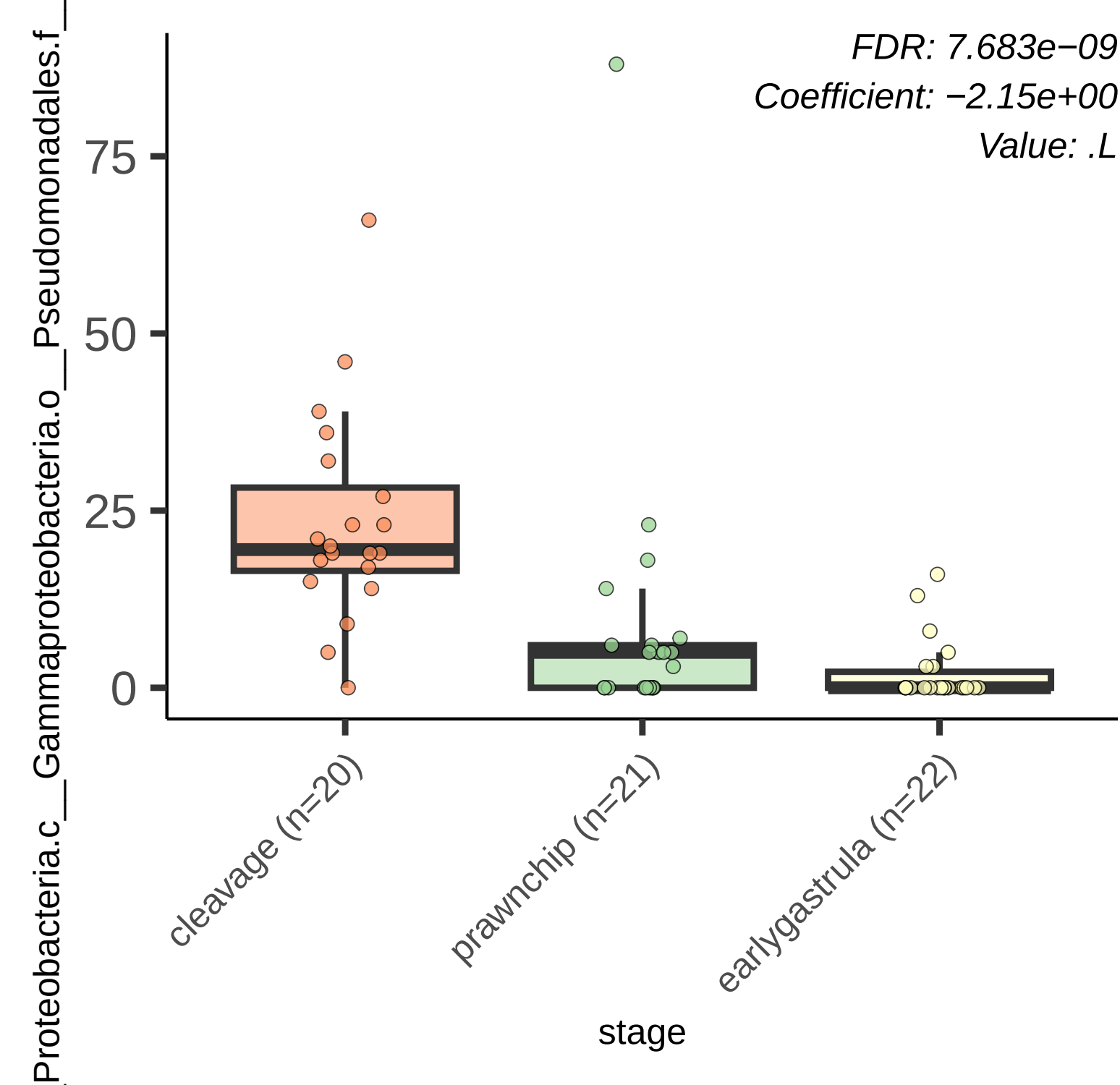
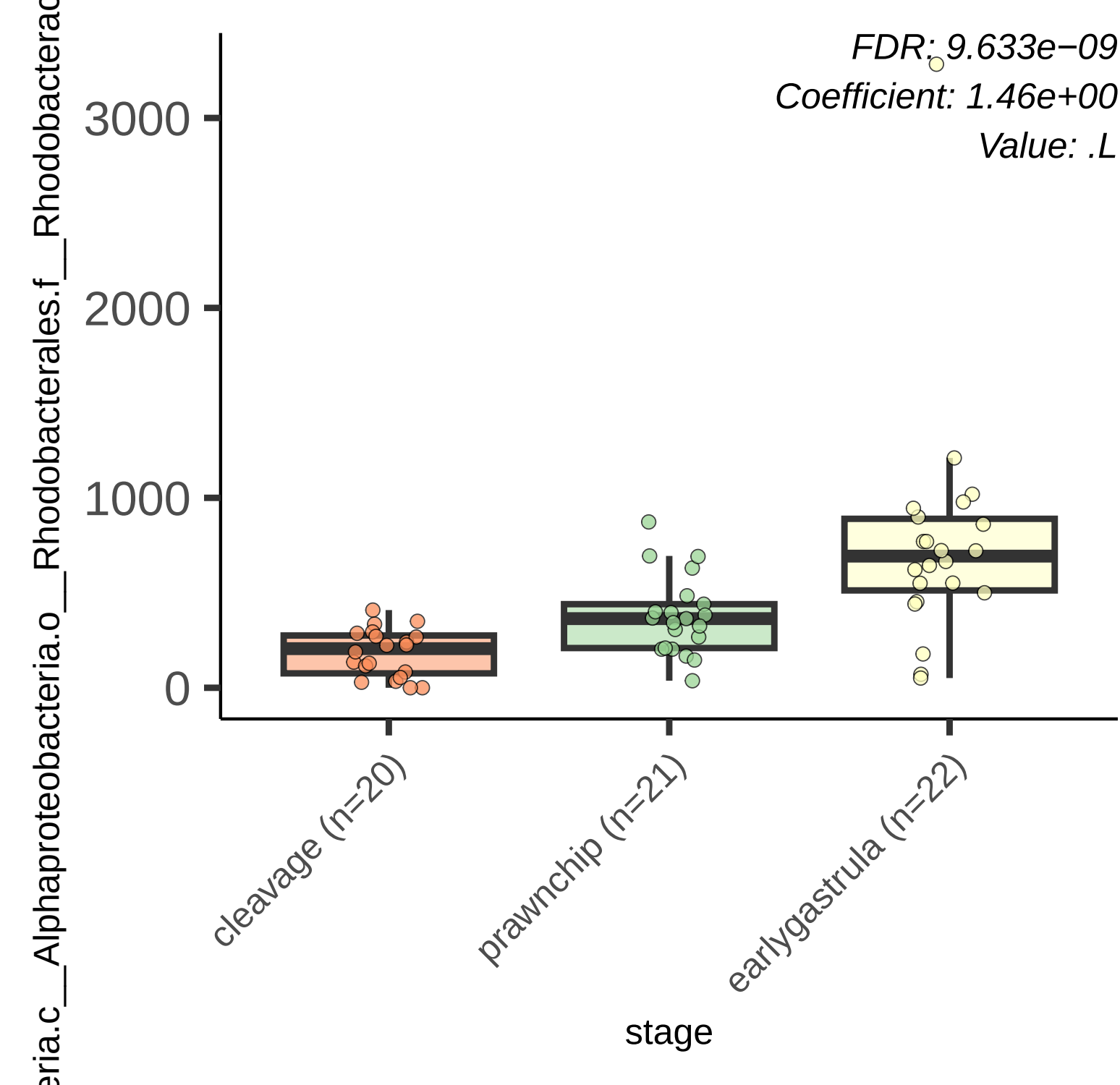
Coefficient: 1.34e+00
Value: .L
2000
1000
0
cleavage (n=20)
prawnchip (n=21)
earlygastrula (n=22)
stage
acteria.c__Gammaproteobacteria.o__Oceanospirillales.f__Nitrincola


FDR: 7.683e−09
Coefficient: −2.15e+00
Value: .L
75
50
25
0
cleavage (n=20)
prawnchip (n=21)
earlygastrula (n=22)
stage
Proteobacteria.c__Gammaproteobacteria.o__Pseudomonadales.f__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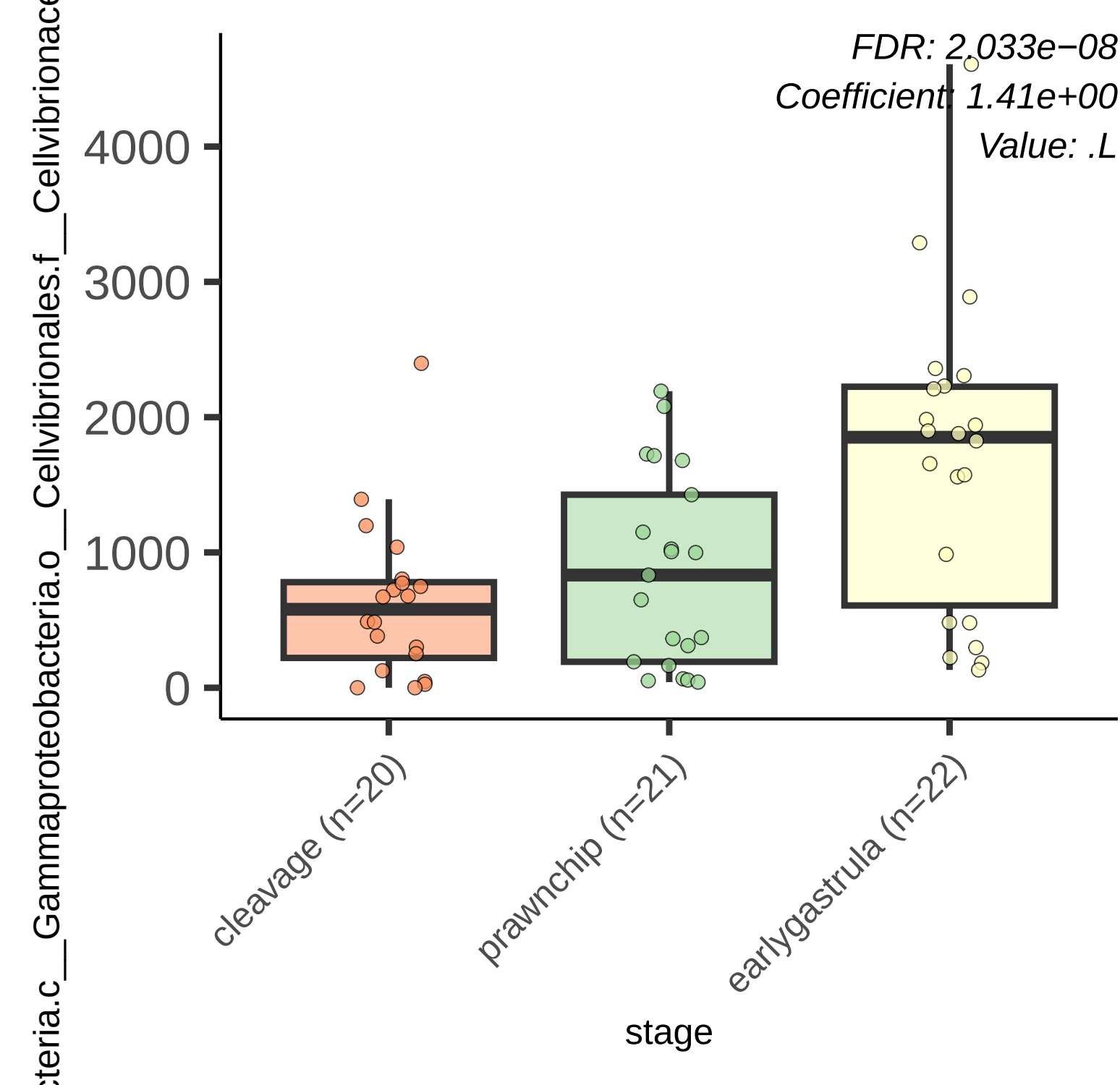
2000
1000
0
cleavage (n=20)
prawnchip (n=21)
earlygastrula (n=22)
stage
eria.c__Alphaproteobacteria.o__Rhodobacterales.f__Rhodobacterac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3000
2000
1000
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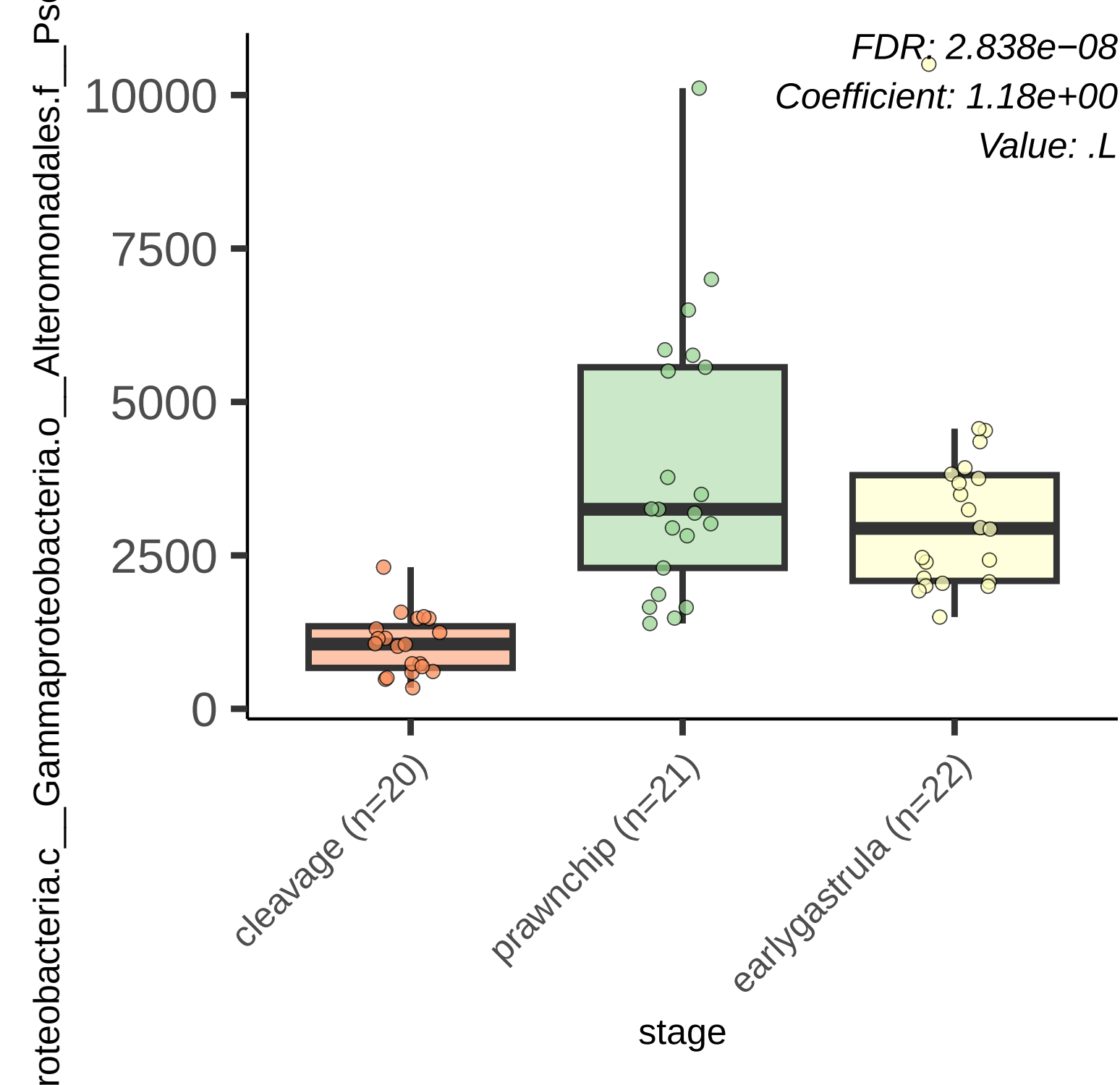
prawnchip (n=21)
earlygastrula (n=22)
stage
cteria.c__Gammaproteobacteria.o__Cellvibrionales.f__Cellvibrionace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5000
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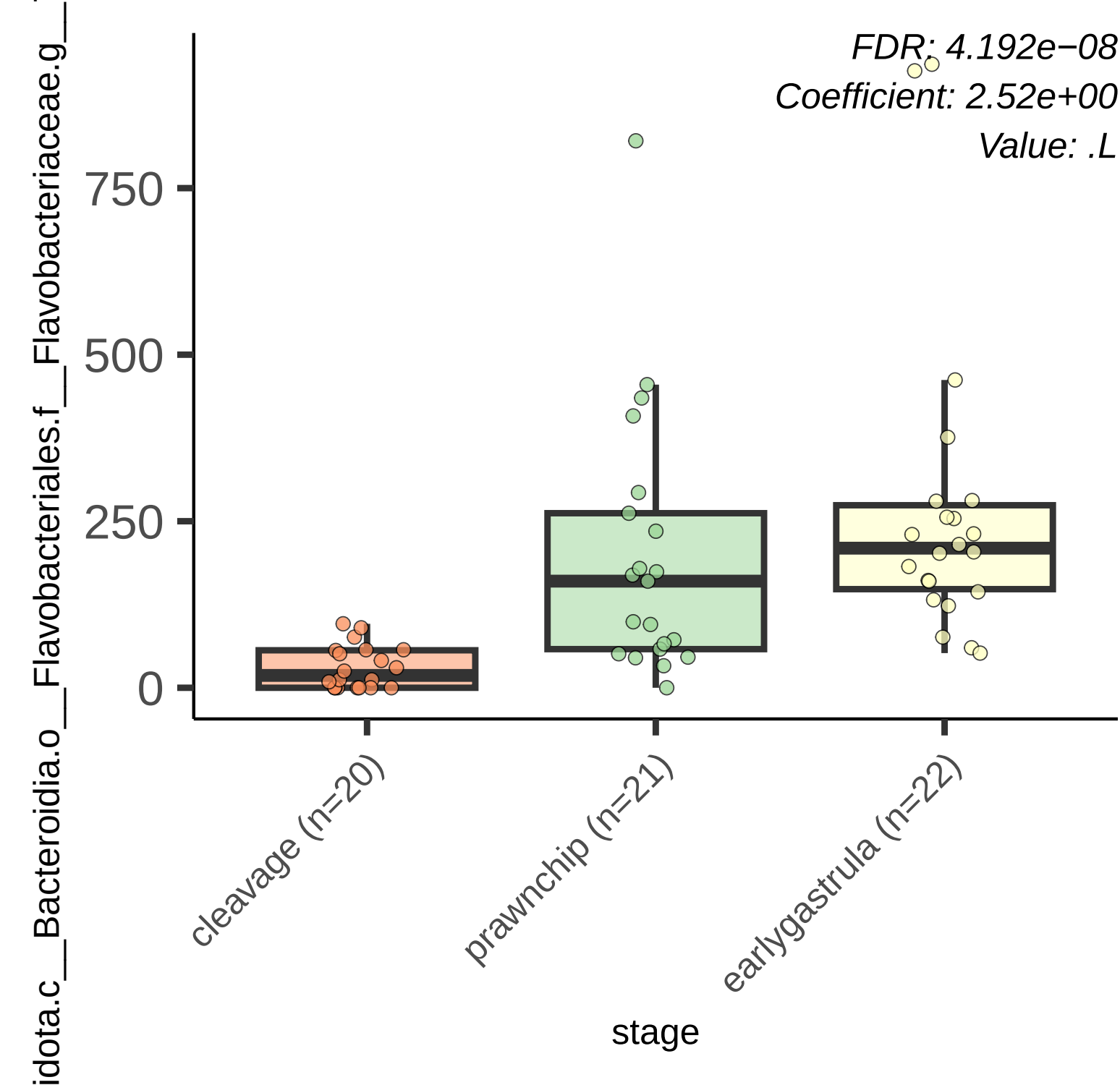
prawnchip (n=21)
earlygastrula (n=22)
stage
roteobacteria.c__Gammaproteobacteria.o__Alteromonadales.f__Ps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250
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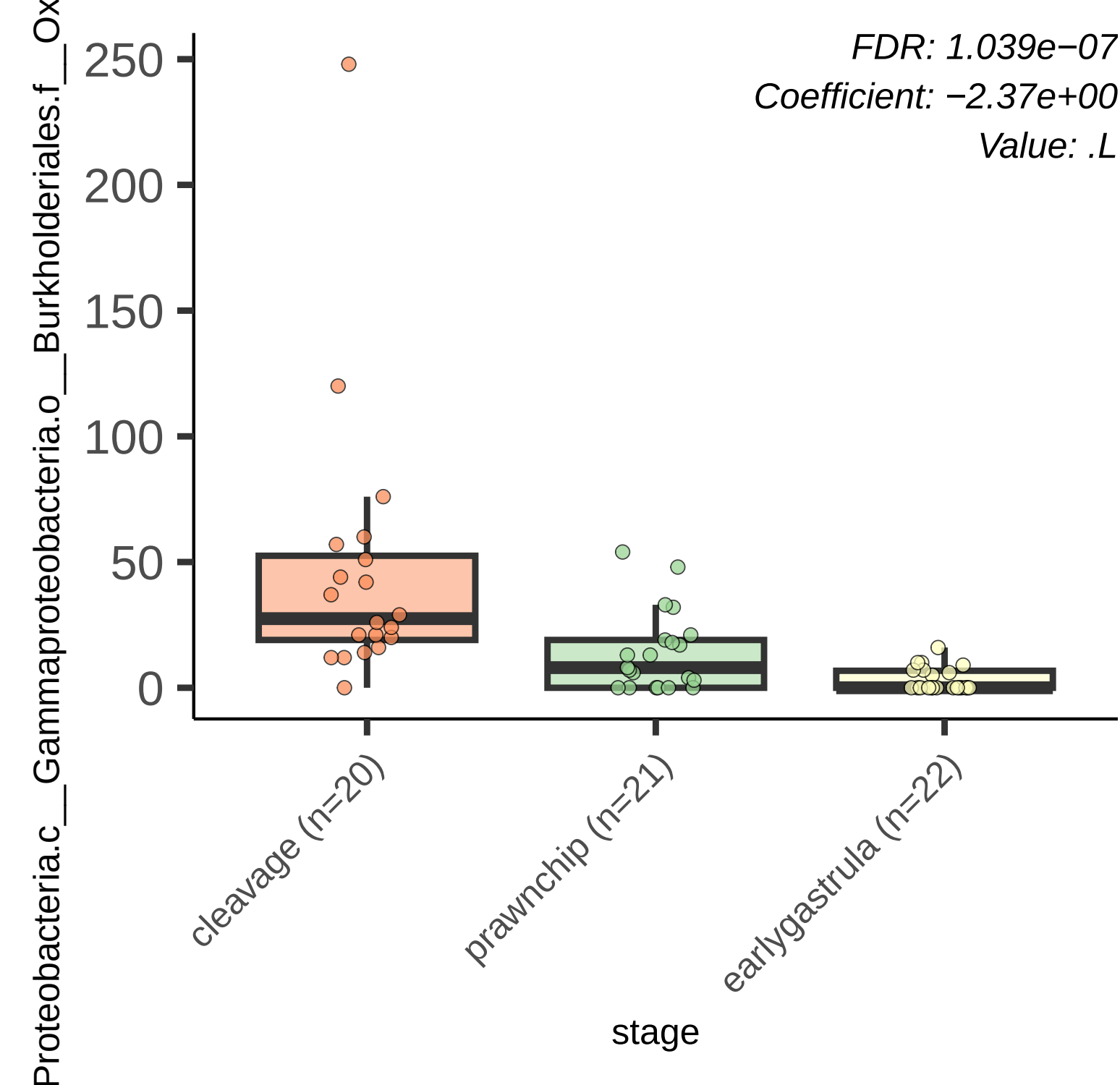
prawnchip (n=21)
earlygastrula (n=22)
stage
idota.c__Bacteroidia.o__Flavobacteriales.f__Flavobacteriaceae.g__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150
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50
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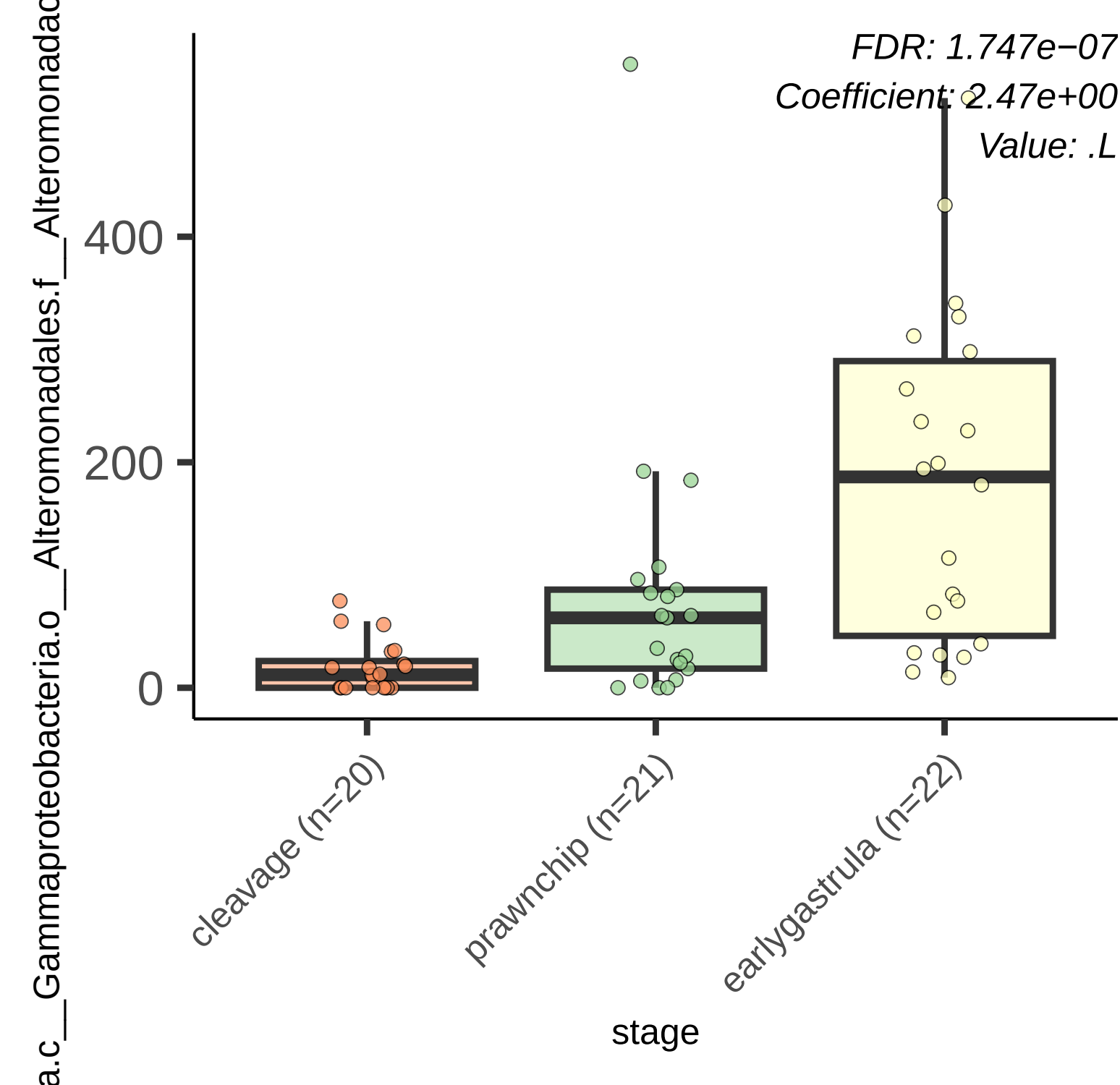
prawnchip (n=21)
earlygastrula (n=22)
stage
Proteobacteria.c__Gammaproteobacteria.o__Burkholderiales.f__Ox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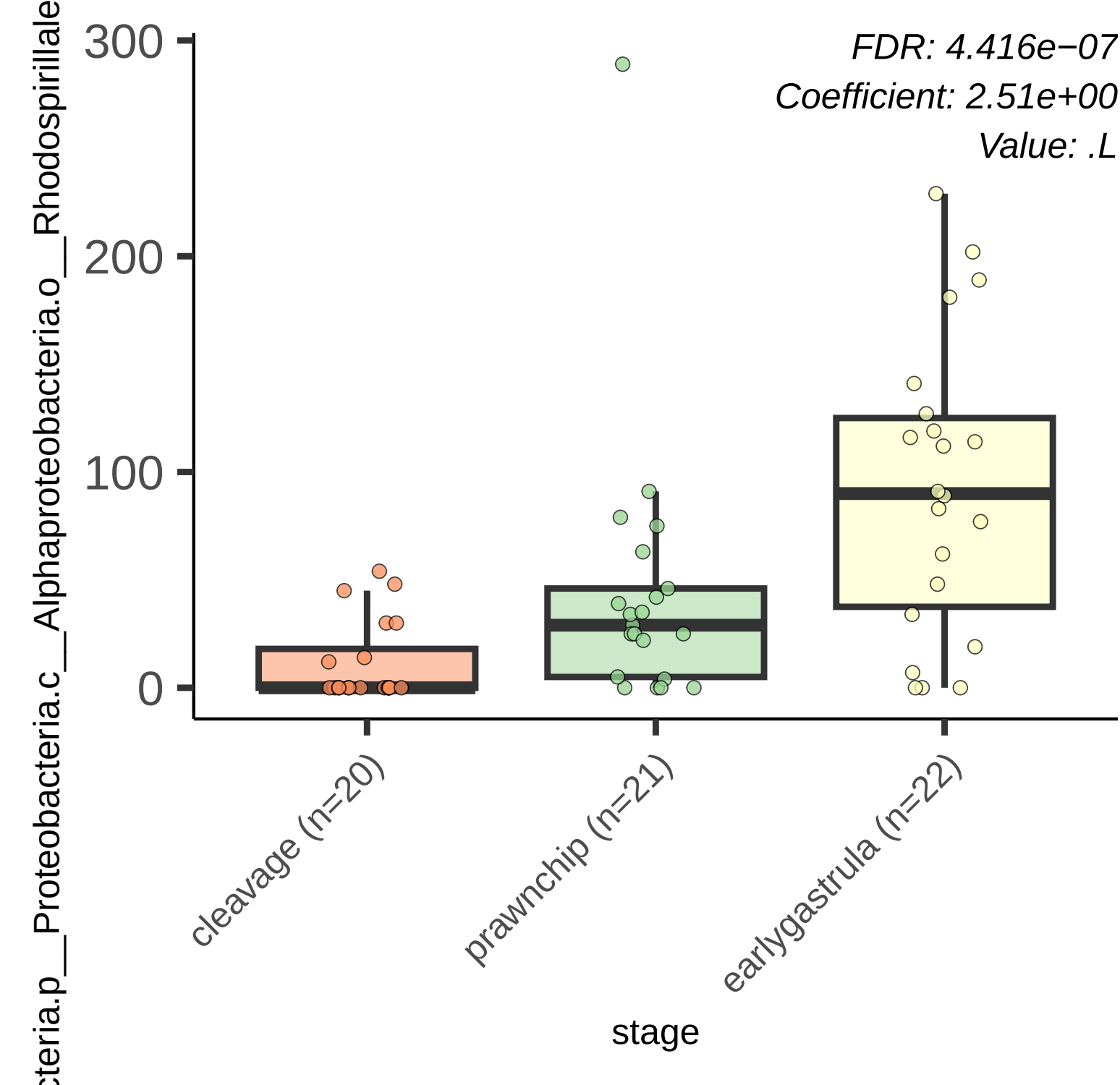
400
200
0
cleavage (n=20)
prawnchip (n=21)
earlygastrula (n=22)
stage
a.c__Gammaproteobacteria.o__Alteromonadales.f__Alteromonadac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100
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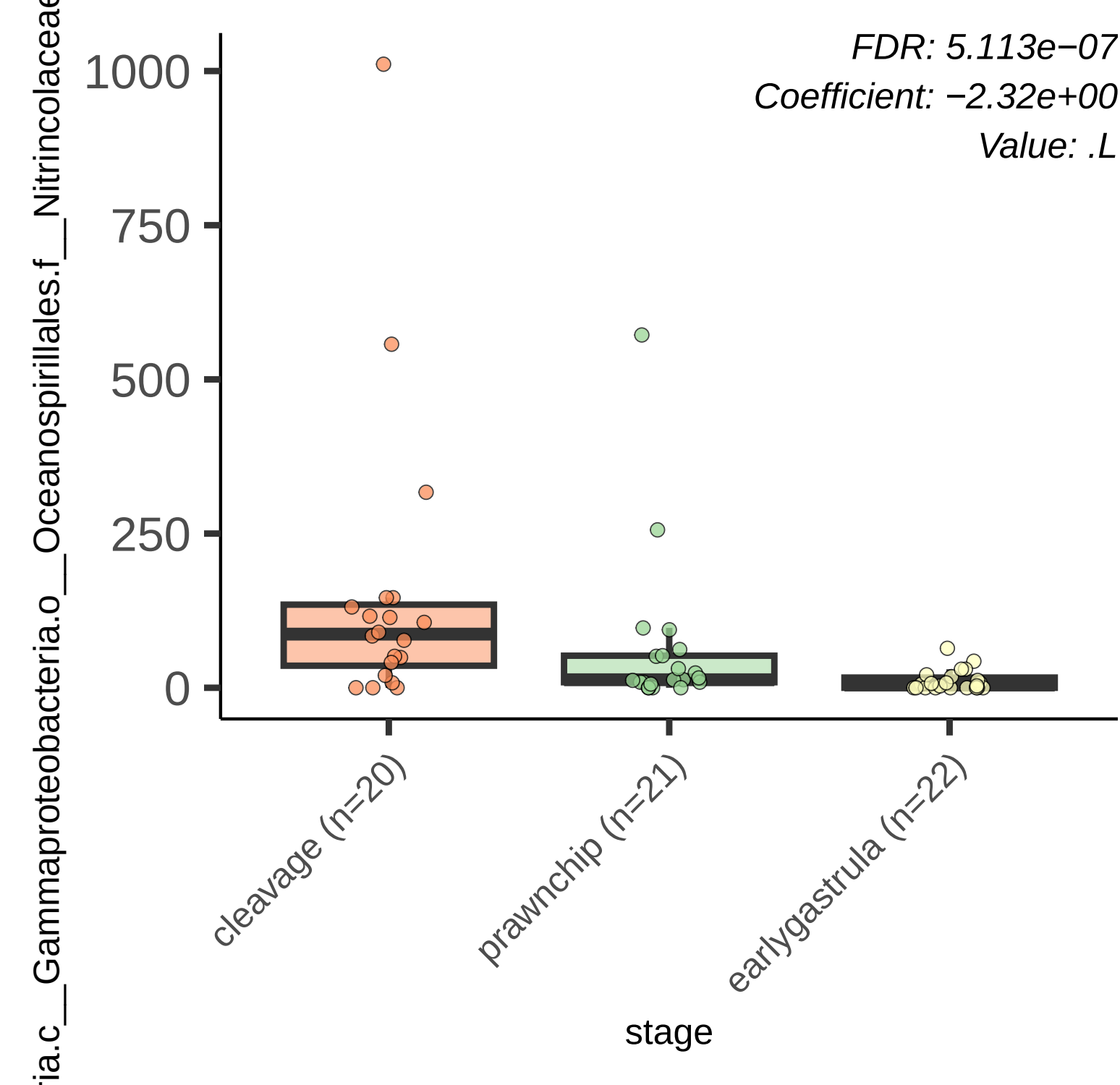
earlygastrula (n=22)
stage
cteria.p__Proteobacteria.c__Alphaproteobacteria.o__Rhodospirillale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500
250
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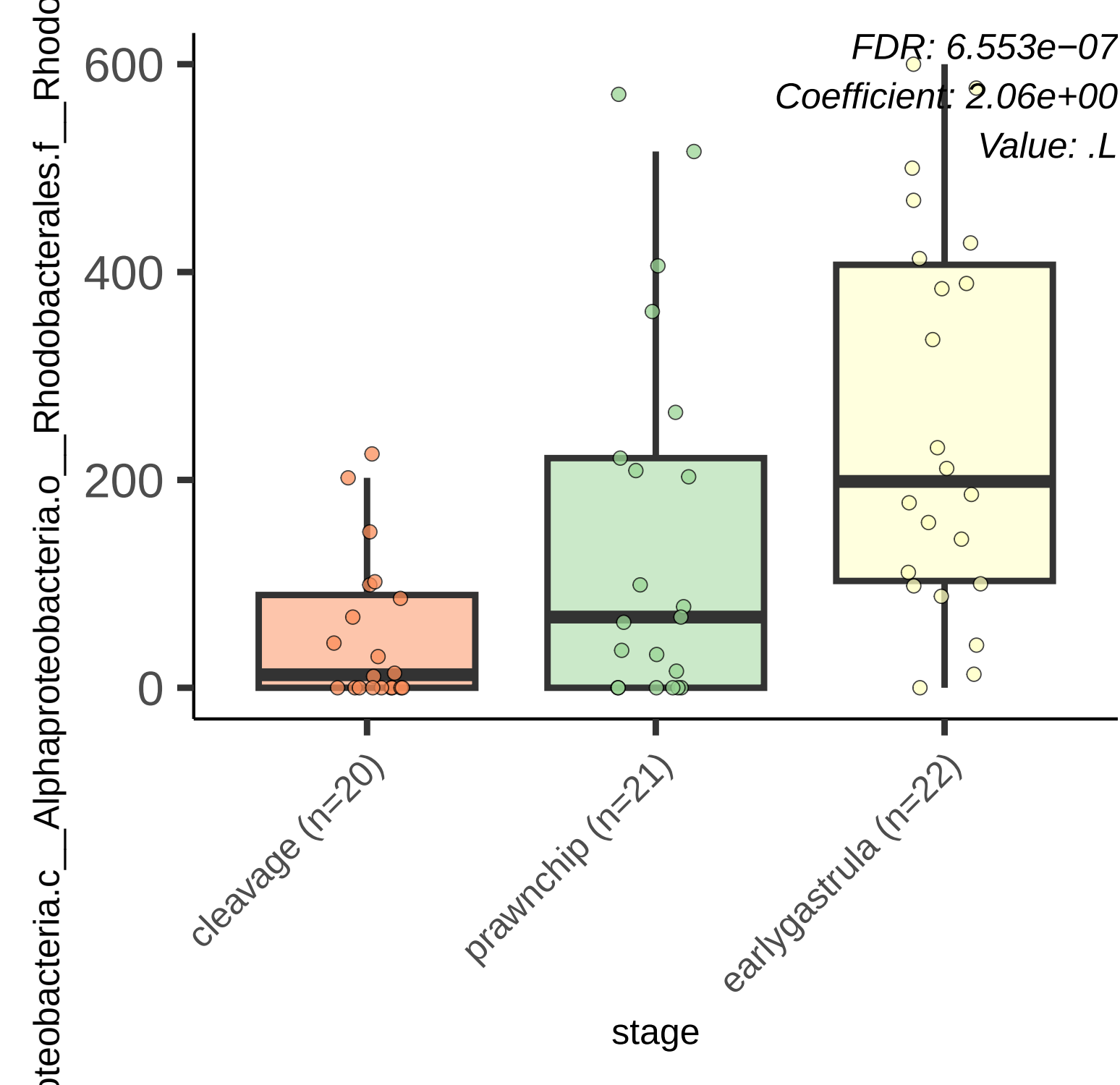
earlygastrula (n=22)
stage
ia.c__Gammaproteobacteria.o__Oceanospirillales.f__Nitrincolacea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400
200
0
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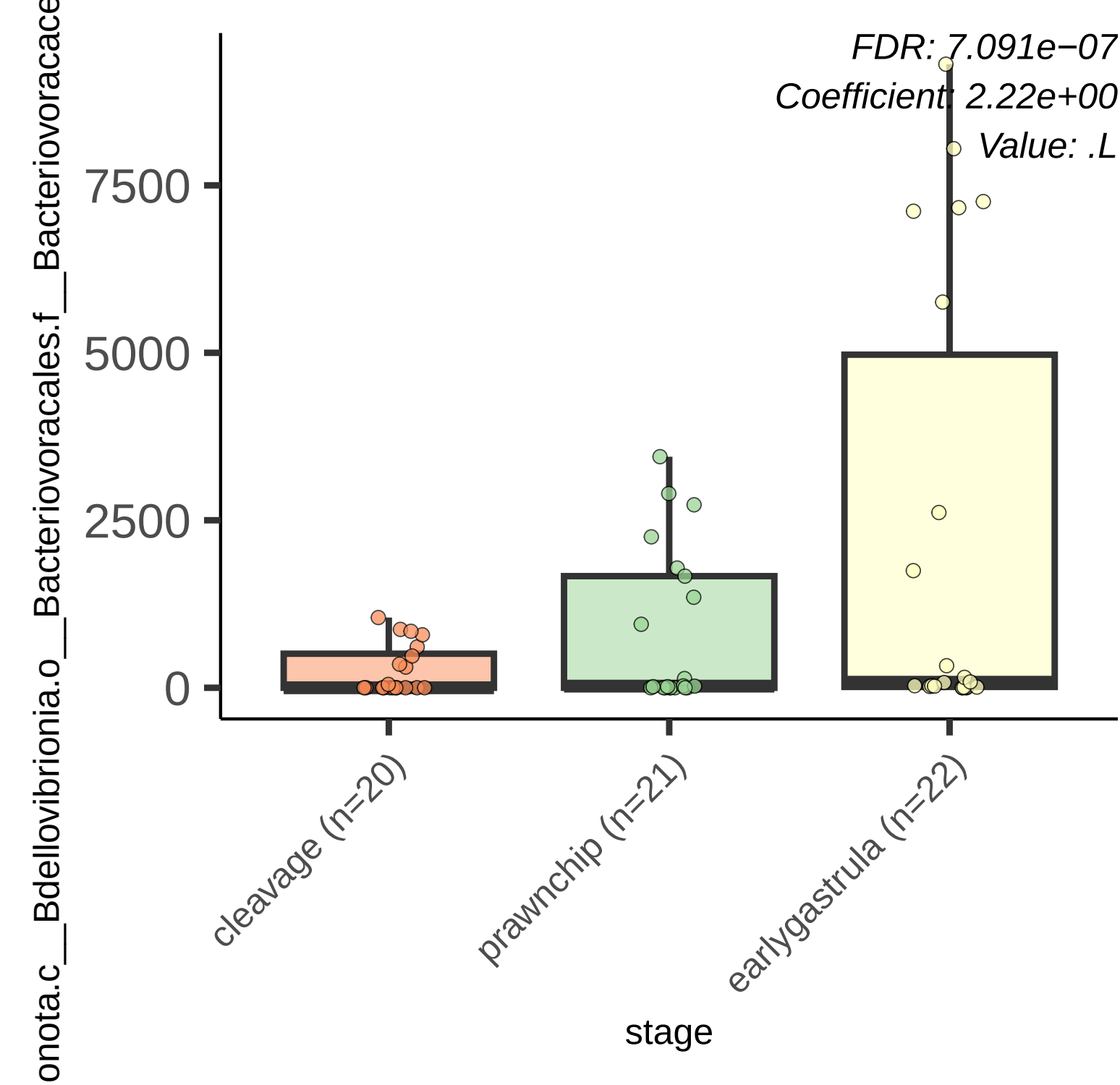
prawnchip (n=21)
earlygastrula (n=22)
stage
teobacteria.c__Alphaproteobacteria.o__Rhodobacterales.f__Rhodo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2500
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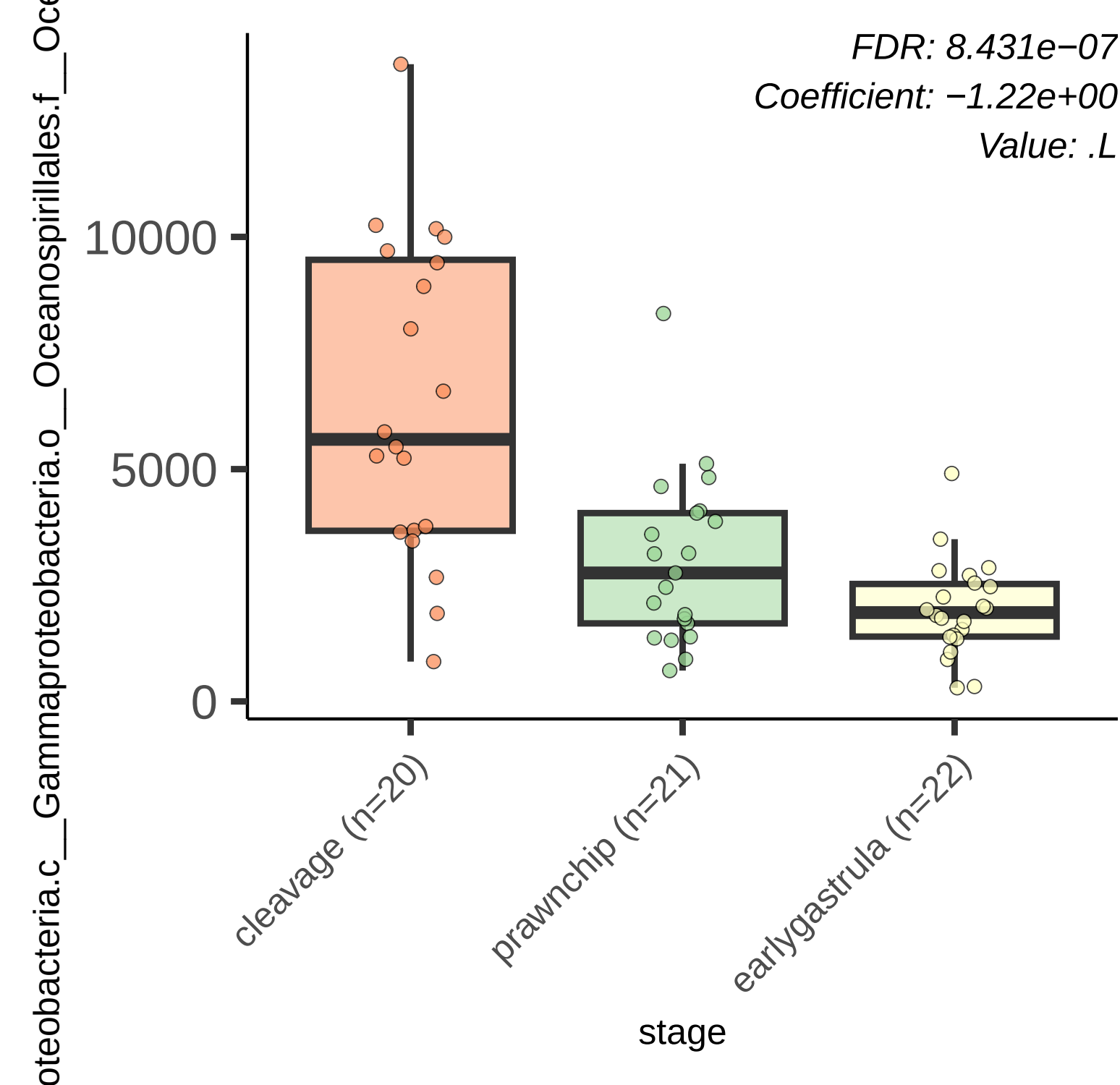
prawnchip (n=21)
earlygastrula (n=22)
stage
onota.c__Bdellovibrionia.o__Bacteriovoracales.f__Bacteriovoracace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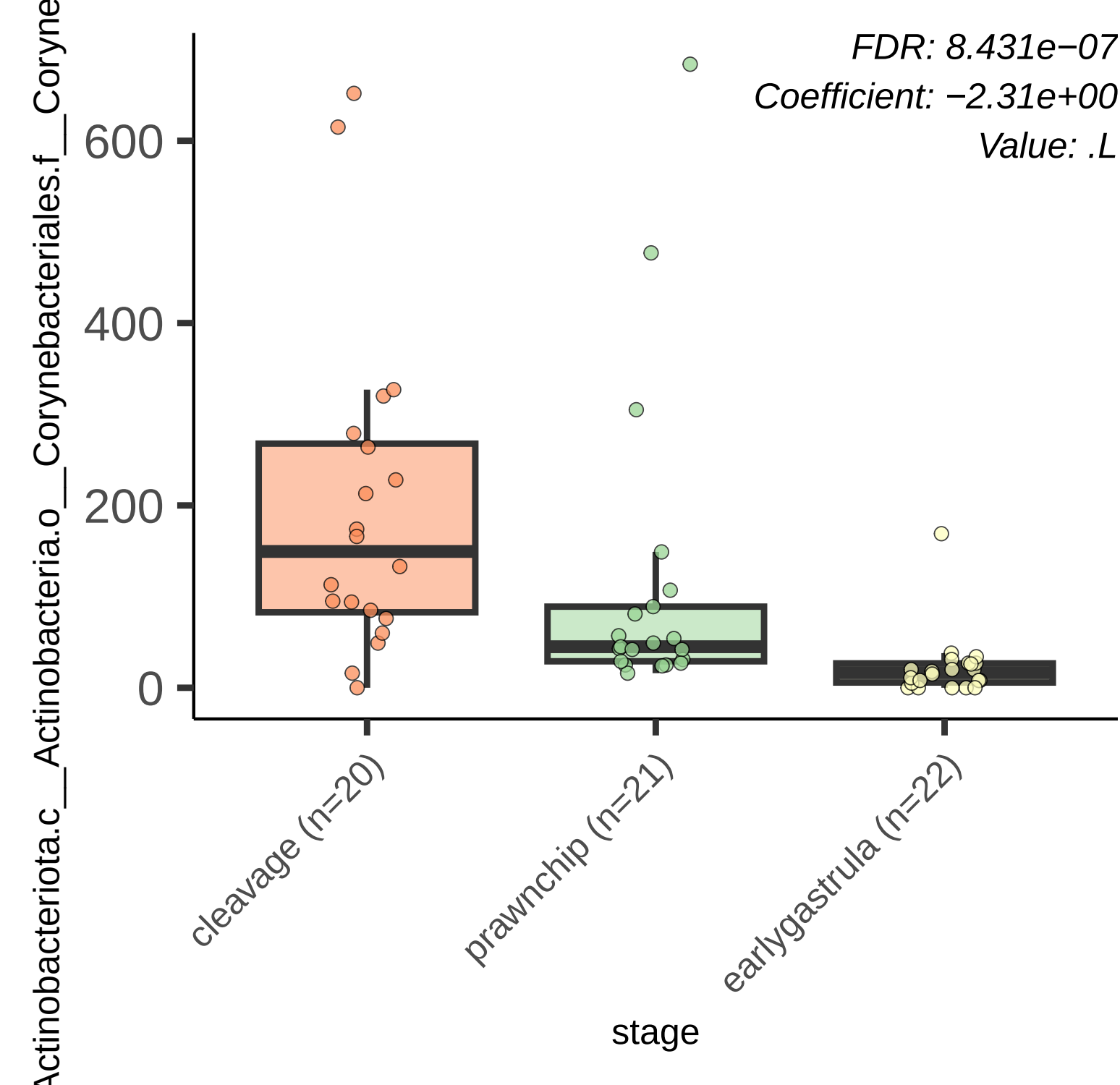
0
cleavage (n=20)
prawnchip (n=21)
earlygastrula (n=22)
stage
oteobacteria.c__Gammaproteobacteria.o__Oceanospirillales.f__Oce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200
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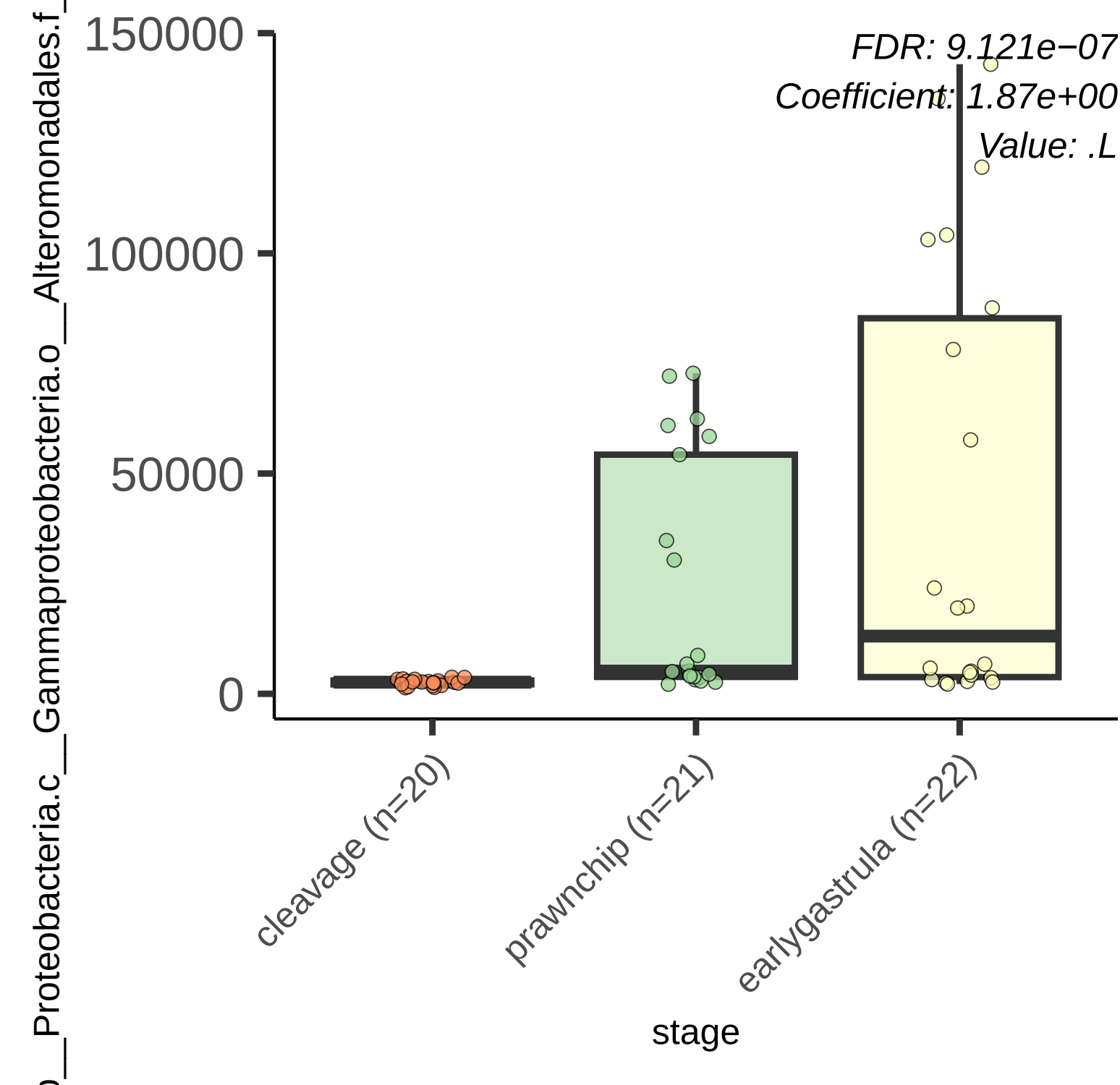
prawnchip (n=21)
earlygastrula (n=22)
stage
Actinobacteriota.c__Actinobacteria.o__Corynebacteriales.f__Coryne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50000
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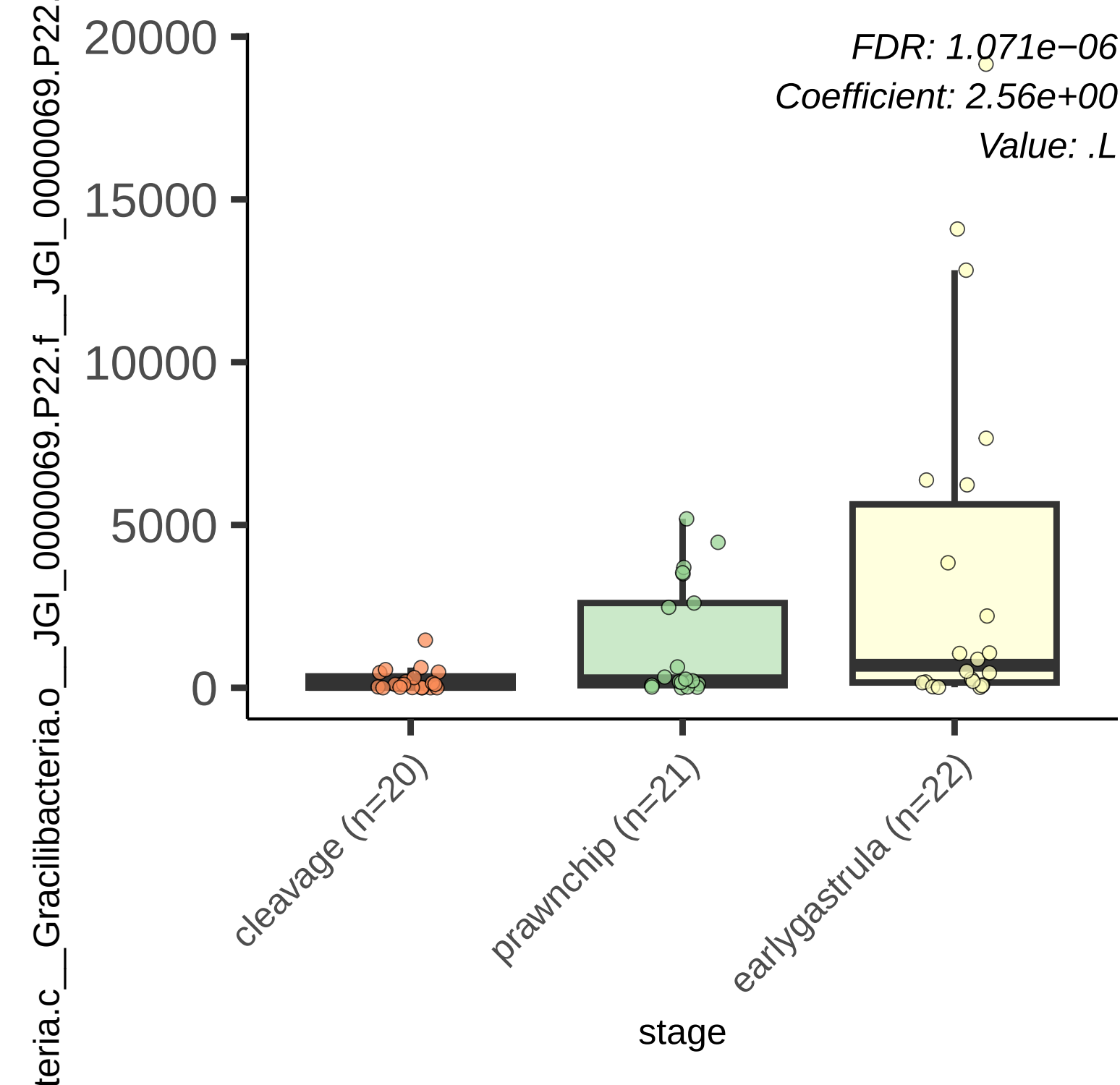
prawnchip (n=21)
earlygastrula (n=22)
stage
o__Proteobacteria.c__Gammaproteobacteria.o__Alteromonadales.f__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15000
10000
5000
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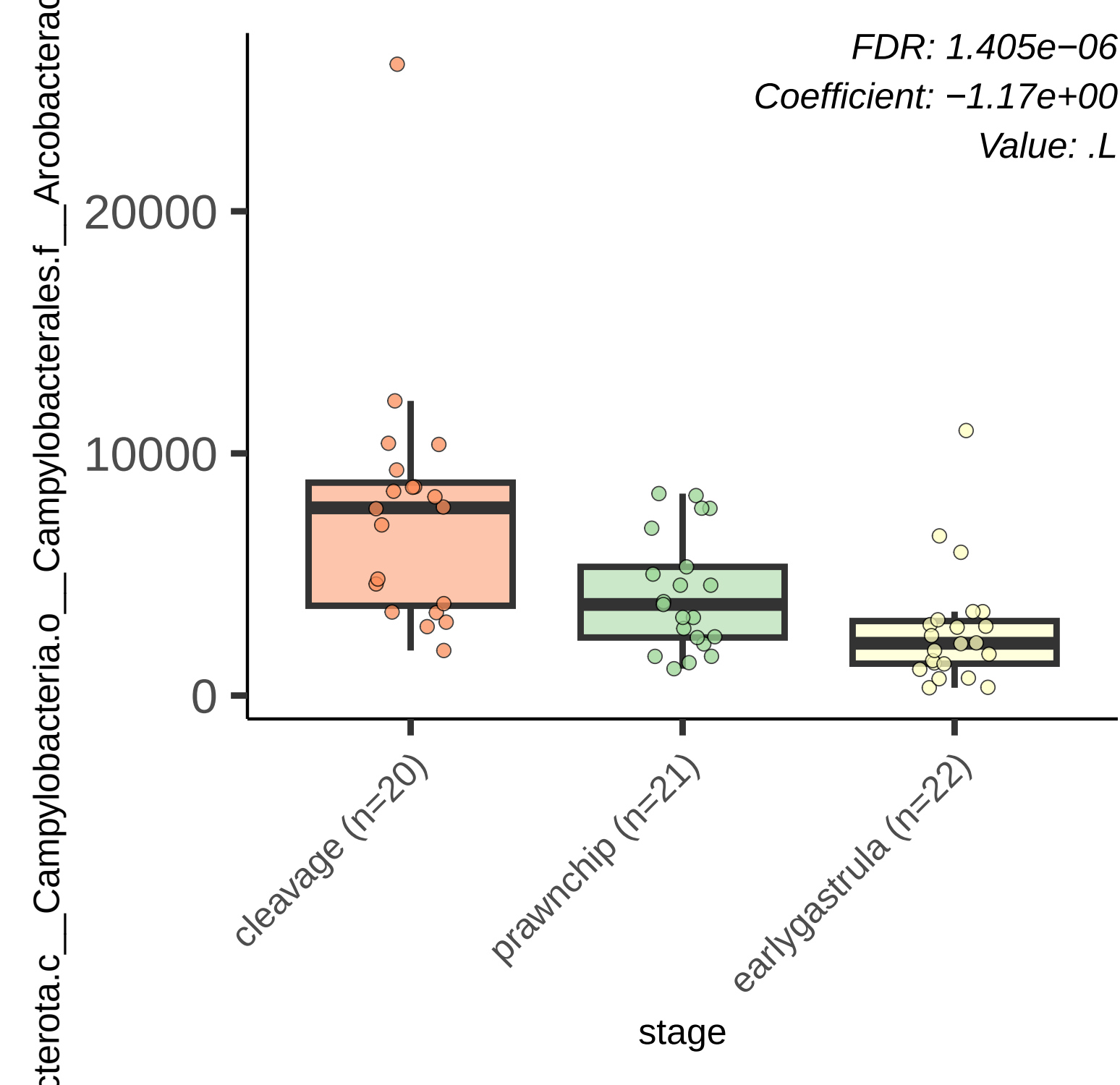
prawnchip (n=21)
earlygastrula (n=22)
stage
teria.c__Gracilibacteria.o__JGI_0000069.P22.f__JGI_0000069.P22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10000
0
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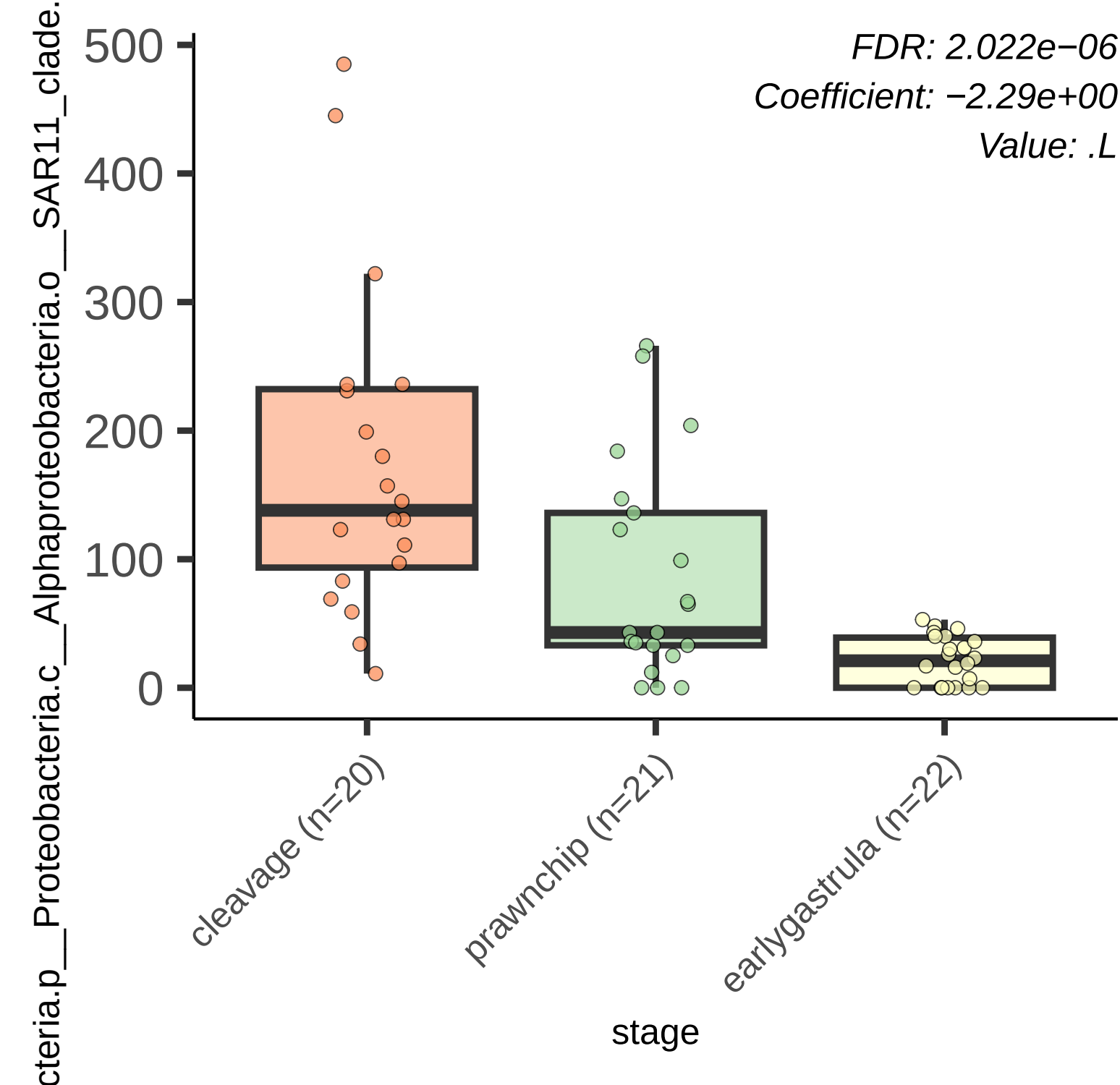
earlygastrula (n=22)
stage
cterota.c__Campylobacteria.o__Campylobacterales.f__Arcobacterac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200
100
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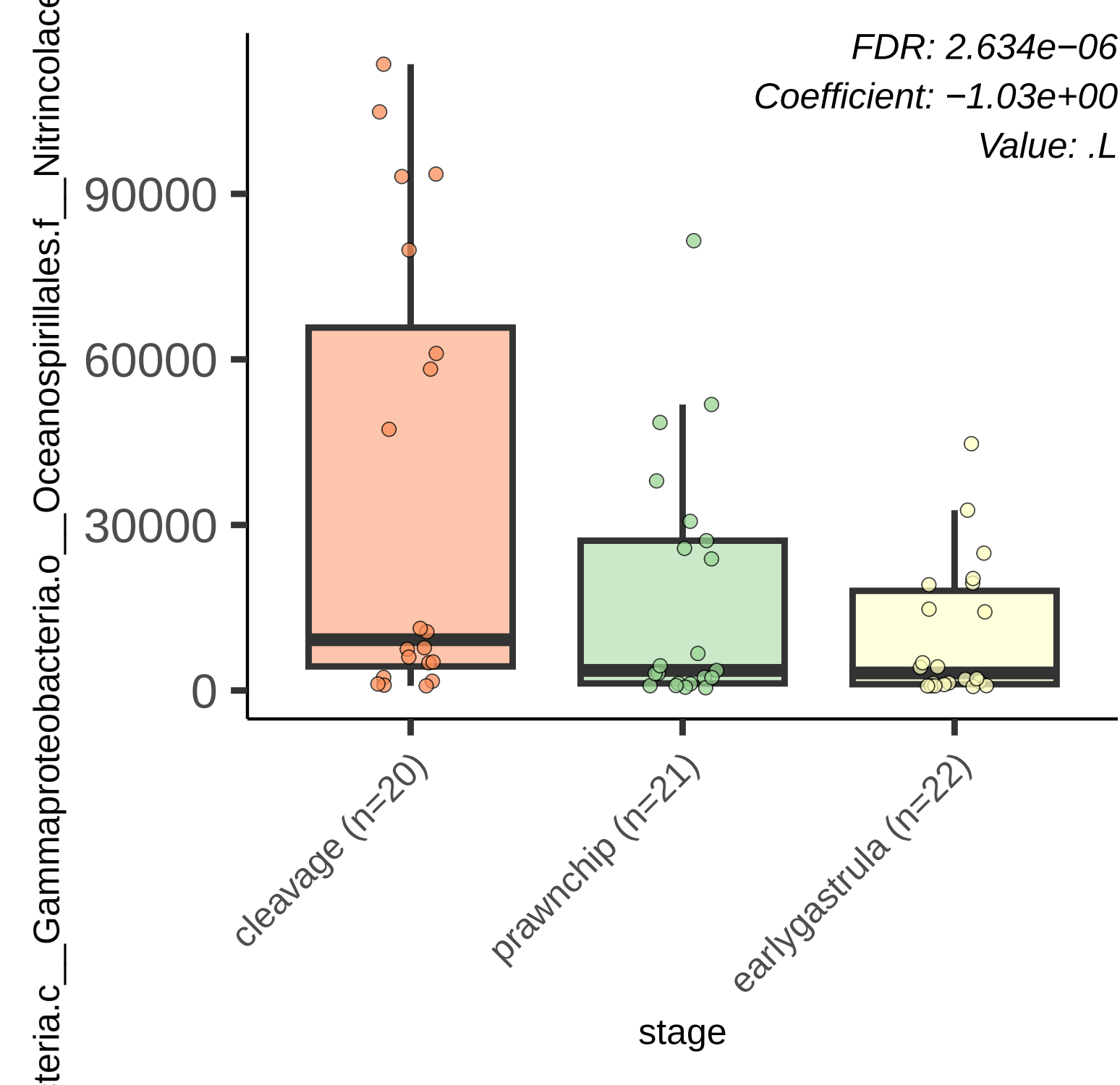
earlygastrula (n=22)
stage
cteria.p__Proteobacteria.c__Alphaproteobacteria.o__SAR11_clade.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60000
30000
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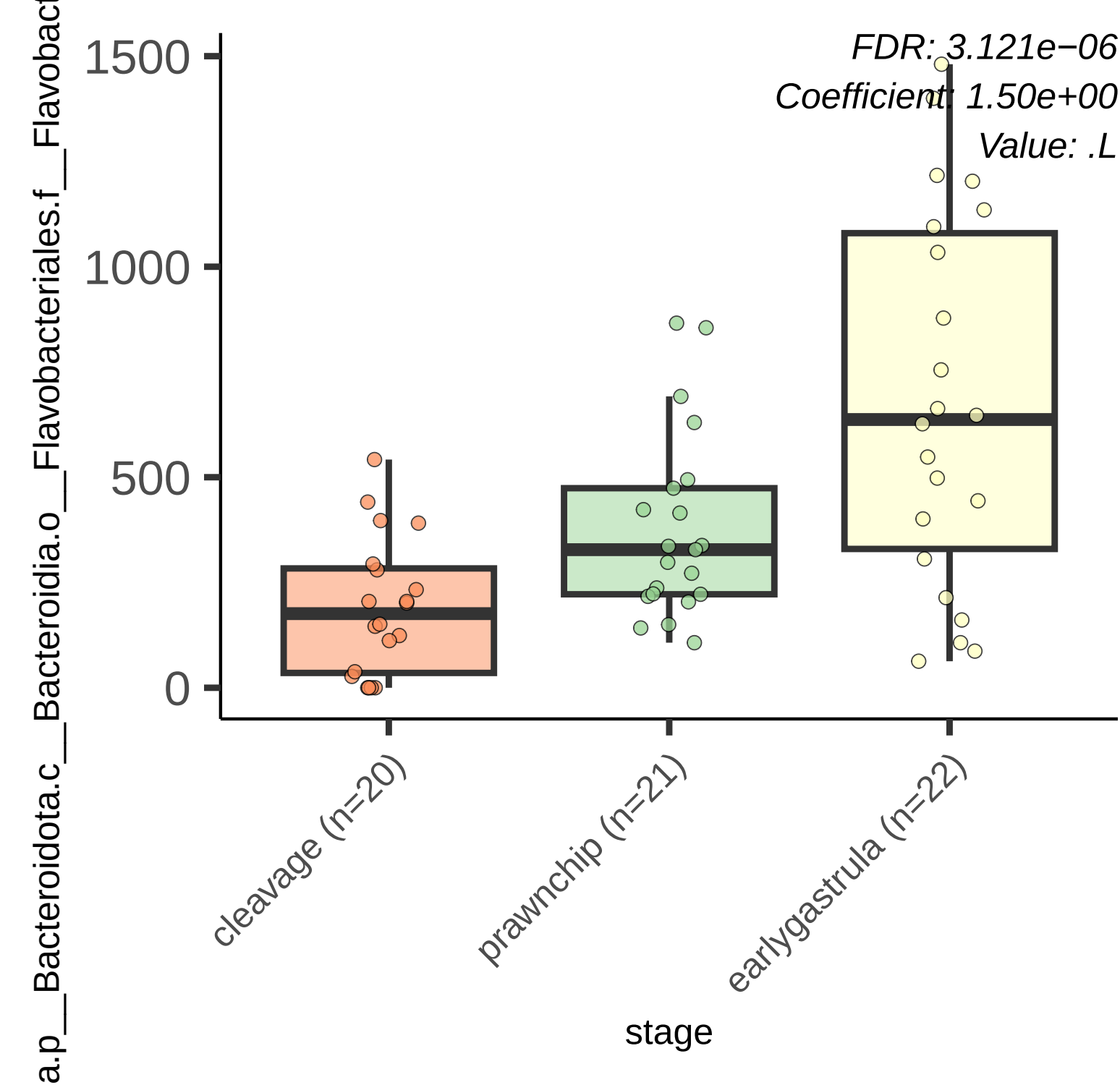
cleavage (n=20)
prawnchip (n=21)
earlygastrula (n=22)
stage
teria.c__Gammaproteobacteria.o__Oceanospirillales.f__Nitrincolace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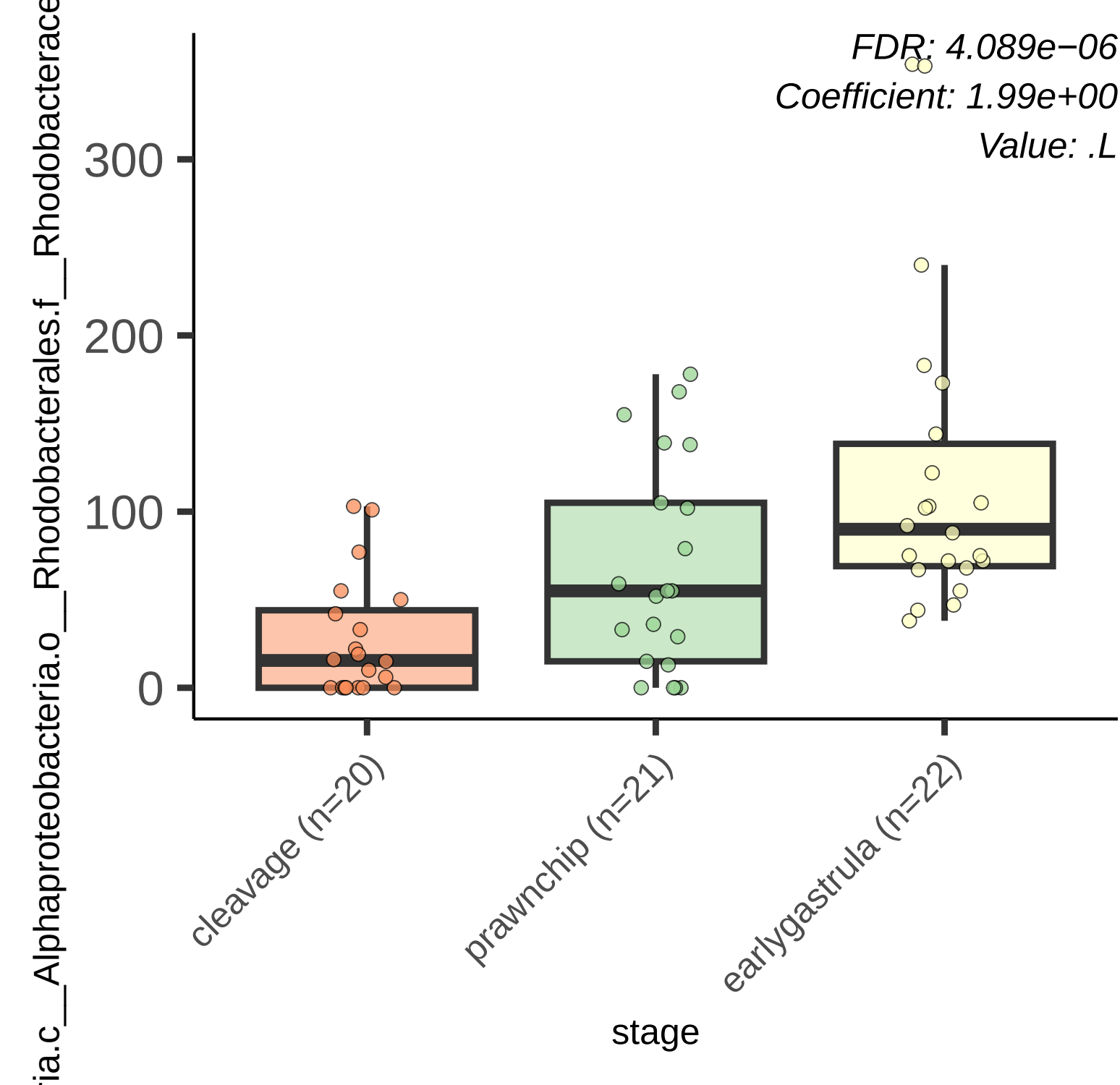
1000
500
0
cleavage (n=20)
prawnchip (n=21)
earlygastrula (n=22)
stage
a.p__Bacteroidota.c__Bacteroidia.o__Flavobacteriales.f__Flavobact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Coefficient: 1.99e+00
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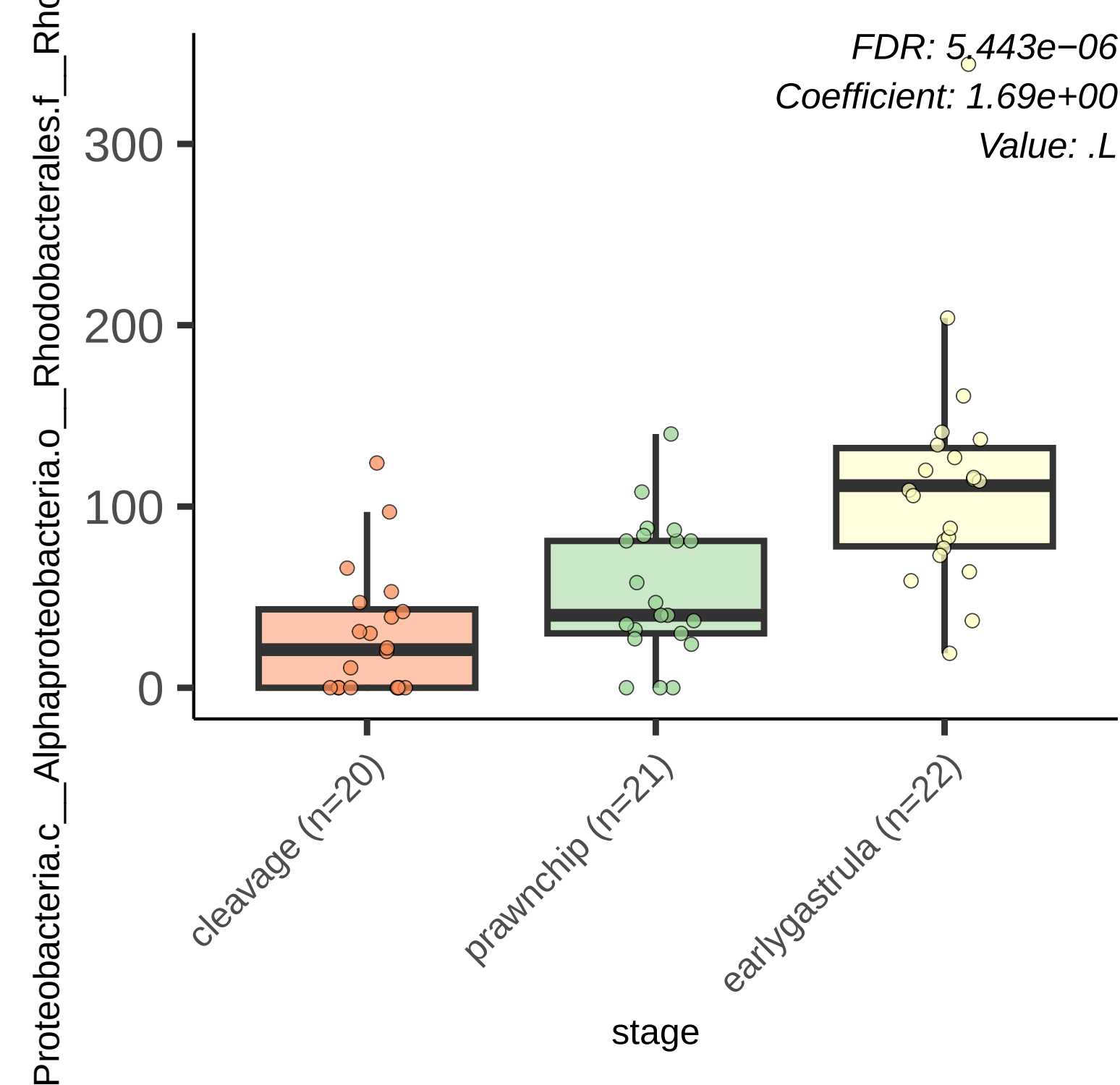
300
200
100
0
cleavage (n=20)
prawnchip (n=21)
earlygastrula (n=22)
stage
ia.c__Alphaproteobacteria.o__Rhodobacterales.f__Rhodobacterace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100
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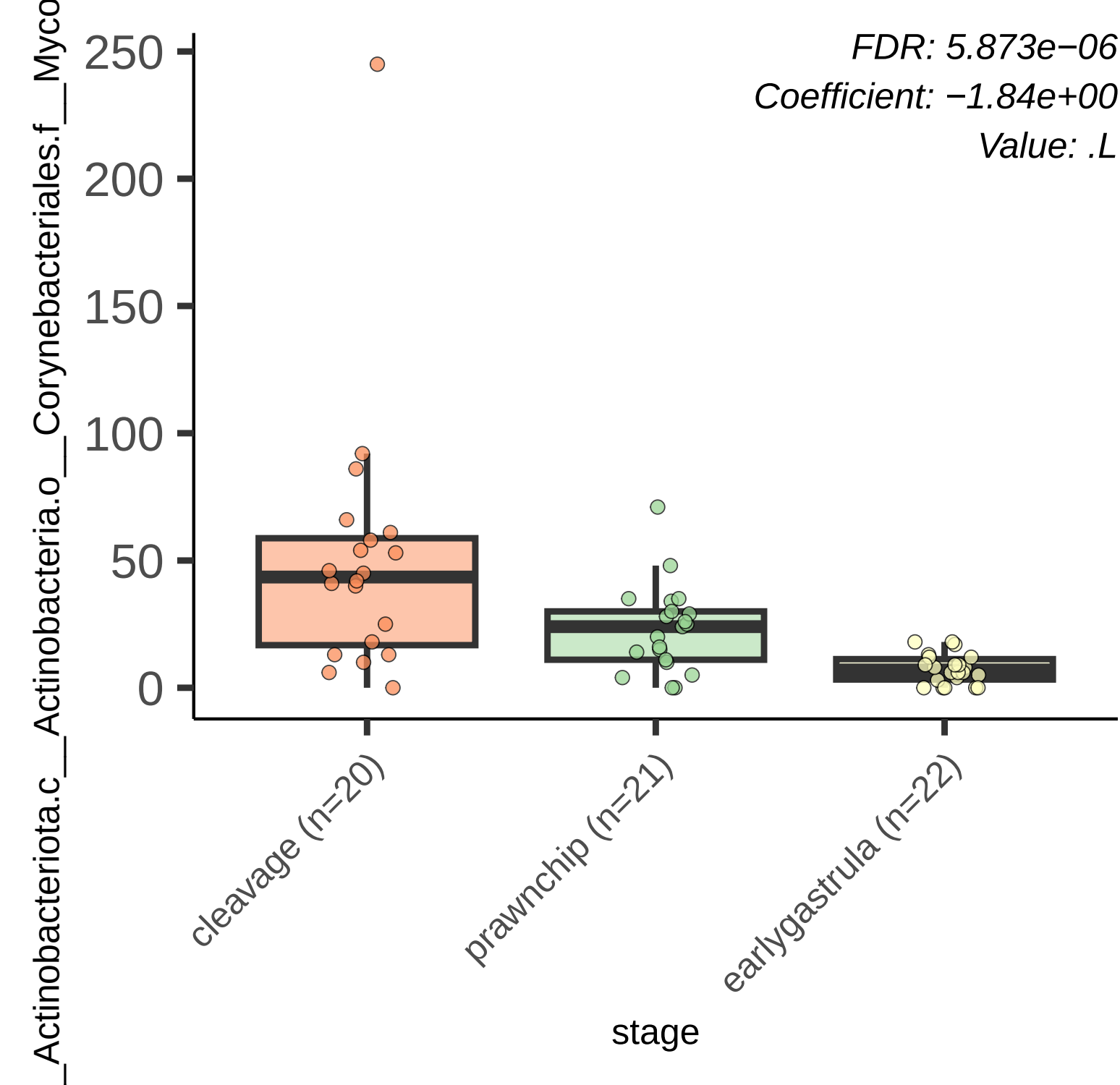
prawnchip (n=21)
earlygastrula (n=22)
stage
Proteobacteria.c__Alphaproteobacteria.o__Rhodobacterales.f__Rho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100
50
0
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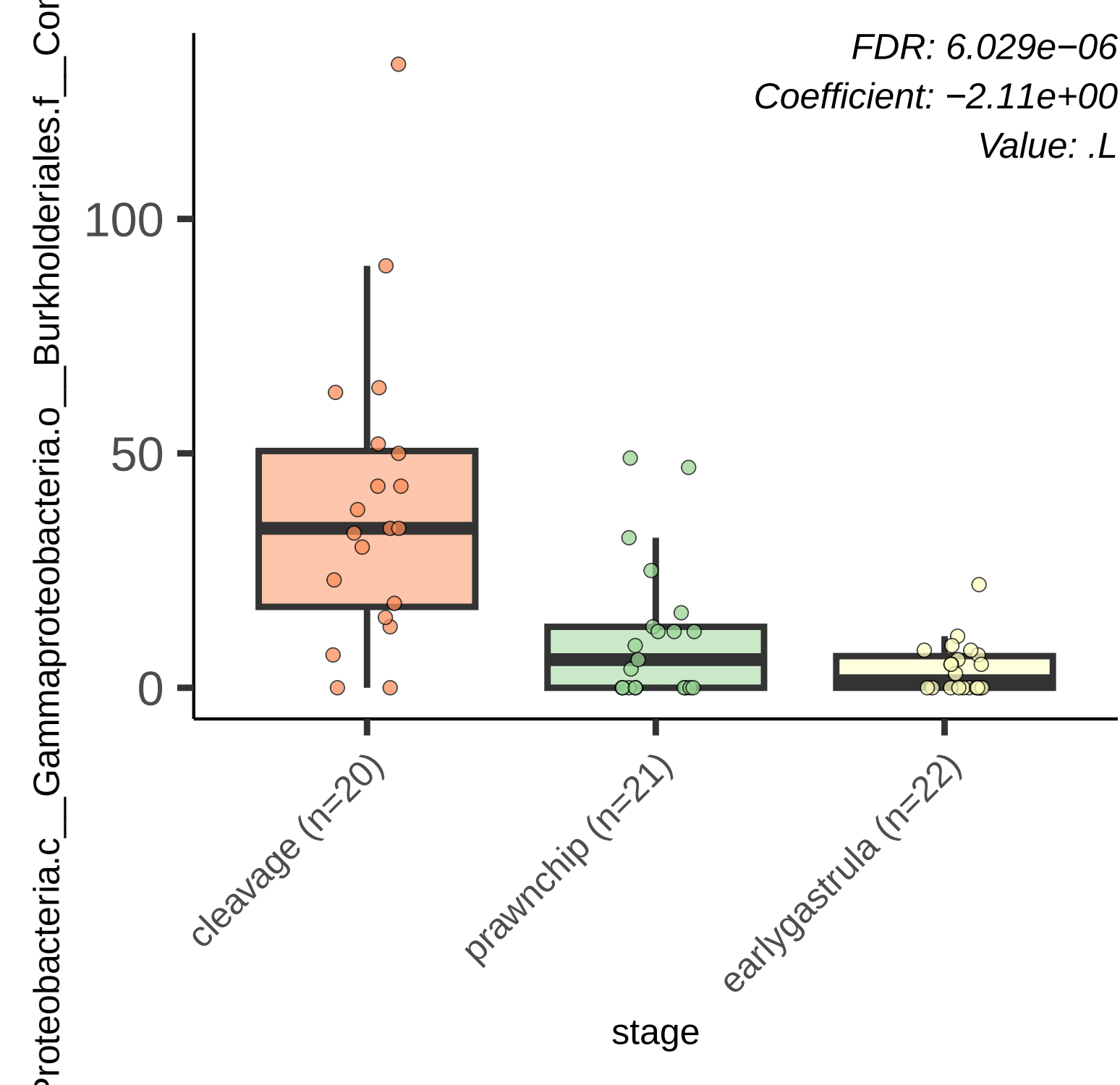
earlygastrula (n=22)
stage
_Actinobacteriota.c__Actinobacteria.o__Corynebacteriales.f__Myco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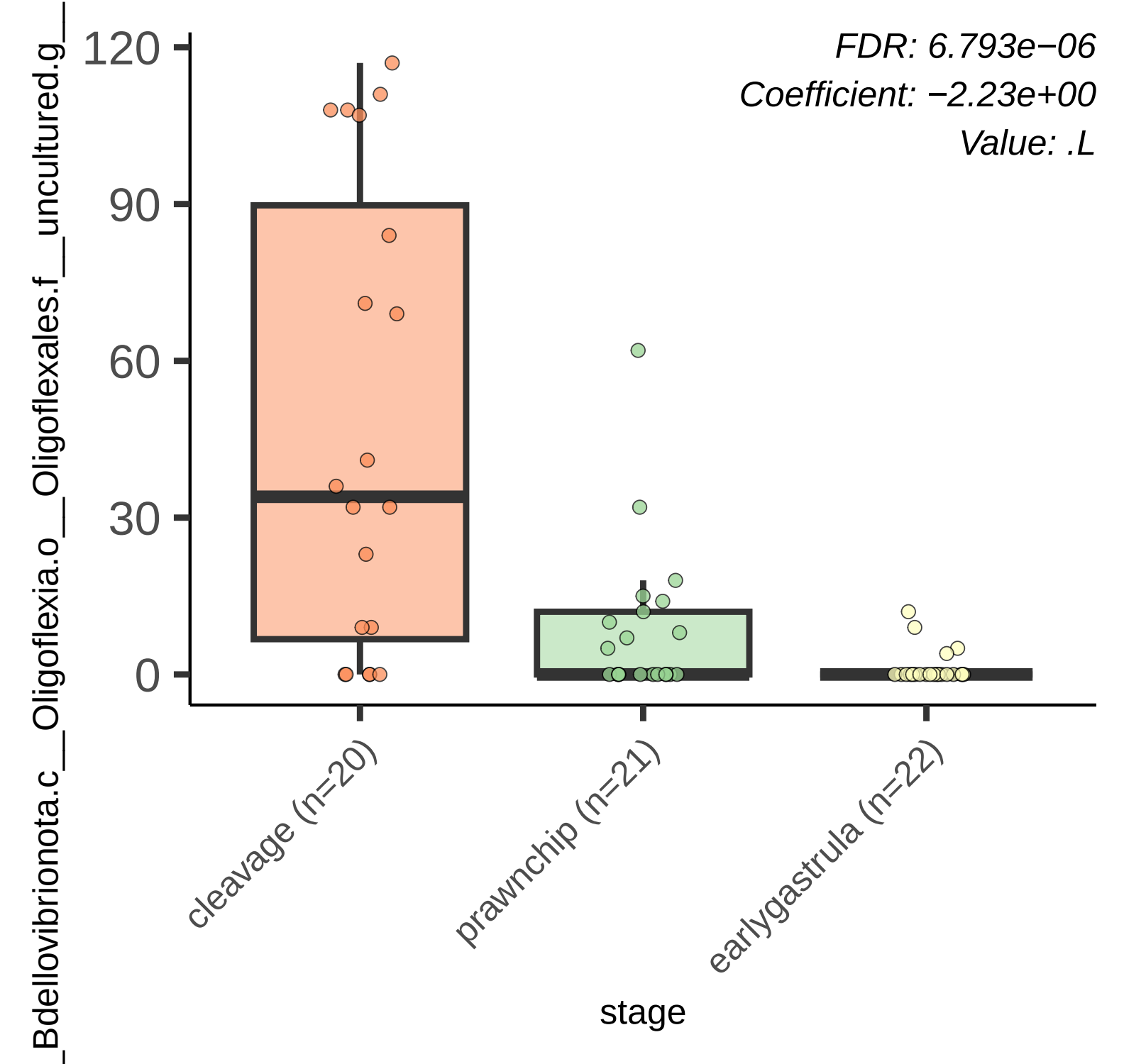
50
0
cleavage (n=20)
prawnchip (n=21)
earlygastrula (n=22)
stage
Proteobacteria.c__Gammaproteobacteria.o__Burkholderiales.f__Com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60
30
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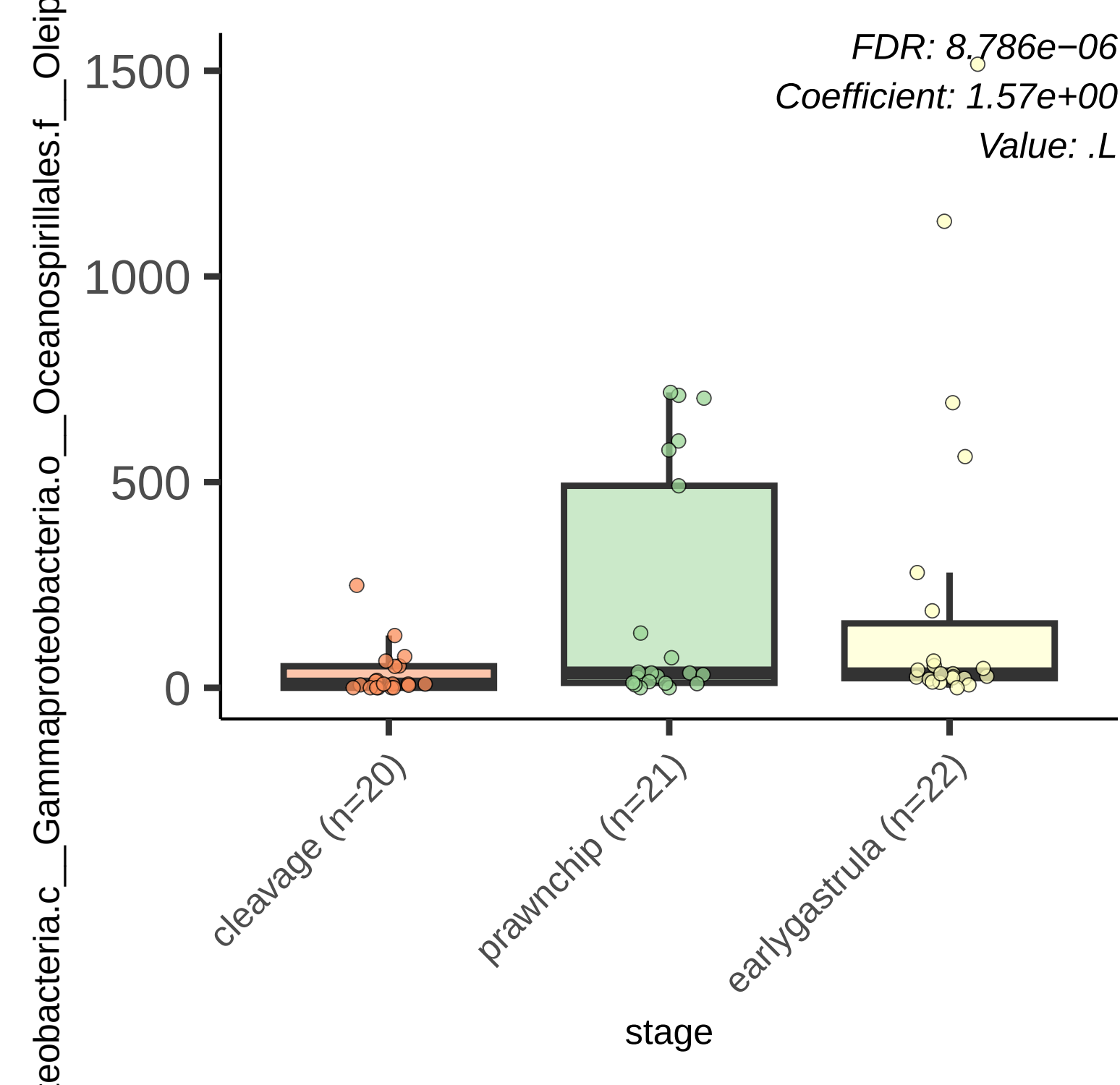
cleavage (n=20)
prawnchip (n=21)
earlygastrula (n=22)
stage
_Bdellovibrionota.c__Oligoflexia.o__Oligoflexales.f__uncultured.g__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500
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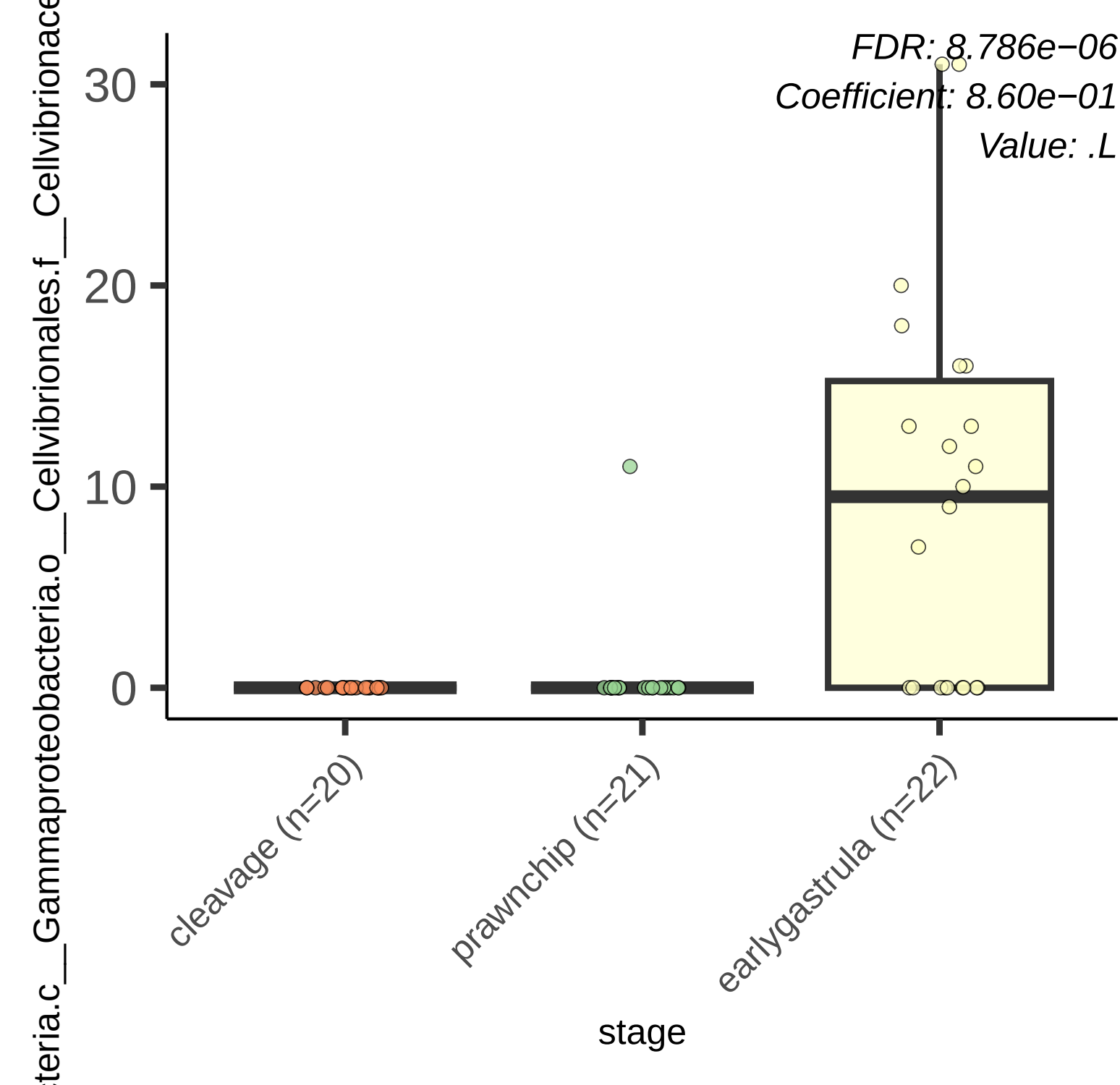
prawnchip (n=21)
earlygastrula (n=22)
stage
teobacteria.c__Gammaproteobacteria.o__Oceanospirillales.f__Oleip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10
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prawnchip (n=21)
earlygastrula (n=22)
stage
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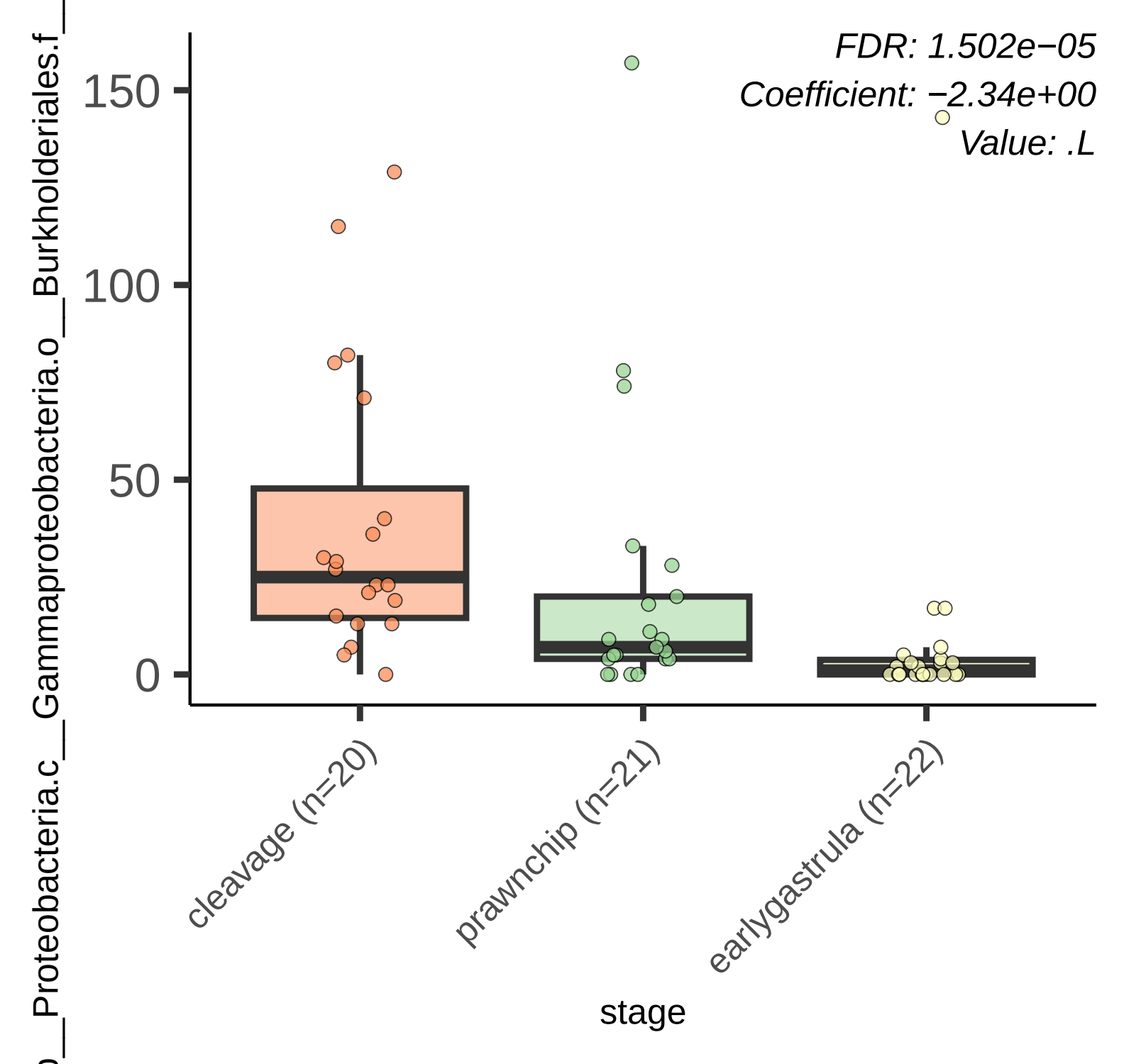
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100
50
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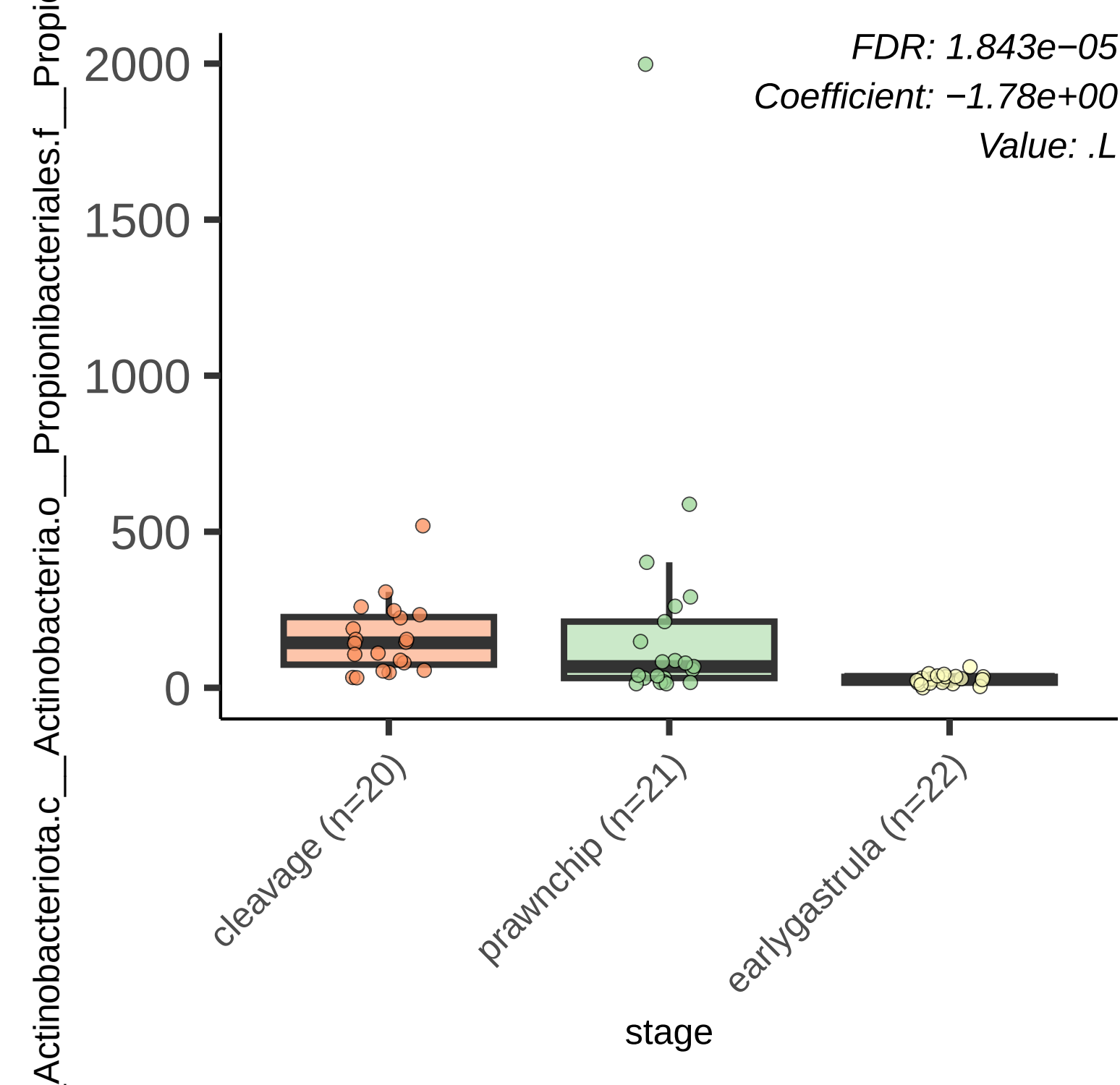
cleavage (n=20)
prawnchip (n=21)
earlygastrula (n=22)
stage
Proteobacteria.c__Gammaproteobacteria.o__Burkholderiales.f__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1000
500
0
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prawnchip (n=21)
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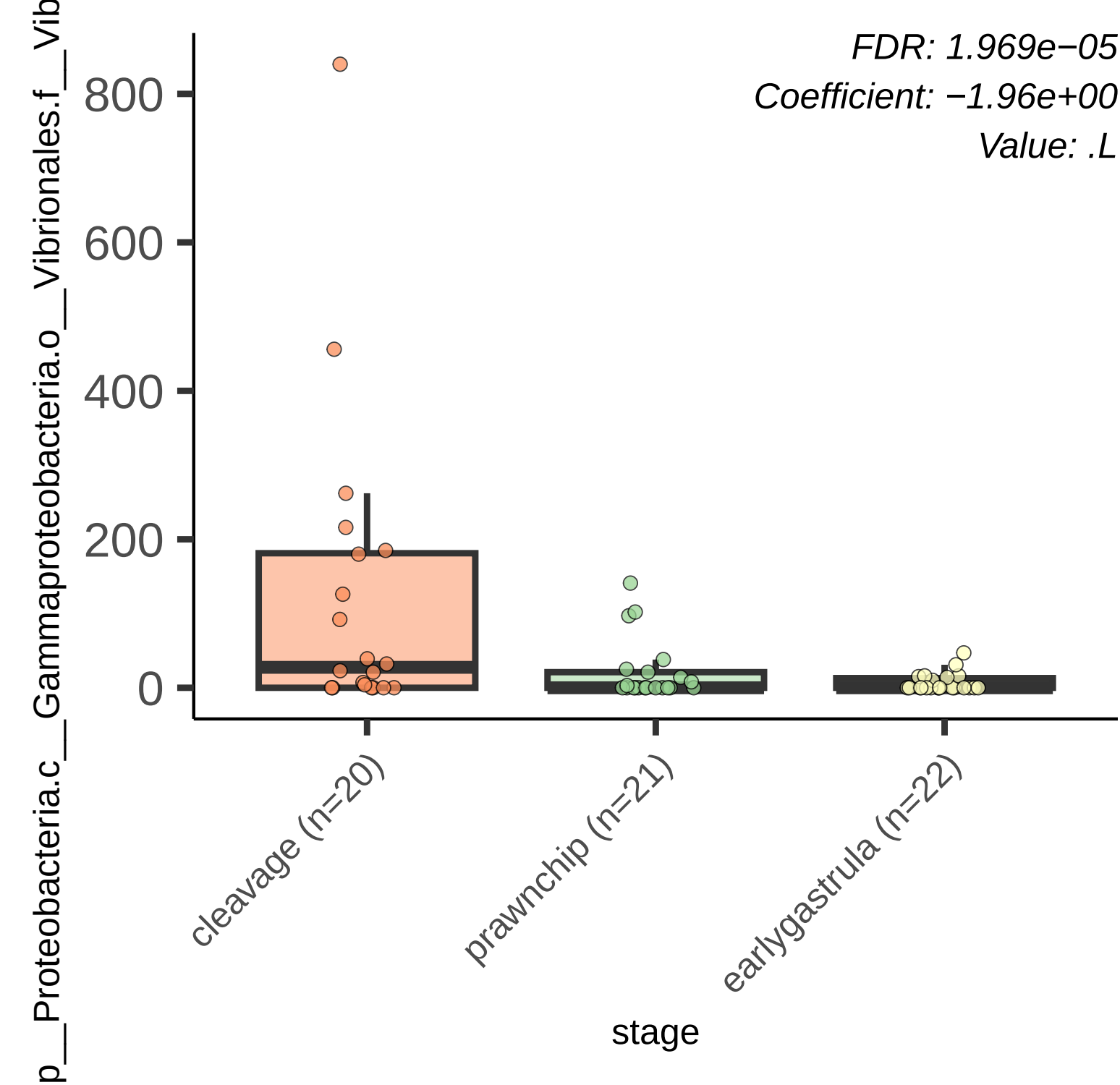
stage
Actinobacteriota.c__Actinobacteria.o__Propionibacteriales.f__Propi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400
200
0
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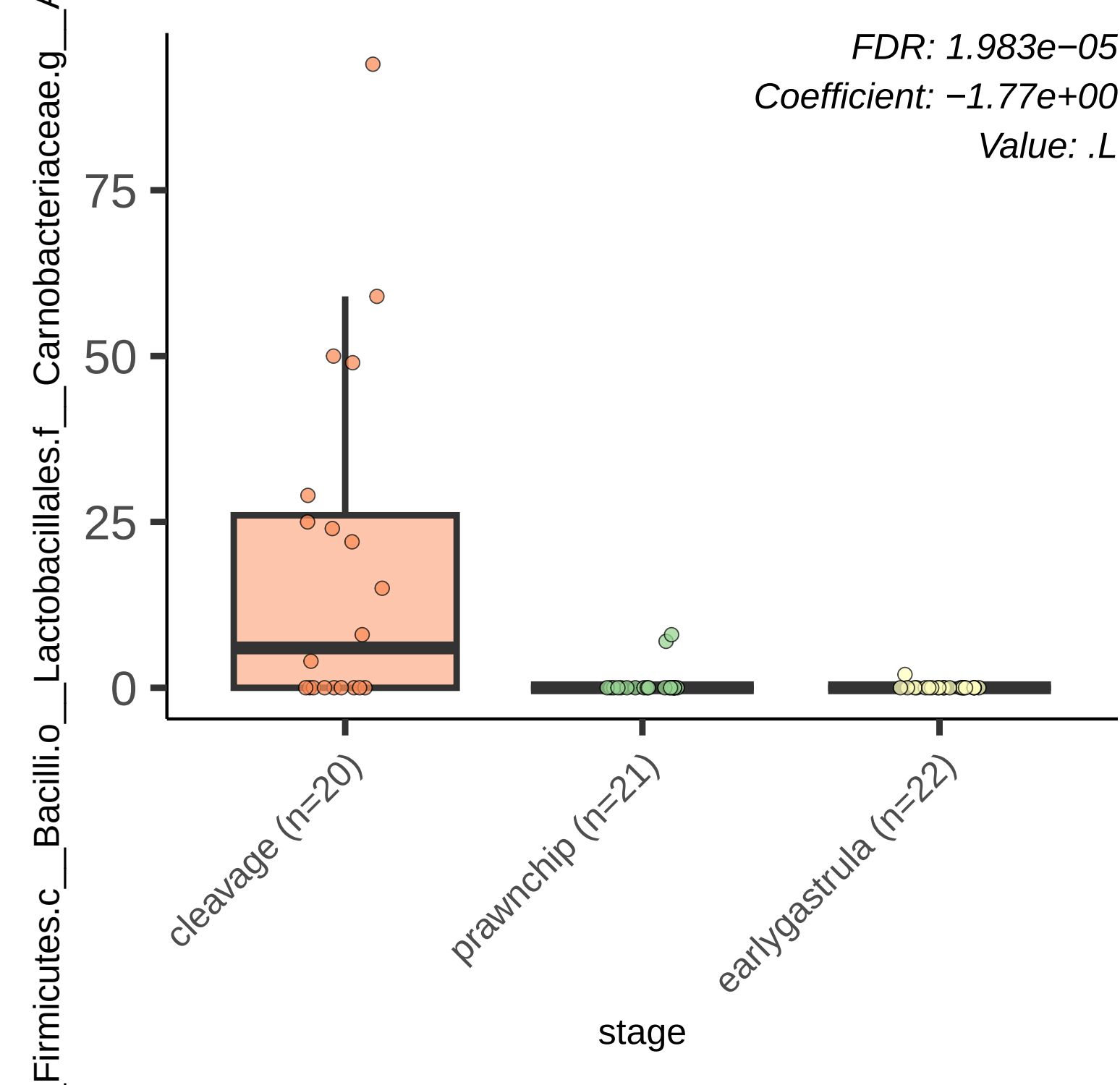
prawnchip (n=21)
earlygastrula (n=22)
stage
p__Proteobacteria.c__Gammaproteobacteria.o__Vibrionales.f__Vib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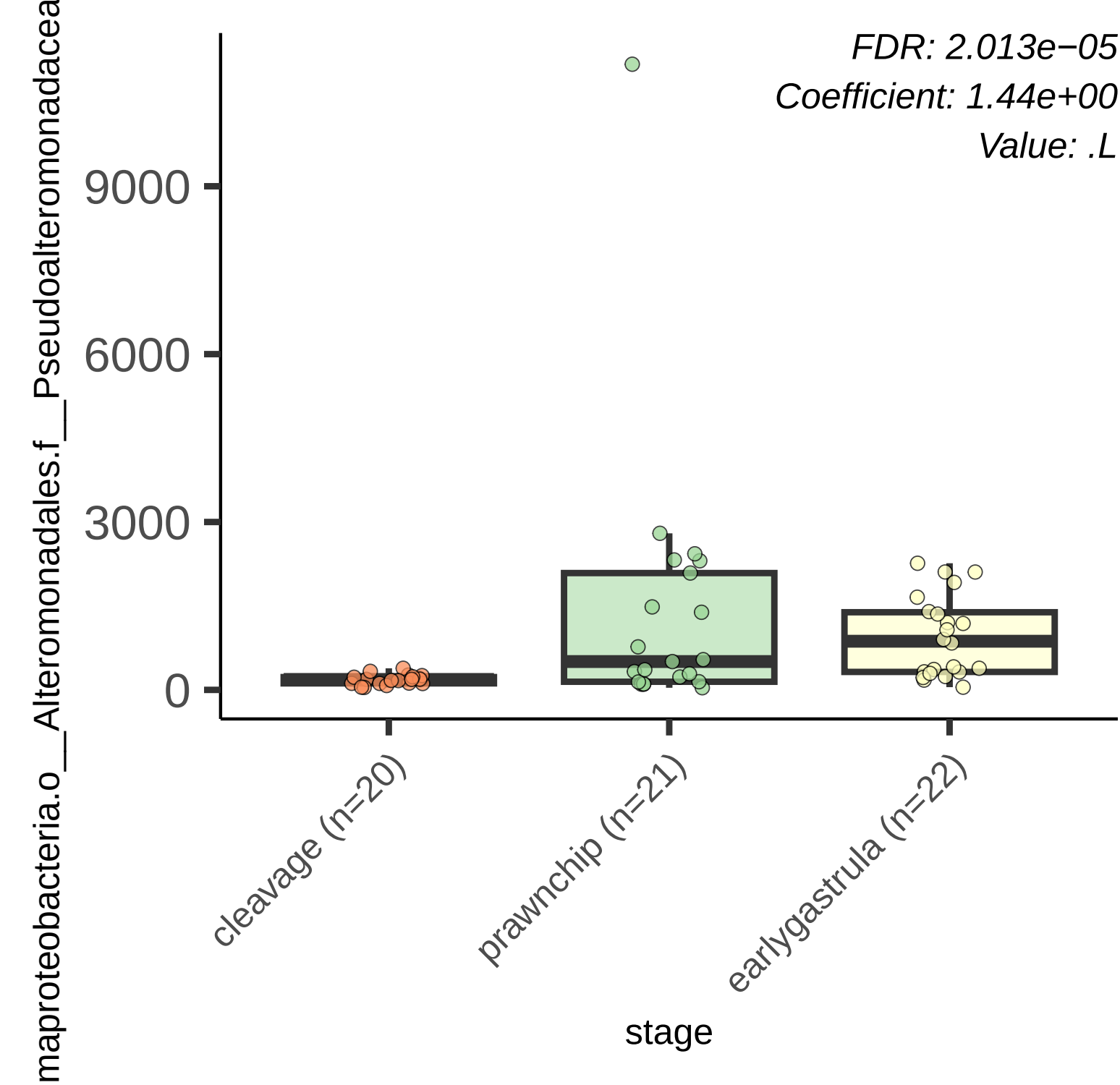
50
25
0
cleavage (n=20)
prawnchip (n=21)
earlygastrula (n=22)
stage
Firmicutes.c__Bacilli.o__Lactobacillales.f__Carnobacteriaceae.g__A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6000
3000
0
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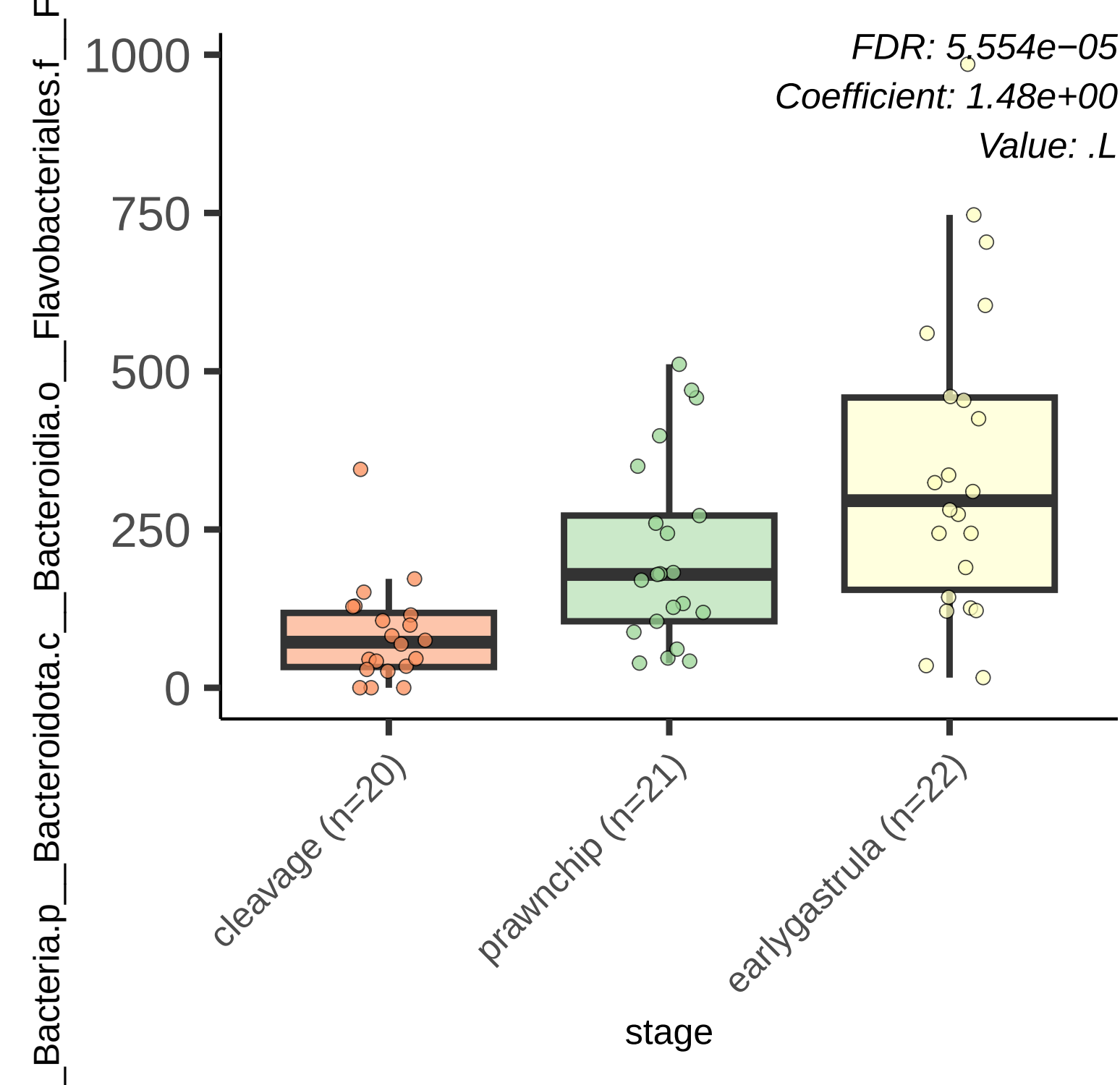
prawnchip (n=21)
earlygastrula (n=22)
stage
maproteobacteria.o__Alteromonadales.f__Pseudoalteromonadacea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Coefficient: 1.48e+00
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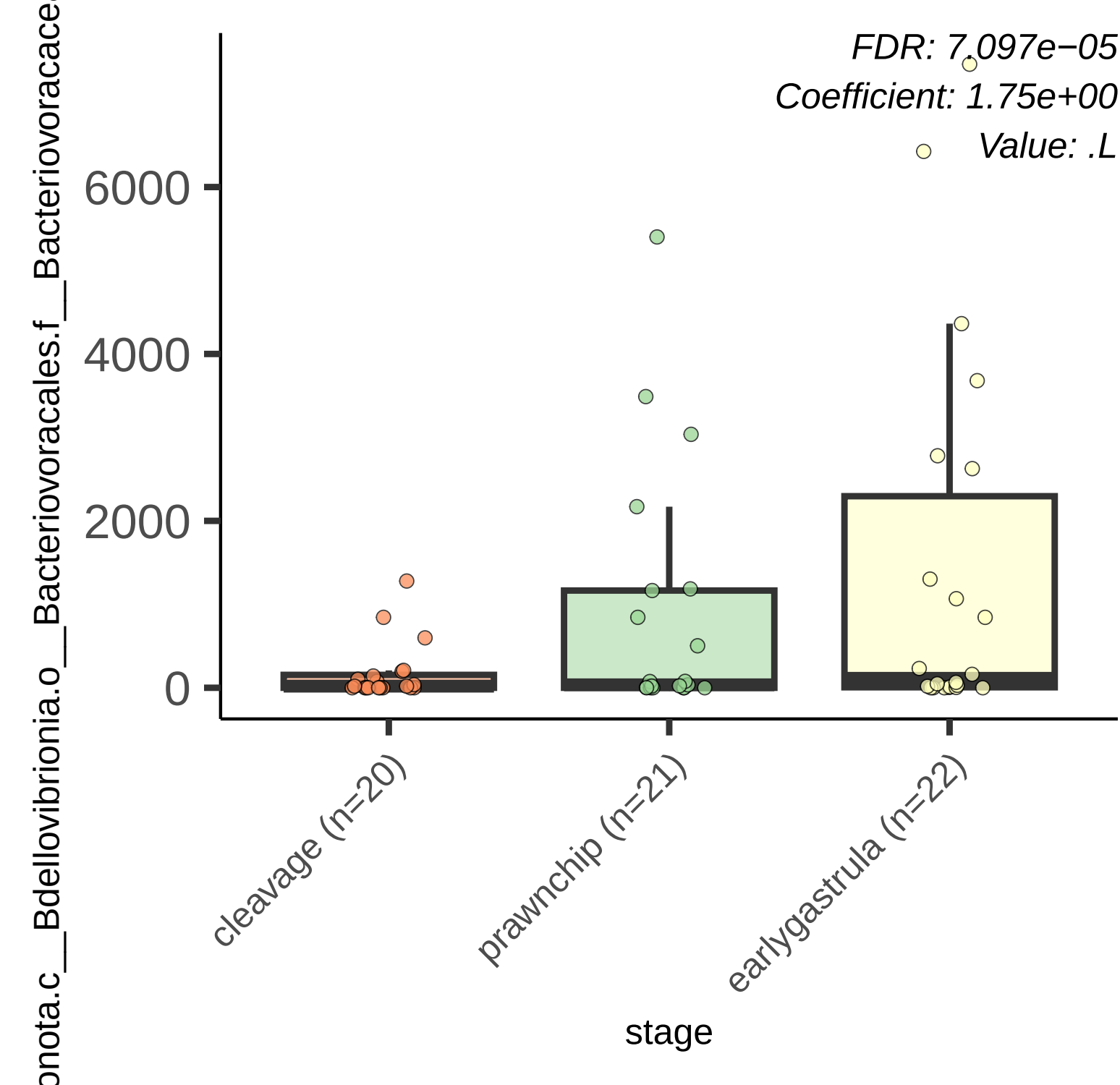
1000
750
500
250
0
cleavage (n=20)
prawnchip (n=21)
earlygastrula (n=22)
stage
Bacteria.p__Bacteroidota.c__Bacteroidia.o__Flavobacteriales.f__F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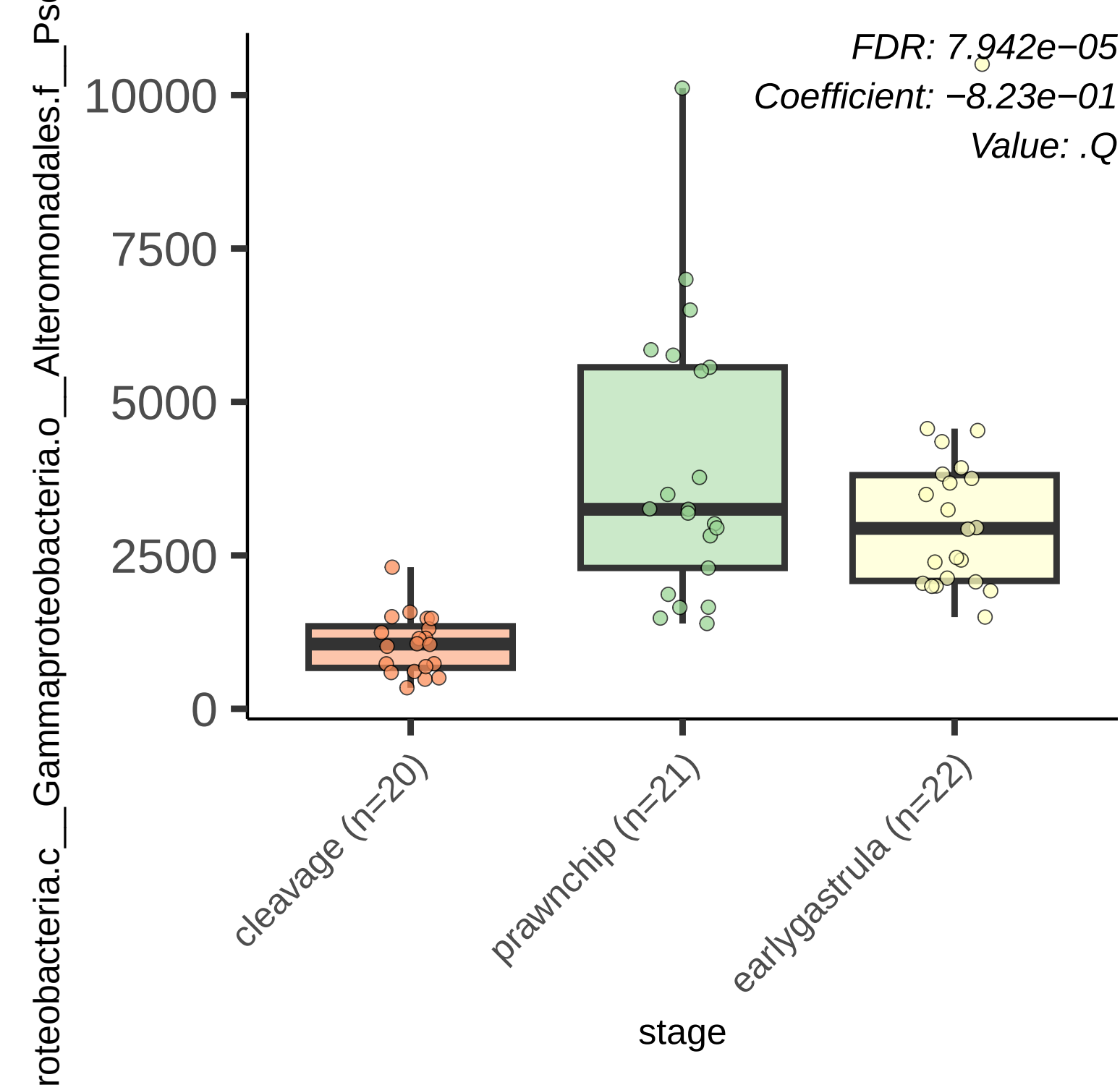
4000
2000
0
cleavage (n=20)
prawnchip (n=21)
earlygastrula (n=22)
stage
onota.c__Bdellovibrionia.o__Bacteriovoracales.f__Bacteriovoracace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Coefficient: −8.23e−01
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5000
2500
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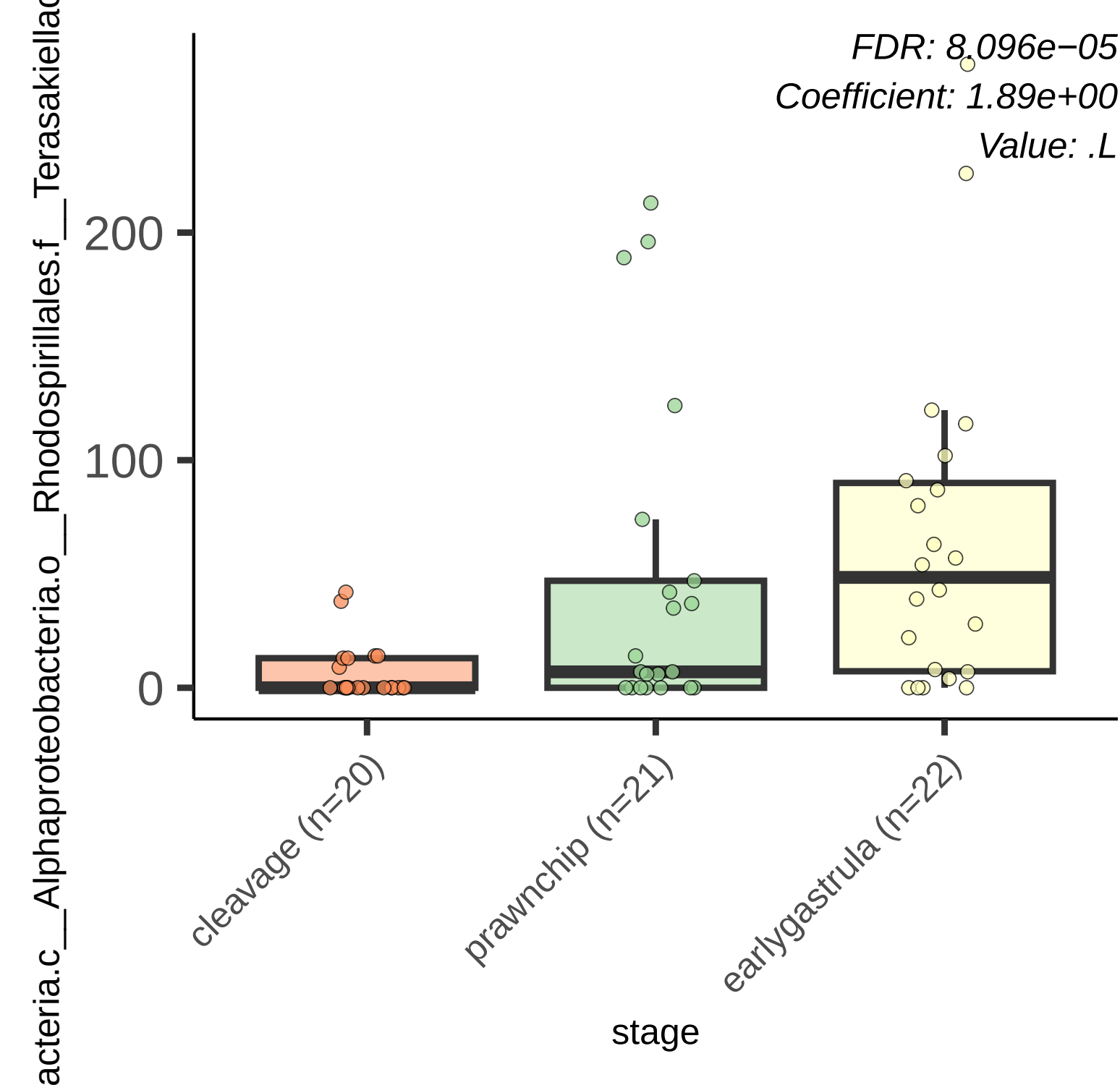
prawnchip (n=21)
earlygastrula (n=22)
stage
roteobacteria.c__Gammaproteobacteria.o__Alteromonadales.f__Ps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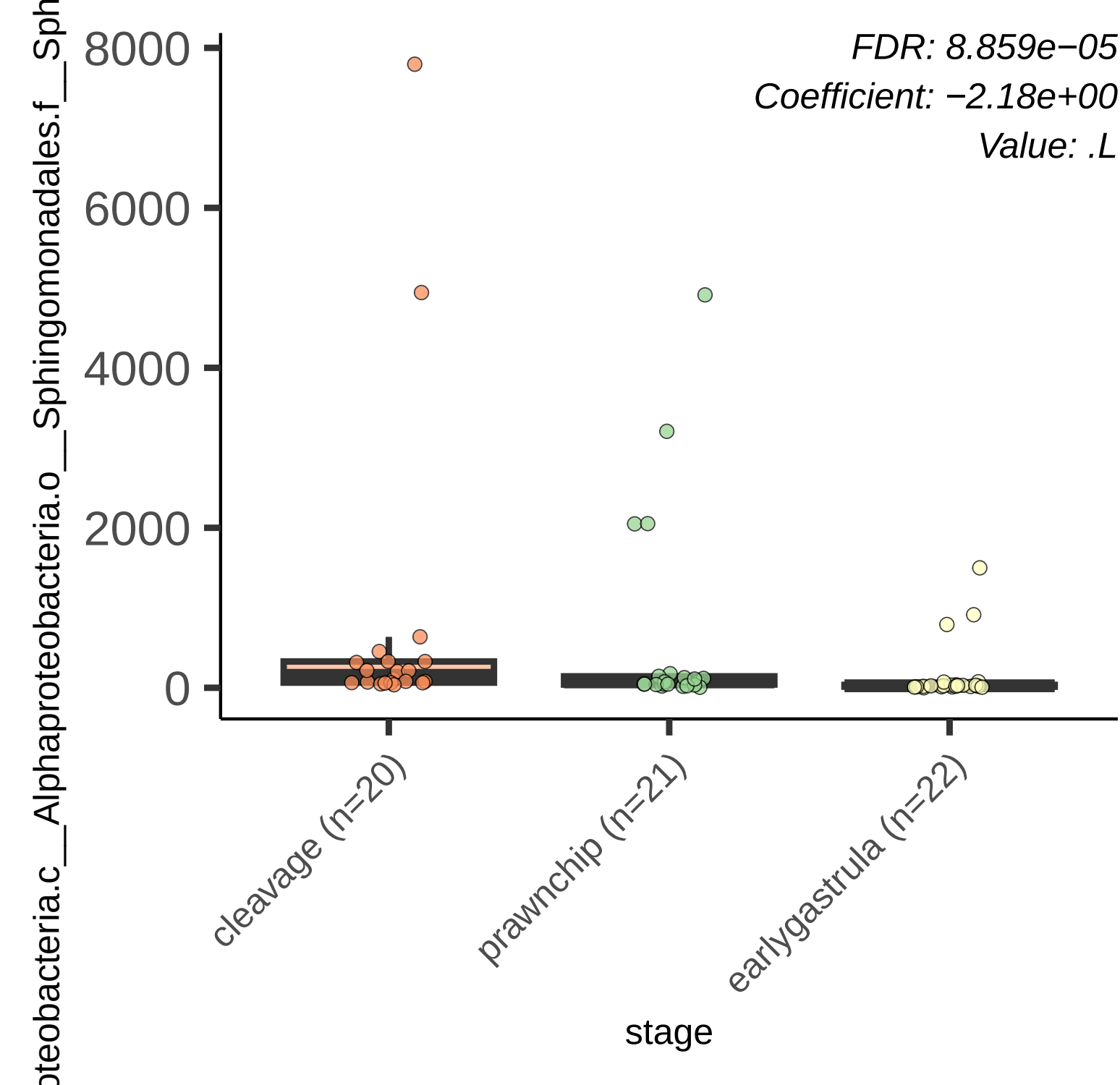
100
0
cleavage (n=20)
prawnchip (n=21)
earlygastrula (n=22)
stage
acteria.c__Alphaproteobacteria.o__Rhodospirillales.f__Terasakiellac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4000
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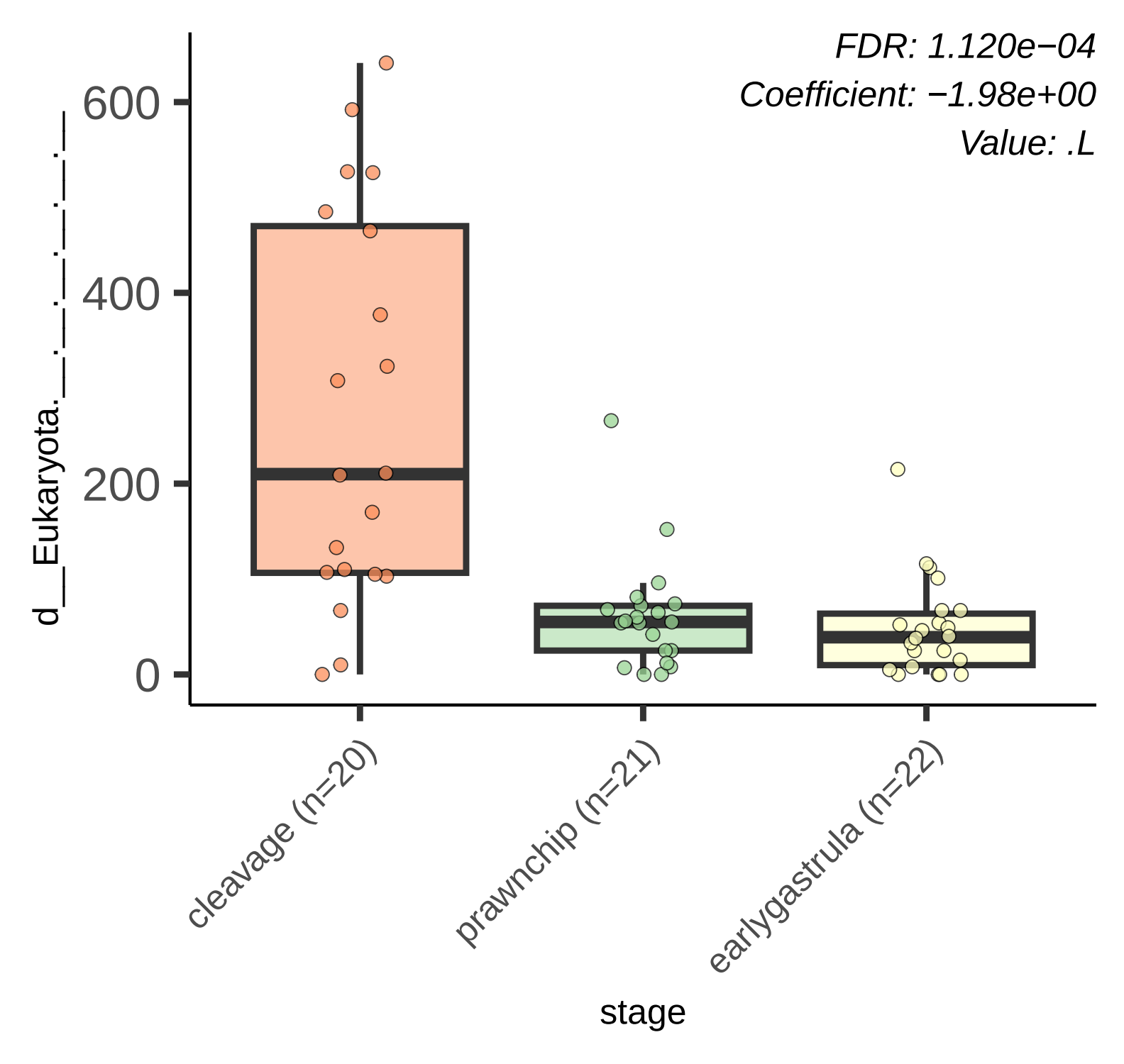
earlygastrula (n=22)
stage
oteobacteria.c__Alphaproteobacteria.o__Sphingomonadales.f__Sph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400
200
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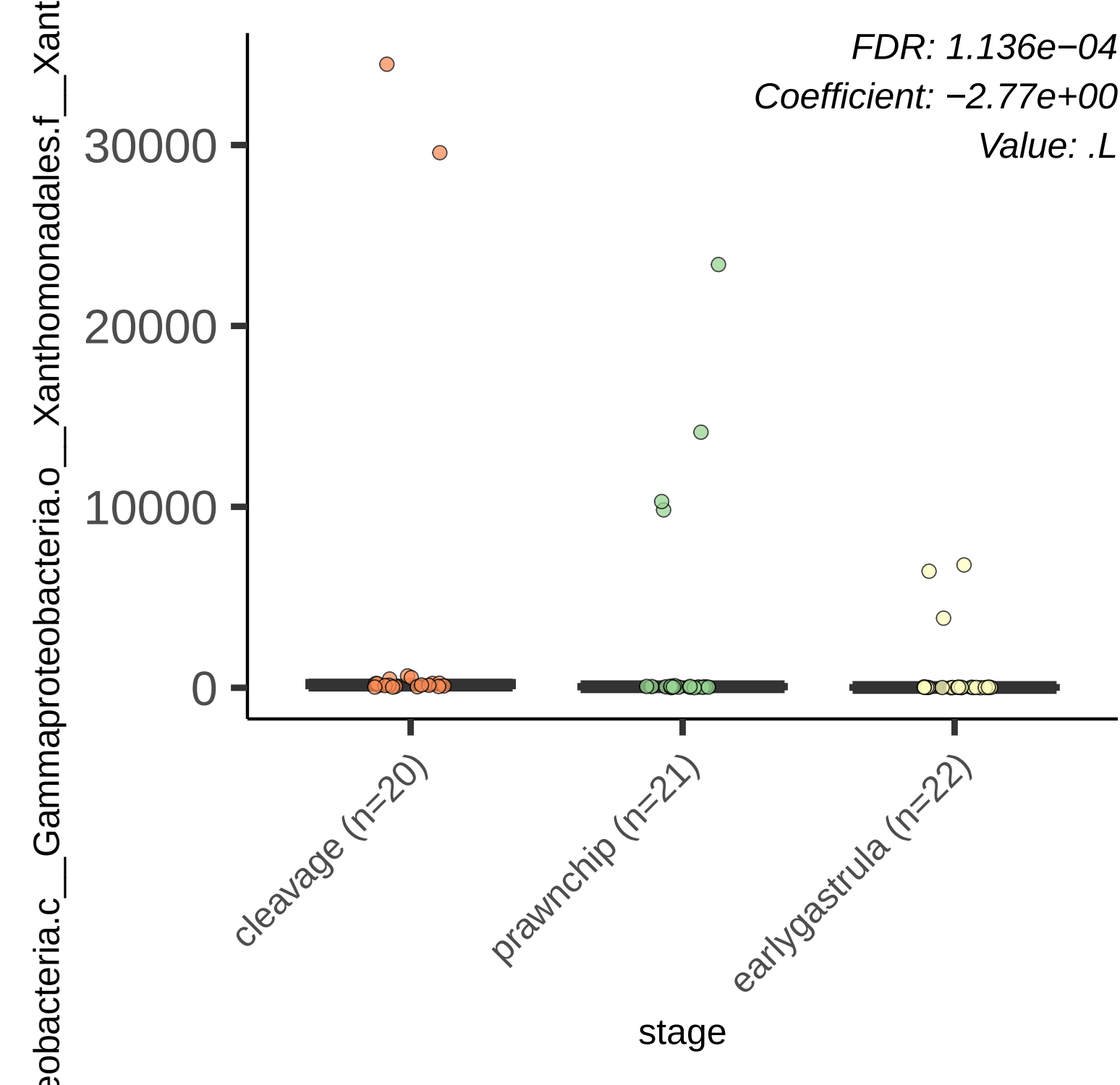
cleavage (n=20)
prawnchip (n=21)
earlygastrula (n=22)
stage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30000
20000
10000
0
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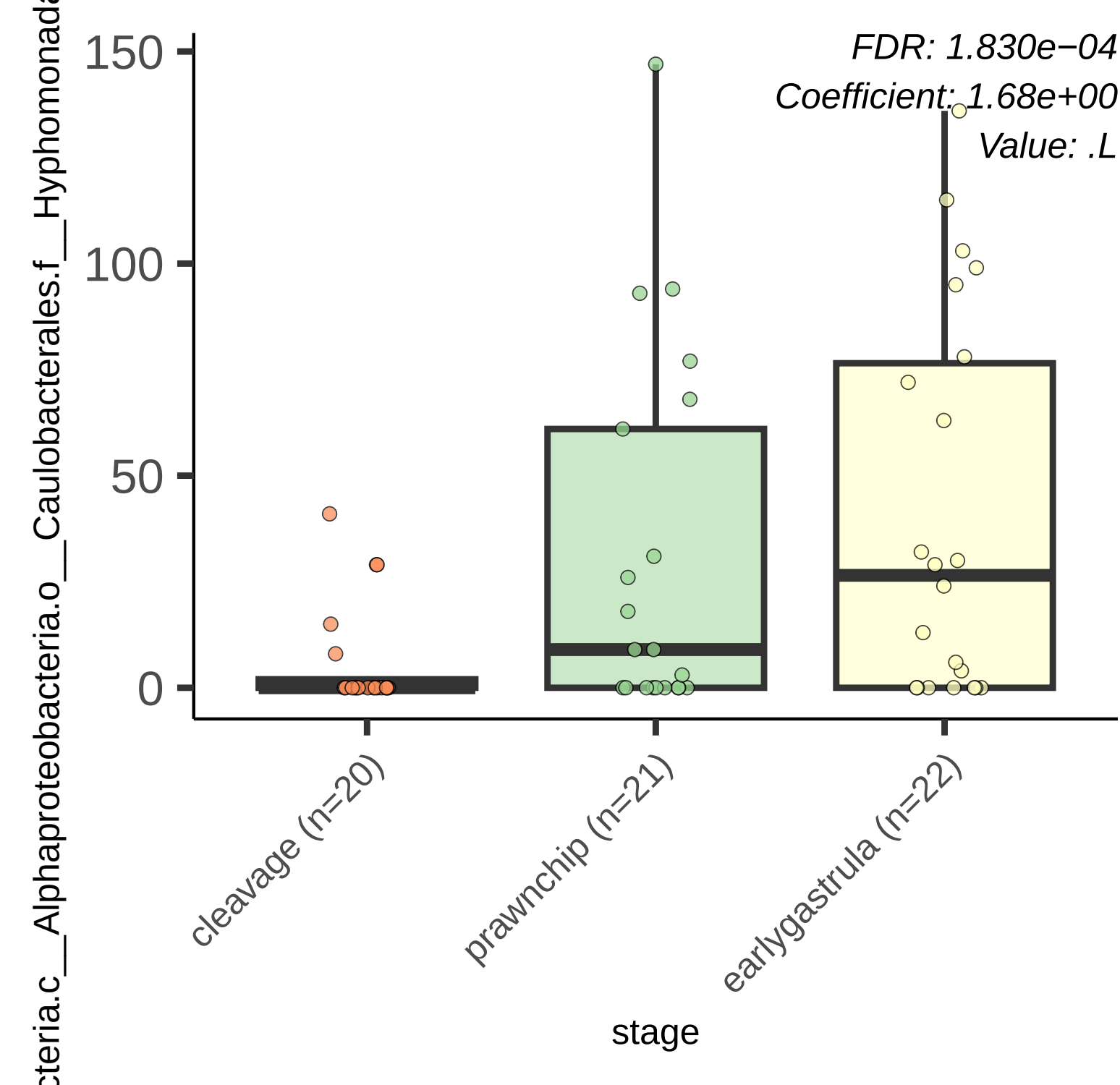
prawnchip (n=21)
earlygastrula (n=22)
stage
eobacteria.c__Gammaproteobacteria.o__Xanthomonadales.f__Xant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50
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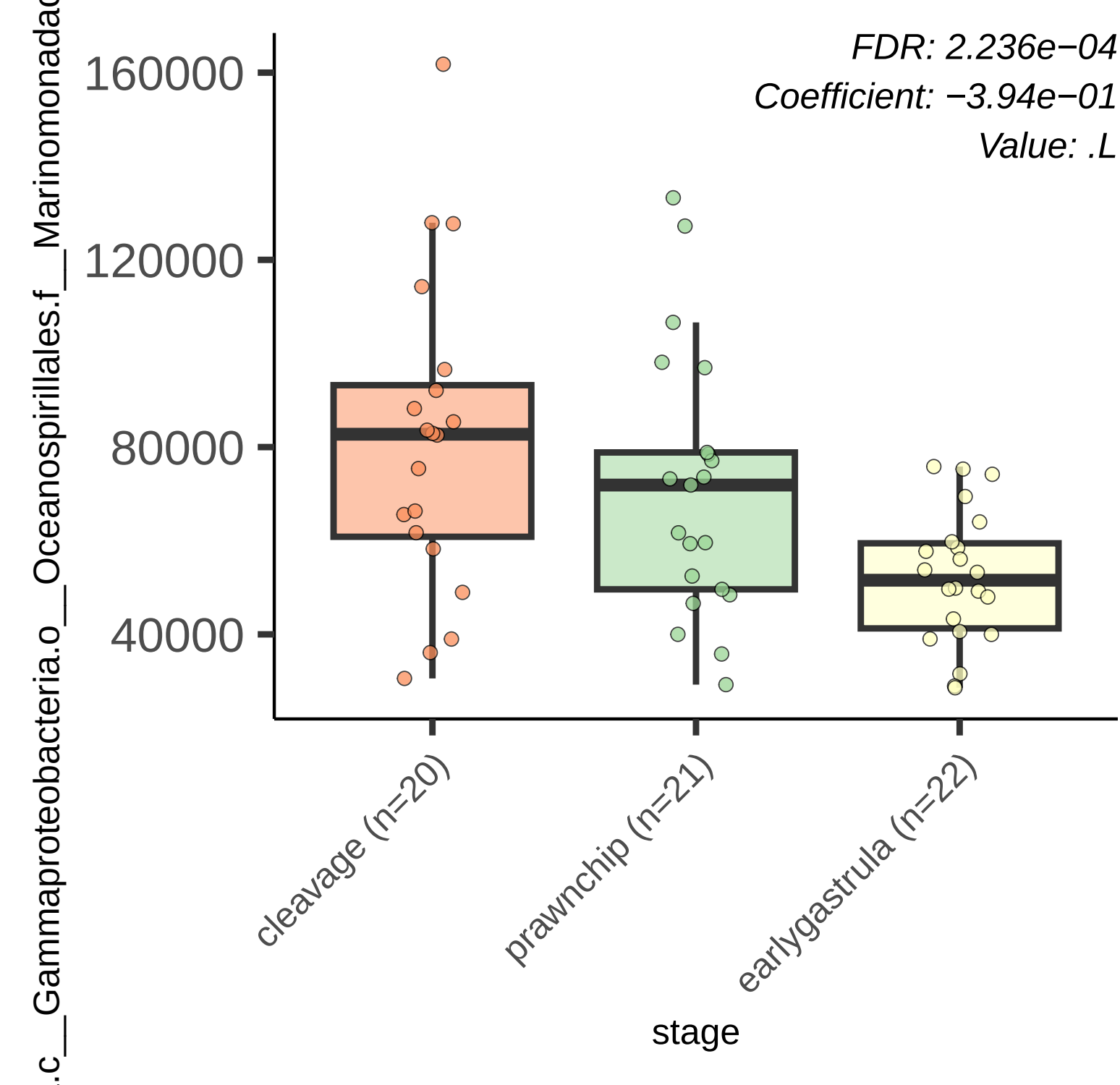
earlygastrula (n=22)
stage
cteria.c__Alphaproteobacteria.o__Caulobacterales.f__Hyphomonada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Value: .L
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120000
80000
40000
cleavage (n=20)
prawnchip (n=21)
earlygastrula (n=22)
stage
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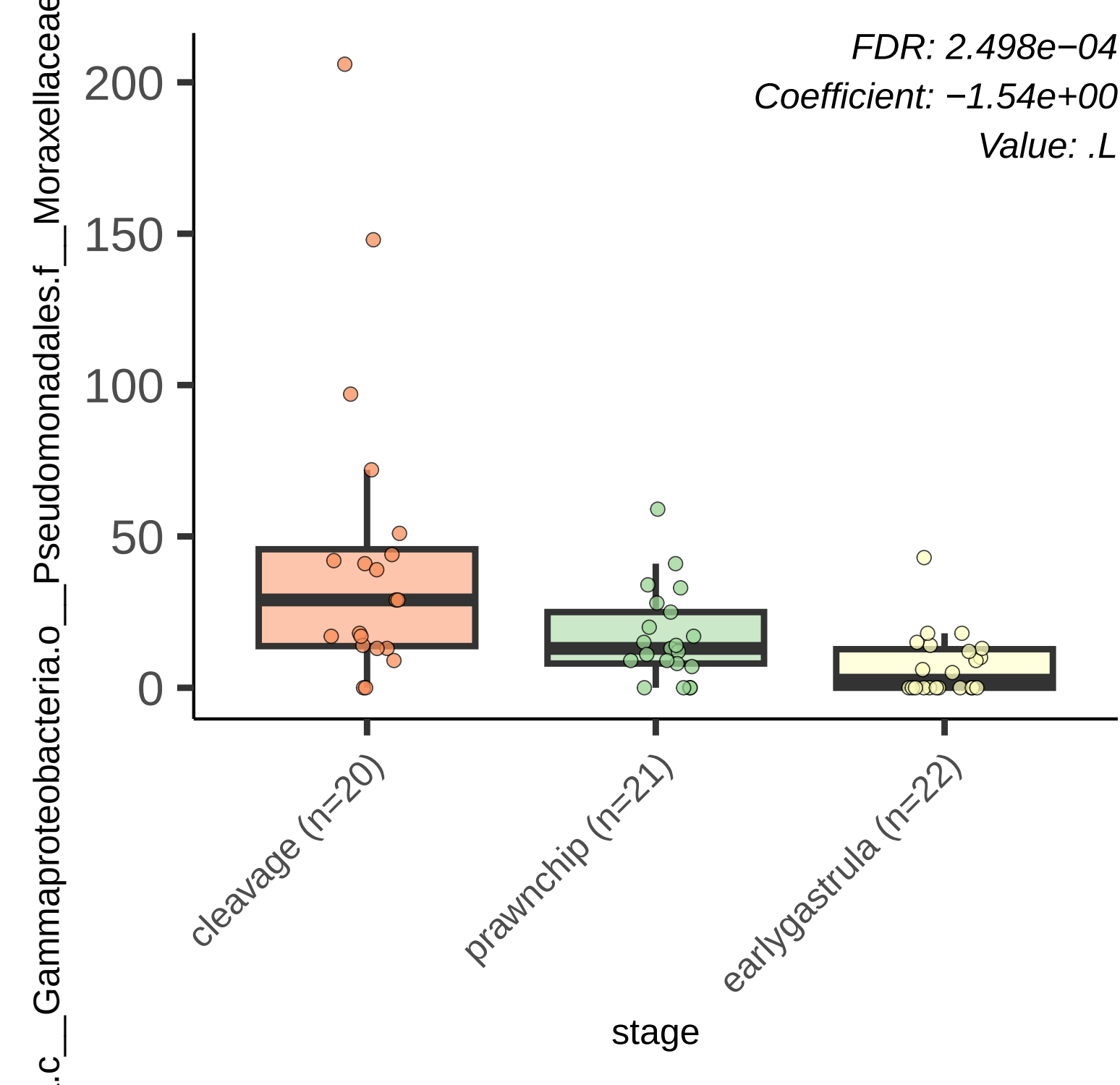
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150
100
50
0
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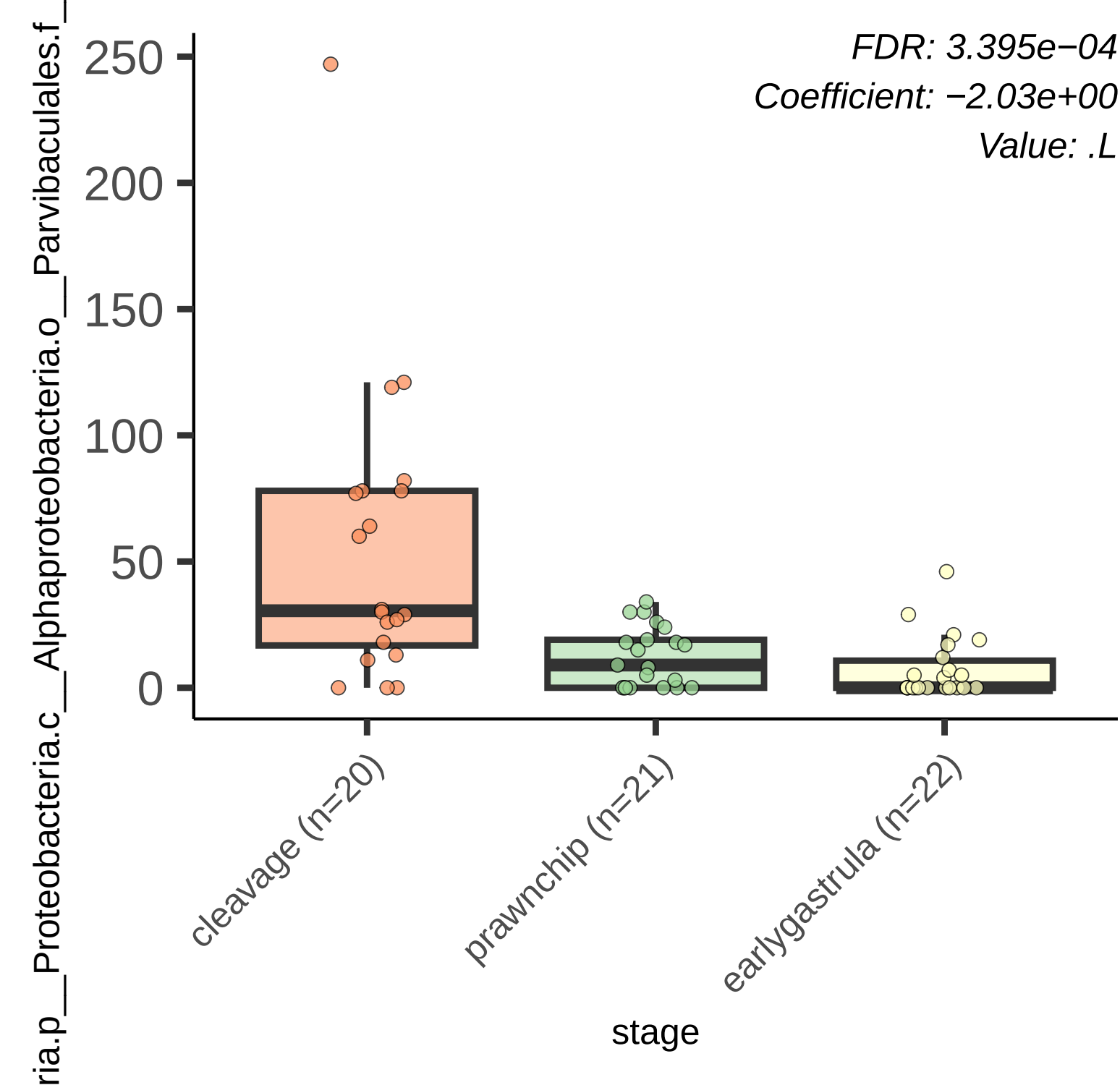
prawnchip (n=21)
earlygastrula (n=22)
stage
...c__Gammaproteobacteria.o__Pseudomonadales.f__Moraxellaceae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Value: .L
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200
150
100
50
0
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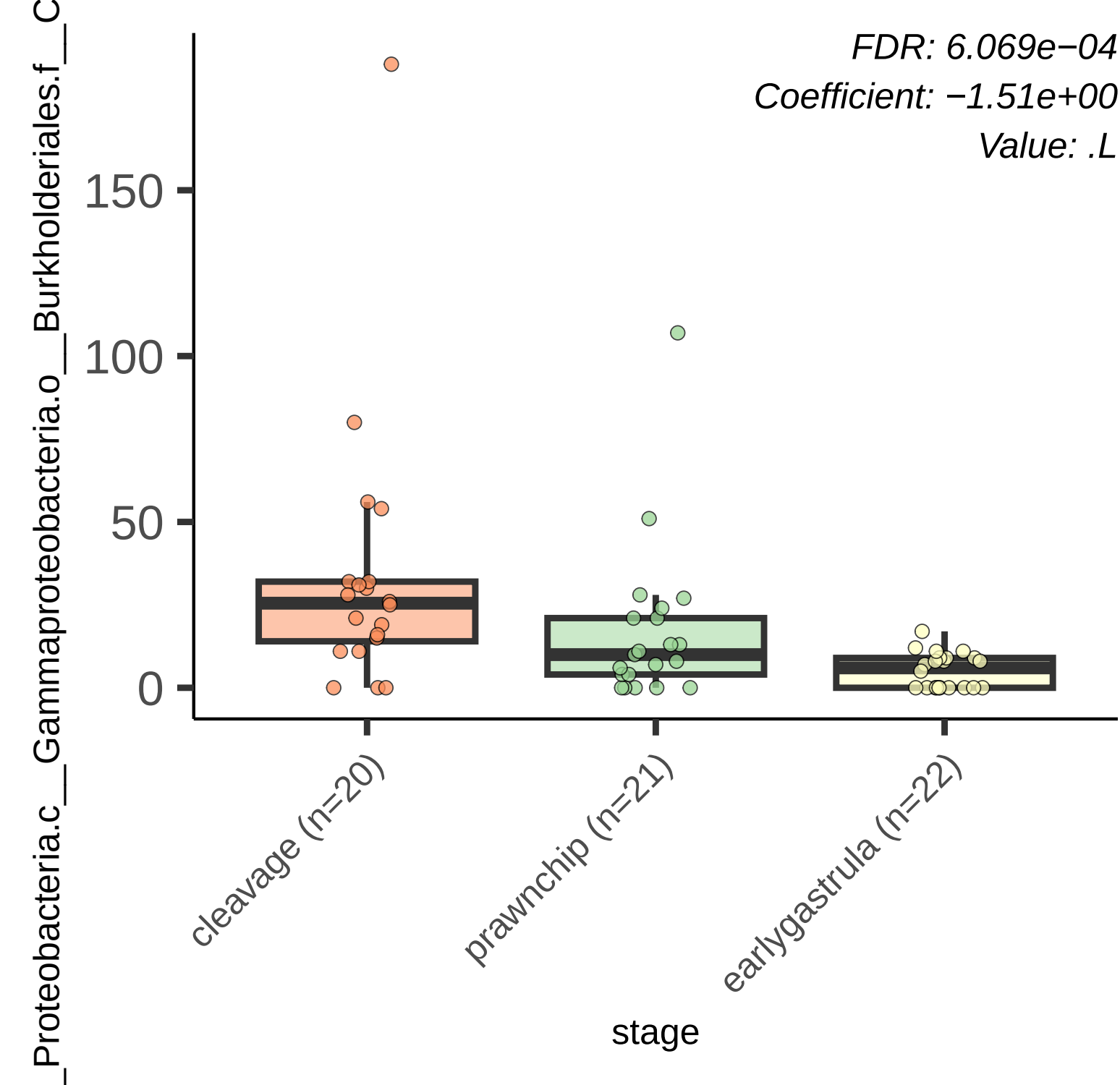
earlygastrula (n=22)
stage
ria.p__Proteobacteria.c__Alphaproteobacteria.o__Parvibaculales.f__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100
50
0
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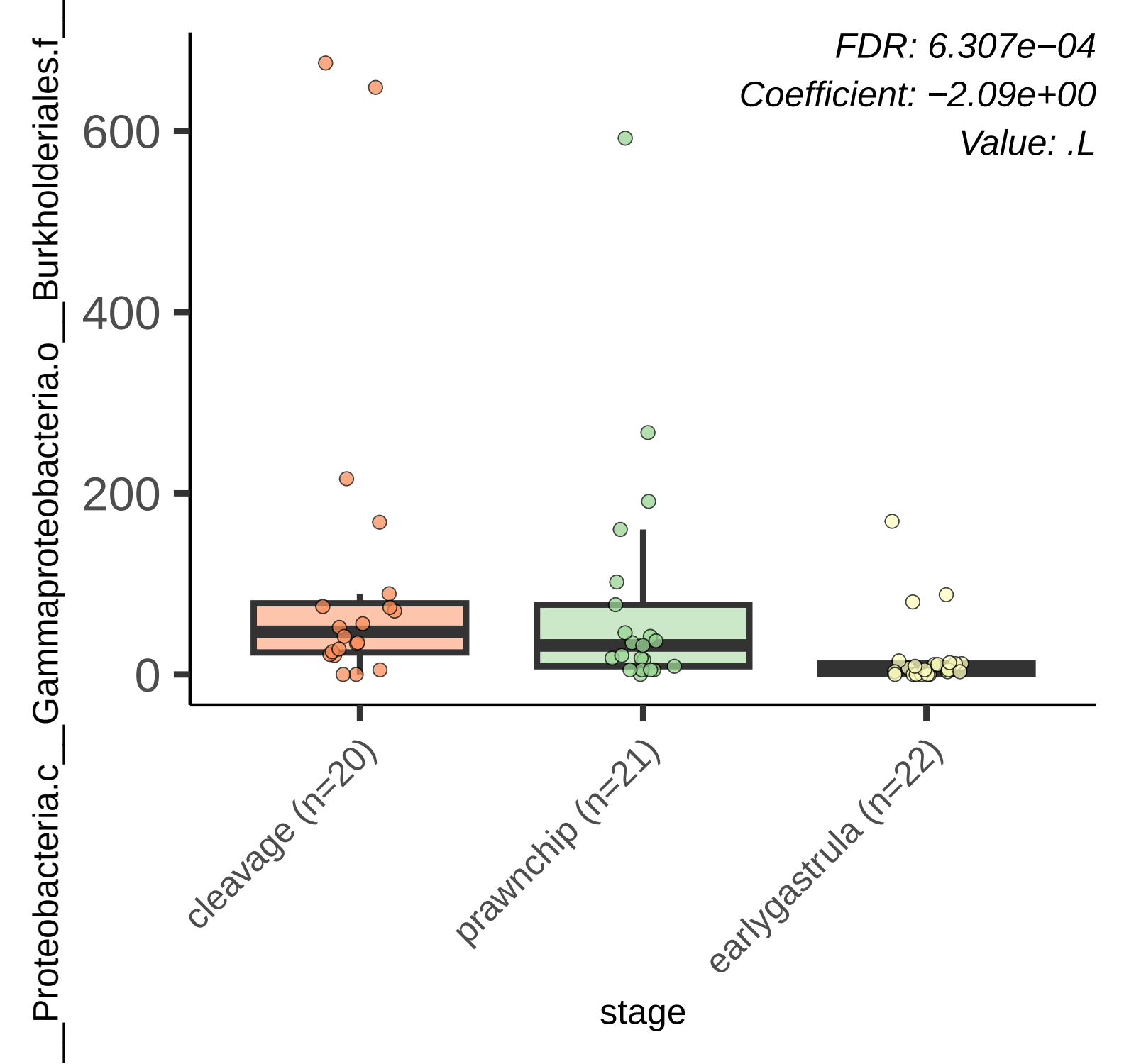
prawnchip (n=21)
earlygastrula (n=22)
stage
_Proteobacteria.c__Gammaproteobacteria.o__Burkholderiales.f__C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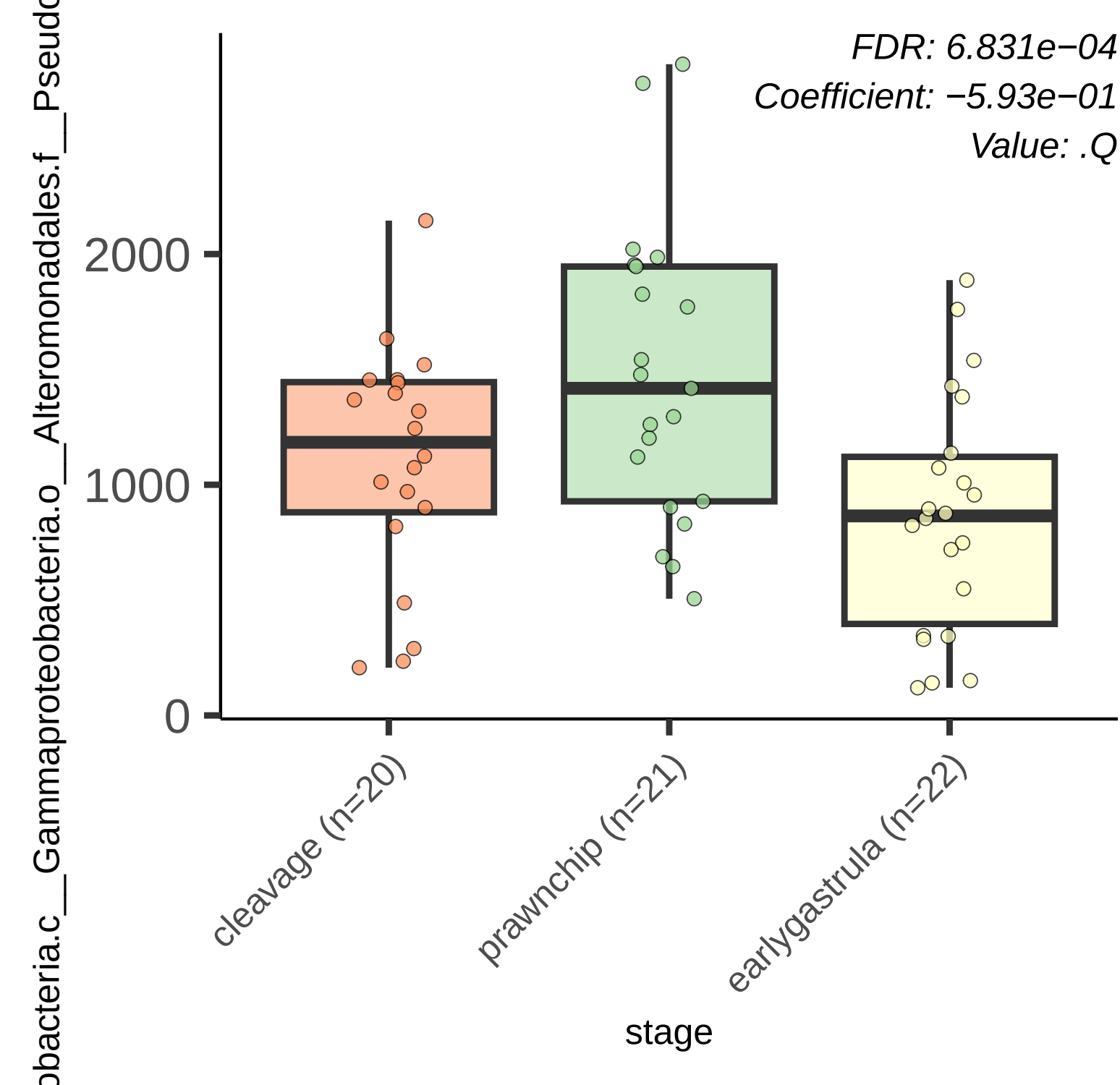
600
400
200
0
cleavage (n=20)
prawnchip (n=21)
earlygastrula (n=22)
stage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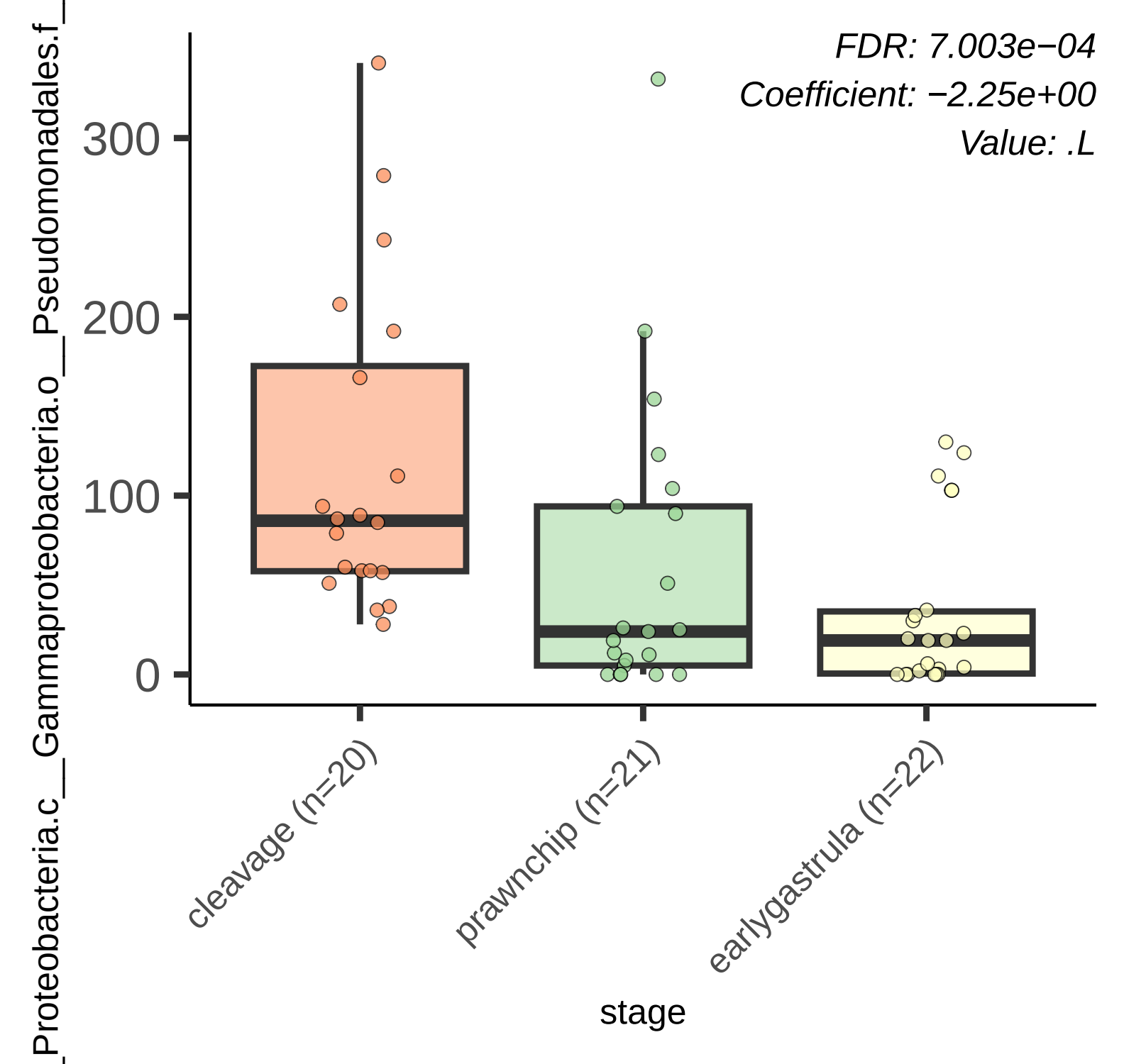
2000
1000
0
cleavage (n=20)
prawnchip (n=21)
earlygastrula (n=22)
stage
obacteria.c__Gammaproteobacteria.o__Alteromonadales.f__Pseudo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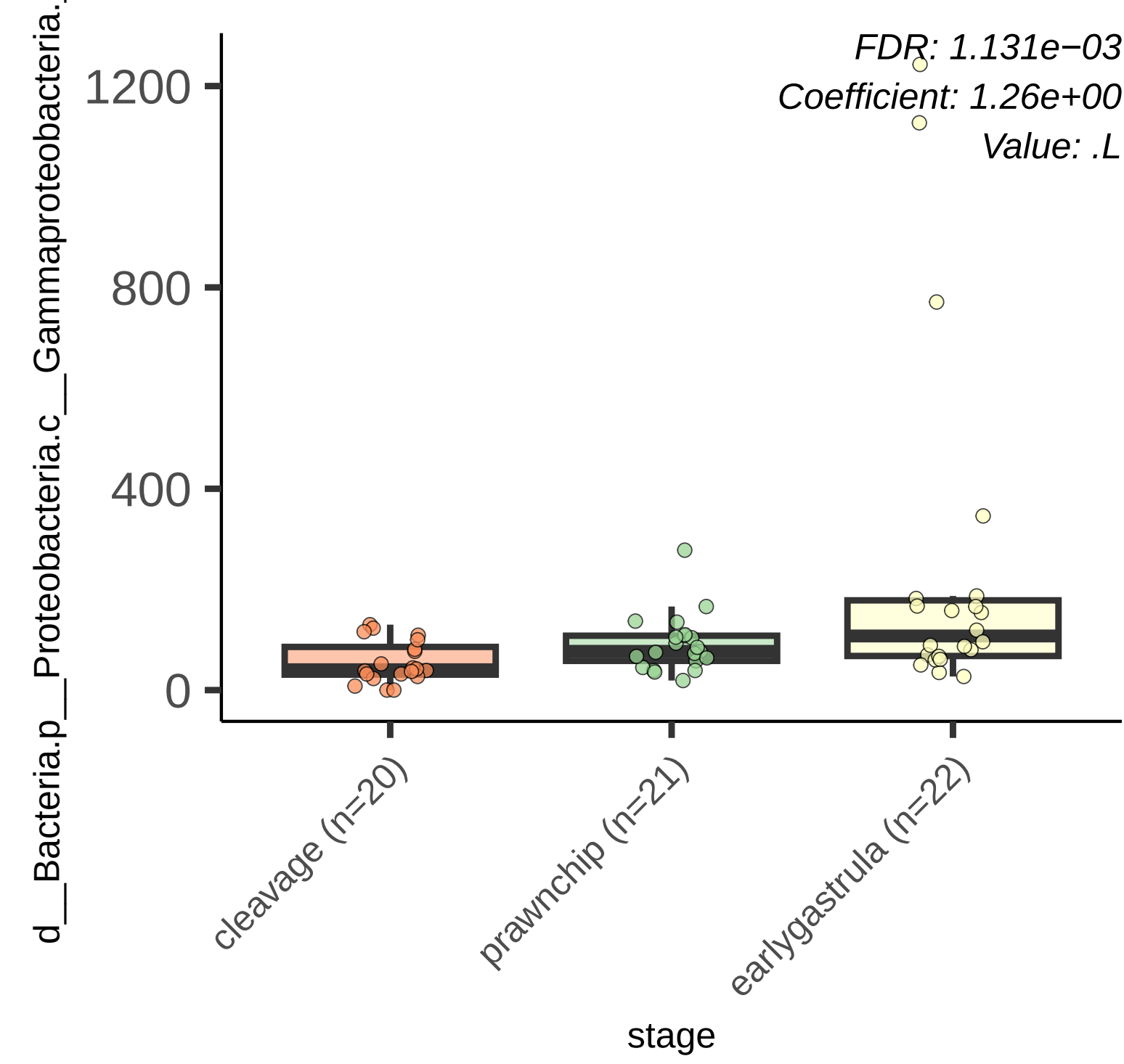
200
100
0
cleavage (n=20)
prawnchip (n=21)
earlygastrula (n=22)
stage
_Proteobacteria.c__Gammaproteobacteria.o__Pseudomonadales.f__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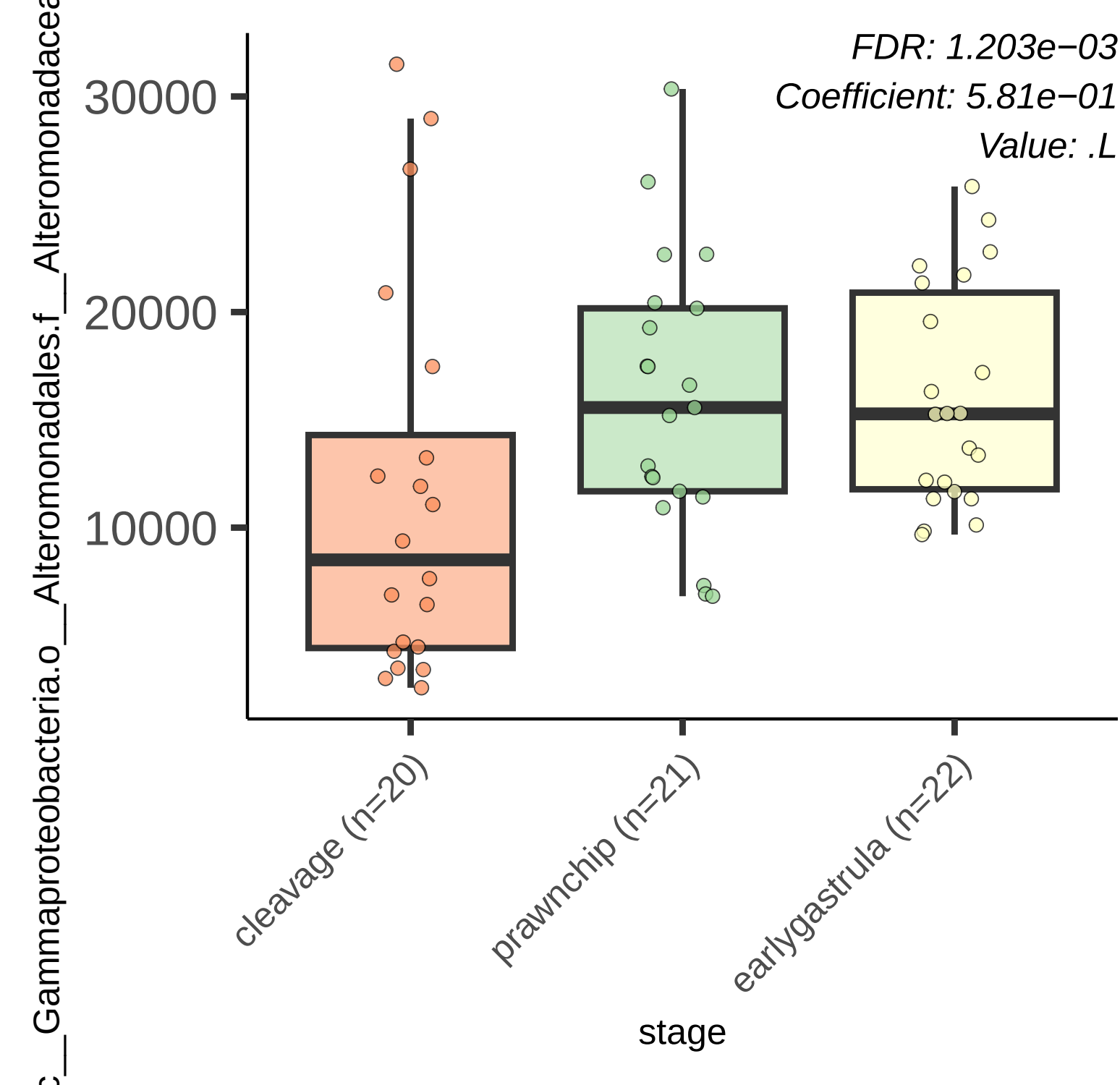
1200
800
400
0
cleavage (n=20)
prawnchip (n=21)
earlygastrula (n=22)
stage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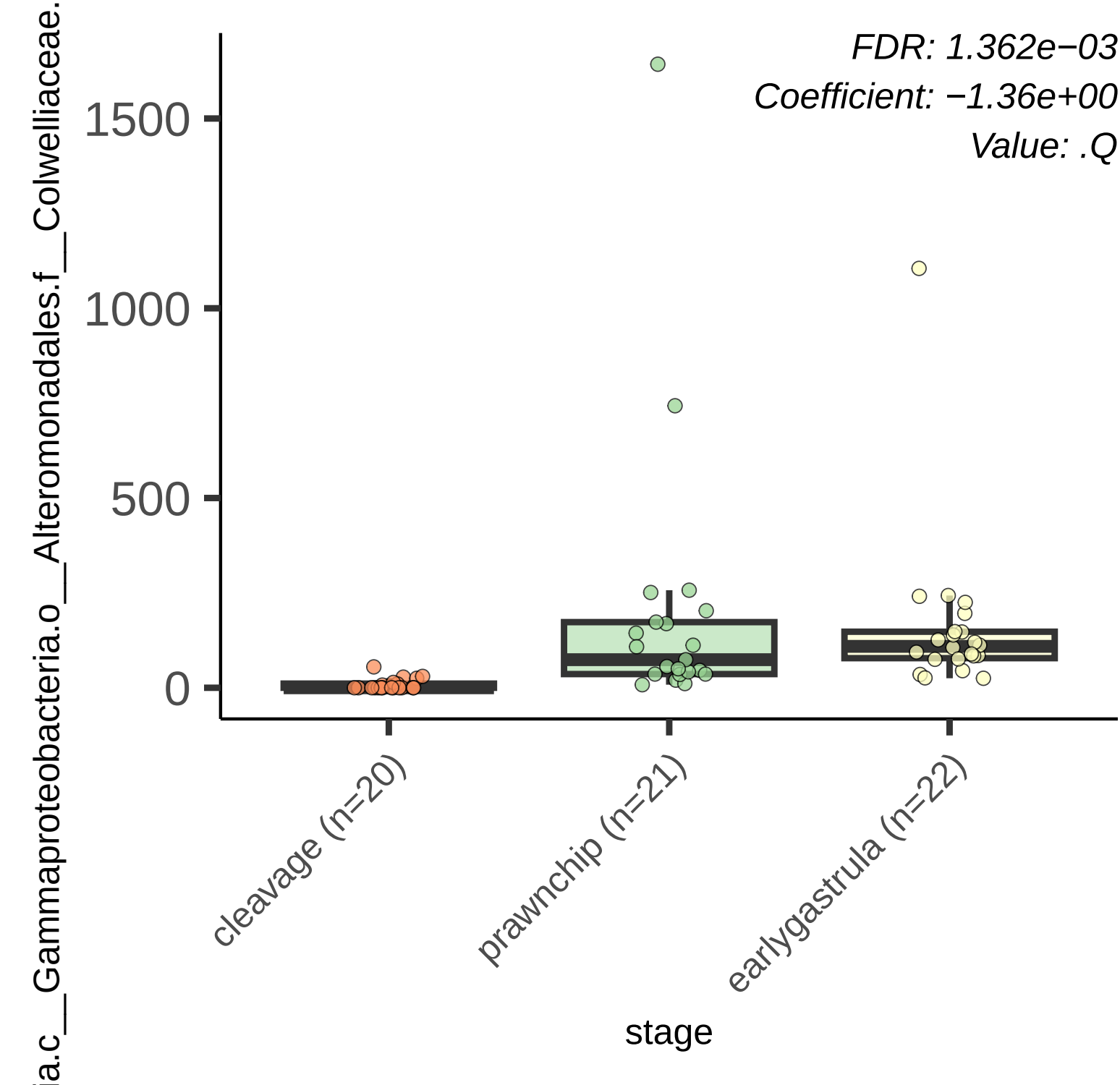
30000
20000
10000
cleavage (n=20)
prawnchip (n=21)
earlygastrula (n=22)
stage
c__Gammaproteobacteria.o__Alteromonadales.f__Alteromonadacea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1500
1000
500
0
cleavage (n=20)
prawnchip (n=21)
earlygastrula (n=22)
stage
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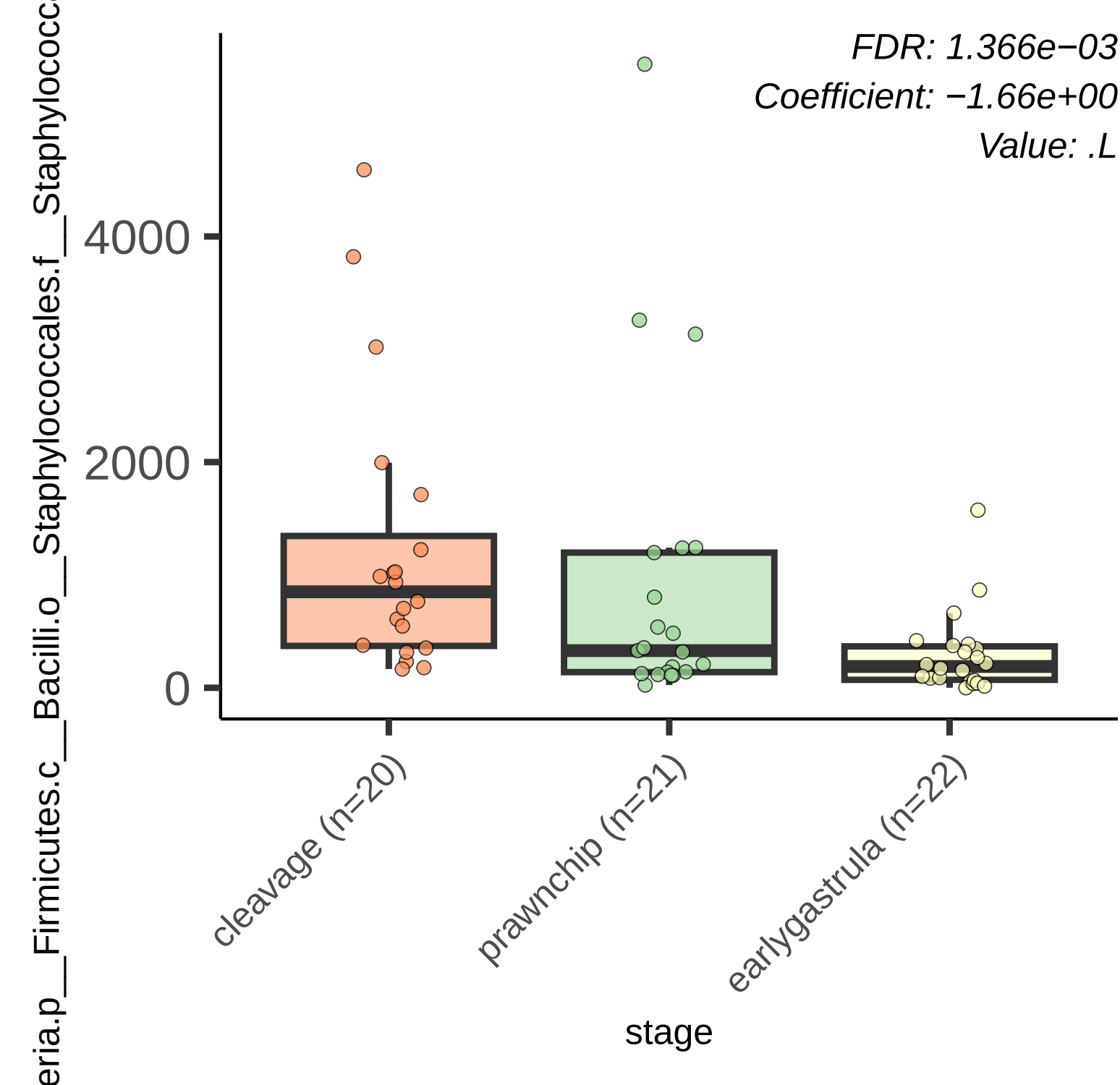
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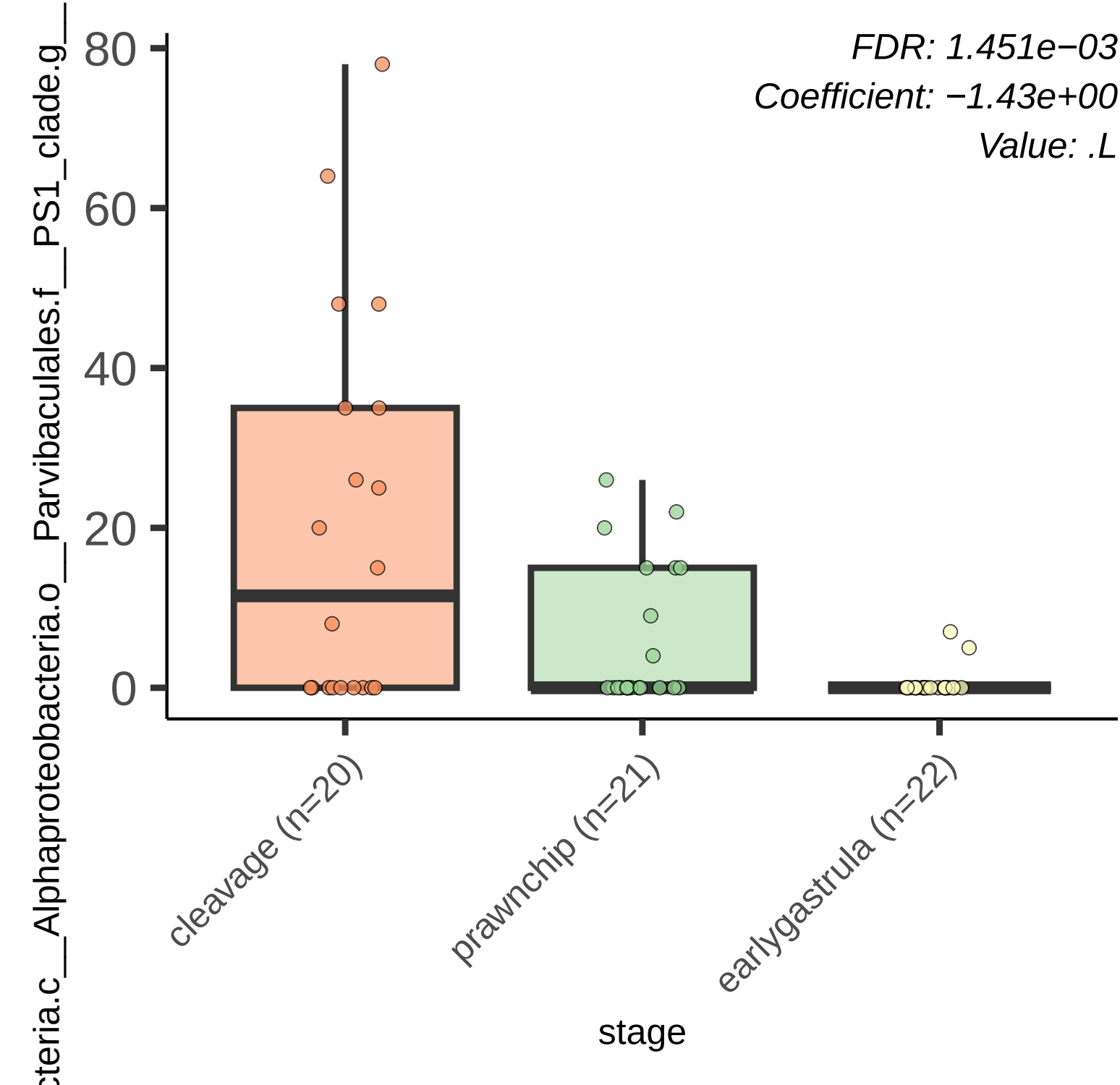
2000
0
cleavage (n=20)
prawnchip (n=21)
earlygastrula (n=22)
stage
eria.p__Firmicutes.c__Bacilli.o__Staphylococcales.f__Staphylococc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80
60
40
20
0
cleavage (n=20)
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earlygastrula (n=22)
stage
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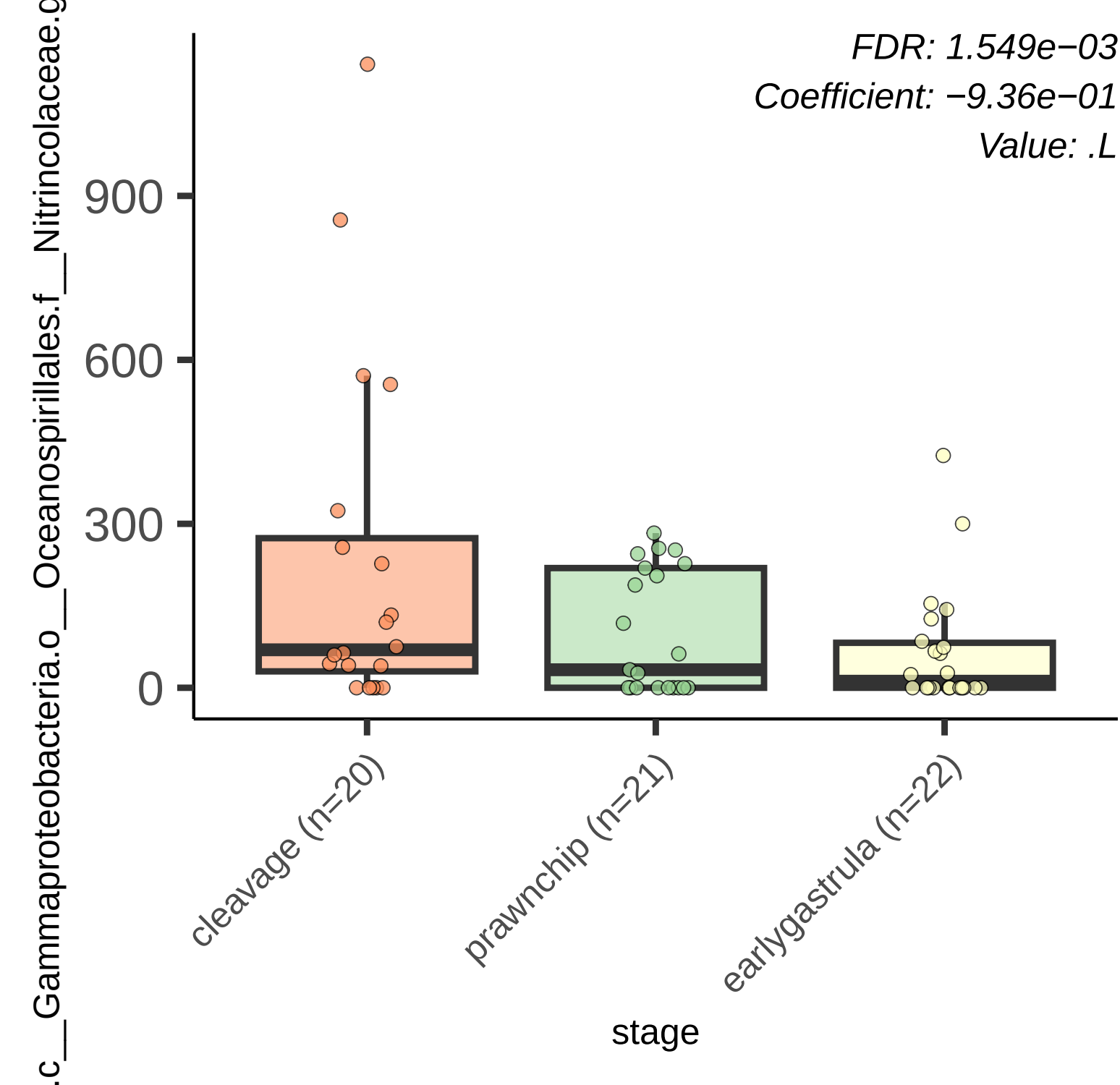
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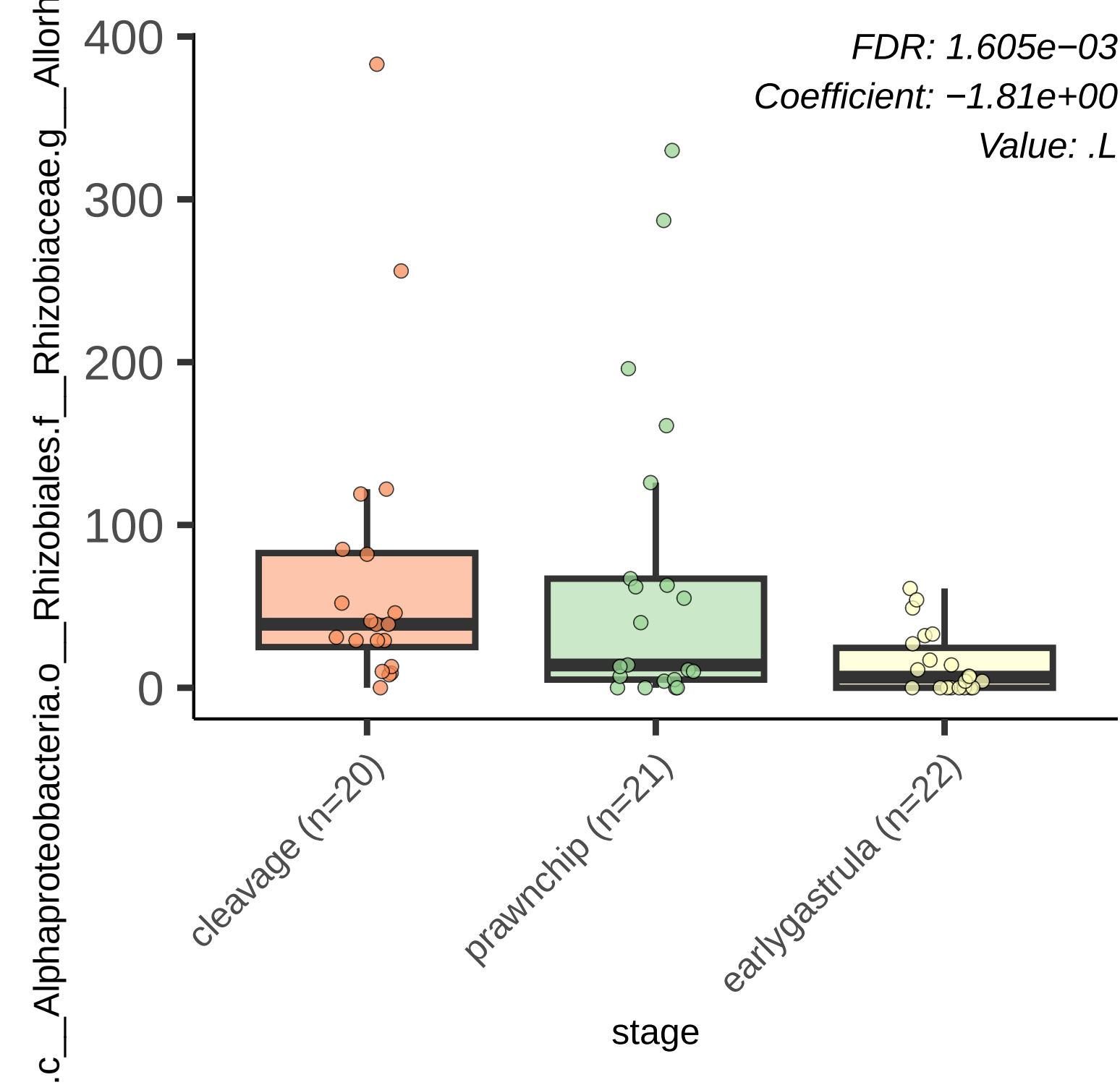
600
300
0
cleavage (n=20)
prawnchip (n=21)
earlygastrula (n=22)
stage
.c__Gammaproteobacteria.o__Oceanospirillales.f__Nitrincolaceae.g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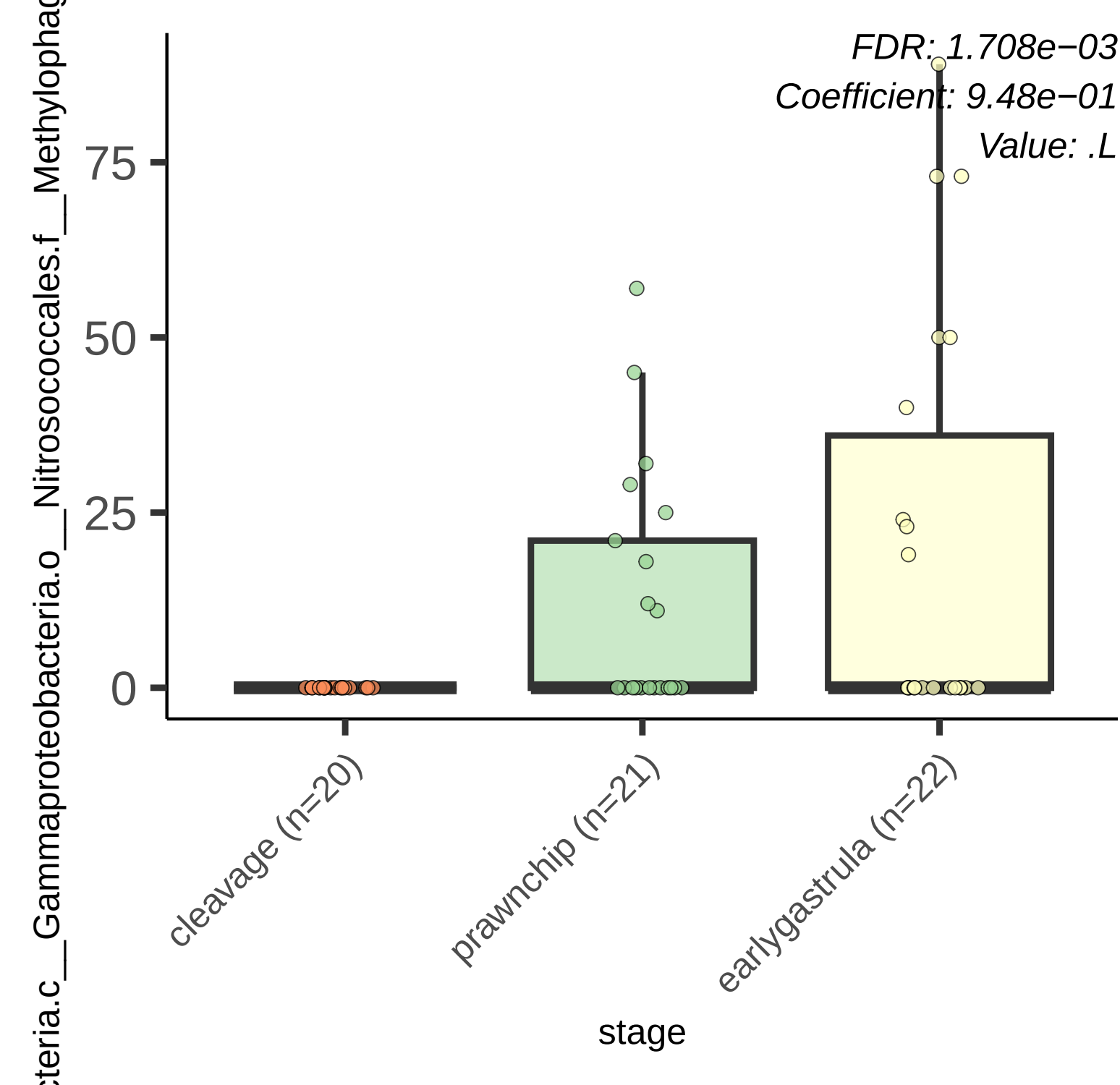
300
200
100
0
cleavage (n=20)
prawnchip (n=21)
earlygastrula (n=22)
stage
.c__Alphaproteobacteria.o__Rhizobiales.f__Rhizobiaceae.g__Allorh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25
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stage
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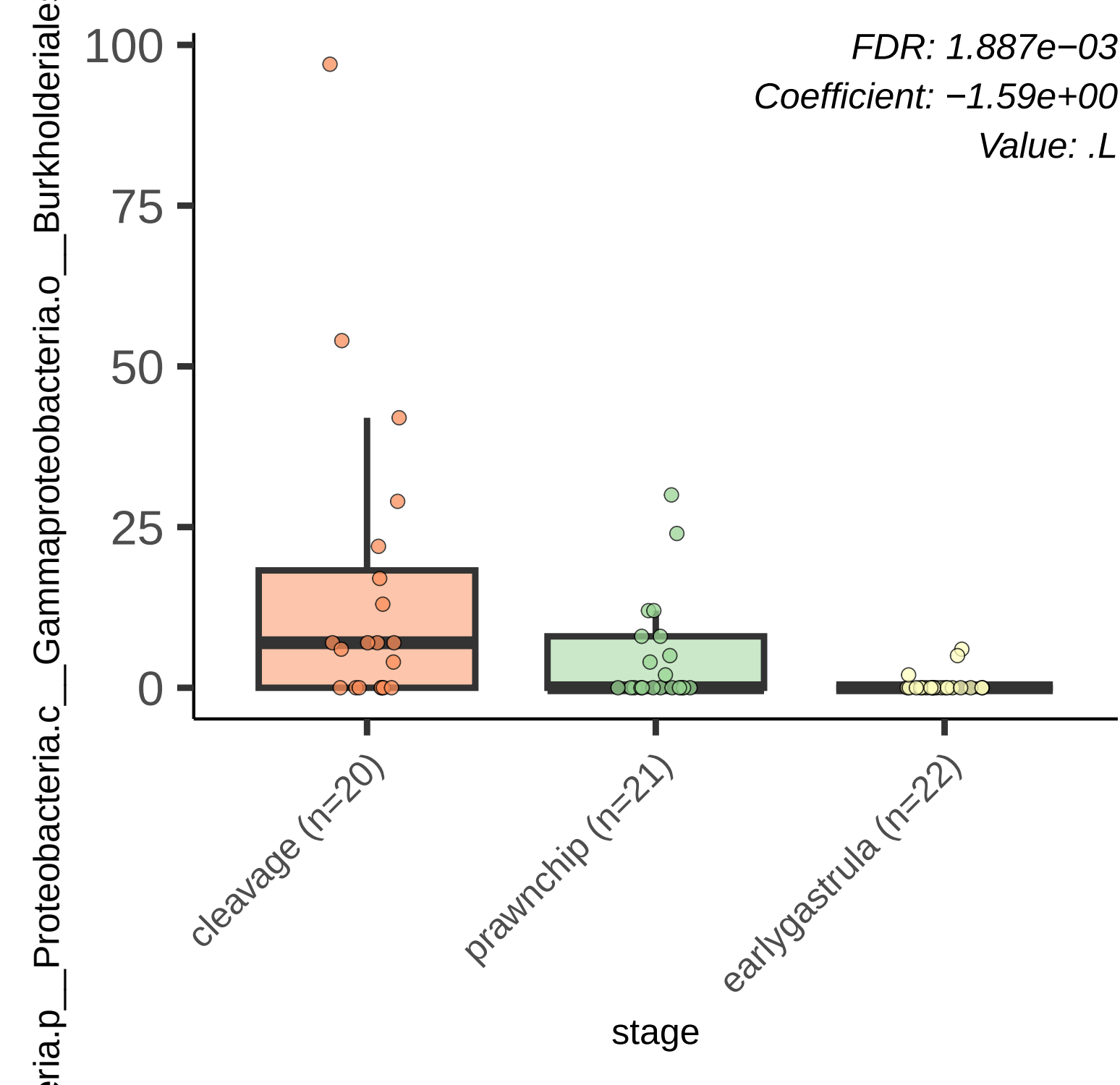
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50
25
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prawnchip (n=21)
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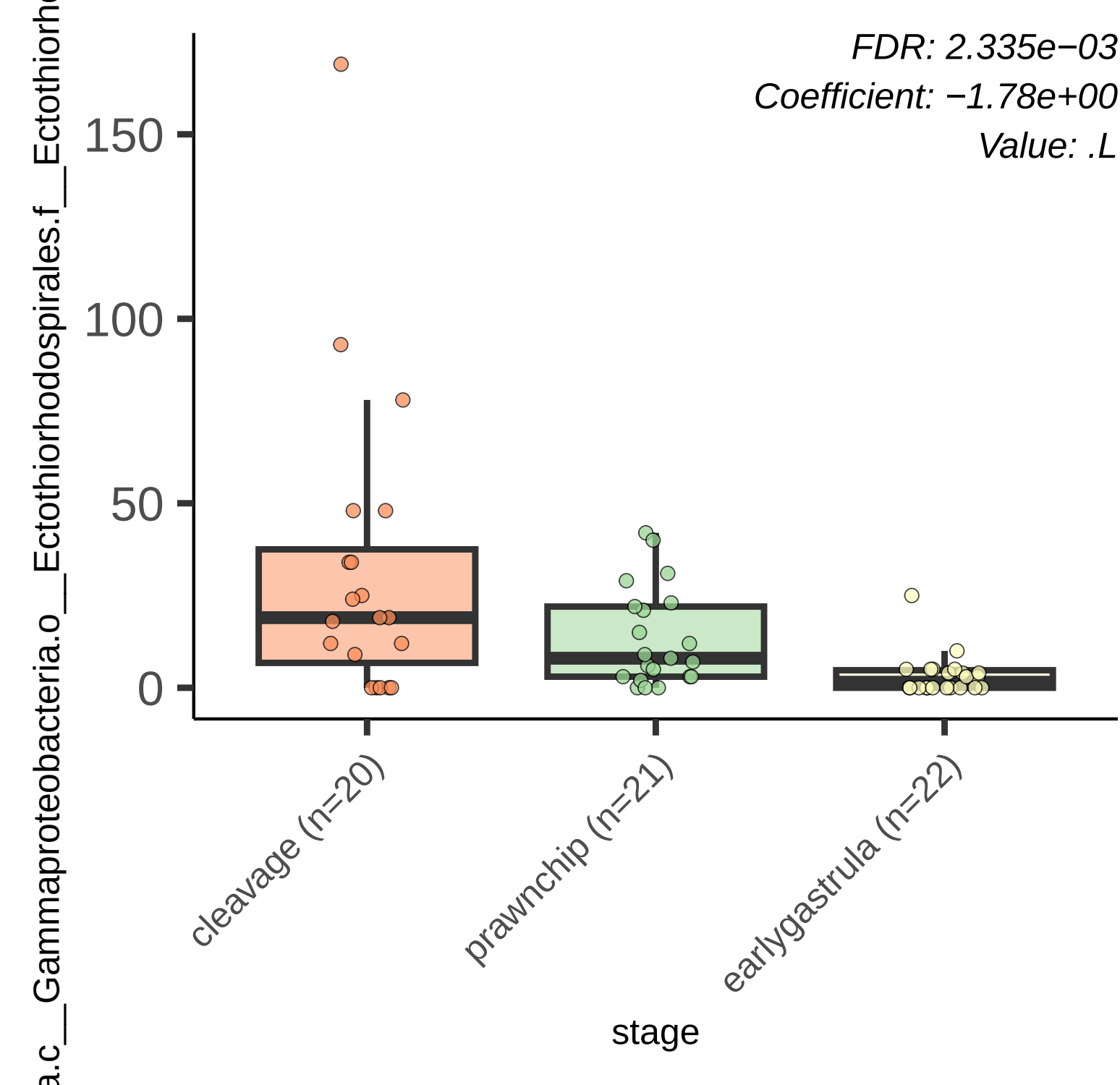
stage
ria.p__Proteobacteria.c__Gammaproteobacteria.o__Burkholderiales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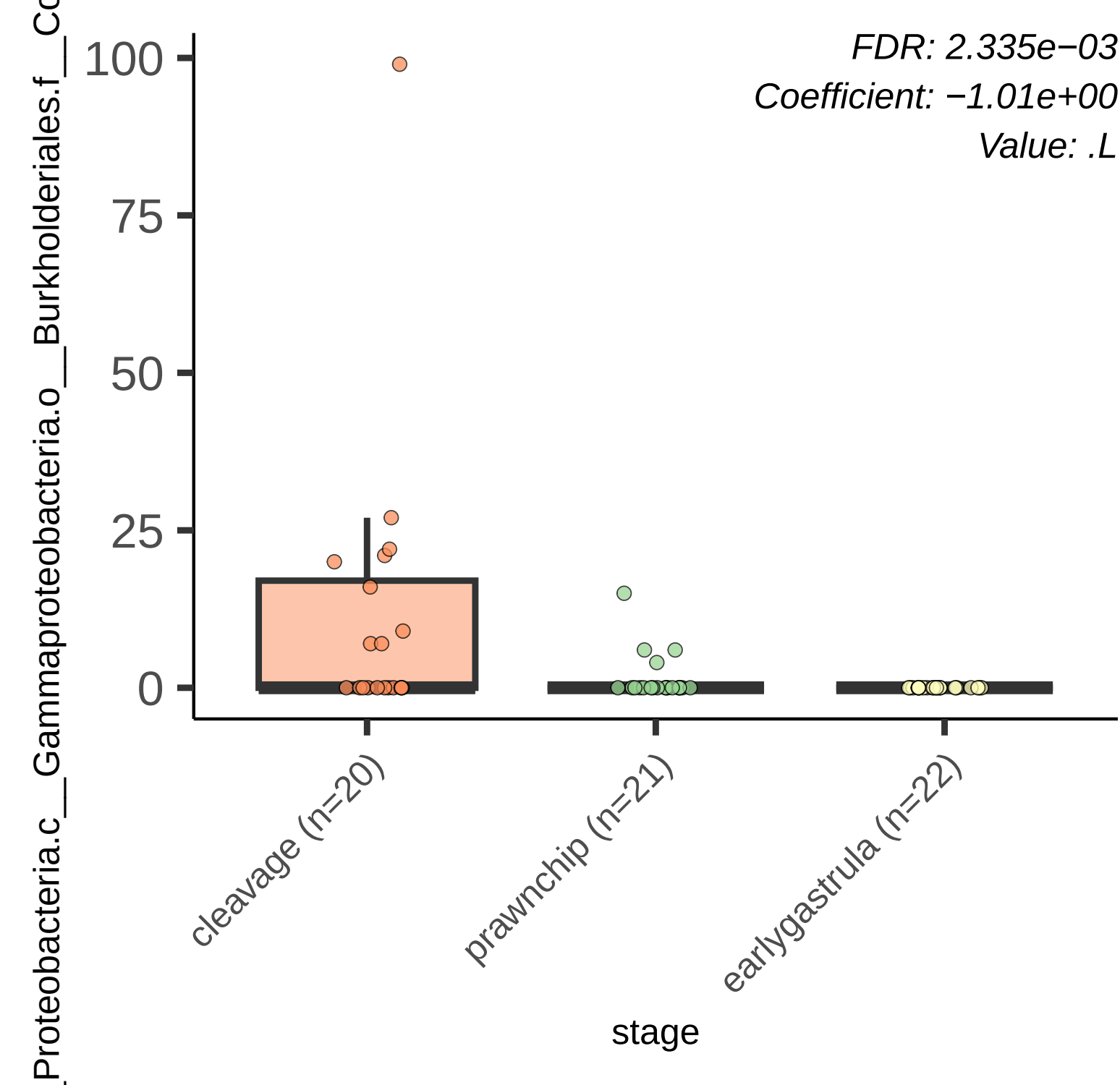
100
50
0
cleavage (n=20)
prawnchip (n=21)
earlygastrula (n=22)
stage
a.c__Gammaproteobacteria.o__Ectothiorhodospirales.f__Ectothiorh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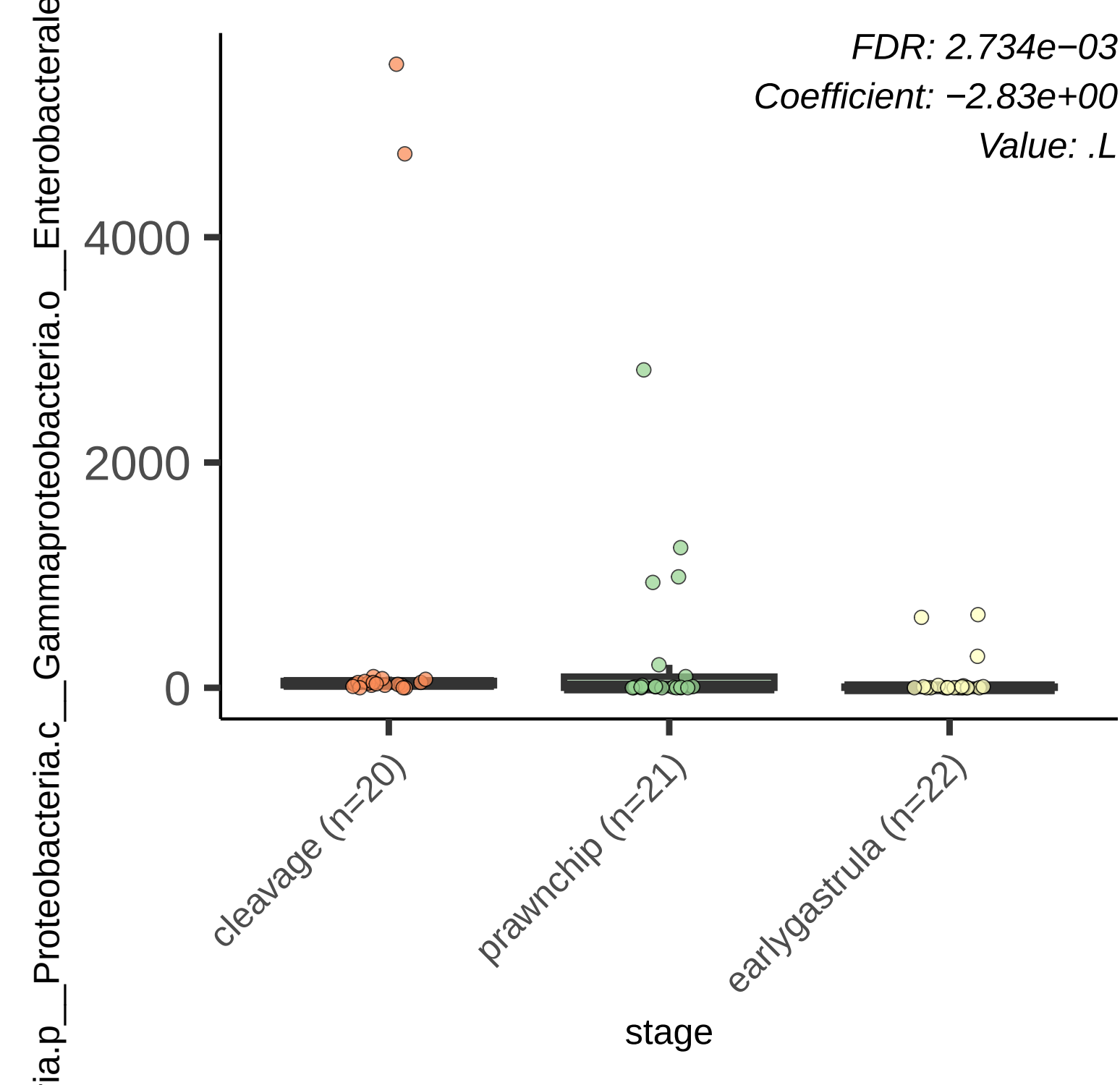
100
75
50
25
0
cleavage (n=20)
prawnchip (n=21)
earlygastrula (n=22)
stage
Proteobacteria.c__Gammaproteobacteria.o__Burkholderiales.f__Co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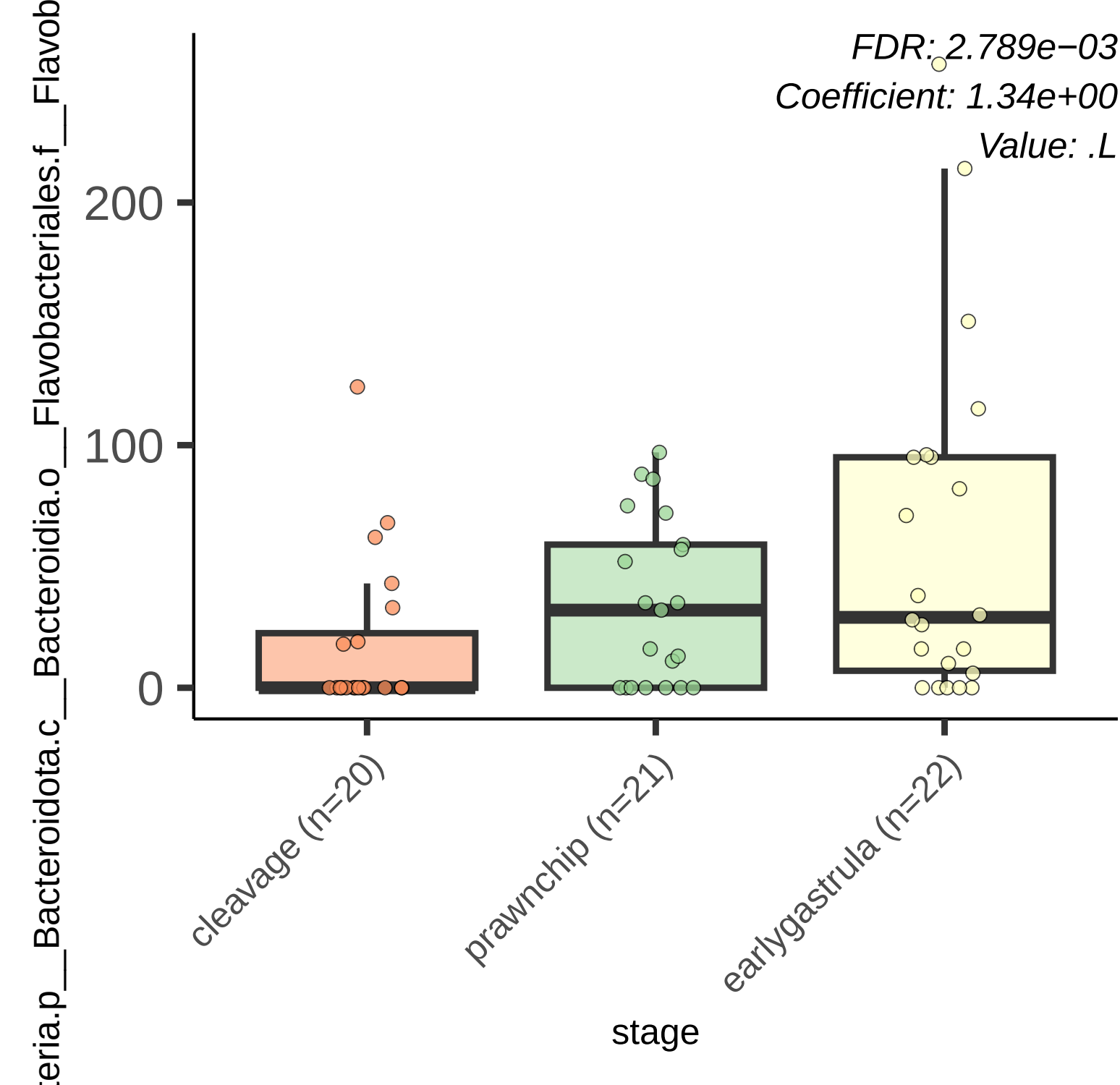
2000
0
cleavage (n=20)
prawnchip (n=21)
earlygastrula (n=22)
stage
ria.p__Proteobacteria.c__Gammaproteobacteria.o__Enterobacterale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Coefficient: 1.34e+00
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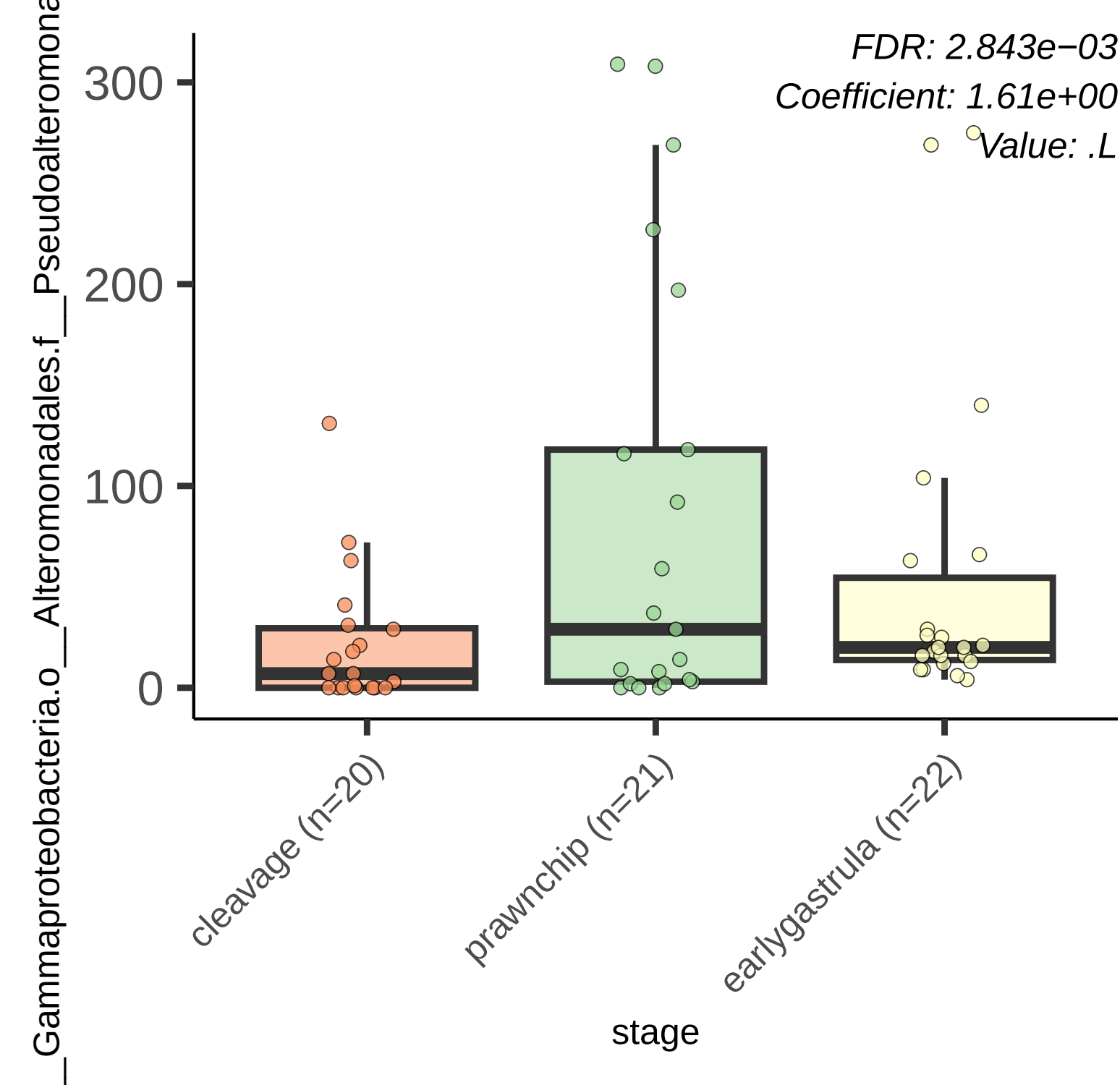
200
100
0
cleavage (n=20)
prawnchip (n=21)
earlygastrula (n=22)
stage
teria.p__Bacteroidota.c__Bacteroidia.o__Flavobacteriales.f__Flavob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200
100
0
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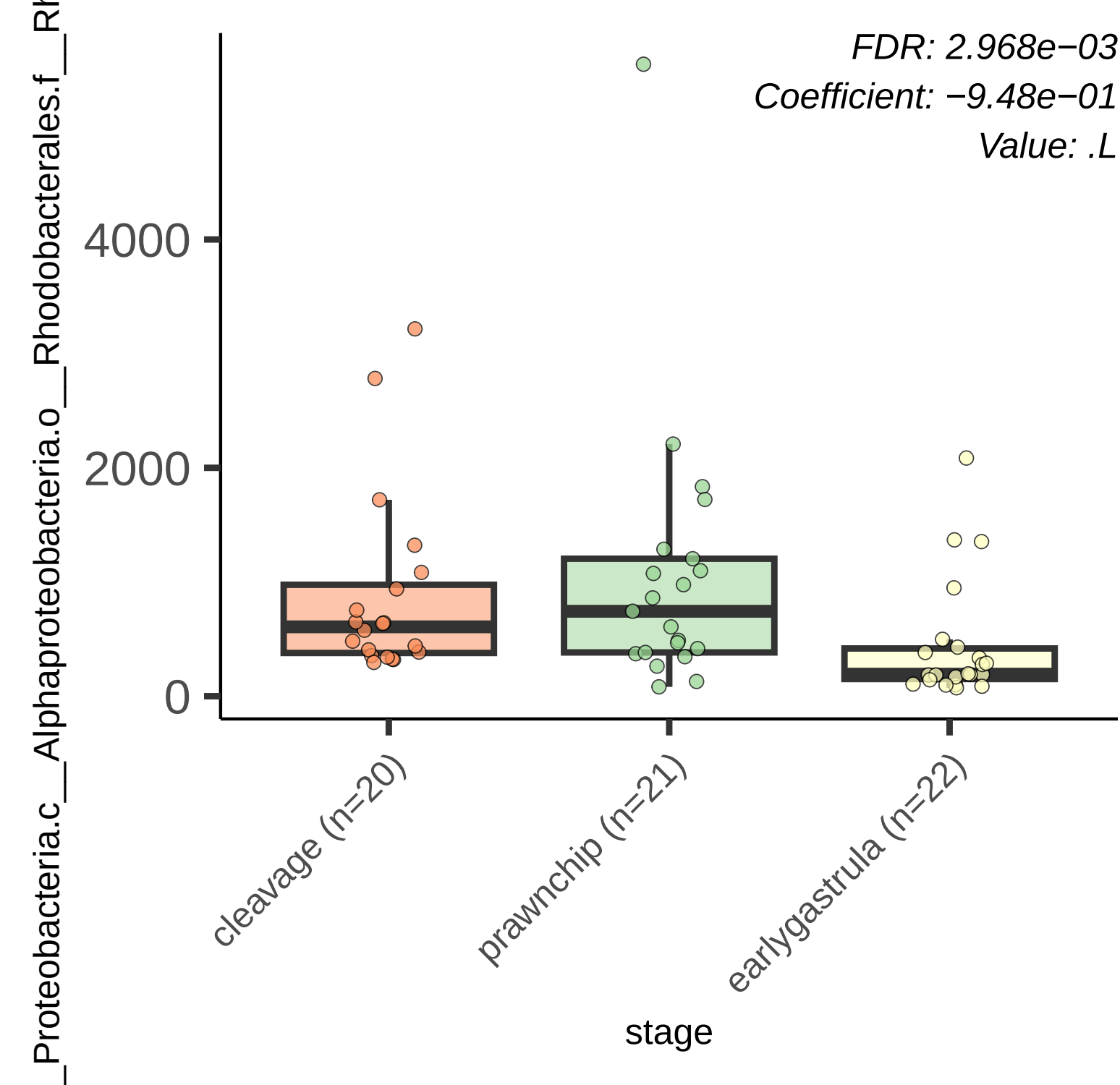
prawnchip (n=21)
earlygastrula (n=22)
stage
__Gammaproteobacteria.o__Alteromonadales.f__Pseudoalteromona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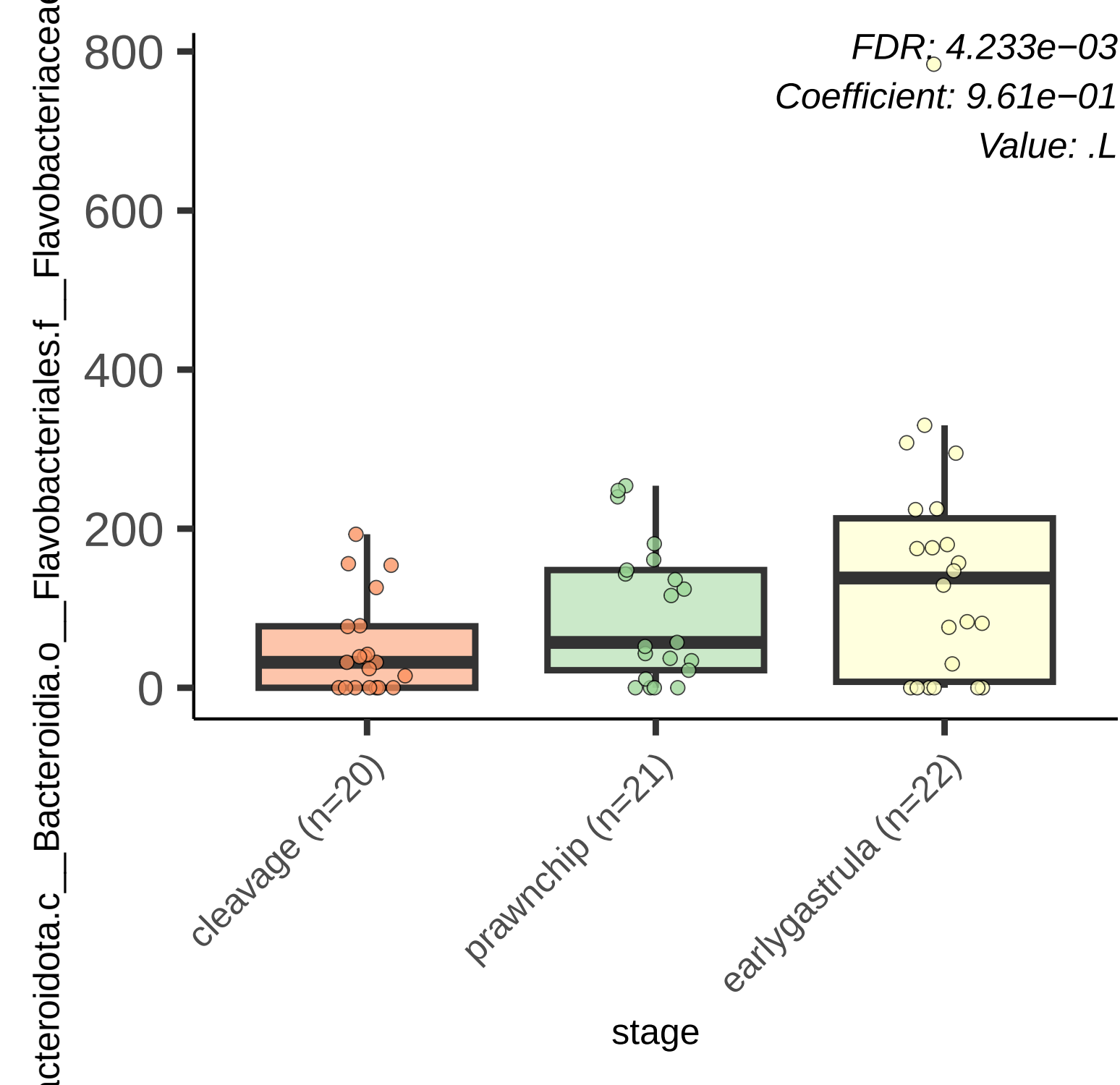
2000
0
cleavage (n=20)
prawnchip (n=21)
earlygastrula (n=22)
stage
_Proteobacteria.c__Alphaproteobacteria.o__Rhodobacterales.f__Rh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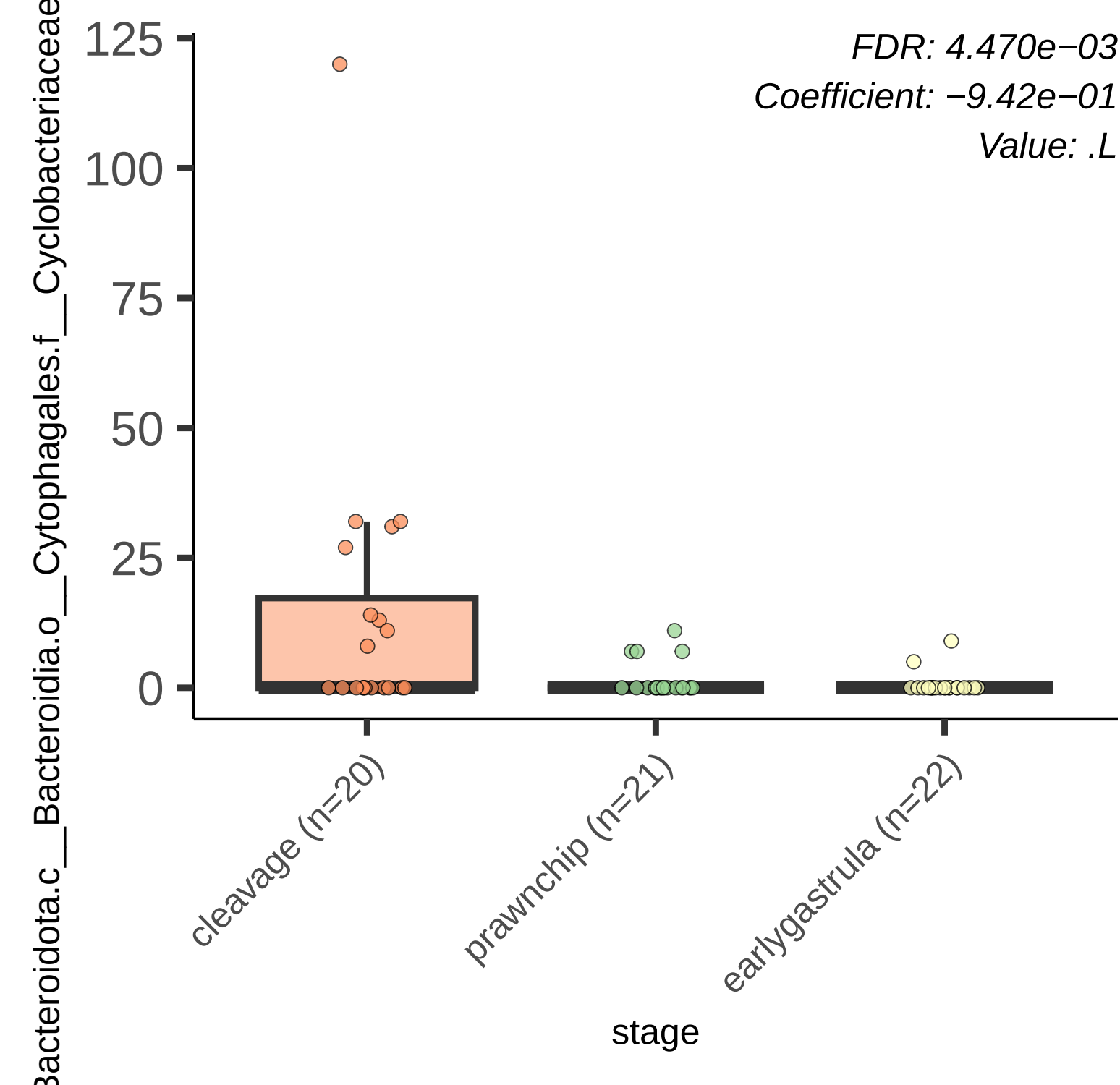
Coefficient: 9.61e−01
Value: .L
800
600
400
200
0
cleavage (n=20)
prawnchip (n=21)
earlygastrula (n=22)
stage
cteroidota.c__Bacteroidia.o__Flavobacteriales.f__Flavobacteriacea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Coefficient: −9.42e−01
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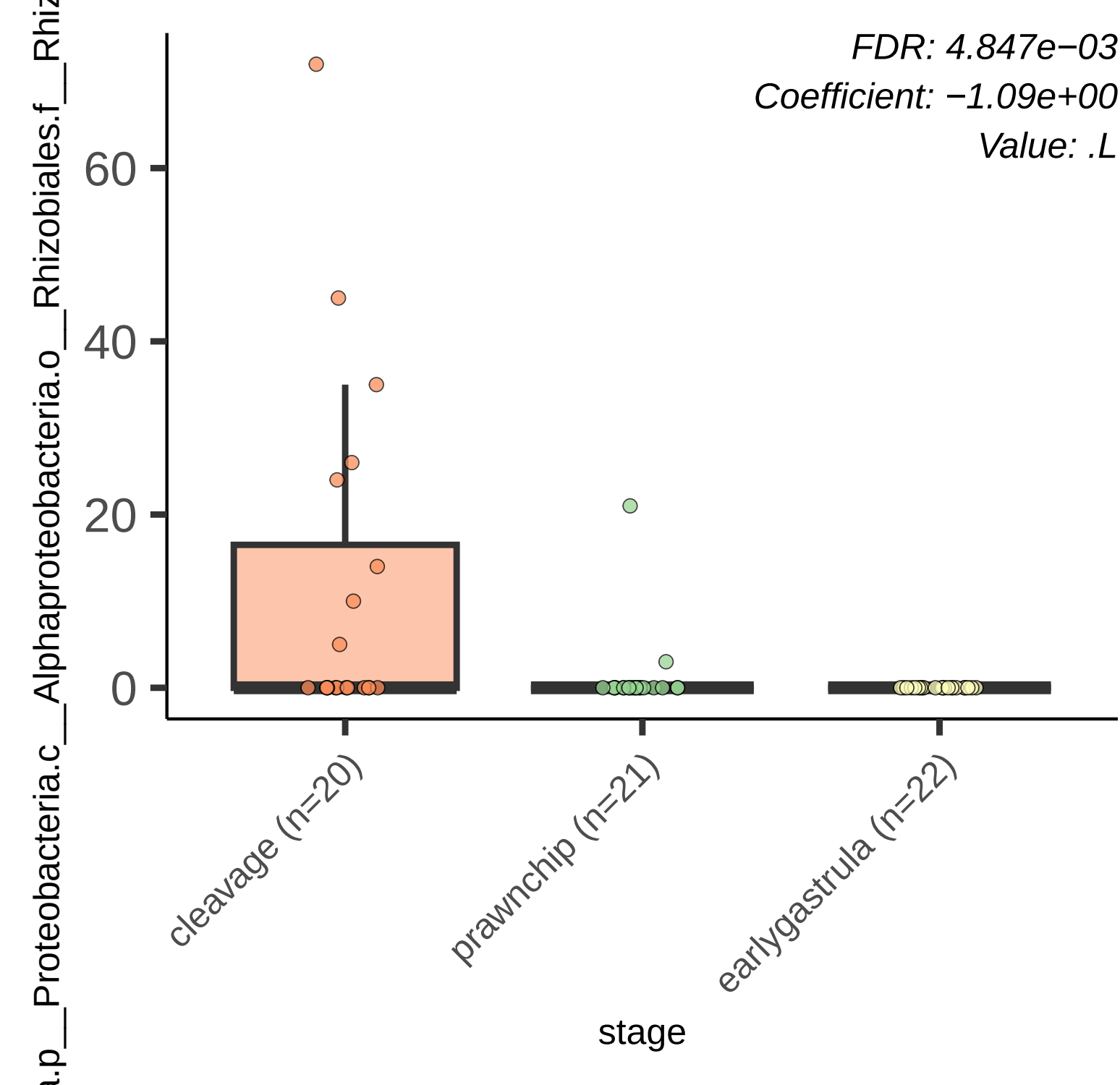
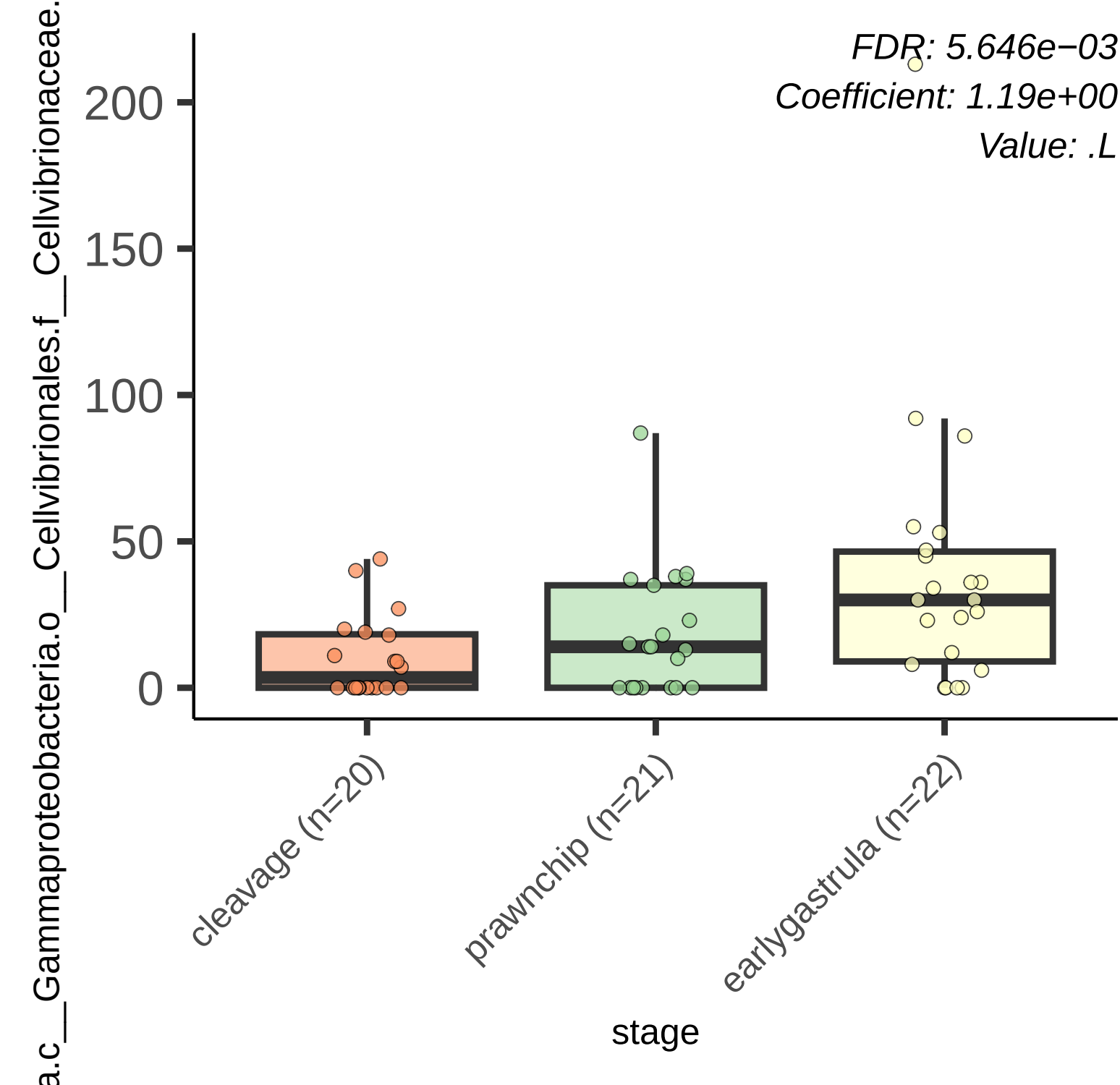
125
100
75
50
25
0
cleavage (n=20)
prawnchip (n=21)
earlygastrula (n=22)
stage
Bacteroidota.c__Bacteroidia.o__Cytophagales.f__Cyclobacteriaceae


FDR: 4.847e−03
Coefficient: −1.09e+00
Value: .L
60
40
20
0
cleavage (n=20)
prawnchip (n=21)
earlygastrula (n=22)
stage
a.p__Proteobacteria.c__Alphaproteobacteria.o__Rhizobiales.f__Rhiz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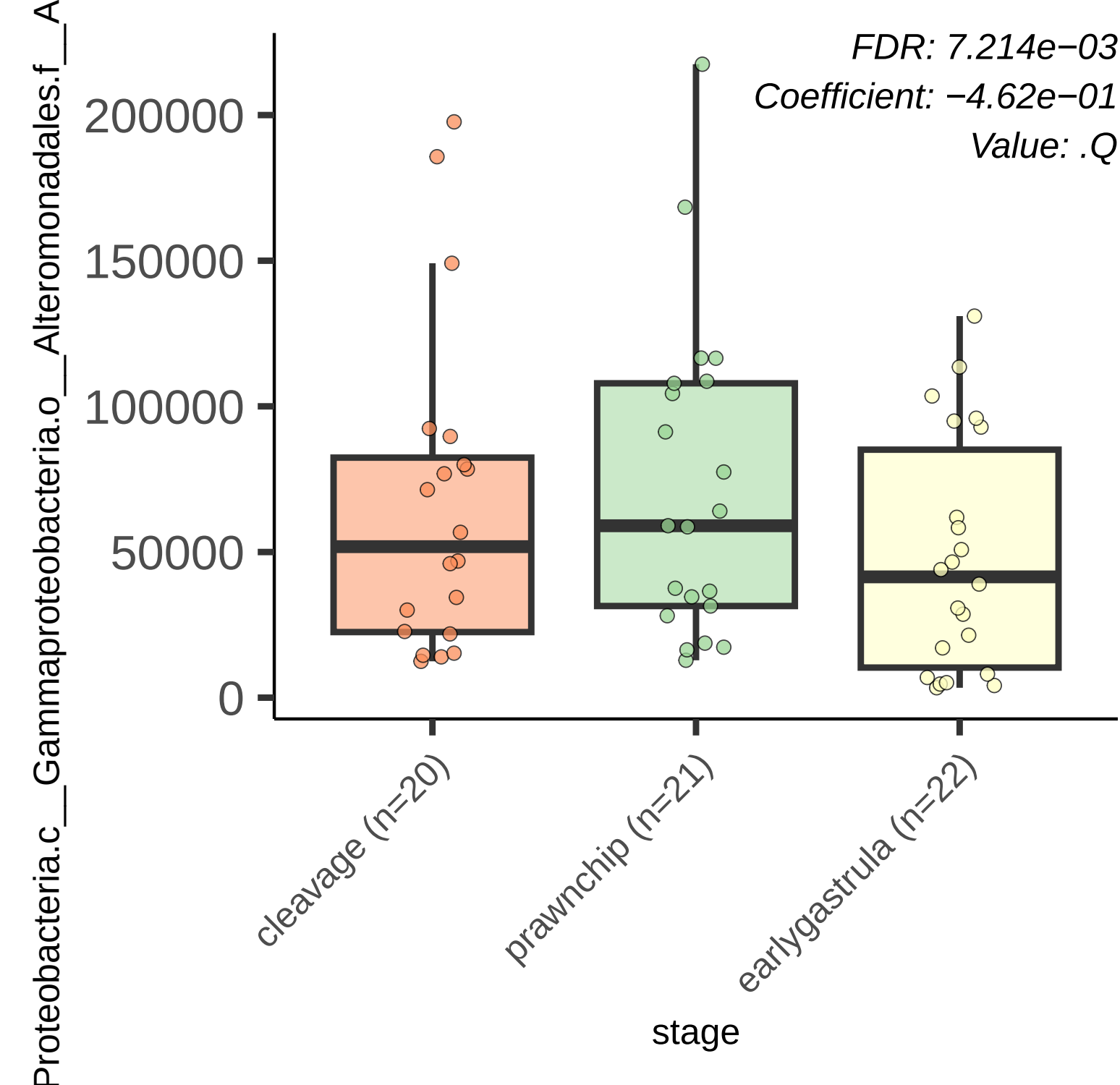
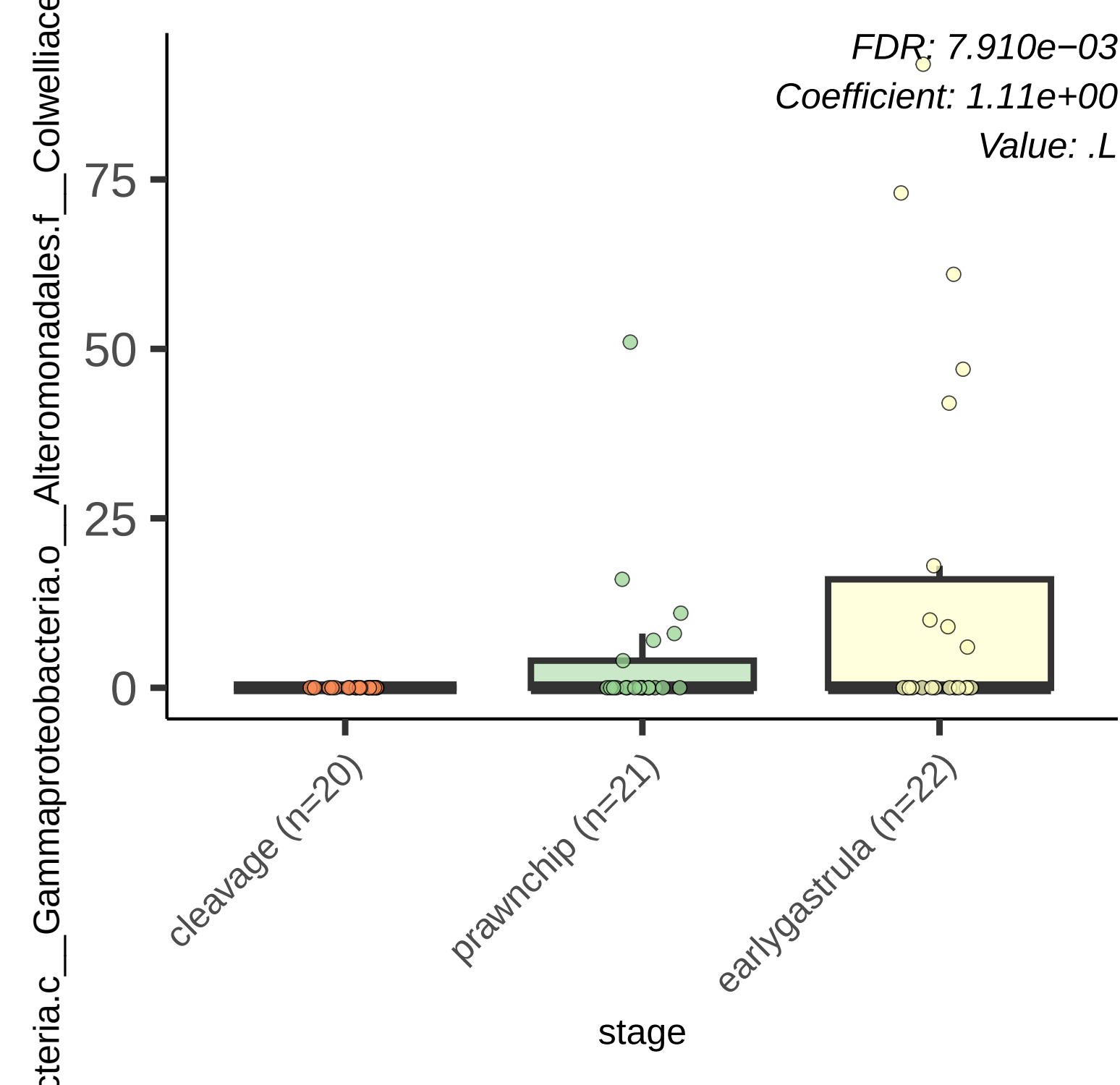
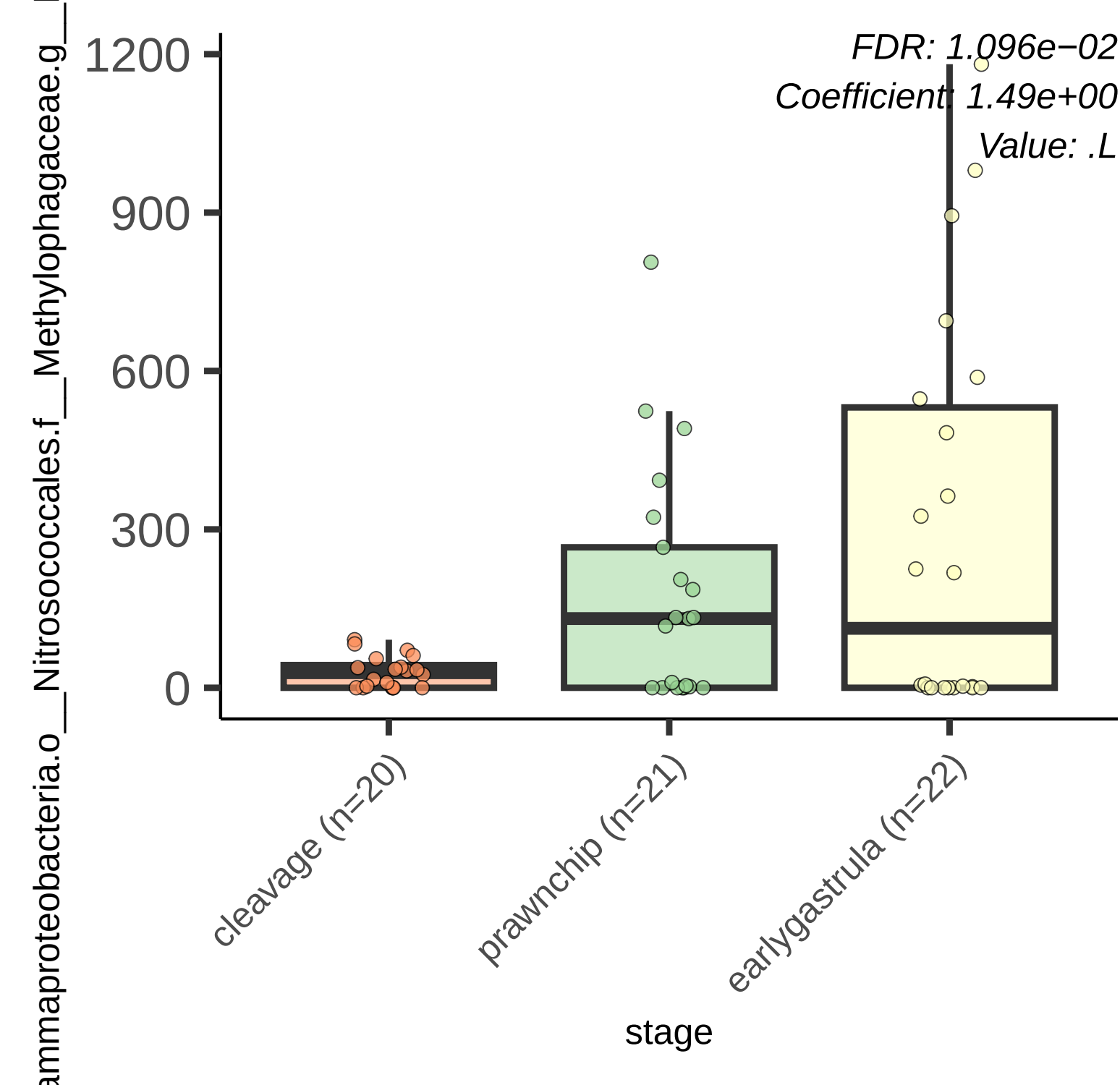
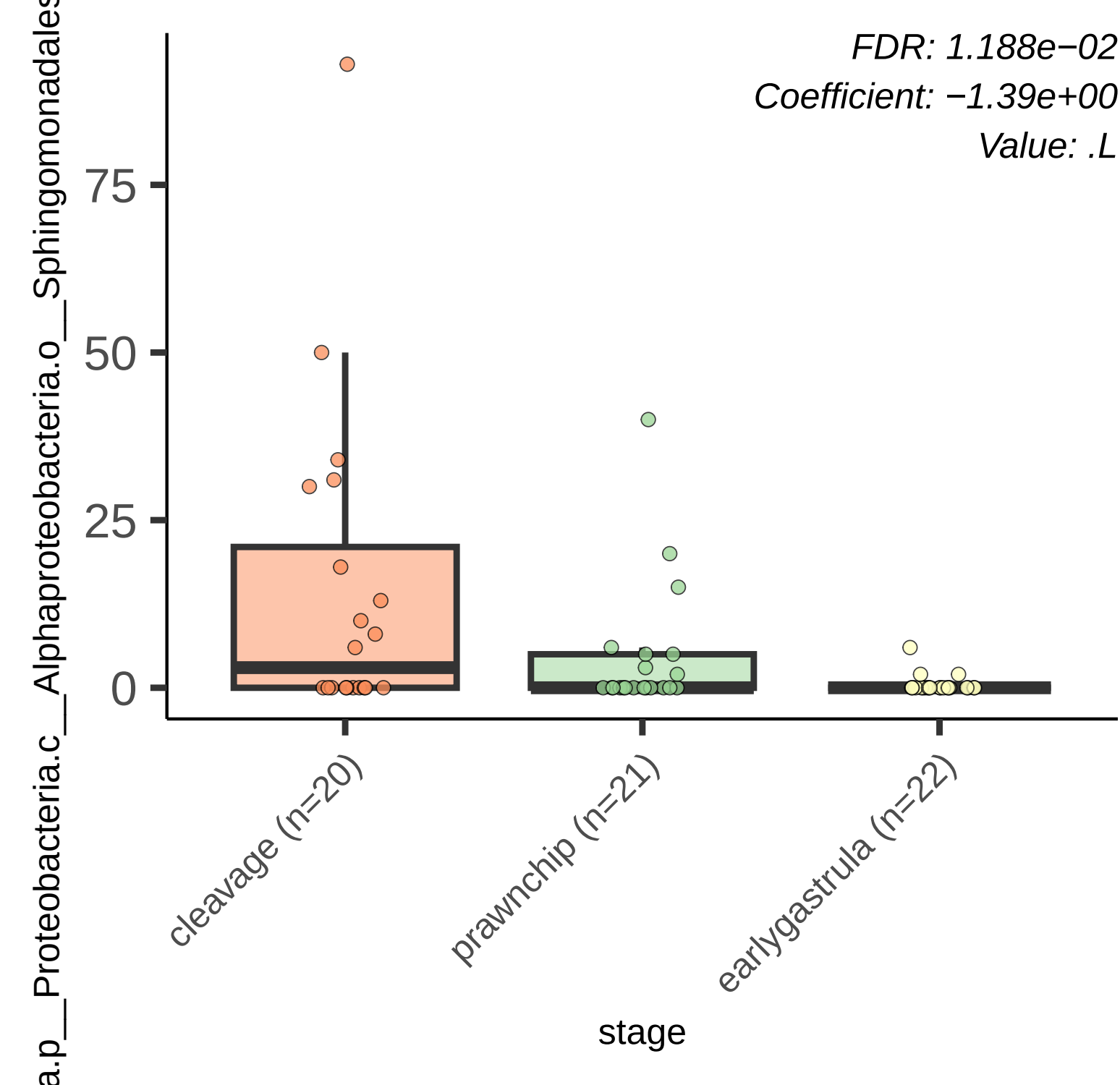
Coefficient: 1.19e+00
Value: .L
200
150
100
50
0
cleavage (n=20)
prawnchip (n=21)
earlygastrula (n=22)
stage
a.c__Gammaproteobacteria.o__Cellvibrionales.f__Cellvibrionaceae.


FDR: 7.214e−03
Coefficient: −4.62e−01
Value: .Q
200000
150000
100000
50000
0
cleavage (n=20)
prawnchip (n=21)
earlygastrula (n=22)
stage
Proteobacteria.c__Gammaproteobacteria.o__Alteromonadales.f__A


FDR: 7.910e−03
Coefficient: 1.11e+00
Value: .L
75
50
25
0
cleavage (n=20)
prawnchip (n=21)
earlygastrula (n=22)
stage
cteria.c__Gammaproteobacteria.o__Alteromonadales.f__Colwelliace


FDR: 1.096e−02
Coefficient: 1.49e+00
Value: .L
1200
900
600
300
0
cleavage (n=20)
prawnchip (n=21)
earlygastrula (n=22)
stage
ammaproteobacteria.o__Nitrosococcales.f__Methylophagaceae.g__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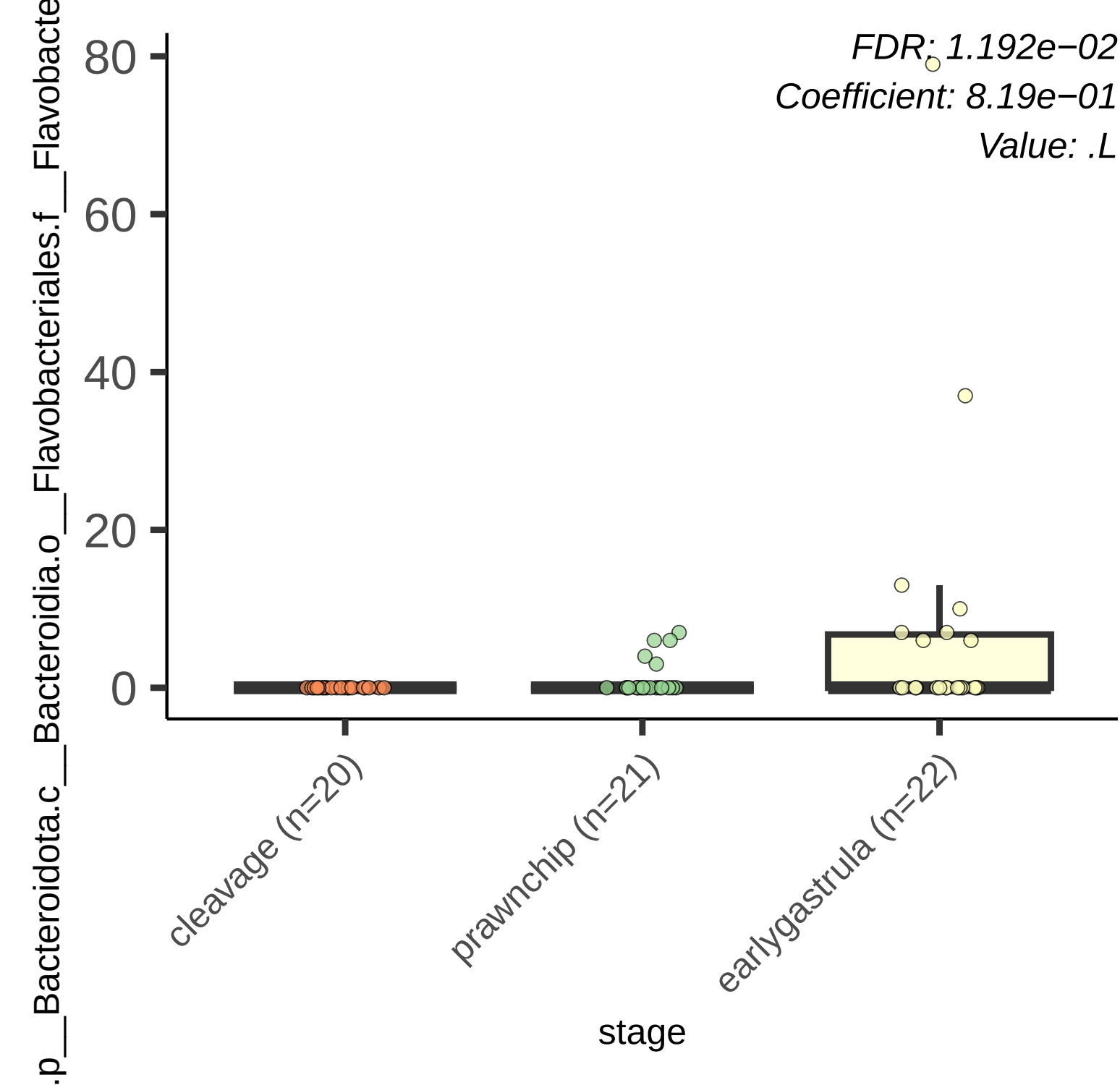
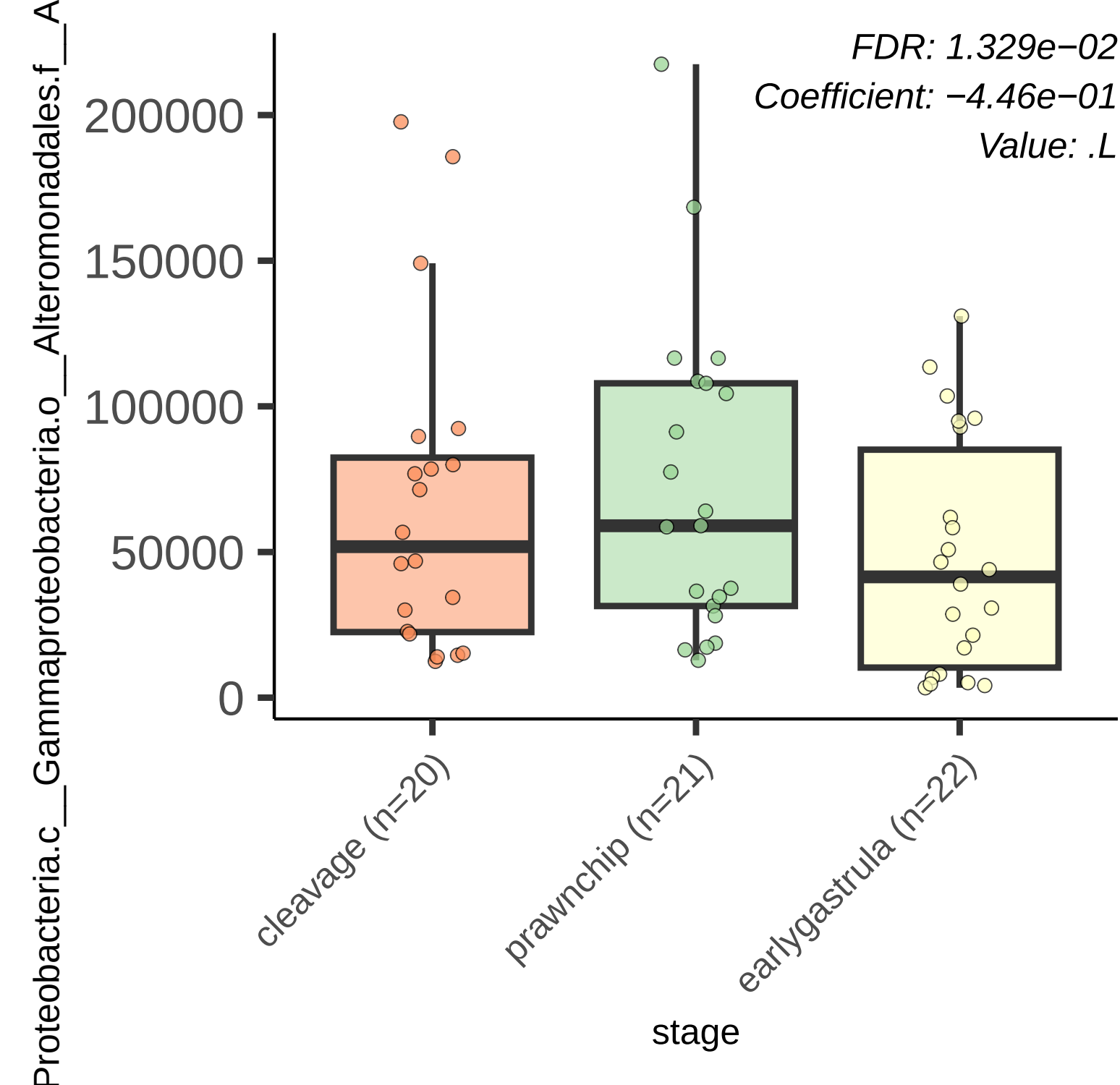
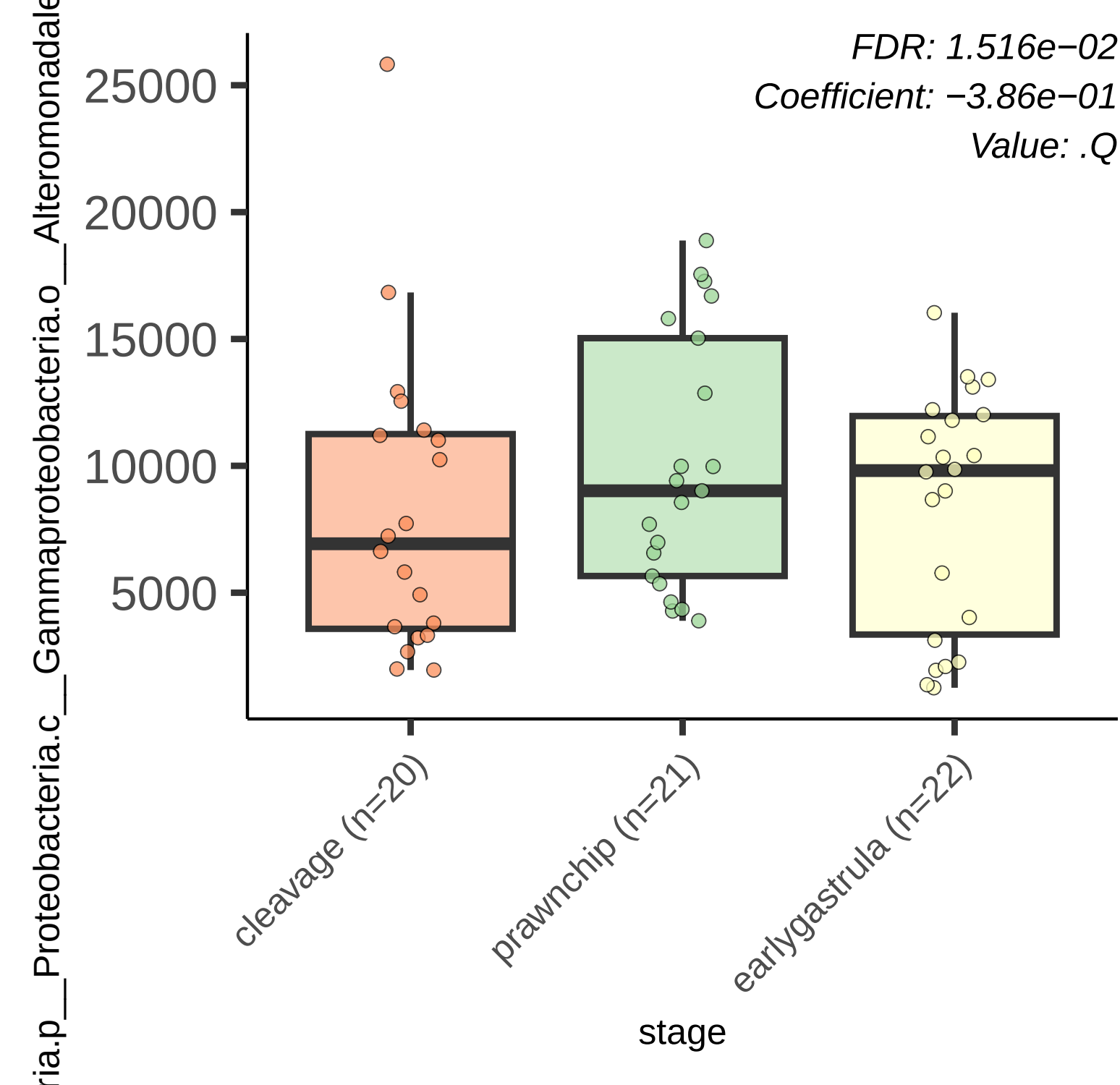
Coefficient: −1.39e+00
Value: .L
75
50
25
0
cleavage (n=20)
prawnchip (n=21)
earlygastrula (n=22)
stage
a.p__Proteobacteria.c__Alphaproteobacteria.o__Sphingomonadales


FDR: 1.192e−02
Coefficient: 8.19e−01
Value: .L
80
60
40
20
0
cleavage (n=20)
prawnchip (n=21)
earlygastrula (n=22)
stage
.p__Bacteroidota.c__Bacteroidia.o__Flavobacteriales.f__Flavobacte


FDR: 1.329e−02
Coefficient: −4.46e−01
Value: .L
200000
150000
100000
50000
0
cleavage (n=20)
prawnchip (n=21)
earlygastrula (n=22)
stage
Proteobacteria.c__Gammaproteobacteria.o__Alteromonadales.f__A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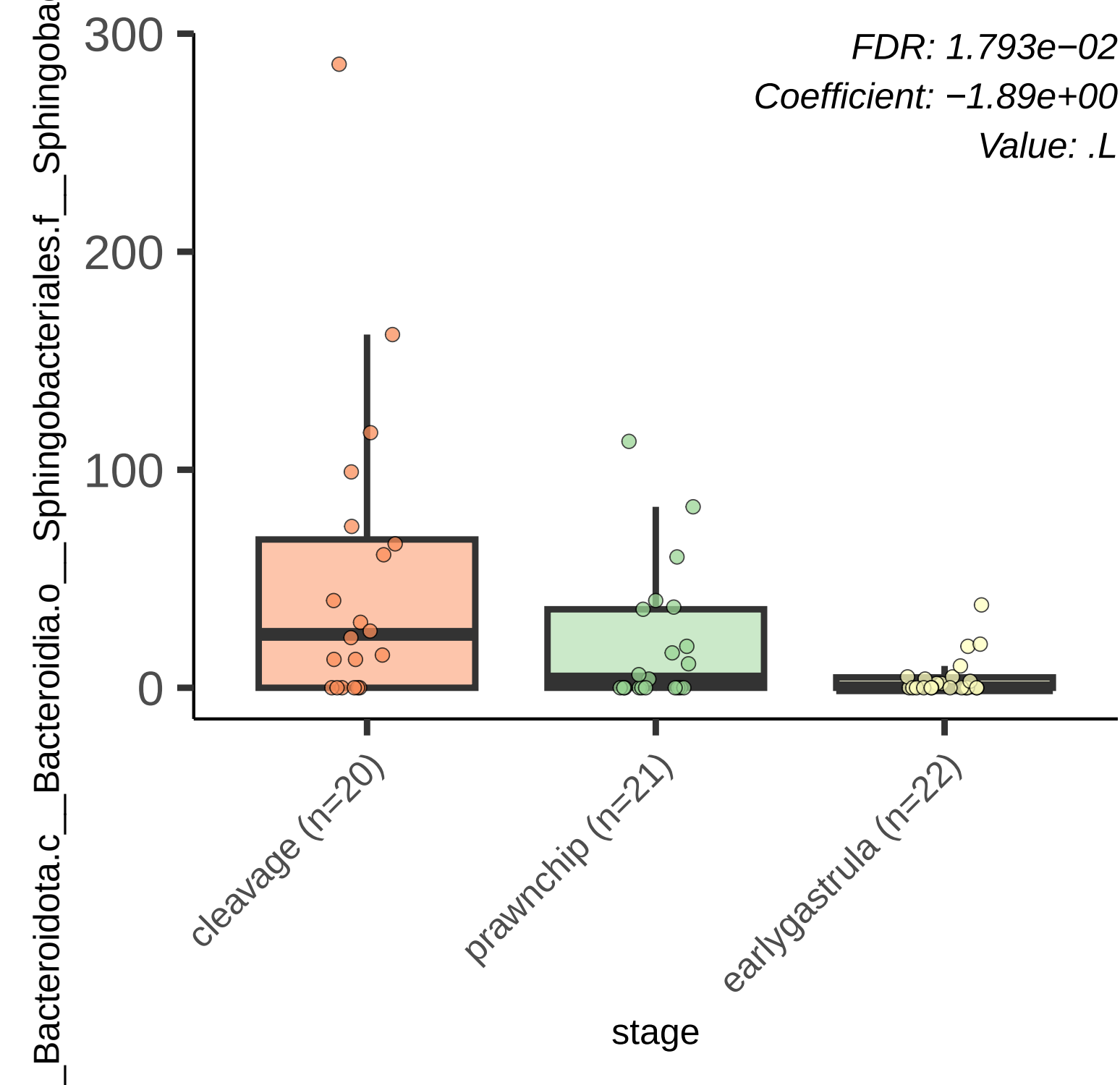
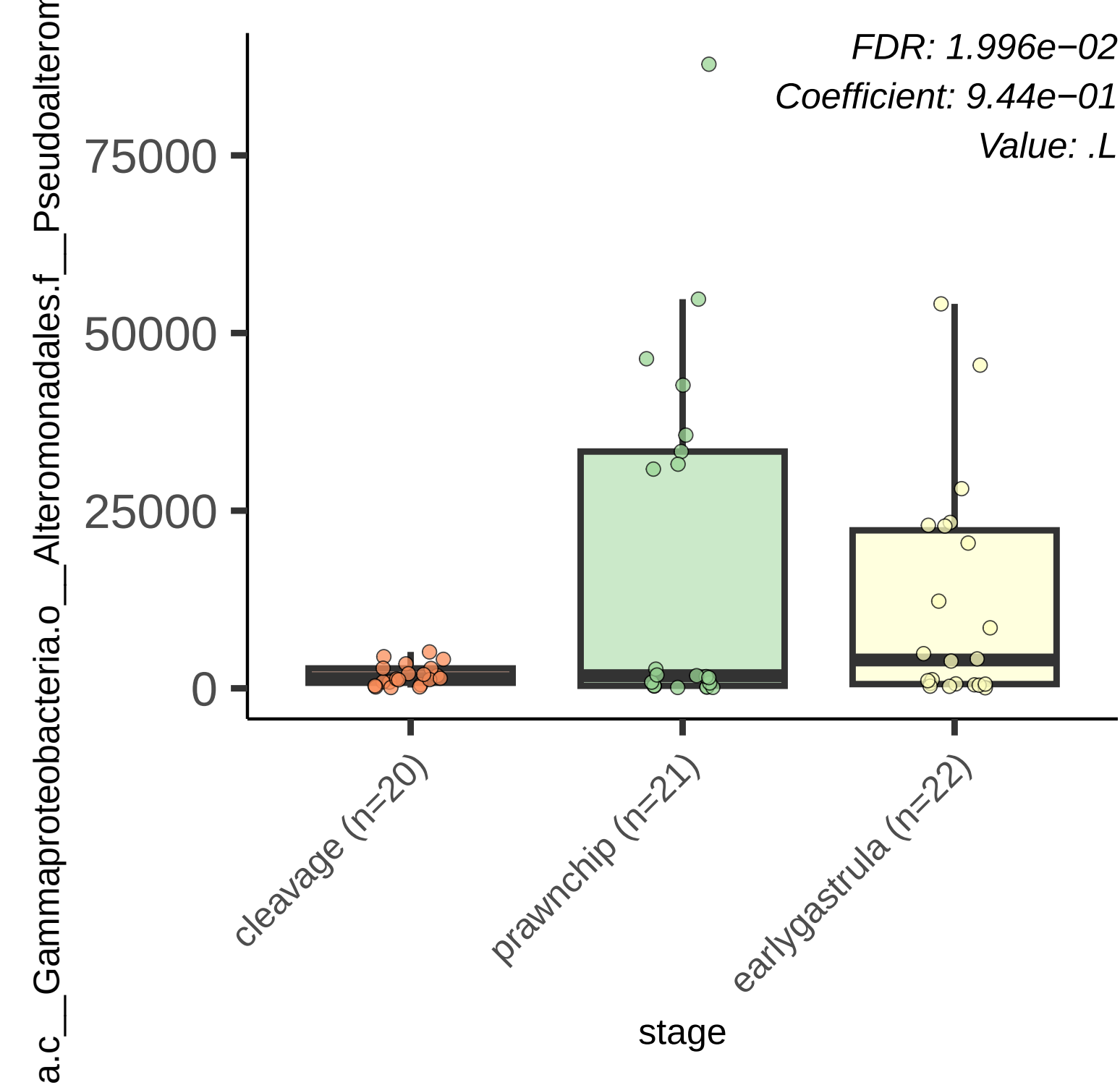
Coefficient: −3.86e−01
Value: .Q
25000
20000
15000
10000
5000
cleavage (n=20)
prawnchip (n=21)
earlygastrula (n=22)
stage
ria.p__Proteobacteria.c__Gammaproteobacteria.o__Alteromonadale


FDR: 1.793e−02
Coefficient: −1.89e+00
Value: .L
300
200
100
0
cleavage (n=20)
prawnchip (n=21)
earlygastrula (n=22)
stage
__Bacteroidota.c__Bacteroidia.o__Sphingobacteriales.f__Sphingoba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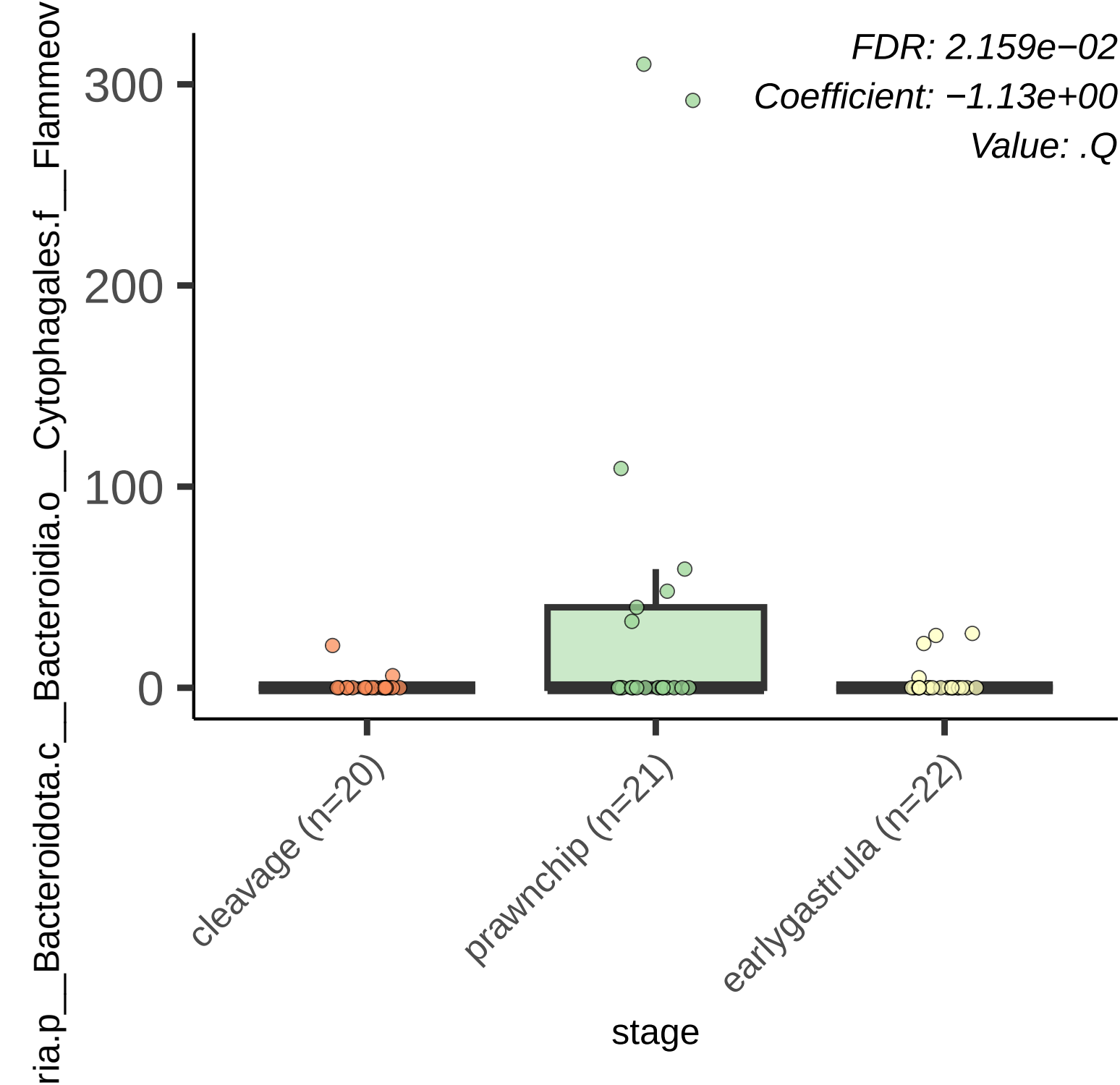
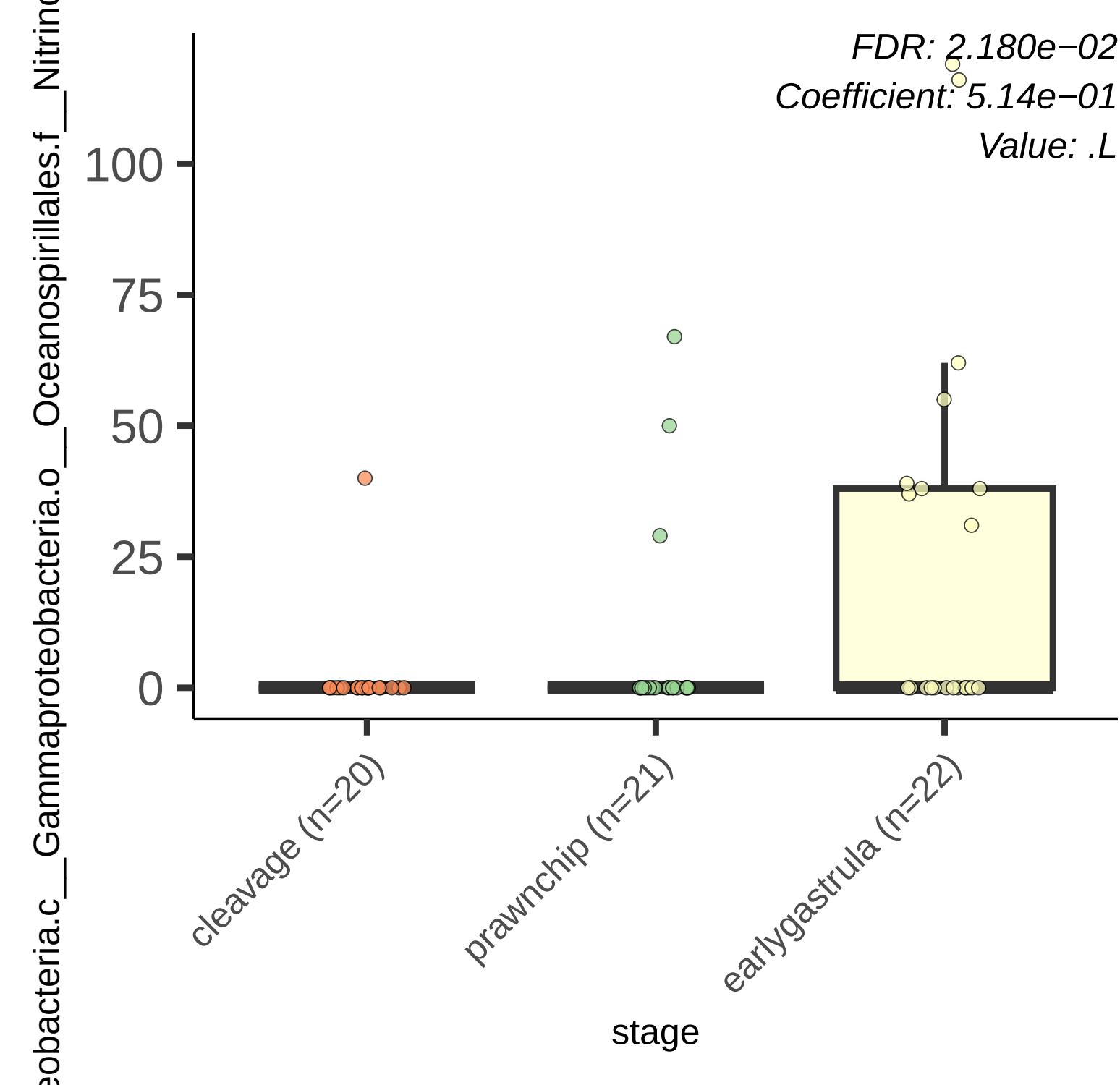
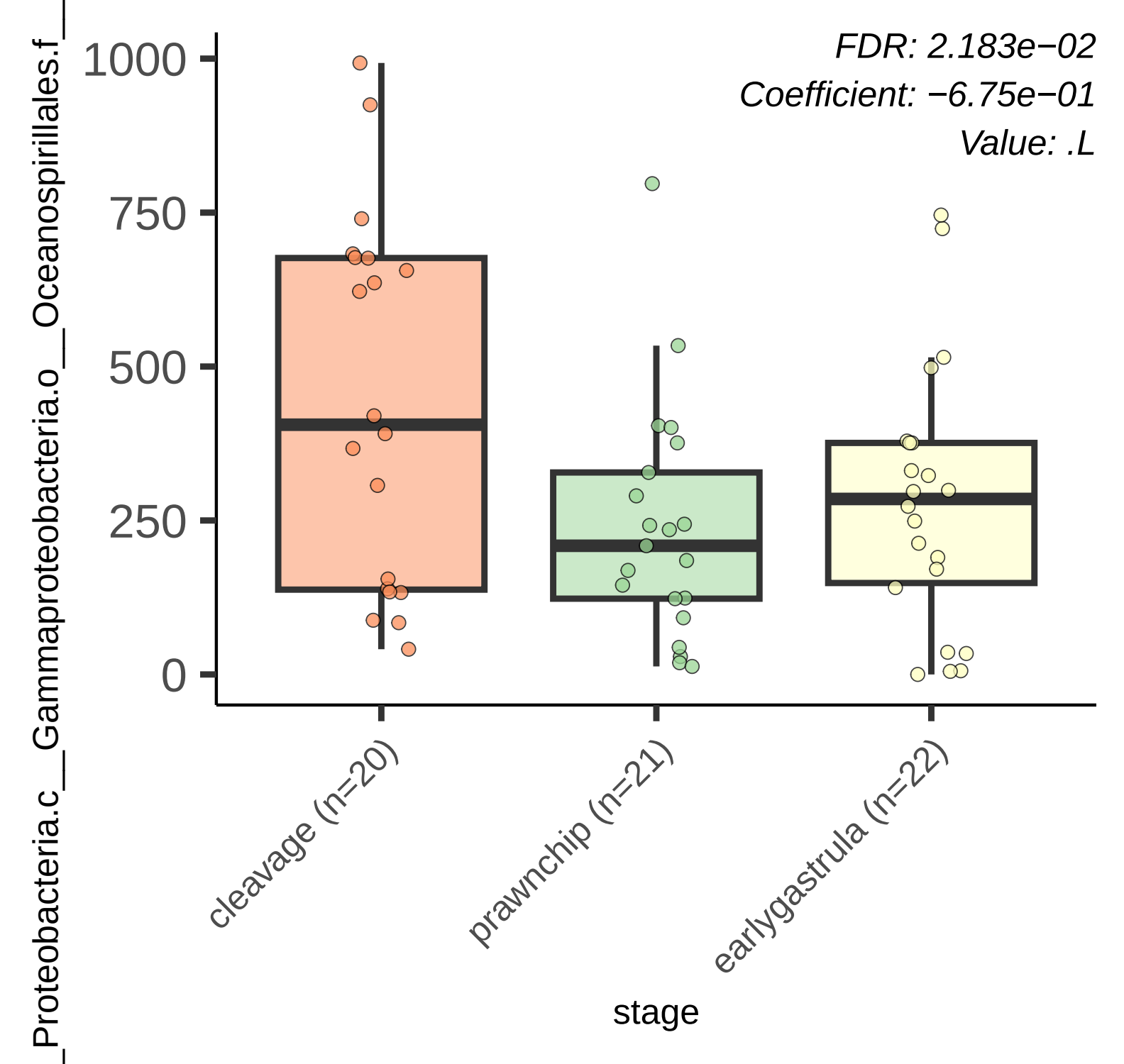
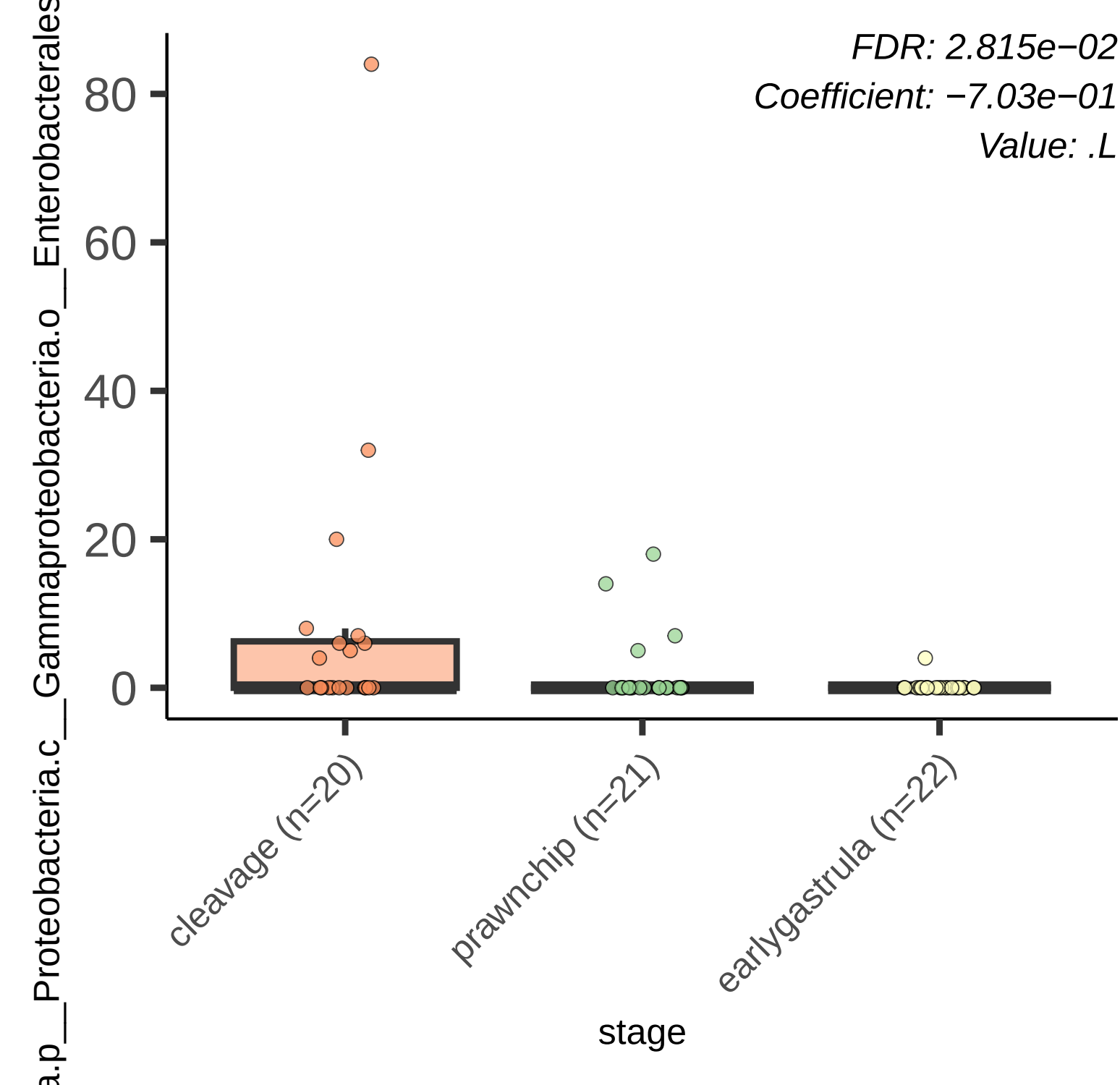
Coefficient: 9.44e−01
Value: .L
75000
50000
25000
0
cleavage (n=20)
prawnchip (n=21)
earlygastrula (n=22)
stage
a.c__Gammaproteobacteria.o__Alteromonadales.f__Pseudoalterom

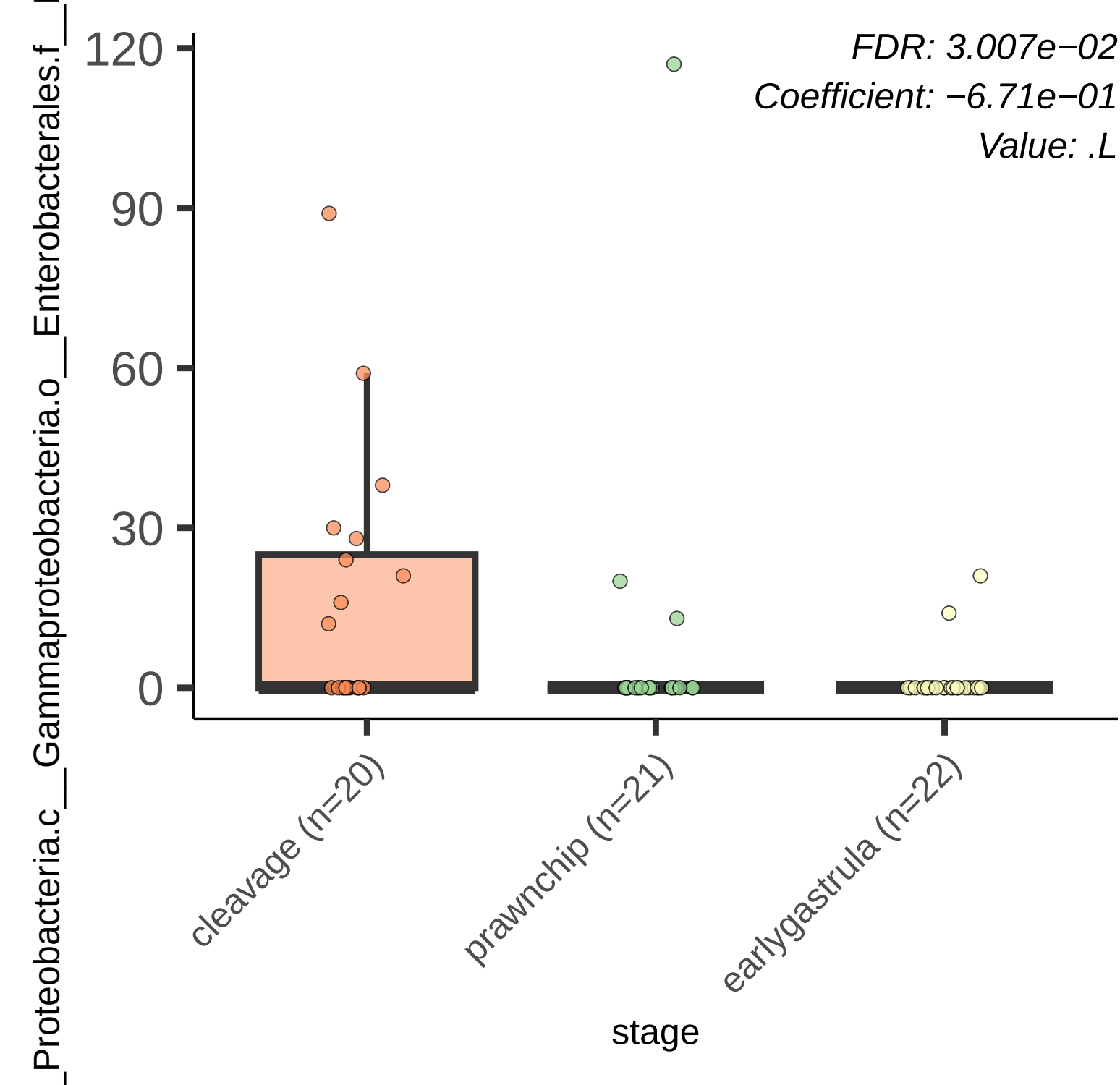

FDR: 2.159e−02
Coefficient: −1.13e+00
Value: .Q
300
200
100
0
cleavage (n=20)
prawnchip (n=21)
earlygastrula (n=22)
stage
ria.p__Bacteroidota.c__Bacteroidia.o__Cytophagales.f__Flammeov

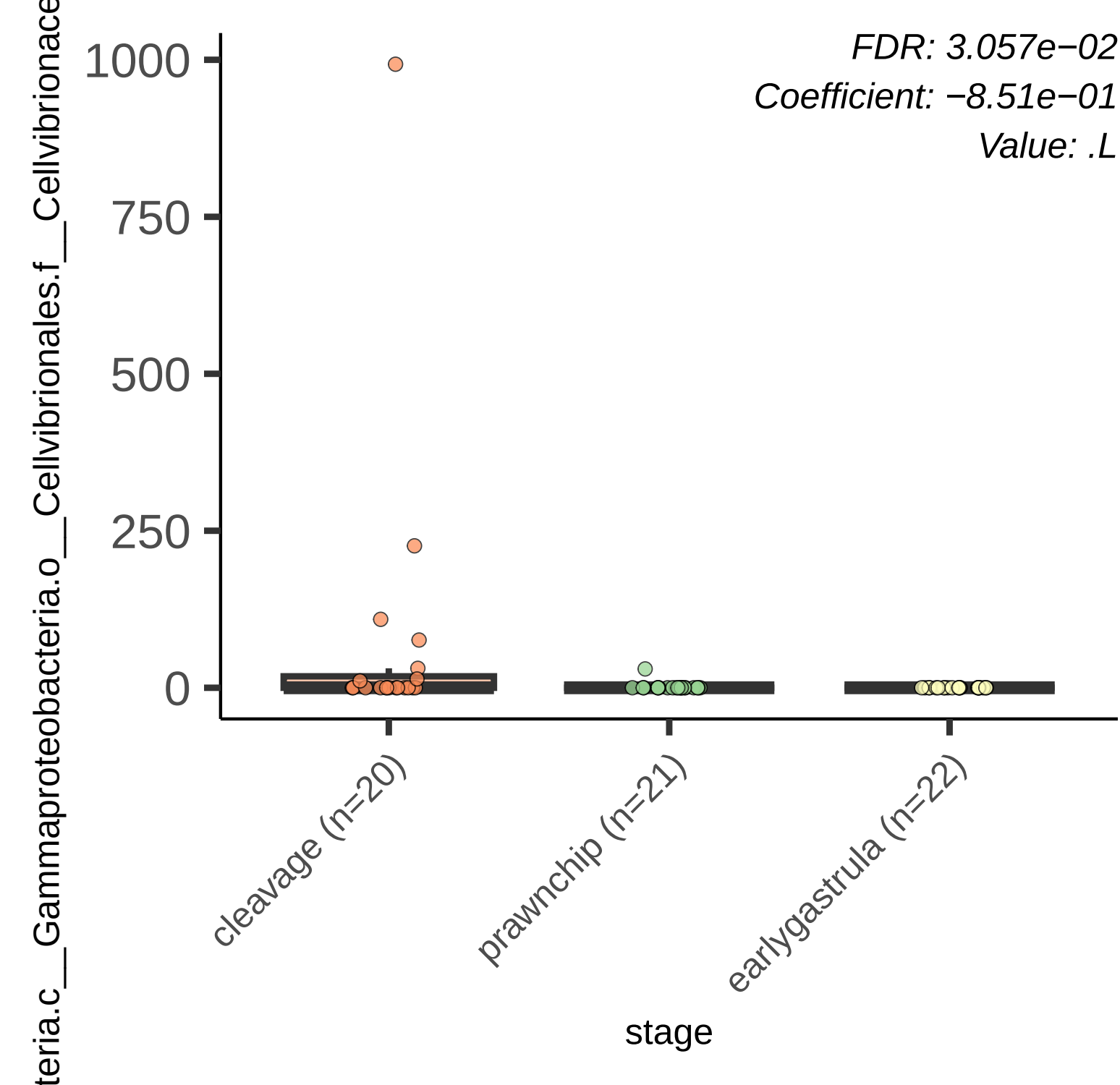

FDR: 2.180e−02
Coefficient: 5.14e−01
Value: .L
100
75
50
25
0
cleavage (n=20)
prawnchip (n=21)
earlygastrula (n=22)
stage
eobacteria.c__Gammaproteobacteria.o__Oceanospirillales.f__Nitrinc


FDR: 2.183e−02
Coefficient: −6.75e−01
Value: .L
1000
750
500
250
0
cleavage (n=20)
prawnchip (n=21)
earlygastrula (n=22)
stage
_Proteobacteria.c__Gammaproteobacteria.o__Oceanospirillales.f__

FDR: 2.479e−02
Coefficient: 8.42e−01
Value: .L
300
200
100
0
cleavage (n=20)
prawnchip (n=21)
earlygastrula (n=22)
stage
teobacteria.c__Gammaproteobacteria.o__Alteromonadales.f__Colw


FDR: 2.815e−02
Coefficient: −7.03e−01
Value: .L
80
60
40
20
0
cleavage (n=20)
prawnchip (n=21)
earlygastrula (n=22)
stage
a.p__Proteobacteria.c__Gammaproteobacteria.o__Enterobacterales


FDR: 3.007e−02
Coefficient: −6.71e−01
Value: .L
120
90
60
30
0
cleavage (n=20)
prawnchip (n=21)
earlygastrula (n=22)
stage
_Proteobacteria.c__Gammaproteobacteria.o__Enterobacterales.f__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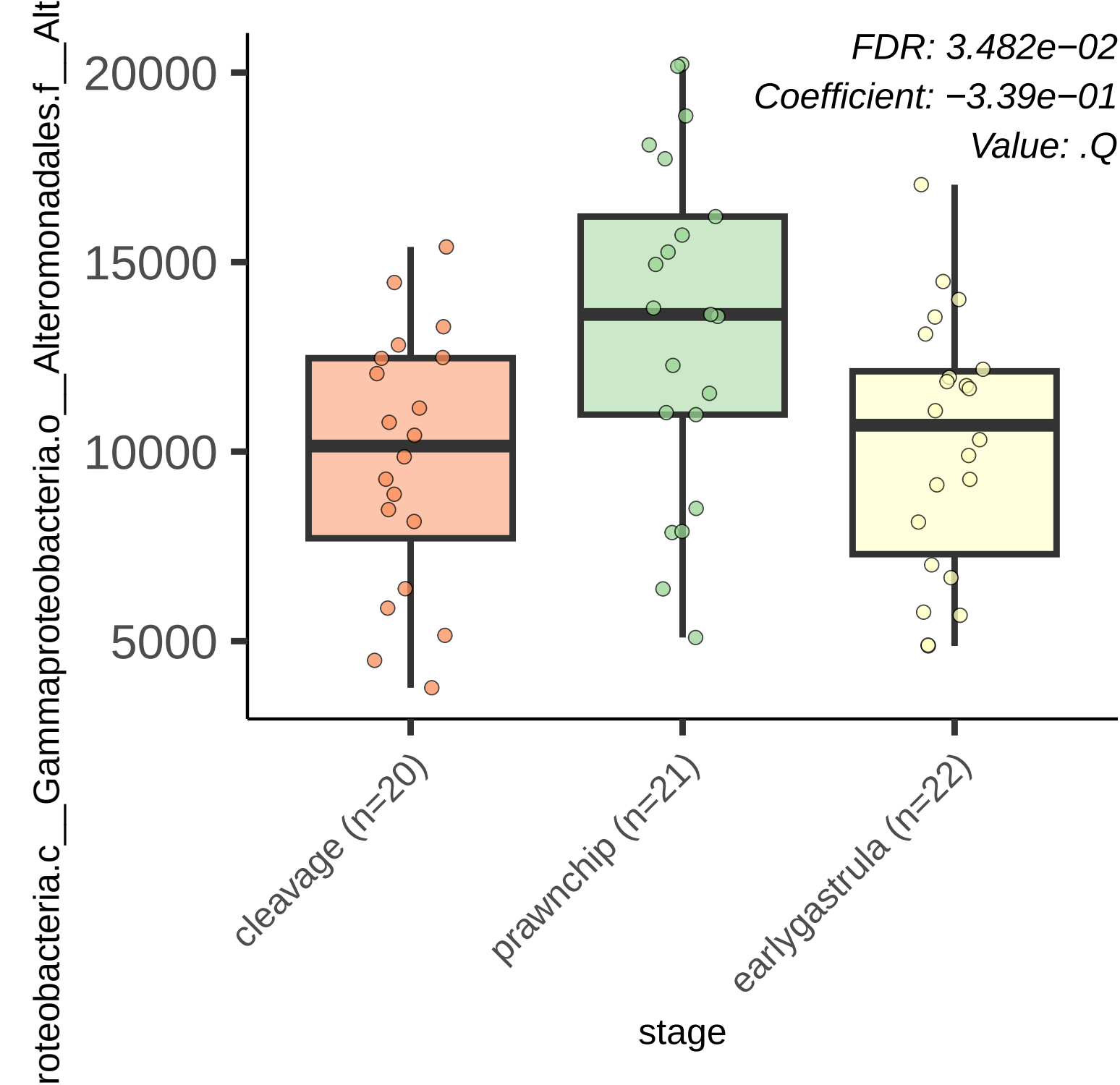
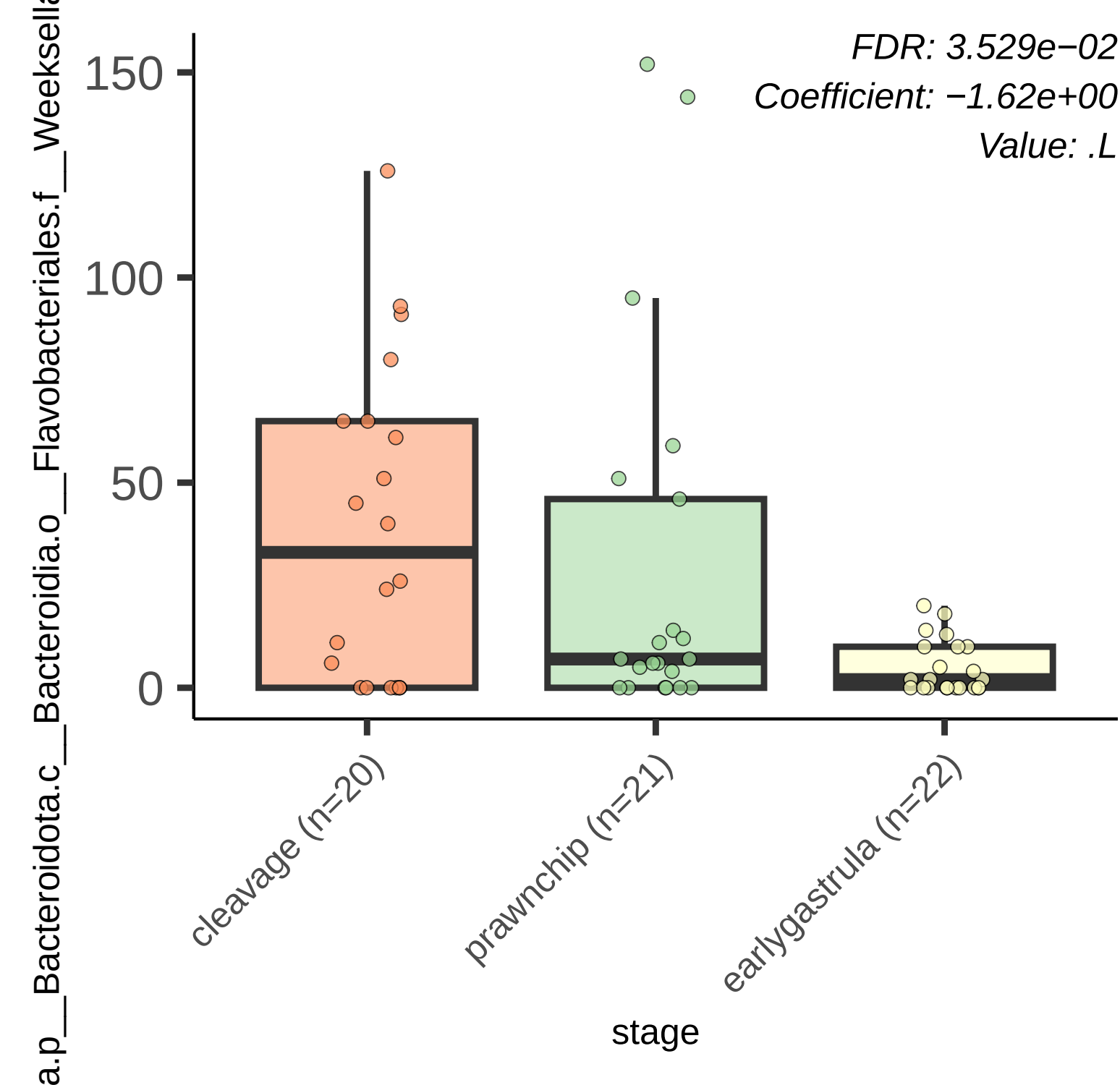
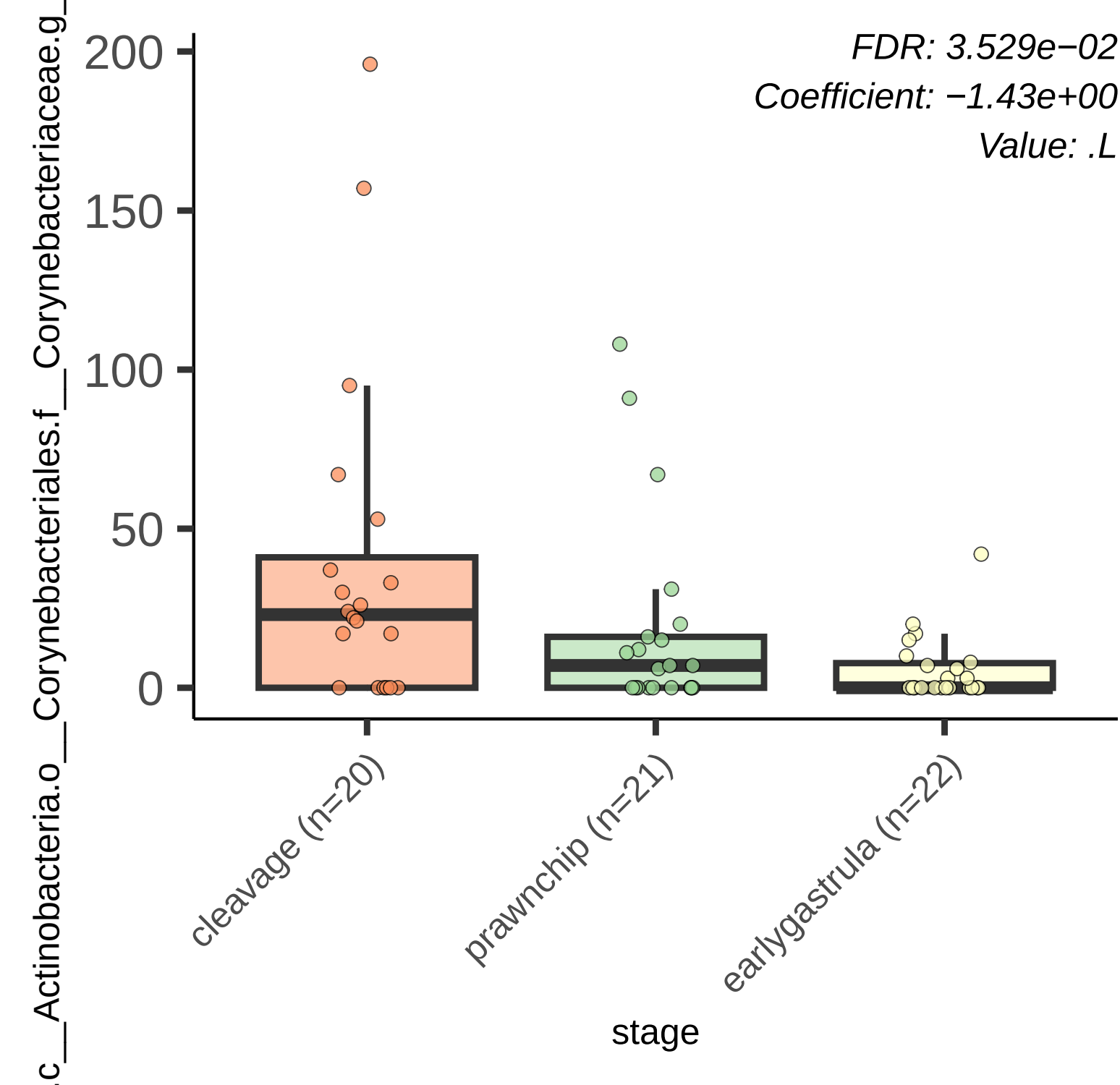
Coefficient: −8.51e−01
Value: .L
1000
750
500
250
0
cleavage (n=20)
prawnchip (n=21)
earlygastrula (n=22)
stage
teria.c__Gammaproteobacteria.o__Cellvibrionales.f__Cellvibrionace


FDR: 3.482e−02
Coefficient: −3.39e−01
Value: .Q
20000
15000
10000
5000
cleavage (n=20)
prawnchip (n=21)
earlygastrula (n=22)
stage
roteobacteria.c__Gammaproteobacteria.o__Alteromonadales.f__Alt


FDR: 3.529e−02
Coefficient: −1.62e+00
Value: .L
150
100
50
0
cleavage (n=20)
prawnchip (n=21)
earlygastrula (n=22)
stage
a.p__Bacteroidota.c__Bacteroidia.o__Flavobacteriales.f__Weeksella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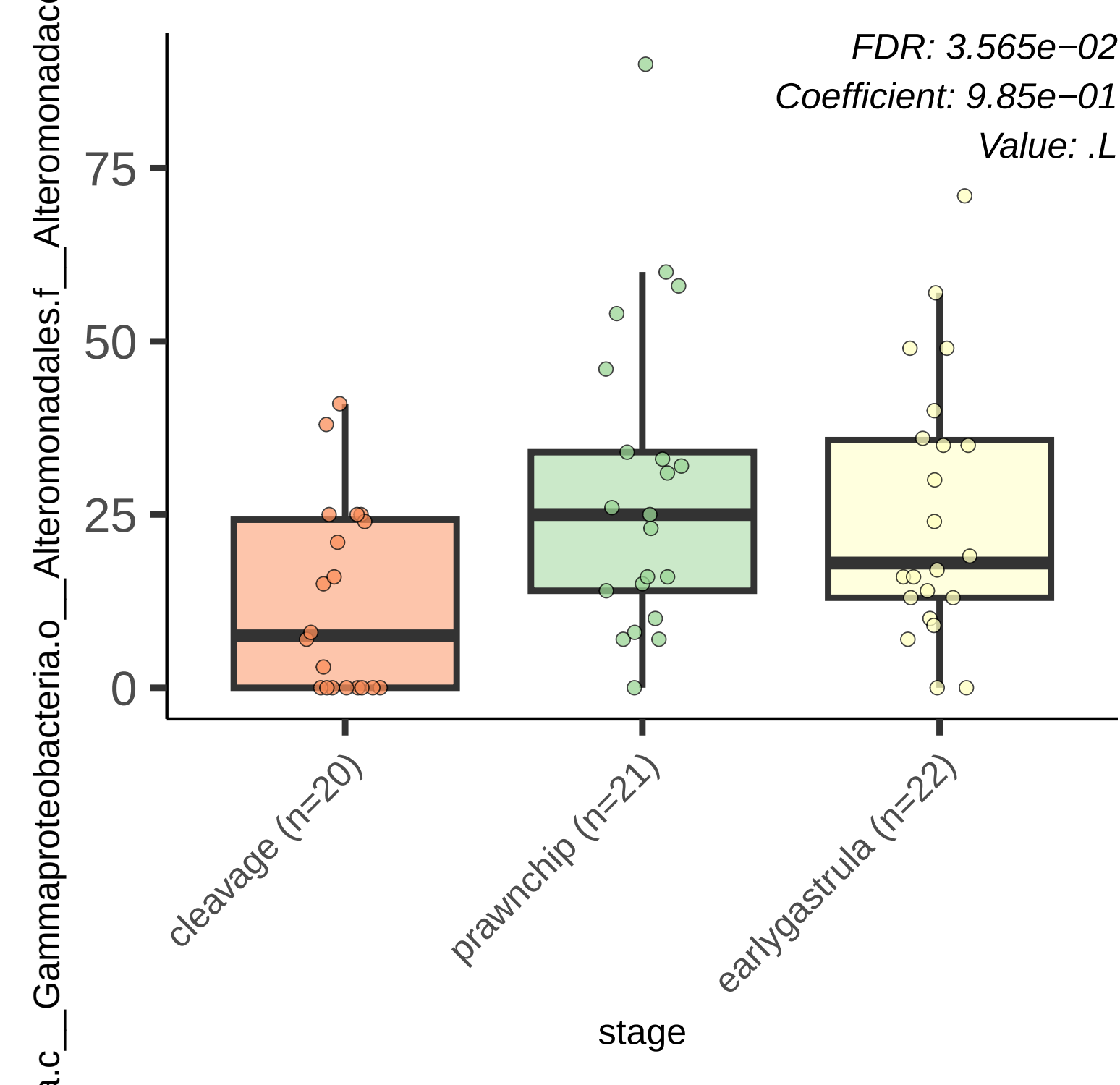
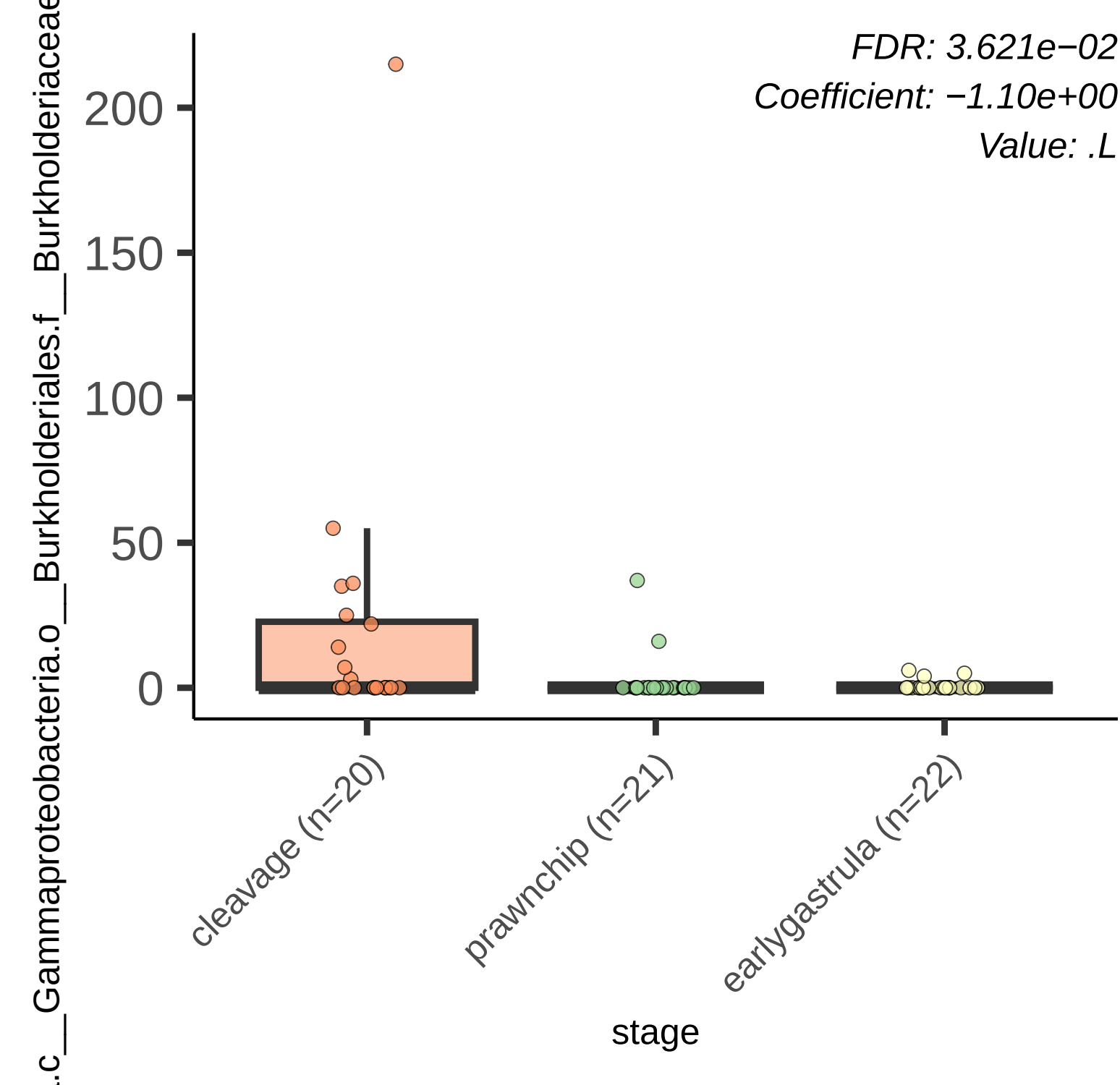
Coefficient: −1.43e+00
Value: .L
200
150
100
50
0
cleavage (n=20)
prawnchip (n=21)
earlygastrula (n=22)
stage
c__Actinobacteria.o__Corynebacteriales.f__Corynebacteriaceae.g_


FDR: 3.565e−02
Coefficient: 9.85e−01
Value: .L
75
50
25
0
cleavage (n=20)
prawnchip (n=21)
earlygastrula (n=22)
stage
a.c__Gammaproteobacteria.o__Alteromonadales.f__Alteromonadac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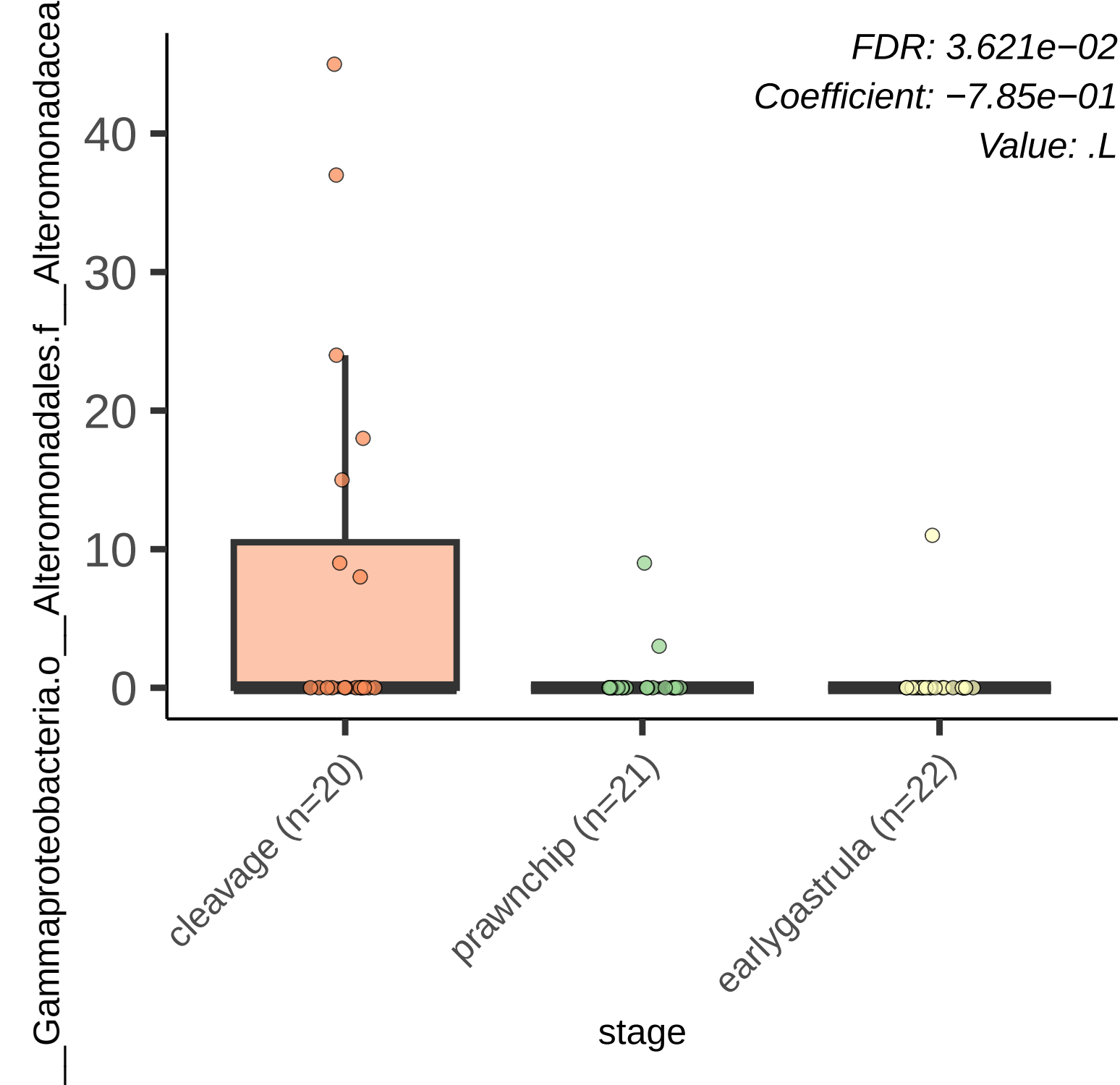
Coefficient: −1.10e+00
Value: .L
200
150
100
50
0
cleavage (n=20)
prawnchip (n=21)
earlygastrula (n=22)
stage
...c__Gammaproteobacteria.o__Burkholderiales.f__Burkholderiaceae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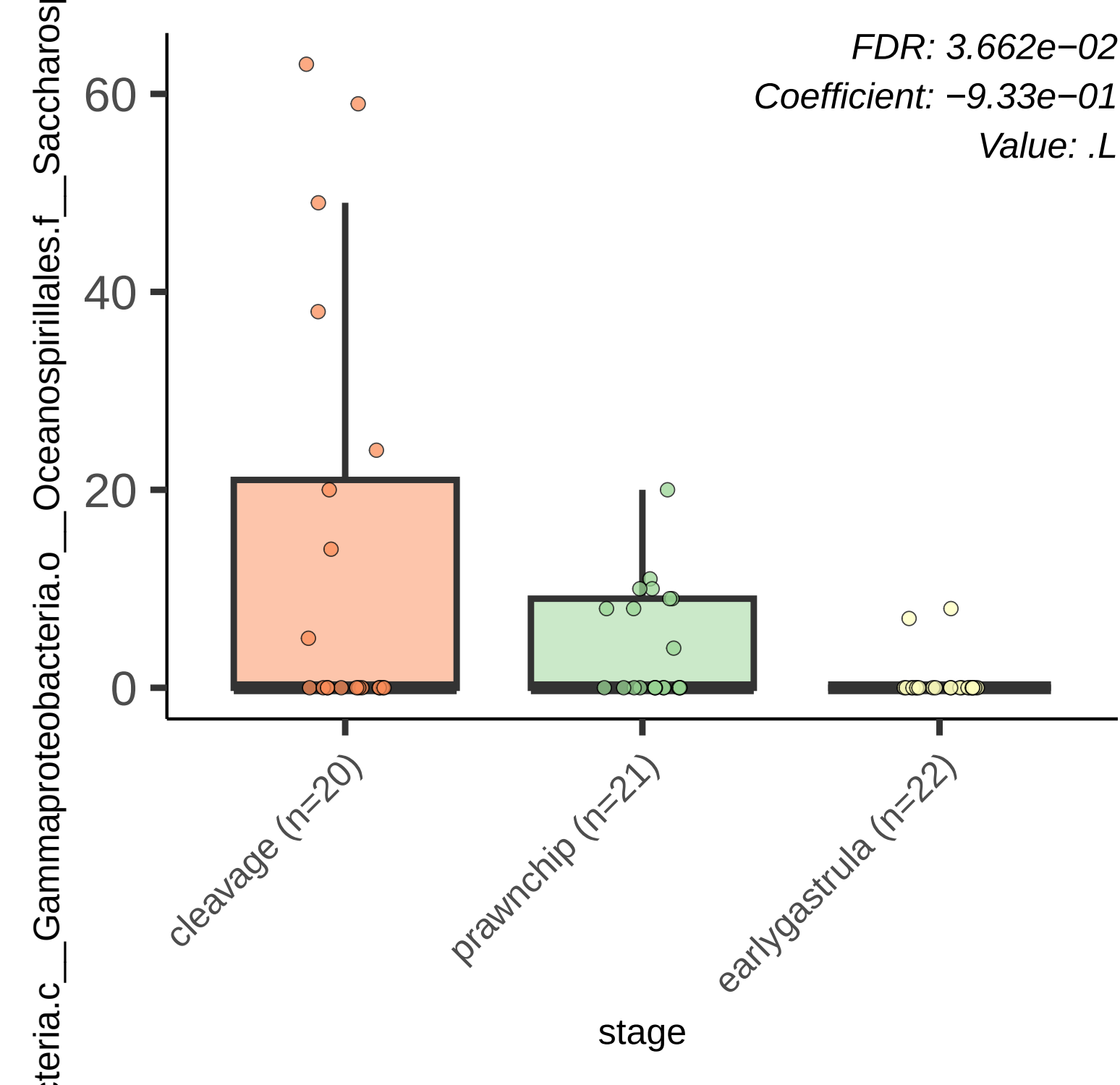
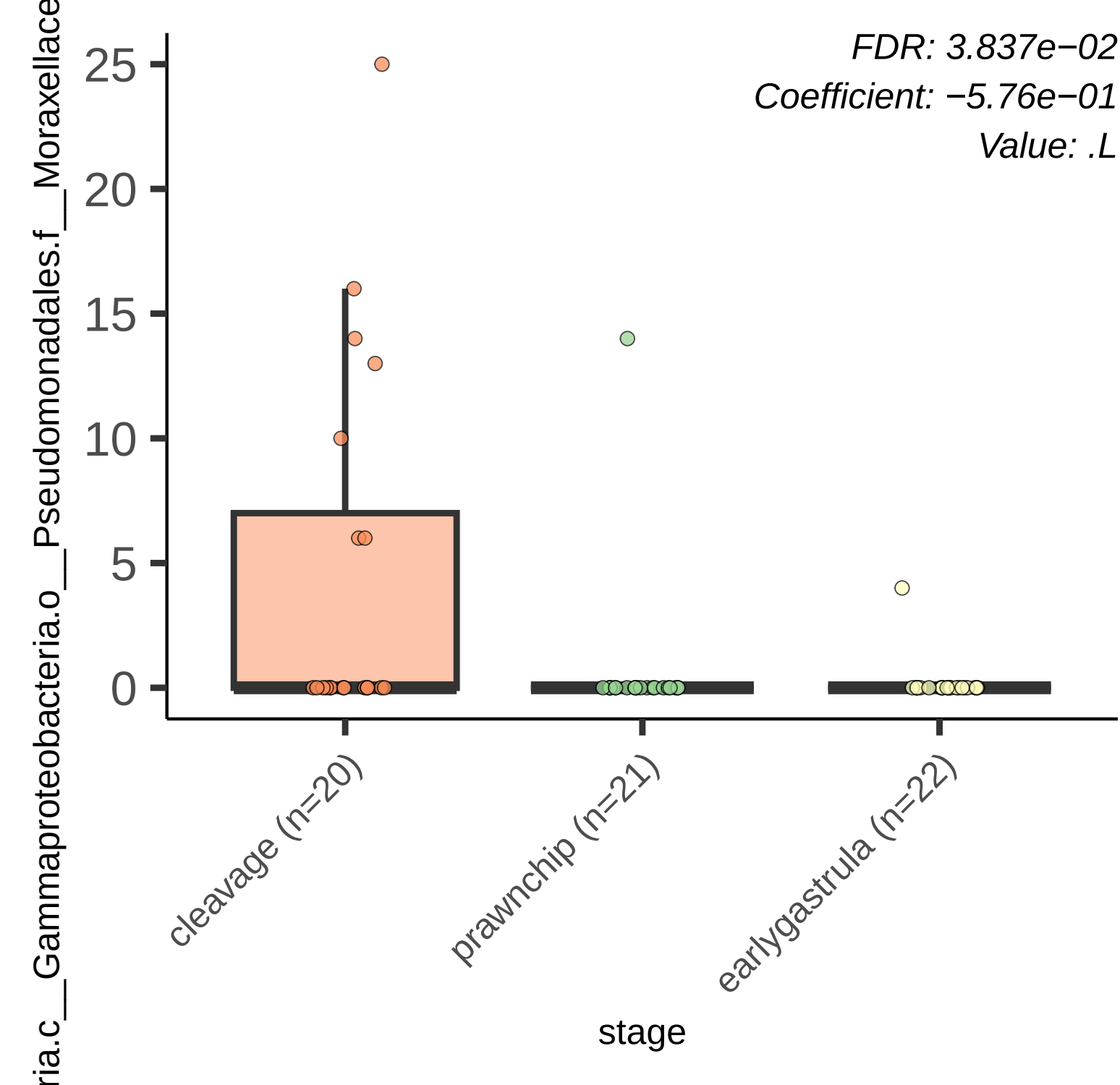
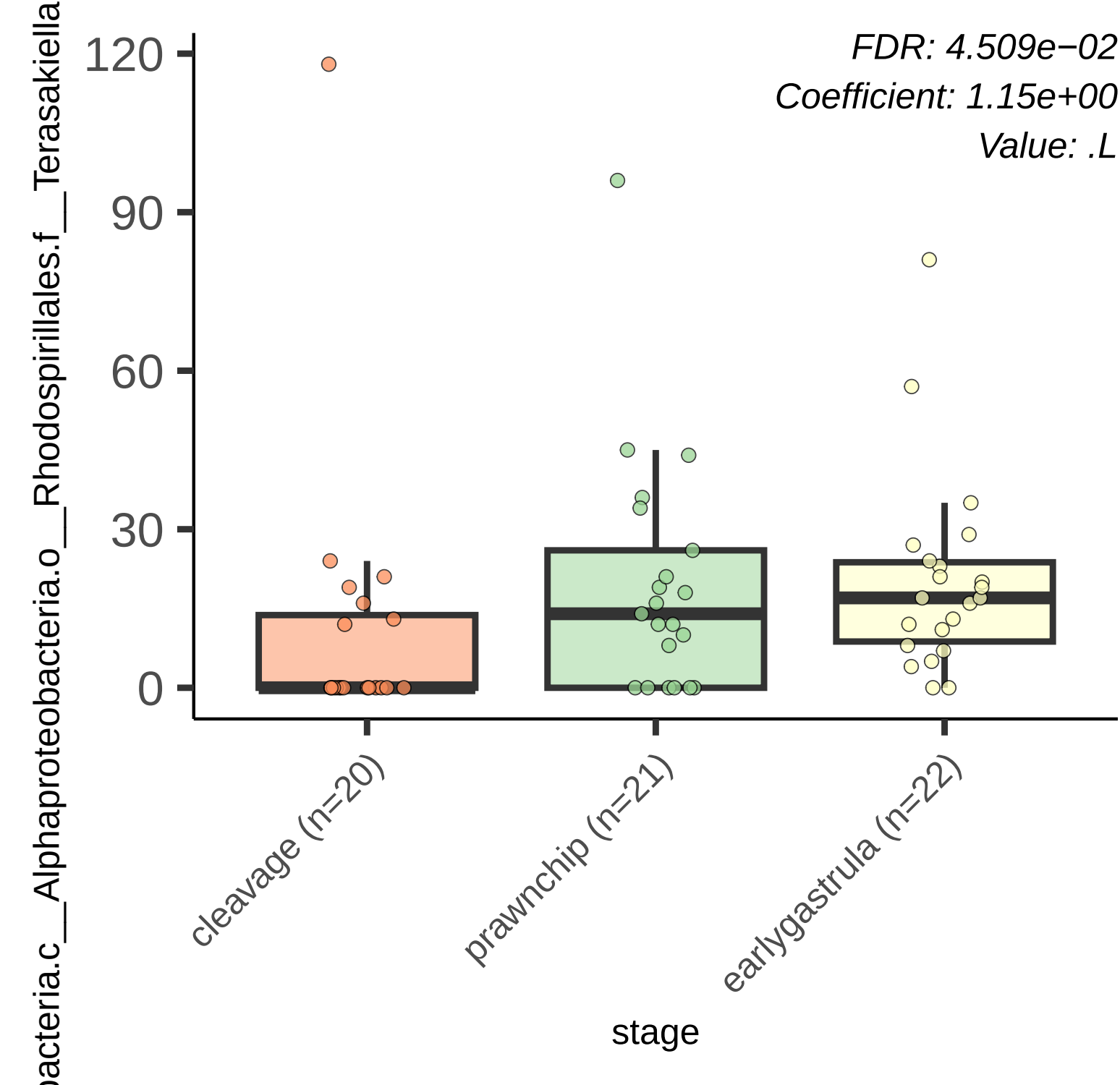
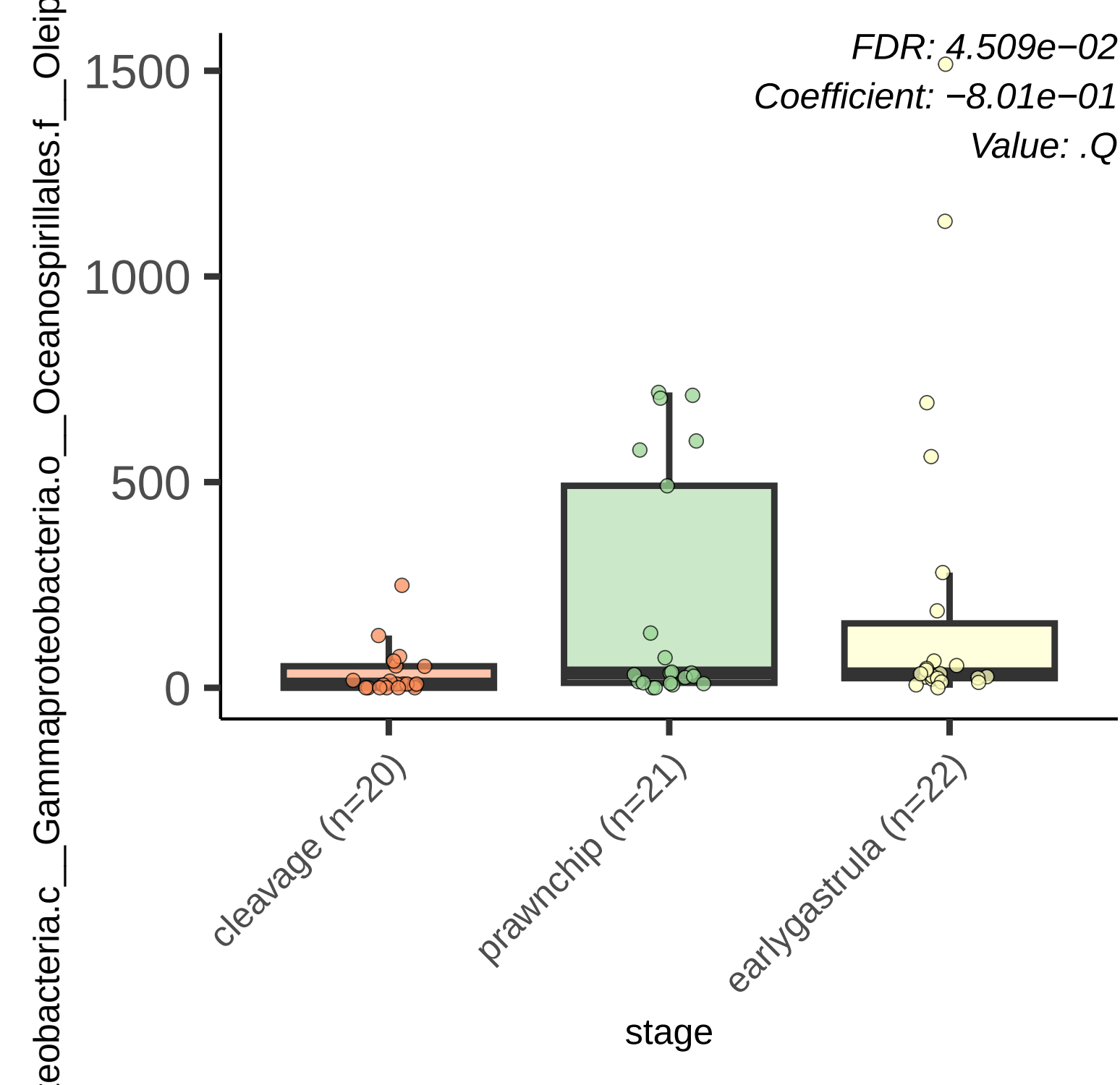
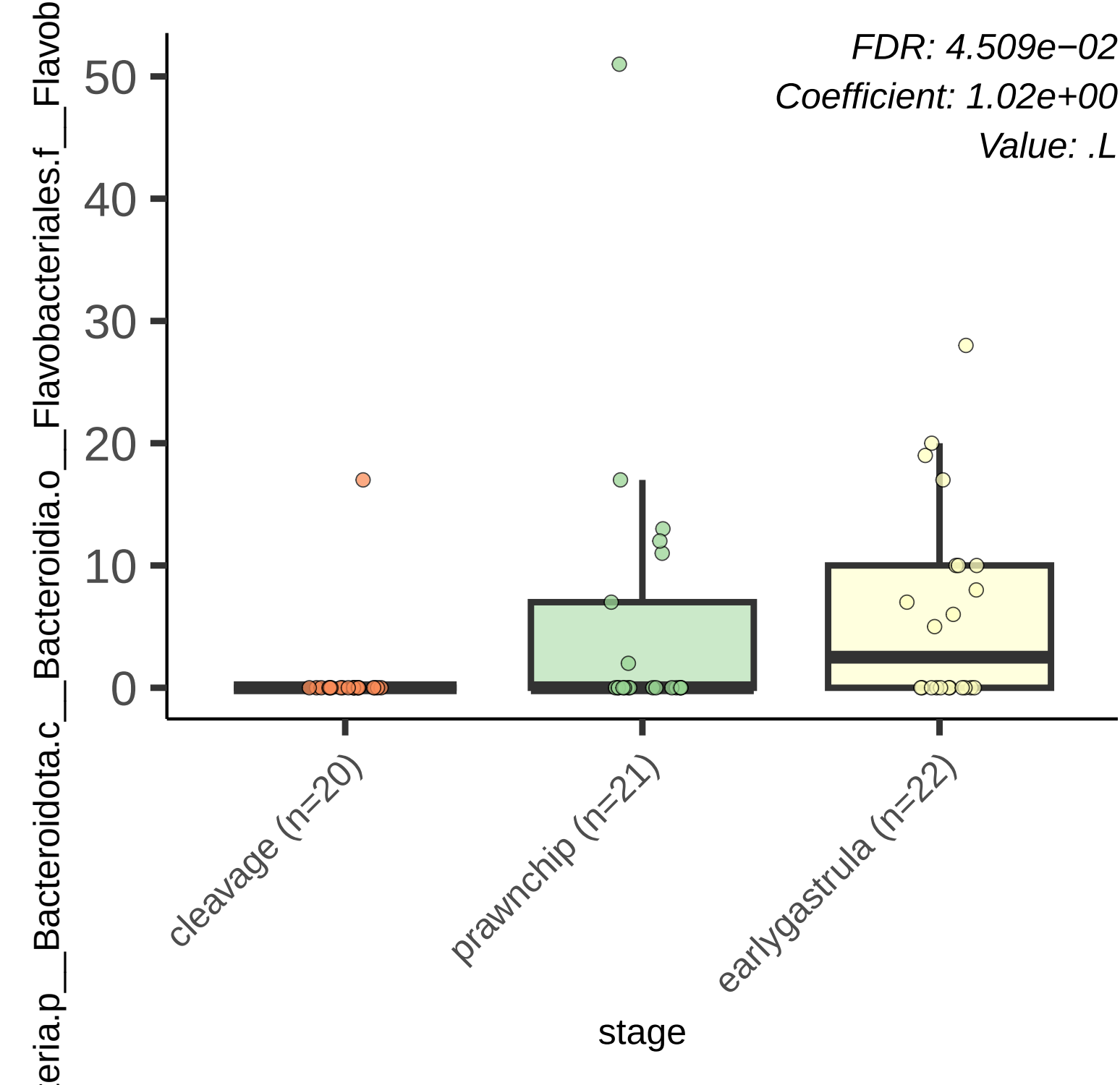
Coefficient: −7.85e−01
Value: .L
40
30
20
10
0
cleavage (n=20)
prawnchip (n=21)
earlygastrula (n=22)
stage
__Gammaproteobacteria.o__Alteromonadales.f__Alteromonadacea

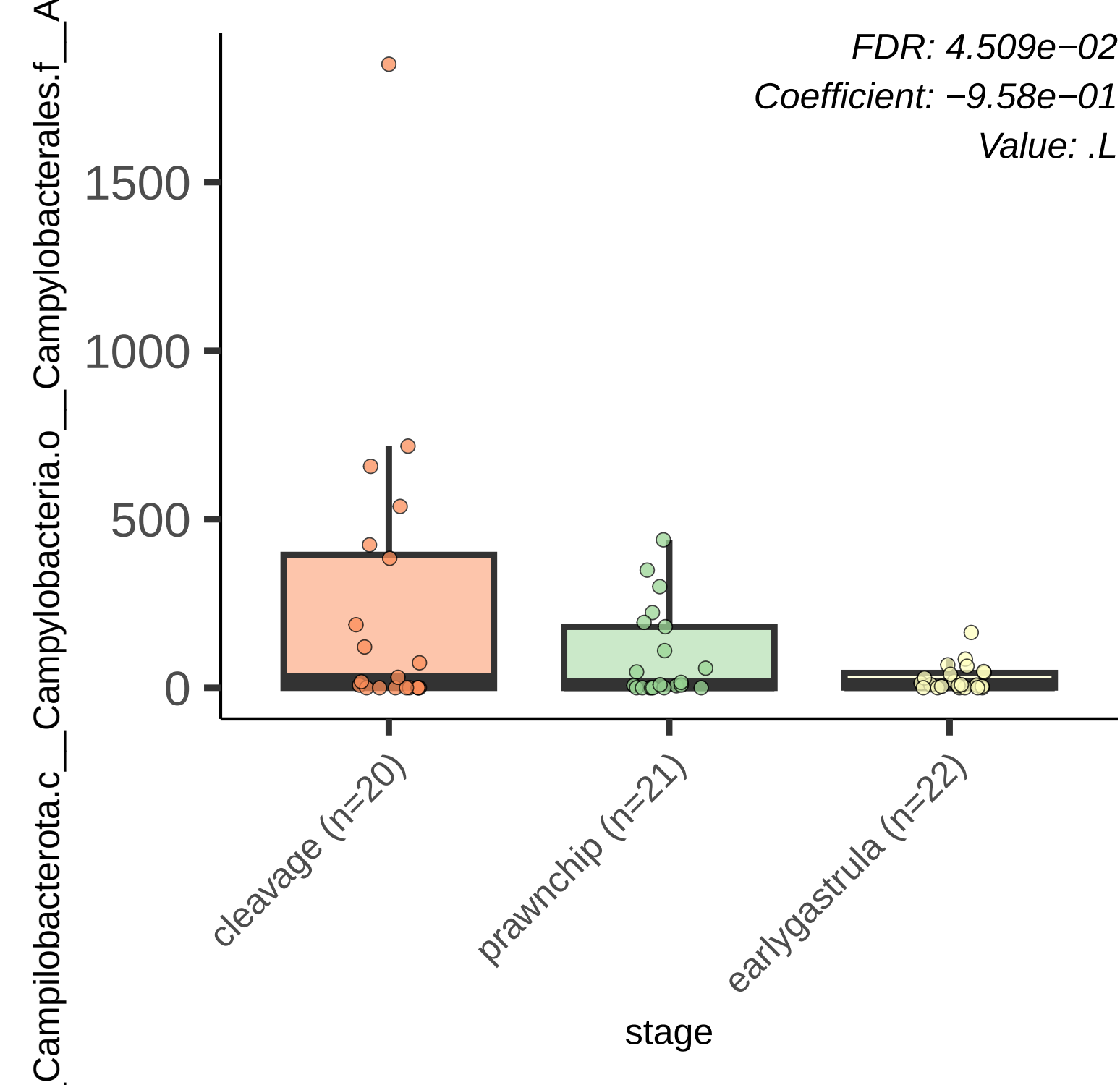

FDR: 3.662e−02
Coefficient: −9.33e−01
Value: .L
60
40
20
0
cleavage (n=20)
prawnchip (n=21)
earlygastrula (n=22)
stage
cteria.c__Gammaproteobacteria.o__Oceanospirillales.f__Saccharos

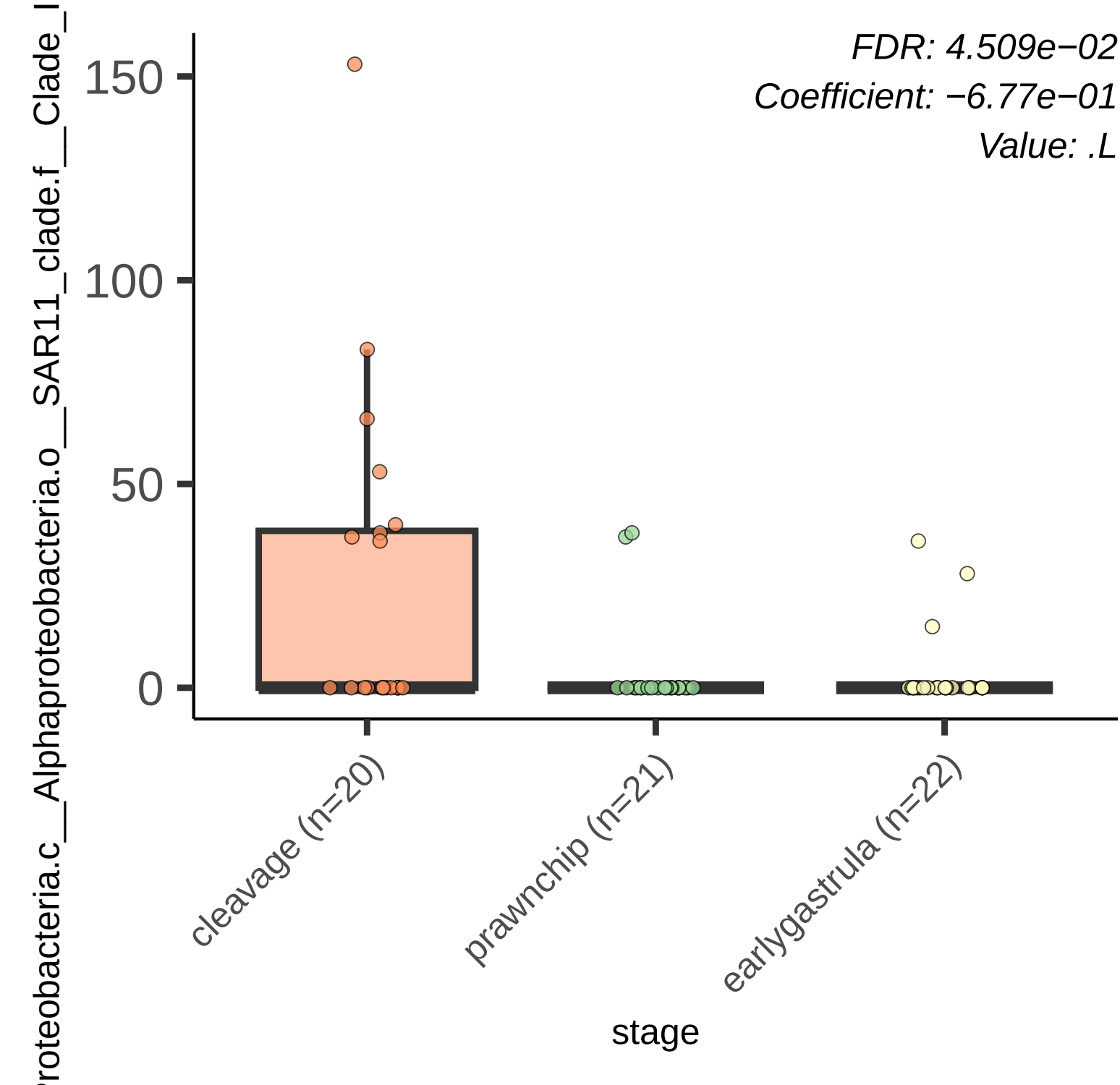

FDR: 3.837e−02
Coefficient: −5.76e−01
Value: .L
25
20
15
10
5
0
cleavage (n=20)
prawnchip (n=21)
earlygastrula (n=22)
stage
ria.c__Gammaproteobacteria.o__Pseudomonadales.f__Moraxellace

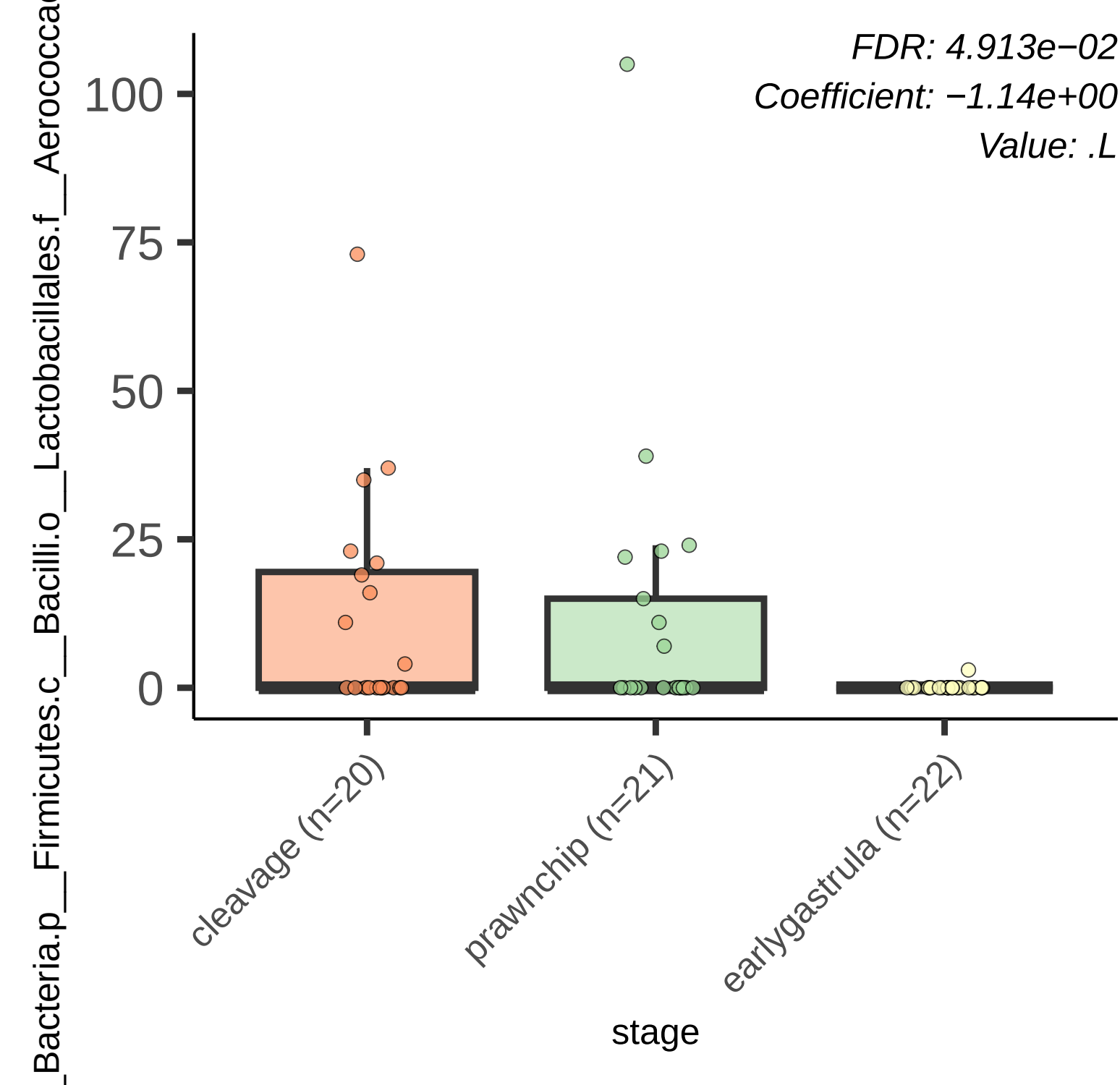

FDR: 4.509e−02
Coefficient: 1.15e+00
Value: .L
120
90
60
30
0
cleavage (n=20)
prawnchip (n=21)
earlygastrula (n=22)
stage
bacteria.c__Alphaproteobacteria.o__Rhodospirillales.f__Terasakiella

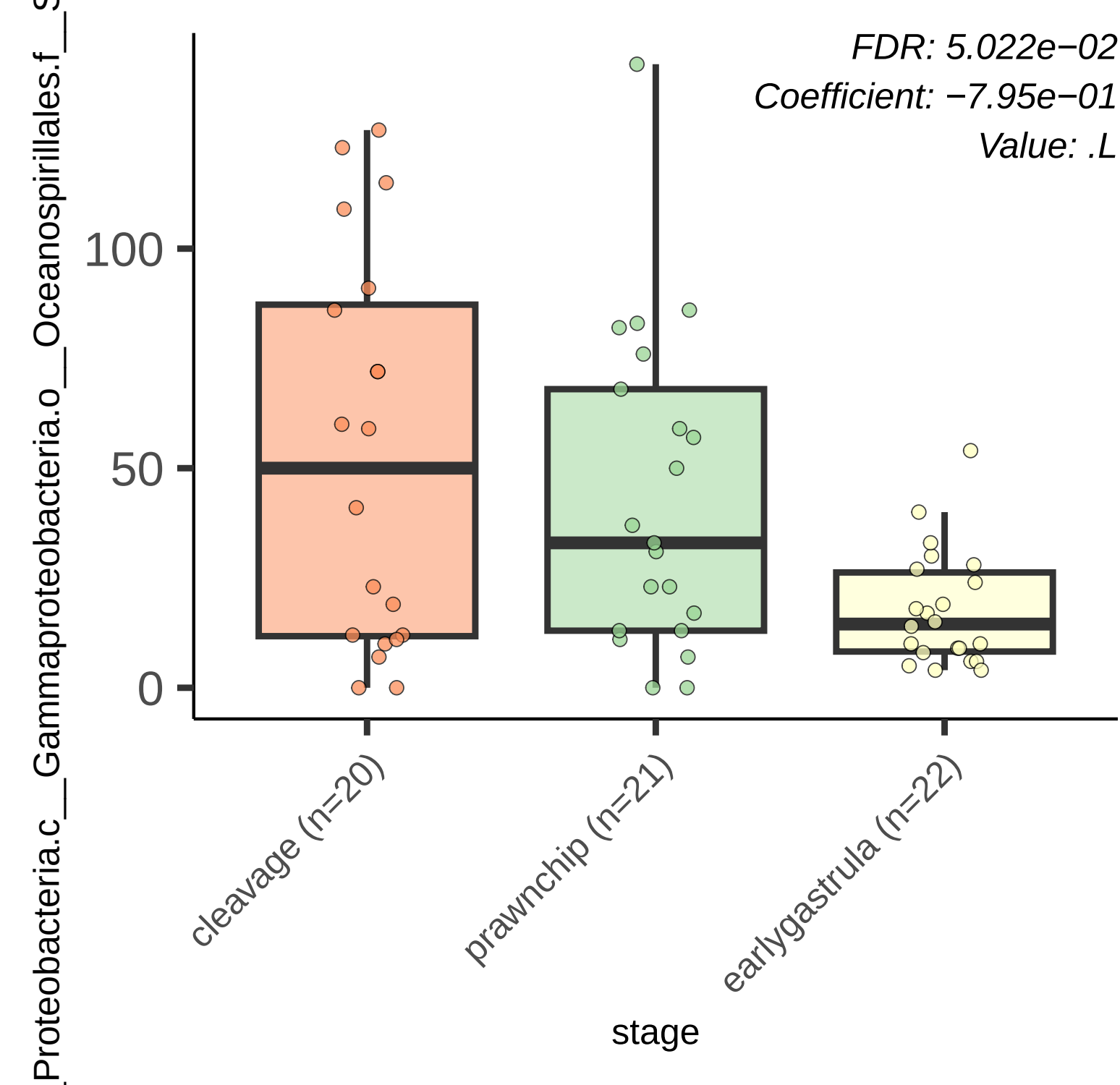

FDR: 4.509e−02
Coefficient: −8.01e−01
Value: .Q
1500
1000
500
0
cleavage (n=20)
prawnchip (n=21)
earlygastrula (n=22)
stage
teobacteria.c__Gammaproteobacteria.o__Oceanospirillales.f__Oleip


FDR: 4.509e−02
Coefficient: 1.02e+00
Value: .L
50
40
30
20
10
0
cleavage (n=20)
prawnchip (n=21)
earlygastrula (n=22)
stage
teria.p__Bacteroidota.c__Bacteroidia.o__Flavobacteriales.f__Flavob


FDR: 4.509e−02
Coefficient: −9.58e−01
Value: .L
1500
1000
500
0
cleavage (n=20)
prawnchip (n=21)
earlygastrula (n=22)
stage
Campilobacterota.c__Campylobacteria.o__Campylobacterales.f__A


FDR: 4.509e−02
Coefficient: −6.77e−01
Value: .L
150
100
50
0
cleavage (n=20)
prawnchip (n=21)
earlygastrula (n=22)
stage
roteobacteria.c__Alphaproteobacteria.o__SAR11_clade.f__Clade_I


FDR: 4.913e−02
Coefficient: −1.14e+00
Value: .L
100
75
50
25
0
cleavage (n=20)
prawnchip (n=21)
earlygastrula (n=22)
stage
_Bacteria.p__Firmicutes.c__Bacilli.o__Lactobacillales.f__Aerococca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Value: .L
100
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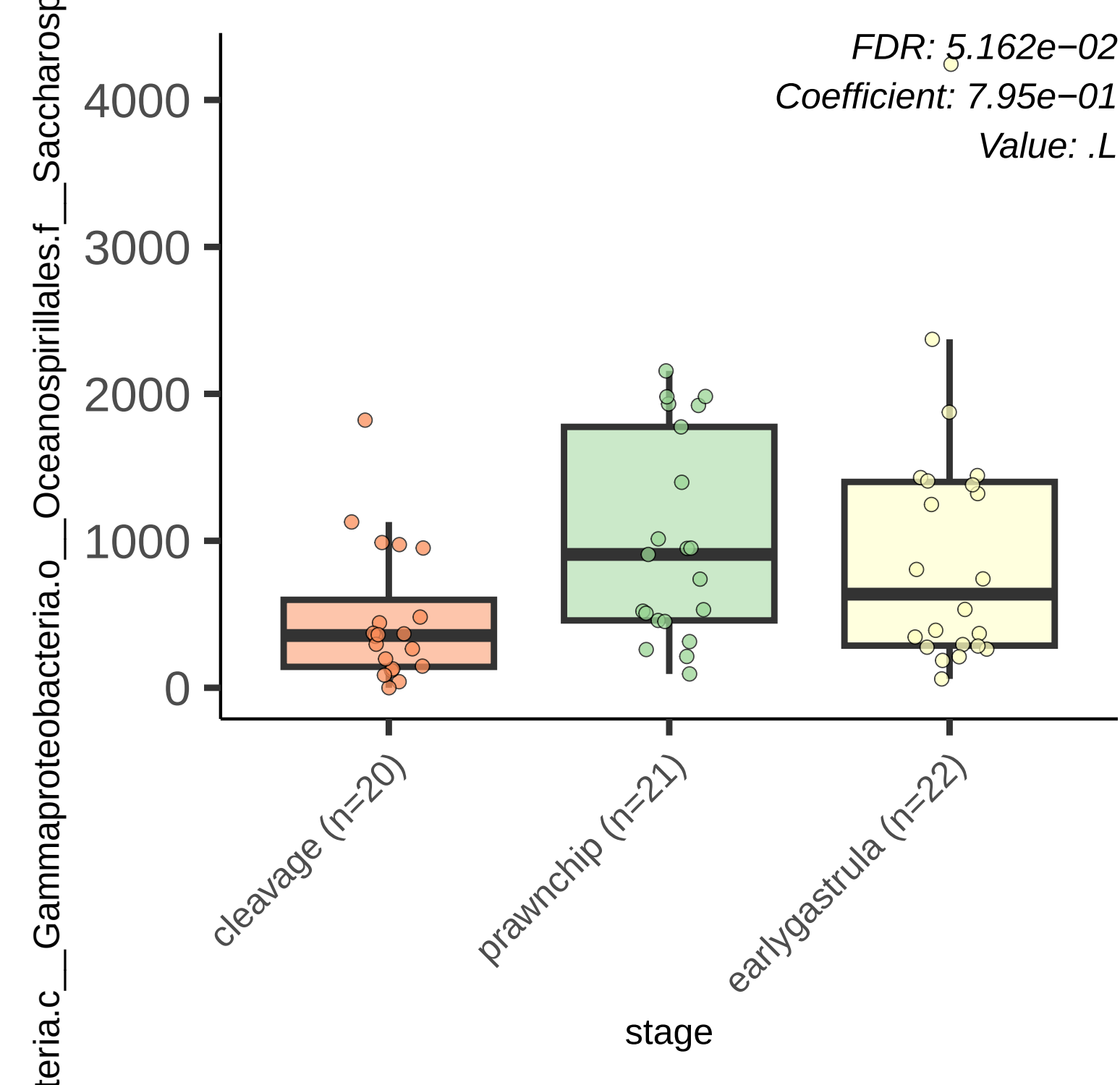
0
cleavage (n=20)
prawnchip (n=21)
earlygastrula (n=22)
stage
Proteobacteria.c__Gammaproteobacteria.o__Oceanospirillales.f__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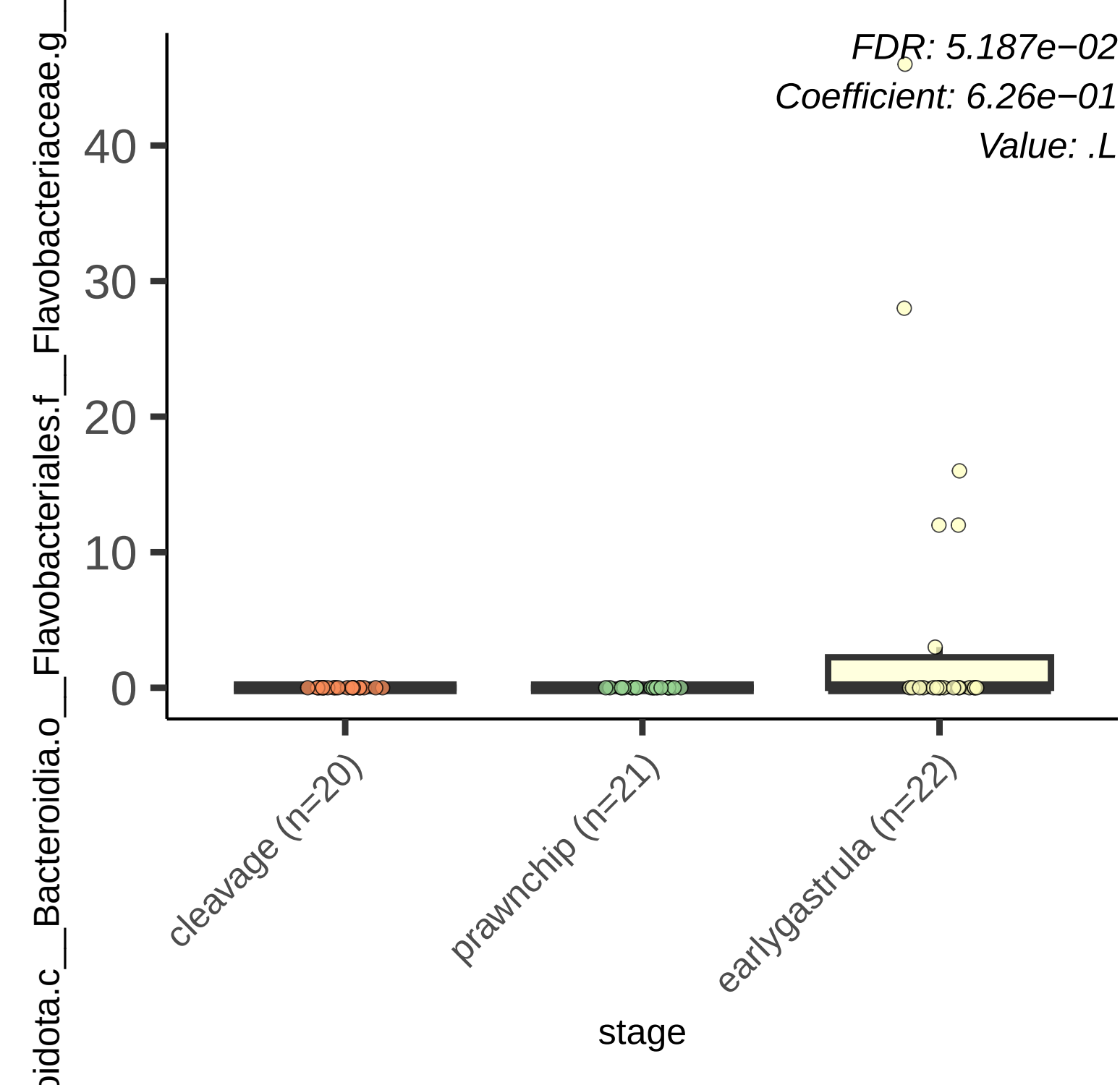
2000
1000
0
cleavage (n=20)
prawnchip (n=21)
earlygastrula (n=22)
stage
teria.c__Gammaproteobacteria.o__Oceanospirillales.f__Saccharosp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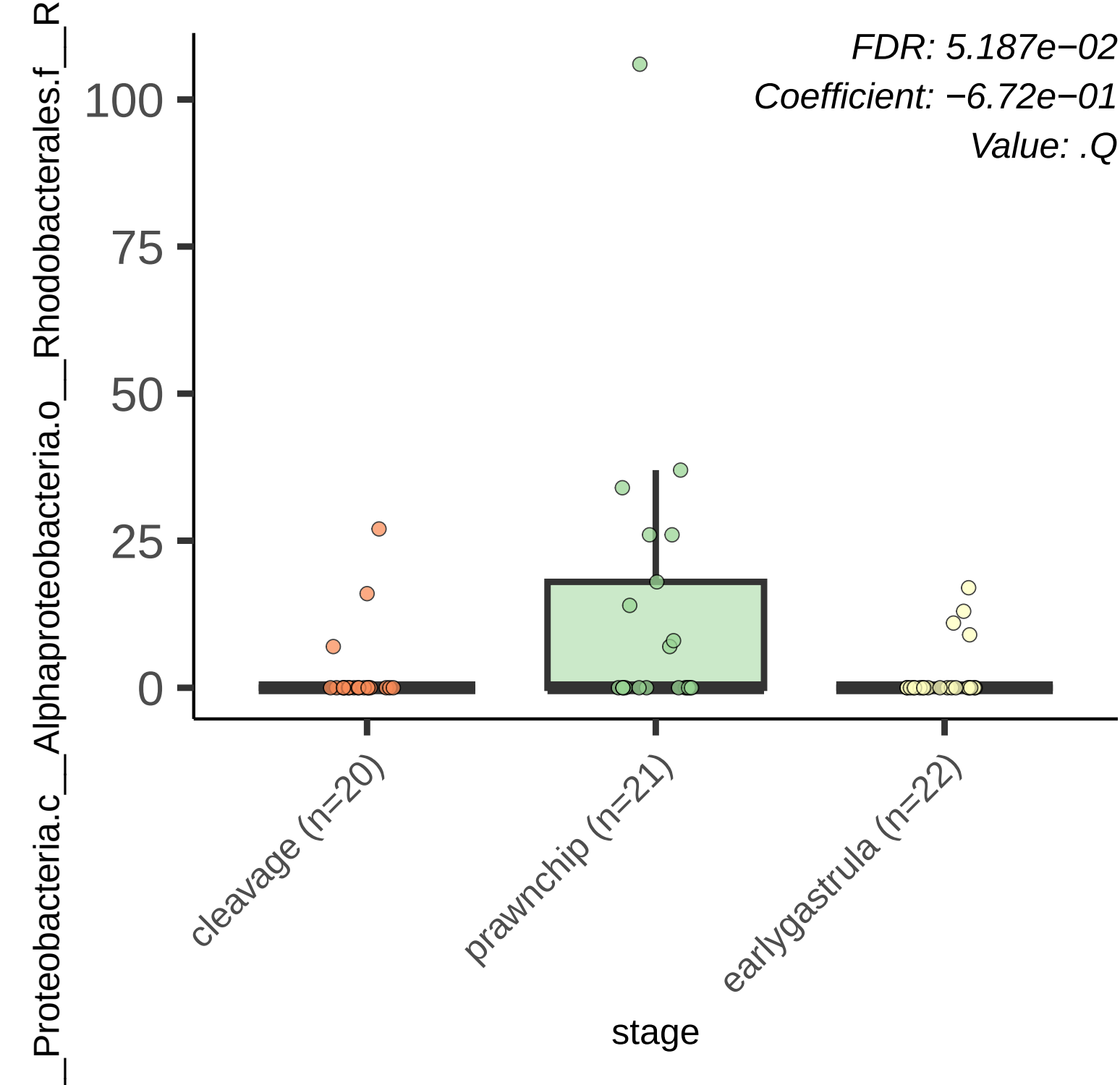
30
20
10
0
cleavage (n=20)
prawnchip (n=21)
earlygastrula (n=22)
stage
oidota.c__Bacteroidia.o__Flavobacteriales.f__Flavobacteriaceae.g__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100
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50
25
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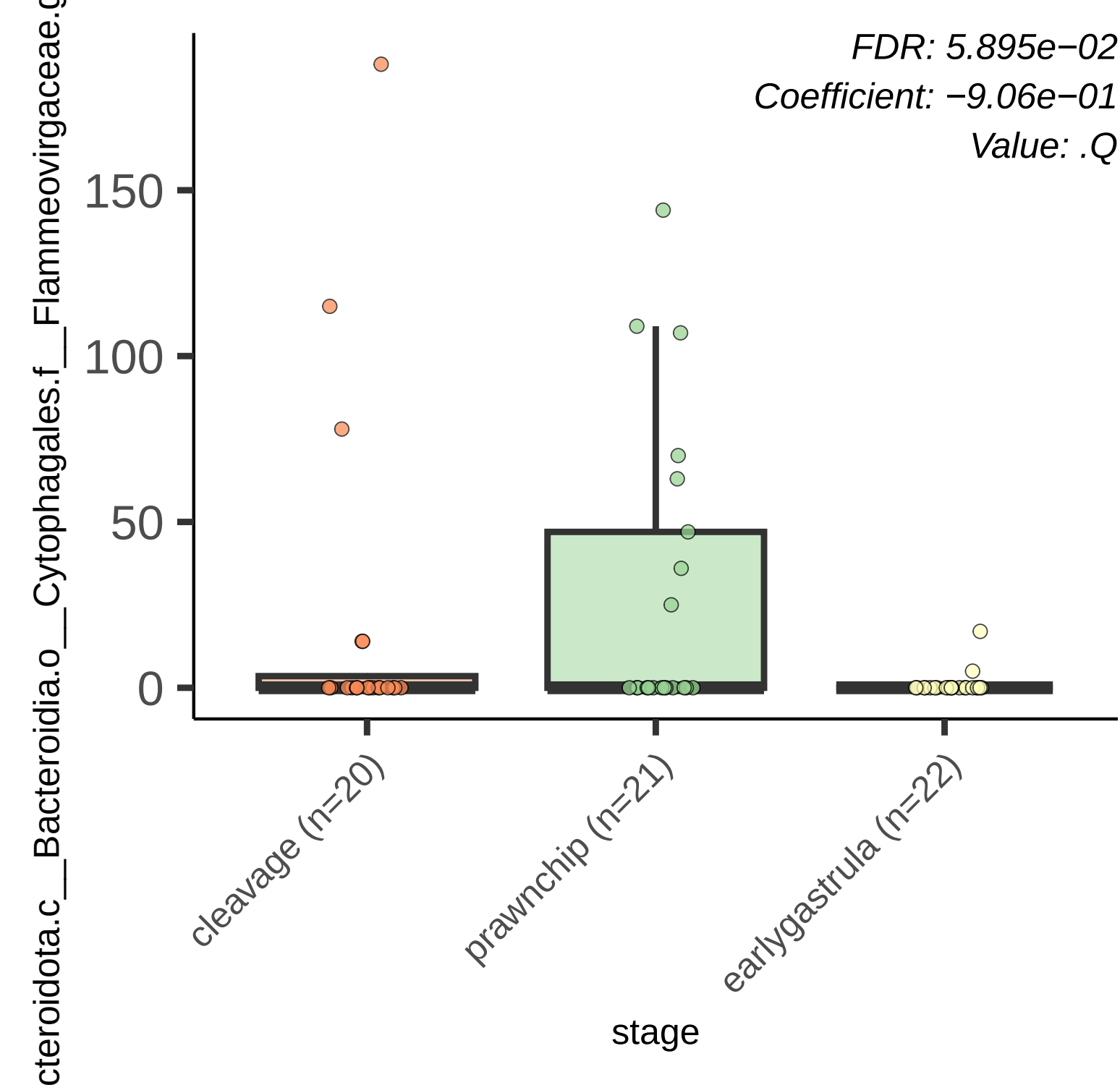
earlygastrula (n=22)
stage
d__Proteobacteria.c__Alphaproteobacteria.o__Rhodobacterales.f__R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50
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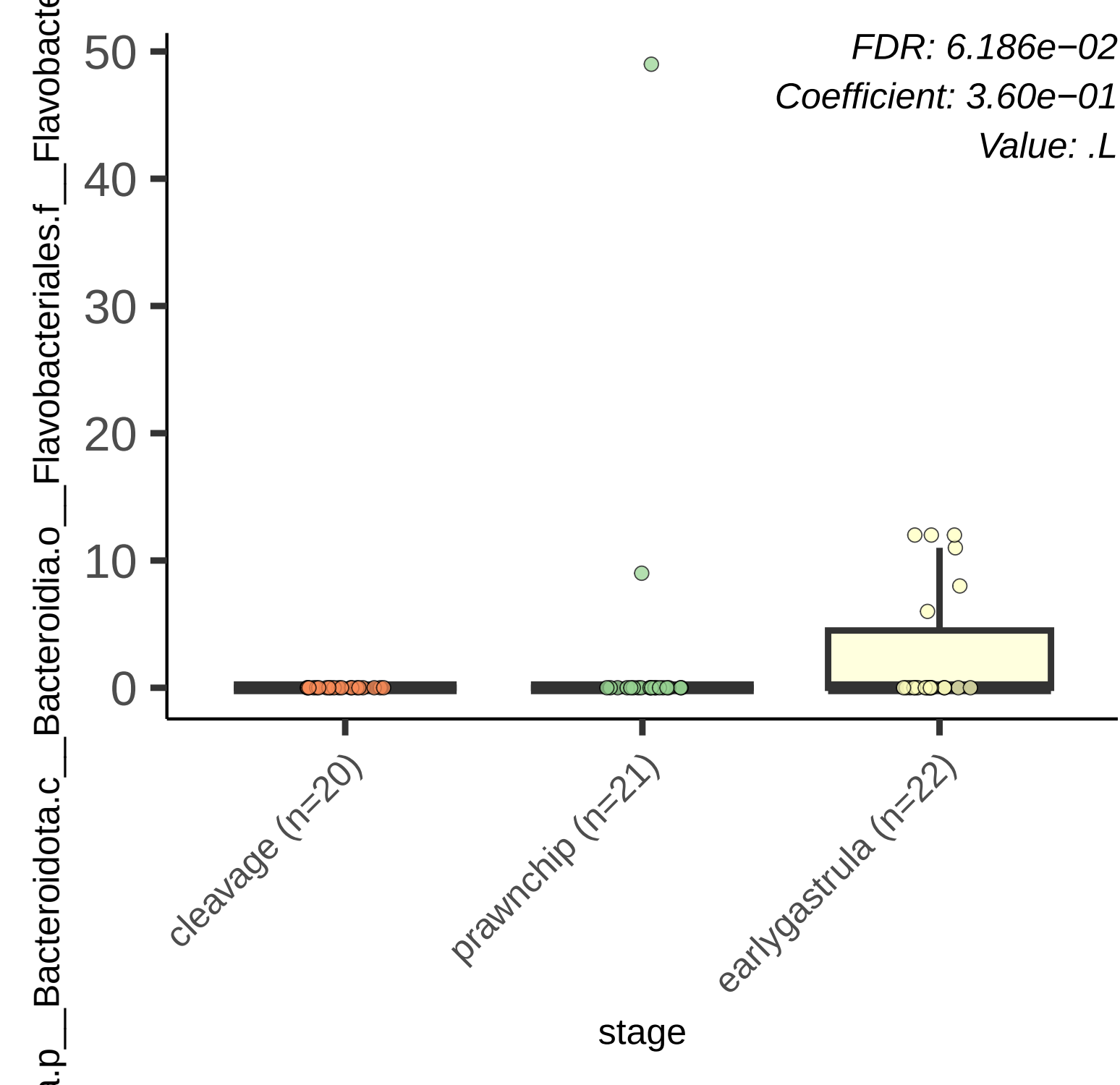
prawnchip (n=21)
earlygastrula (n=22)
stage
cteroidota.c__Bacteroidia.o__Cytophagales.f__Flammeovirgaceae.g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40
30
20
10
0
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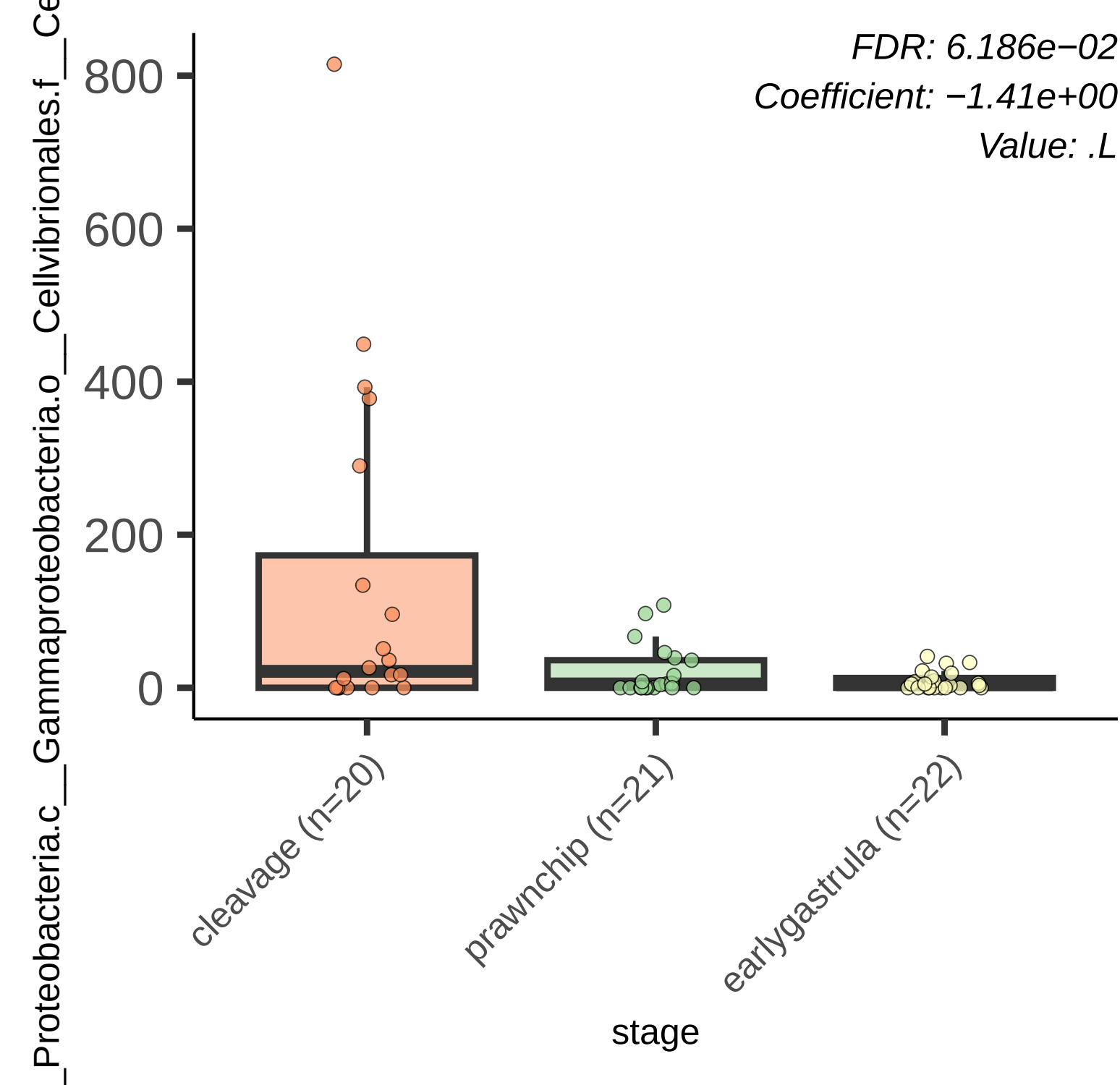
earlygastrula (n=22)
stage
a.p__Bacteroidota.c__Bacteroidia.o__Flavobacteriales.f__Flavobacte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600
400
200
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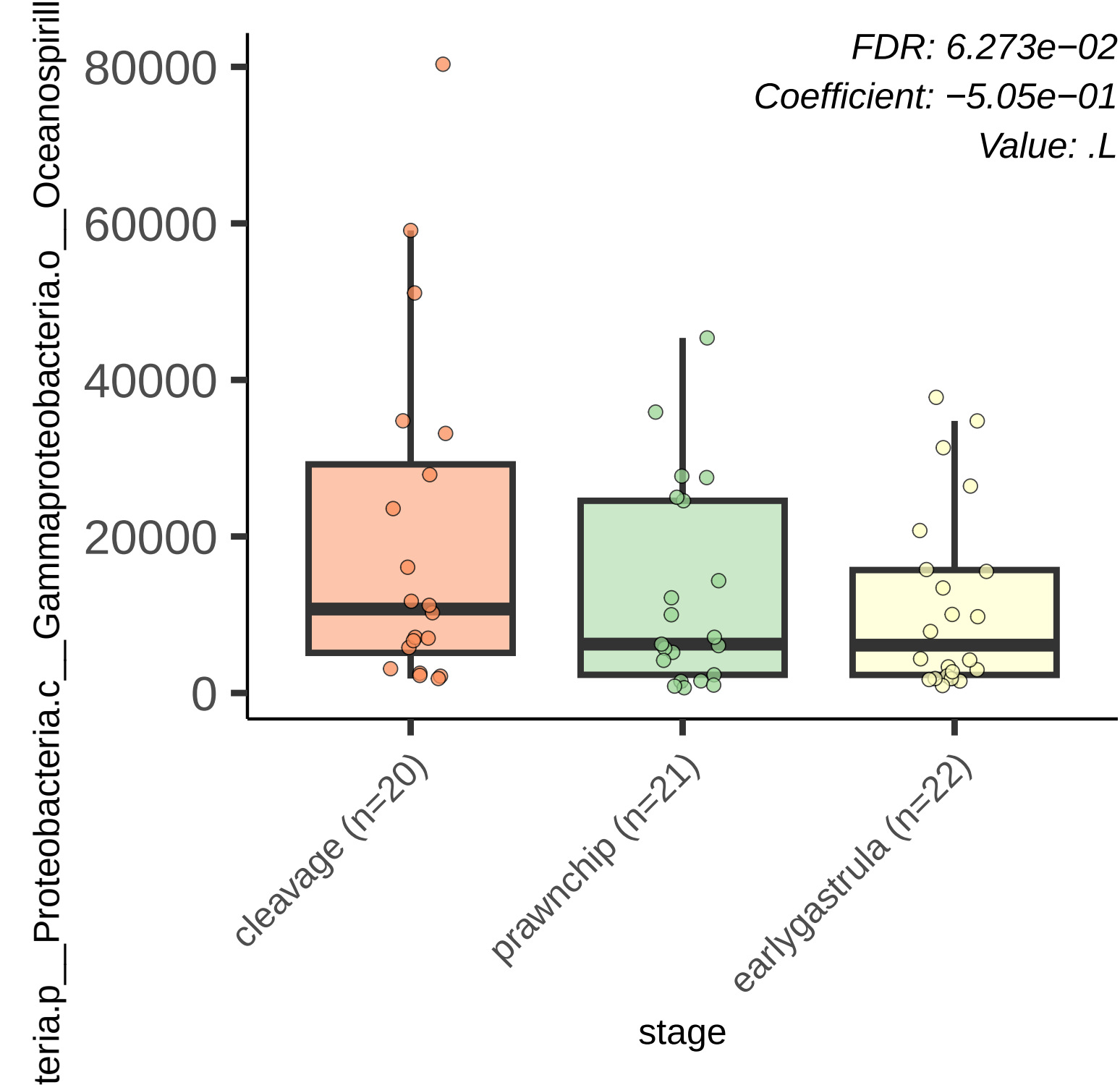
prawnchip (n=21)
earlygastrula (n=22)
stage
_Proteobacteria.c__Gammaproteobacteria.o__Cellvibrionales.f__Ce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60000
40000
20000
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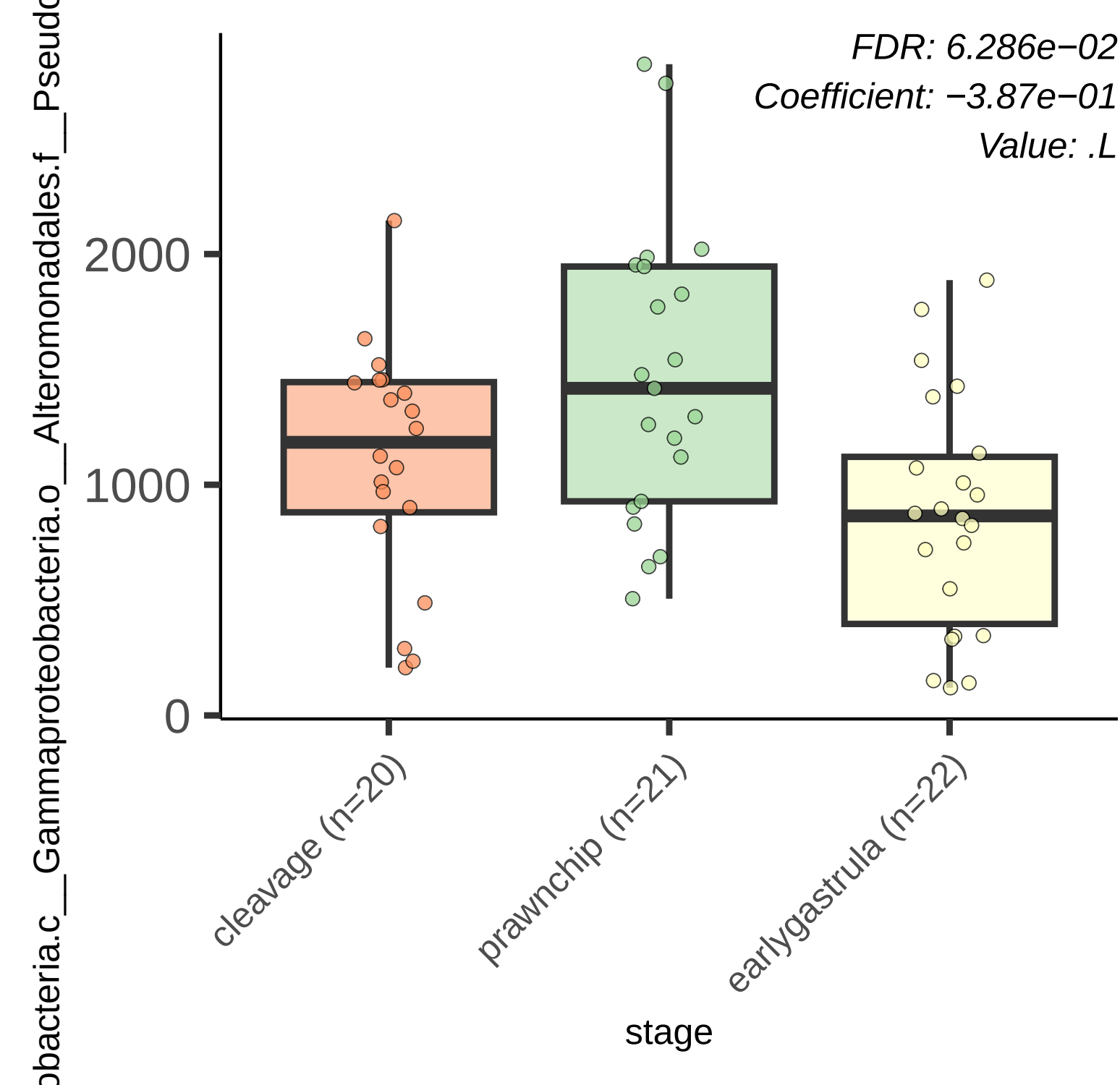
prawnchip (n=21)
earlygastrula (n=22)
stage
teria.p__Proteobacteria.c__Gammaproteobacteria.o__Oceanospirill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1000
0
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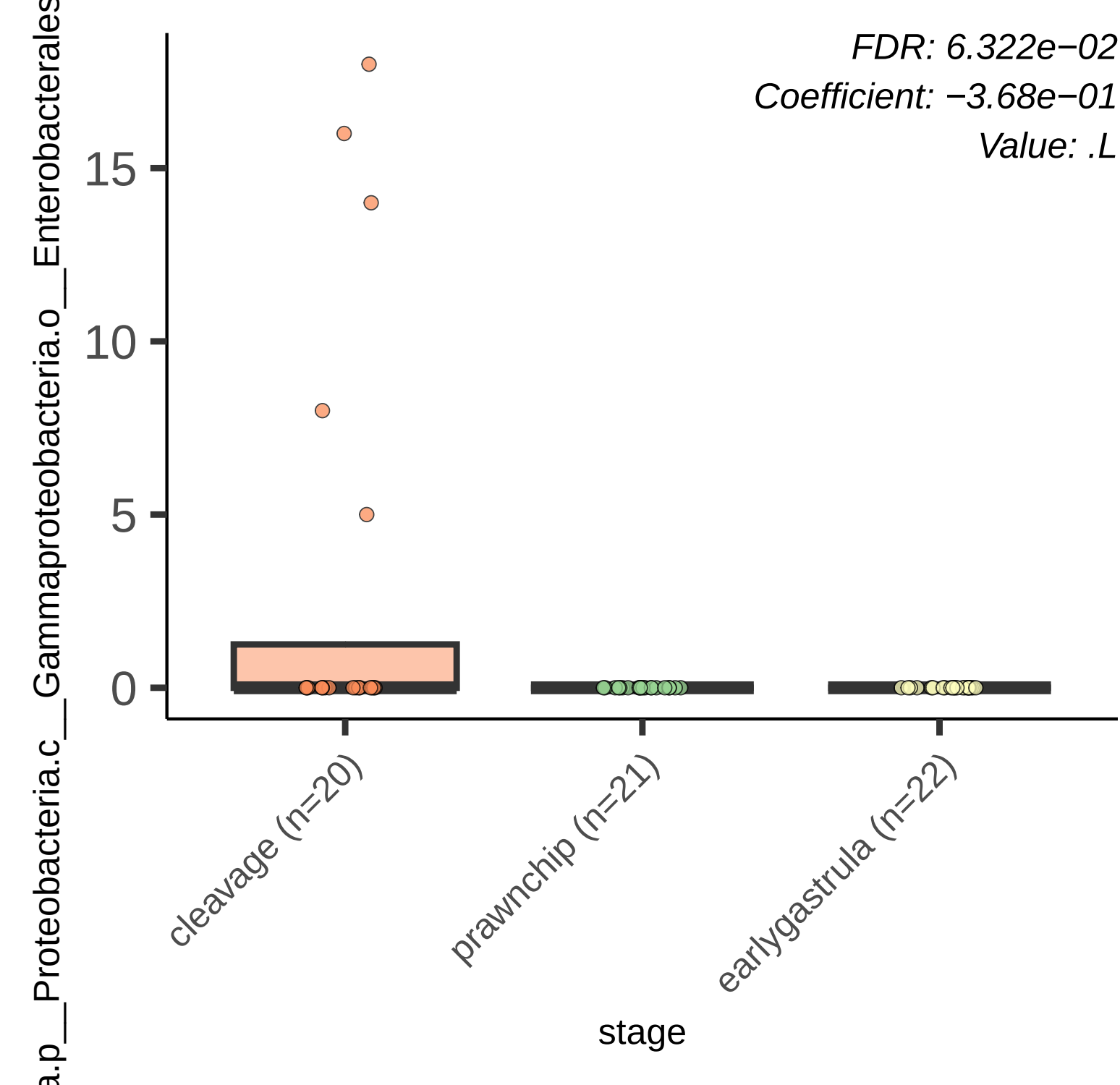
prawnchip (n=21)
earlygastrula (n=22)
stage
obacteria.c__Gammaproteobacteria.o__Alteromonadales.f__Pseudo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10
5
0
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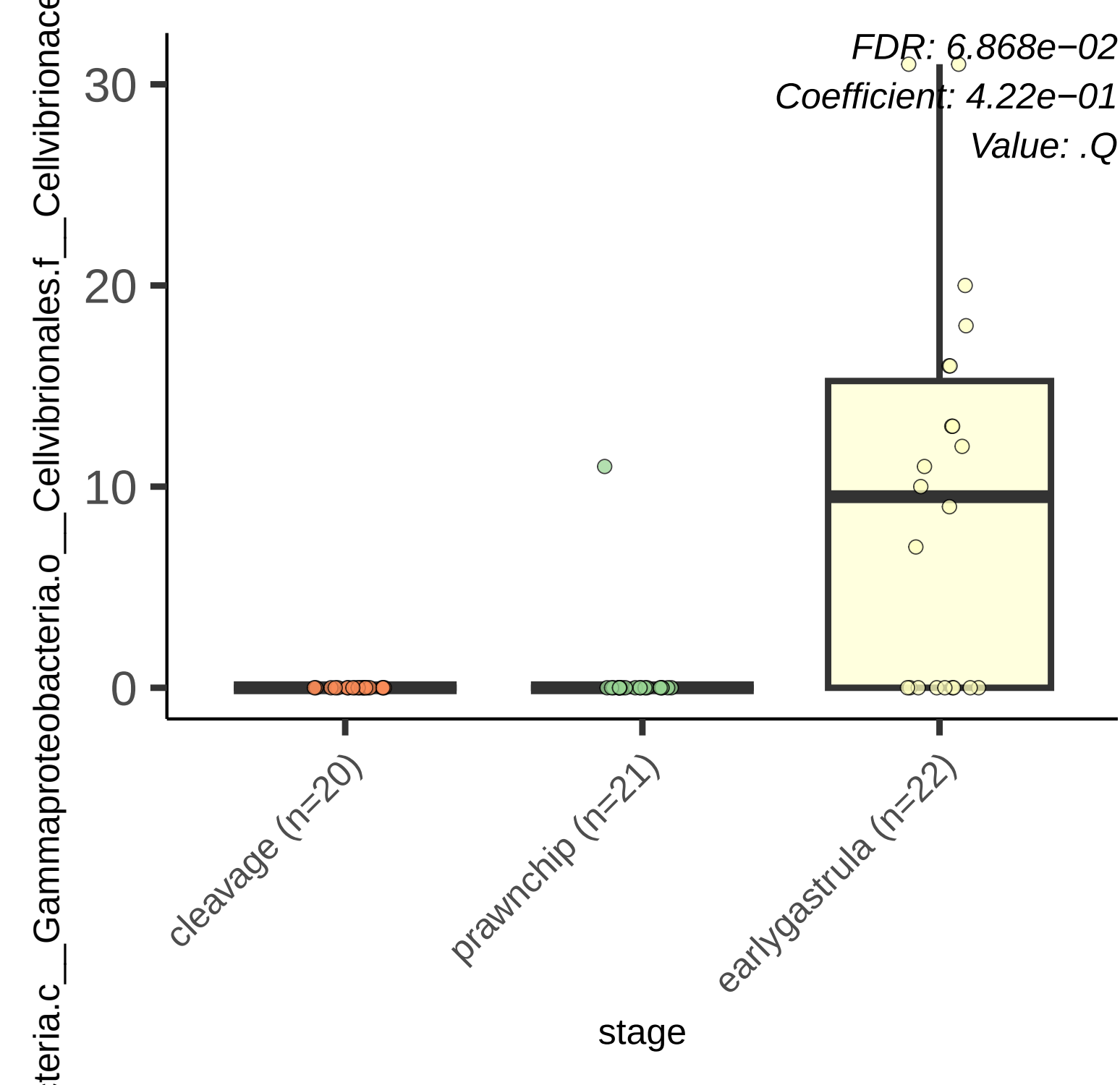
prawnchip (n=21)
earlygastrula (n=22)
stage
a.p__Proteobacteria.c__Gammaproteobacteria.o__Enterobacterales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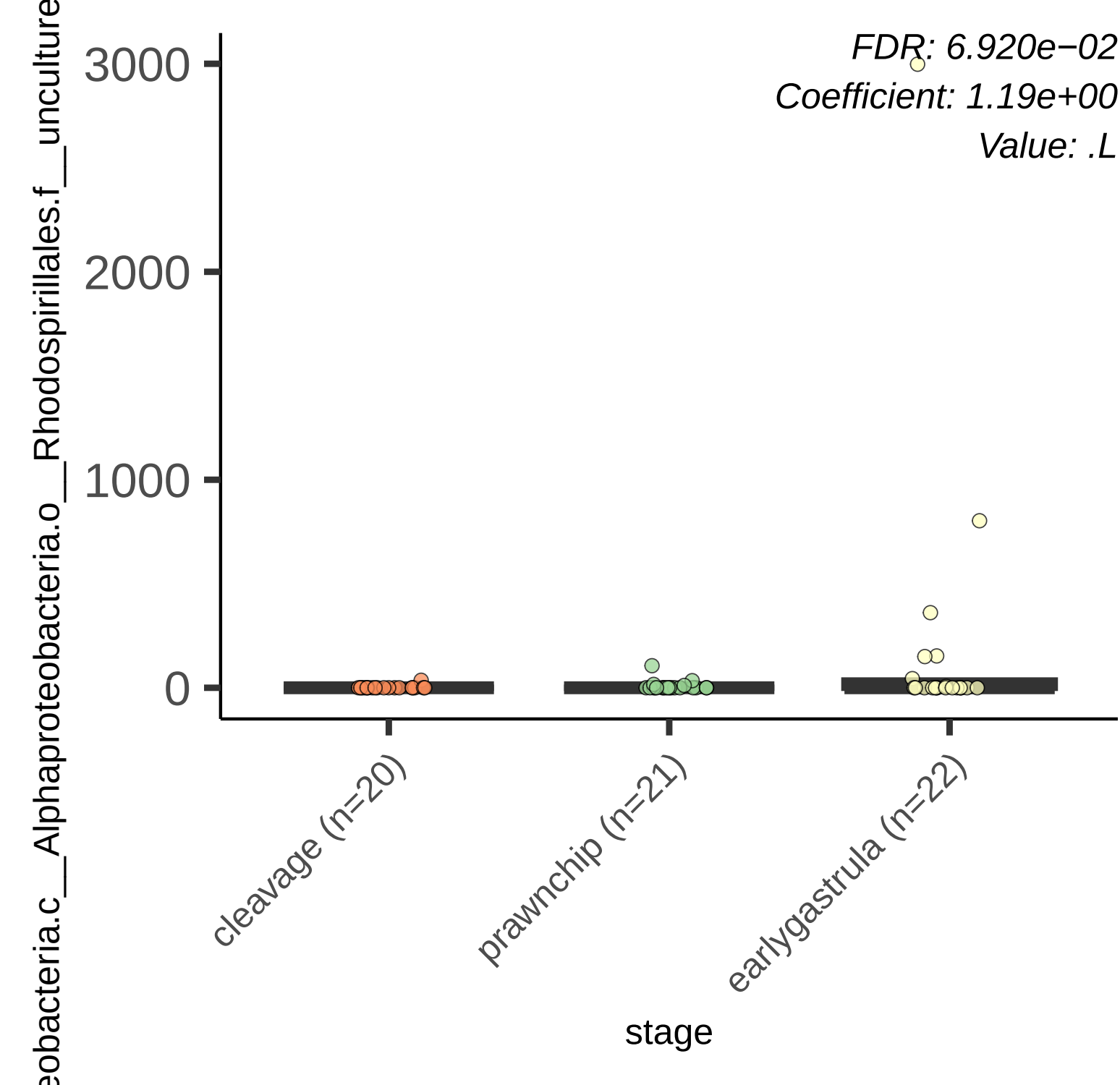
20
10
0
cleavage (n=20)
prawnchip (n=21)
earlygastrula (n=22)
stage
teria.c__Gammaproteobacteria.o__Cellvibrionales.f__Cellvibrionace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Coefficient: 1.19e+00
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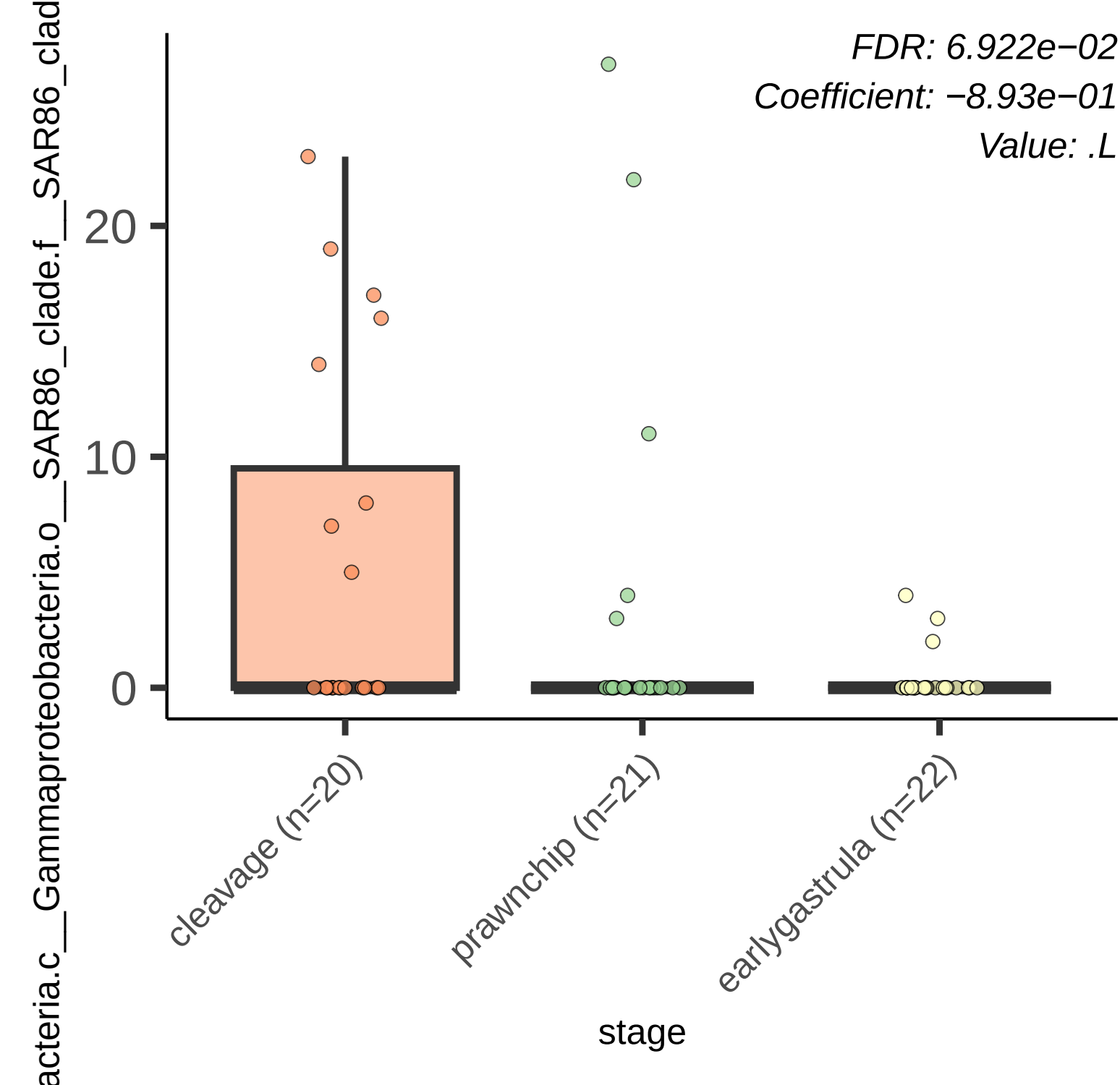
2000
1000
0
cleavage (n=20)
prawnchip (n=21)
earlygastrula (n=22)
stage
eobacteria.c__Alphaproteobacteria.o__Rhodospirillales.f__uncultured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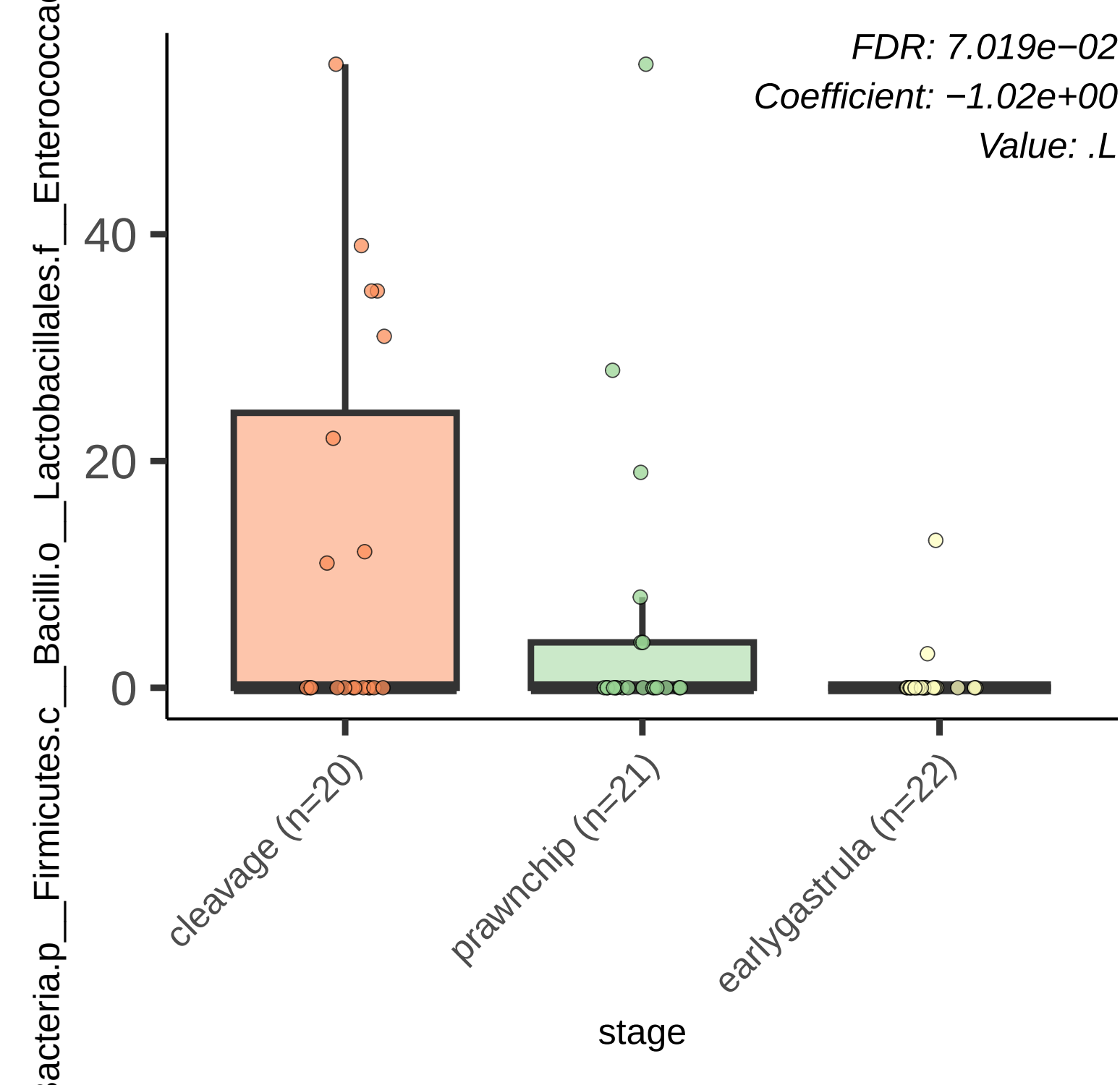
10
0
cleavage (n=20)
prawnchip (n=21)
earlygastrula (n=22)
stage
acteria.c__Gammaproteobacteria.o__SAR86_clade.f__SAR86_clad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20
0
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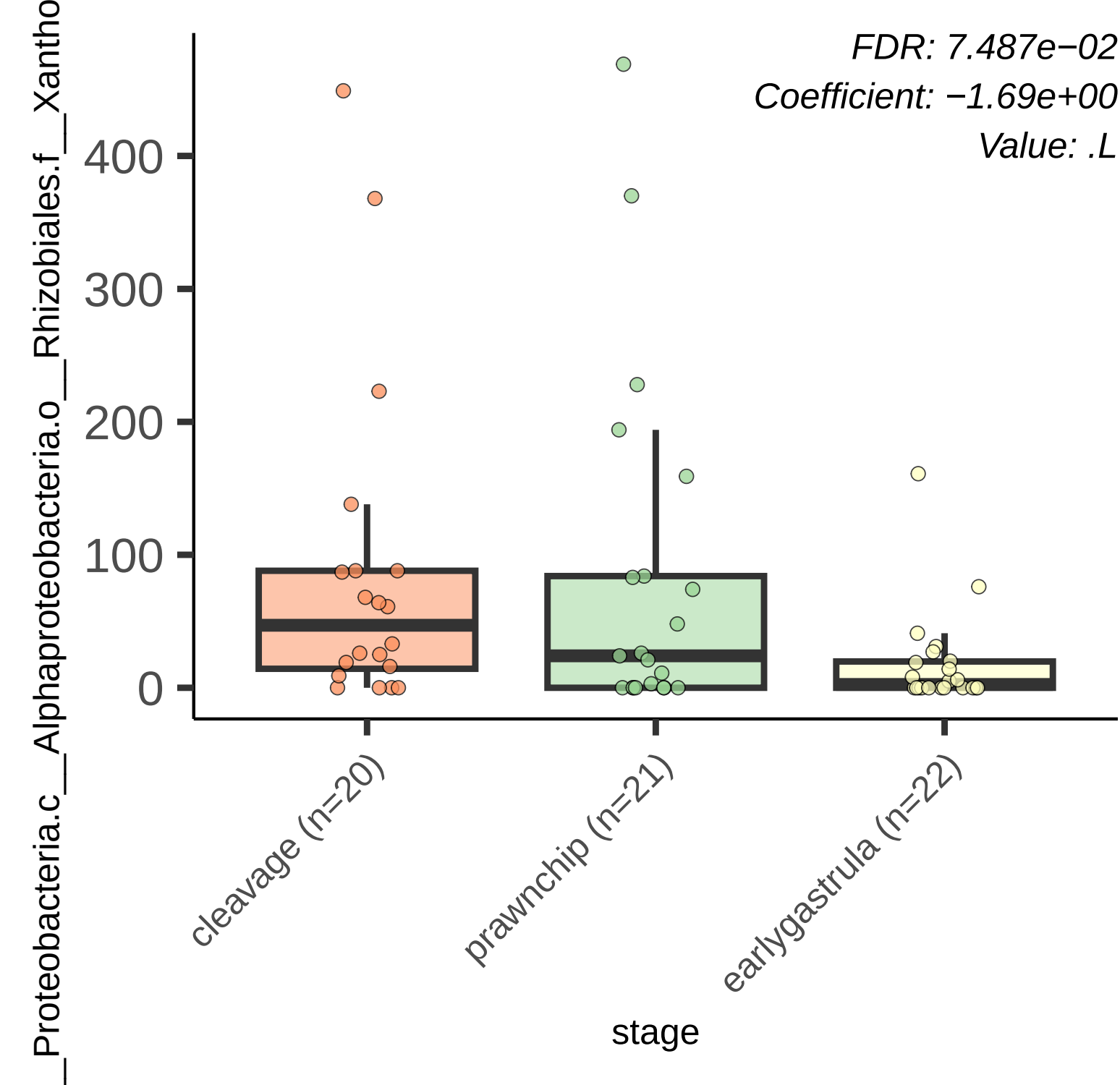
prawnchip (n=21)
earlygastrula (n=22)
stage
acteria.p__Firmicutes.c__Bacilli.o__Lactobacillales.f__Enterococca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300
200
100
0
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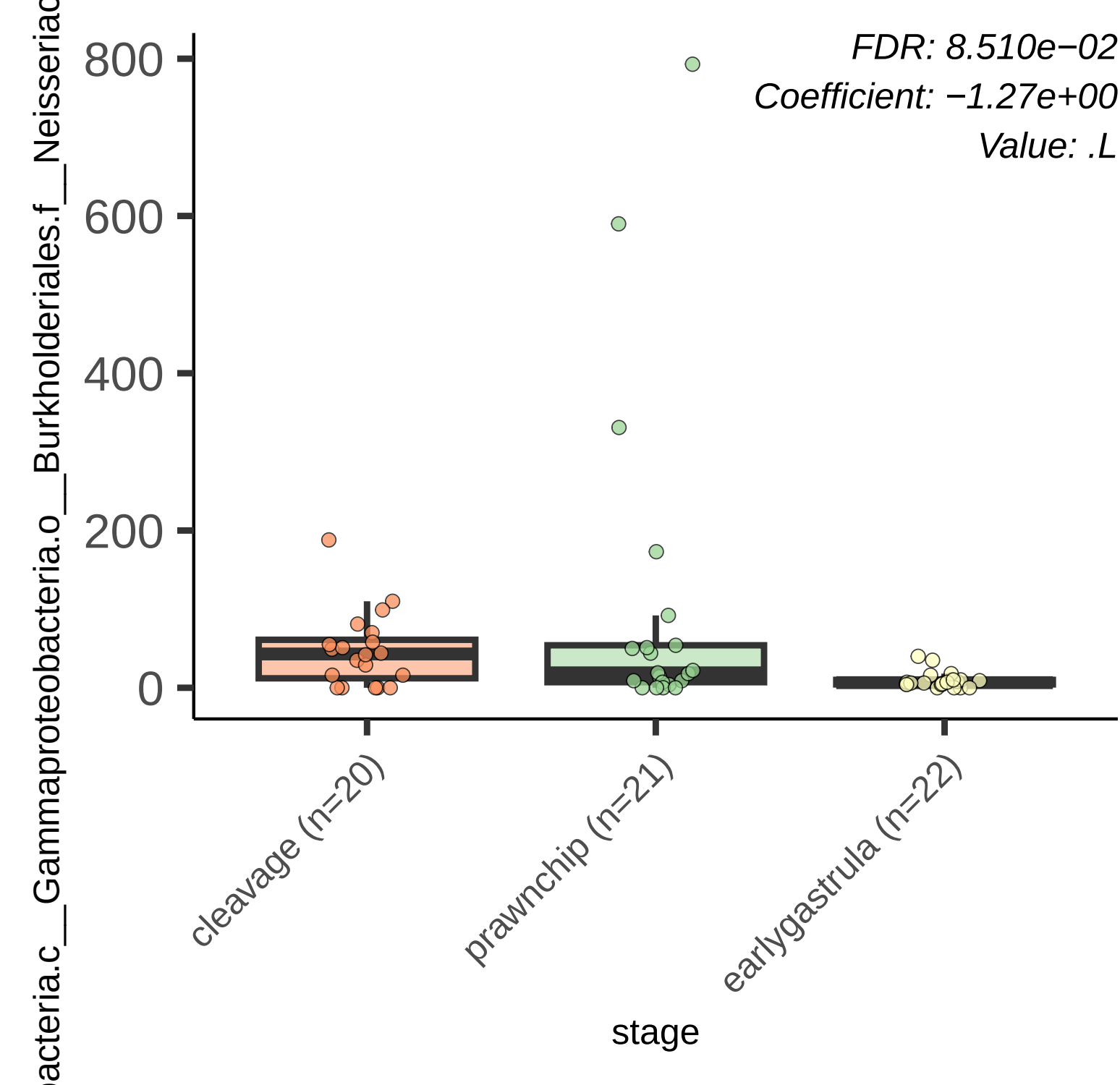
prawnchip (n=21)
earlygastrula (n=22)
stage
__Proteobacteria.c__Alphaproteobacteria.o__Rhizobiales.f__Xantho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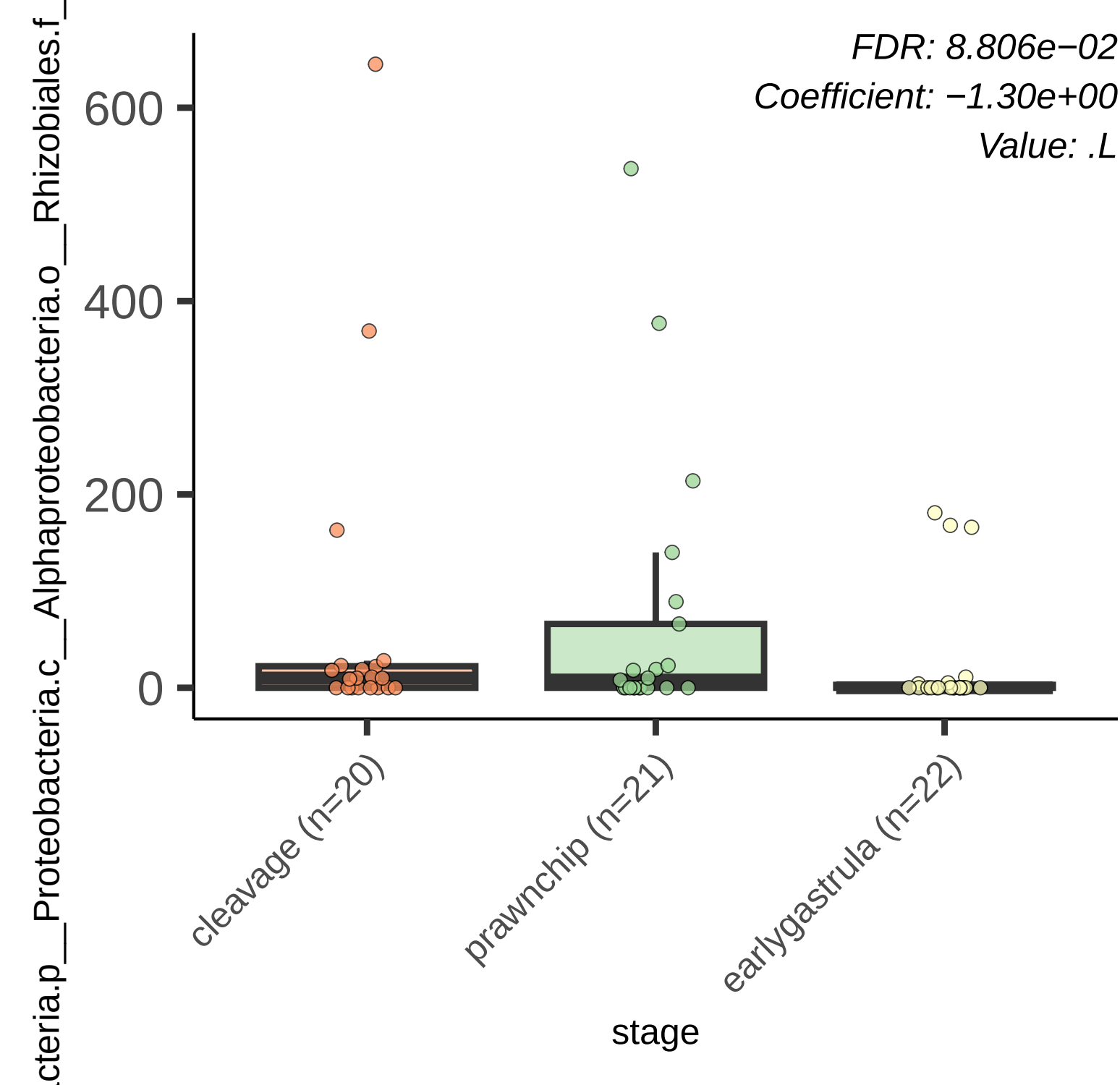
800
600
400
200
0
cleavage (n=20)
prawnchip (n=21)
earlygastrula (n=22)
stage
bacteria.c__Gammaproteobacteria.o__Burkholderiales.f__Neisseriac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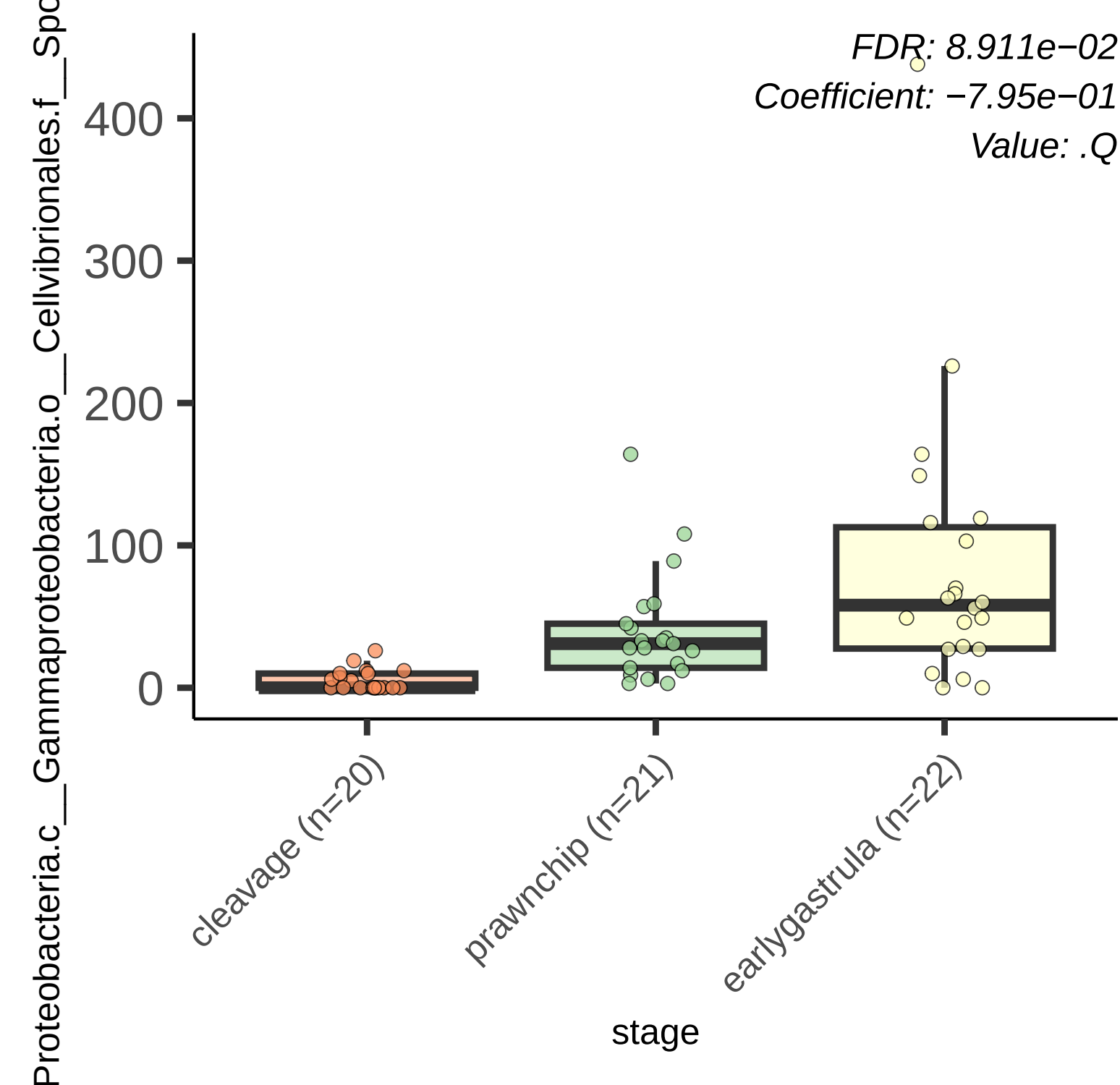
600
400
200
0
cleavage (n=20)
prawnchip (n=21)
earlygastrula (n=22)
stage
cteria.p__Proteobacteria.c__Alphaproteobacteria.o__Rhizobiales.f__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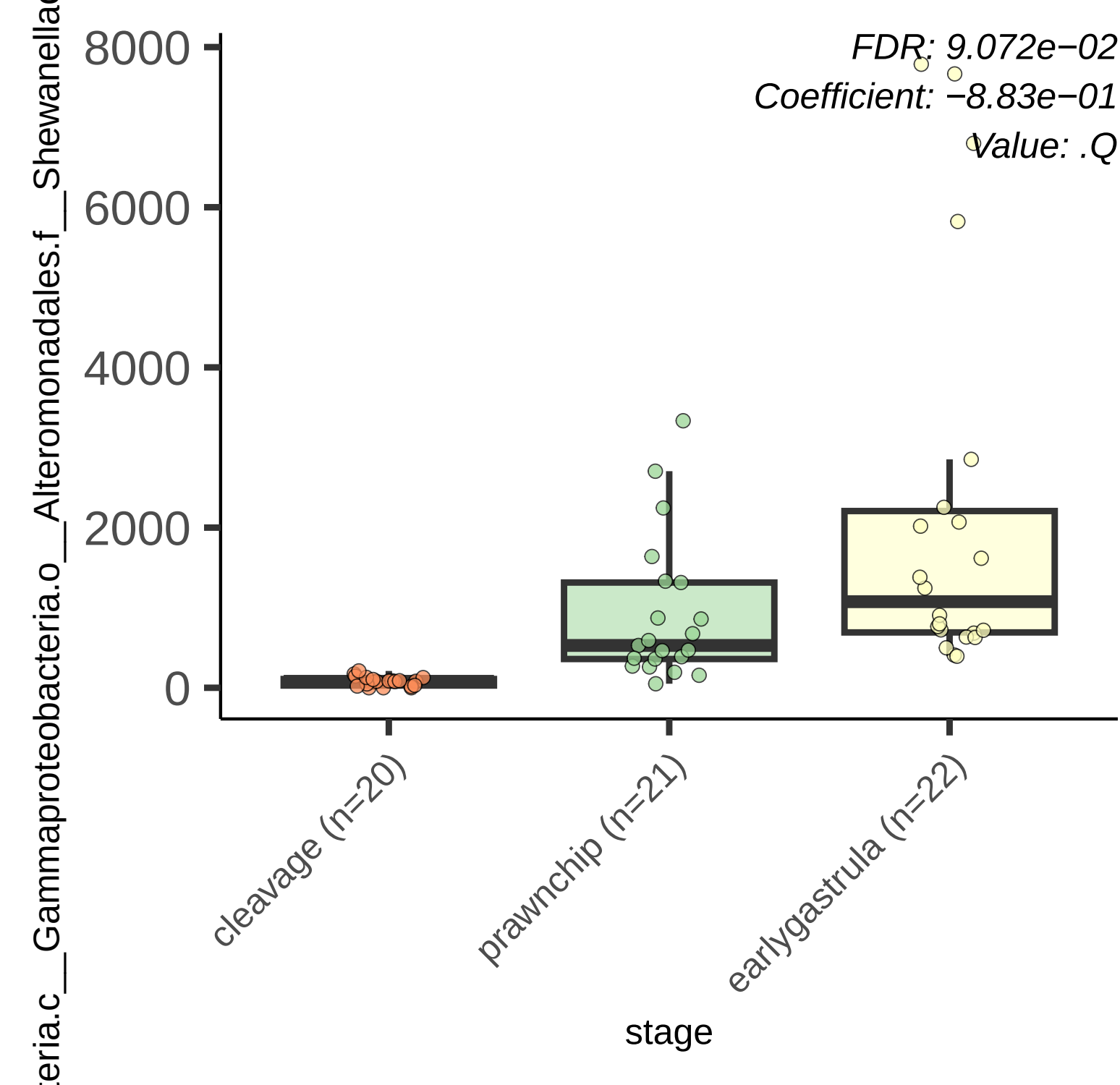
400
300
200
100
0
cleavage (n=20)
prawnchip (n=21)
earlygastrula (n=22)
stage
Proteobacteria.c__Gammaproteobacteria.o__Cellvibrionales.f__Spo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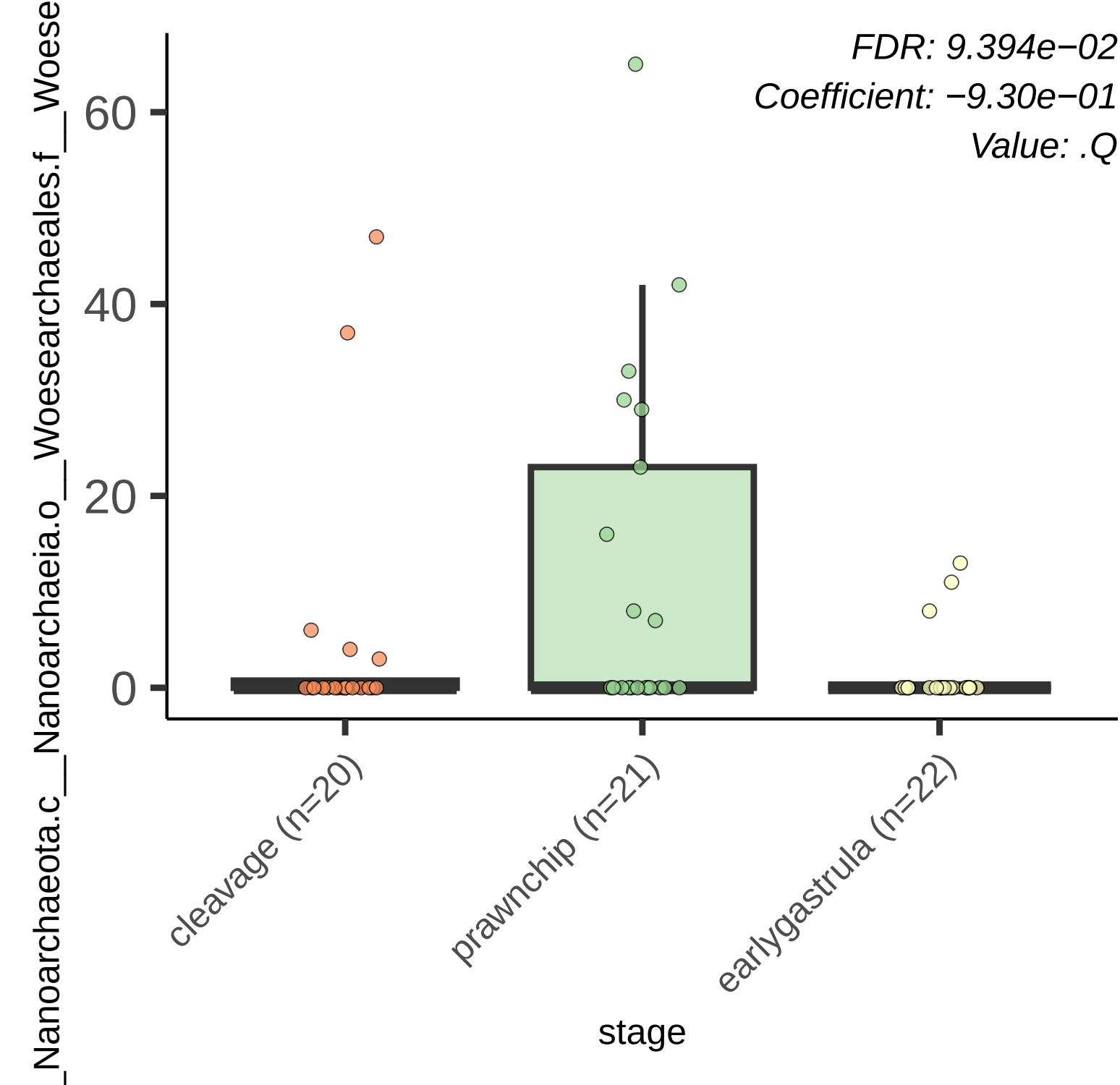
6000
4000
2000
0
cleavage (n=20)
prawnchip (n=21)
earlygastrula (n=22)
stage
teria.c__Gammaproteobacteria.o__Alteromonadales.f__Shewanellac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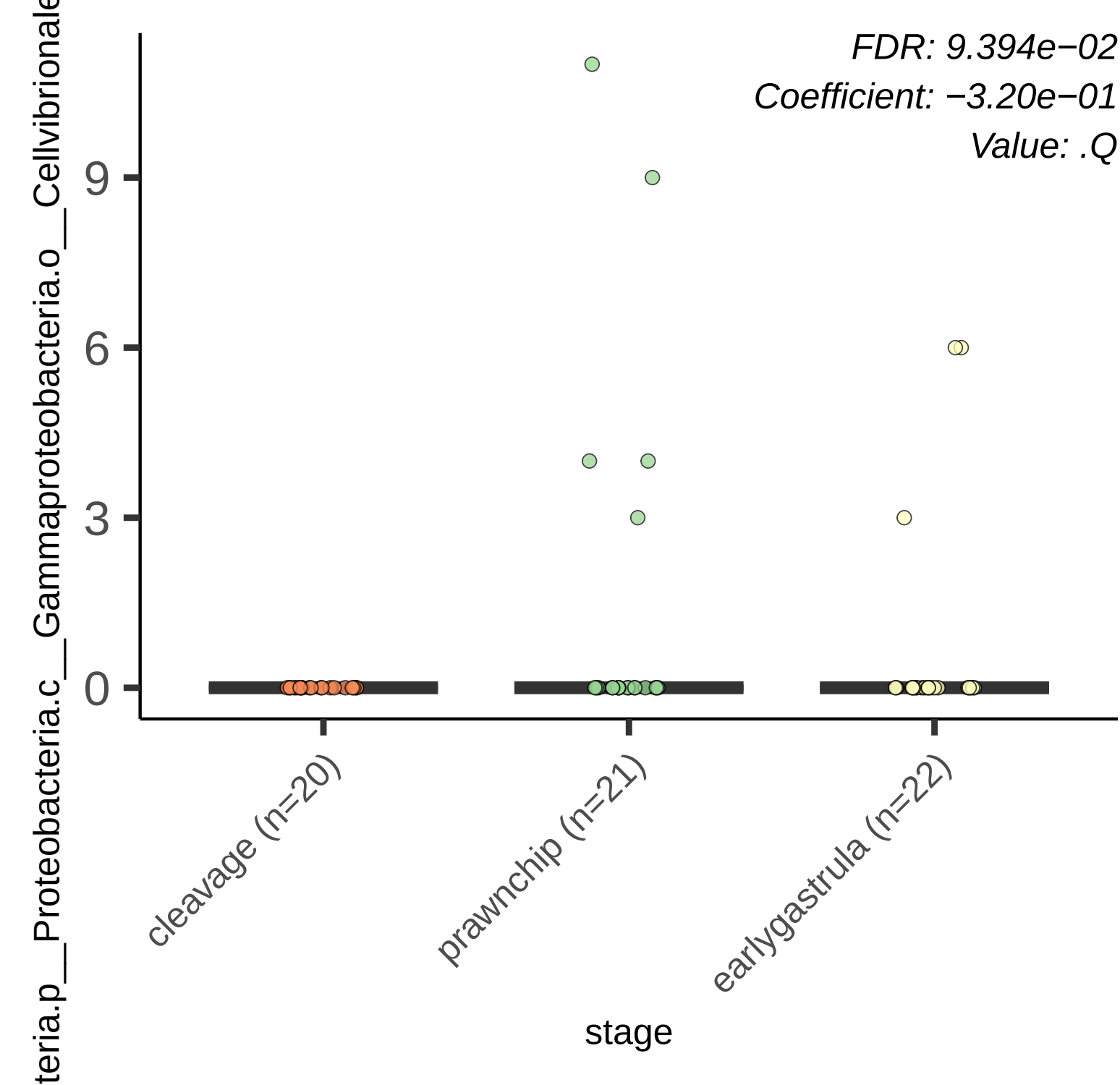
40
20
0
cleavage (n=20)
prawnchip (n=21)
earlygastrula (n=22)
stage
_Nanoarchaeota.c__Nanoarchaeia.o__Woesearchaeales.f__Woese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9
6
3
0
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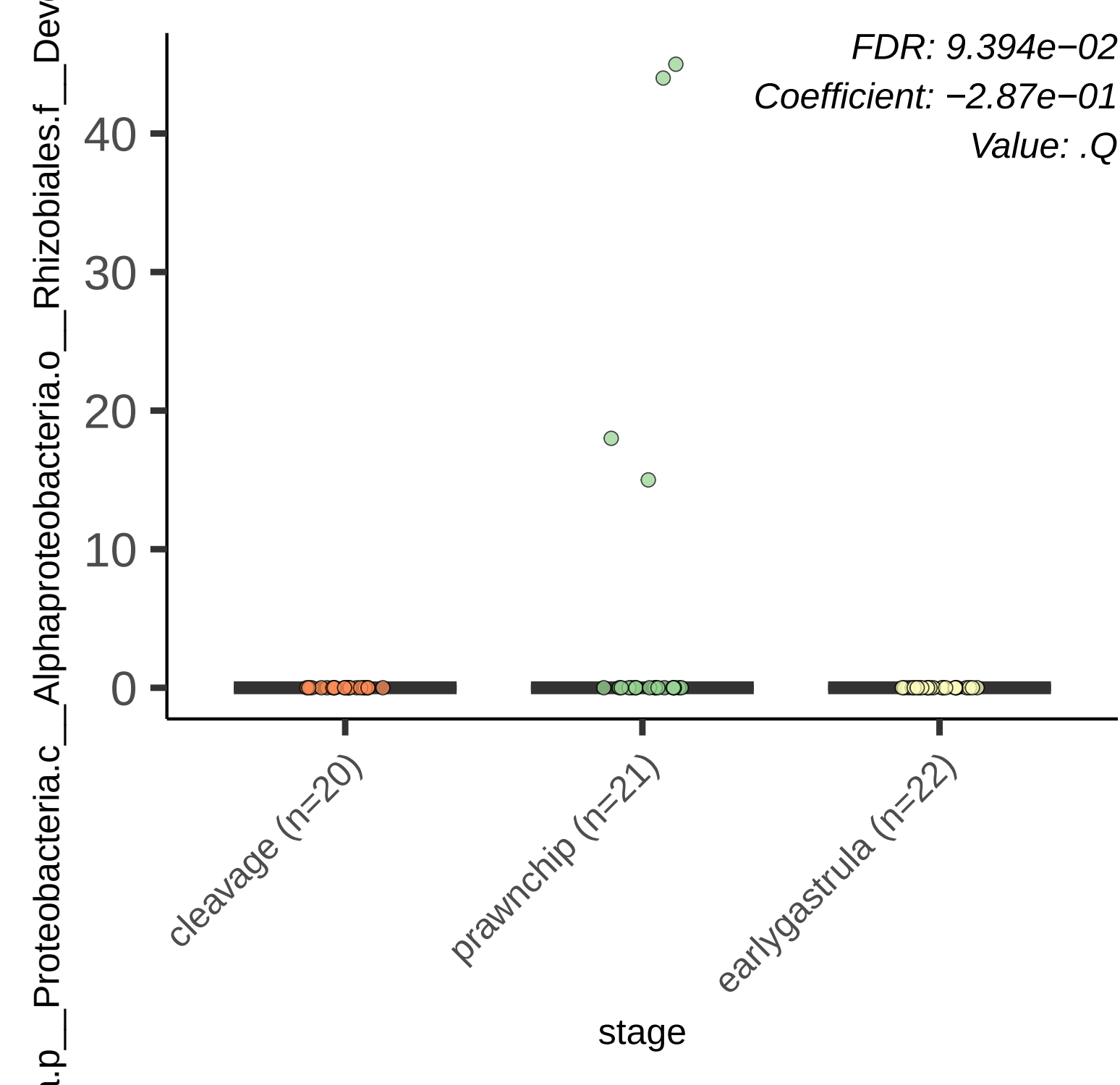
prawnchip (n=21)
earlygastrula (n=22)
stage
teria.p__Proteobacteria.c__Gammaproteobacteria.o__Cellvibrionale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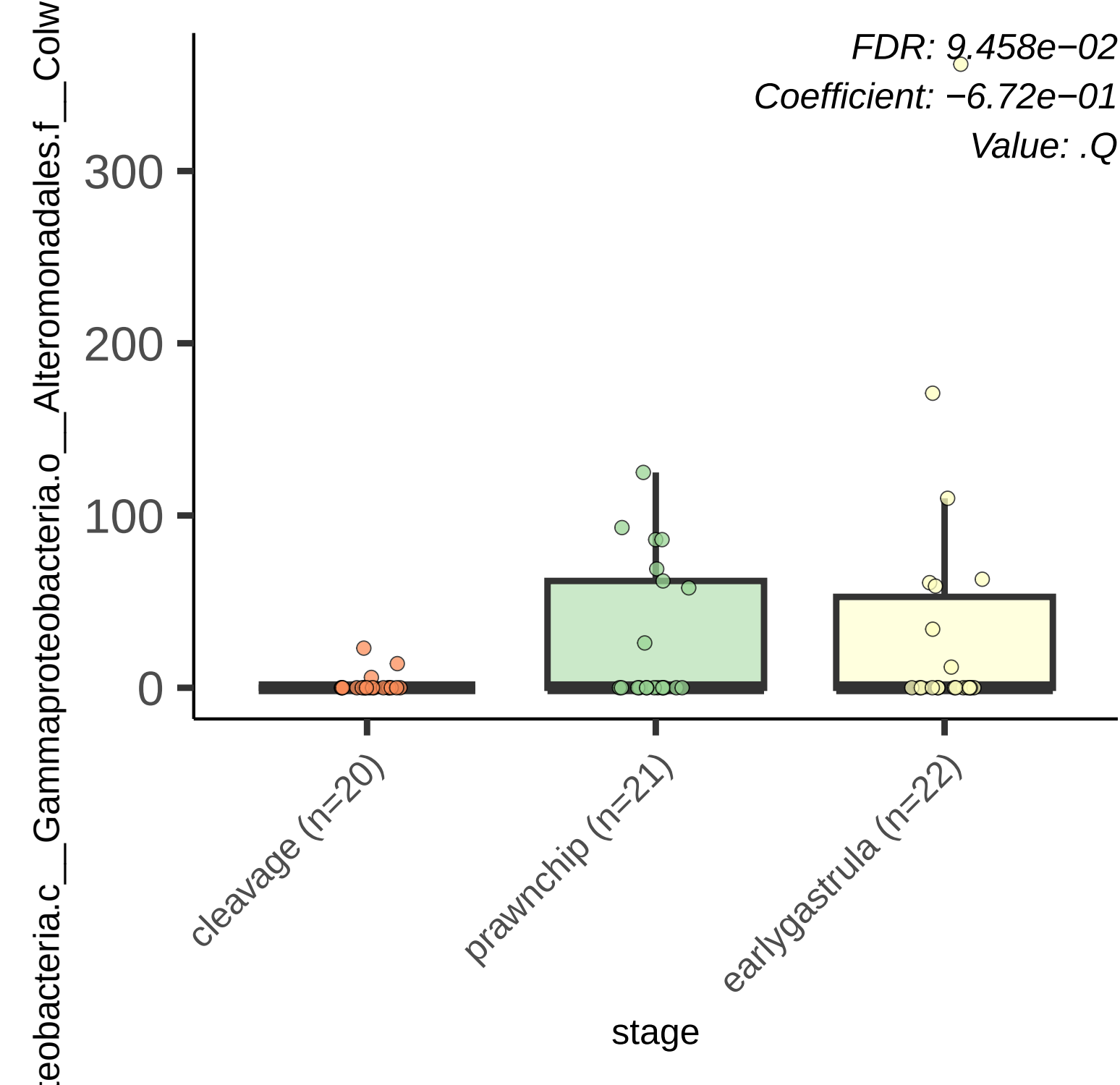
30
20
10
0
cleavage (n=20)
prawnchip (n=21)
earlygastrula (n=22)
stage
d__Bacteria.p__Proteobacteria.c__Alphaproteobacteria.o__Rhizobiales.f__Dev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200
100
0
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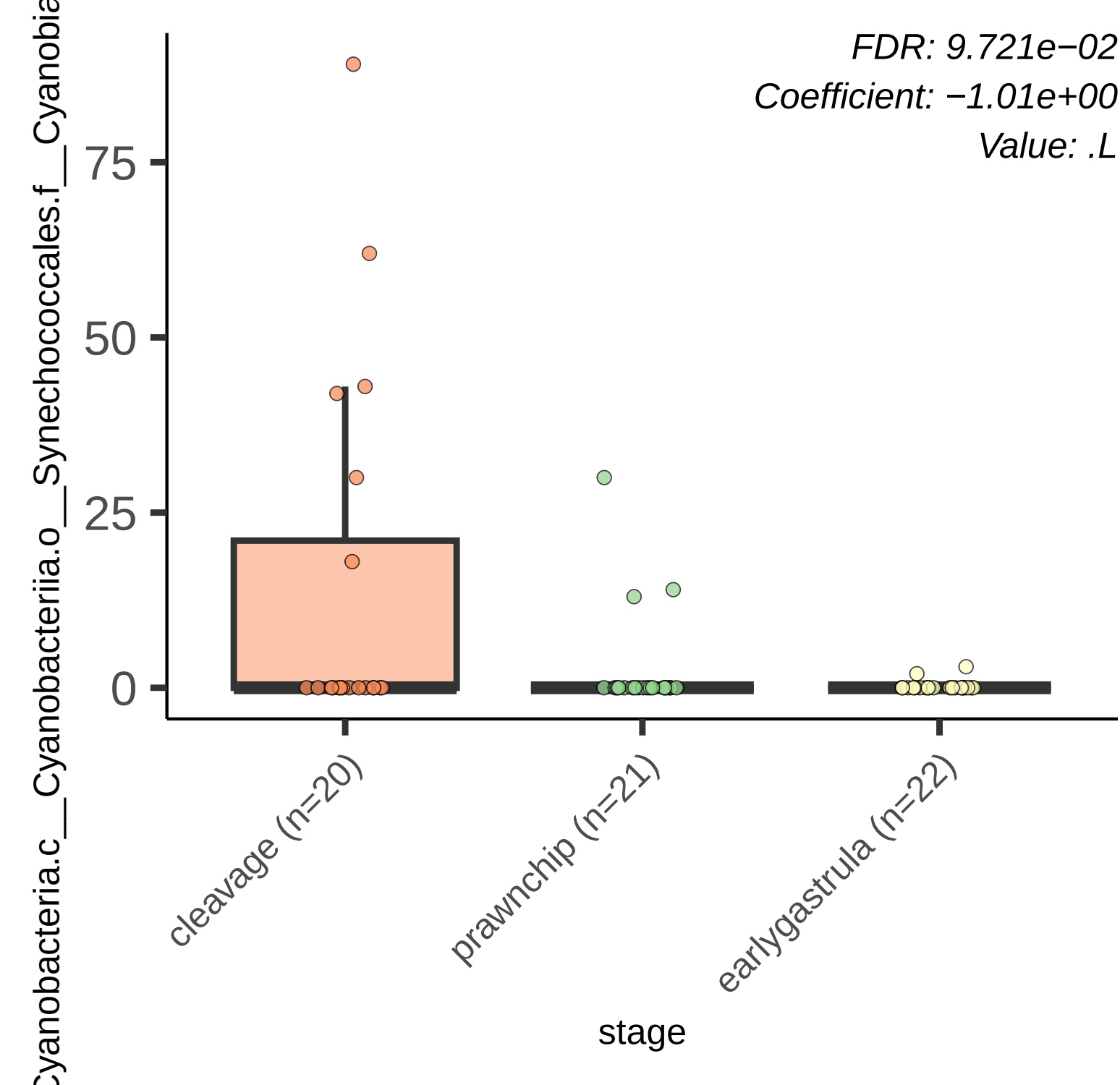
prawnchip (n=21)
earlygastrula (n=22)
stage
teobacteria.c__Gammaproteobacteria.o__Alteromonadales.f__Colw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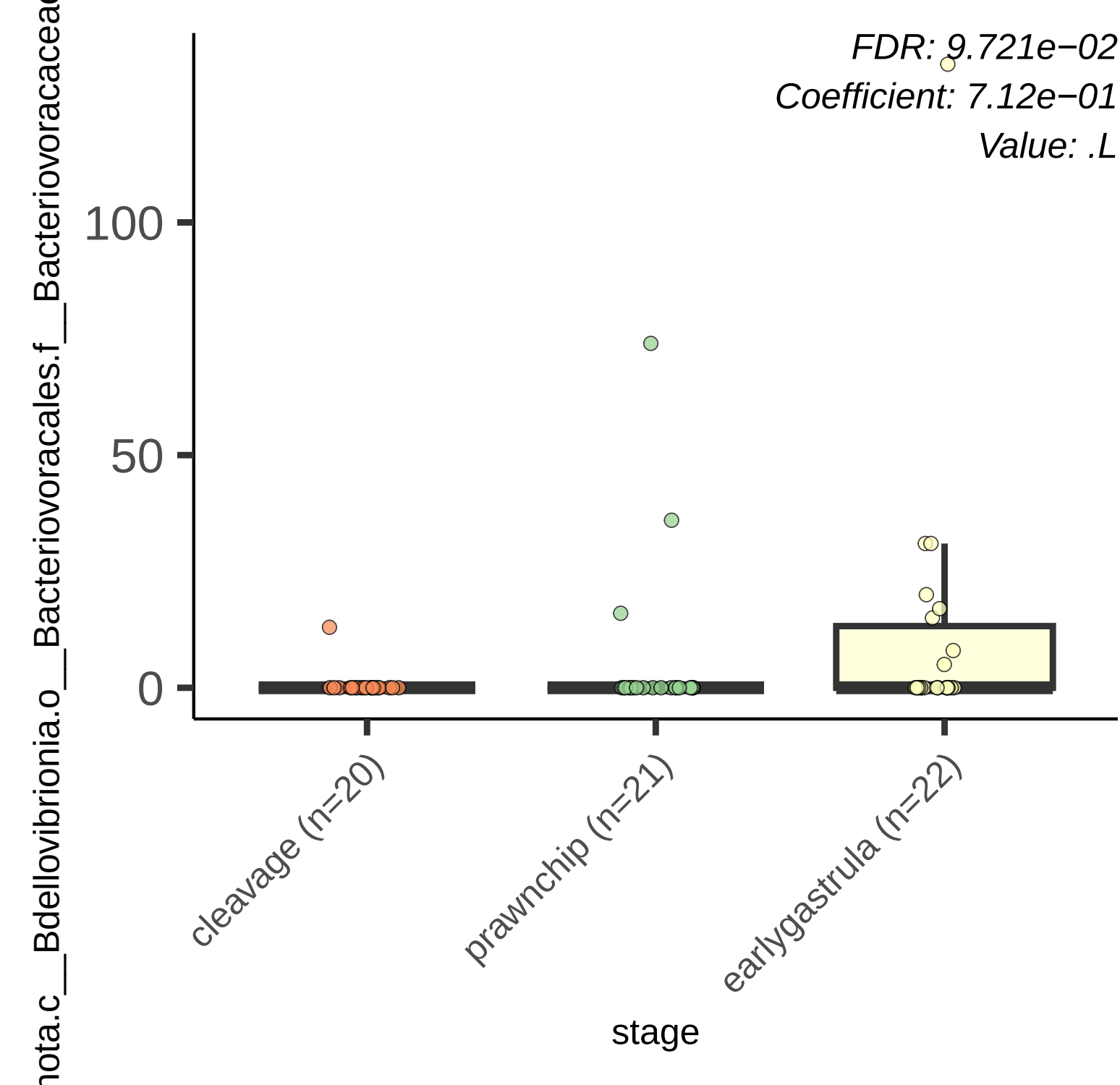
50
25
0
cleavage (n=20)
prawnchip (n=21)
earlygastrula (n=22)
stage
Cyanobacteria.c__Cyanobacteriia.o__Synechococcales.f__Cyanobia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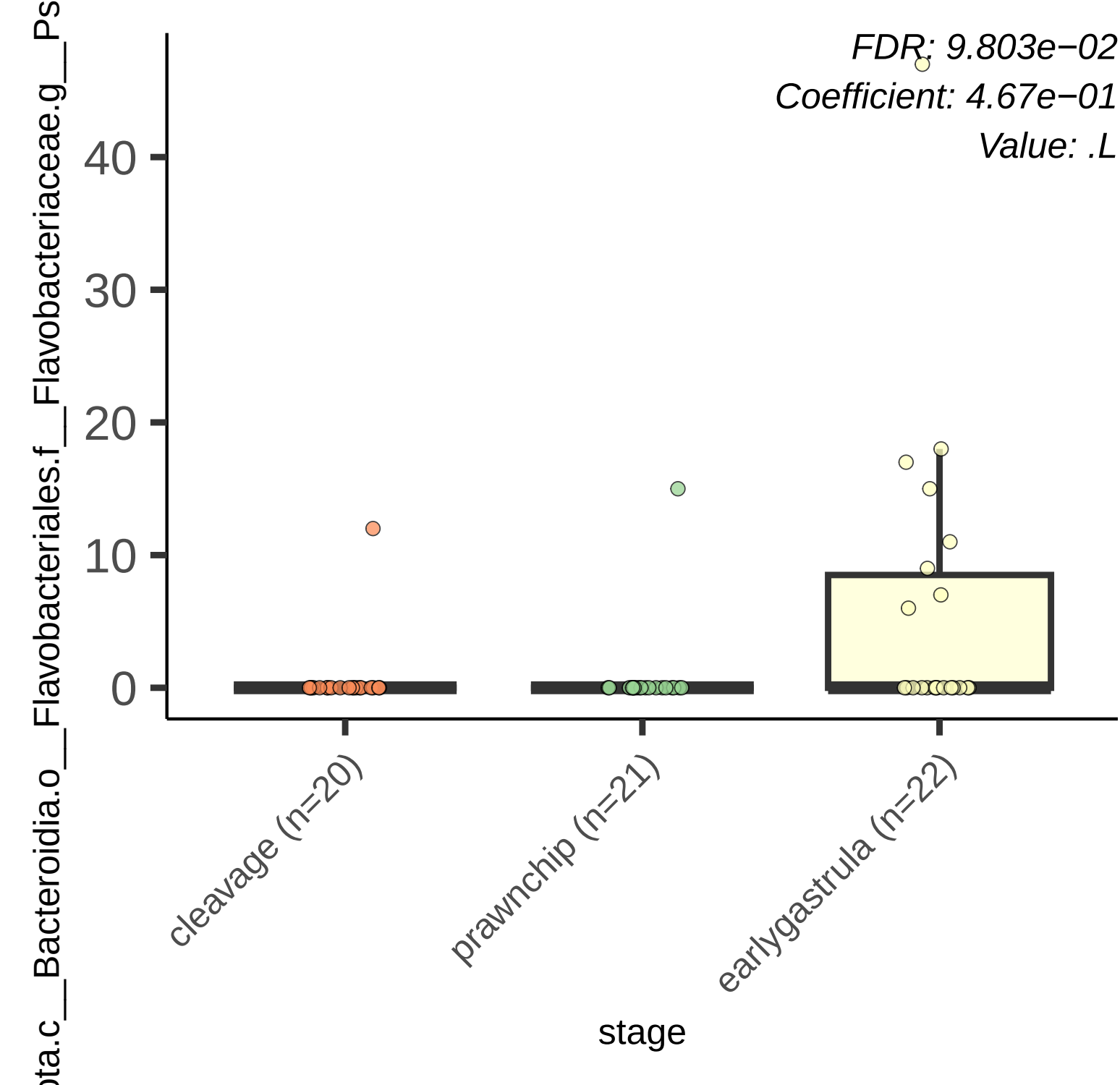
50
0
cleavage (n=20)
prawnchip (n=21)
earlygastrula (n=22)
stage
nota.c__Bdellovibrionia.o__Bacteriovoracales.f__Bacteriovoracacea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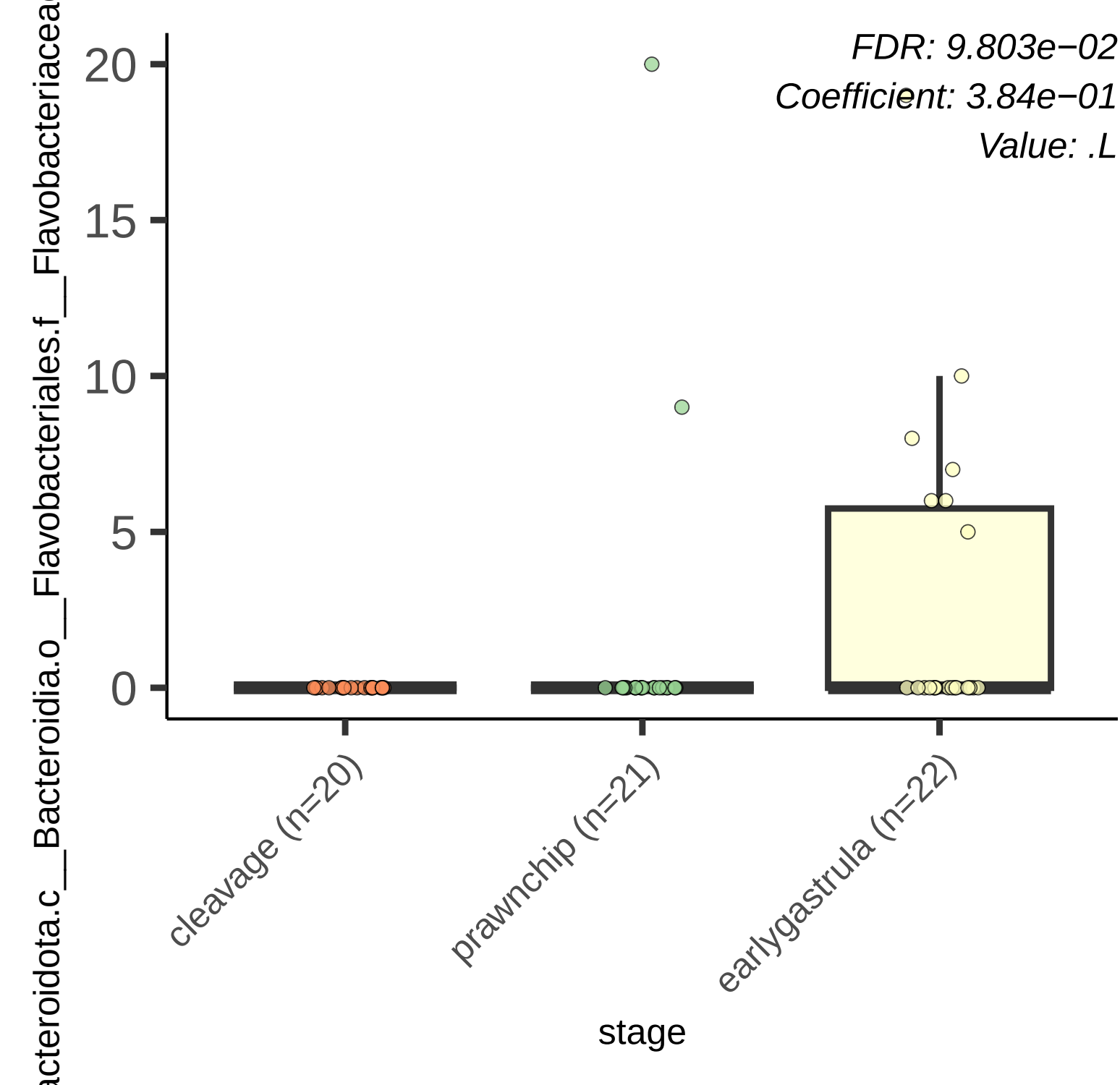
30
20
10
0
cleavage (n=20)
prawnchip (n=21)
earlygastrula (n=22)
stage
ota.c__Bacteroidia.o__Flavobacteriales.f__Flavobacteriaceae.g__Ps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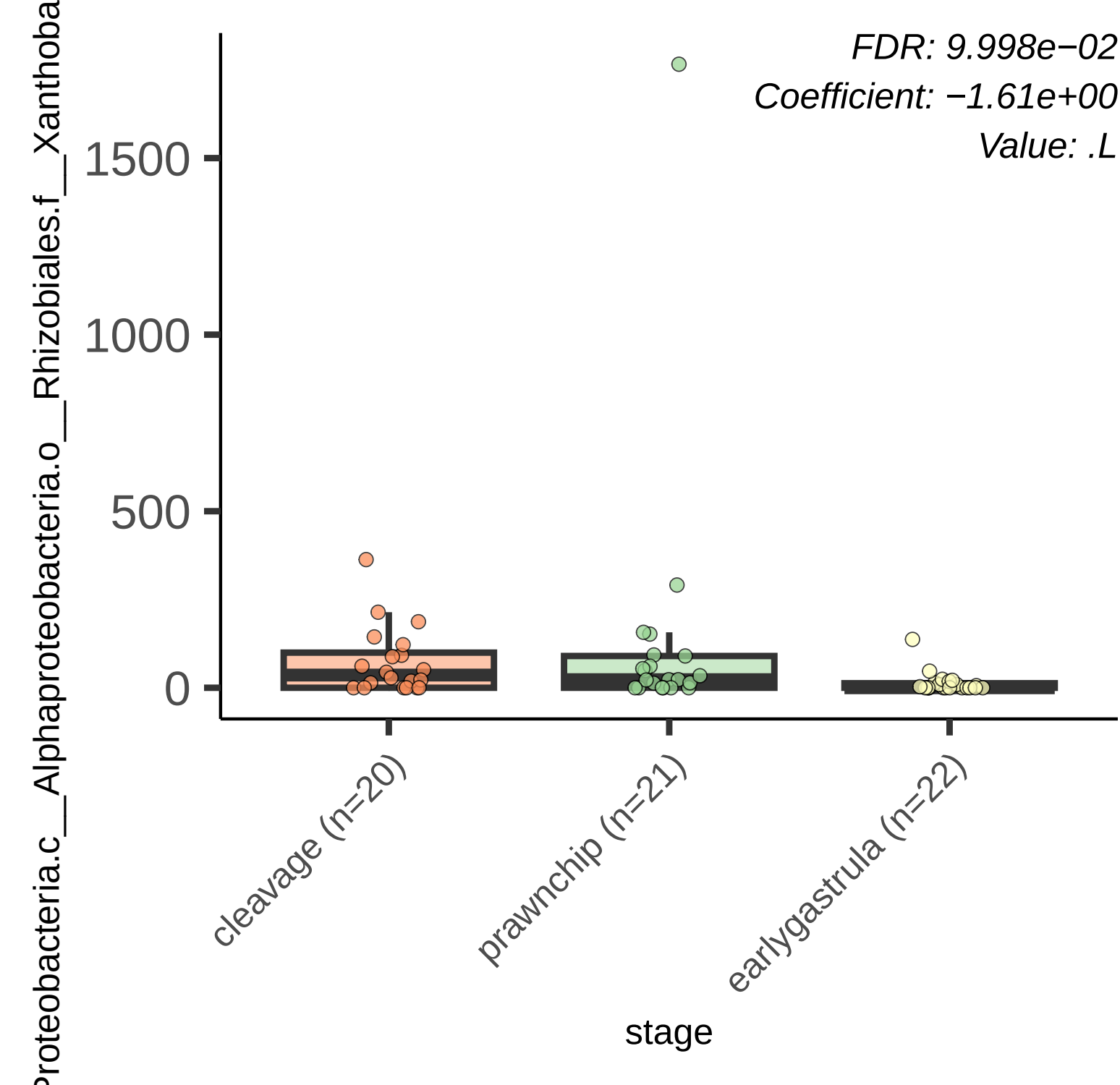
20
15
10
5
0
cleavage (n=20)
prawnchip (n=21)
earlygastrula (n=22)
stage
acteroidota.c__Bacteroidia.o__Flavobacteriales.f__Flavobacteriacea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Coefficient: −1.61e+00
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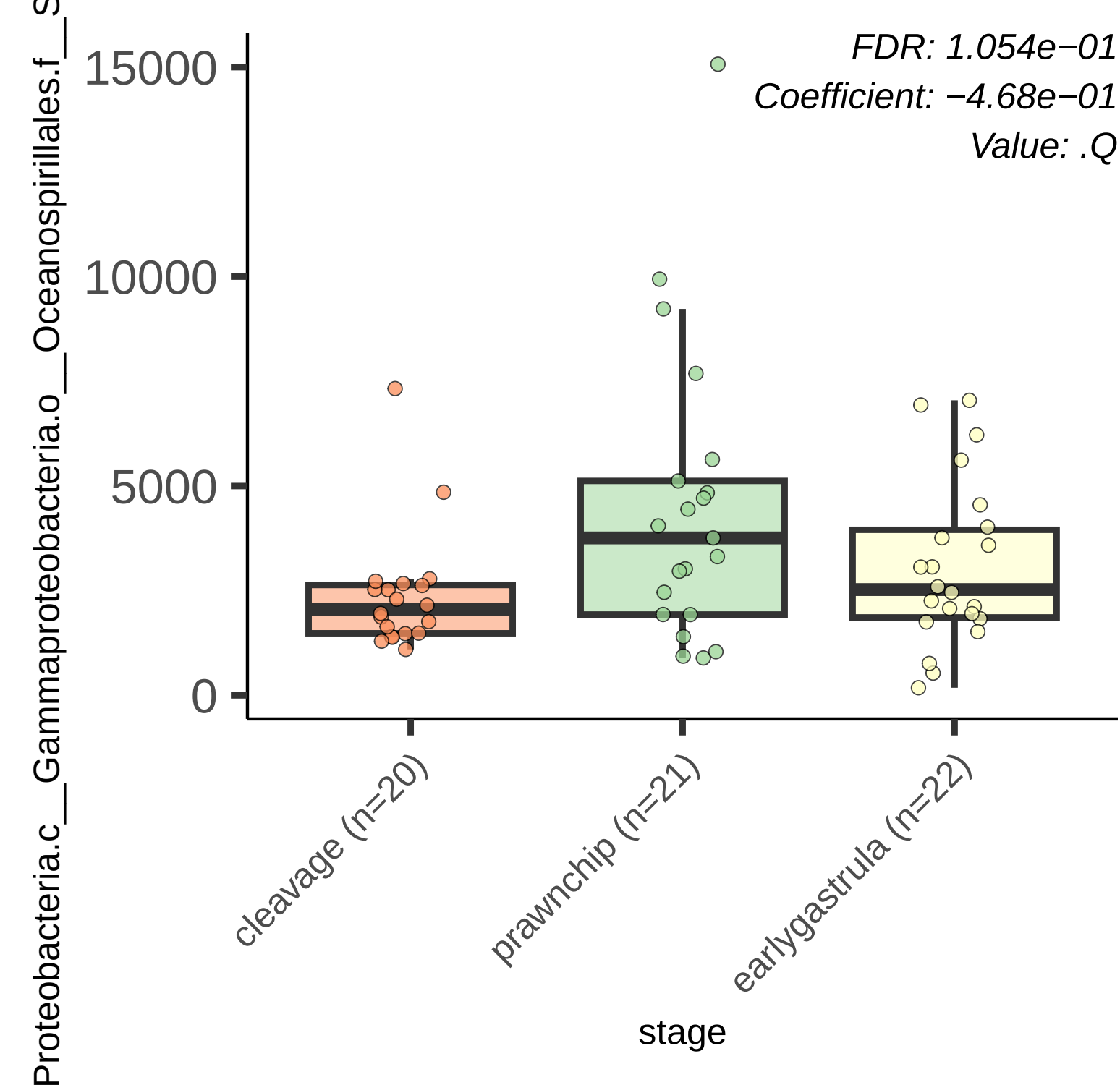
1500
1000
500
0
cleavage (n=20)
prawnchip (n=21)
earlygastrula (n=22)
stage
Proteobacteria.c__Alphaproteobacteria.o__Rhizobiales.f__Xanthoba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Coefficient: −4.68e−01
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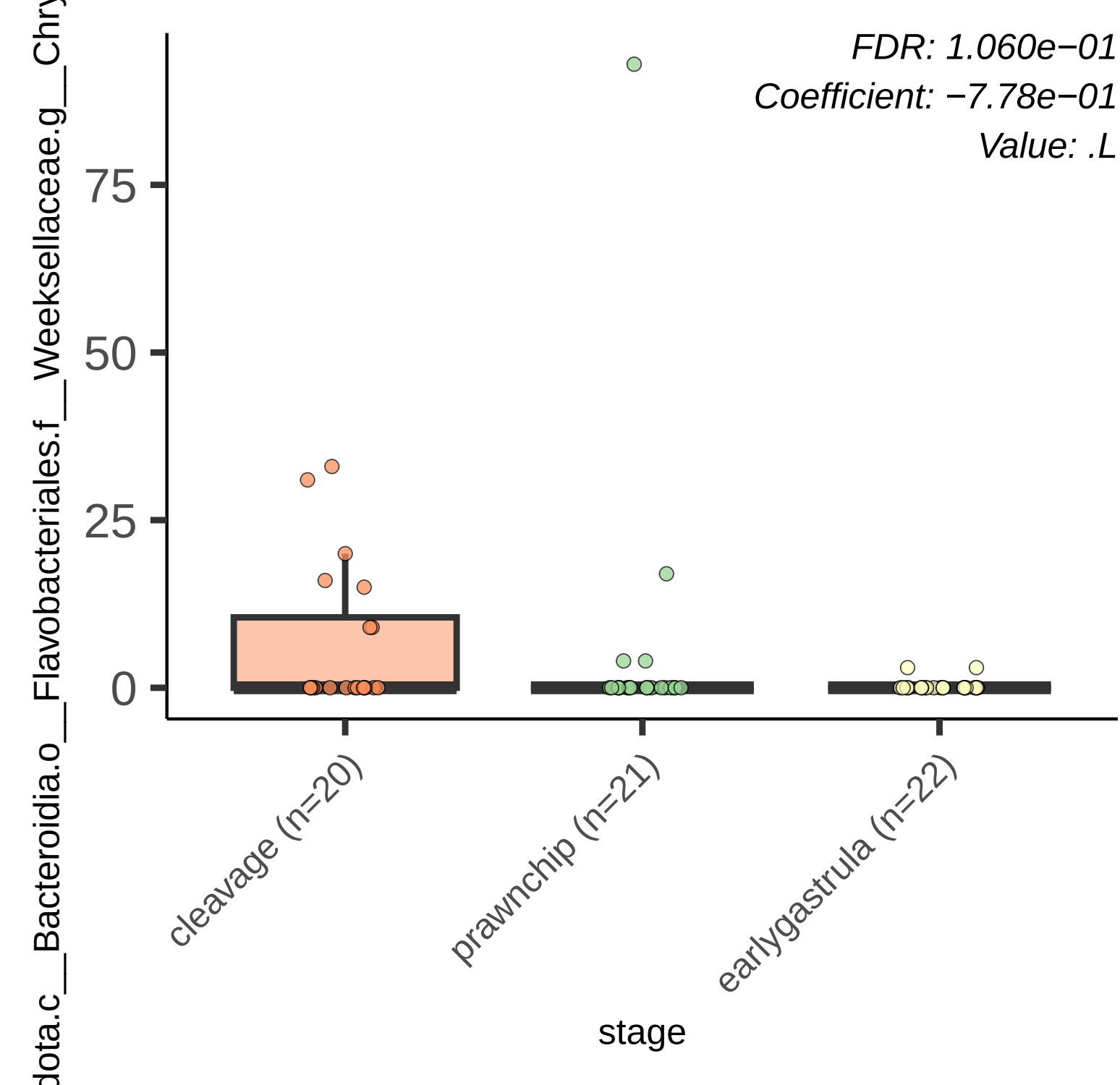
10000
5000
0
cleavage (n=20)
prawnchip (n=21)
earlygastrula (n=22)
stage
Proteobacteria.c__Gammaproteobacteria.o__Oceanospirillales.f__S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Coefficient: −7.78e−01
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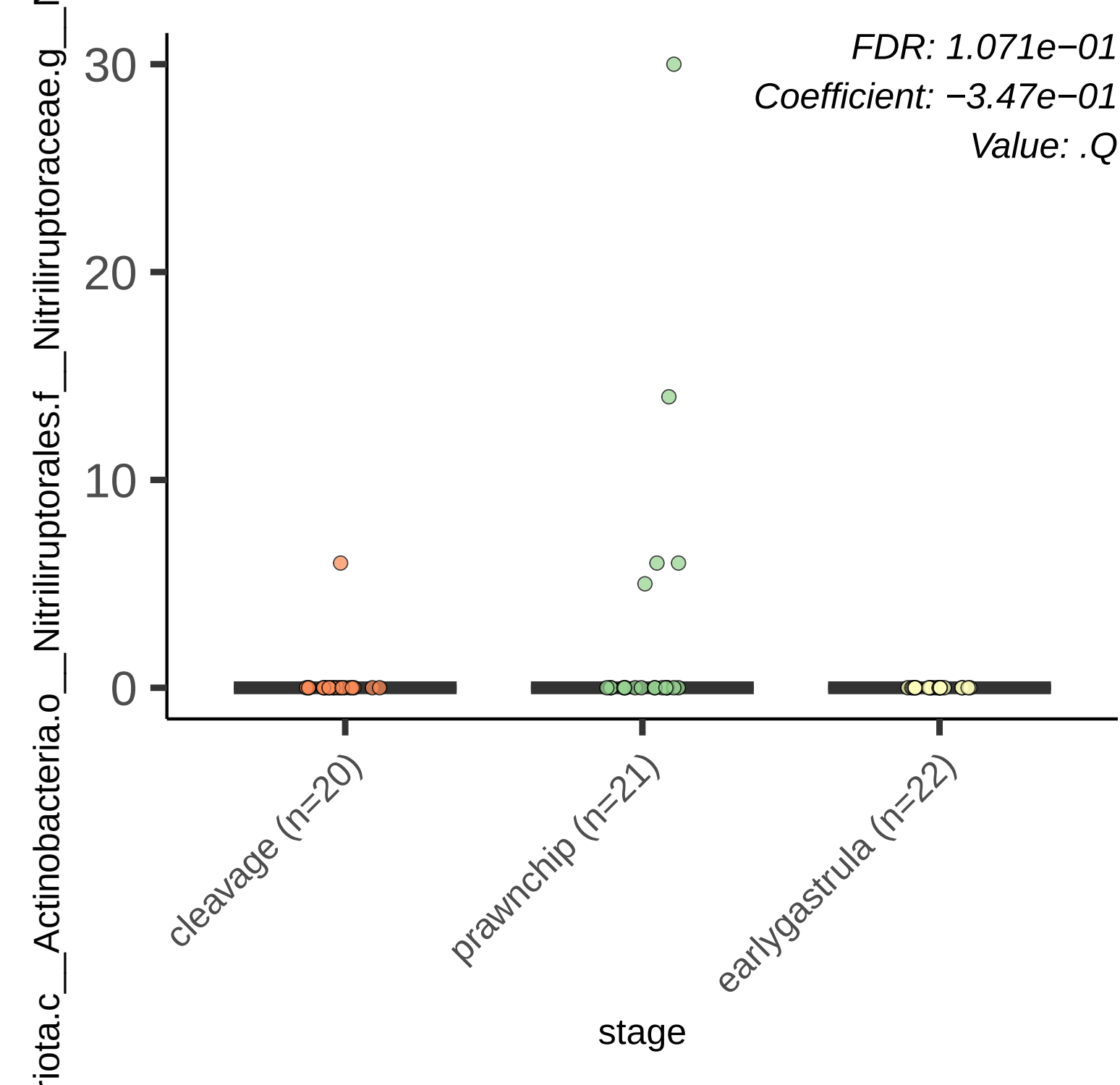
50
25
0
cleavage (n=20)
prawnchip (n=21)
earlygastrula (n=22)
stage
dota.c__Bacteroidia.o__Flavobacteriales.f__Weeksellaceae.g__Chry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10
0
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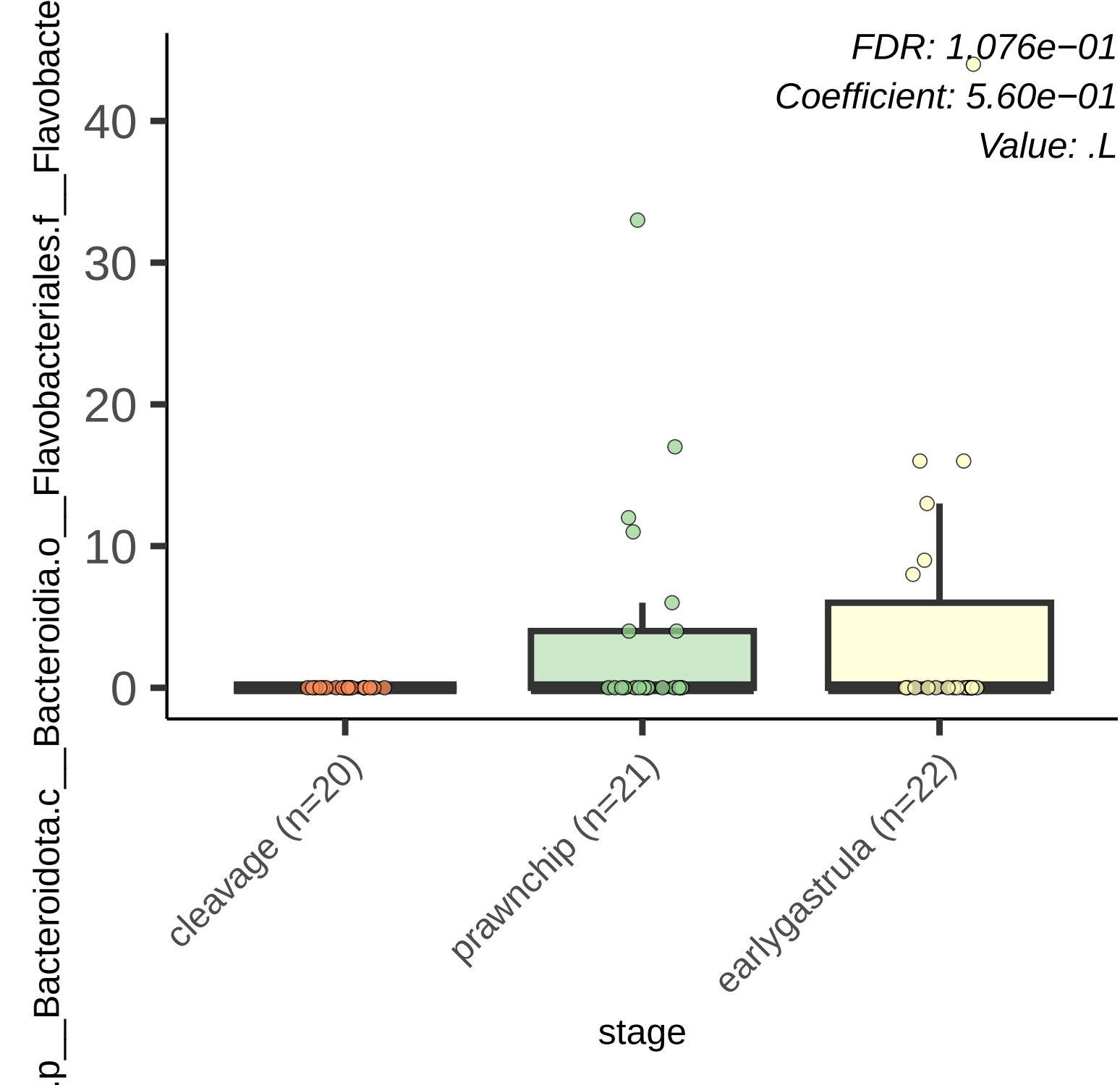
prawnchip (n=21)
earlygastrula (n=22)
stage
riota.c__Actinobacteria.o__Nitriliruptorales.f__Nitriliruptoraceae.g__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Value: .L
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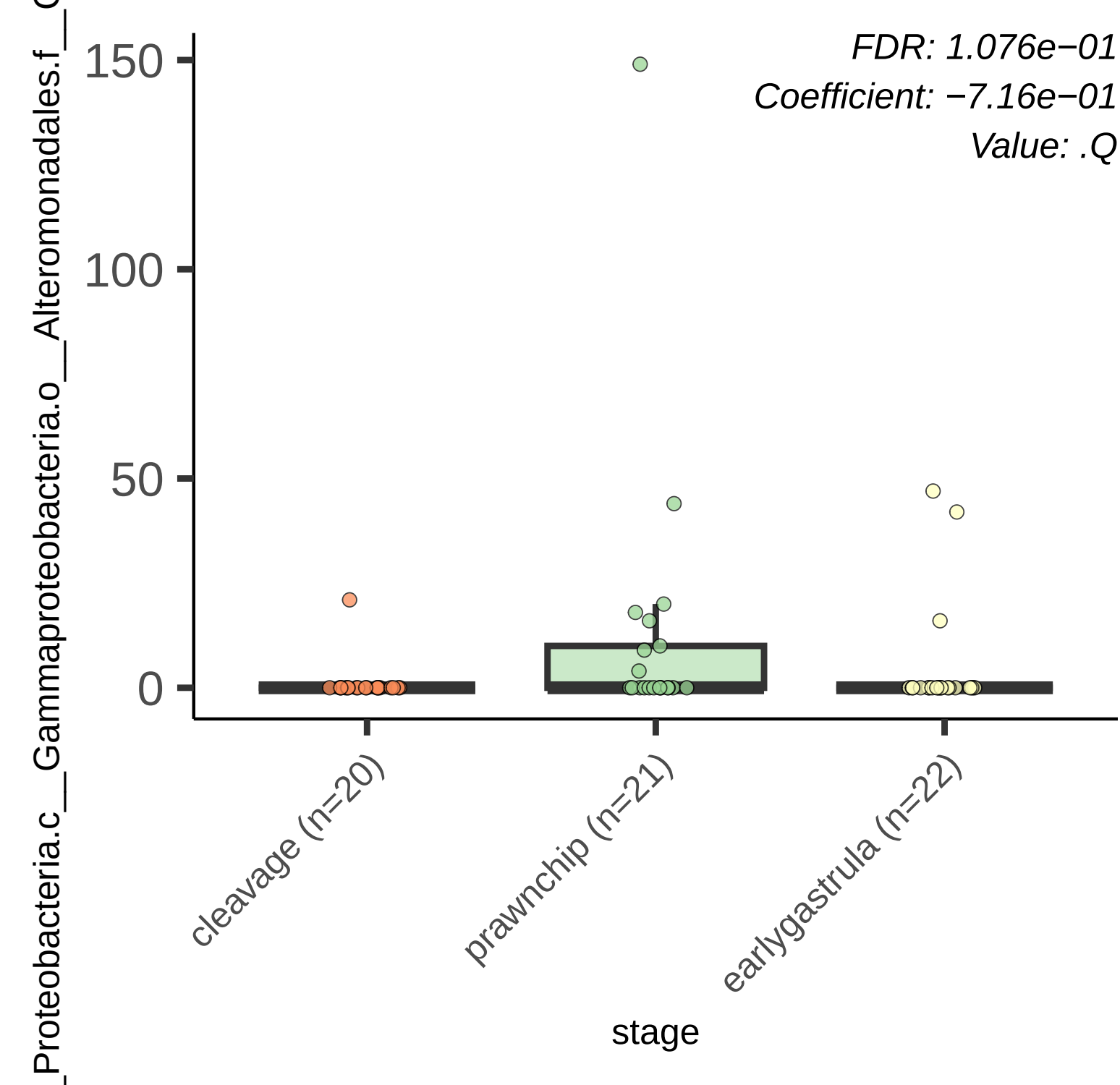
30
20
10
0
cleavage (n=20)
prawnchip (n=21)
earlygastrula (n=22)
stage
.p__Bacteroidota.c__Bacteroidia.o__Flavobacteriales.f__Flavobacte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50
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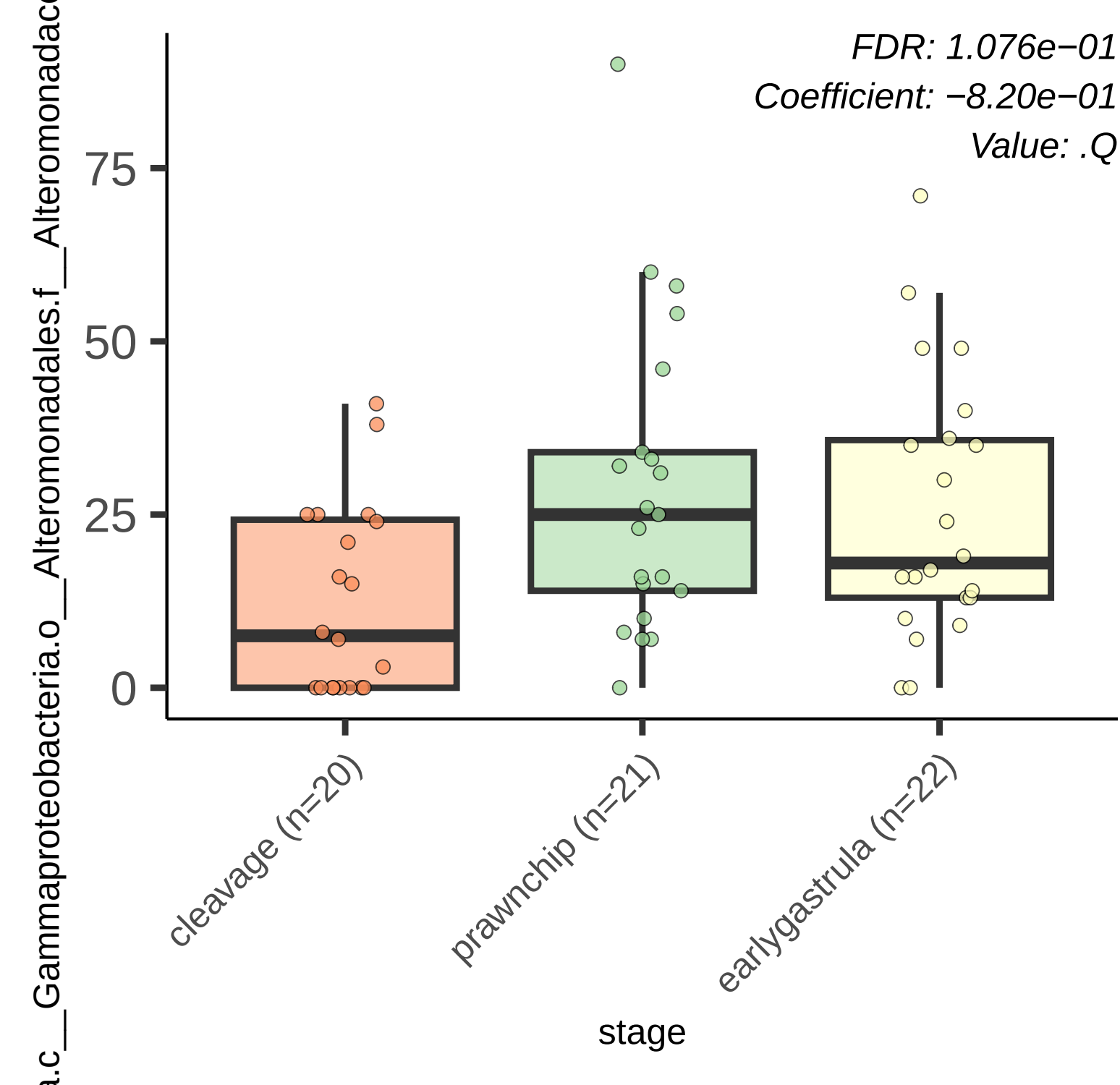
prawnchip (n=21)
earlygastrula (n=22)
stage
_Proteobacteria.c__Gammaproteobacteria.o__Alteromonadales.f__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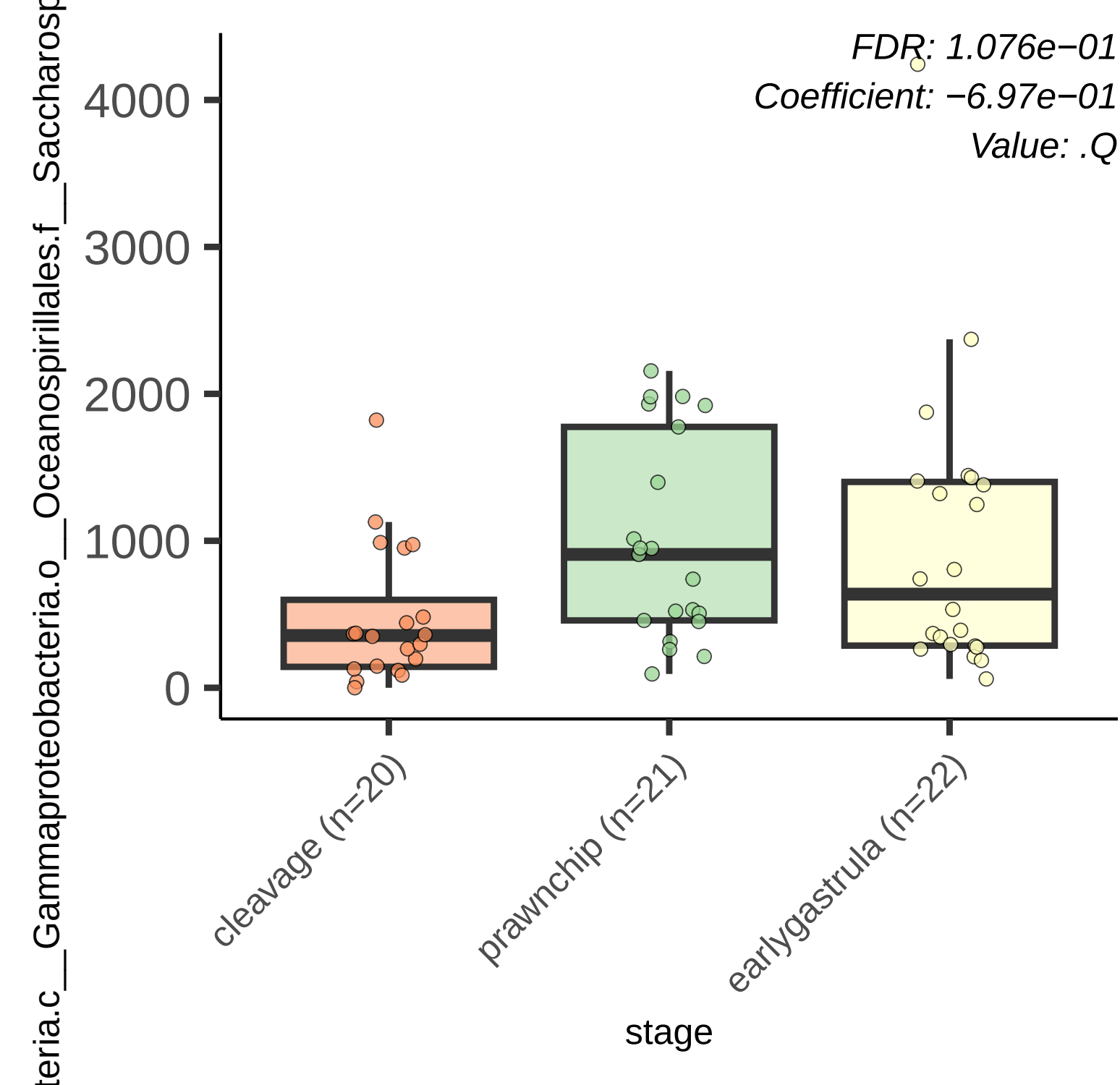
25
0
cleavage (n=20)
prawnchip (n=21)
earlygastrula (n=22)
stage
a.c__Gammaproteobacteria.o__Alteromonadales.f__Alteromonadac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2000
1000
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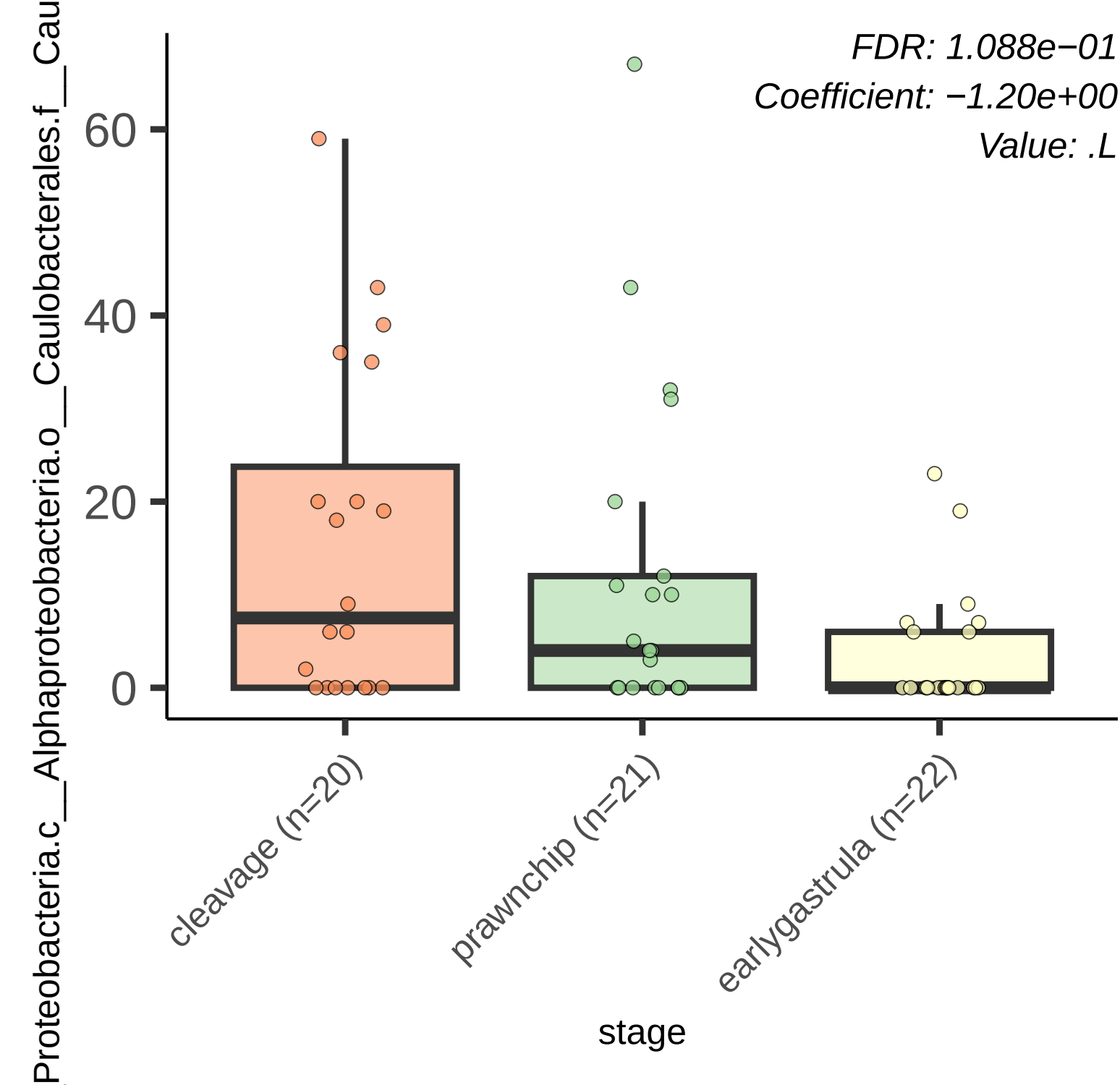
prawnchip (n=21)
earlygastrula (n=22)
stage
teria.c__Gammaproteobacteria.o__Oceanospirillales.f__Saccharosp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20
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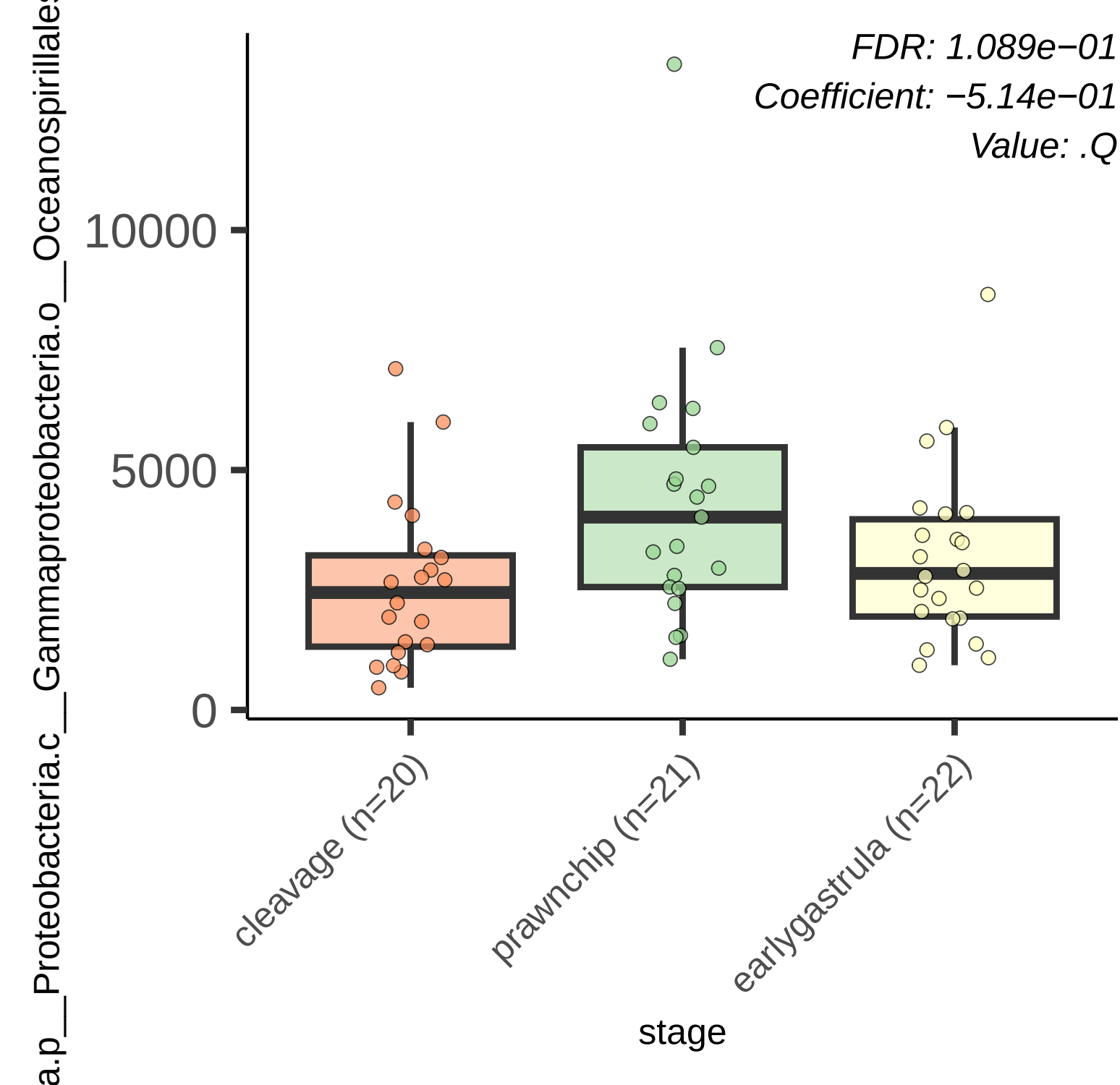
cleavage (n=20)
prawnchip (n=21)
earlygastrula (n=22)
stage
Proteobacteria.c__Alphaproteobacteria.o__Caulobacterales.f__Cau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5000
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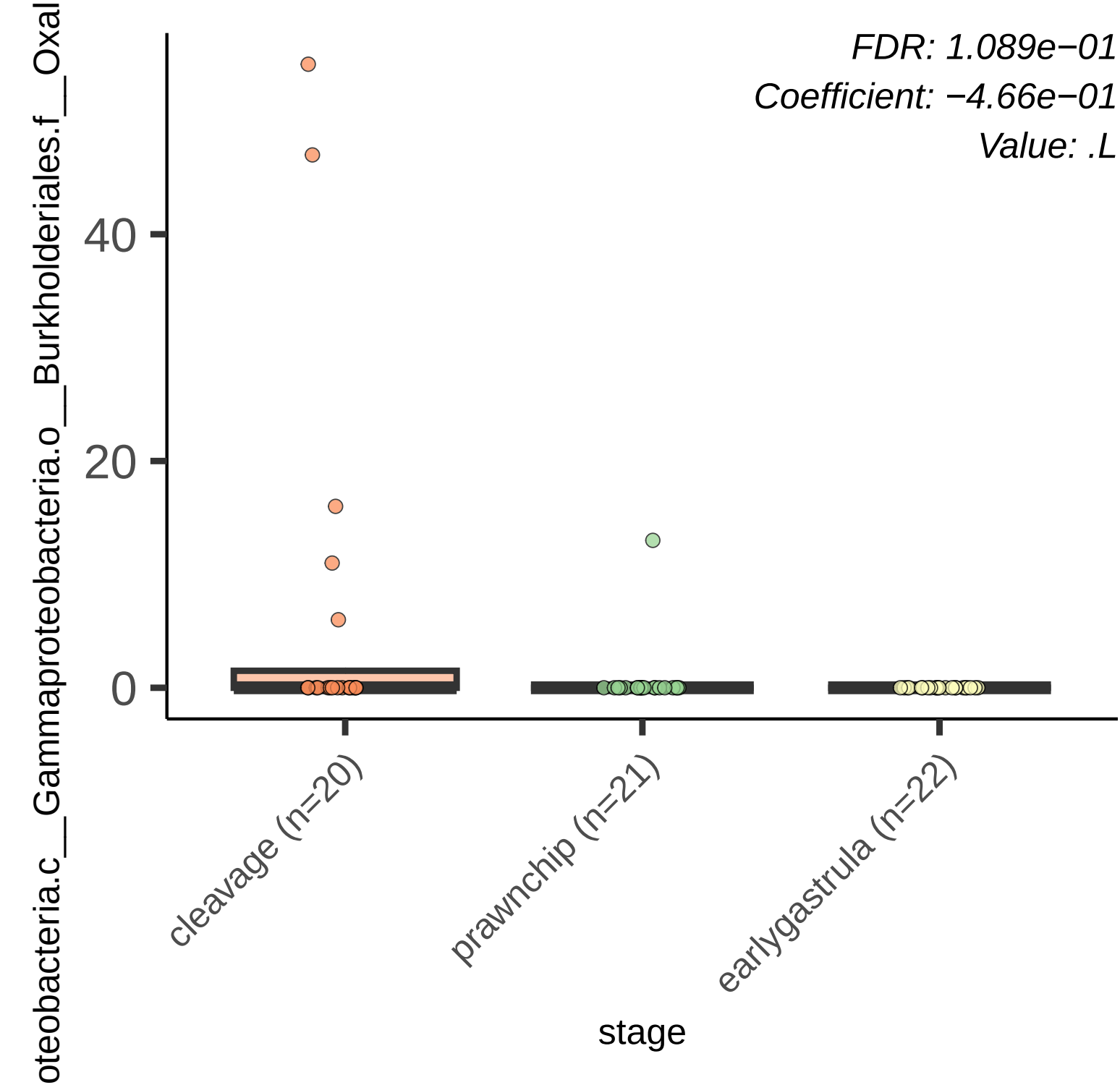
prawnchip (n=21)
earlygastrula (n=22)
stage
a.p__Proteobacteria.c__Gammaproteobacteria.o__Oceanospirillales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20
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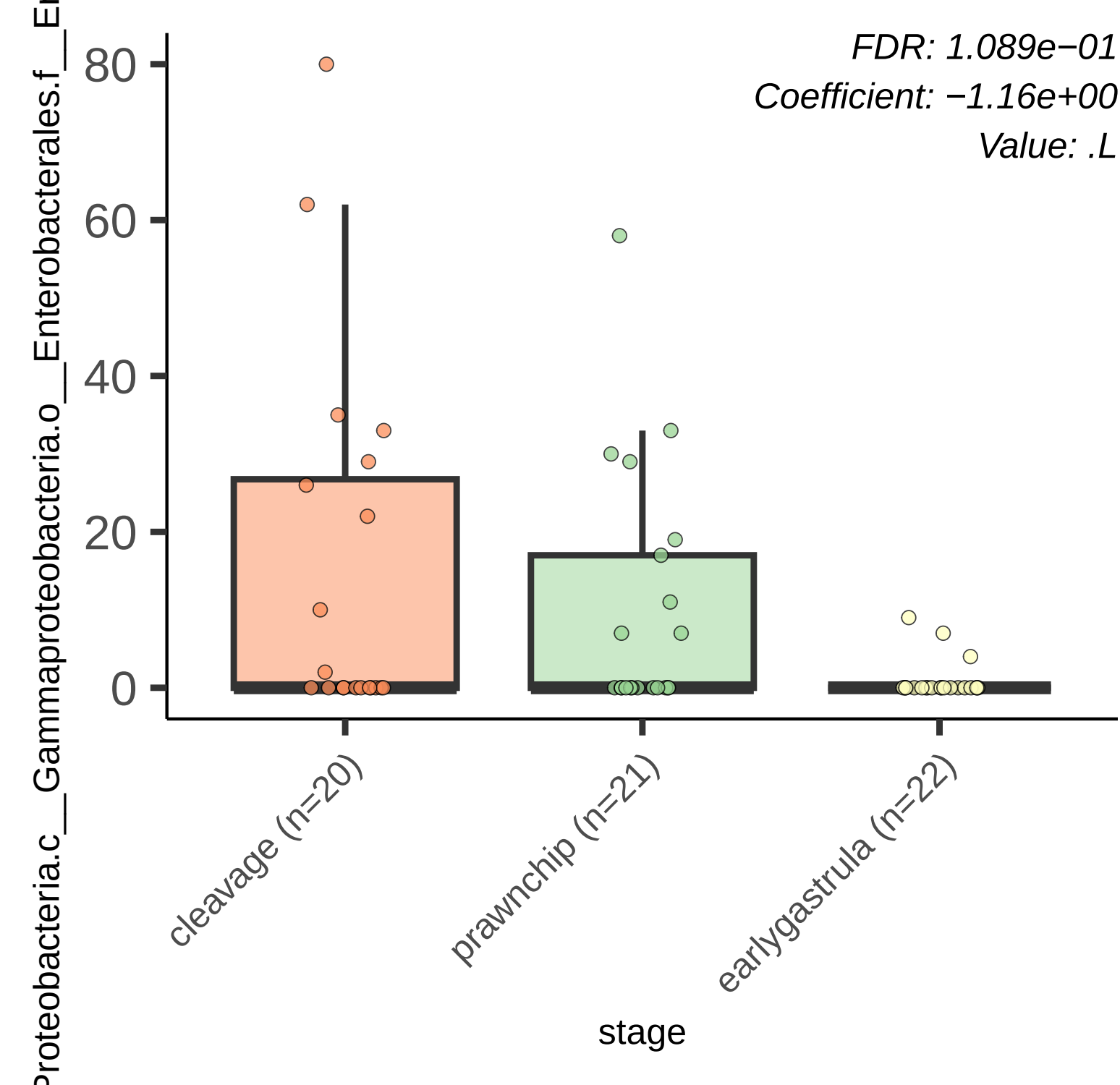
cleavage (n=20)
prawnchip (n=21)
earlygastrula (n=22)
stage
oteobacteria.c__Gammaproteobacteria.o__Burkholderiales.f__Oxal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60
40
20
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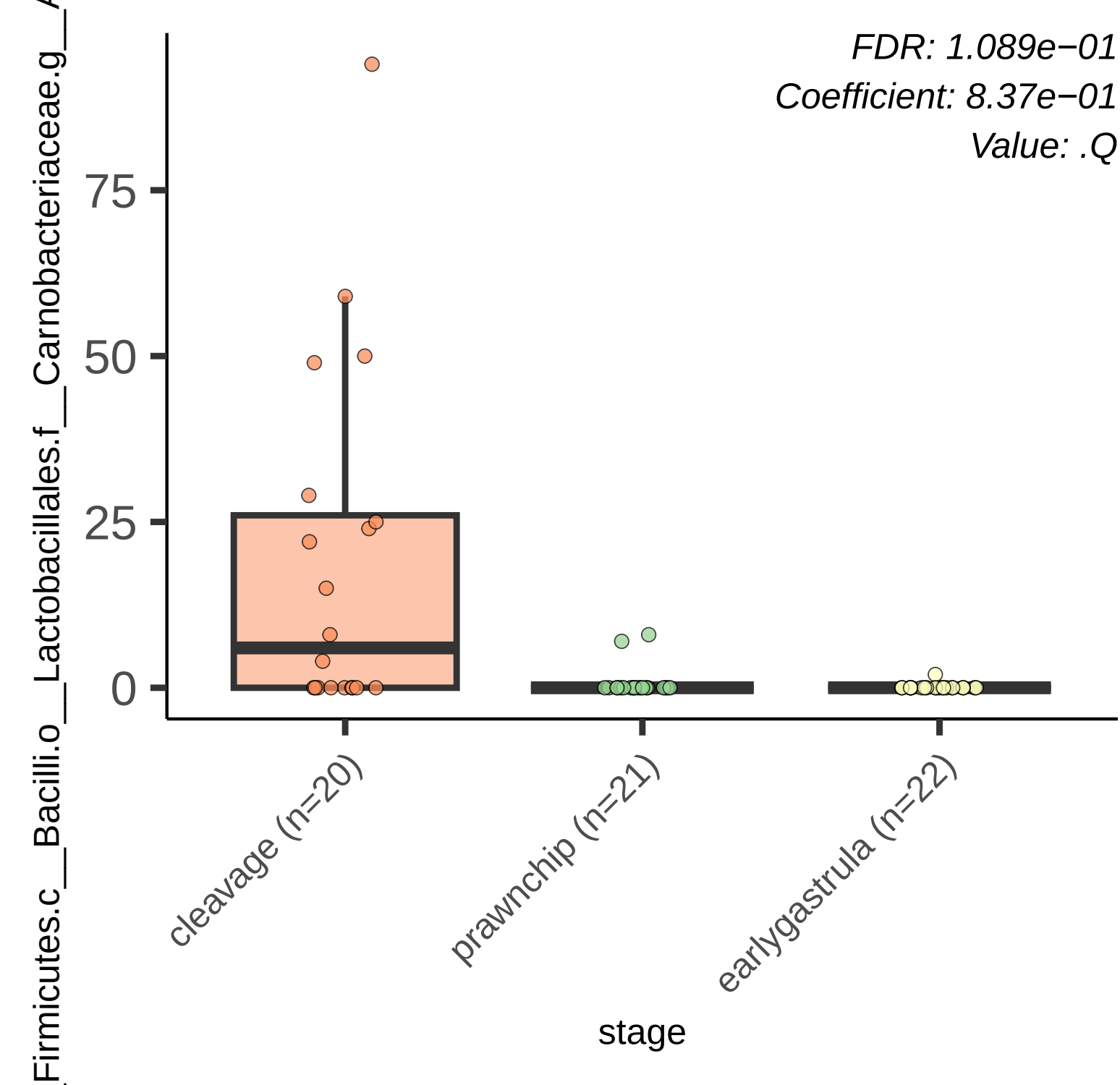
cleavage (n=20)
prawnchip (n=21)
earlygastrula (n=22)
stage
Proteobacteria.c__Gammaproteobacteria.o__Enterobacterales.f__E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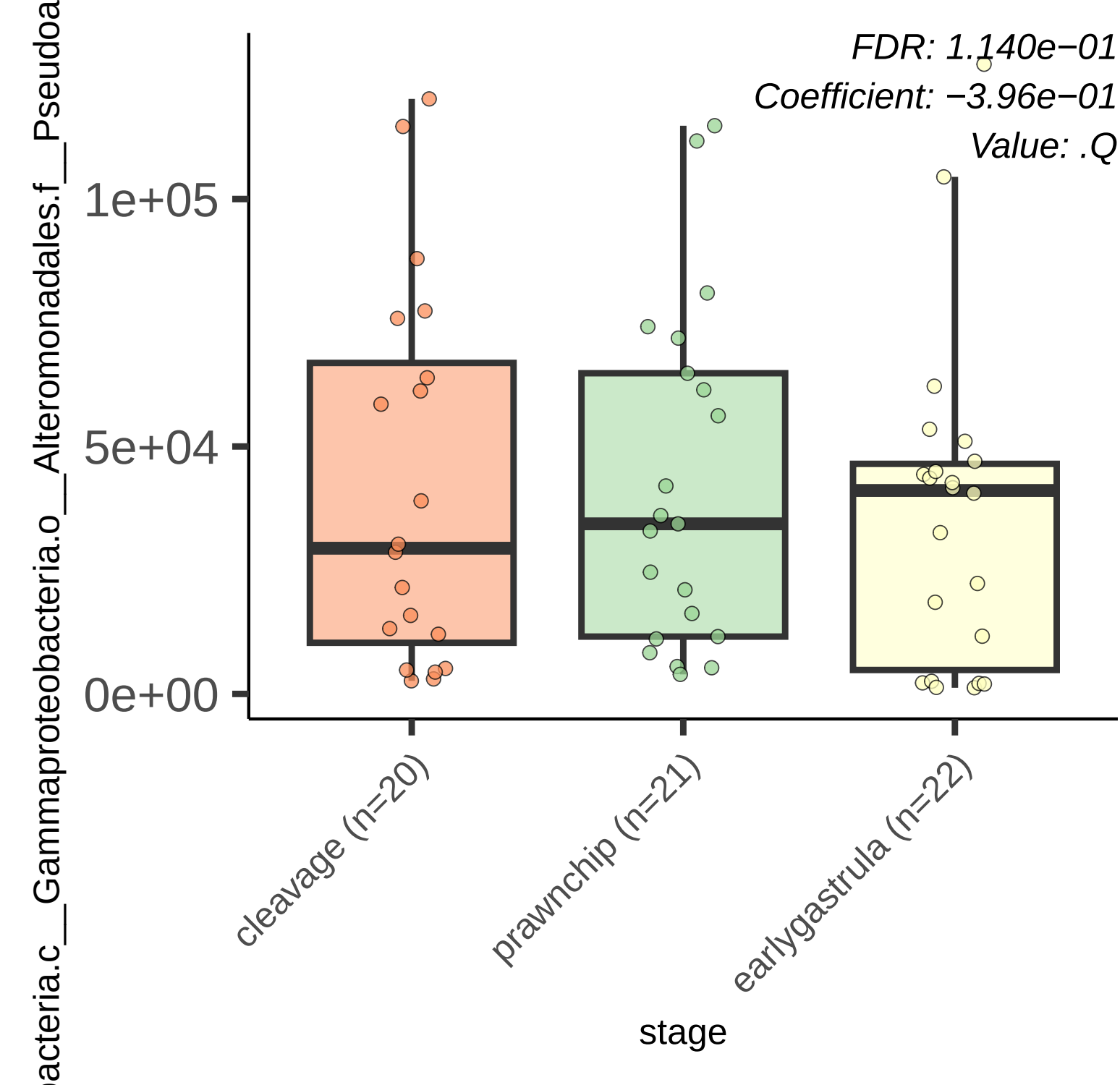
50
25
0
cleavage (n=20)
prawnchip (n=21)
earlygastrula (n=22)
stage
Firmicutes.c__Bacilli.o__Lactobacillales.f__Carnobacteriaceae.g__A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5e+04
0e+00
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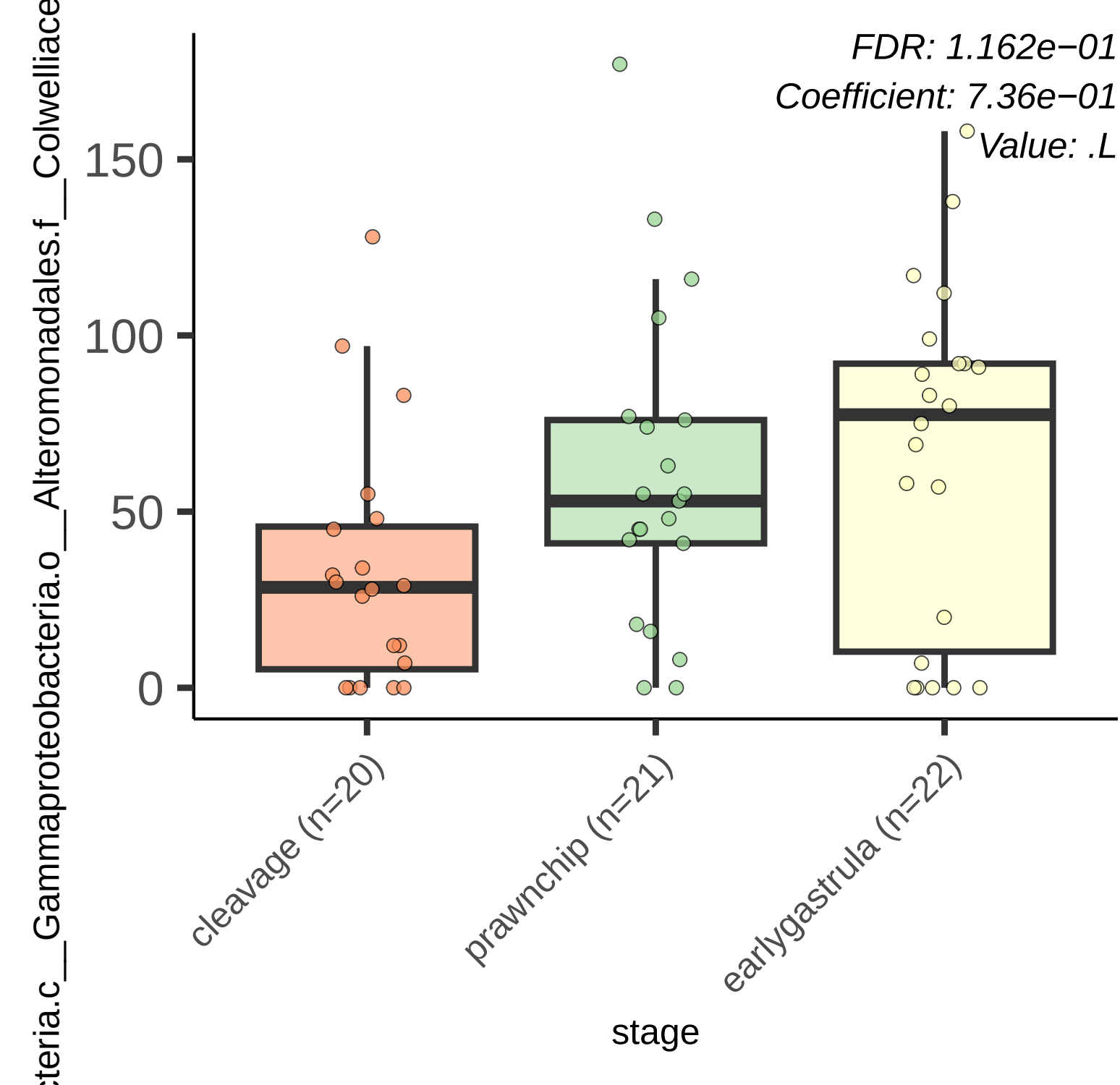
prawnchip (n=21)
earlygastrula (n=22)
stage
bacteria.c__Gammaproteobacteria.o__Alteromonadales.f__Pseudoa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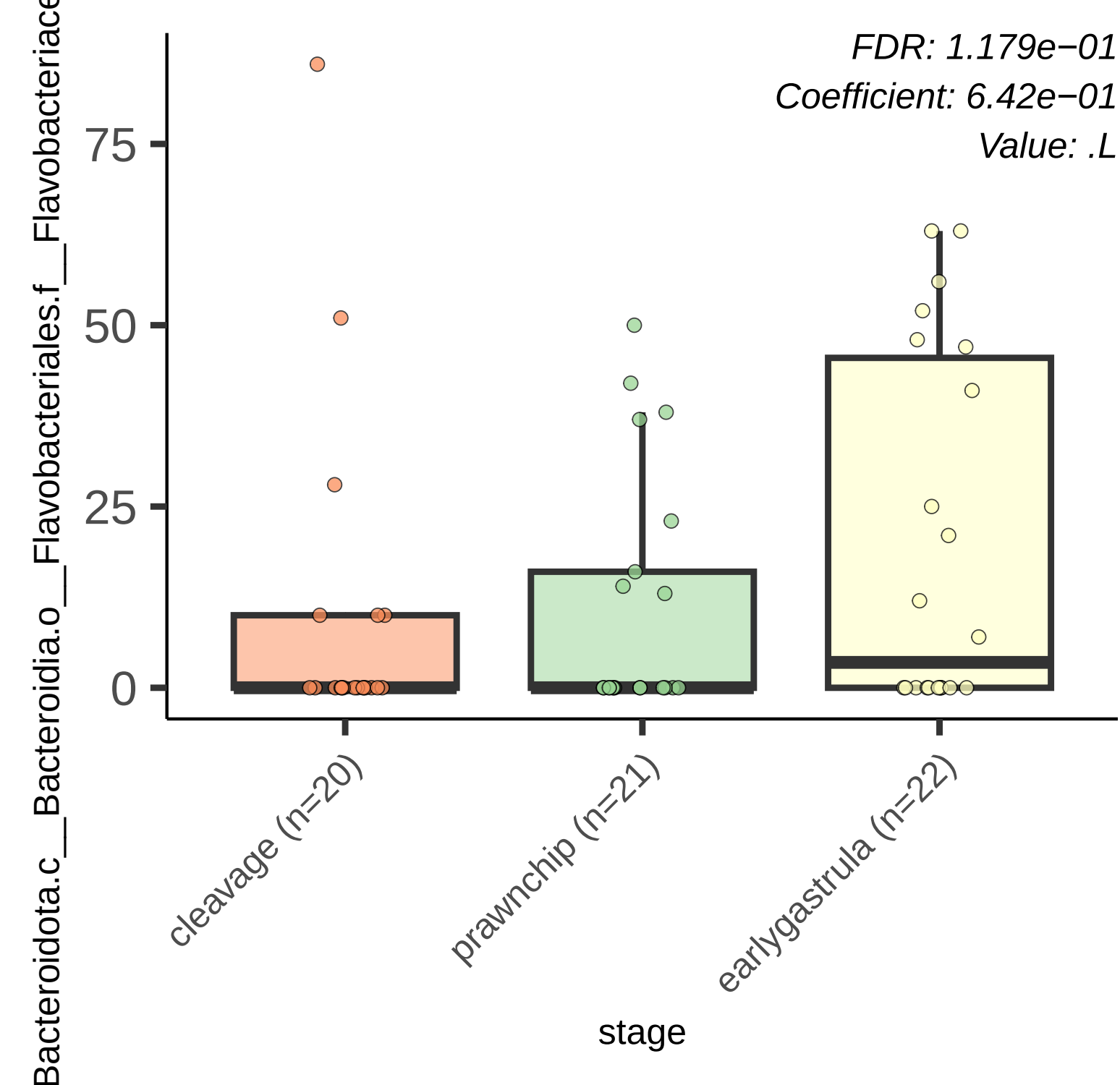
100
50
0
cleavage (n=20)
prawnchip (n=21)
earlygastrula (n=22)
stage
cteria.c__Gammaproteobacteria.o__Alteromonadales.f__Colwelliace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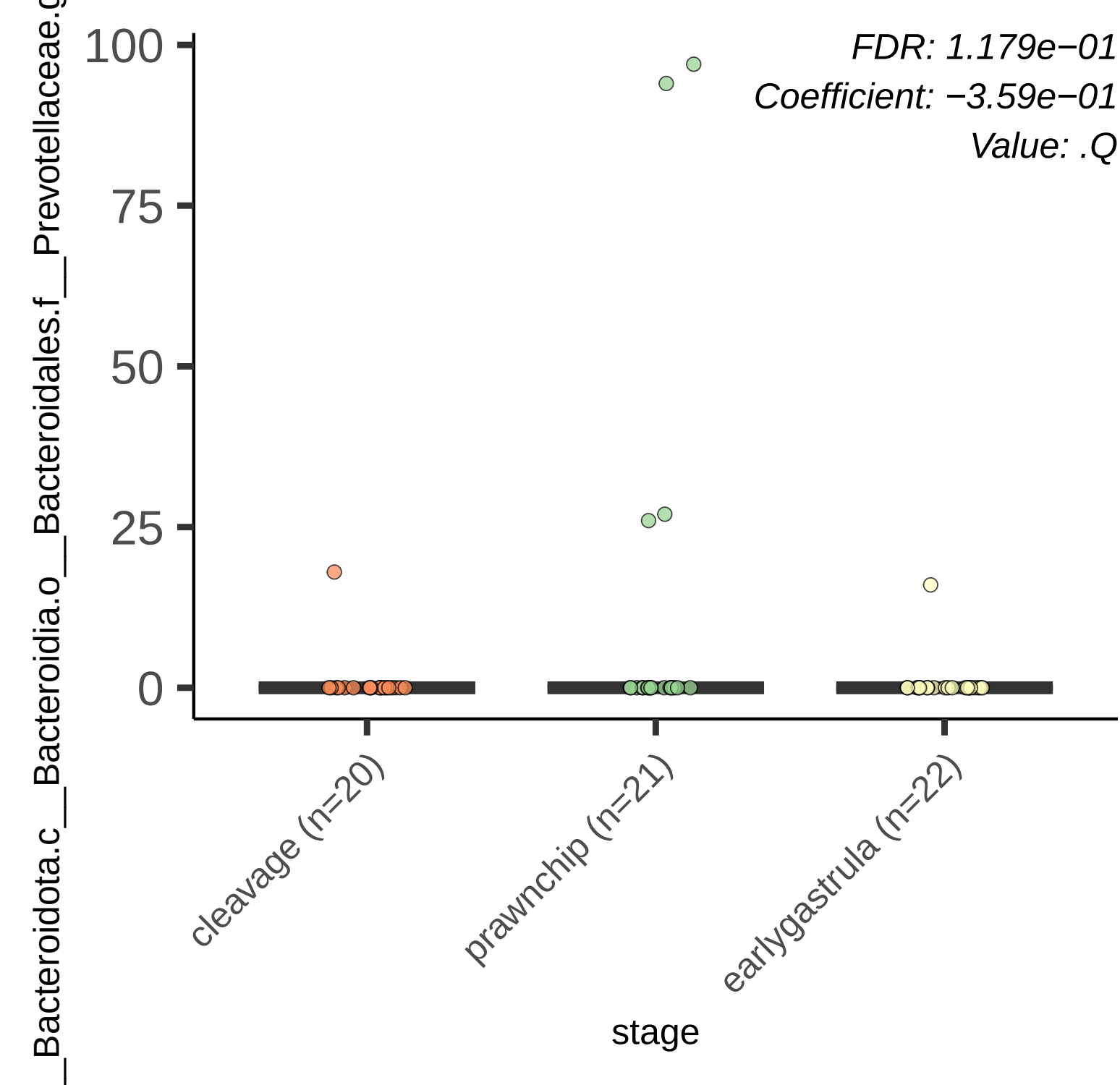
50
25
0
cleavage (n=20)
prawnchip (n=21)
earlygastrula (n=22)
stage
Bacteroidota.c__Bacteroidia.o__Flavobacteriales.f__Flavobacteriace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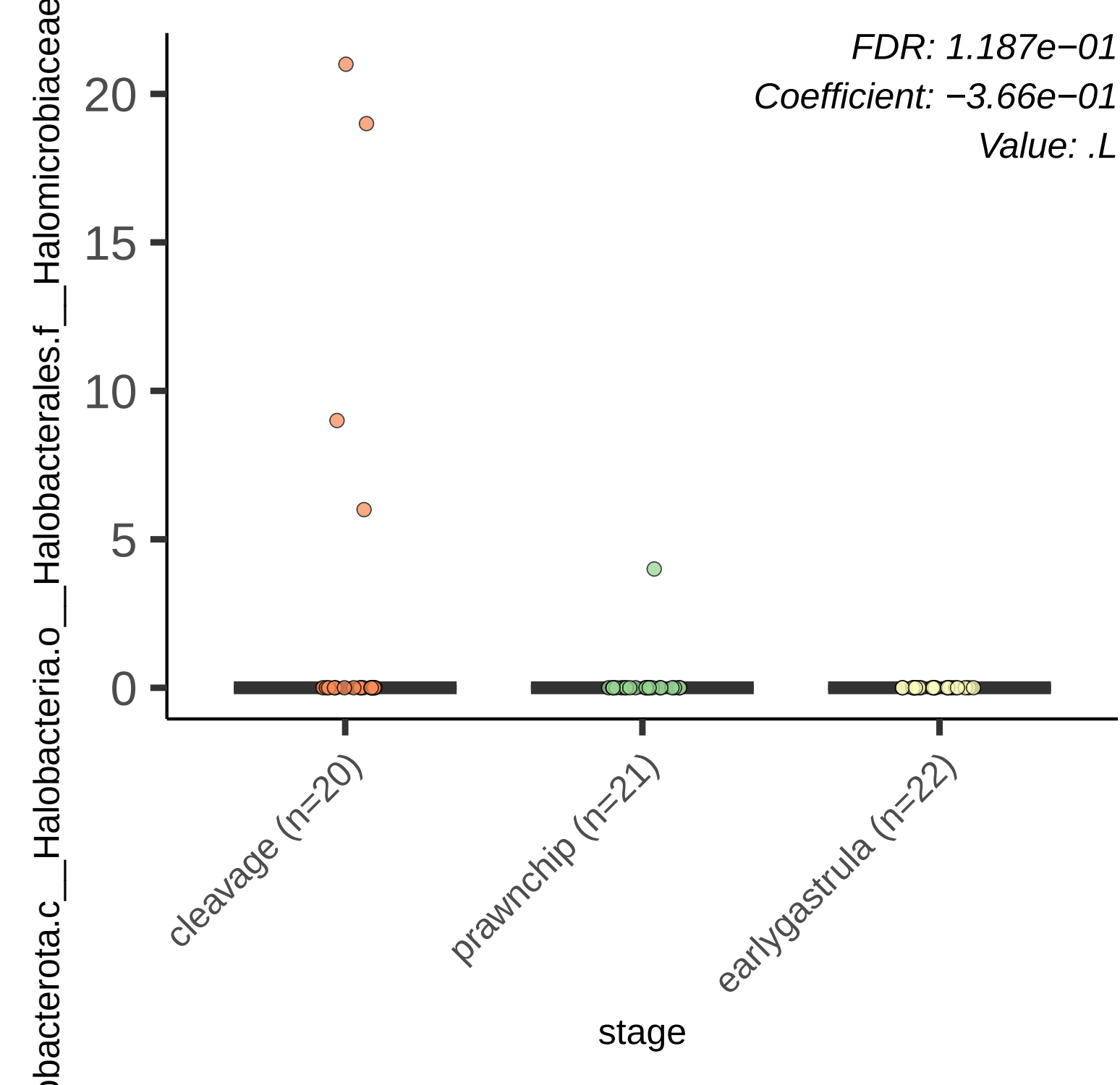
Value: .Q
100
75
50
25
0
cleavage (n=20)
prawnchip (n=21)
earlygastrula (n=22)
stage
__Bacteroidota.c__Bacteroidia.o__Bacteroidales.f__Prevotellaceae.g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Coefficient: −3.66e−01
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15
10
5
0
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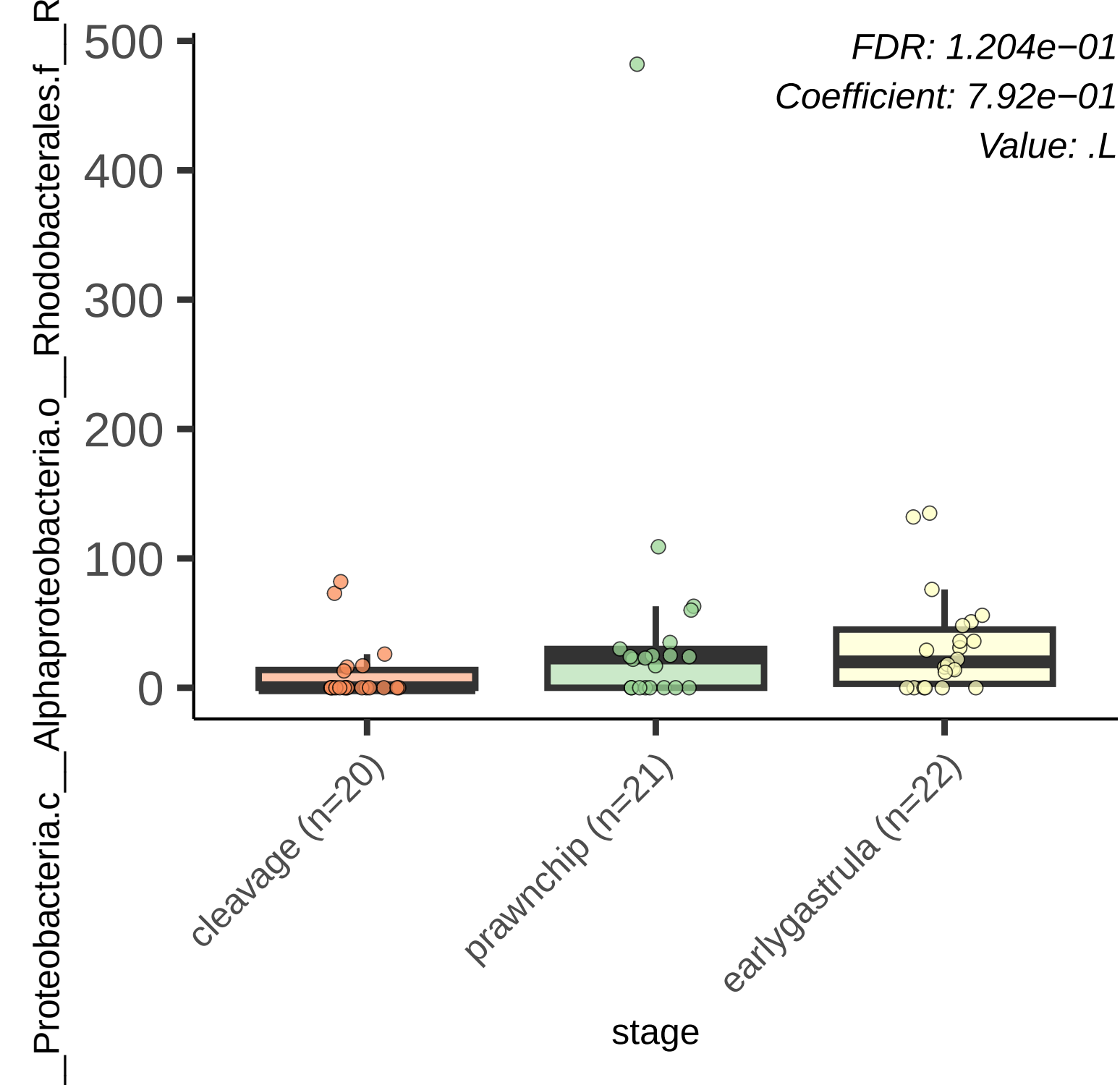
prawnchip (n=21)
earlygastrula (n=22)
stage
obacterota.c__Halobacteria.o__Halobacterales.f__Halomicrobiaceae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Coefficient: 7.92e−01
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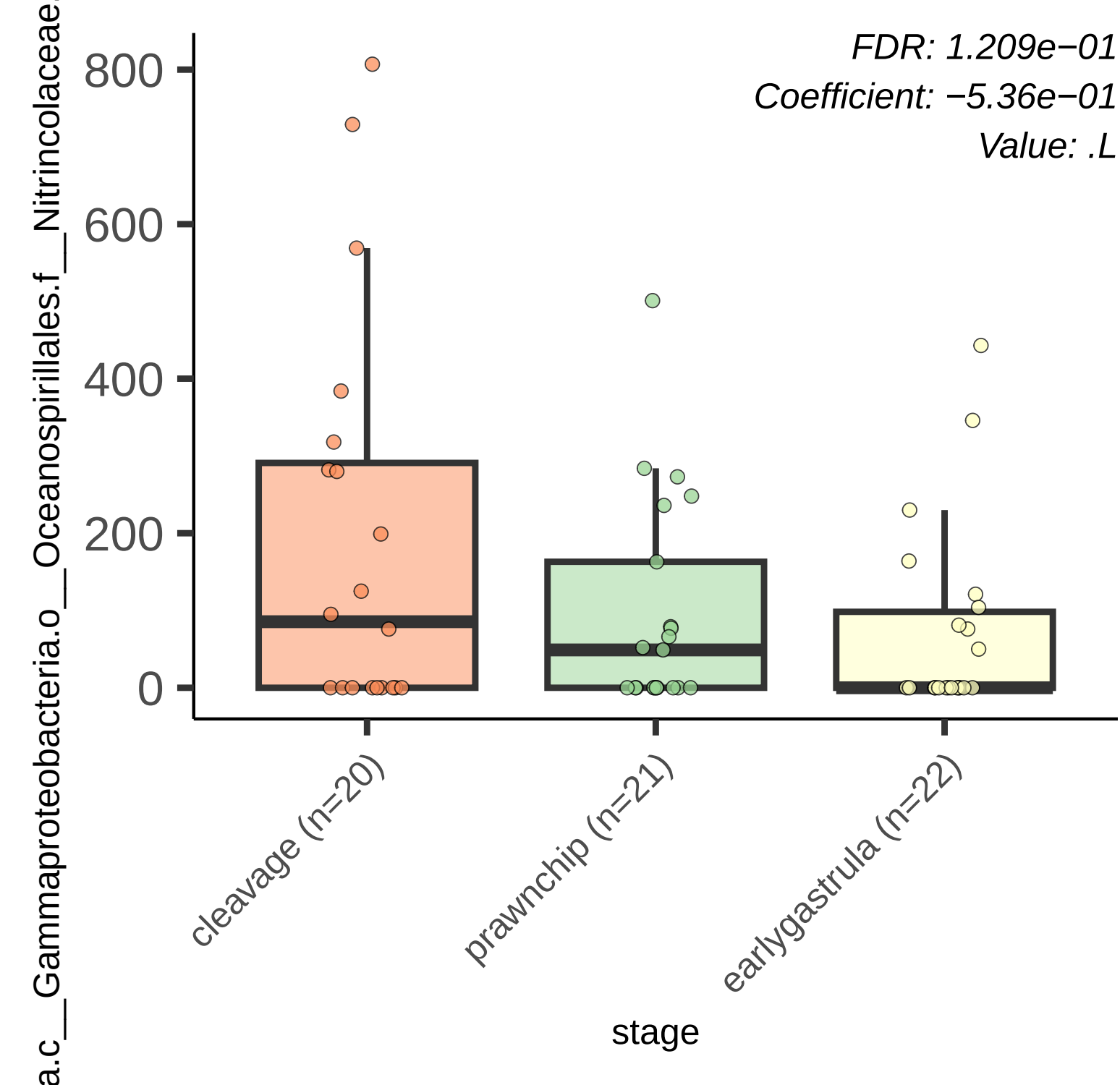
400
300
200
100
0
cleavage (n=20)
prawnchip (n=21)
earlygastrula (n=22)
stage
__Proteobacteria.c__Alphaproteobacteria.o__Rhodobacterales.f__R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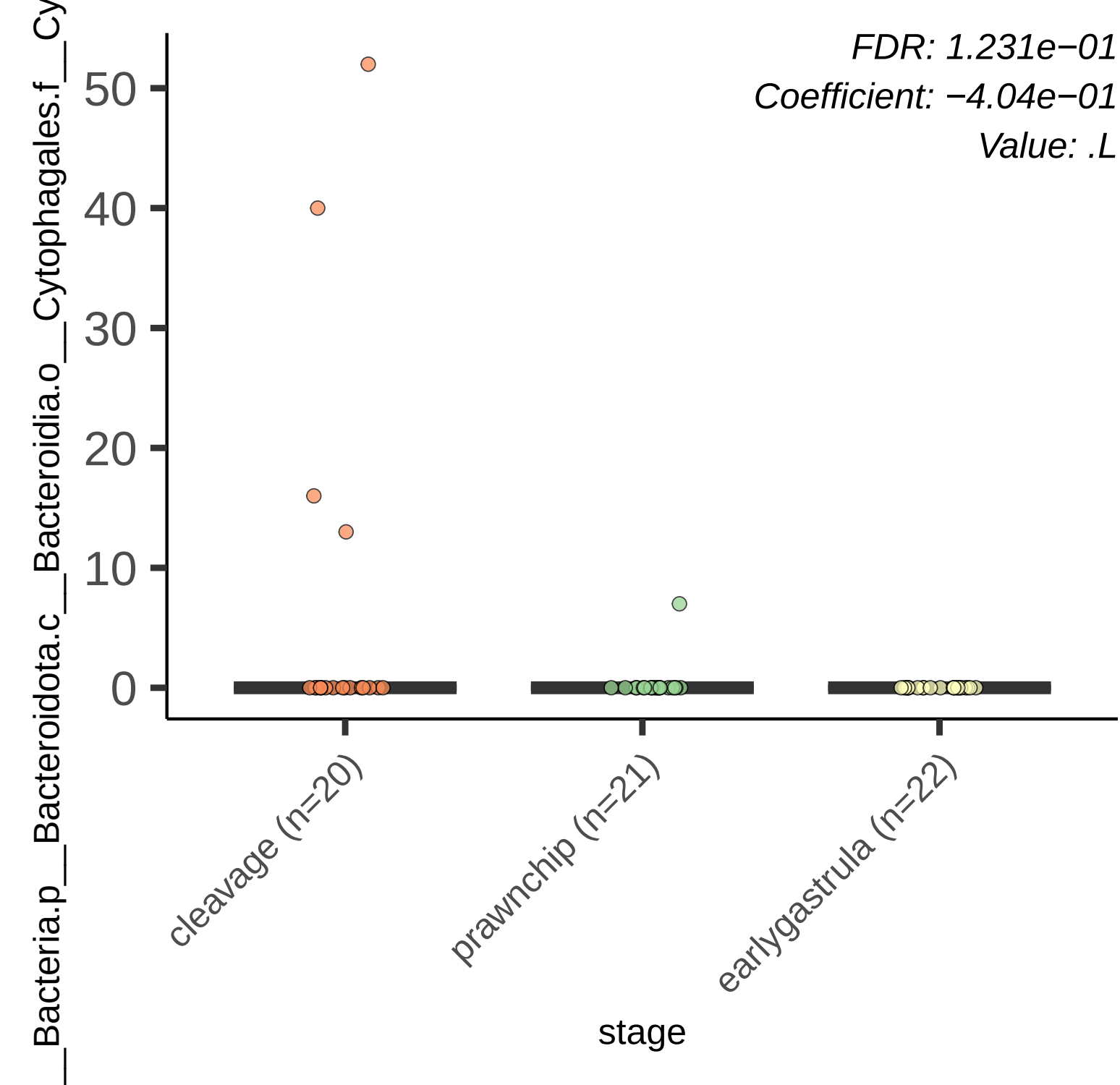
Coefficient: −5.36e−01
Value: .L
800
600
400
200
0
cleavage (n=20)
prawnchip (n=21)
earlygastrula (n=22)
stage
a.c__Gammaproteobacteria.o__Oceanospirillales.f__Nitrincolaceae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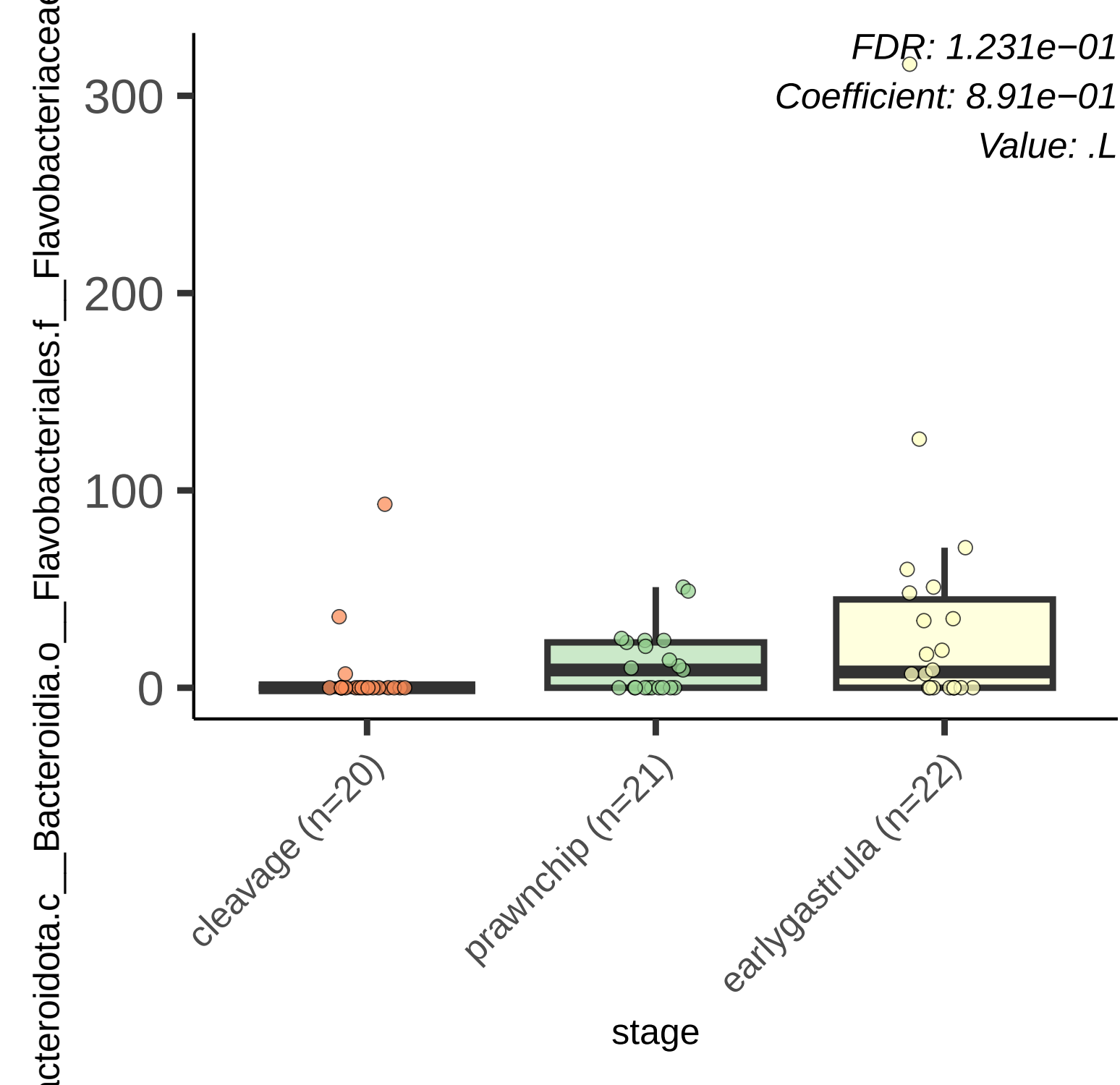
40
30
20
10
0
cleavage (n=20)
prawnchip (n=21)
earlygastrula (n=22)
stage
__Bacteria.p__Bacteroidota.c__Bacteroidia.o__Cytophagales.f__Cy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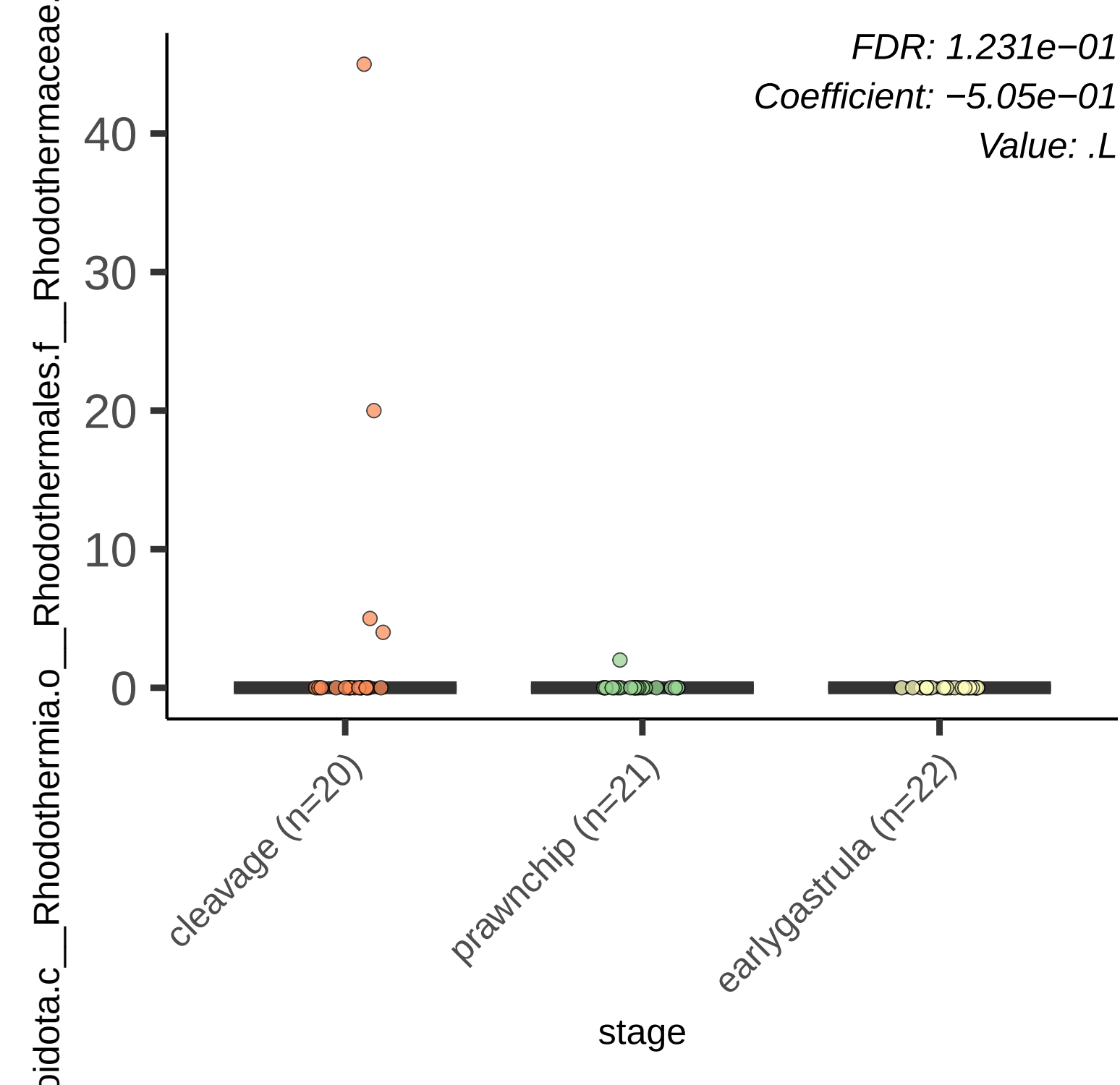
300
200
100
0
cleavage (n=20)
prawnchip (n=21)
earlygastrula (n=22)
stage
cteroidota.c__Bacteroidia.o__Flavobacteriales.f__Flavobacteriacea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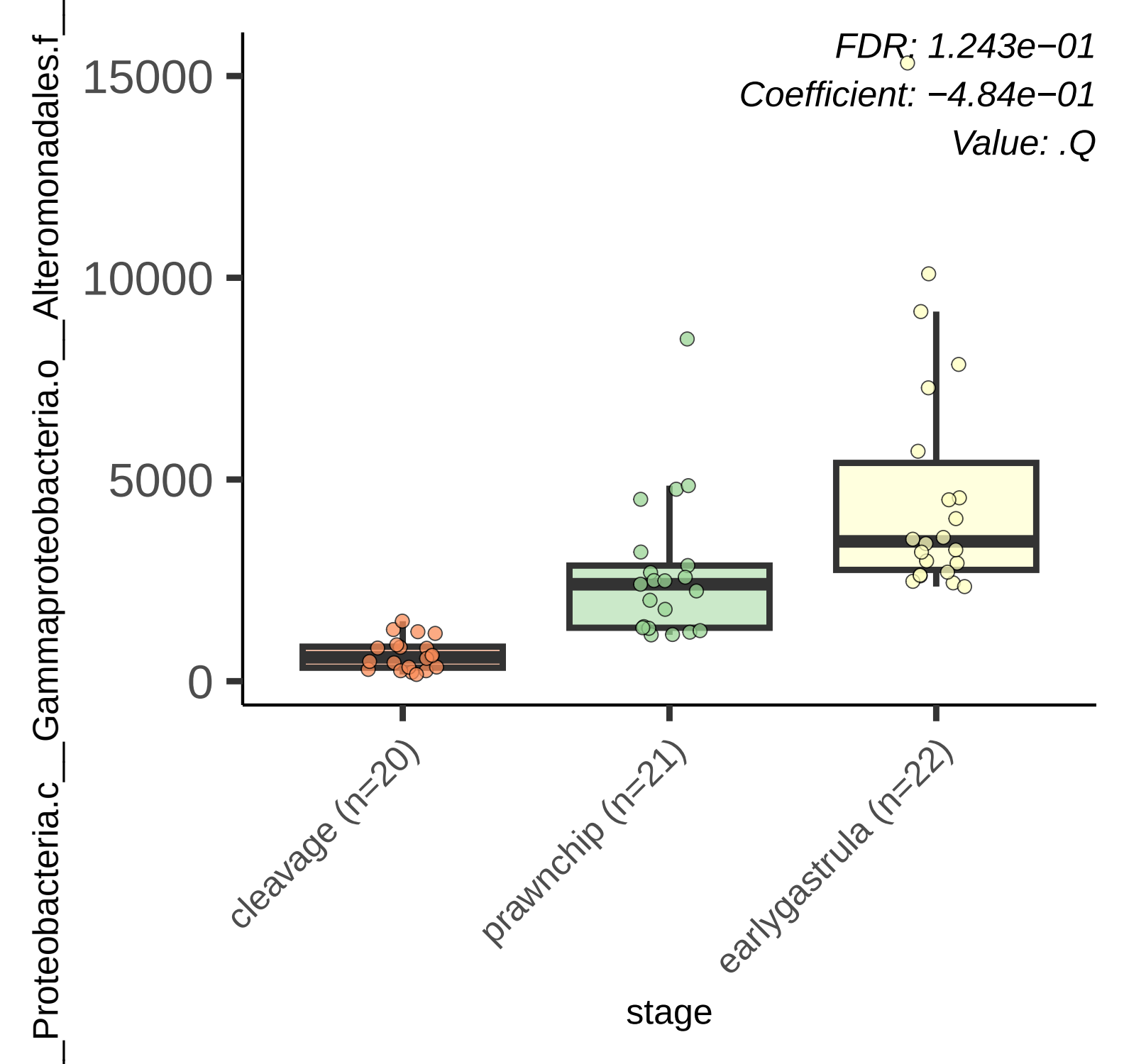
30
20
10
0
cleavage (n=20)
prawnchip (n=21)
earlygastrula (n=22)
stage
poidota.c__Rhodothermia.o__Rhodothermales.f__Rhodothermaceae.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10000
5000
0
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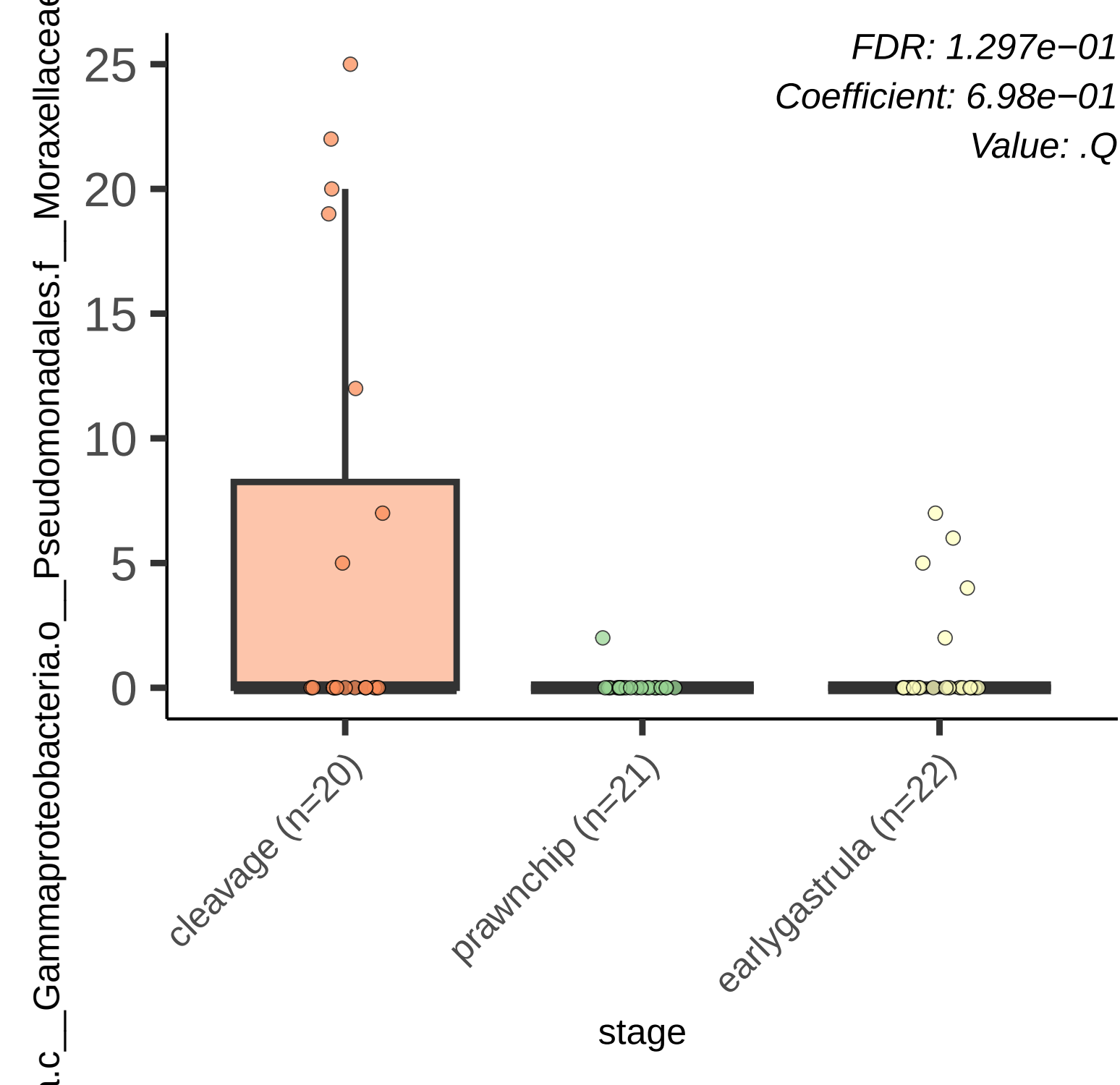
prawnchip (n=21)
earlygastrula (n=22)
stage
__Proteobacteria.c__Gammaproteobacteria.o__Alteromonadales.f__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20
15
10
5
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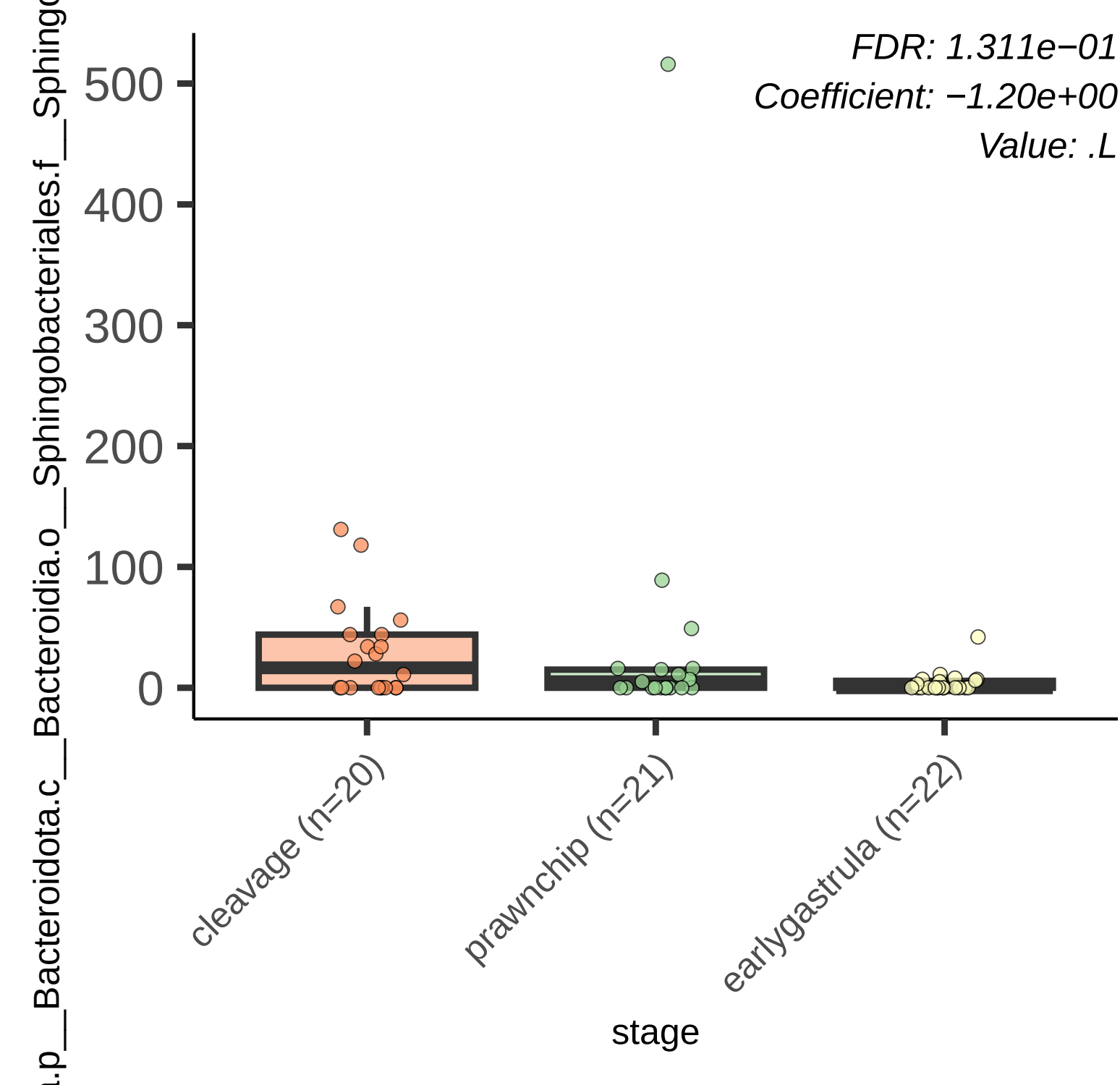
prawnchip (n=21)
earlygastrula (n=22)
stage
c__Gammaproteobacteria.o__Pseudomonadales.f__Moraxellaceae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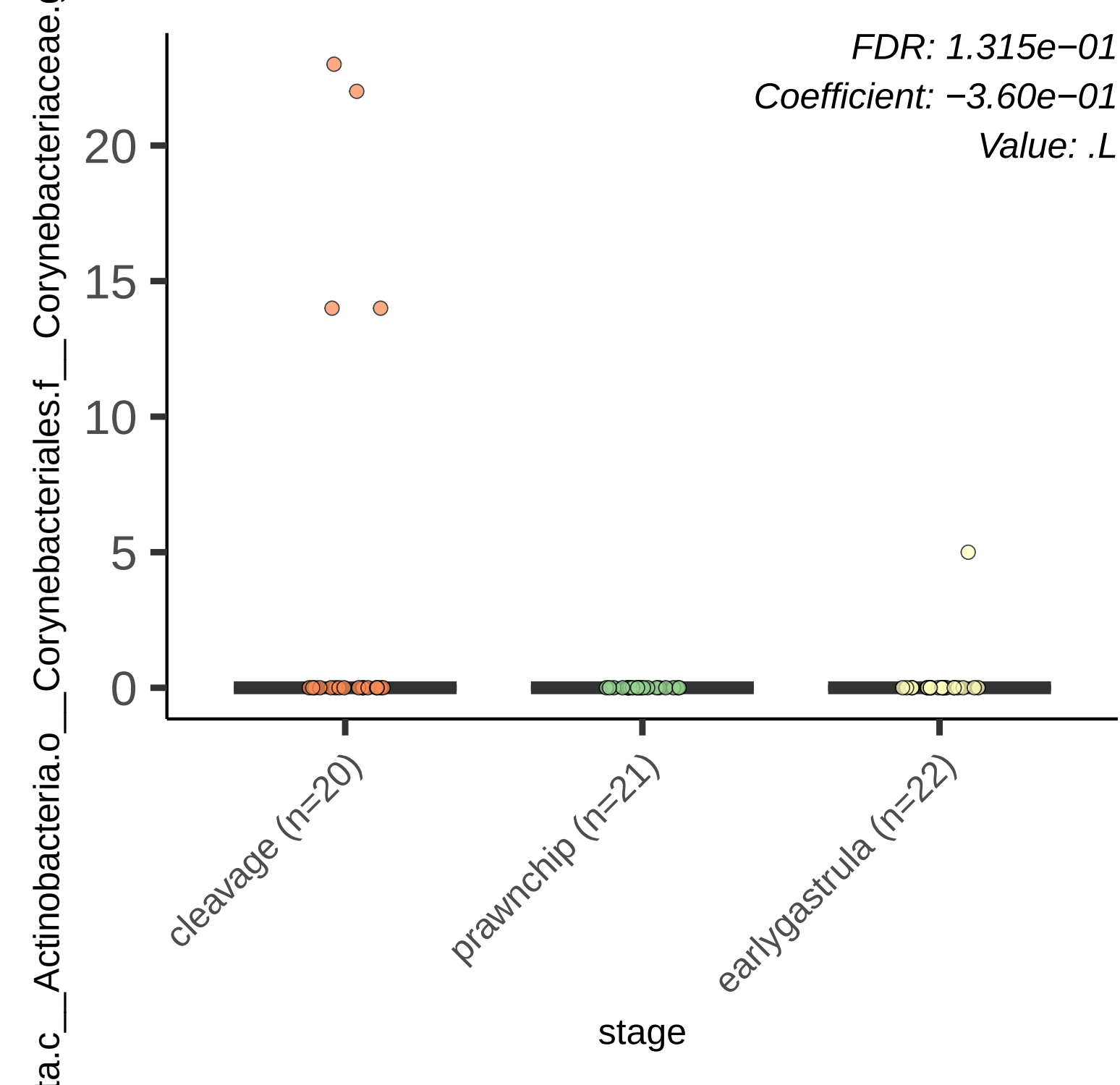
400
300
200
100
0
cleavage (n=20)
prawnchip (n=21)
earlygastrula (n=22)
stage
k__Bacteria.p__Bacteroidota.c__Bacteroidia.o__Sphingobacteriales.f__Sphingo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15
10
5
0
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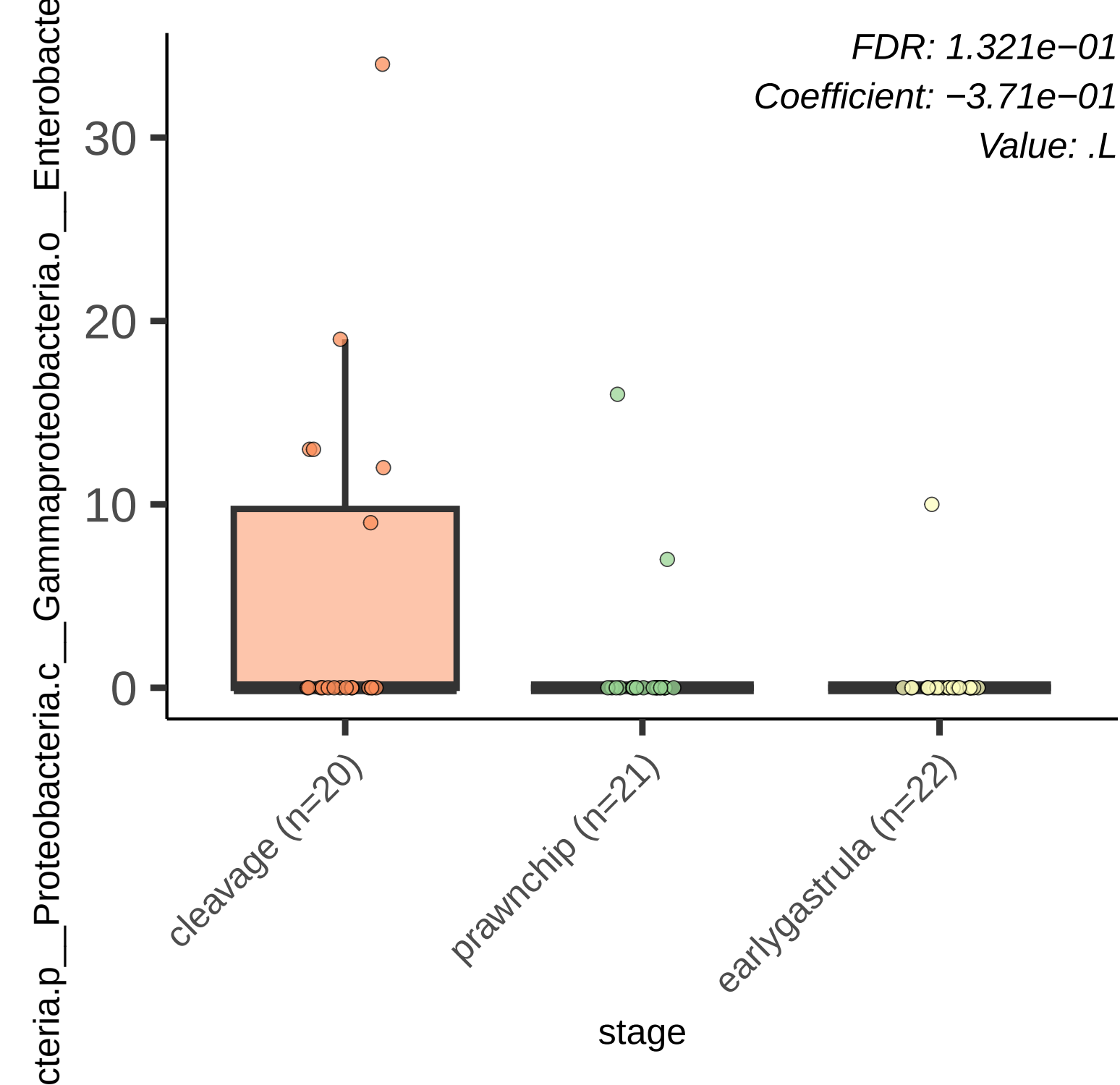
prawnchip (n=21)
earlygastrula (n=22)
stage
ria.c__Actinobacteria.o__Corynebacteriales.f__Corynebacteriaceae.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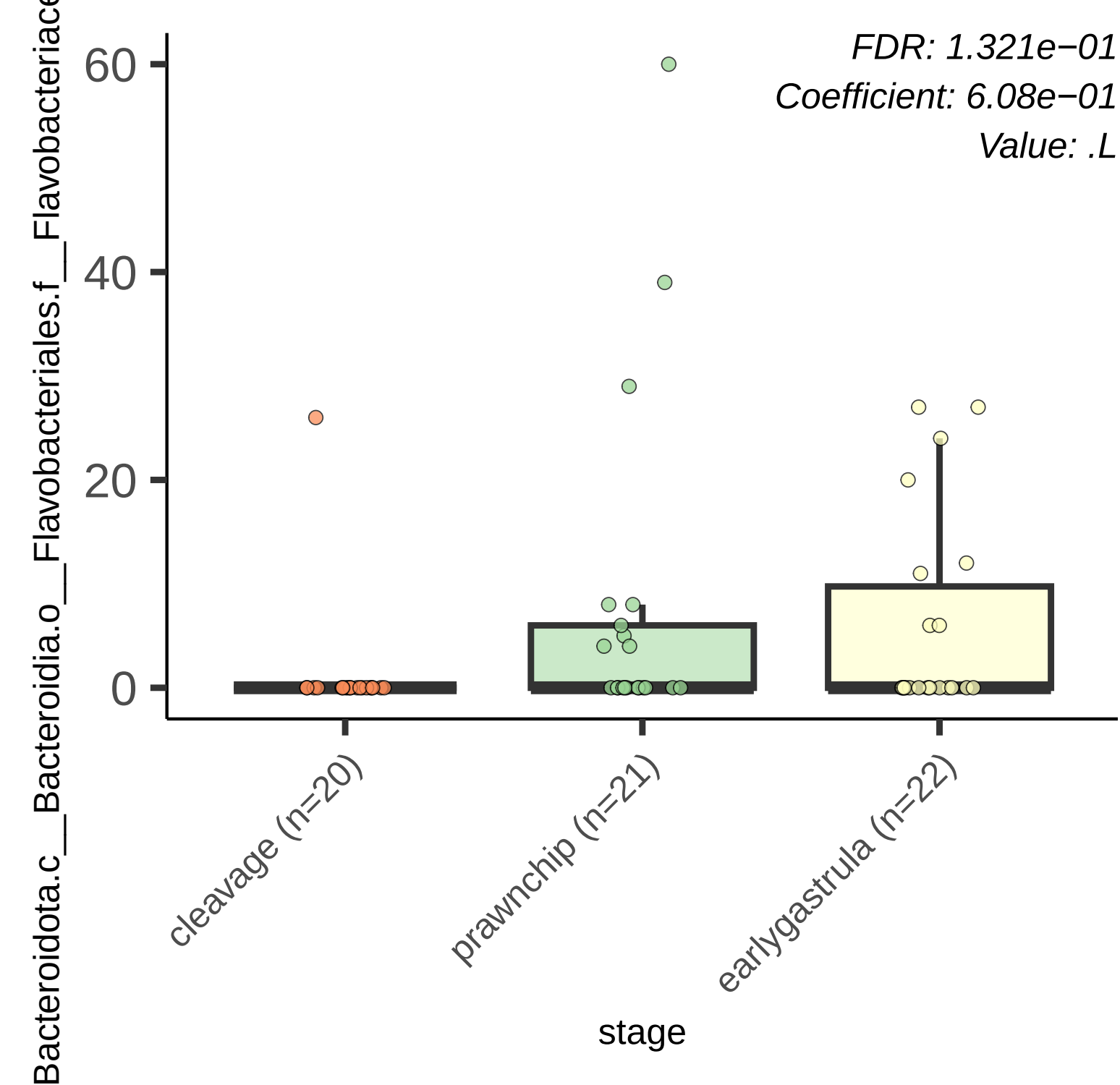
30
20
10
0
cleavage (n=20)
prawnchip (n=21)
earlygastrula (n=22)
stage
cteria.p__Proteobacteria.c__Gammaproteobacteria.o__Enterobacte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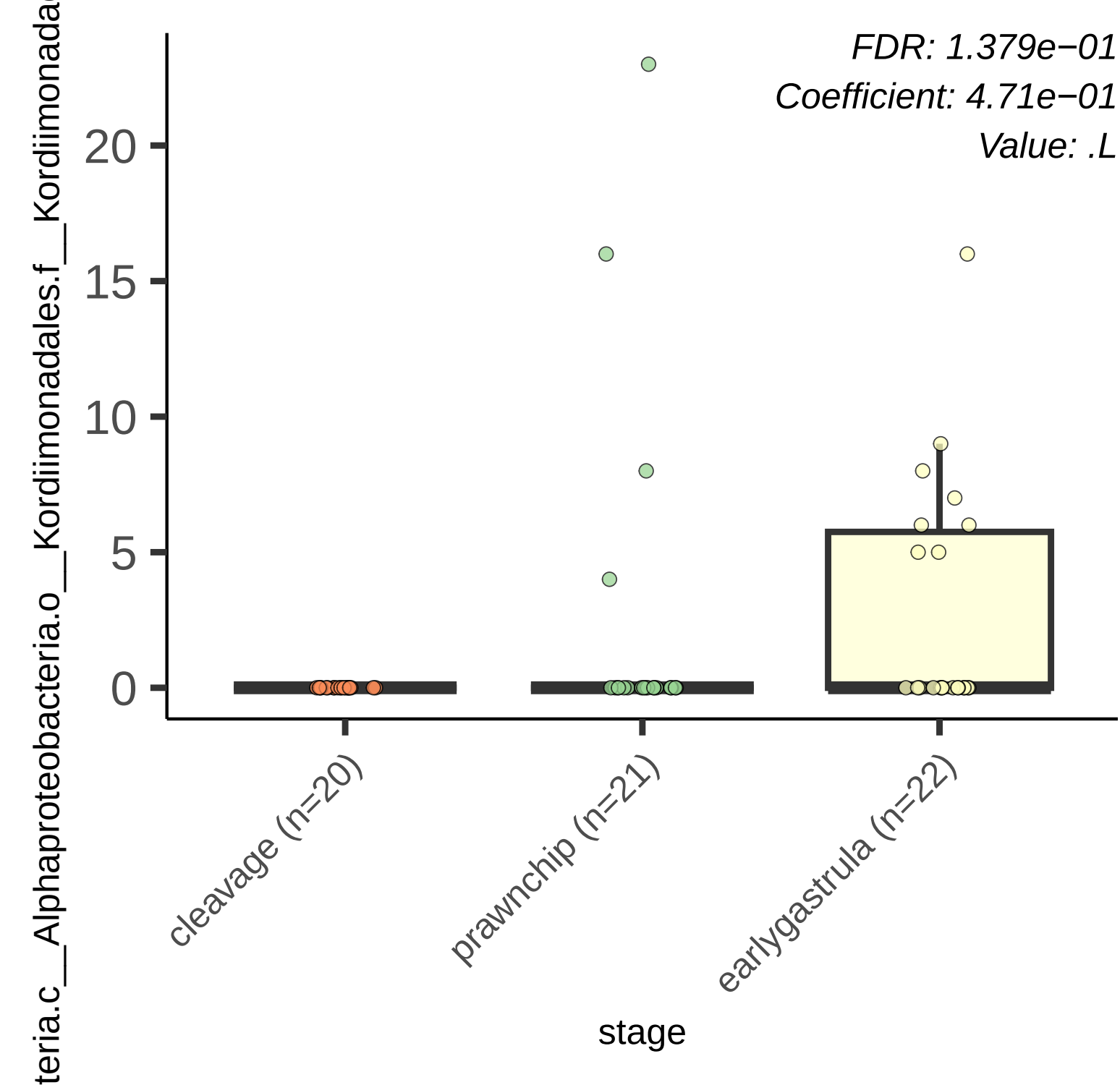
40
20
0
cleavage (n=20)
prawnchip (n=21)
earlygastrula (n=22)
stage
Bacteroidota.c__Bacteroidia.o__Flavobacteriales.f__Flavobacteriace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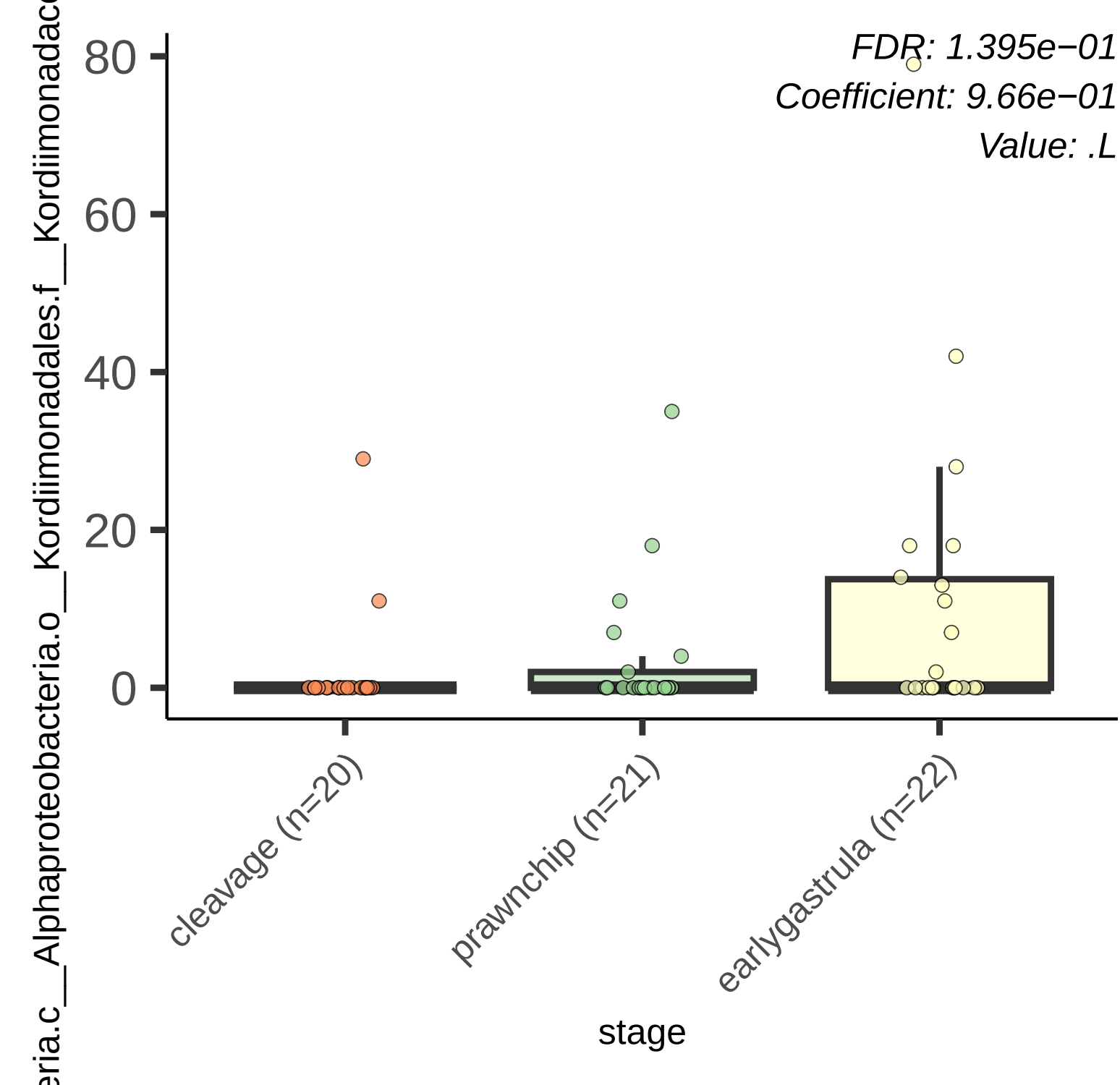
15
10
5
0
cleavage (n=20)
prawnchip (n=21)
earlygastrula (n=22)
stage
teria.c__Alphaproteobacteria.o__Kordiimonadales.f__Kordiimonada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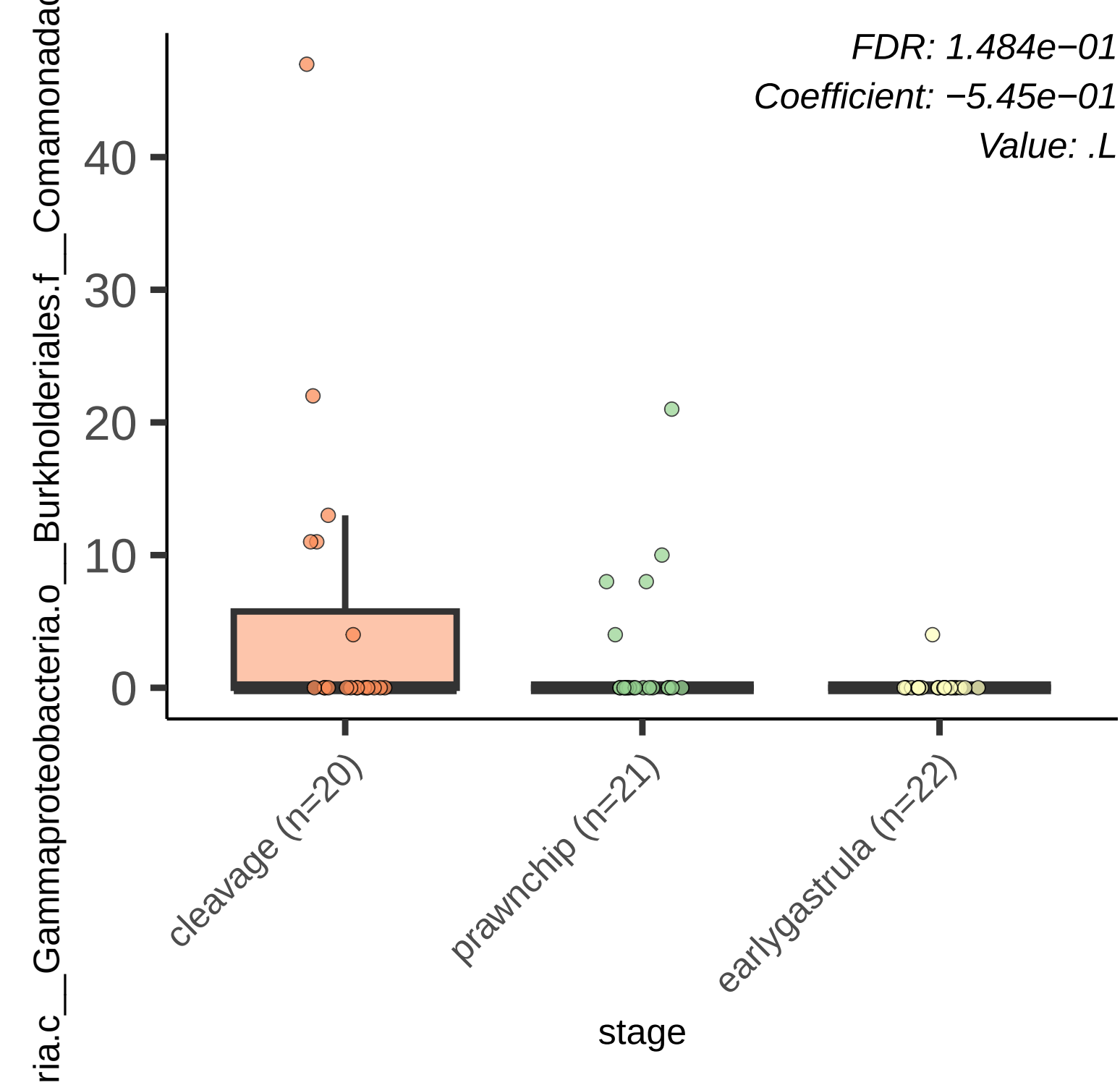
60
40
20
0
cleavage (n=20)
prawnchip (n=21)
earlygastrula (n=22)
stage
eria.c__Alphaproteobacteria.o__Kordiimonadales.f__Kordiimonadac

FDR: 1.478e−01
Coefficient: −7.60e−01
Value: .L
2000
1000
0
cleavage (n=20)
prawnchip (n=21)
earlygastrula (n=22)
stage
teria.c__Gammaproteobacteria.o__Oceanospirillales.f__Nitrincolace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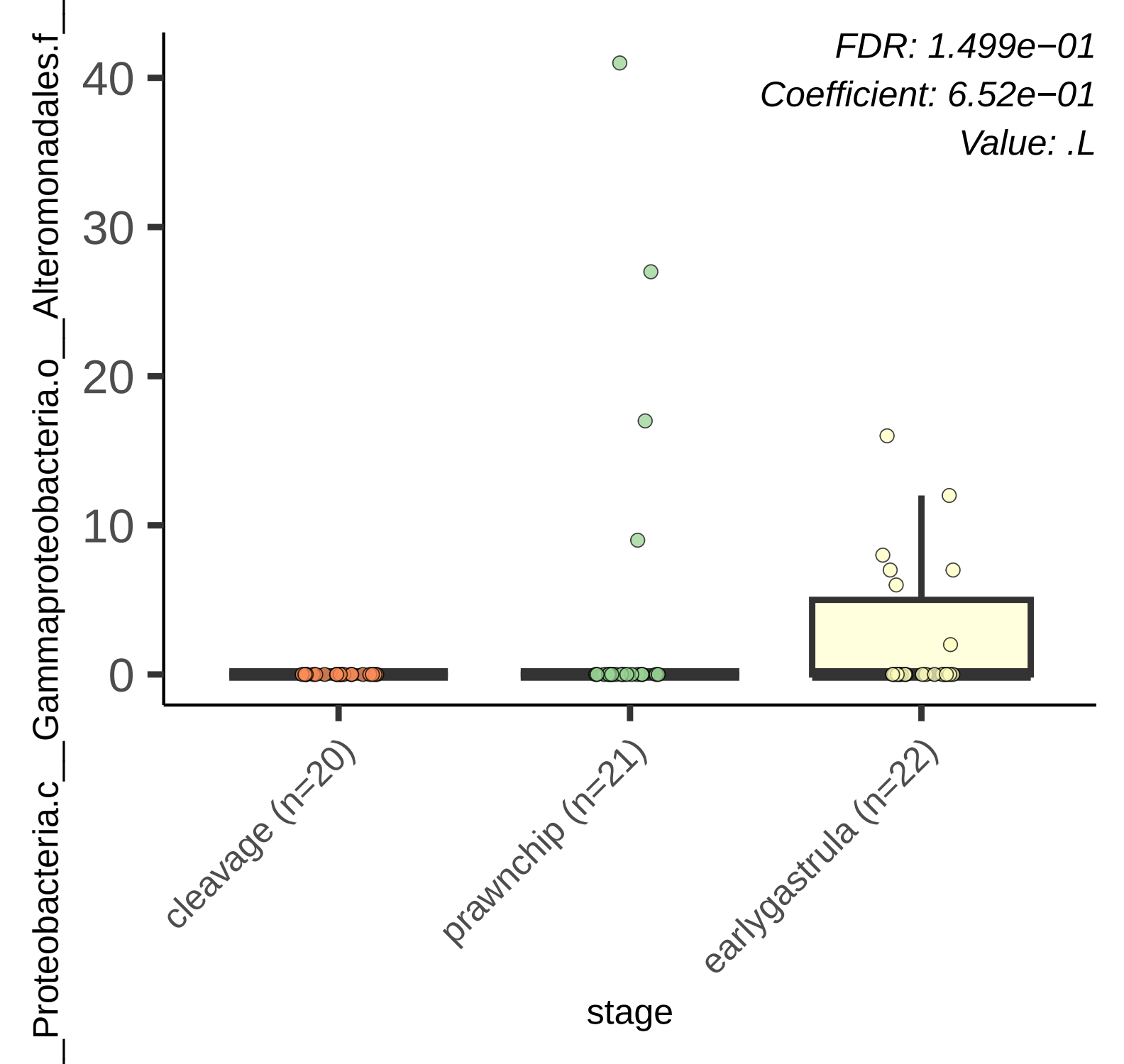
30
20
10
0
cleavage (n=20)
prawnchip (n=21)
earlygastrula (n=22)
stage
ria.c__Gammaproteobacteria.o__Burkholderiales.f__Comamonada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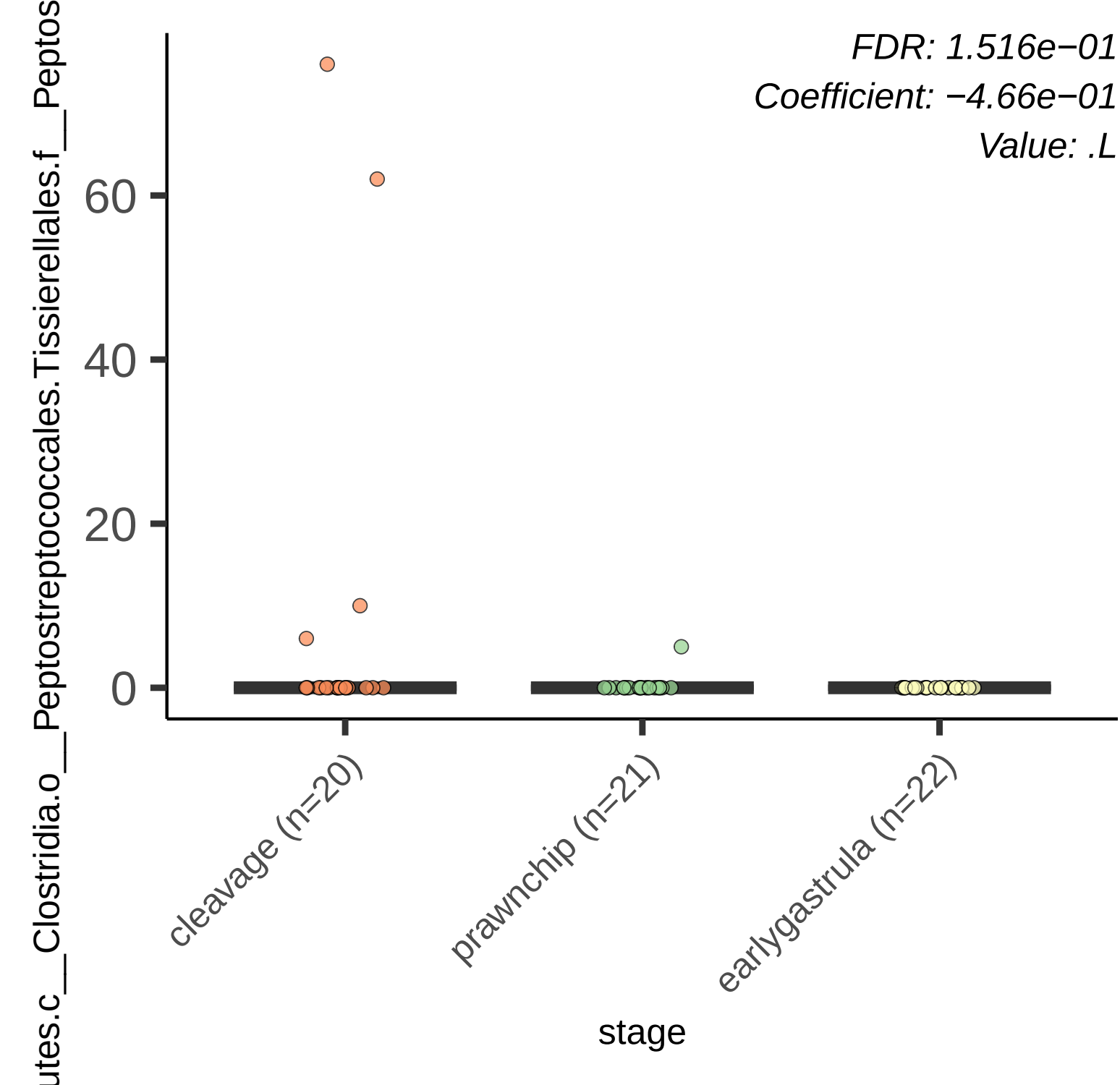
30
20
10
0
cleavage (n=20)
prawnchip (n=21)
earlygastrula (n=22)
stage
__Proteobacteria.c__Gammaproteobacteria.o__Alteromonadales.f__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Coefficient: −4.66e−01
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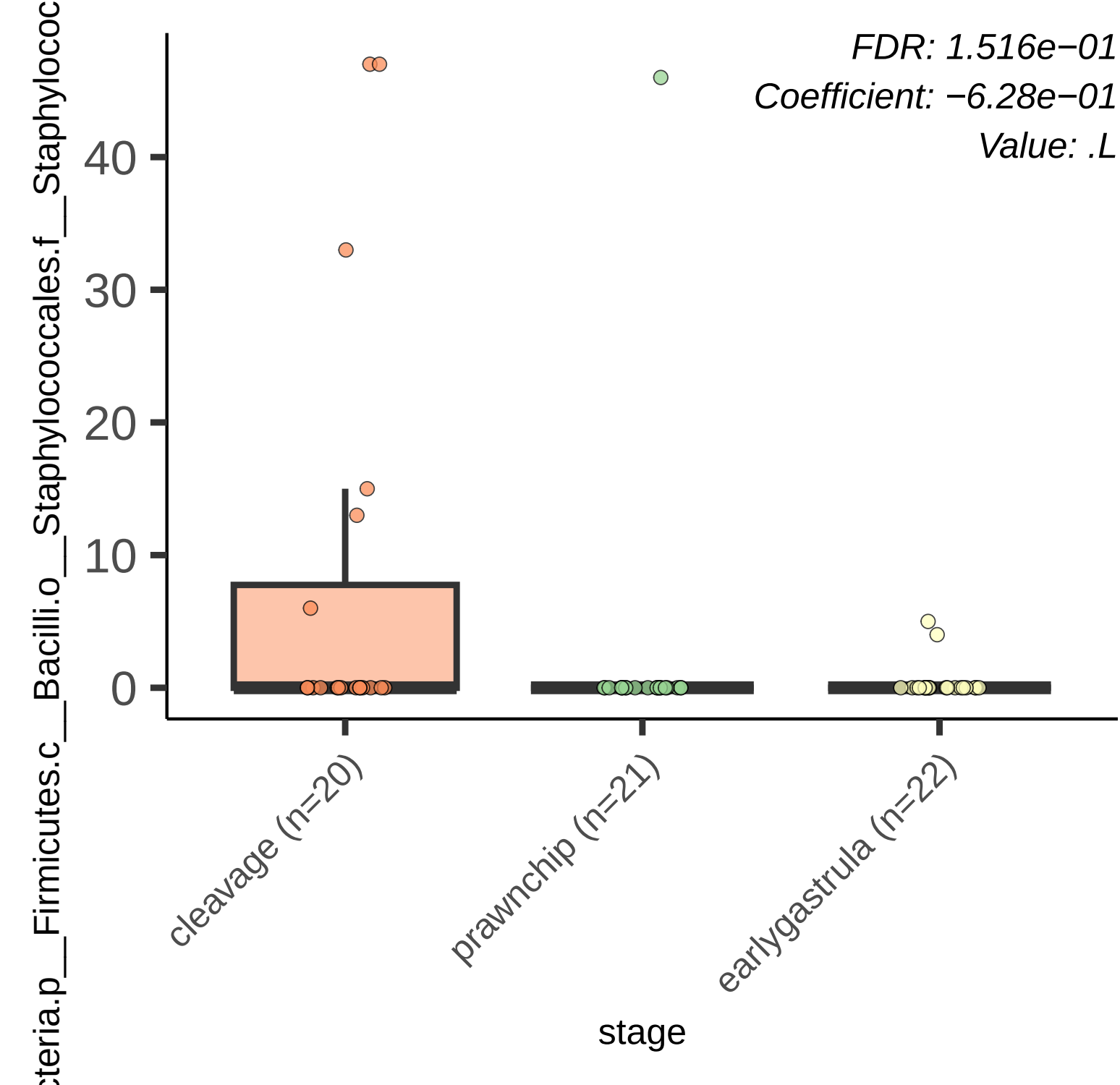
60
40
20
0
cleavage (n=20)
prawnchip (n=21)
earlygastrula (n=22)
stage
utes.c__Clostridia.o__Peptostreptococcales.Tissierellales.f__Peptos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Coefficient: −6.28e−01
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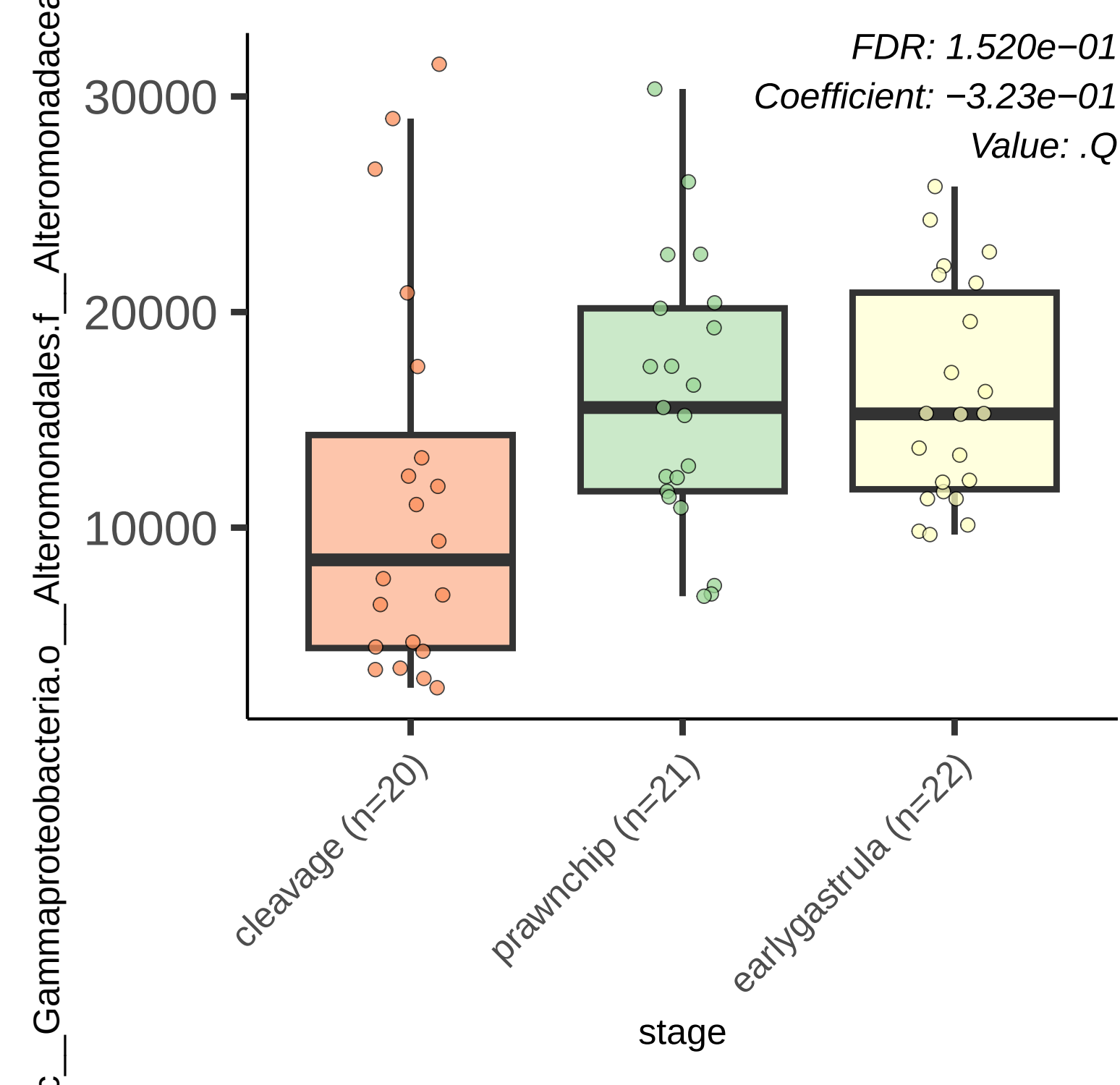
30
20
10
0
cleavage (n=20)
prawnchip (n=21)
earlygastrula (n=22)
stage
cteria.p__Firmicutes.c__Bacilli.o__Staphylococcales.f__Staphyloco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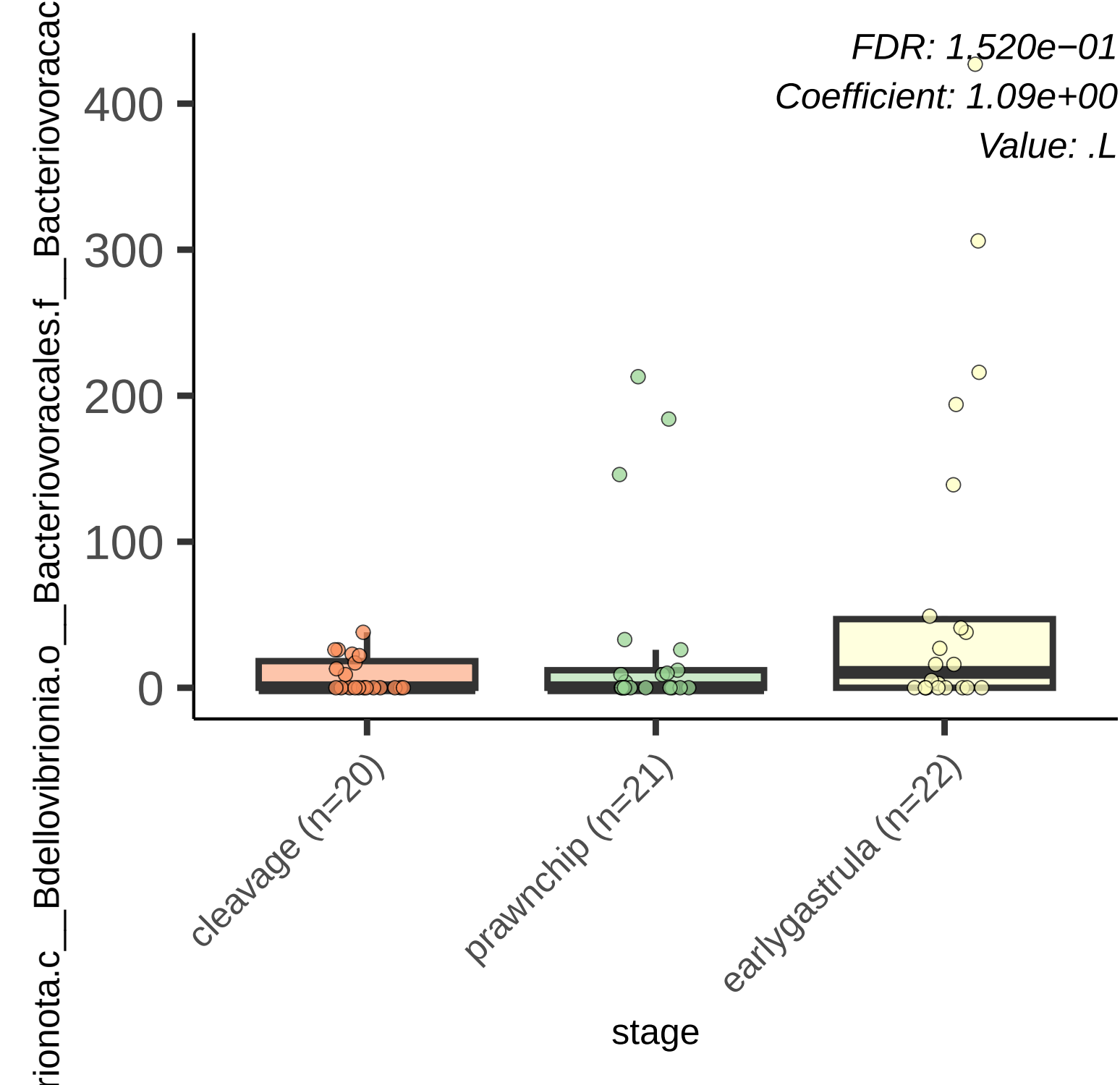
Coefficient: −3.23e−01
Value: .Q
30000
20000
10000
cleavage (n=20)
prawnchip (n=21)
earlygastrula (n=22)
stage
c__Gammaproteobacteria.o__Alteromonadales.f__Alteromonadacea

FDR: 1.520e−01
Coefficient: −1.34e+00
Value: .L
1000
750
500
250
0
cleavage (n=20)
prawnchip (n=21)
earlygastrula (n=22)
stage
a.p__Actinobacteriota.c__Actinobacteria.o__Micrococcales.f__Micro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Coefficient: 1.09e+00
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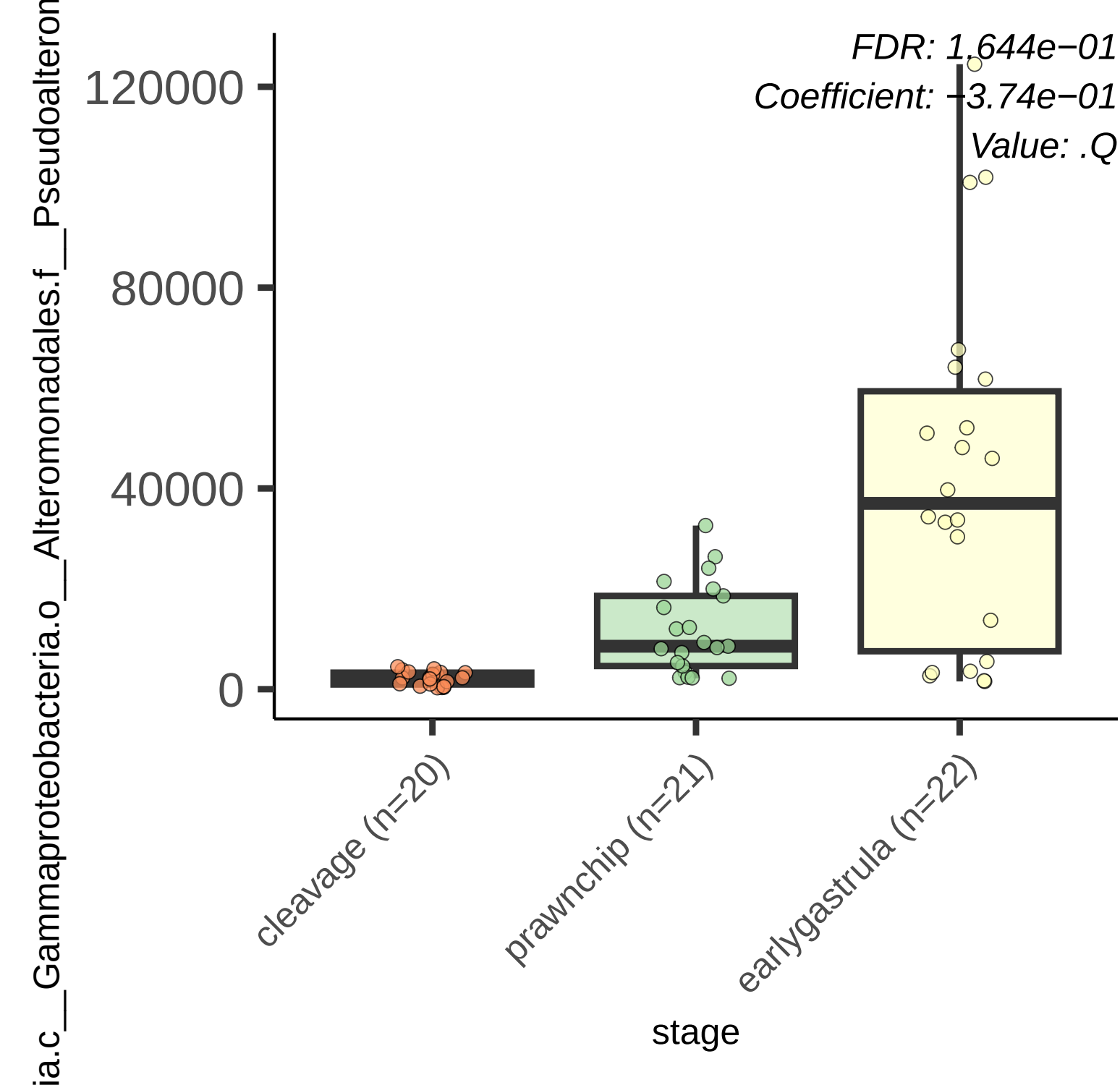
400
300
200
100
0
cleavage (n=20)
prawnchip (n=21)
earlygastrula (n=22)
stage
rionota.c__Bdellovibrionia.o__Bacteriovoracales.f__Bacteriovoracac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80000
40000
0
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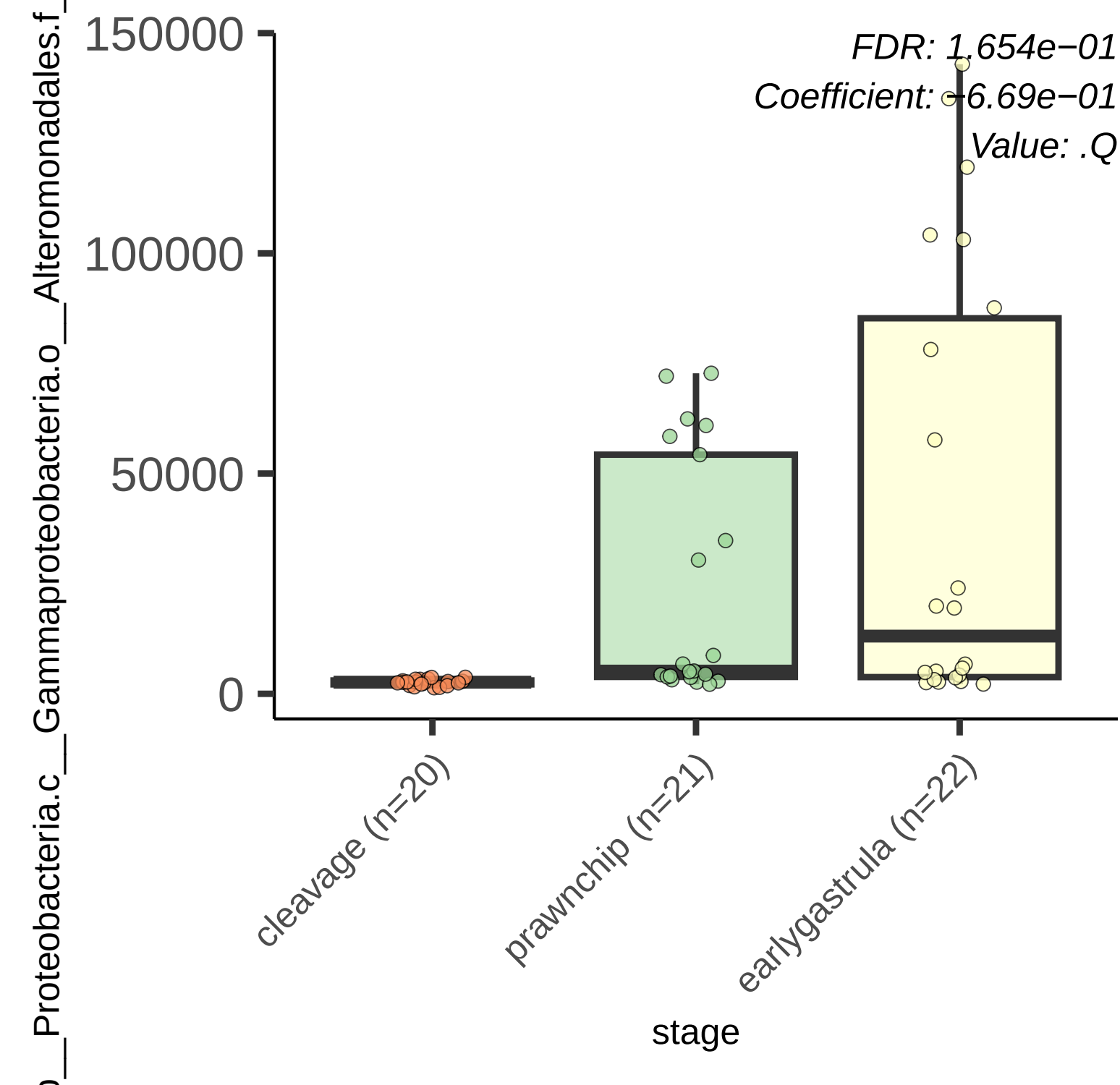
prawnchip (n=21)
earlygastrula (n=22)
stage
ia.c__Gammaproteobacteria.o__Alteromonadales.f__Pseudoalteron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150000
100000
50000
0
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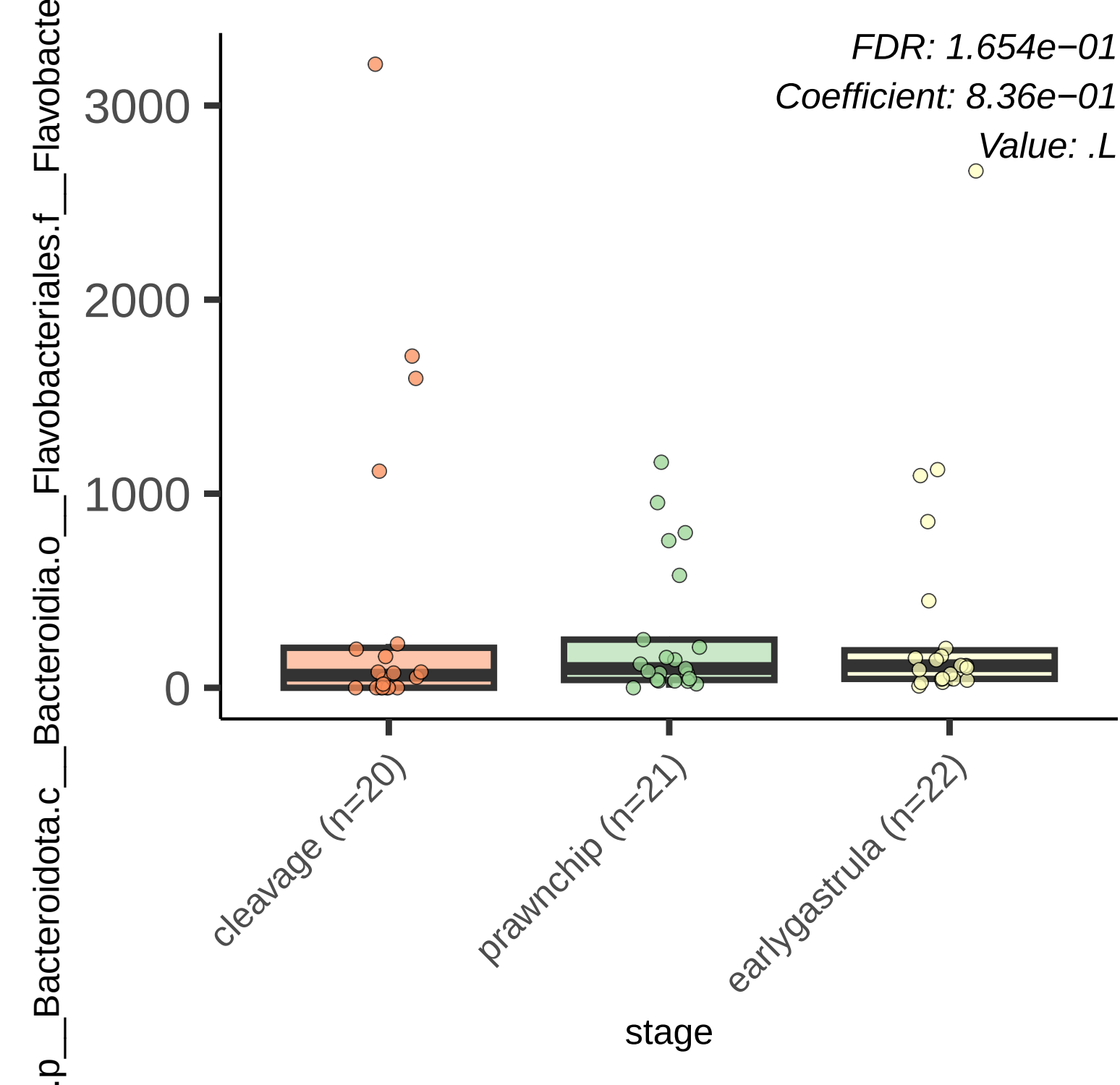
prawnchip (n=21)
earlygastrula (n=22)
stage
o__Proteobacteria.c__Gammaproteobacteria.o__Alteromonadales.f__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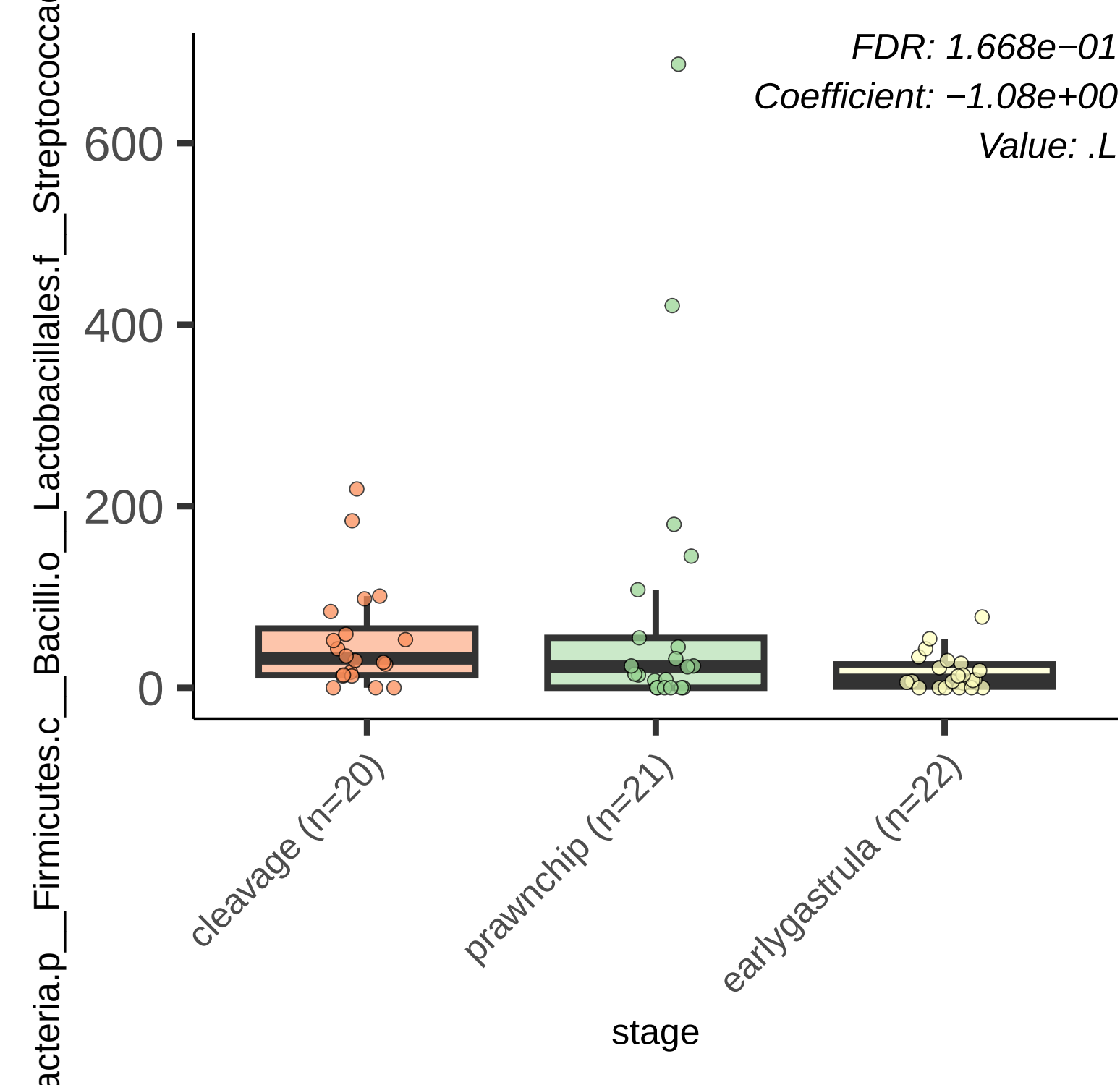
3000
2000
1000
0
cleavage (n=20)
prawnchip (n=21)
earlygastrula (n=22)
stage
.p__Bacteroidota.c__Bacteroidia.o__Flavobacteriales.f__Flavobacte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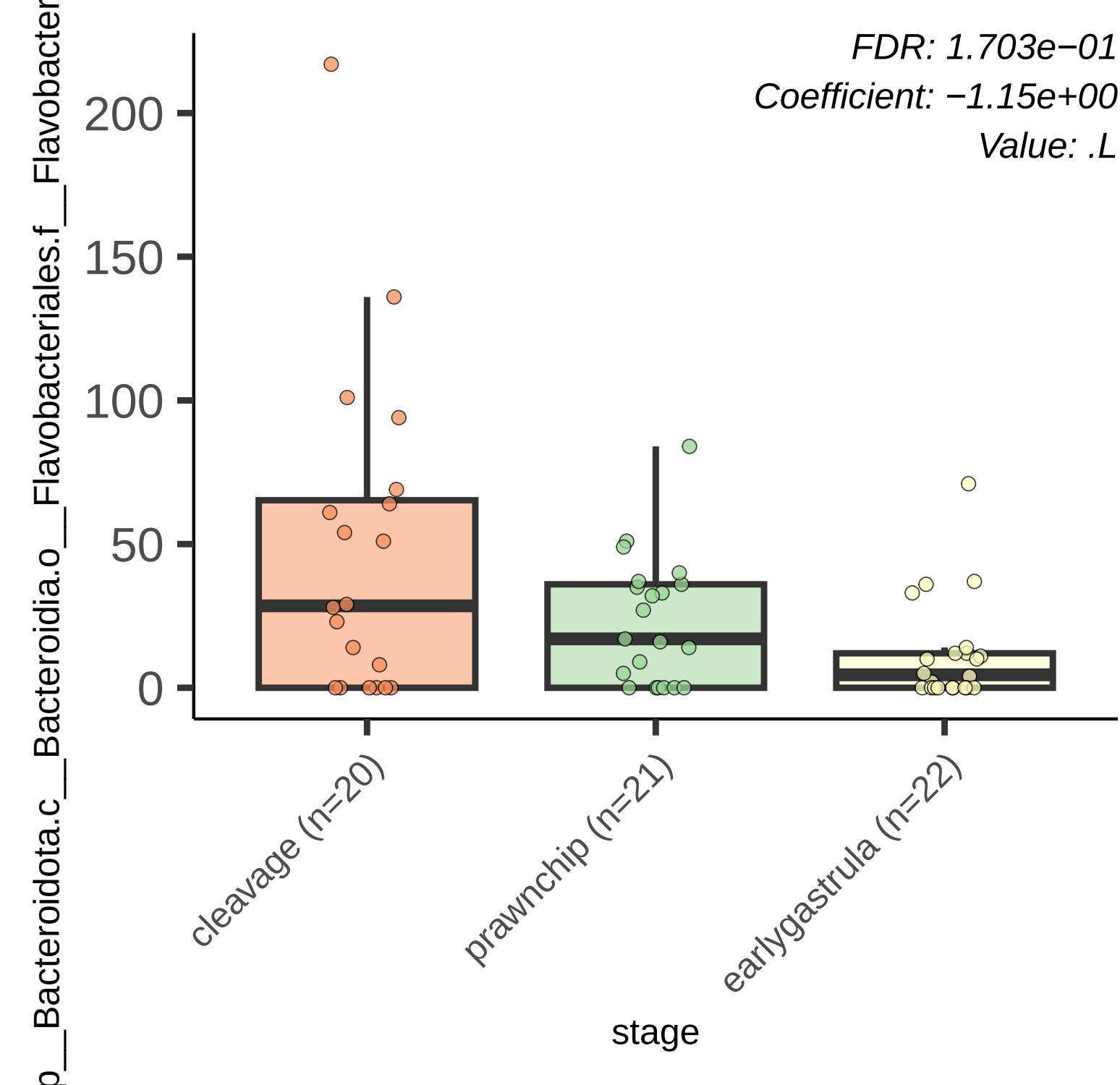
600
400
200
0
cleavage (n=20)
prawnchip (n=21)
earlygastrula (n=22)
stage
acteria.p__Firmicutes.c__Bacilli.o__Lactobacillales.f__Streptococca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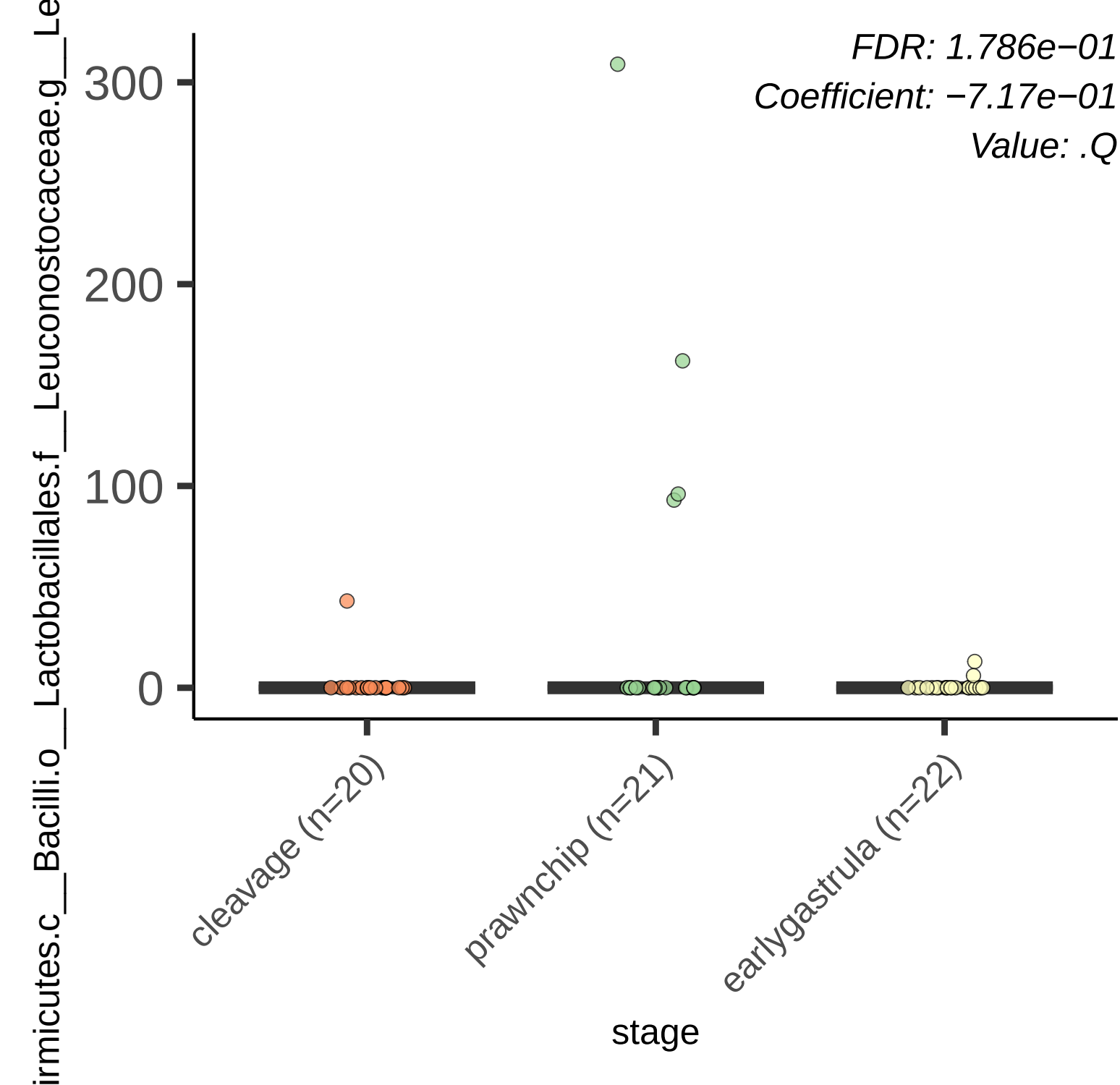
Coefficient: −1.15e+00
Value: .L
200
150
100
50
0
cleavage (n=20)
prawnchip (n=21)
earlygastrula (n=22)
stage
p__Bacteroidota.c__Bacteroidia.o__Flavobacteriales.f__Flavobacter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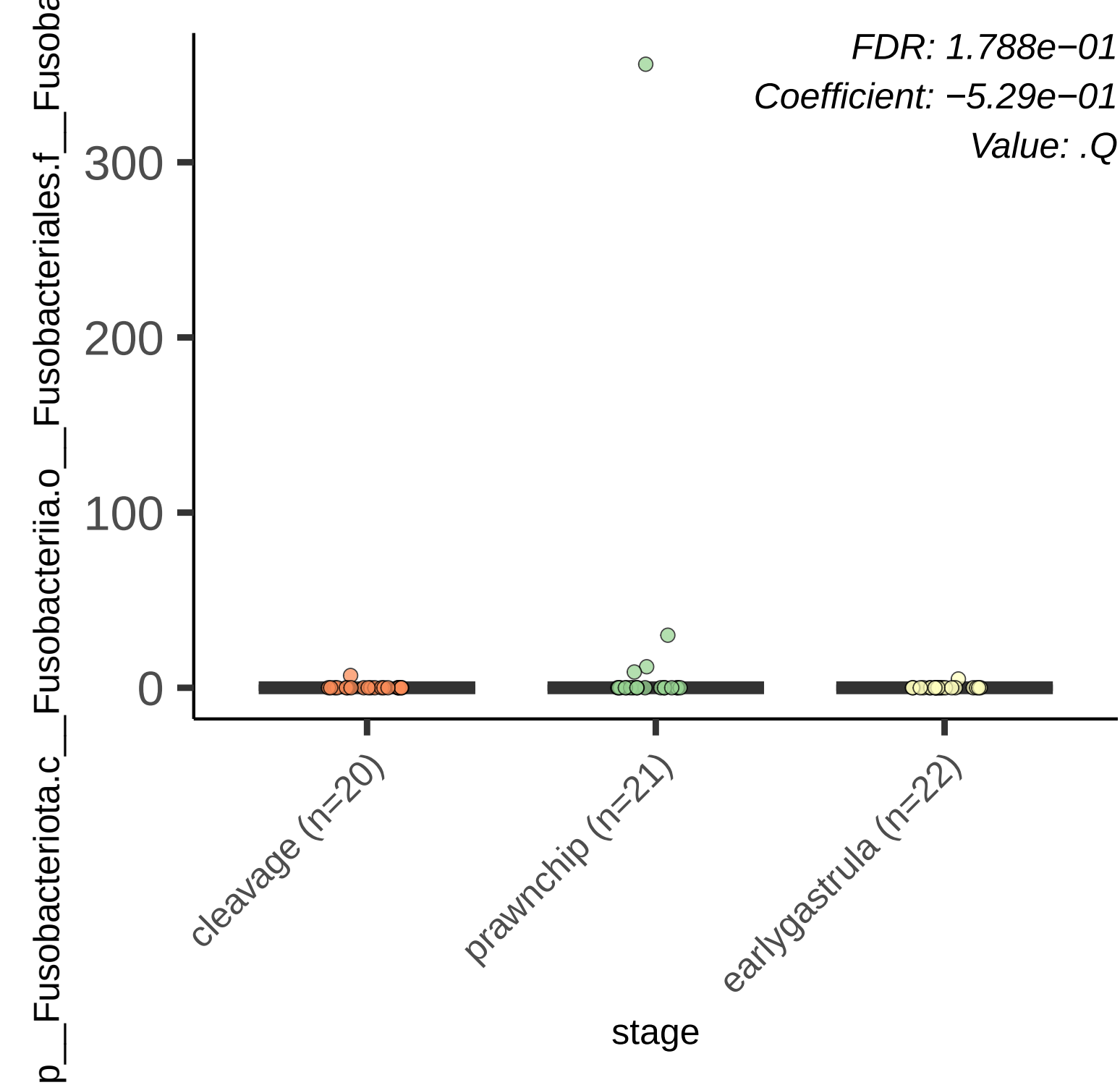
300
200
100
0
cleavage (n=20)
prawnchip (n=21)
earlygastrula (n=22)
stage
Firmicutes.c__Bacilli.o__Lactobacillales.f__Leuconostocaceae.g__Le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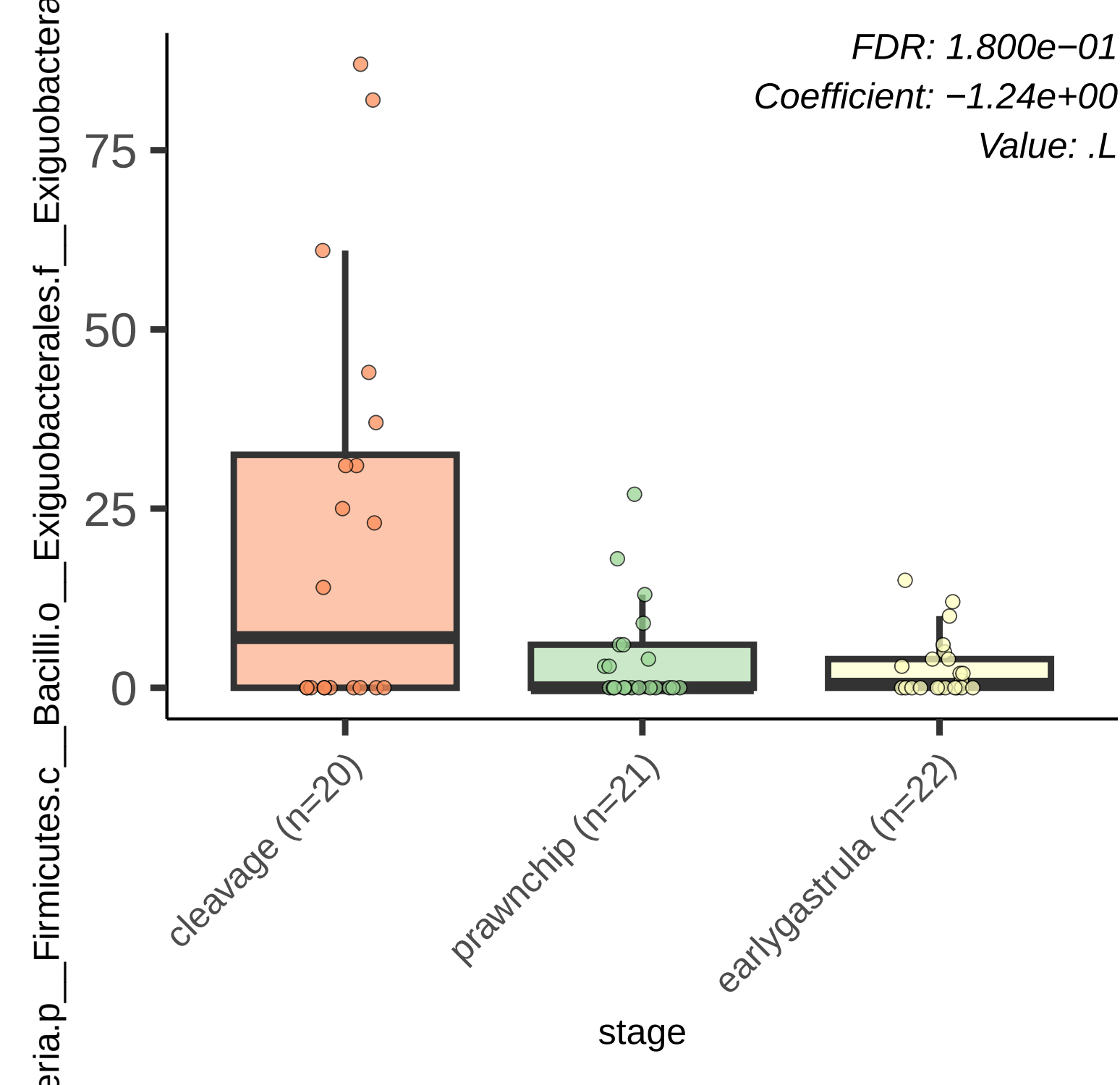
Coefficient: −5.29e−01
Value: .Q
300
200
100
0
cleavage (n=20)
prawnchip (n=21)
earlygastrula (n=22)
stage
p__Fusobacteriota.c__Fusobacteriia.o__Fusobacteriales.f__Fusoba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Coefficient: −1.24e+00
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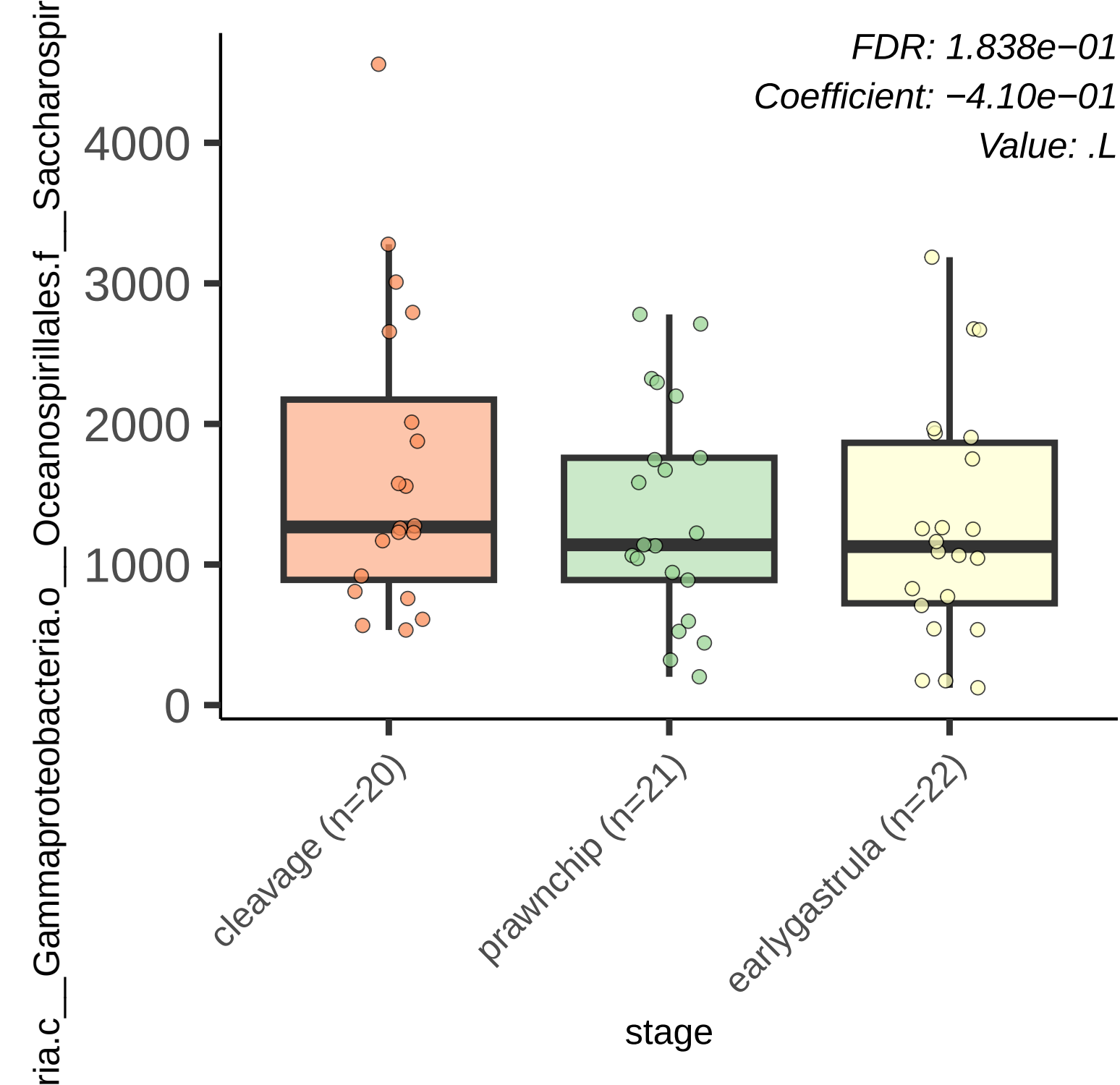
75
50
25
0
cleavage (n=20)
prawnchip (n=21)
earlygastrula (n=22)
stage
eria.p__Firmicutes.c__Bacilli.o__Exiguobacterales.f__Exiguobacte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4000
3000
2000
1000
0
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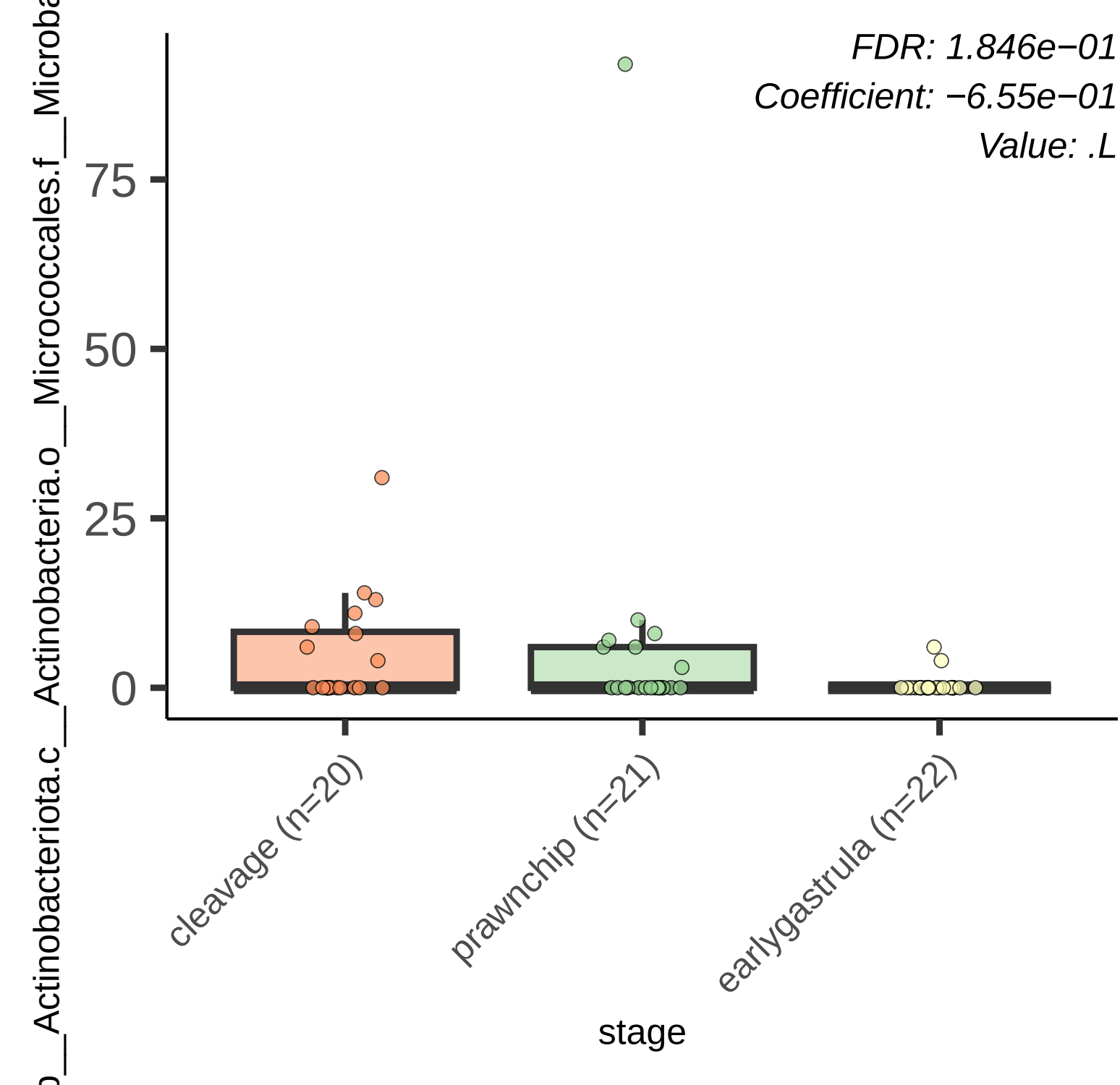
prawnchip (n=21)
earlygastrula (n=22)
stage
ria.c__Gammaproteobacteria.o__Oceanospirillales.f__Saccharospir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Coefficient: −6.55e−01
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50
25
0
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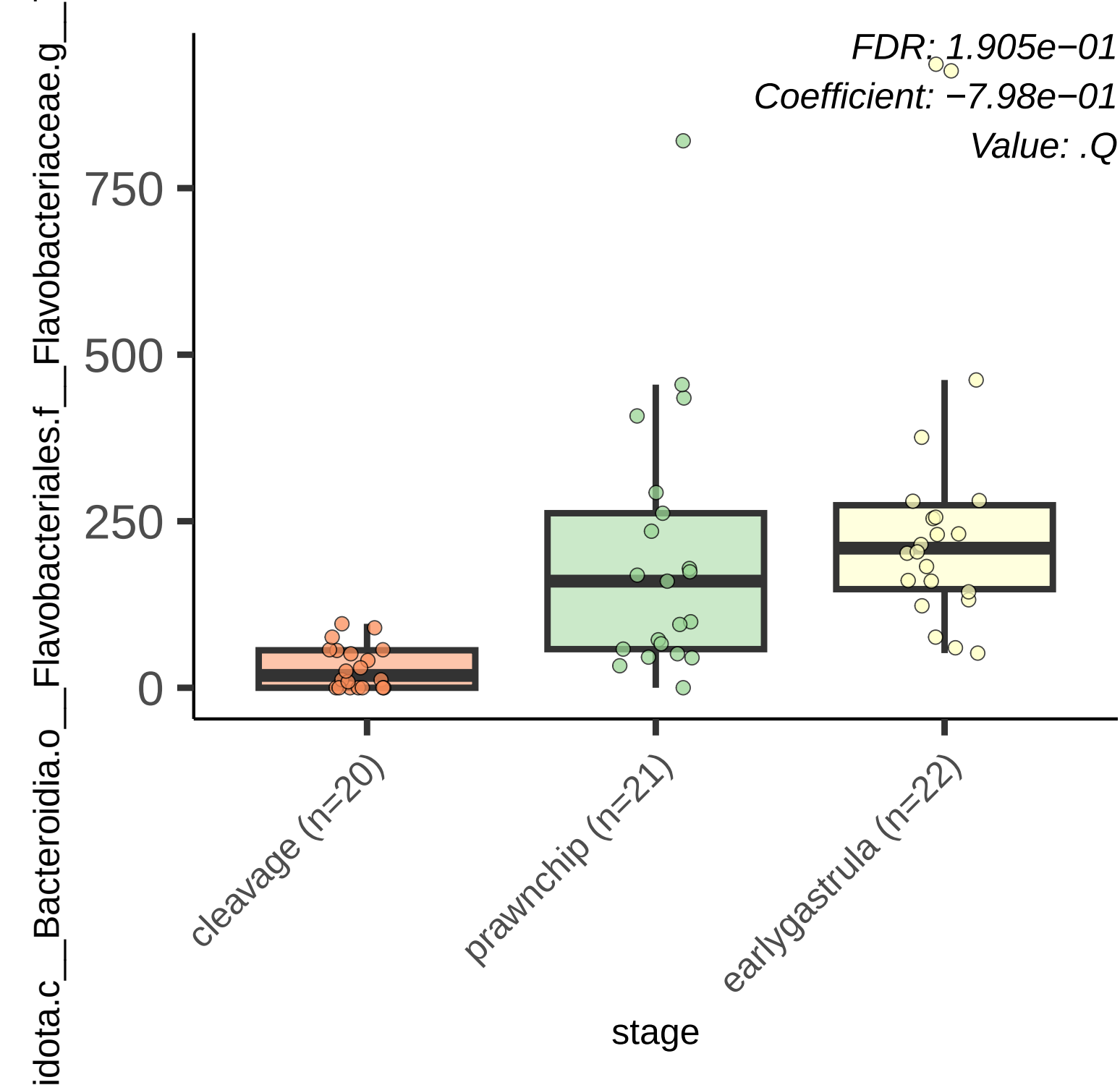
prawnchip (n=21)
earlygastrula (n=22)
stage
p__Actinobacteriota.c__Actinobacteria.o__Micrococcales.f__Microba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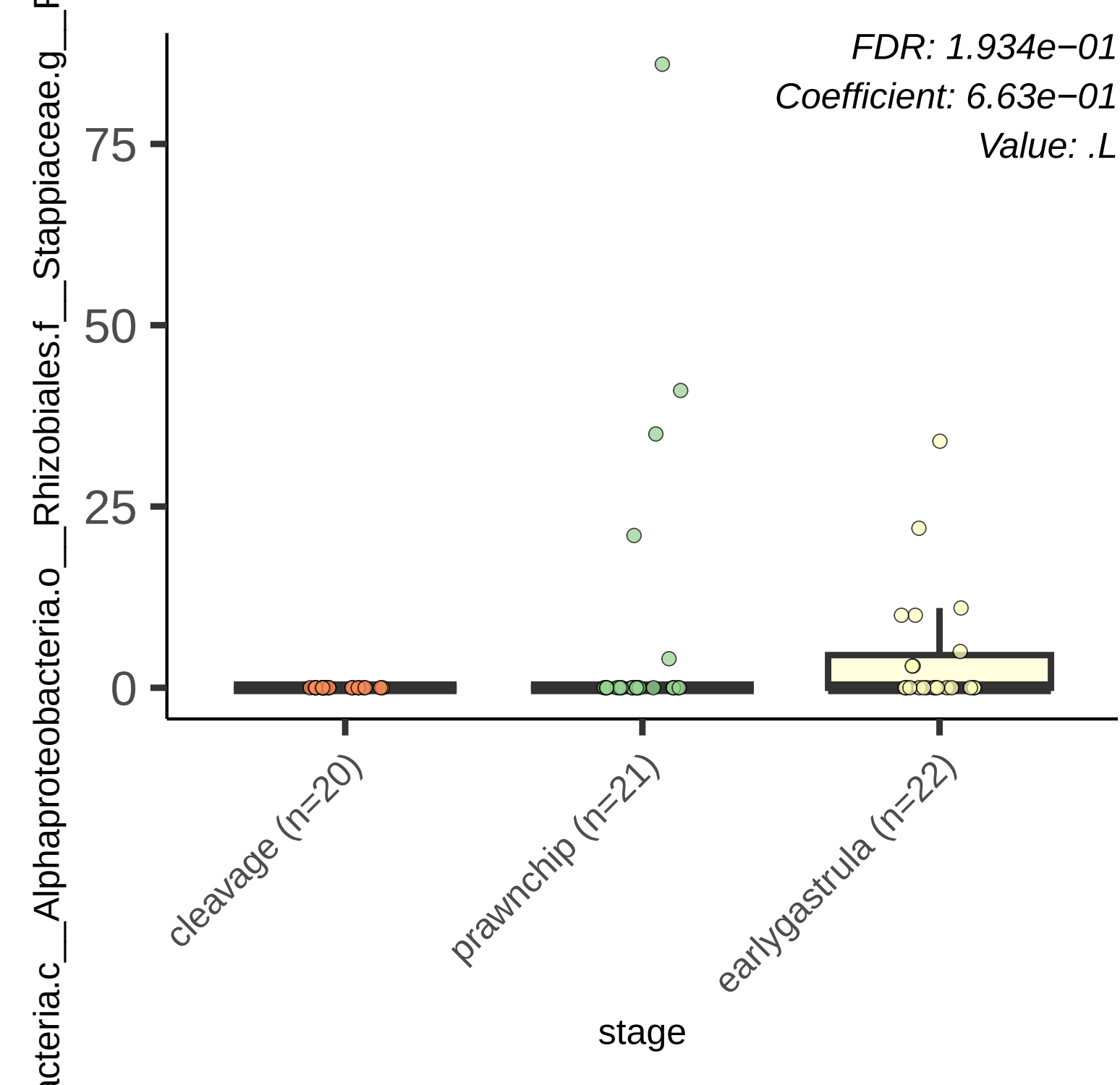
Coefficient: −7.98e−01
Value: .Q
750
500
250
0
cleavage (n=20)
prawnchip (n=21)
earlygastrula (n=22)
stage
idota.c__Bacteroidia.o__Flavobacteriales.f__Flavobacteriaceae.g__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Coefficient: 6.63e−01
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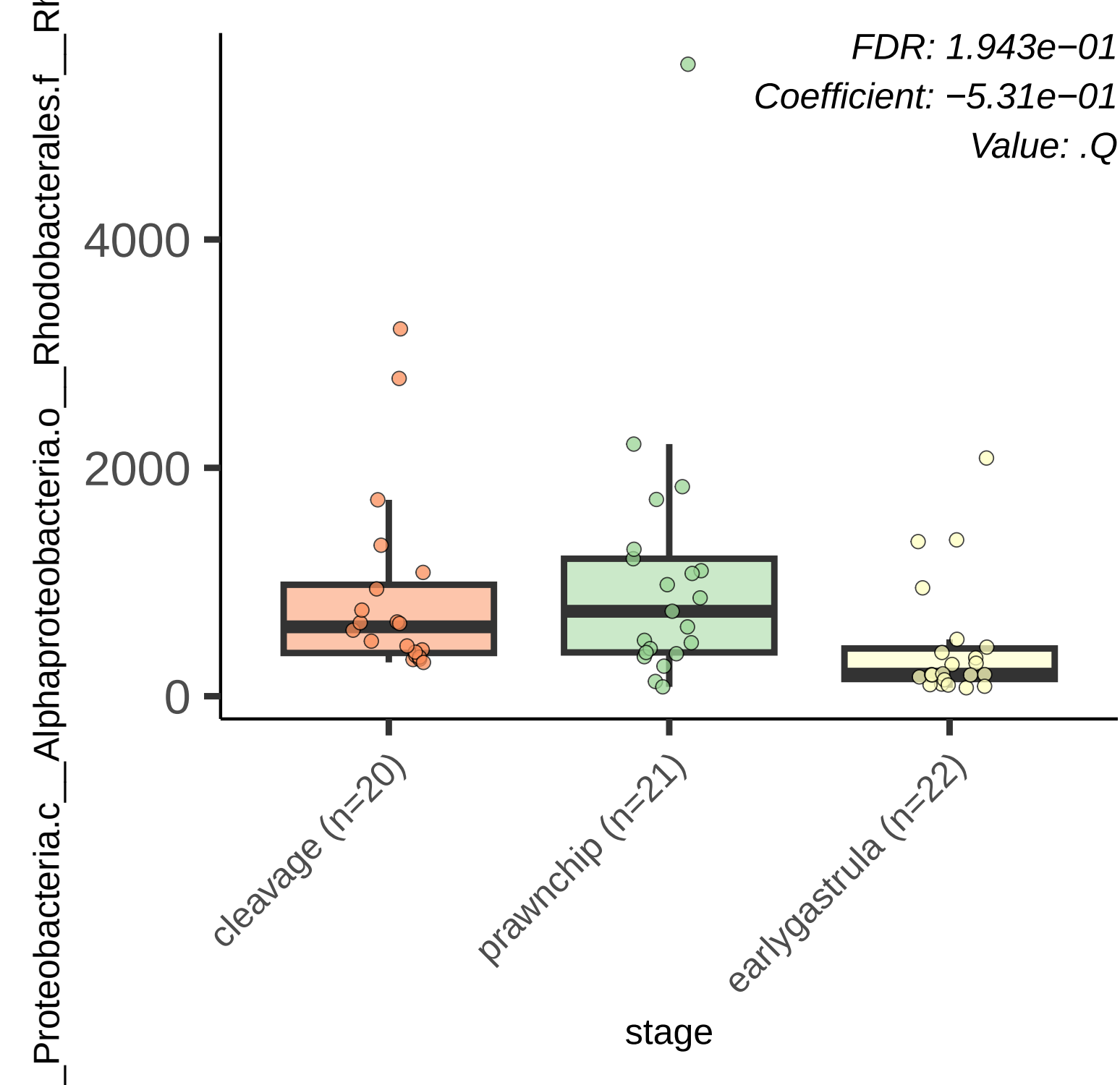
75
50
25
0
cleavage (n=20)
prawnchip (n=21)
earlygastrula (n=22)
stage
cteria.c__Alphaproteobacteria.o__Rhizobiales.f__Stappiaceae.g__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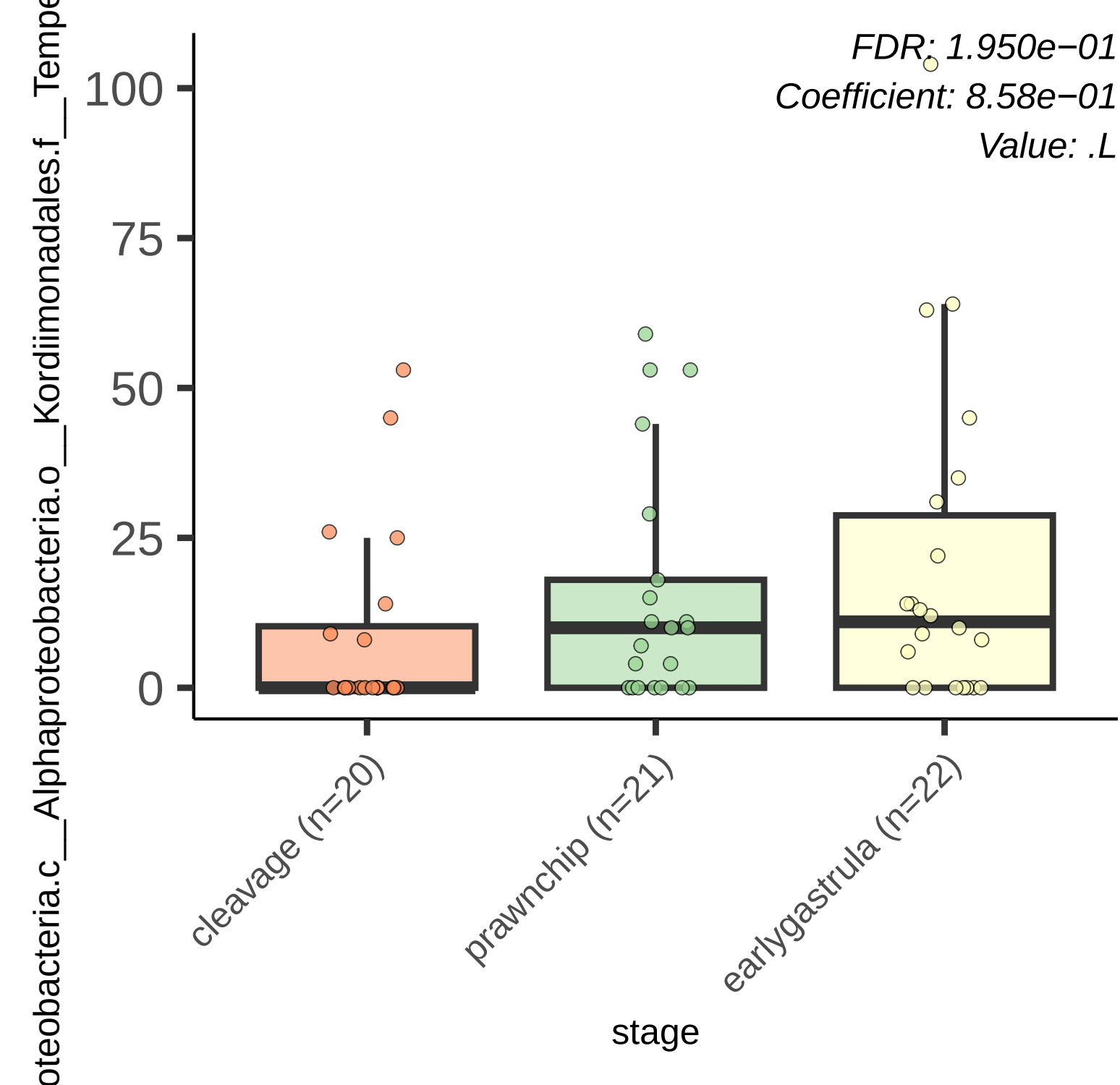
2000
0
cleavage (n=20)
prawnchip (n=21)
earlygastrula (n=22)
stage
_Proteobacteria.c__Alphaproteobacteria.o__Rhodobacterales.f__Rh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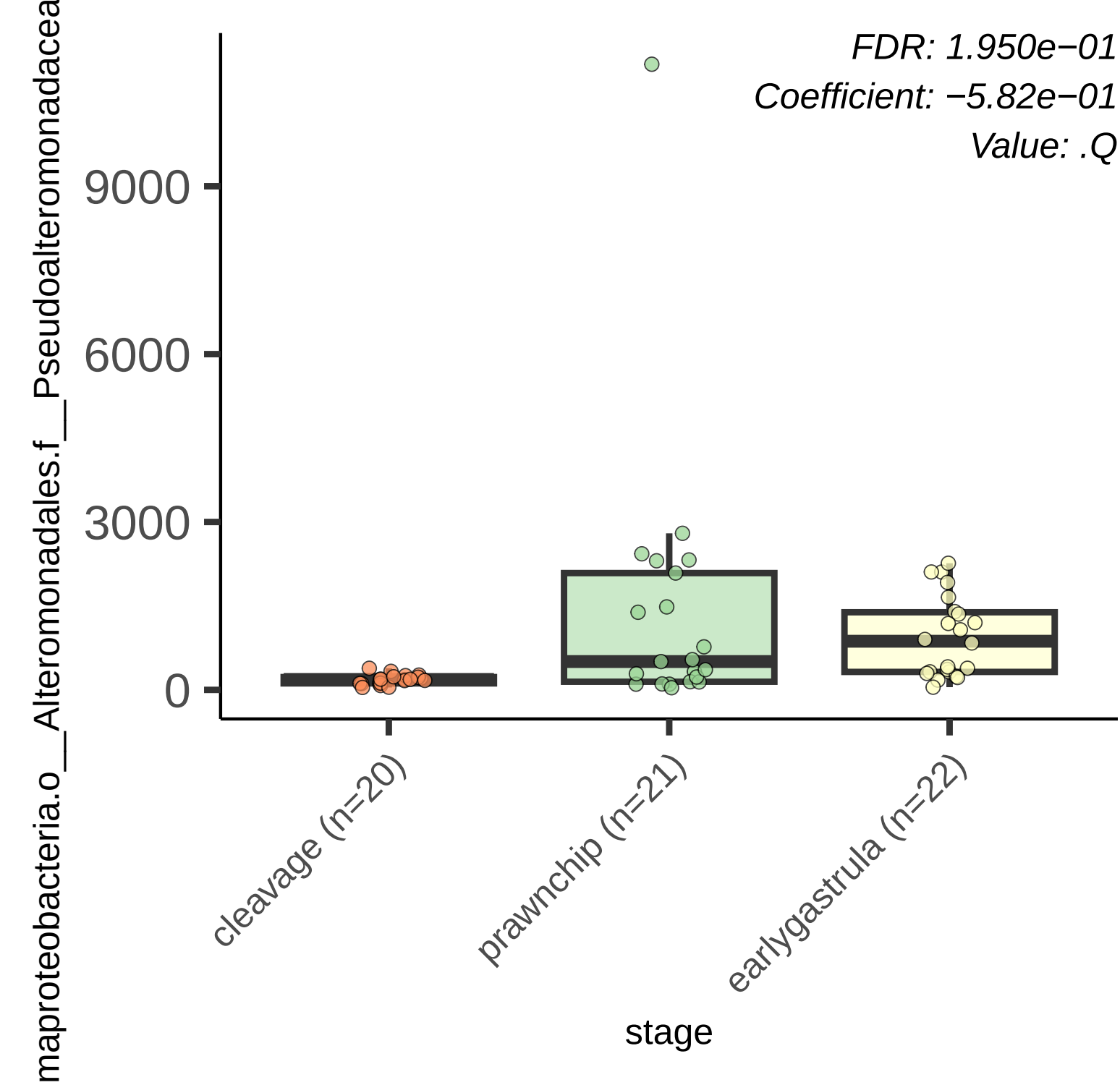
100
75
50
25
0
cleavage (n=20)
prawnchip (n=21)
earlygastrula (n=22)
stage
oteobacteria.c__Alphaproteobacteria.o__Kordiimonadales.f__Tempe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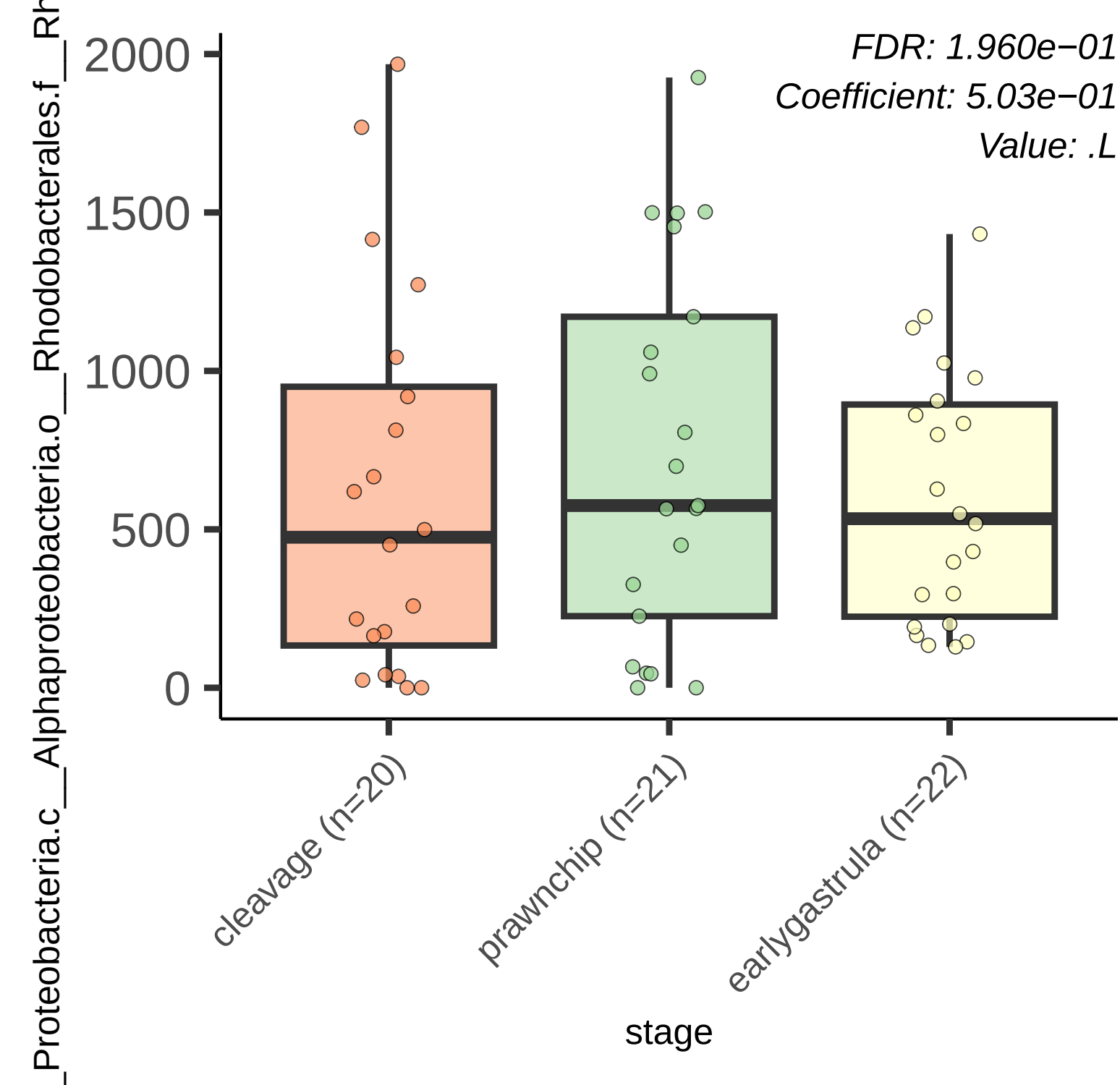
6000
3000
0
cleavage (n=20)
prawnchip (n=21)
earlygastrula (n=22)
stage
mapproteobacteria.o__Alteromonadales.f__Pseudoalteromonadacea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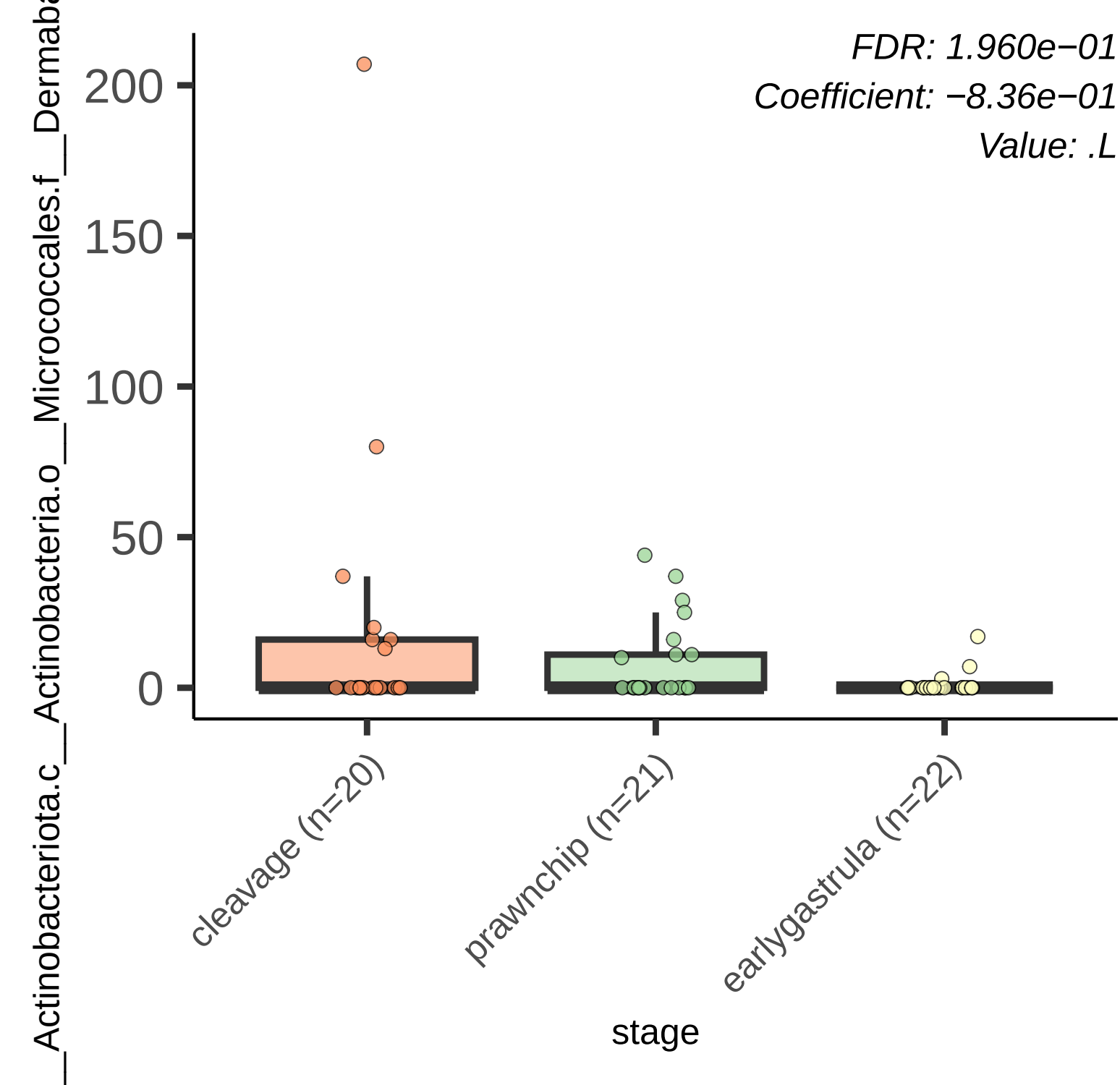
2000
1500
1000
500
0
cleavage (n=20)
prawnchip (n=21)
earlygastrula (n=22)
stage
_Proteobacteria.c__Alphaproteobacteria.o__Rhodobacterales.f__Rh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Coefficient: −8.36e−01
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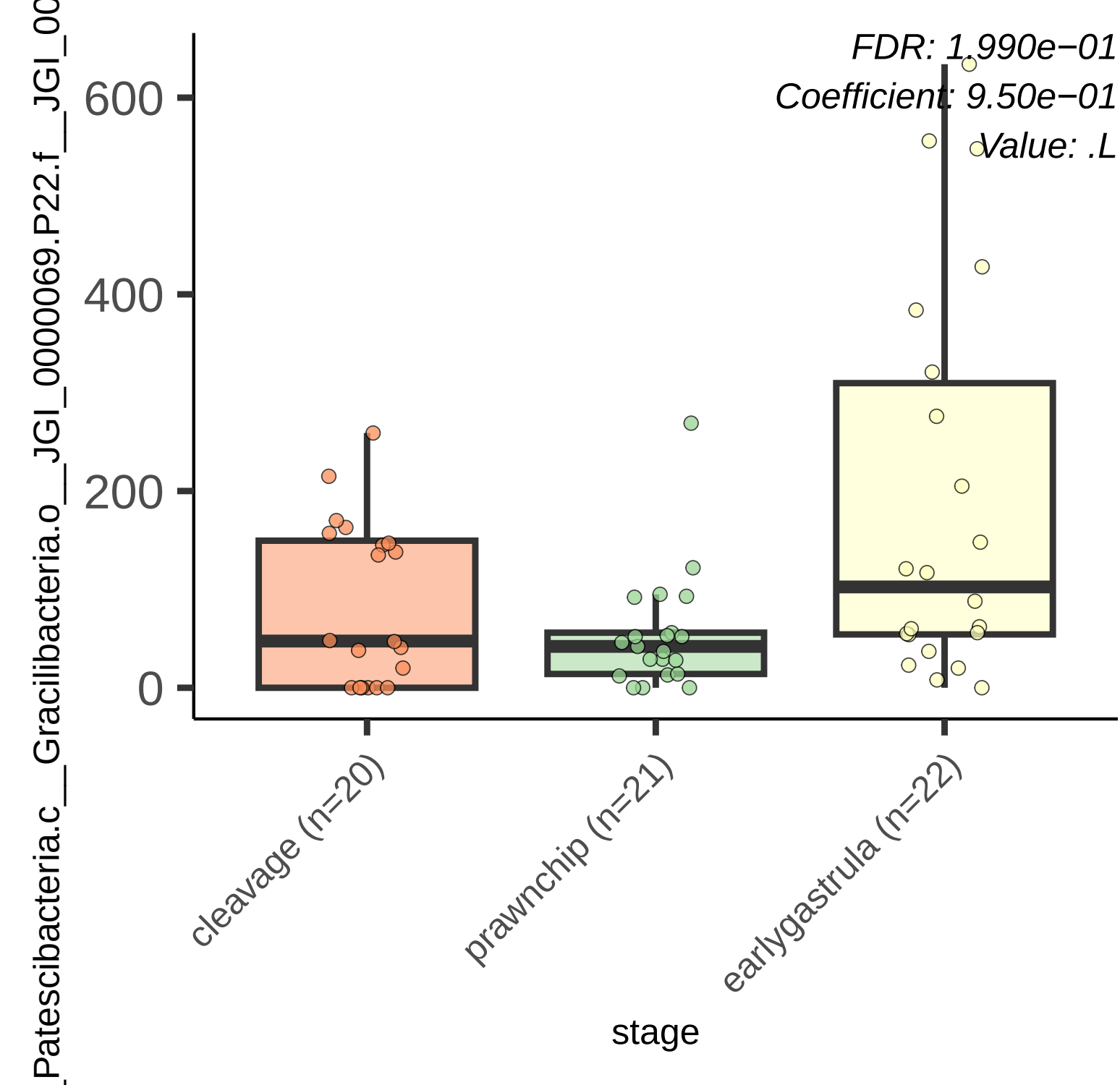
200
150
100
50
0
cleavage (n=20)
prawnchip (n=21)
earlygastrula (n=22)
stage
__Actinobacteriota.c__Actinobacteria.o__Micrococcales.f__Dermaba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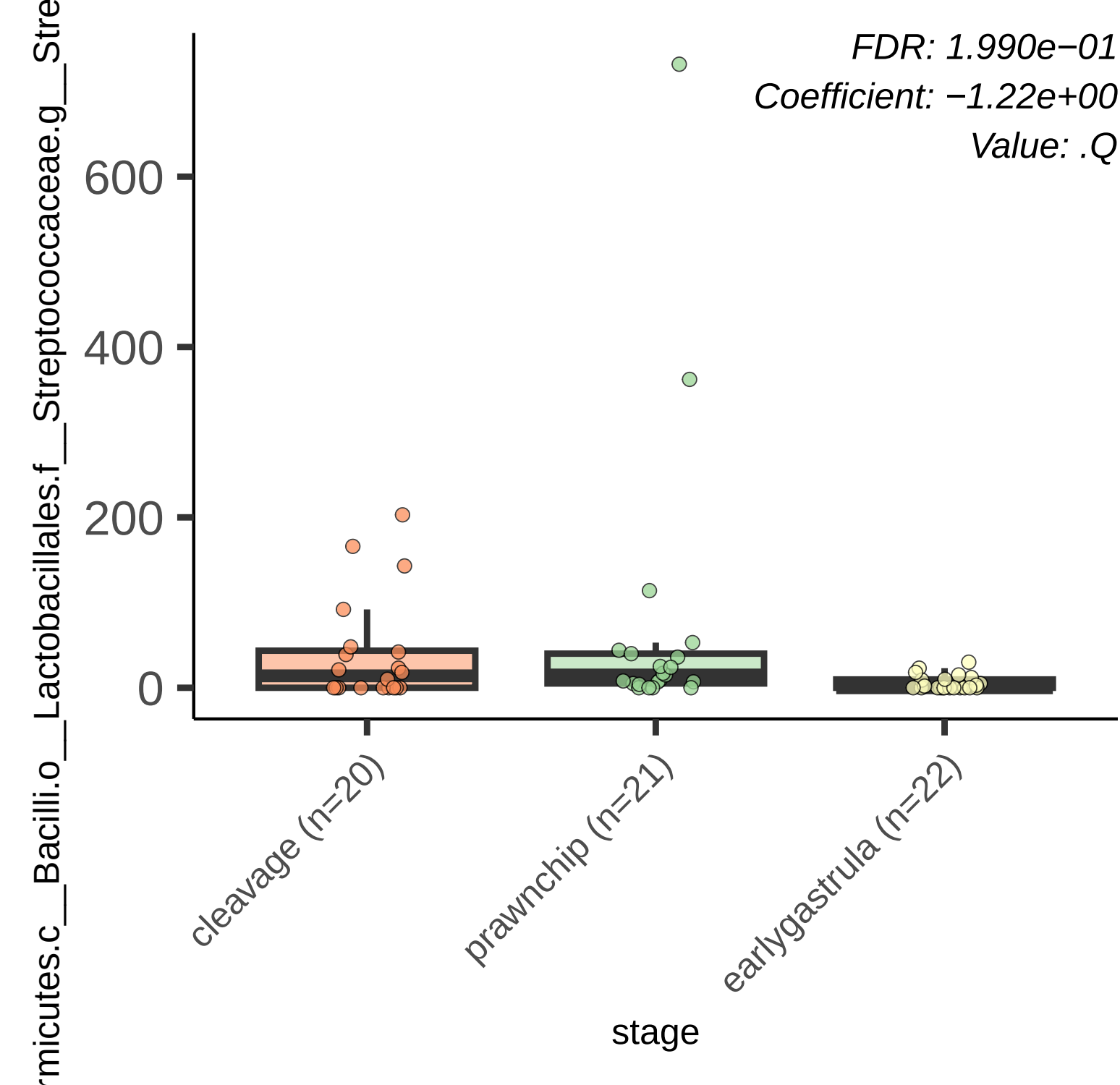
400
200
0
cleavage (n=20)
prawnchip (n=21)
earlygastrula (n=22)
stage
_Patescibacteria.c__Gracilibacteria.o__JGI_0000069.P22.f__JGI_00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Coefficient: −1.22e+00
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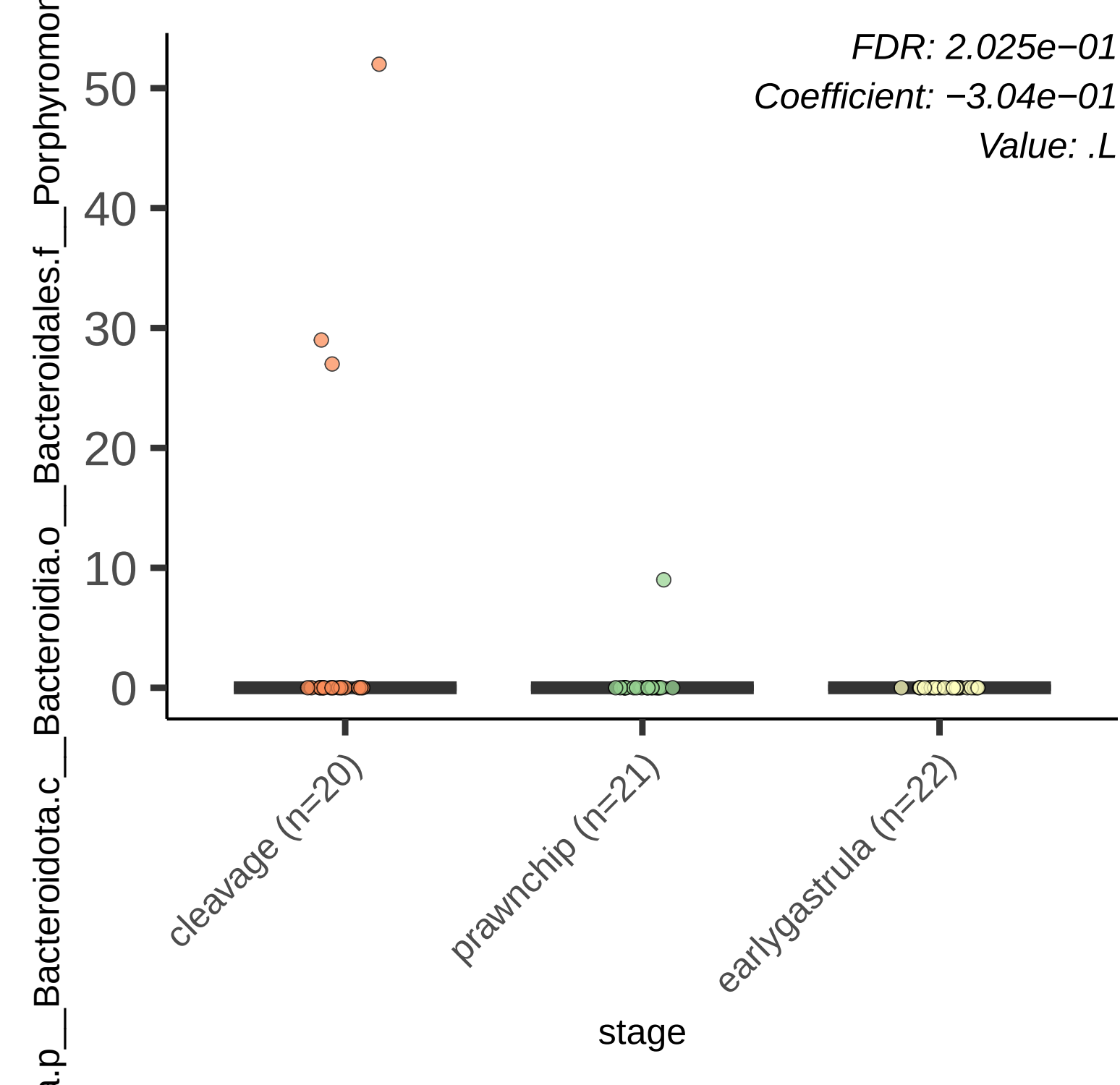
600
400
200
0
cleavage (n=20)
prawnchip (n=21)
earlygastrula (n=22)
stage
rmicutes.c__Bacilli.o__Lactobacillales.f__Streptococcaceae.g__Stre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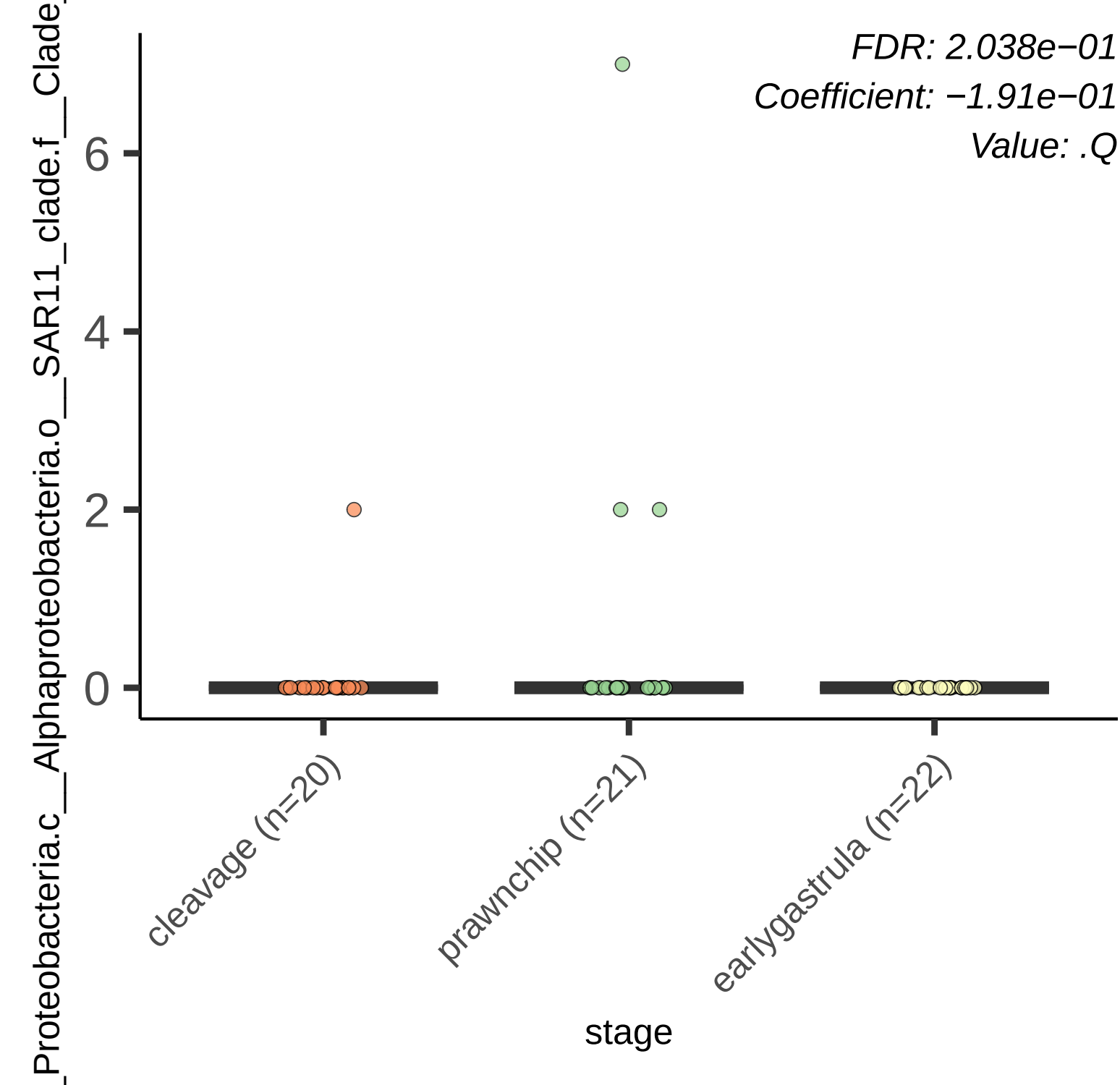
40
30
20
10
0
cleavage (n=20)
prawnchip (n=21)
earlygastrula (n=22)
stage
a.p__Bacteroidota.c__Bacteroidia.o__Bacteroidales.f__Porphyromon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6
4
2
0
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prawnchip (n=21)
earlygastrula (n=22)
stage
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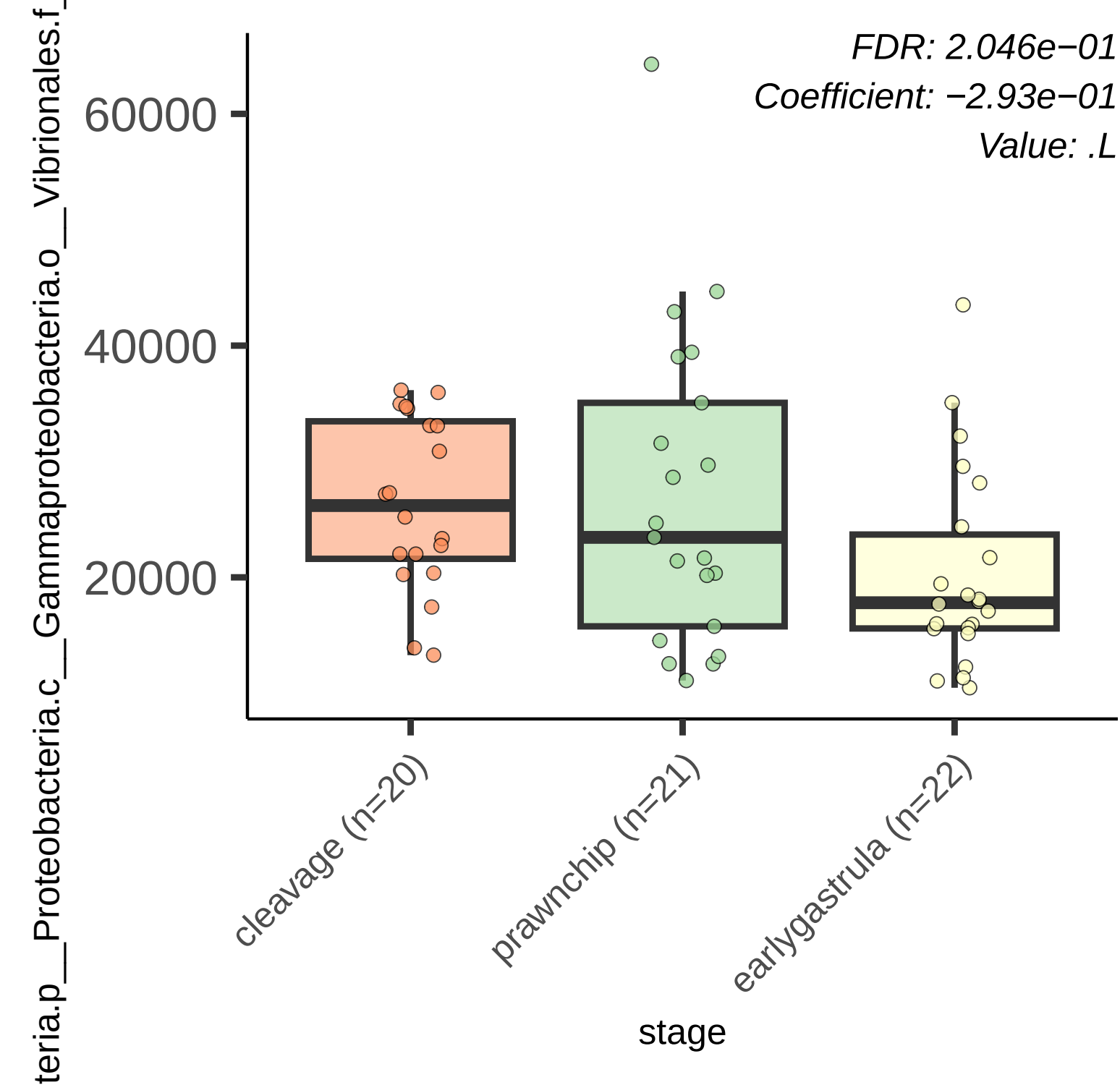
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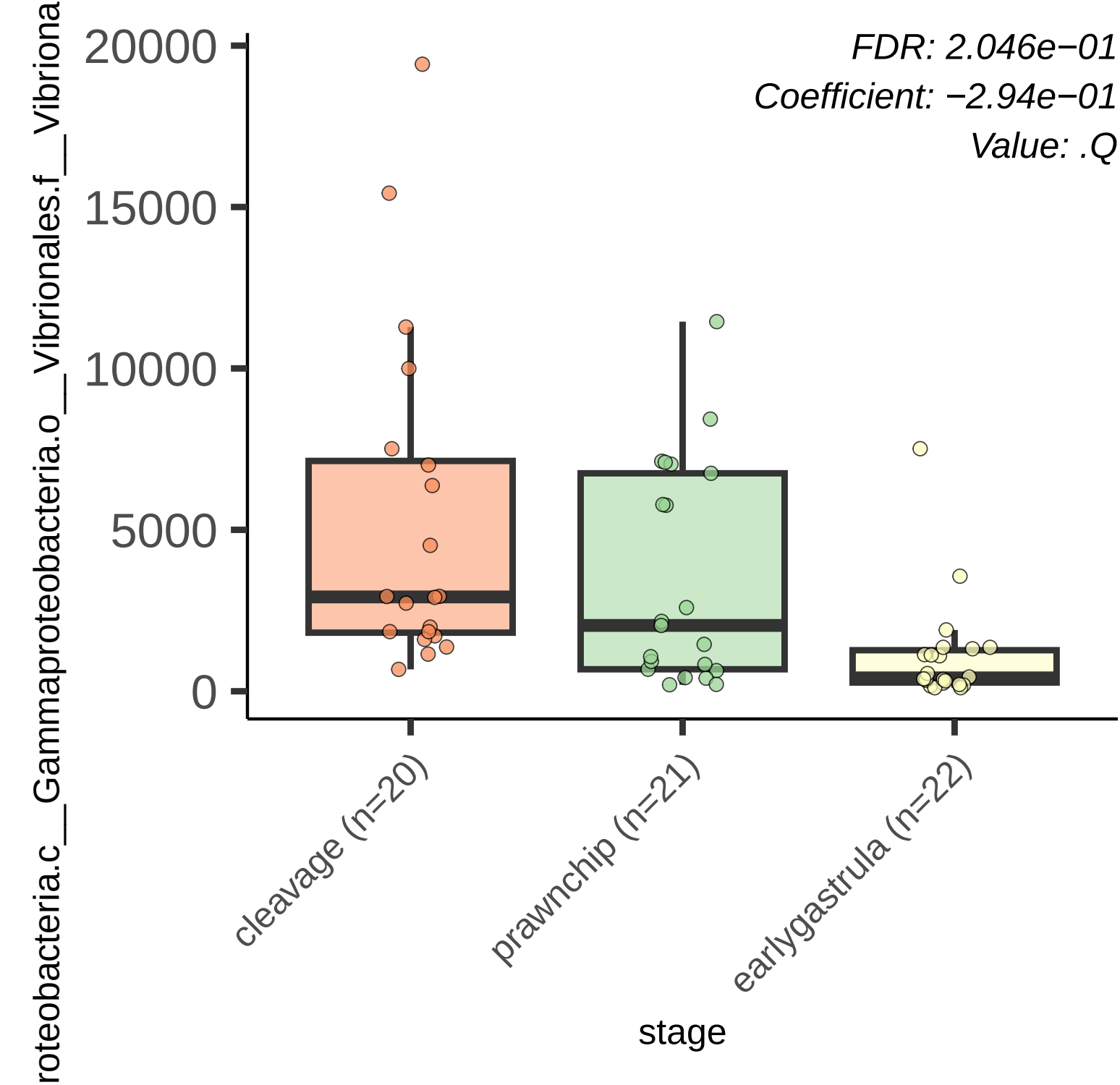
60000
40000
20000
cleavage (n=20)
prawnchip (n=21)
earlygastrula (n=22)
stage
teria.p__Proteobacteria.c__Gammaproteobacteria.o__Vibrionales.f_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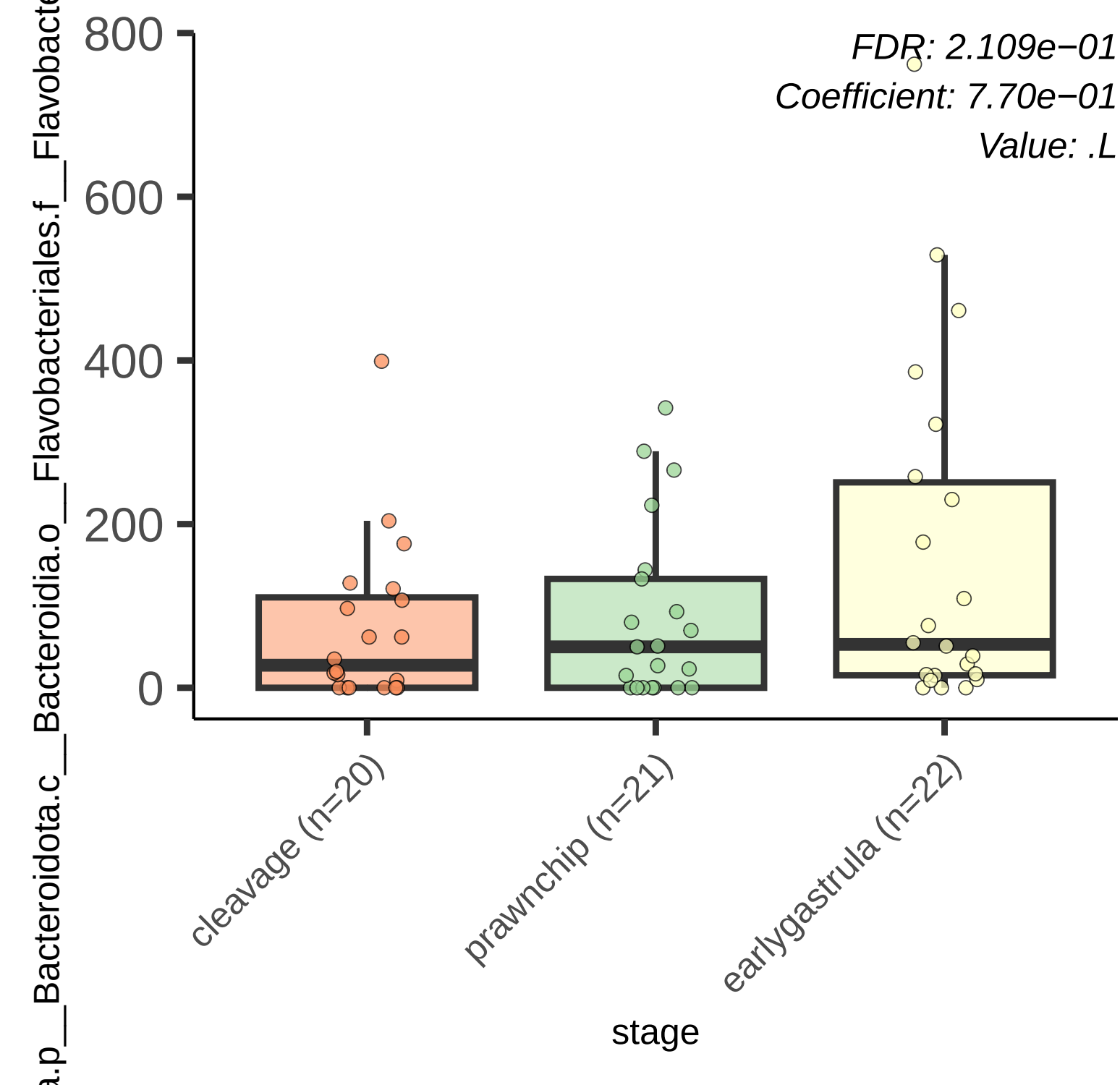
20000
15000
10000
5000
0
cleavage (n=20)
prawnchip (n=21)
earlygastrula (n=22)
stage
roteobacteria.c__Gammaproteobacteria.o__Vibrionales.f__Vibriona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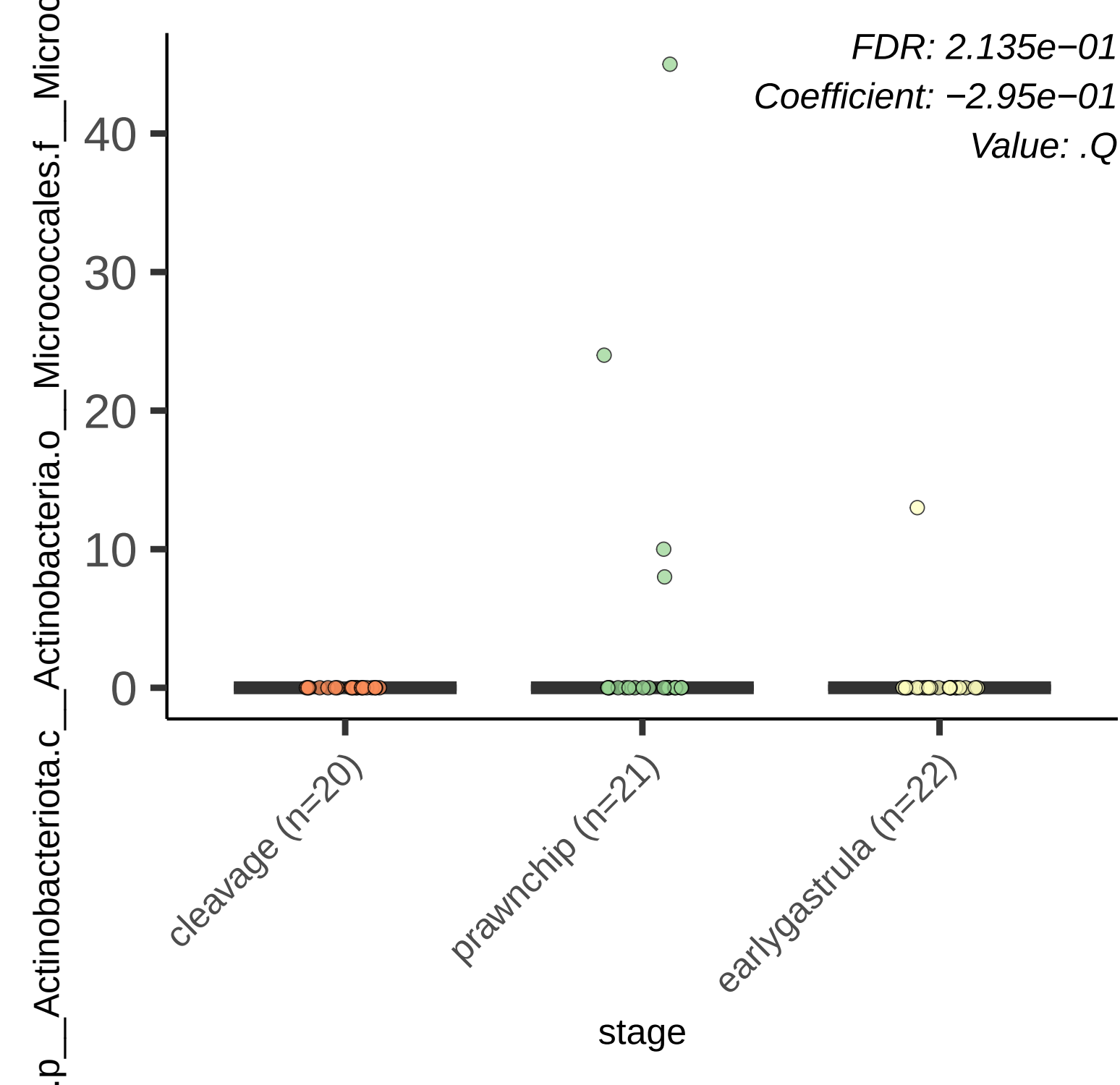
800
600
400
200
0
cleavage (n=20)
prawnchip (n=21)
earlygastrula (n=22)
stage
a.p__Bacteroidota.c__Bacteroidia.o__Flavobacteriales.f__Flavobacte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Coefficient: −2.95e−01
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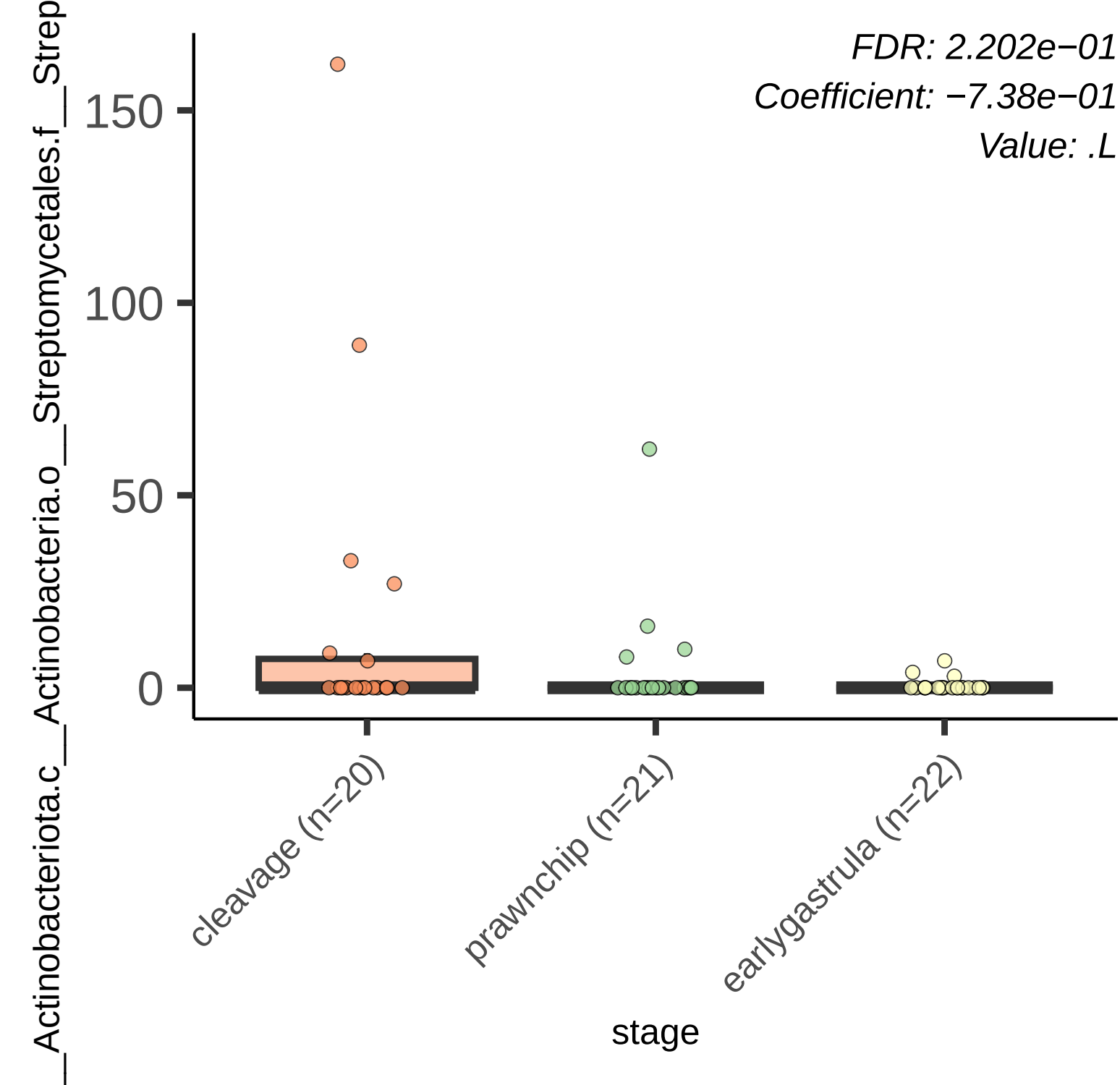
40
30
20
10
0
cleavage (n=20)
prawnchip (n=21)
earlygastrula (n=22)
stage
.p__Actinobacteriota.c__Actinobacteria.o__Micrococcales.f__Microc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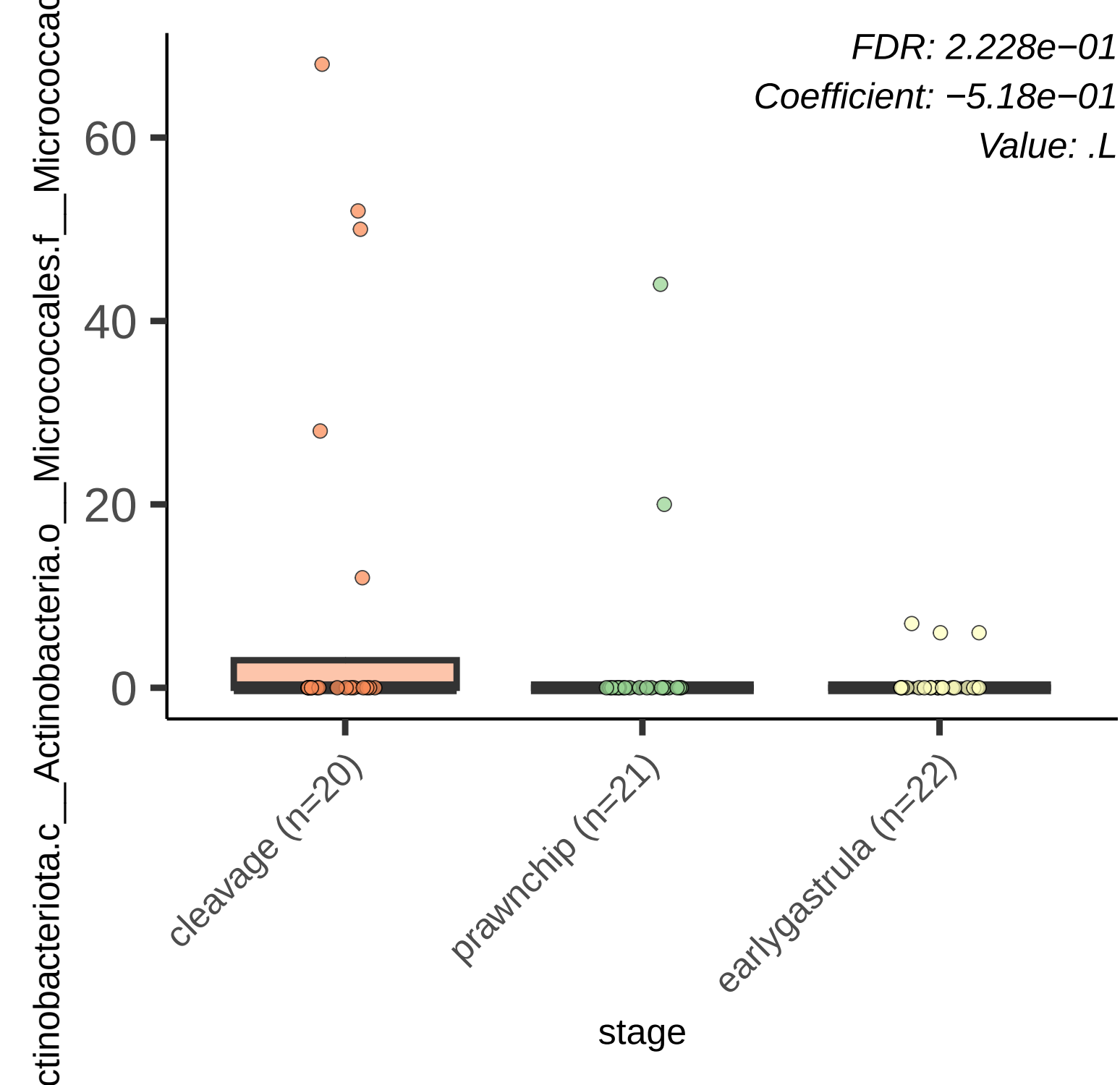
Value: .L
150
100
50
0
cleavage (n=20)
prawnchip (n=21)
earlygastrula (n=22)
stage
__Actinobacteriota.c__Actinobacteria.o__Streptomycetales.f__Strep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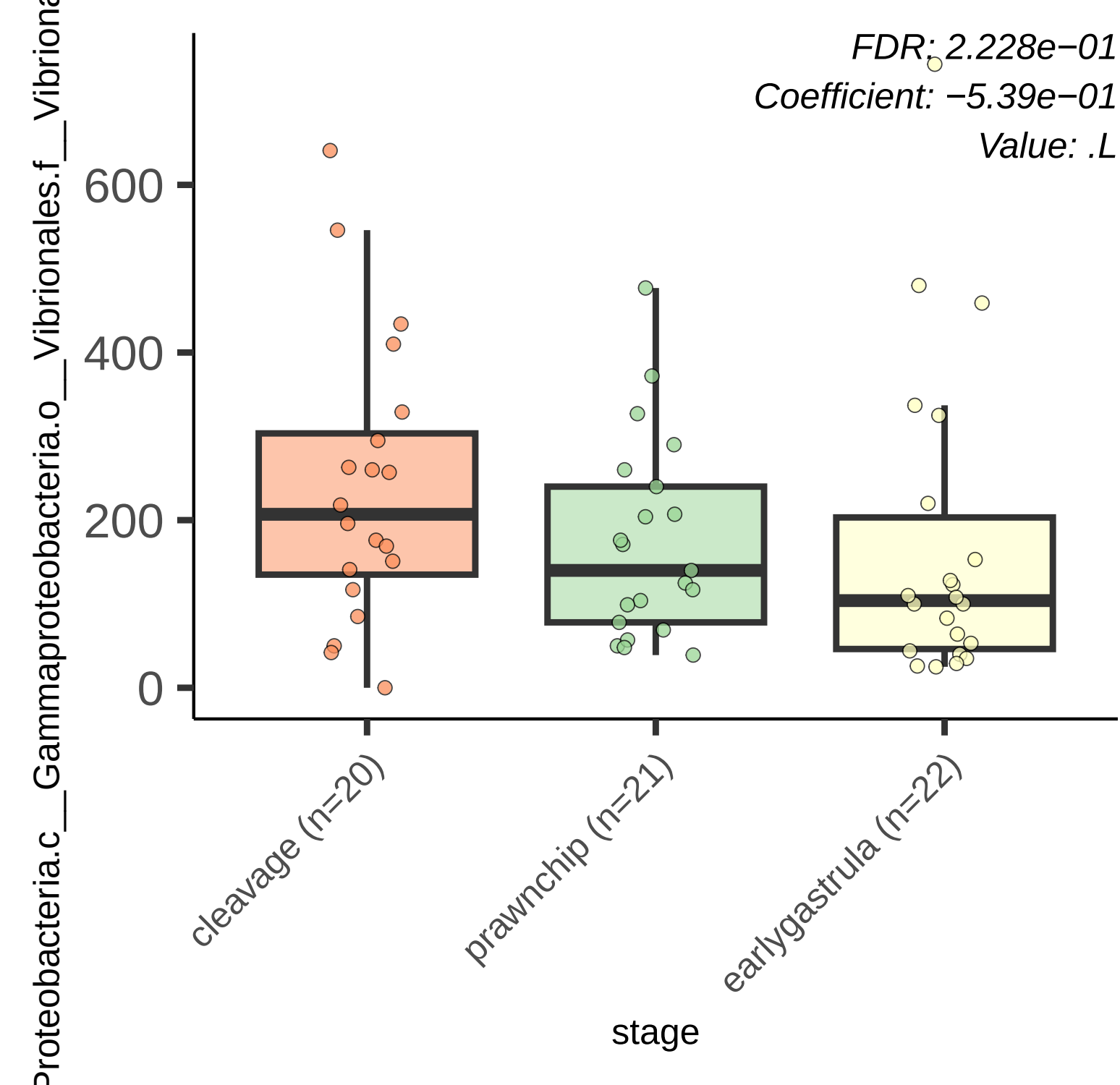
60
40
20
0
cleavage (n=20)
prawnchip (n=21)
earlygastrula (n=22)
stage
ctinobacteriota.c__Actinobacteria.o__Micrococcales.f__Micrococcac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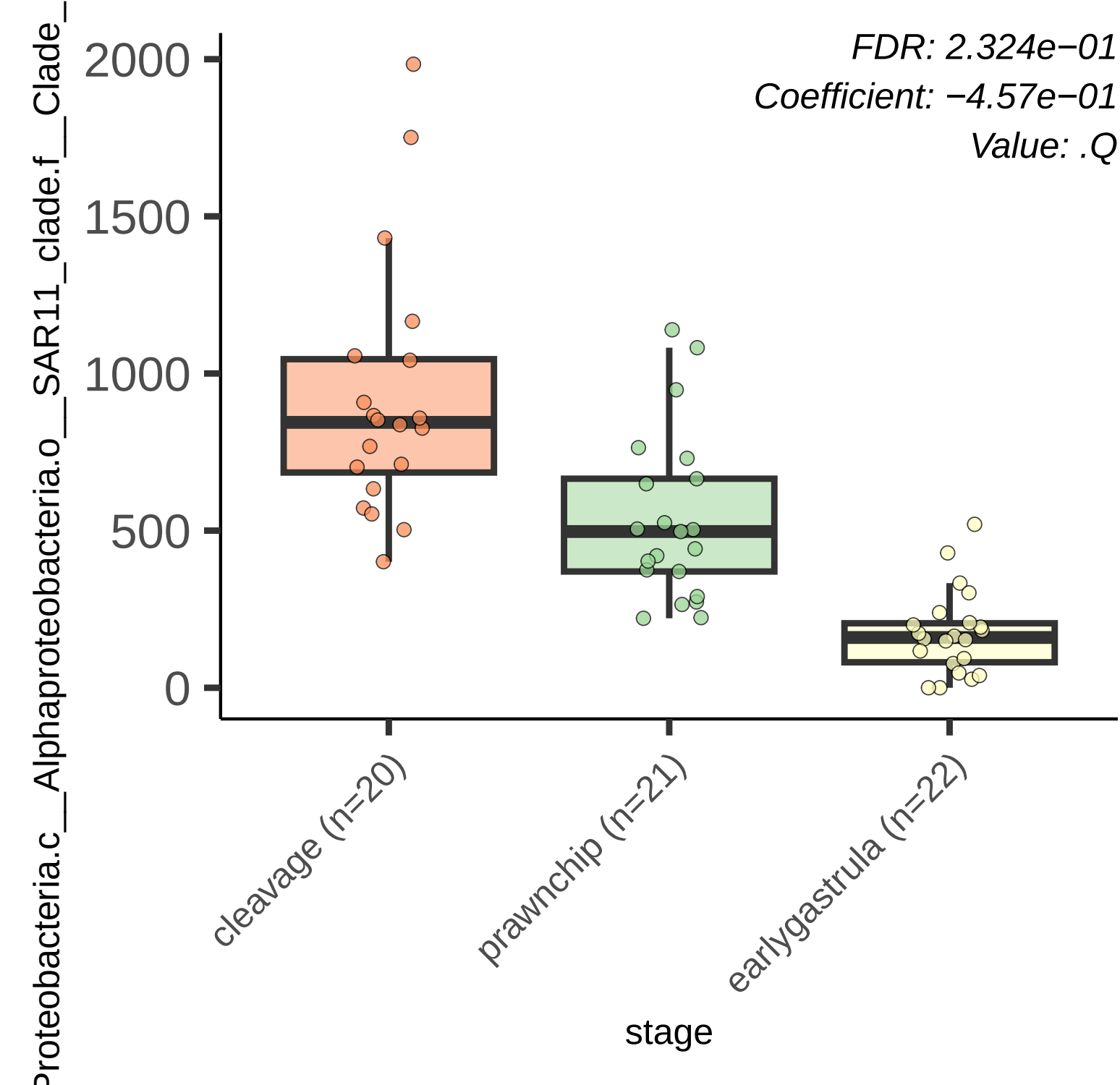
600
400
200
0
cleavage (n=20)
prawnchip (n=21)
earlygastrula (n=22)
stage
Proteobacteria.c__Gammaproteobacteria.o__Vibrionales.f__Vibriona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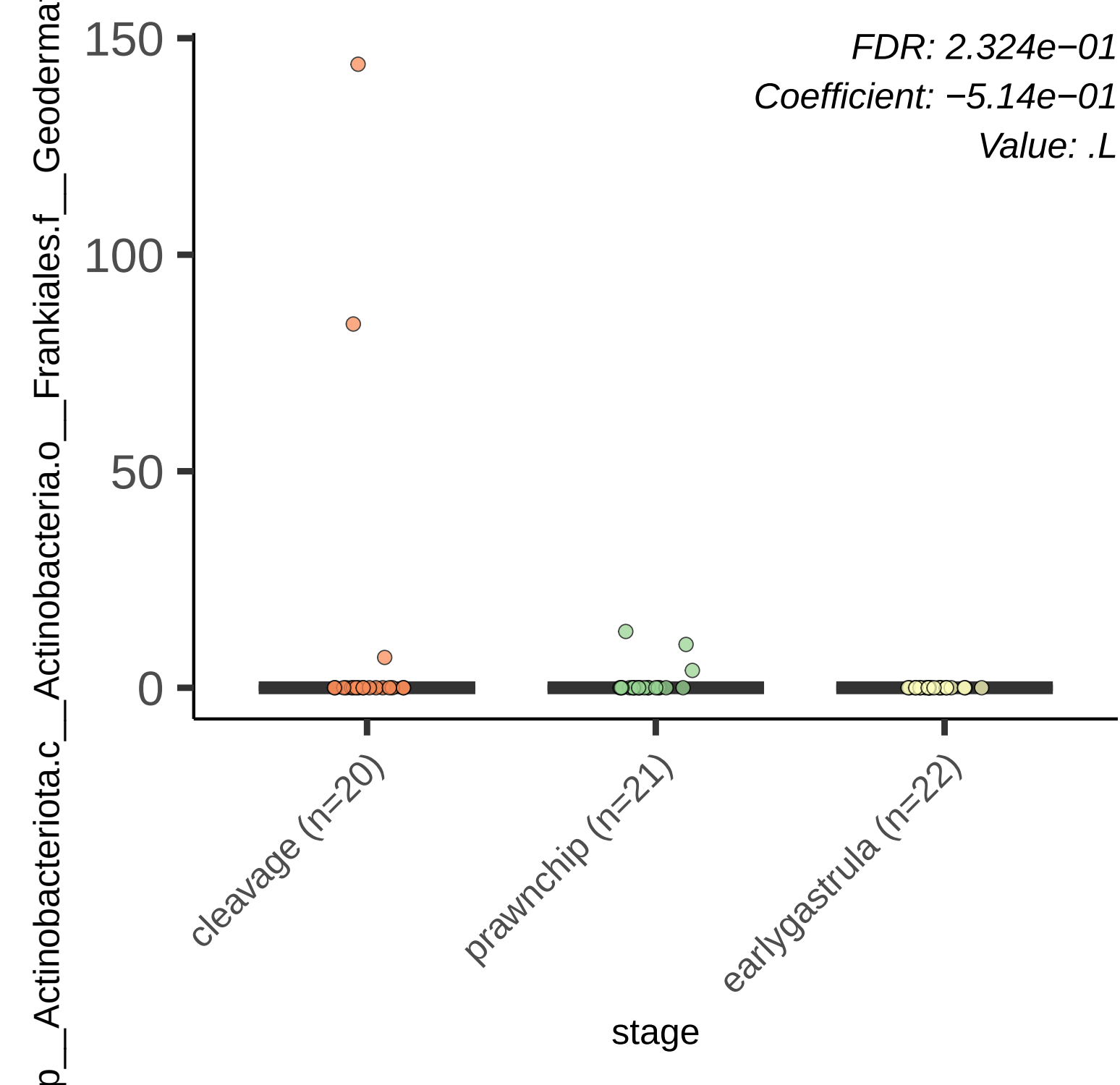
Coefficient: −4.57e−01
Value: .Q
2000
1500
1000
500
0
cleavage (n=20)
prawnchip (n=21)
earlygastrula (n=22)
stage
Proteobacteria.c__Alphaproteobacteria.o__SAR11_clade.f__Clade_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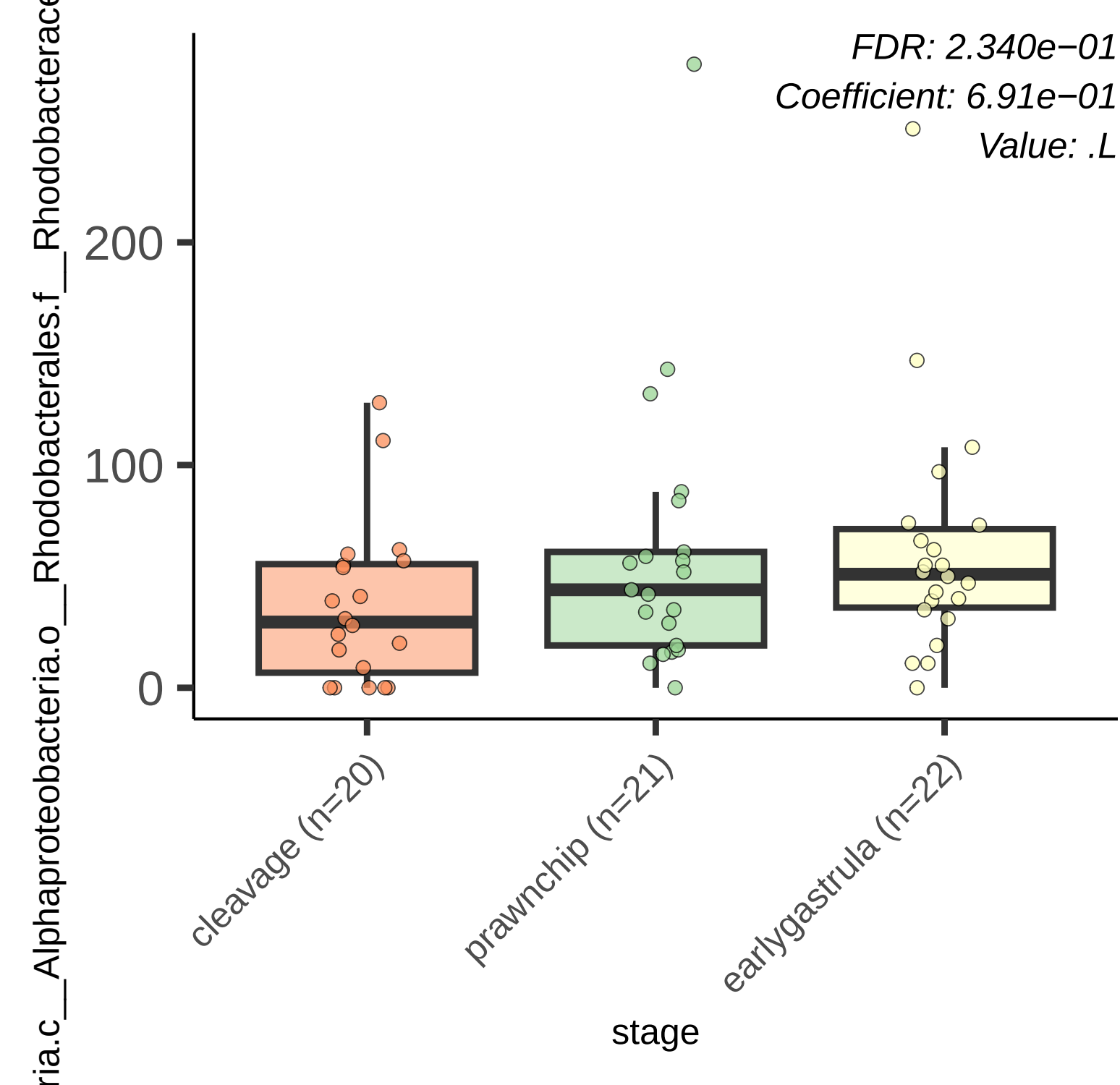
100
50
0
cleavage (n=20)
prawnchip (n=21)
earlygastrula (n=22)
stage
p__Actinobacteriota.c__Actinobacteria.o__Frankiales.f__Geoderma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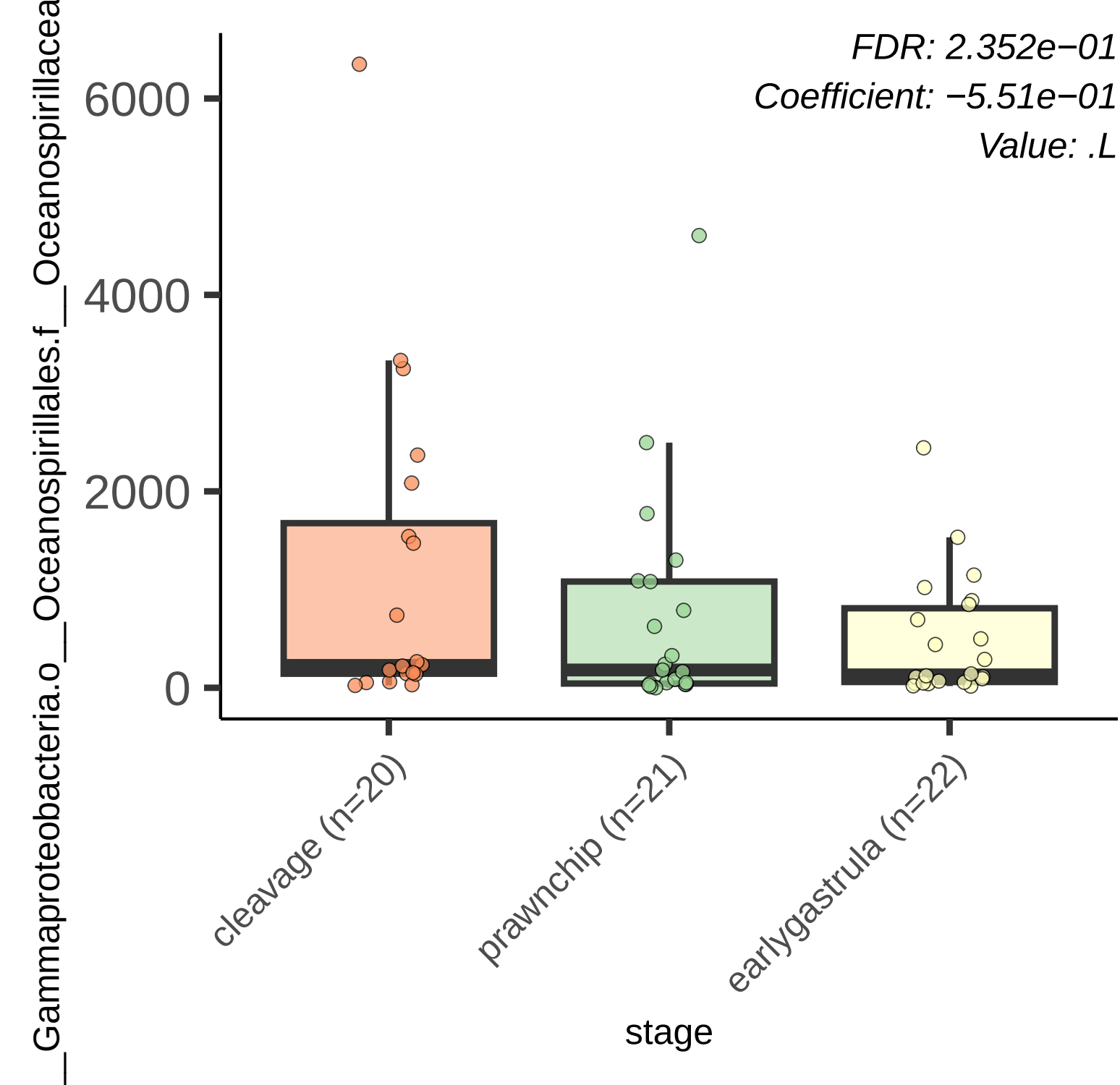
Coefficient: 6.91e−01
Value: .L
200
100
0
cleavage (n=20)
prawnchip (n=21)
earlygastrula (n=22)
stage
ria.c__Alphaproteobacteria.o__Rhodobacterales.f__Rhodobacterace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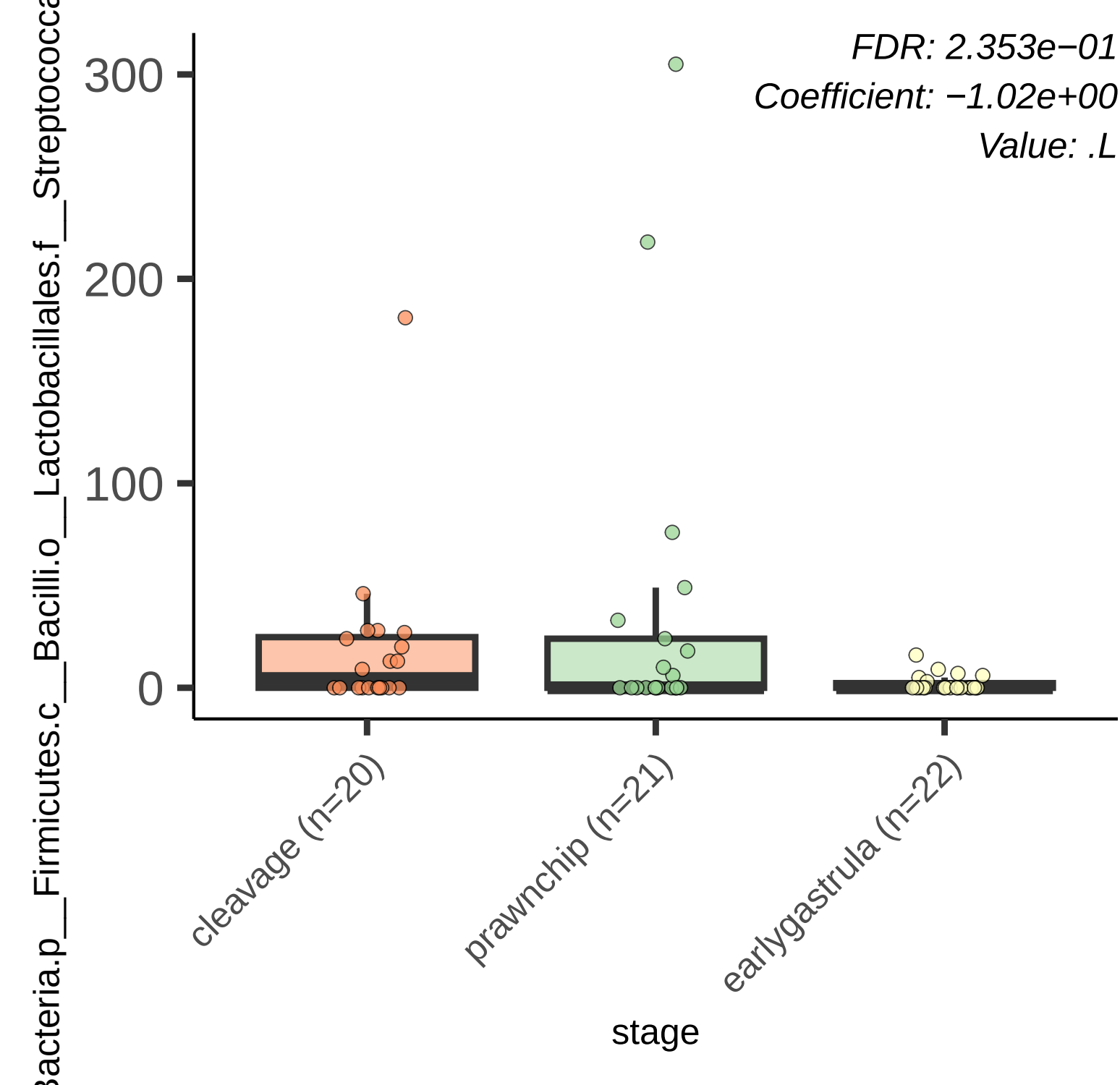
Value: .L
6000
4000
2000
0
cleavage (n=20)
prawnchip (n=21)
earlygastrula (n=22)
stage
__Gammaproteobacteria.o__Oceanospirillales.f__Oceanospirillacea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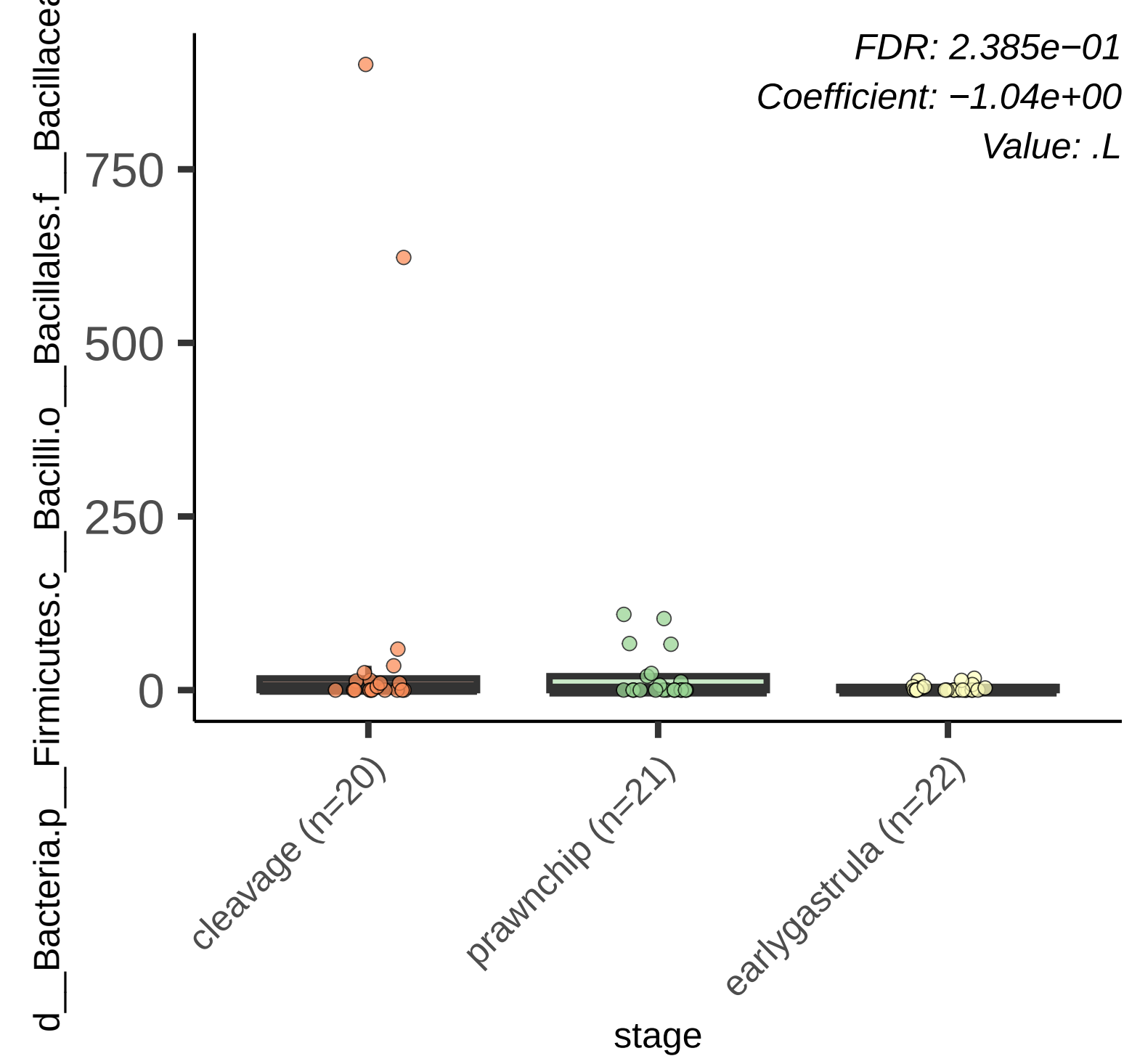
Coefficient: −1.02e+00
Value: .L
300
200
100
0
cleavage (n=20)
prawnchip (n=21)
earlygastrula (n=22)
stage
Bacteria.p__Firmicutes.c__Bacilli.o__Lactobacillales.f__Streptococca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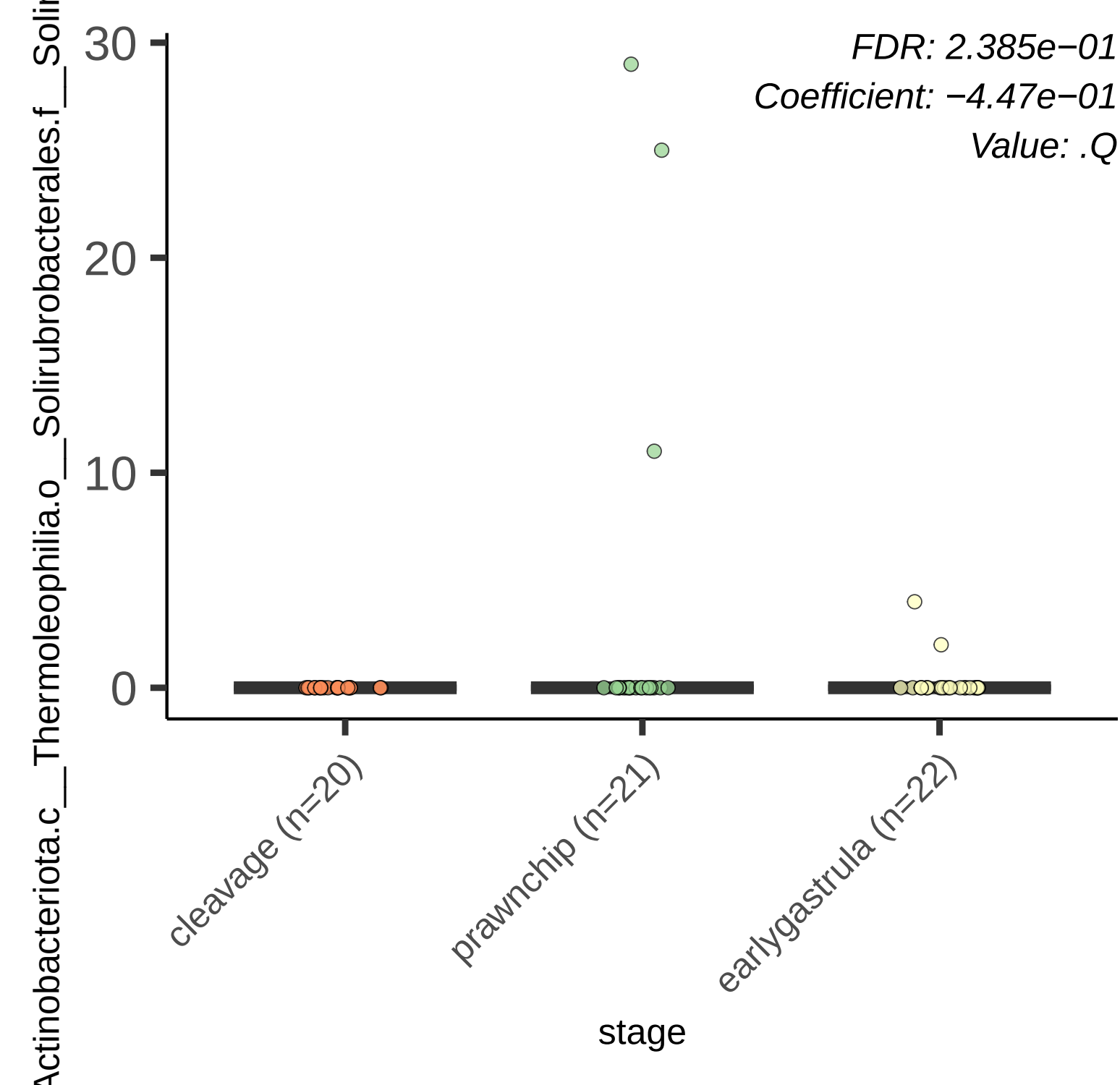
d__Bacteria.p__Firmicutes.c__Bacilli.o__Bacillales.f__Bacillace
750
500
250
0
cleavage (n=20)
prawnchip (n=21)
earlygastrula (n=22)
stage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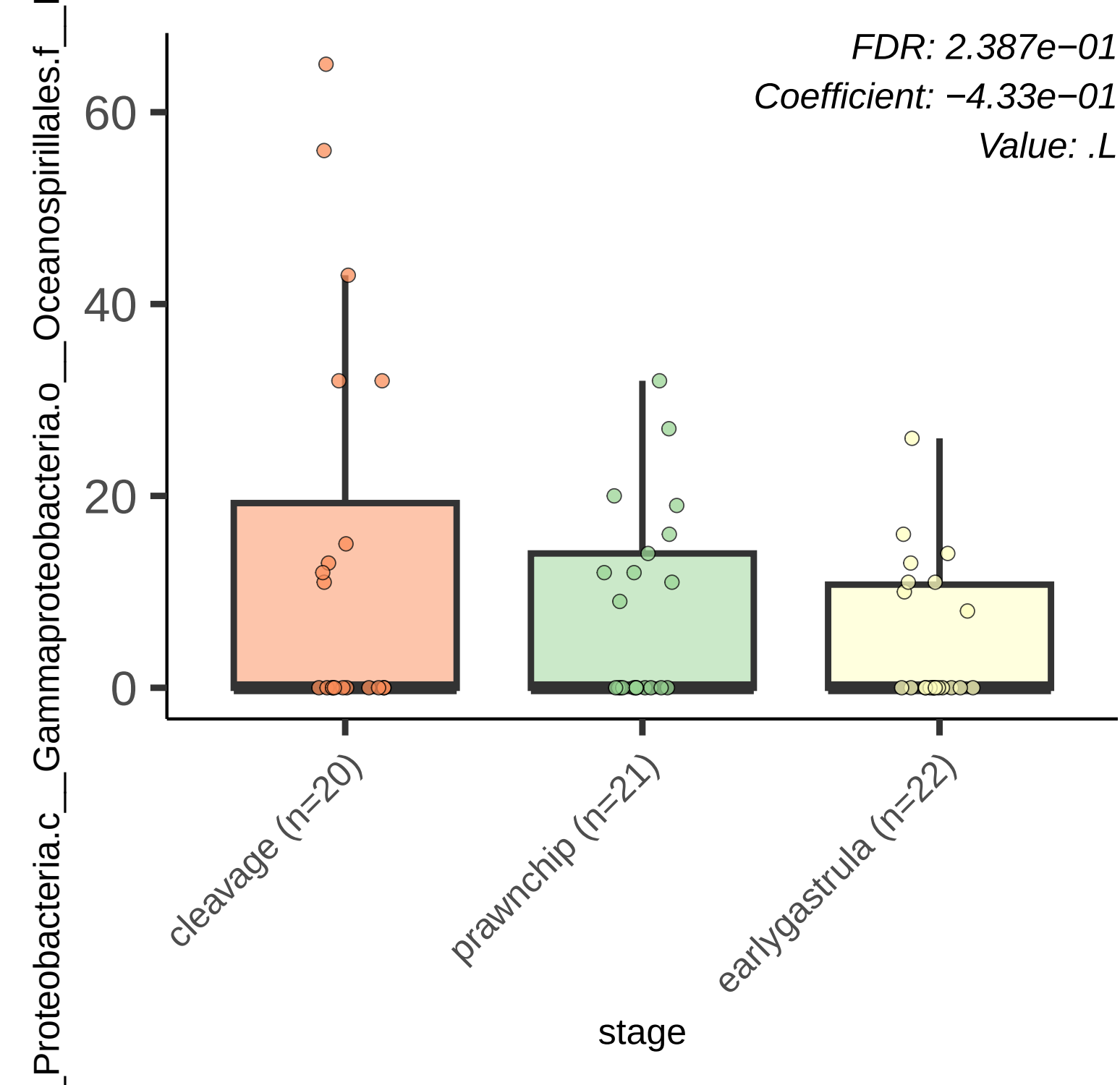
Coefficient: −4.47e−01
Value: .Q
30
20
10
0
cleavage (n=20)
prawnchip (n=21)
earlygastrula (n=22)
stage
Actinobacteriota.c__Thermoleophilia.o__Solirubrobacterales.f__Soli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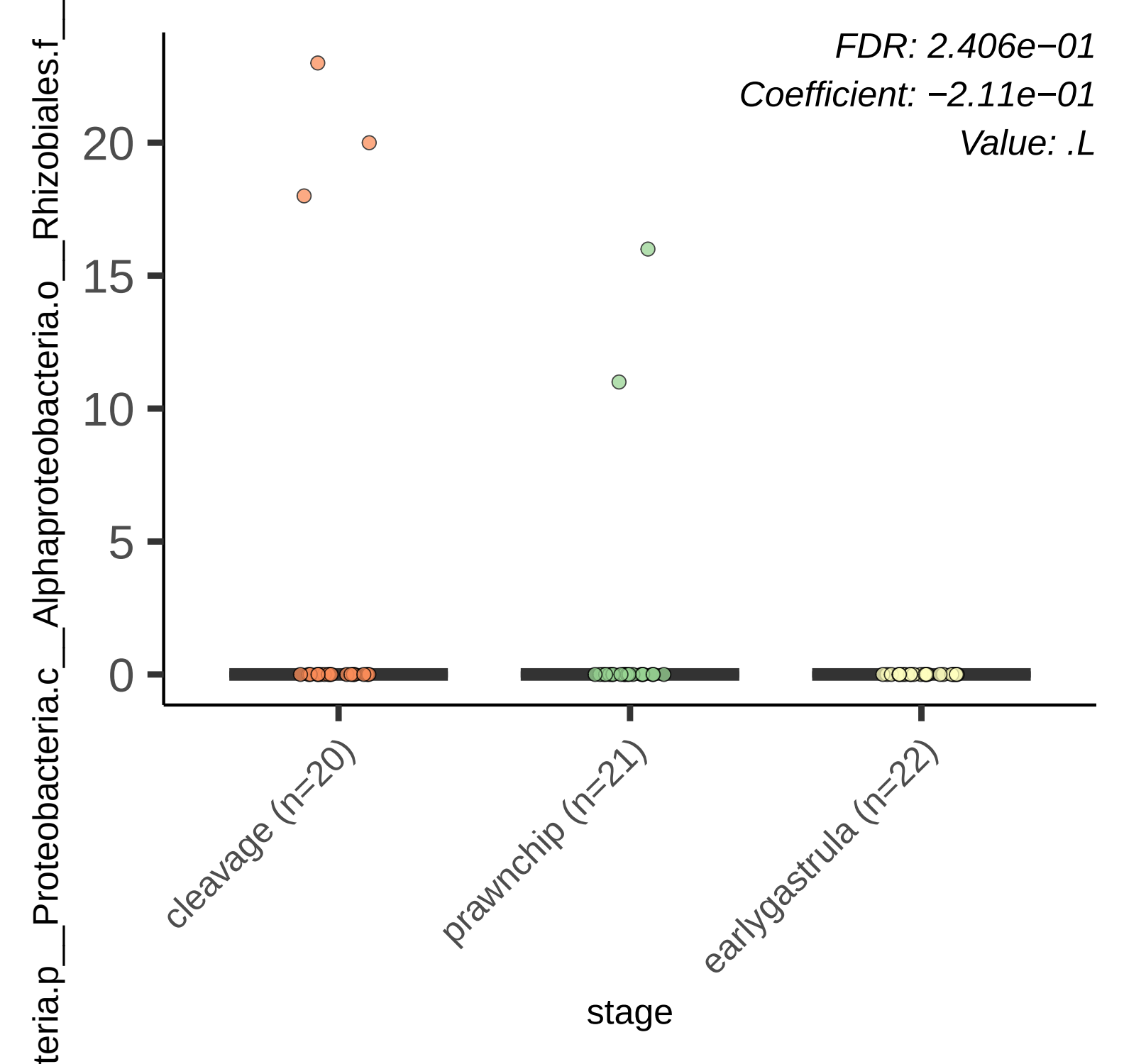
Coefficient: −4.33e−01
Value: .L
60
40
20
0
cleavage (n=20)
prawnchip (n=21)
earlygastrula (n=22)
stage
_Proteobacteria.c__Gammaproteobacteria.o__Oceanospirillales.f__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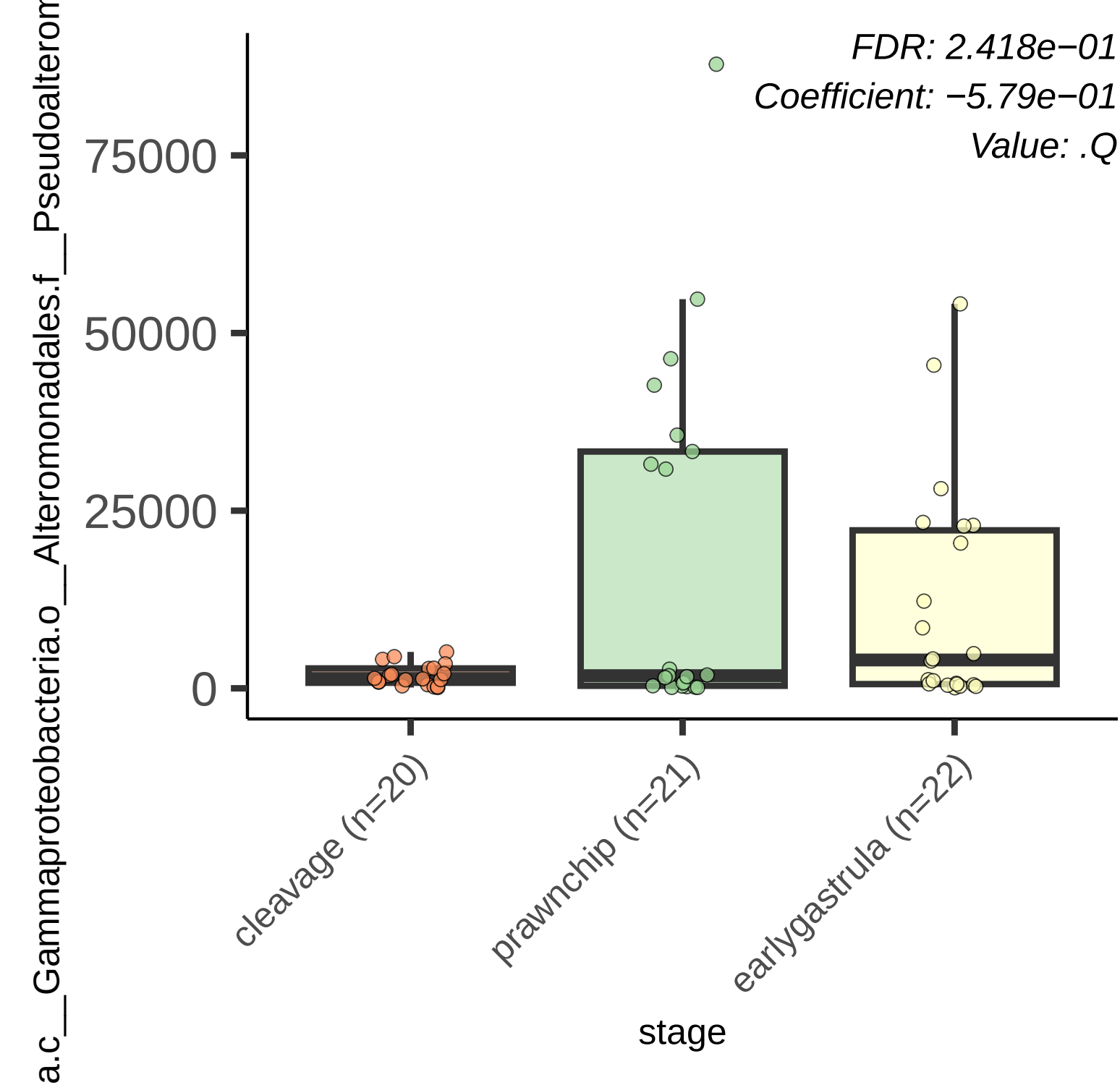
Coefficient: −2.11e−01
Value: .L
20
15
10
5
0
cleavage (n=20)
prawnchip (n=21)
earlygastrula (n=22)
stage
teria.p__Proteobacteria.c__Alphaproteobacteria.o__Rhizobiales.f__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Coefficient: −5.79e−01
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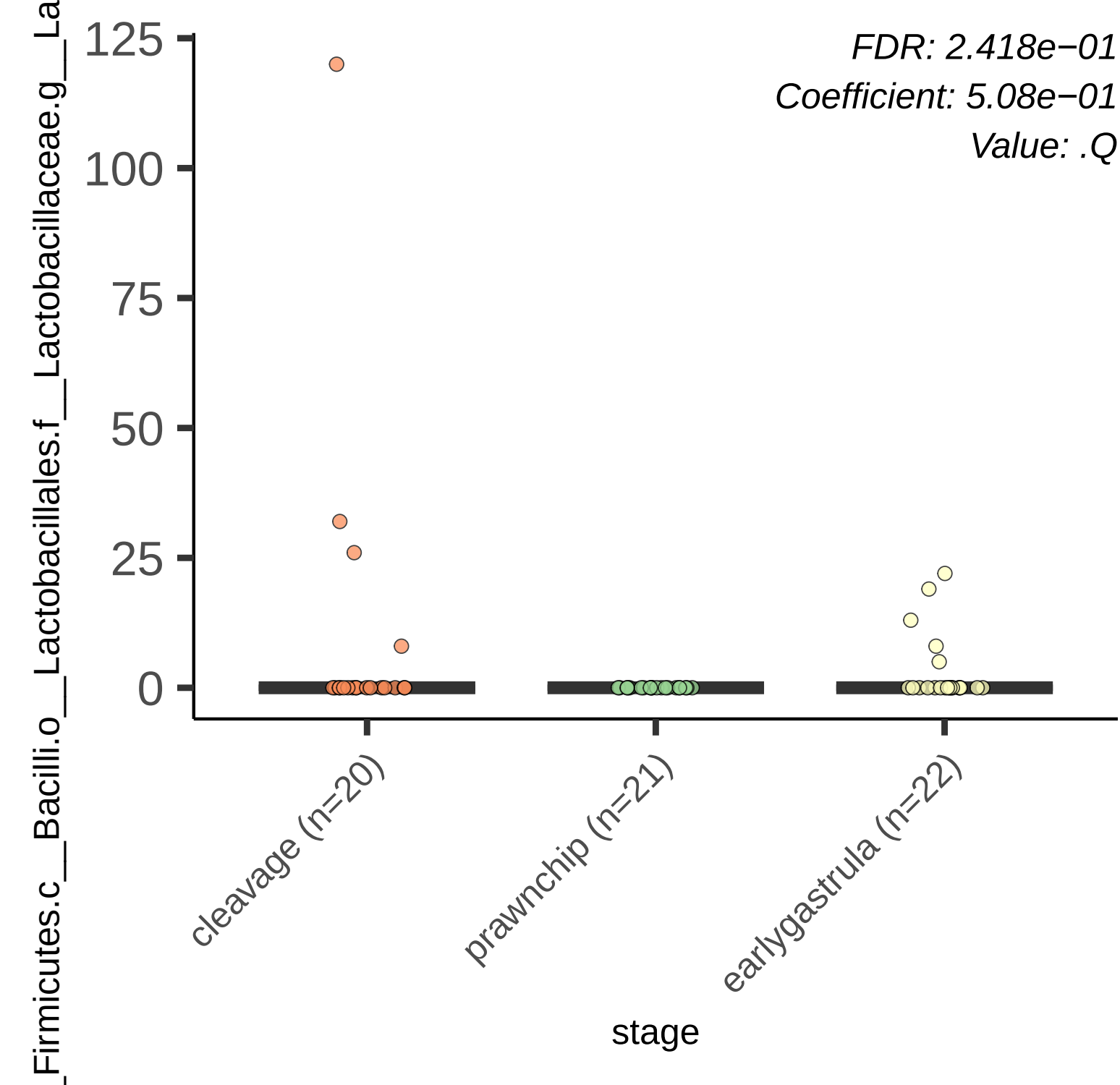
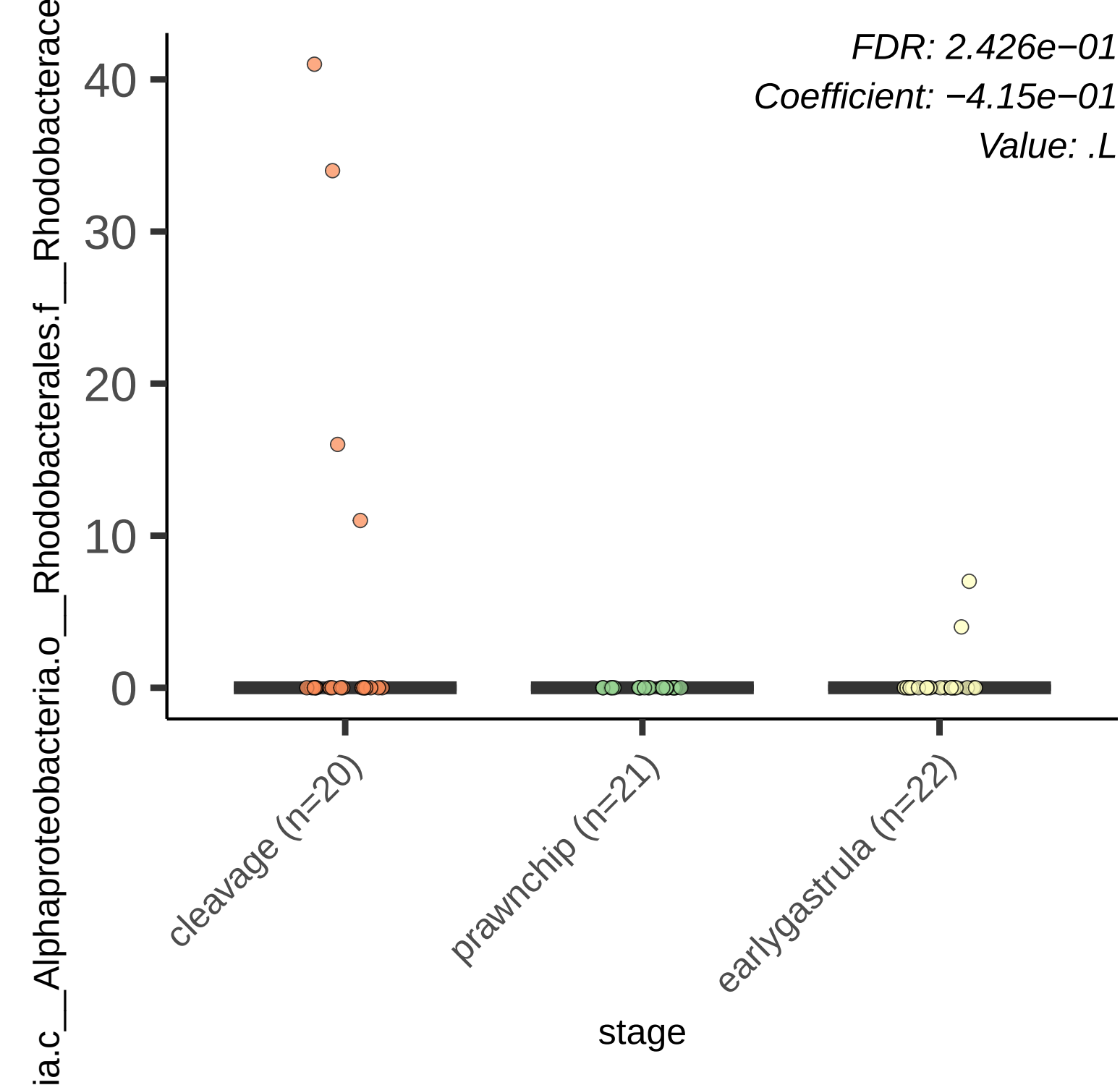
75000
50000
25000
0
cleavage (n=20)
prawnchip (n=21)
earlygastrula (n=22)
stage
a.c__Gammaproteobacteria.o__Alteromonadales.f__Pseudoalterom


FDR: 2.418e−01
Coefficient: 5.08e−01
Value: .Q
125
100
75
50
25
0
cleavage (n=20)
prawnchip (n=21)
earlygastrula (n=22)
stage
Firmicutes.c__Bacilli.o__Lactobacillales.f__Lactobacillaceae.g__La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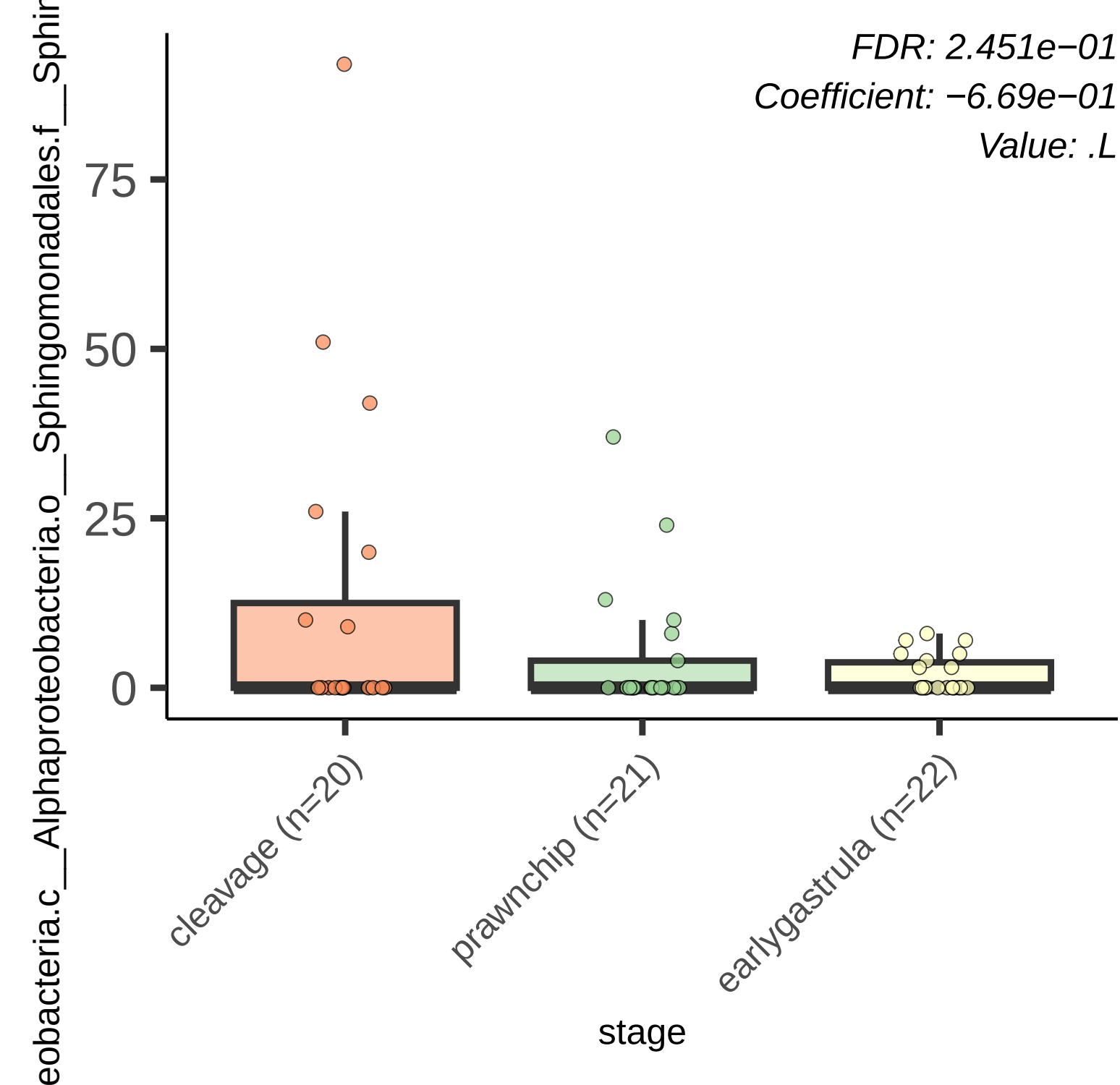
Coefficient: −4.15e−01
Value: .L
40
30
20
10
0
cleavage (n=20)
prawnchip (n=21)
earlygastrula (n=22)
stage
ia.c__Alphaproteobacteria.o__Rhodobacterales.f__Rhodobacterace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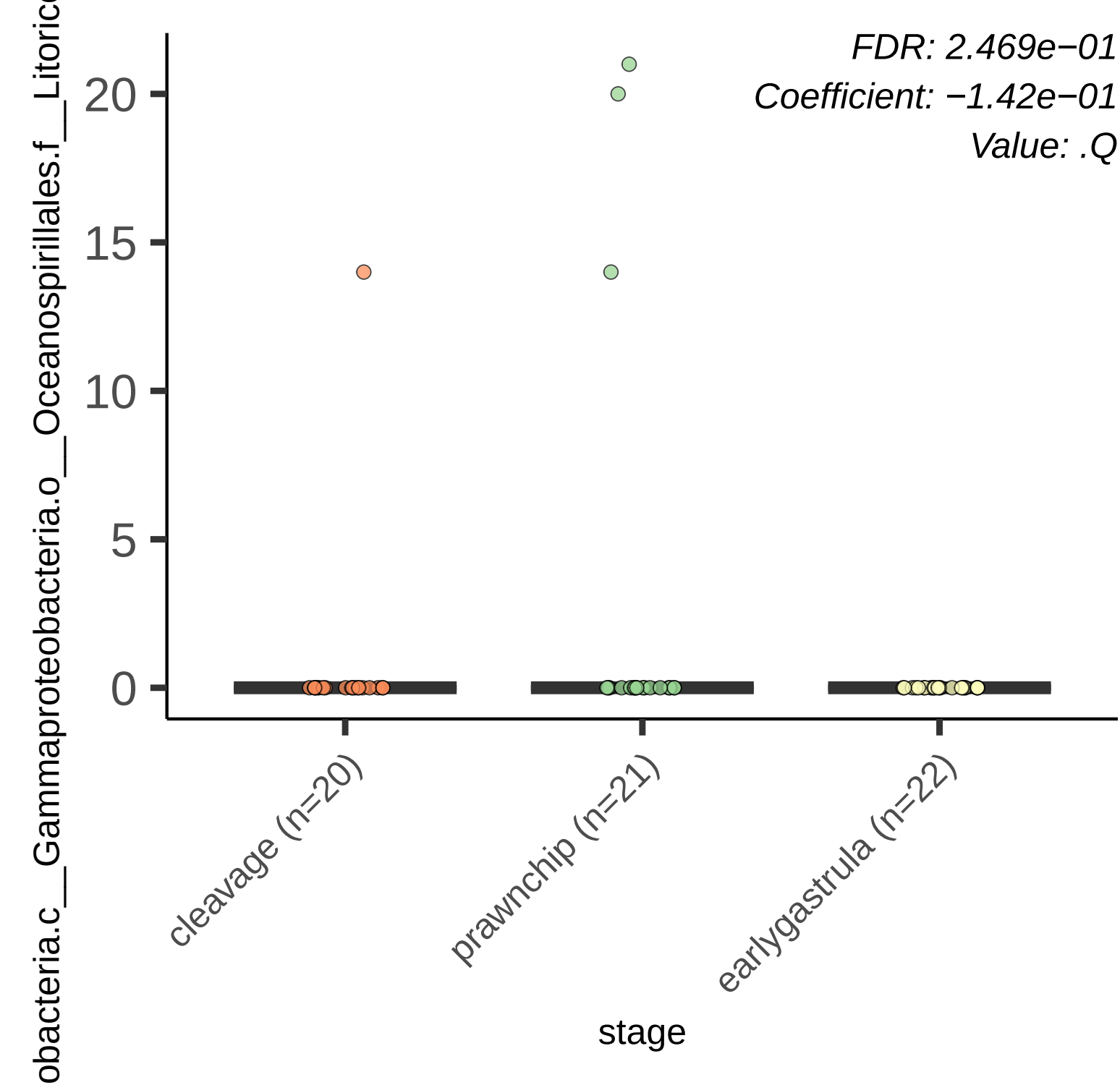
Coefficient: −6.69e−01
Value: .L
75
50
25
0
cleavage (n=20)
prawnchip (n=21)
earlygastrula (n=22)
stage
eobacteria.c__Alphaproteobacteria.o__Sphingomonadales.f__Sphin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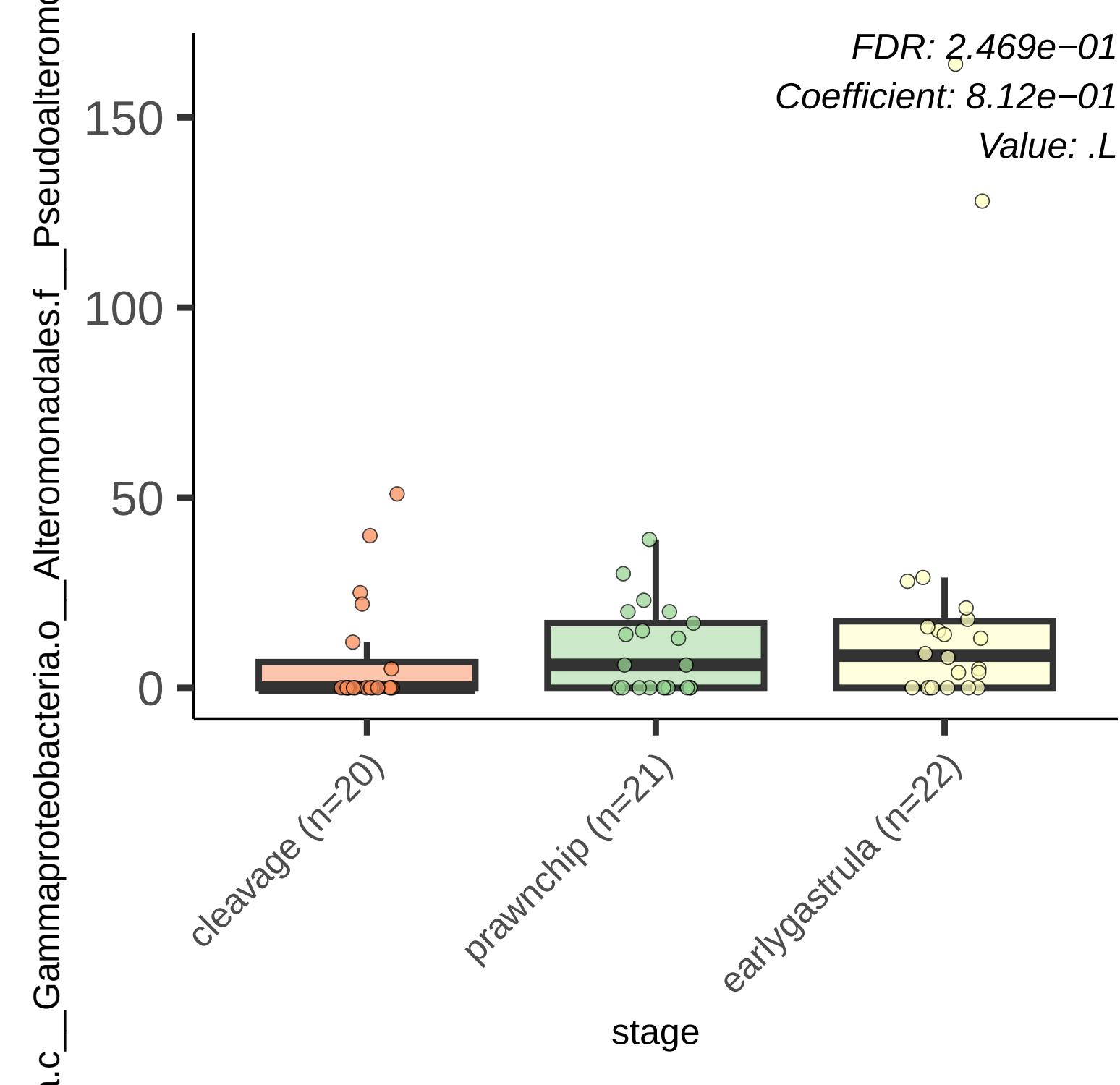
Coefficient: −1.42e−01
Value: .Q
20
15
10
5
0
cleavage (n=20)
prawnchip (n=21)
earlygastrula (n=22)
stage
obacteria.c__Gammaproteobacteria.o__Oceanospirillales.f__Litoric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Coefficient: 8.12e−01
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150
100
50
0
cleavage (n=20)
prawnchip (n=21)
earlygastrula (n=22)
stage
c__Gammaproteobacteria.o__Alteromonadales.f__Pseudoalteromo