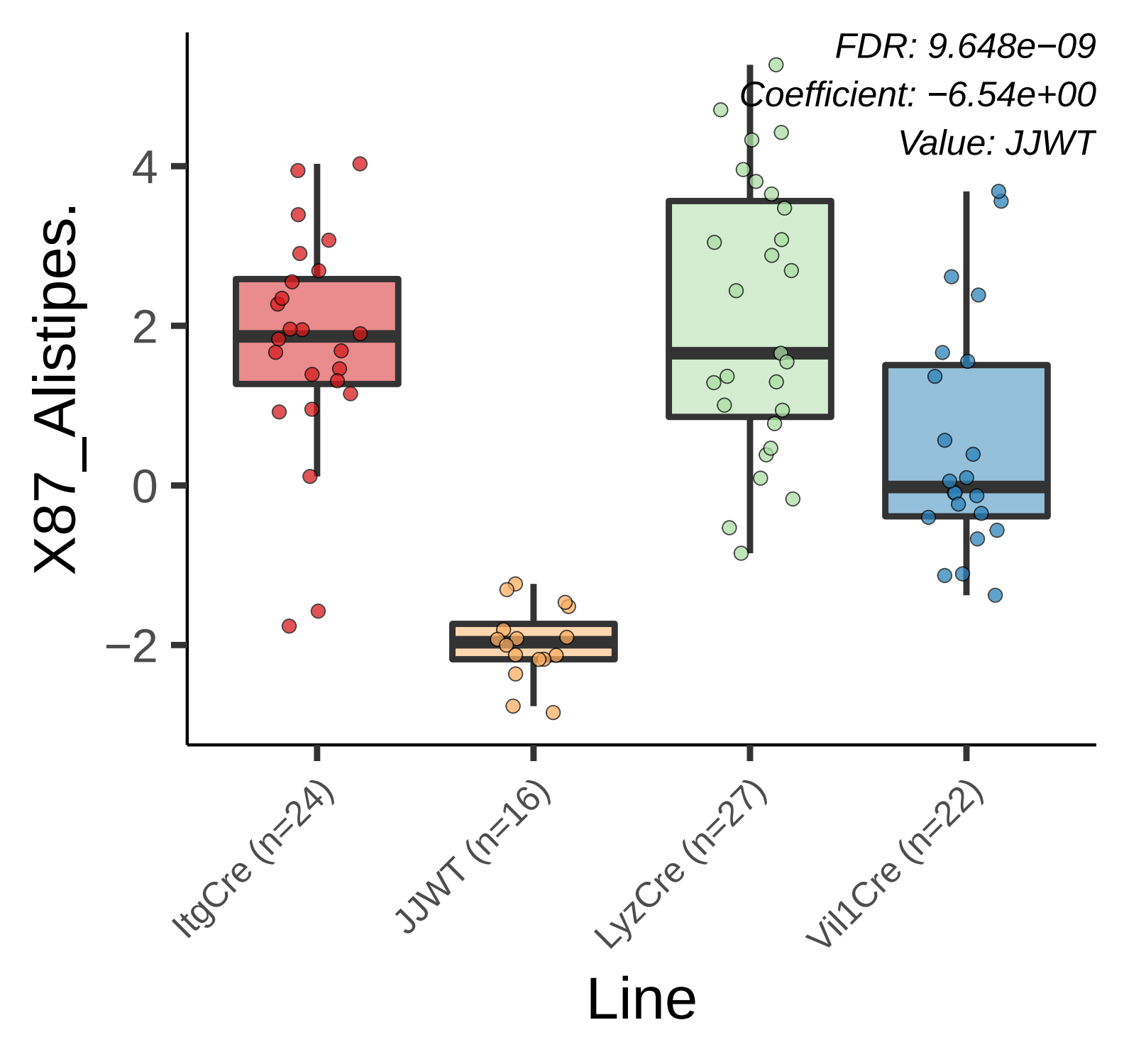
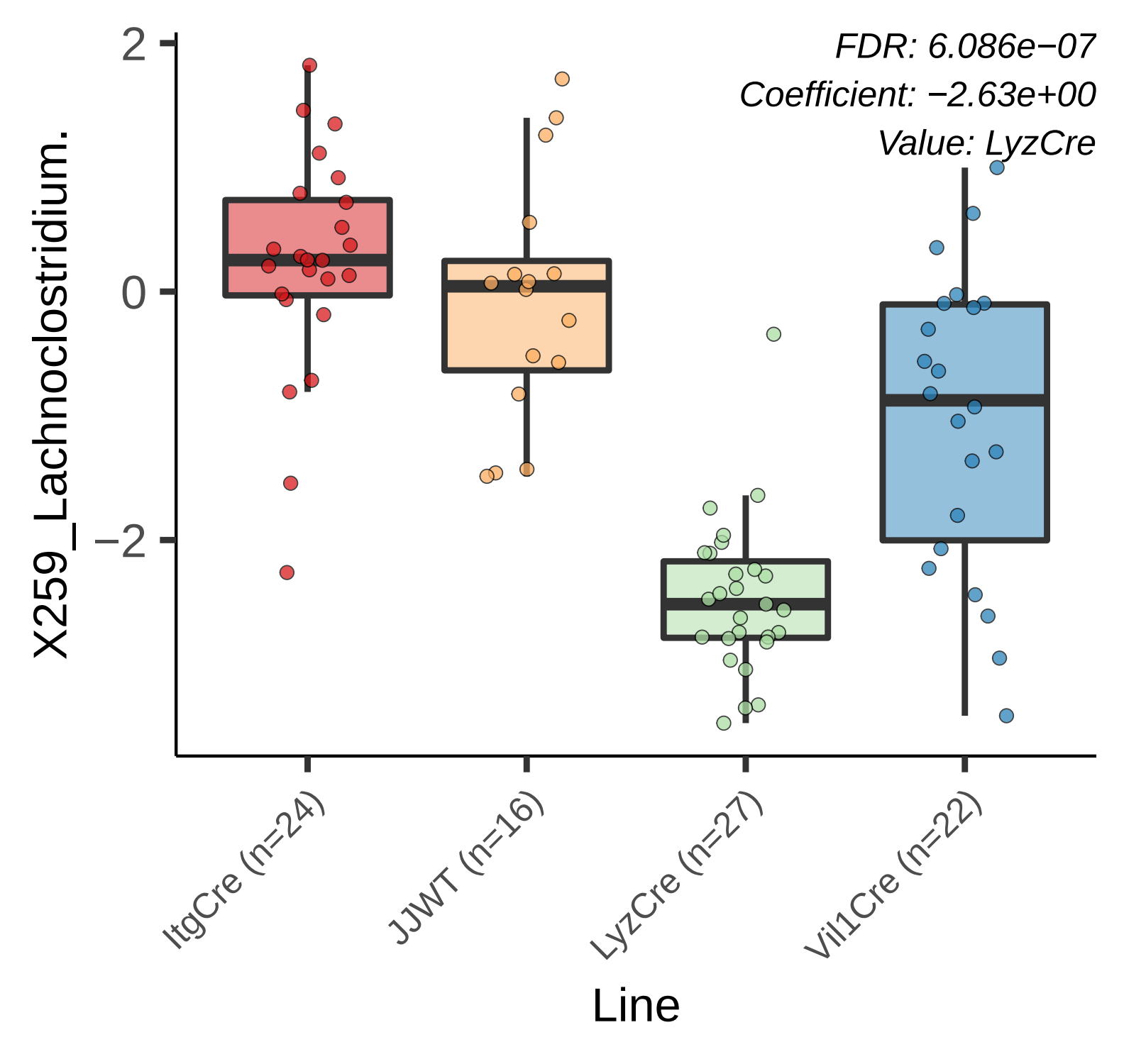
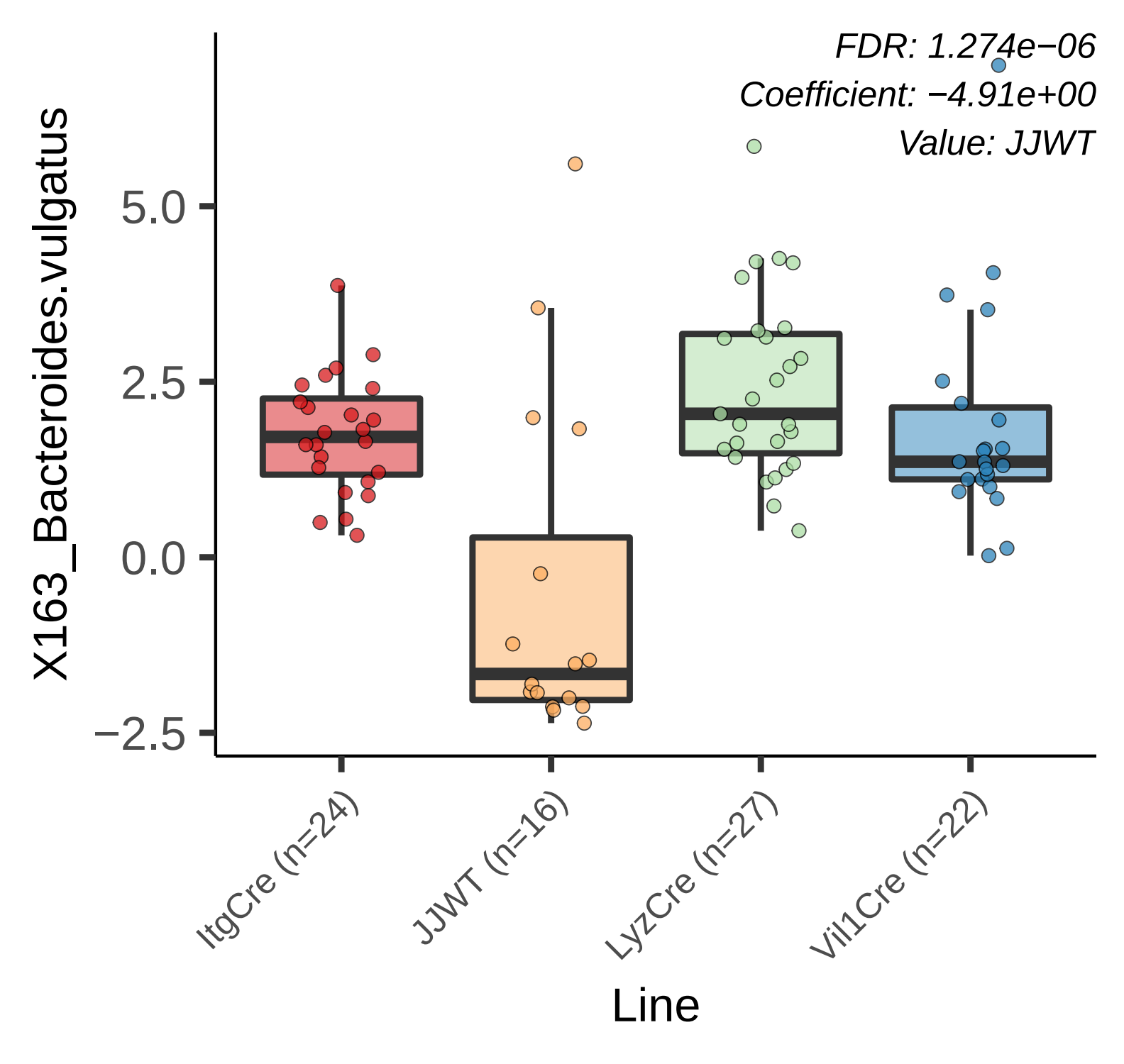
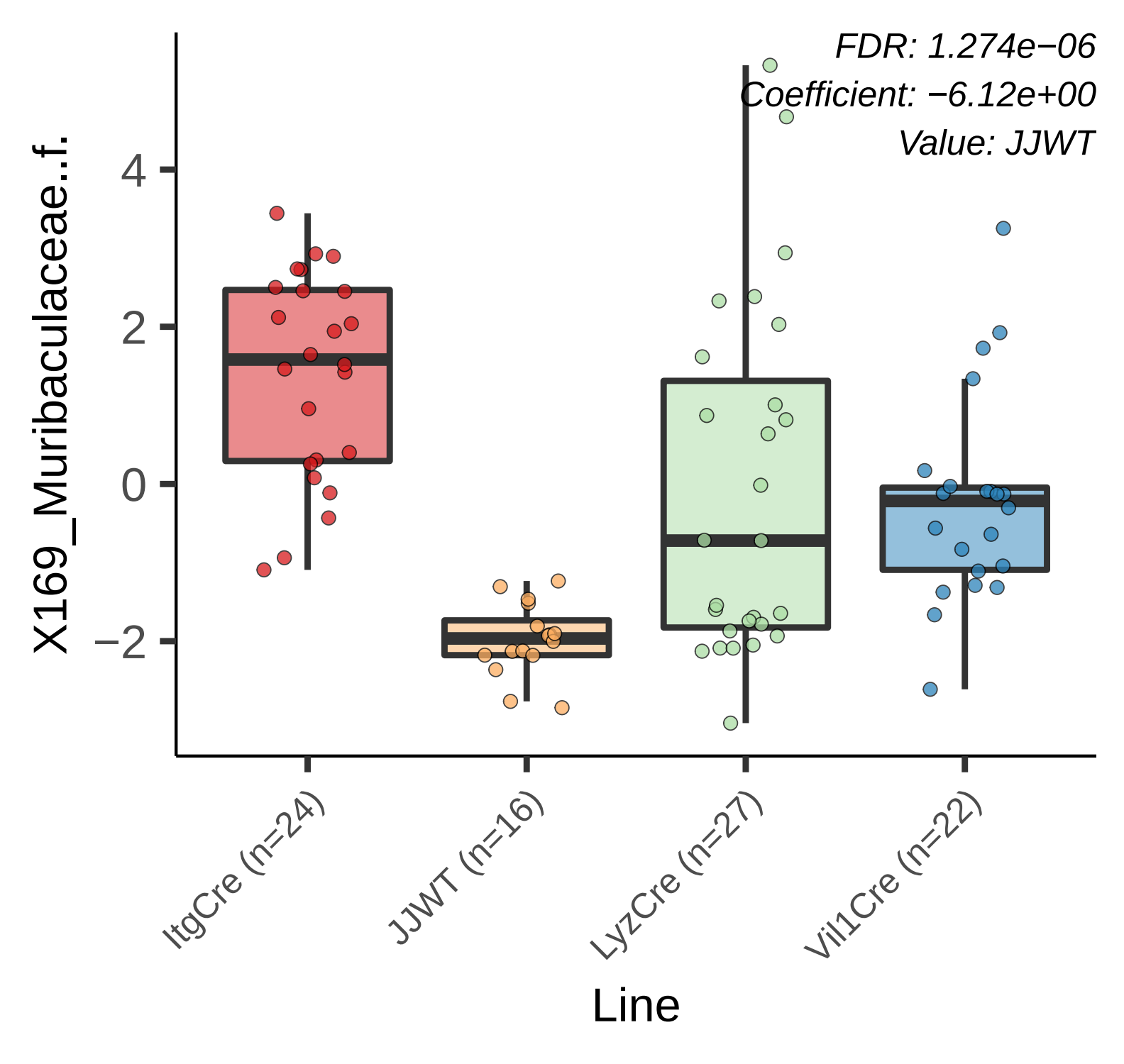
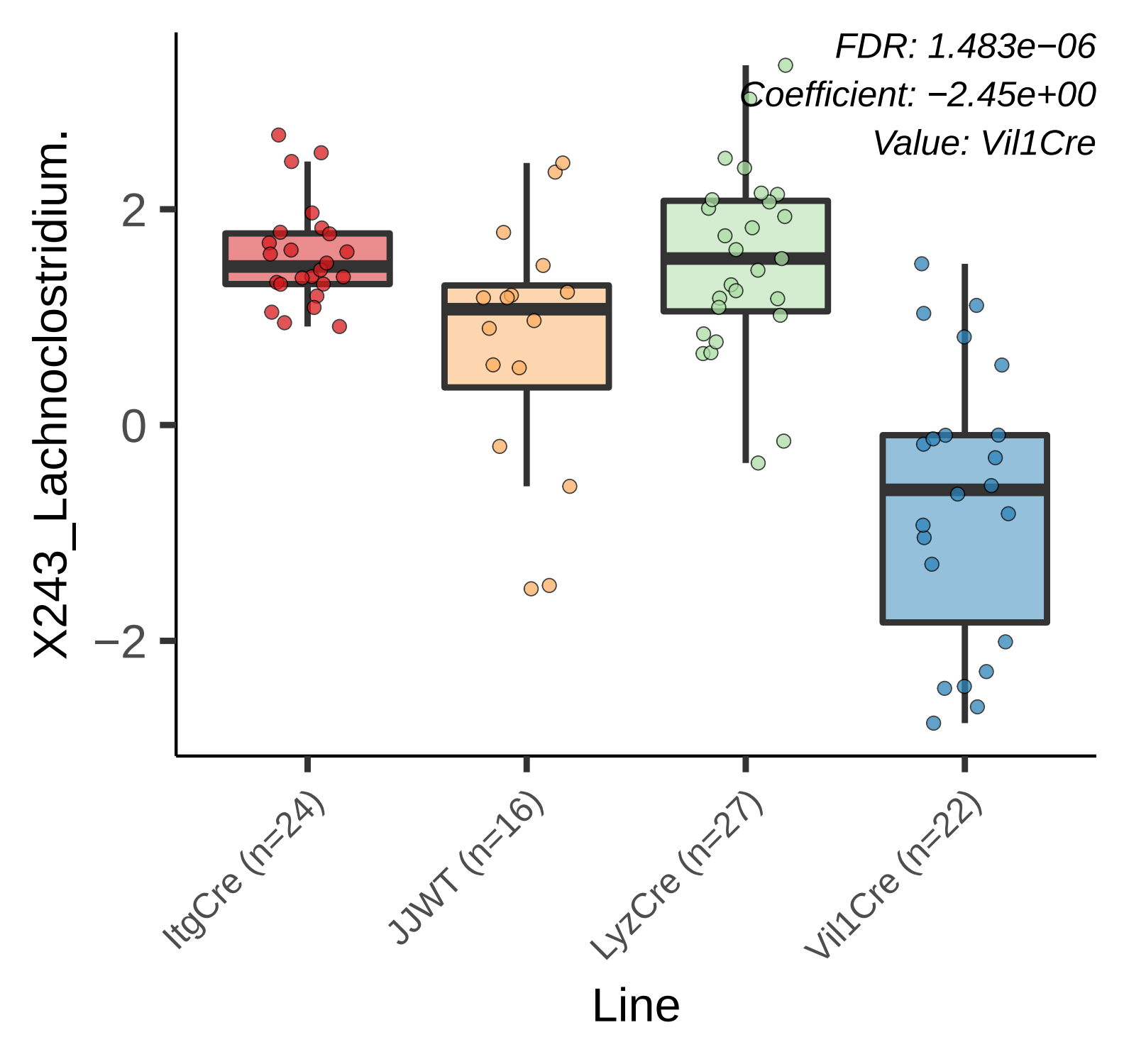
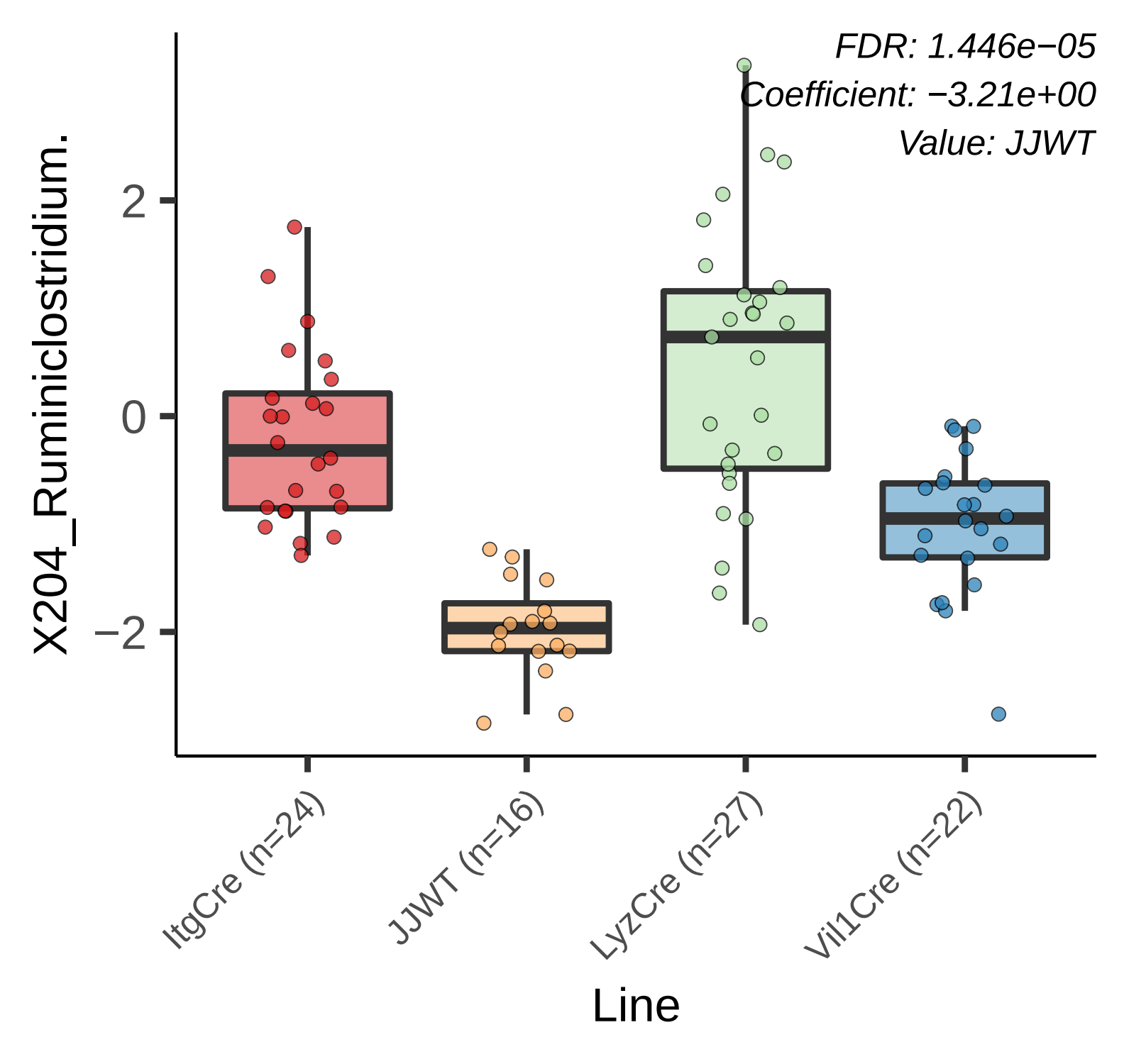

FDR: 9.648e−09
Coefficient: −6.54e+00
Value: JJWT
X87_Alistipes.
4
2
0
−2
ItgCre (n=24)
JJWT (n=16)
LyzCre (n=27)
Vil1Cre (n=22)
Line

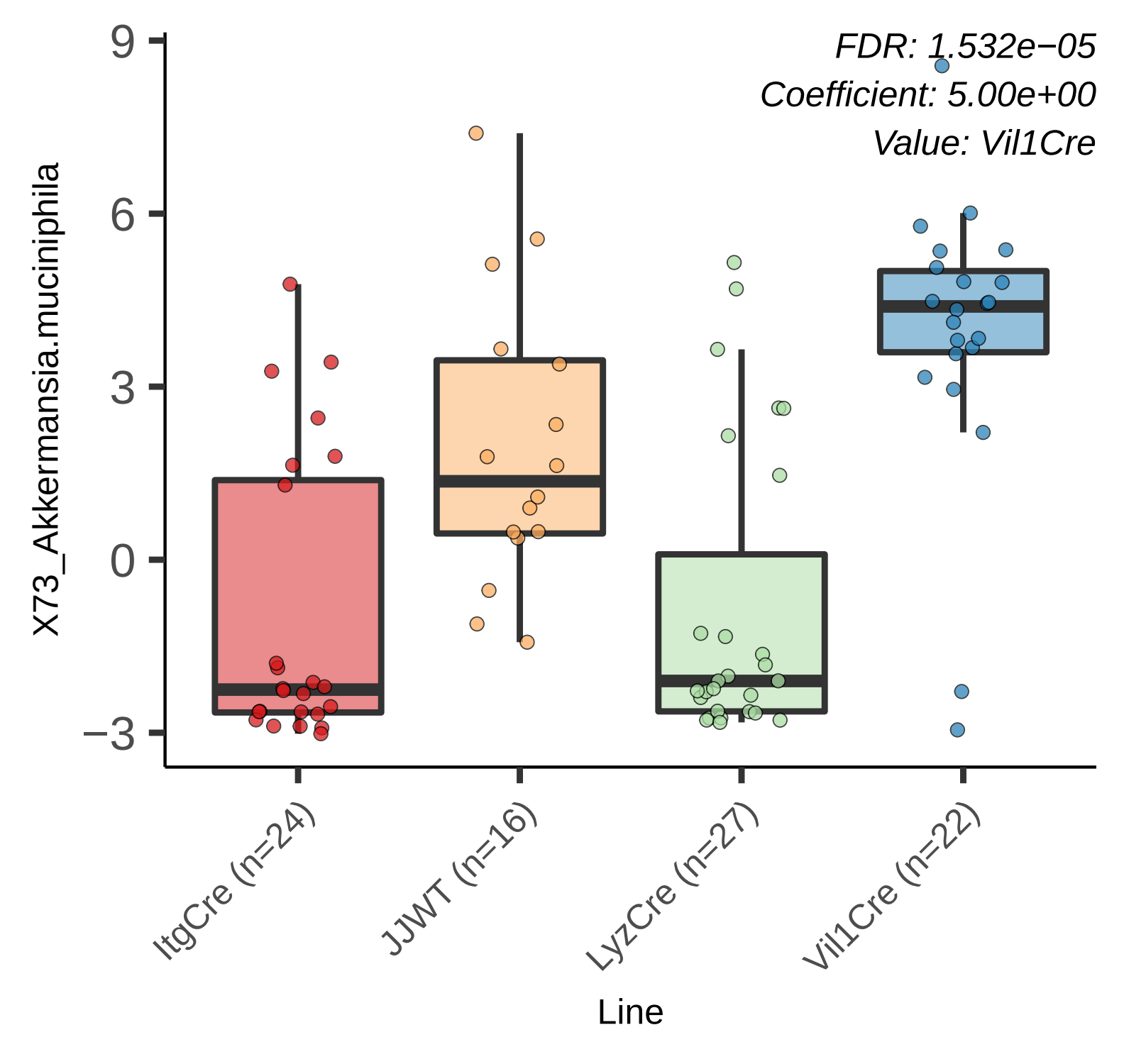

FDR: 6.086e−07
Coefficient: −2.63e+00
Value: LyzCre
X259_Lachnoclostridium.
2
0
−2
ItgCre (n=24)
JJWT (n=16)
LyzCre (n=27)
Vil1Cre (n=22)
Line

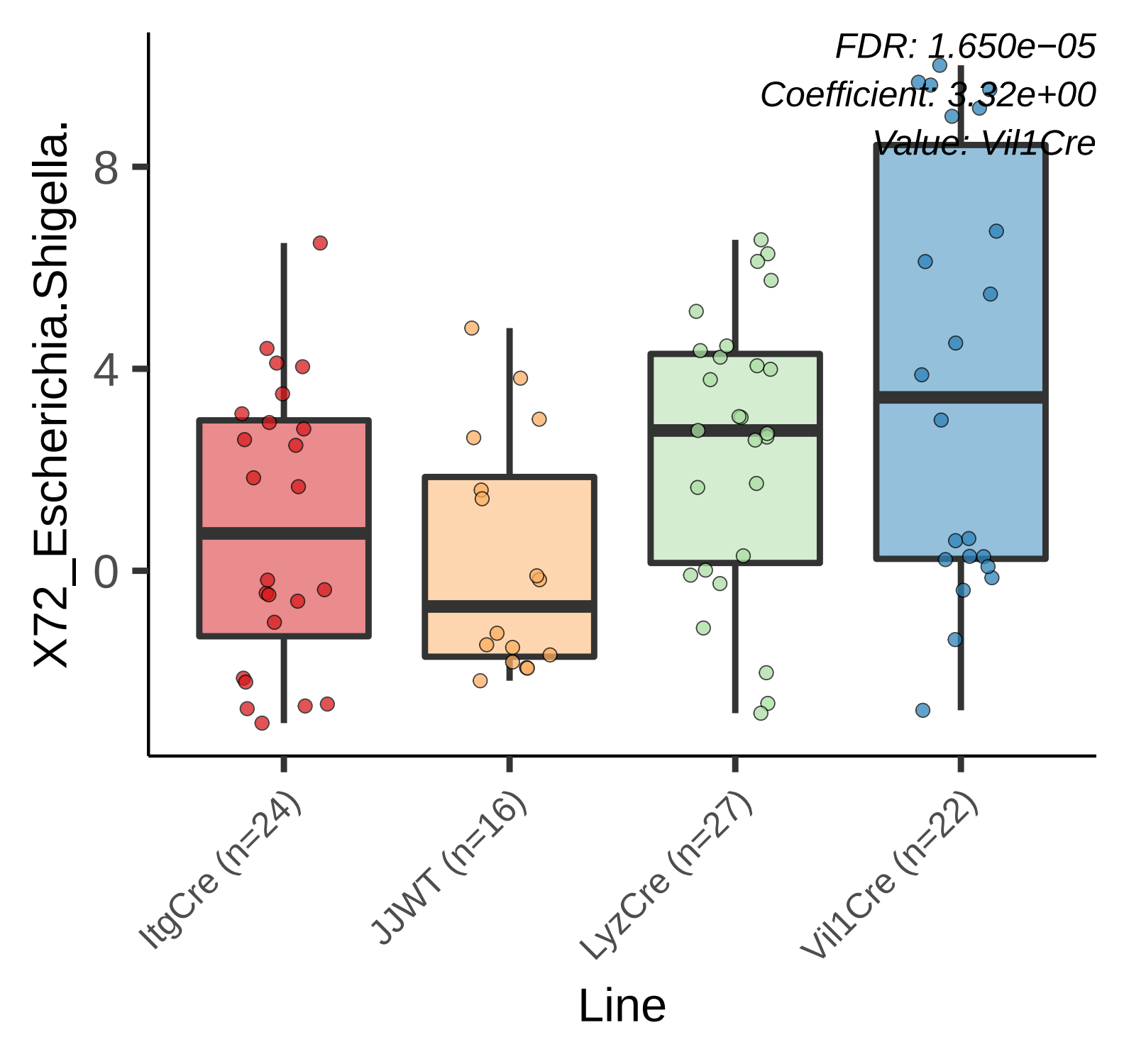

FDR: 1.274e−06
Coefficient: −4.91e+00
Value: JJWT
X163_Bacteroides.vulgatus
5.0
2.5
0.0
−2.5
ItgCre (n=24)
JJWT (n=16)
LyzCre (n=27)
Vil1Cre (n=22)
Line

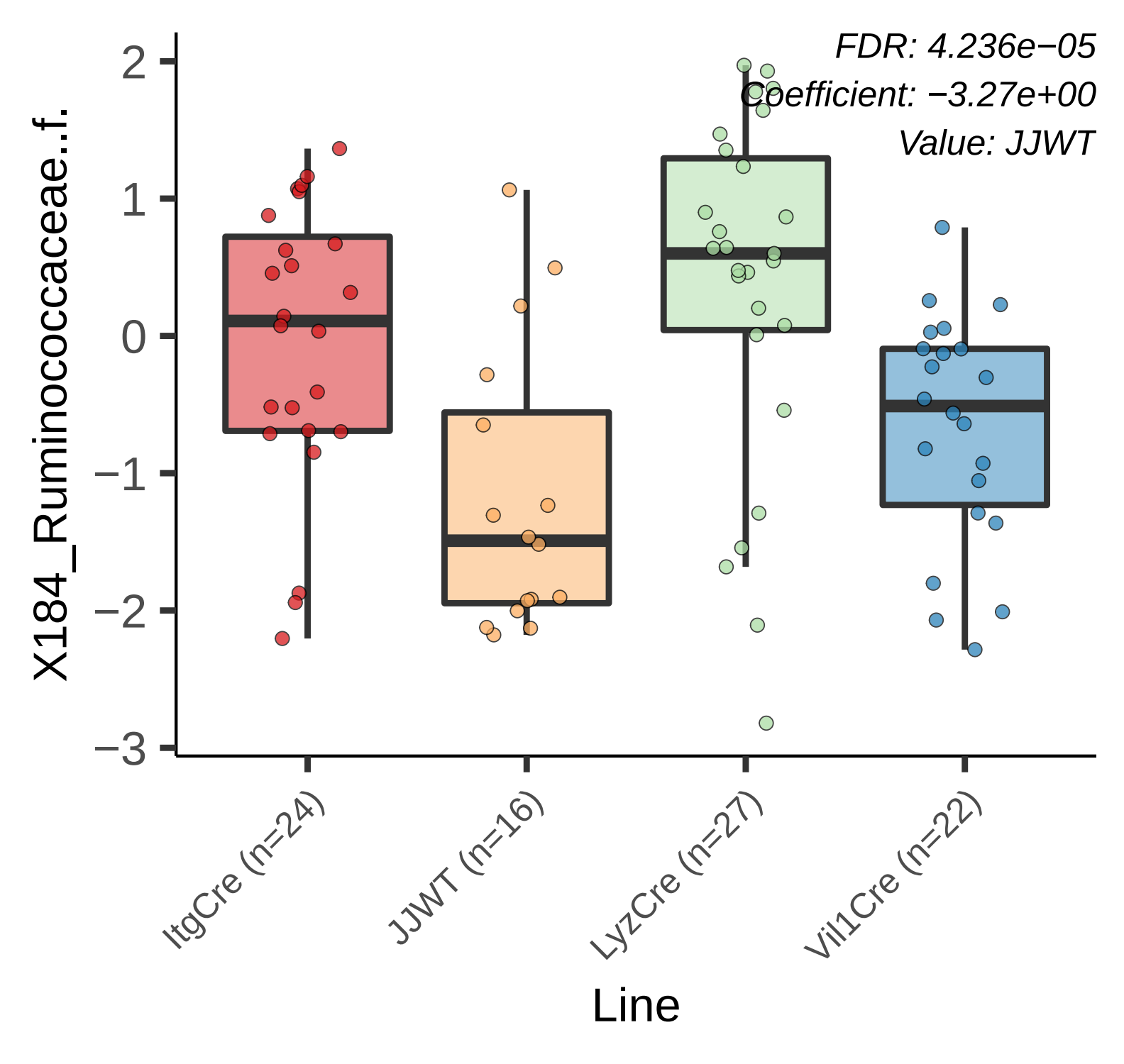

FDR: 1.274e−06
Coefficient: −6.12e+00
Value: JJWT
X169_Muribaculaceae..f.
4
2
0
−2
ItgCre (n=24)
JJWT (n=16)
LyzCre (n=27)
Vil1Cre (n=22)
Line

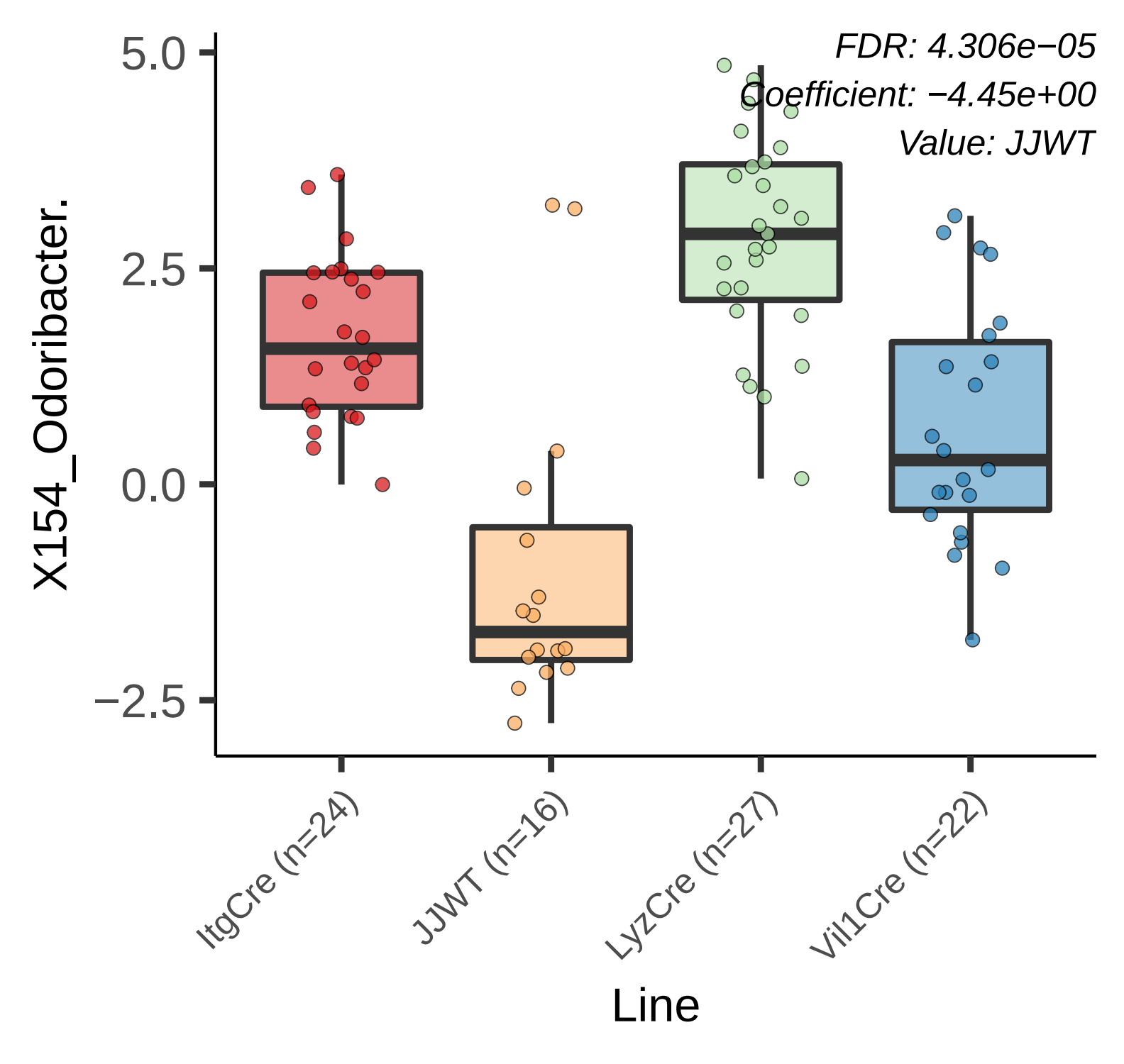

FDR: 1.483e−06
Coefficient: −2.45e+00
Value: Vil1Cre
X243_Lachnoclostridium.
2
0
−2
ItgCre (n=24)
JJWT (n=16)
LyzCre (n=27)
Vil1Cre (n=22)
Line

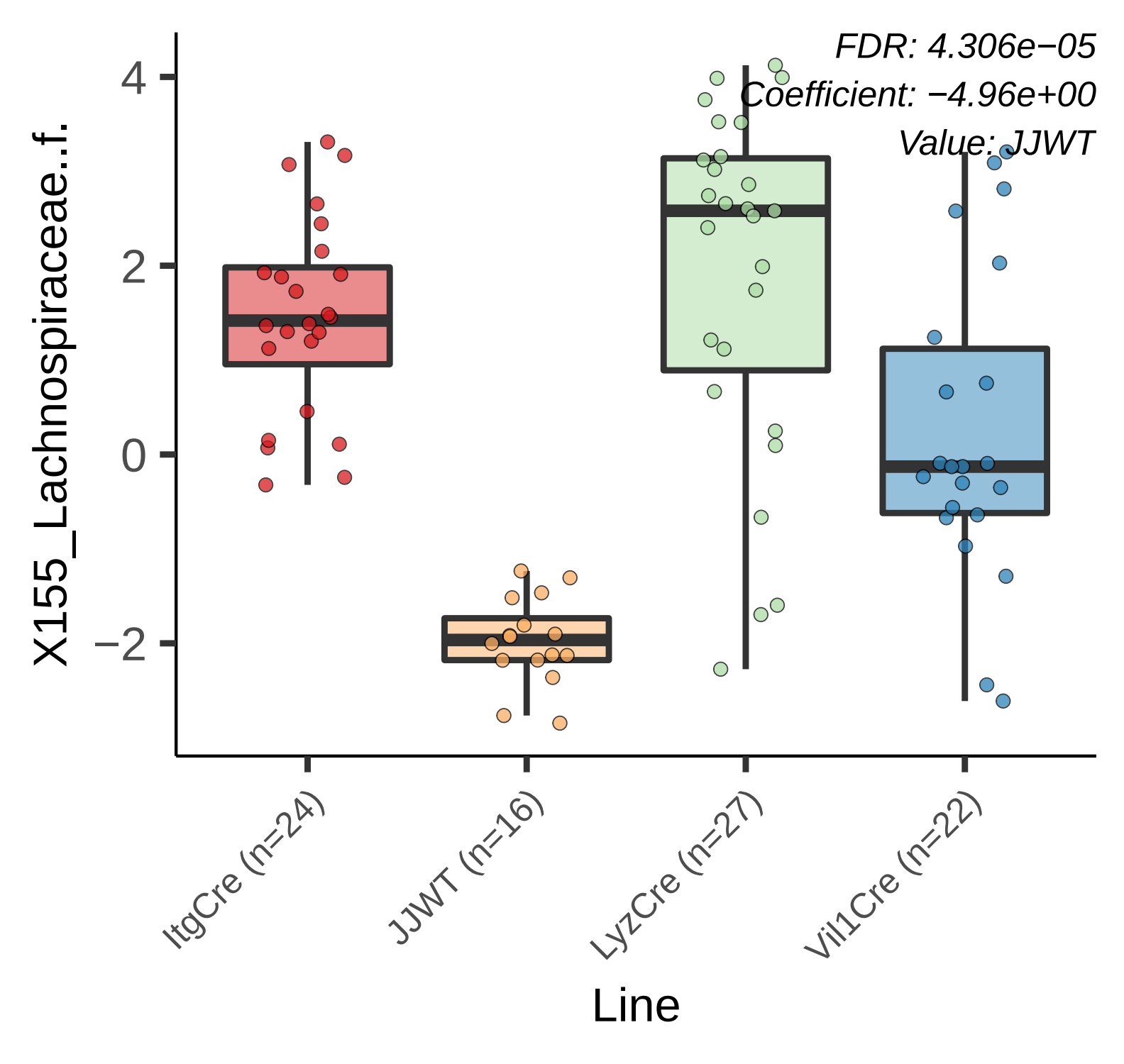

FDR: 1.446e−05
Coefficient: −3.21e+00
Value: JJWT
X204_Ruminiclostridium.
2
0
−2
ItgCre (n=24)
JJWT (n=16)
LyzCre (n=27)
Vil1Cre (n=22)
Line

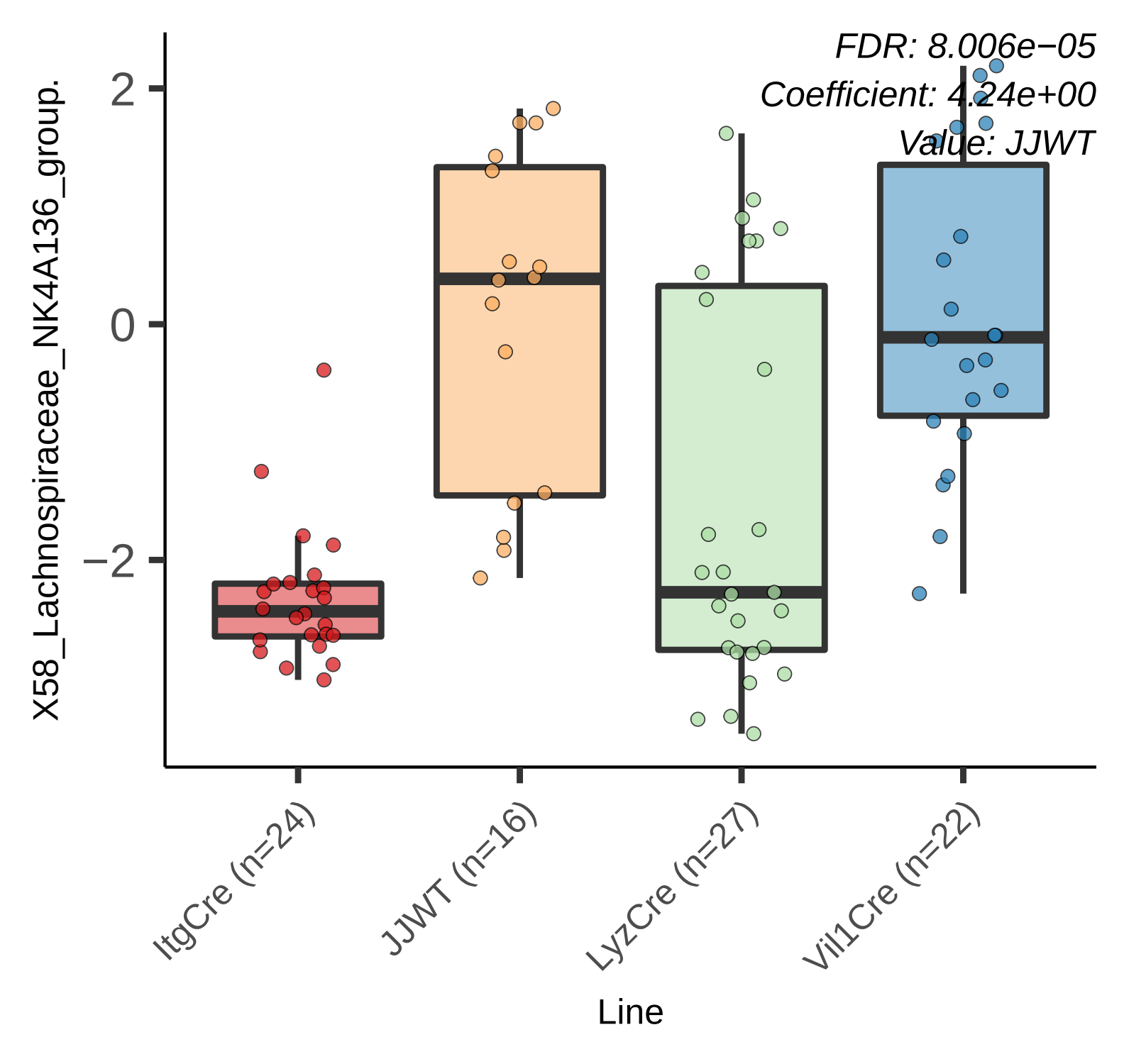

FDR: 1.532e−05
Coefficient: 5.00e+00
Value: Vil1Cre
X73_Akkermansia.muciniphila
9
6
3
0
−3
ItgCre (n=24)
JJWT (n=16)
LyzCre (n=27)
Vil1Cre (n=22)
Line

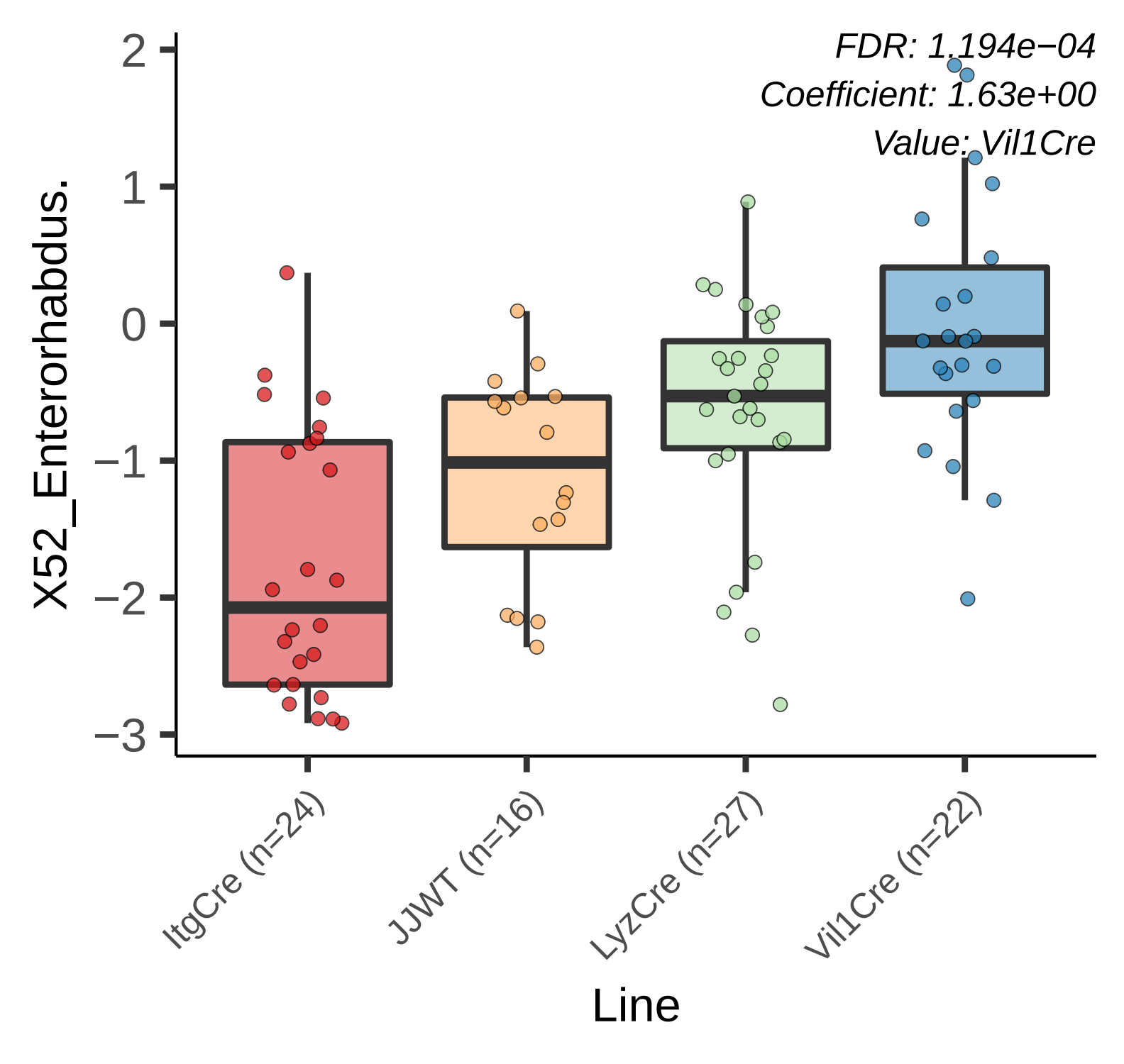

FDR: 1.650e−05
Coefficient: 3.32e+00
Value: Vil1Cre
X72_Escherichia.Shigella.
8
4
0
ItgCre (n=24)
JJWT (n=16)
LyzCre (n=27)
Vil1Cre (n=22)
Line

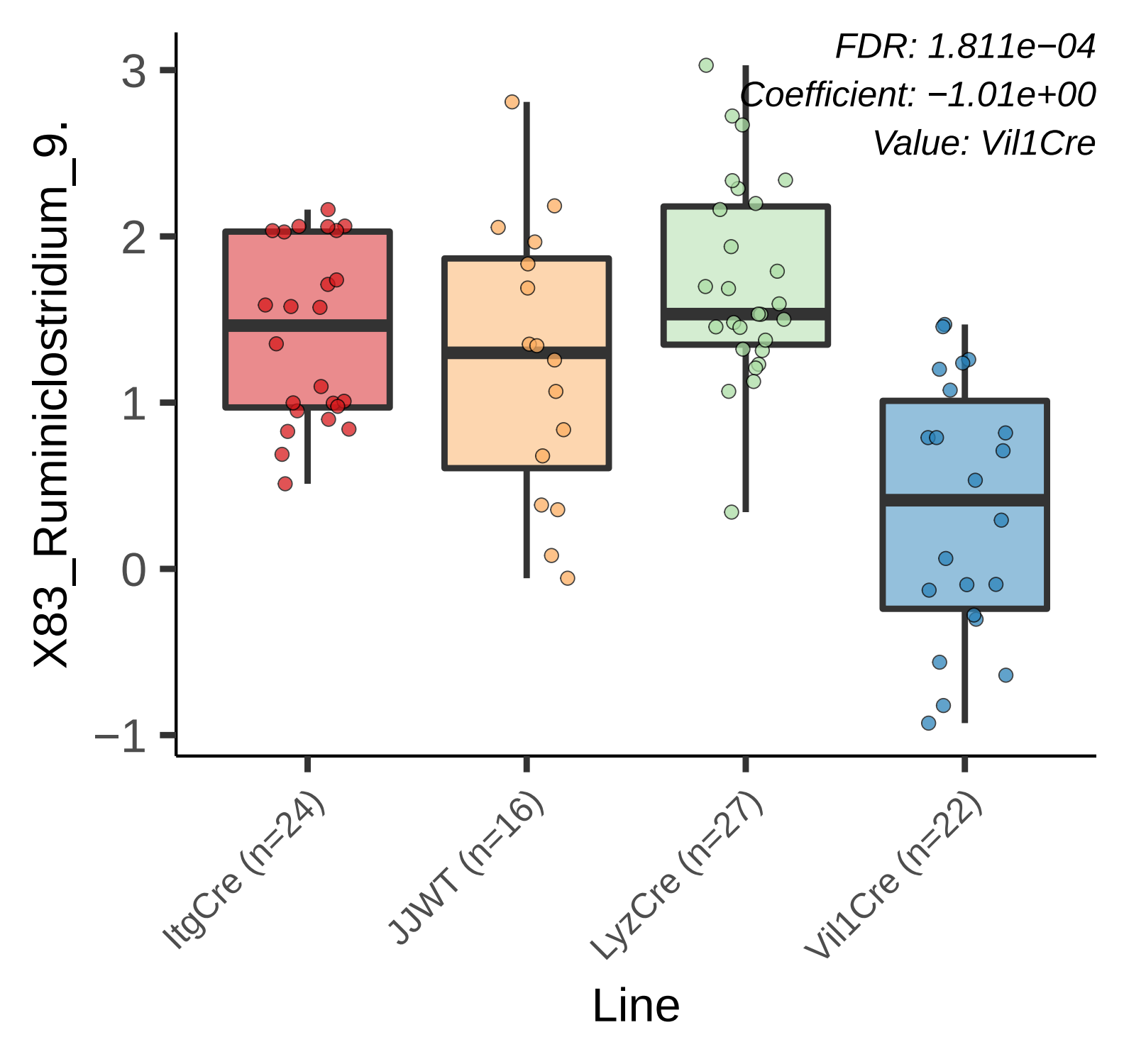

FDR: 4.236e−05
Coefficient: −3.27e+00
Value: JJWT
X184_Ruminococcaceae..f.
2
1
0
−1
−2
−3
ItgCre (n=24)
JJWT (n=16)
LyzCre (n=27)
Vil1Cre (n=22)
Line

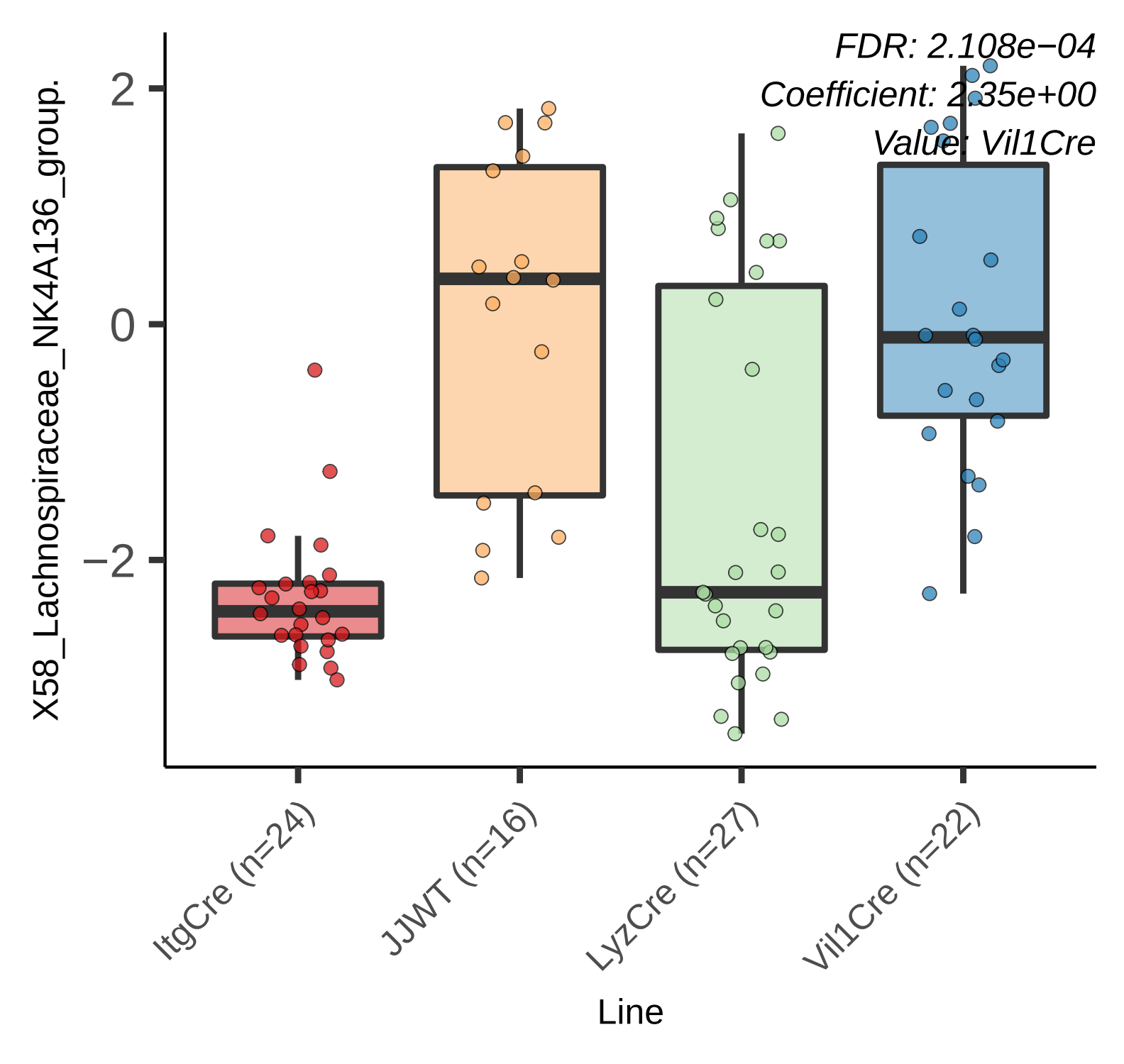

FDR: 4.306e−05
Coefficient: −4.45e+00
Value: JJWT
X154_Odoribacter.
5.0
2.5
0.0
−2.5
ItgCre (n=24)
JJWT (n=16)
LyzCre (n=27)
Vil1Cre (n=22)
Line

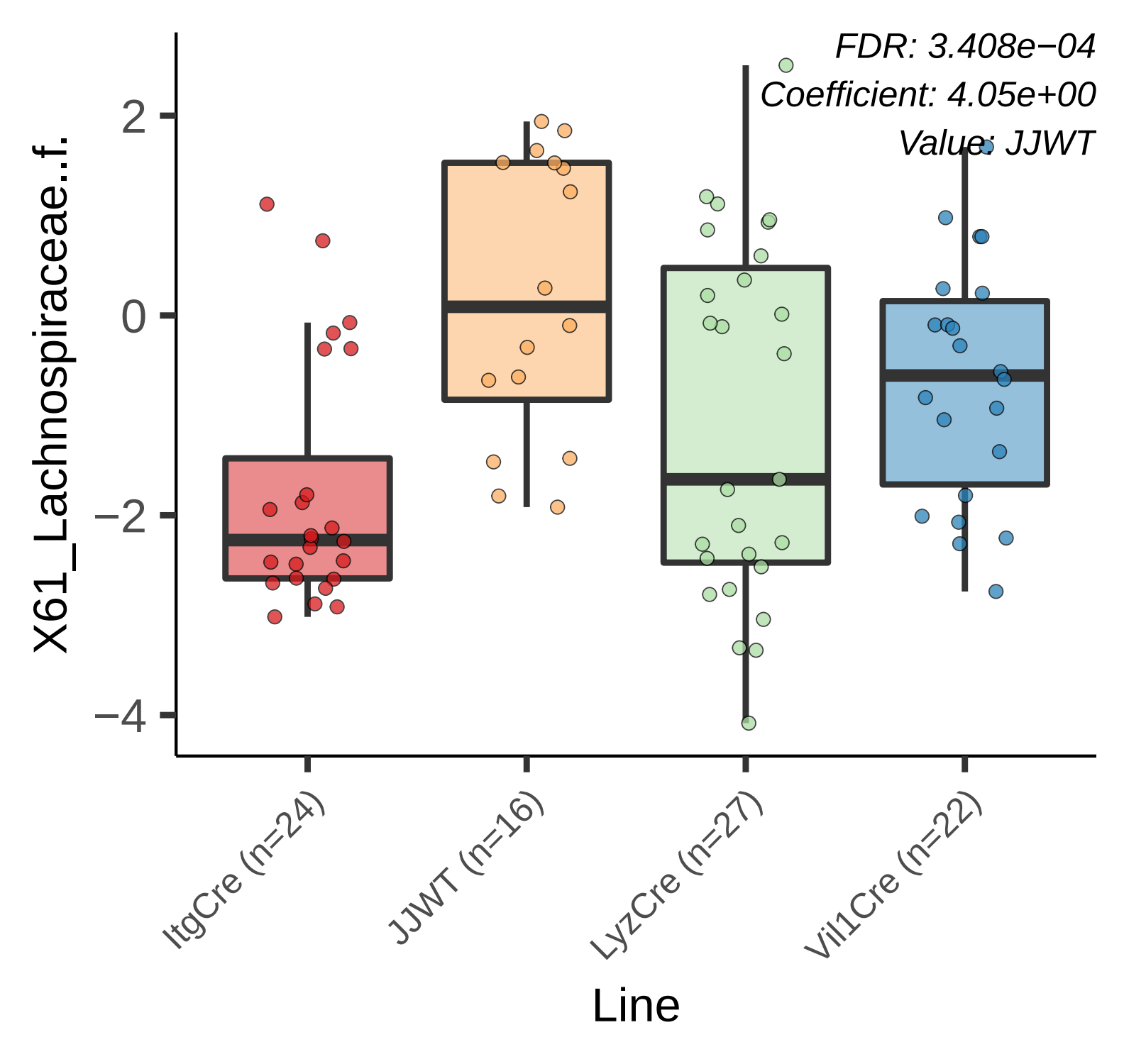

FDR: 4.306e−05
Coefficient: −4.96e+00
Value: JJWT
X155_Lachnospiraceae.f.
4
2
0
−2
ItgCre (n=24)
JJWT (n=16)
LyzCre (n=27)
Vil1Cre (n=22)
Line

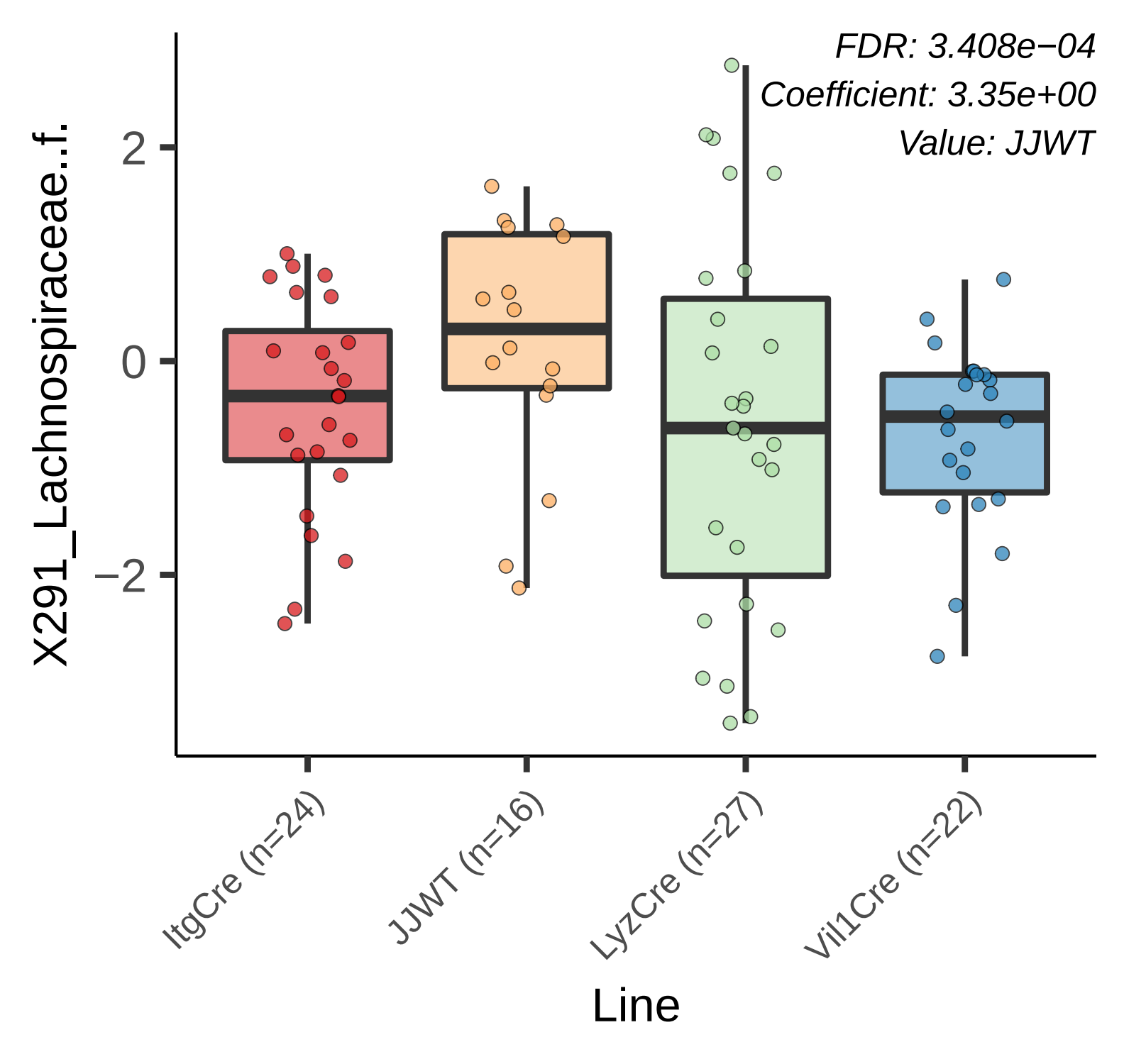

FDR: 8.006e−05
Coefficient: 4.24e+00
Value: JJWT
X58_Lachnospiraceae_NK4A136_group.
2
0
−2
ItgCre (n=24)
JJWT (n=16)
LyzCre (n=27)
Vil1Cre (n=22)
Line

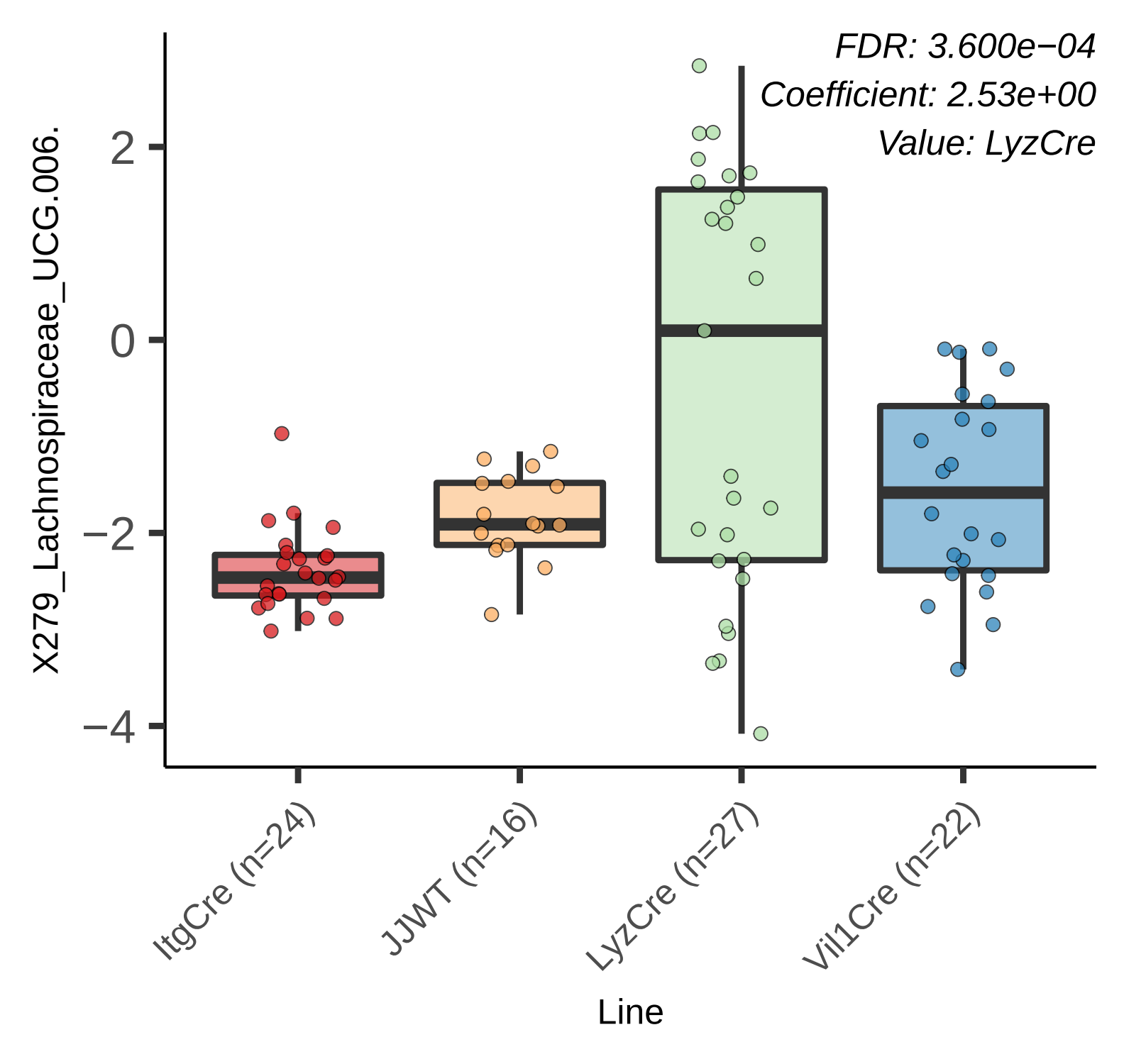

FDR: 1.194e−04
Coefficient: 1.63e+00
Value: Vil1Cre
X52_Enterorhabdus.
2
1
0
−1
−2
−3
ItgCre (n=24)
JJWT (n=16)
LyzCre (n=27)
Vil1Cre (n=22)
Line

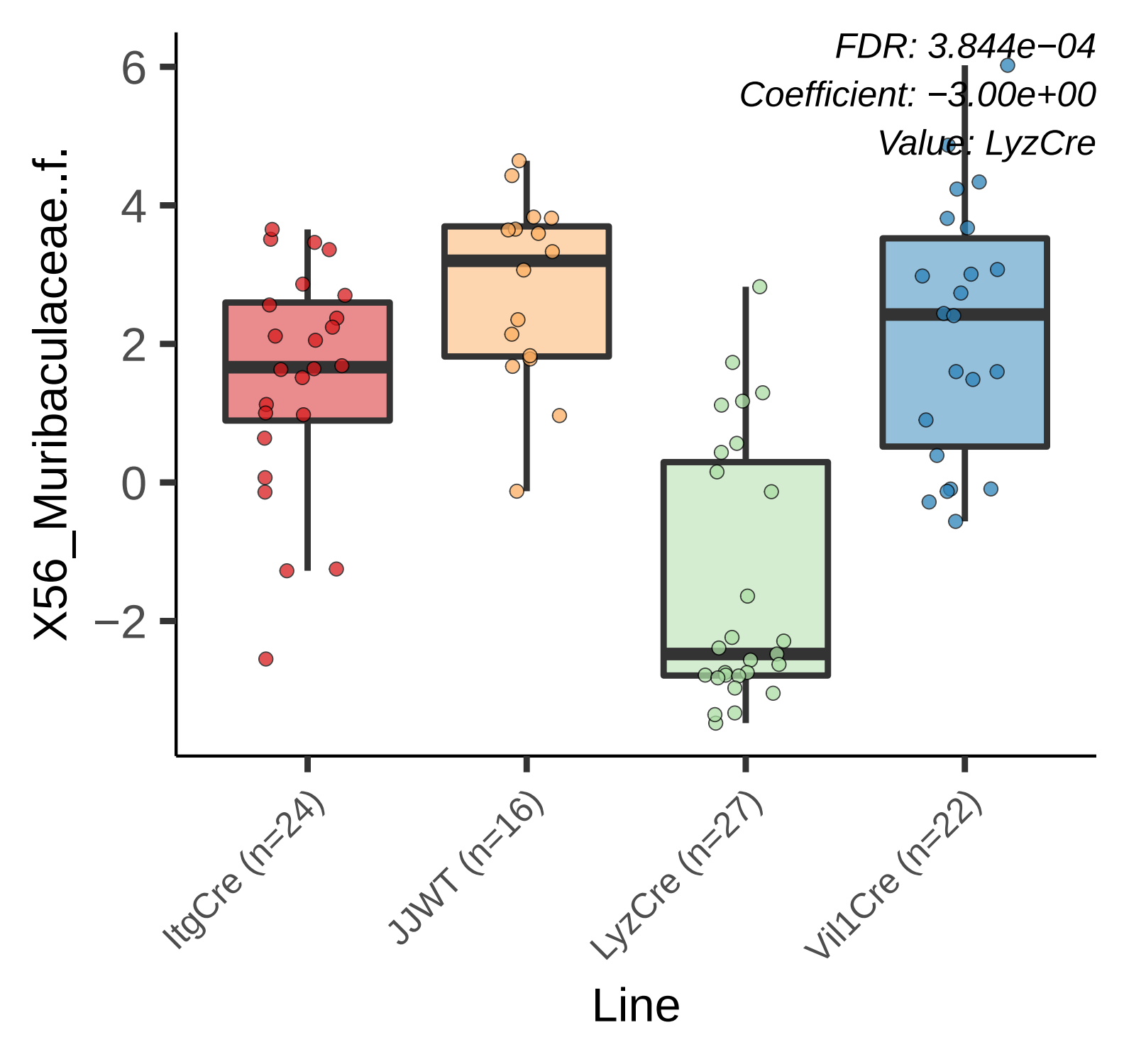

FDR: 1.811e−04
Coefficient: −1.01e+00
Value: Vil1Cre
X83_Ruminiclostridium_9.
3
2
1
0
−1
ItgCre (n=24)
JJWT (n=16)
LyzCre (n=27)
Vil1Cre (n=22)
Line

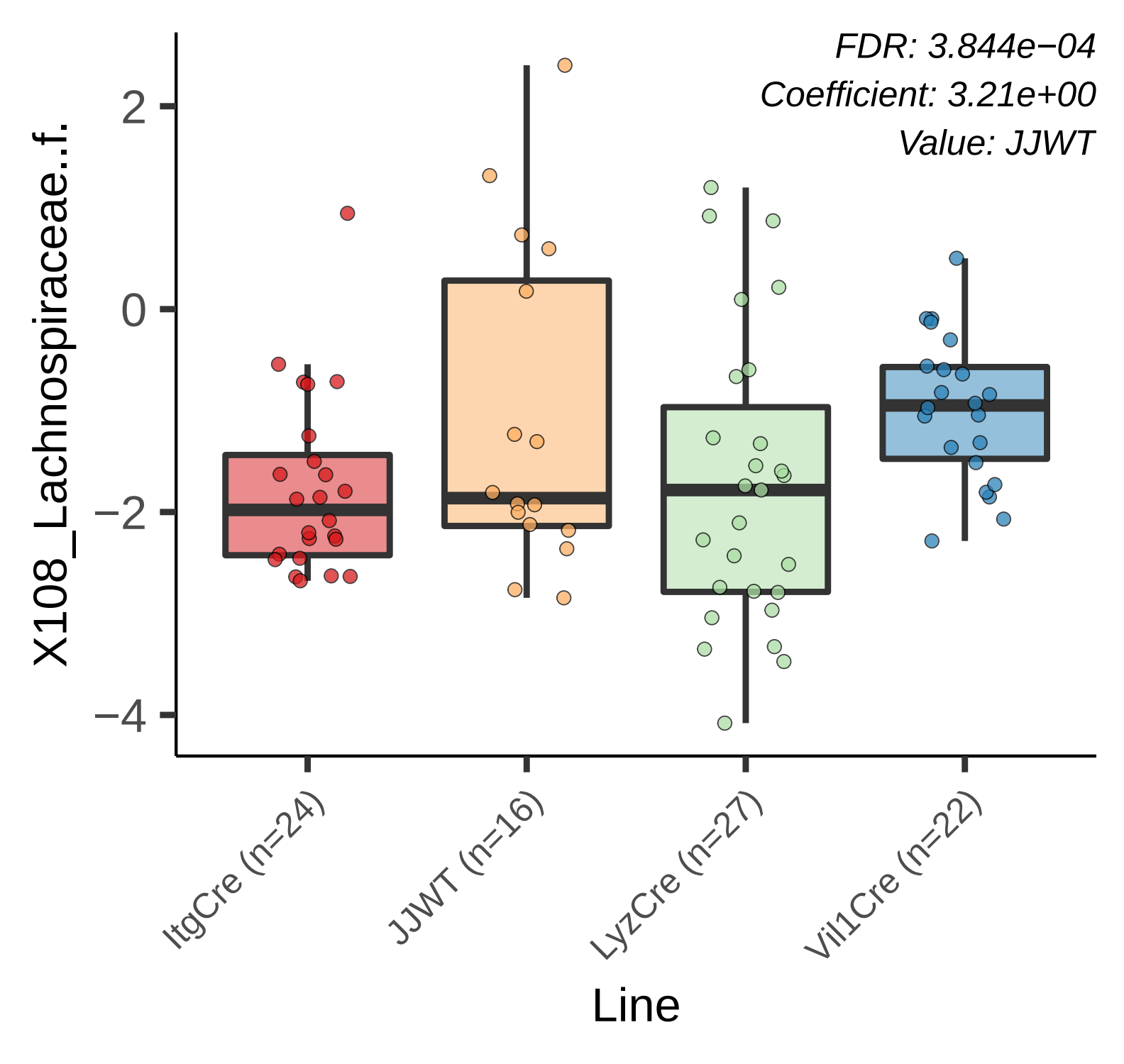

FDR: 2.108e−04
Coefficient: 2.35e+00
Value: Vil1Cre
X58_Lachnospiraceae_NK4A136_group.
2
0
−2
ItgCre (n=24)
JJWT (n=16)
LyzCre (n=27)
Vil1Cre (n=22)
Line

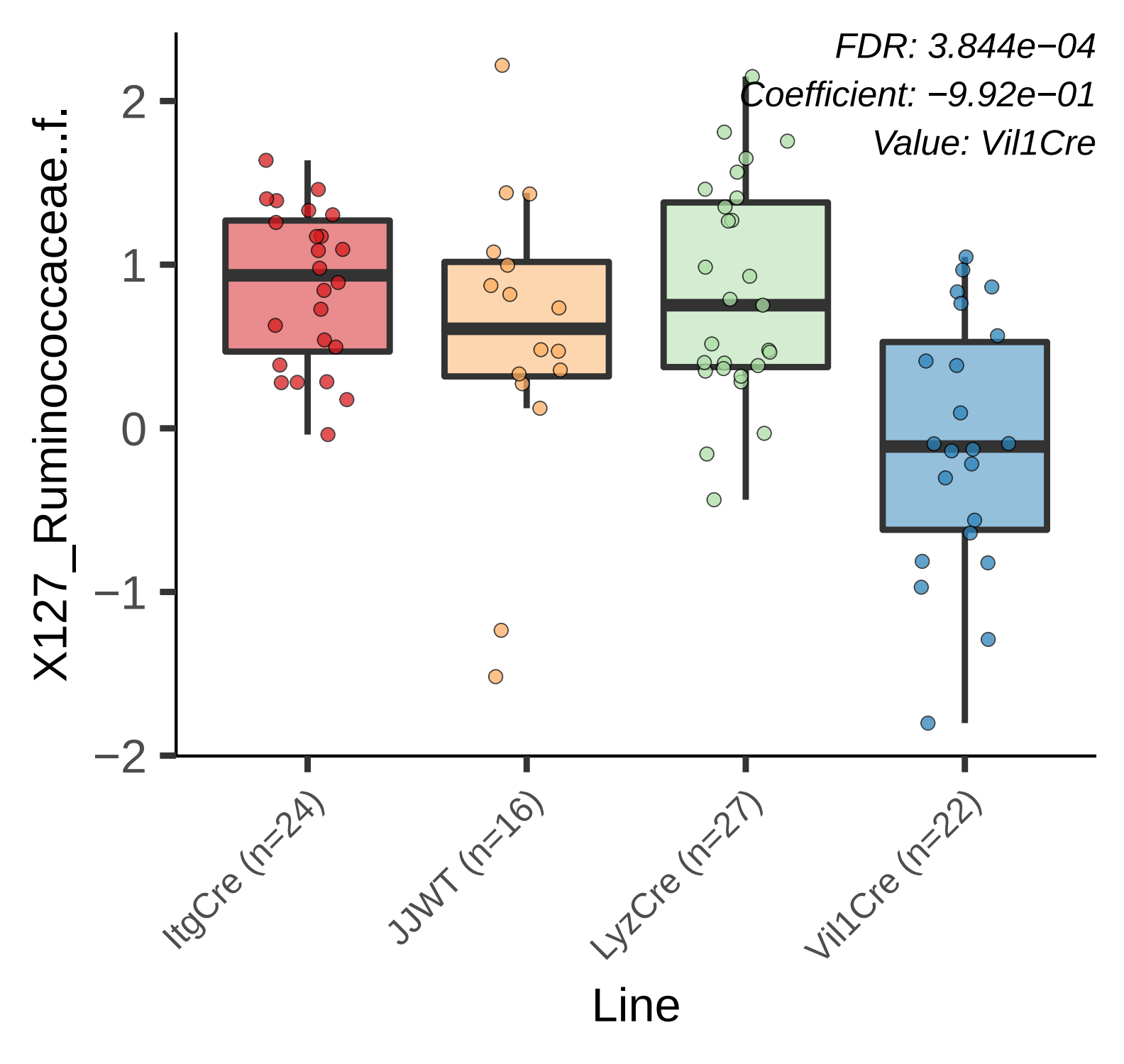

FDR: 3.408e−04
Coefficient: 4.05e+00
Value: JJWT
X61_Lachnospiraceae..f.
2
0
−2
−4
ItgCre (n=24)
JJWT (n=16)
LyzCre (n=27)
Vil1Cre (n=22)
Line

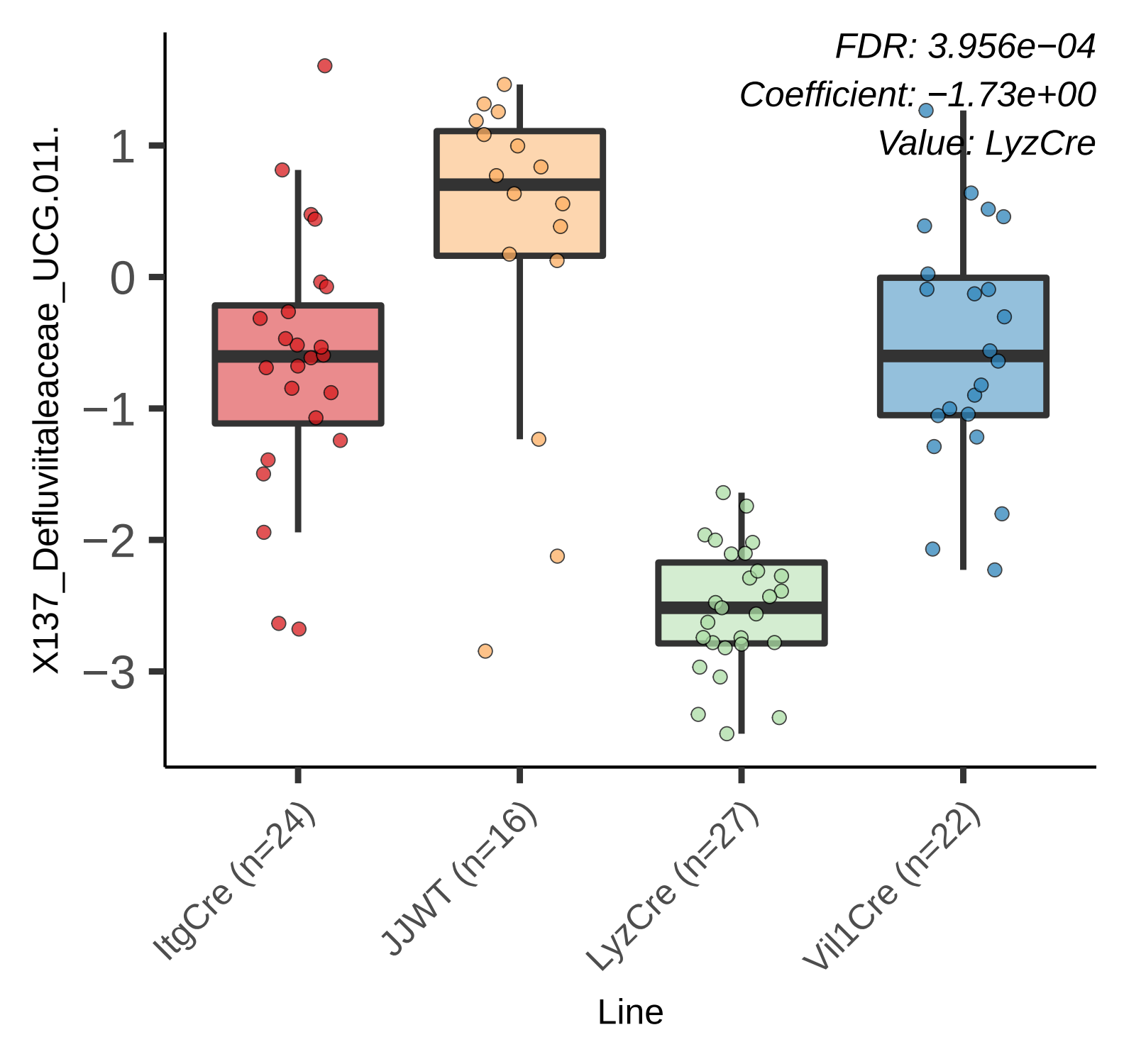

FDR: 3.408e−04
Coefficient: 3.35e+00
Value: JJWT
X291_Lachnospiraceae.f.
2
0
−2
ItgCre (n=24)
JJWT (n=16)
LyzCre (n=27)
Vil1Cre (n=22)
Line

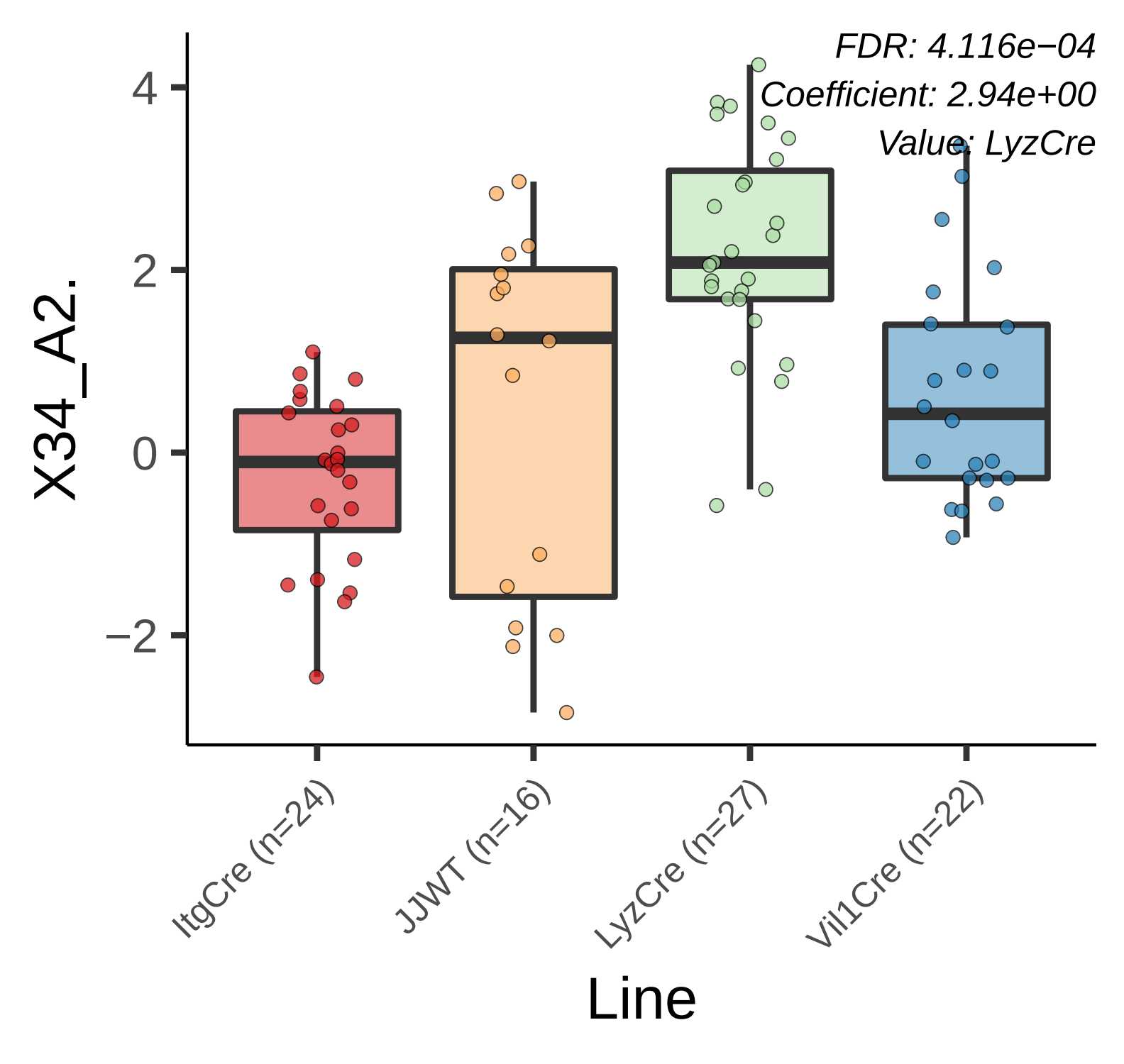

FDR: 3.600e−04
Coefficient: 2.53e+00
Value: LyzCre
X279_Lachnospiraceae_UCG.006.
2
0
−2
−4
ItgCre (n=24)
JJWT (n=16)
LyzCre (n=27)
Vil1Cre (n=22)
Line

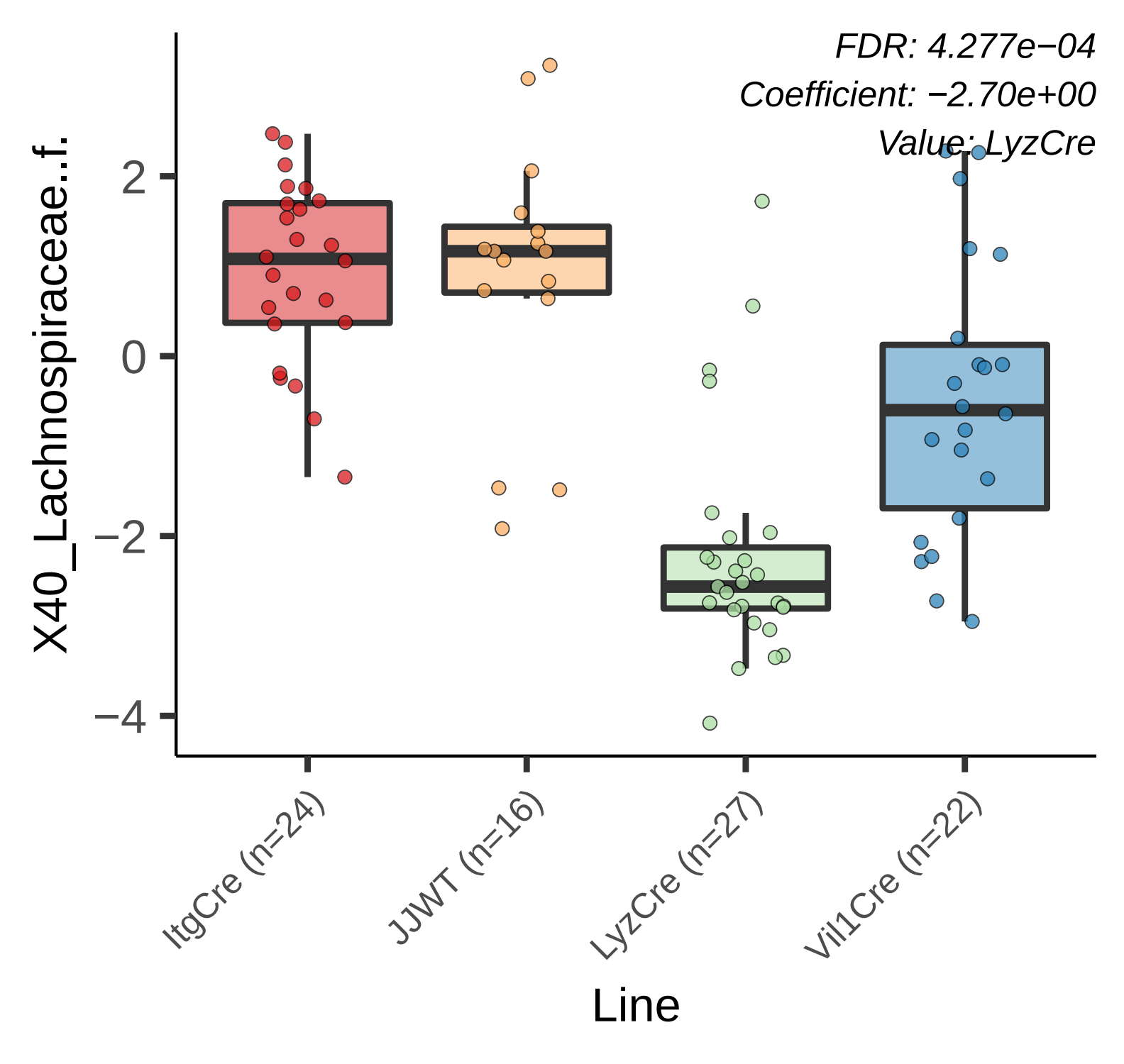

FDR: 3.844e−04
Coefficient: −3.00e+00
Value: LyzCre
X56_Muribaculaceae..f.
6
4
2
0
−2
ItgCre (n=24)
JJWT (n=16)
LyzCre (n=27)
Vil1Cre (n=22)
Line

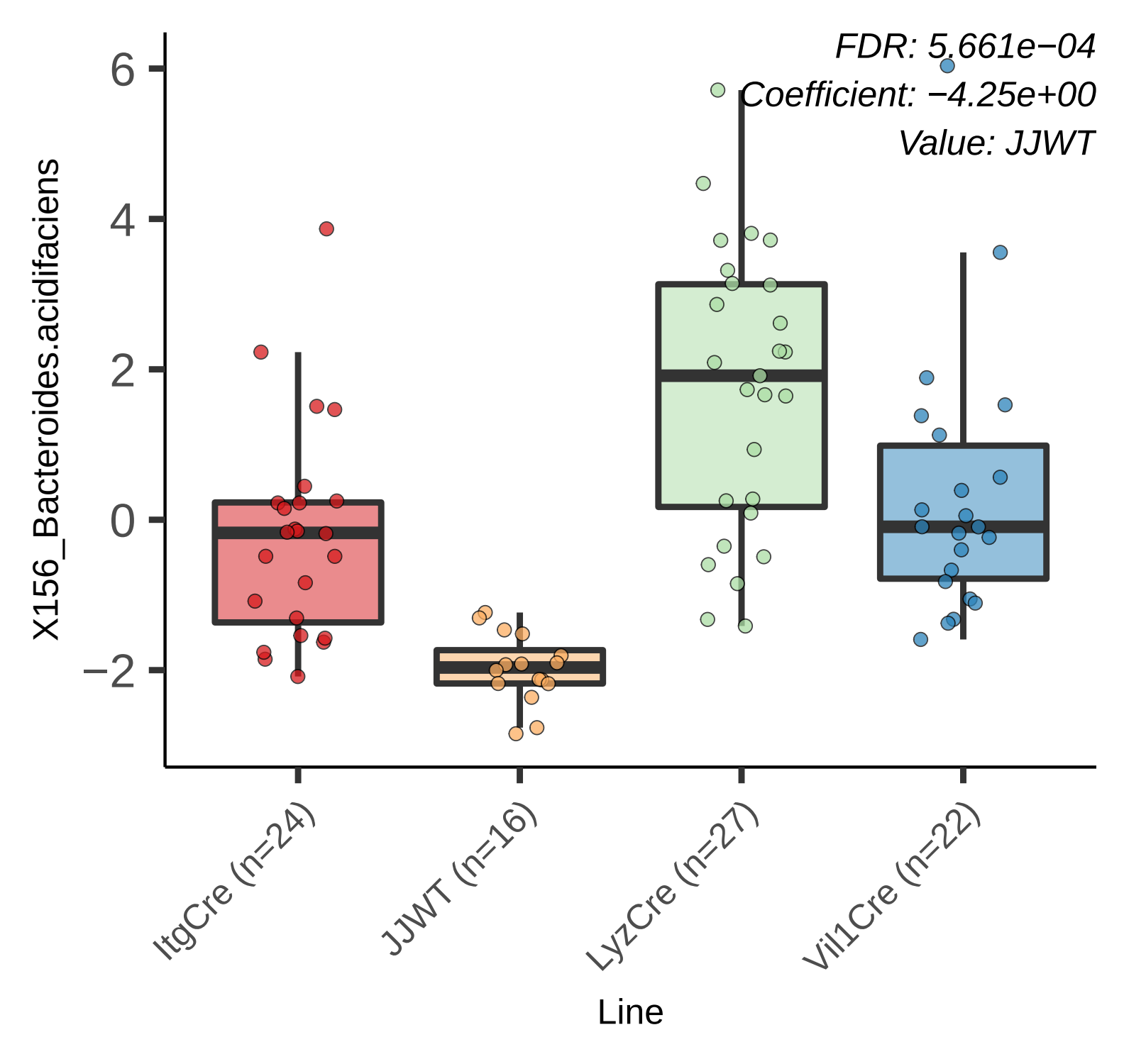

FDR: 3.844e−04
Coefficient: 3.21e+00
Value: JJWT
X108_Lachnospiraceae.f.
2
0
−2
−4
ItgCre (n=24)
JJWT (n=16)
LyzCre (n=27)
Vil1Cre (n=22)
Line

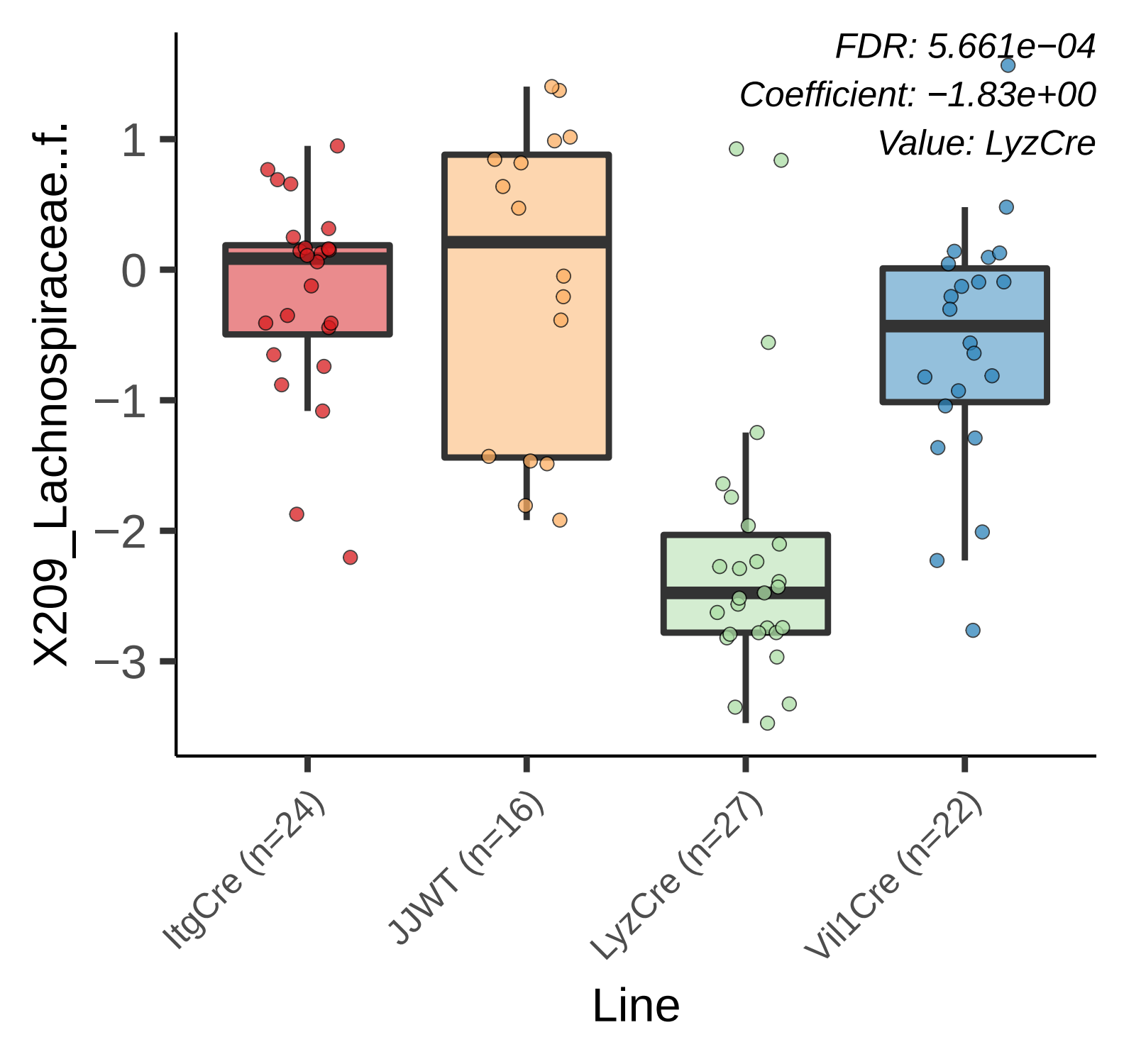

FDR: 3.844e−04
Coefficient: −9.92e−01
Value: Vil1Cre
X127_Ruminococcaceae..f.
2
1
0
−1
−2
ItgCre (n=24)
JJWT (n=16)
LyzCre (n=27)
Vil1Cre (n=22)
Line

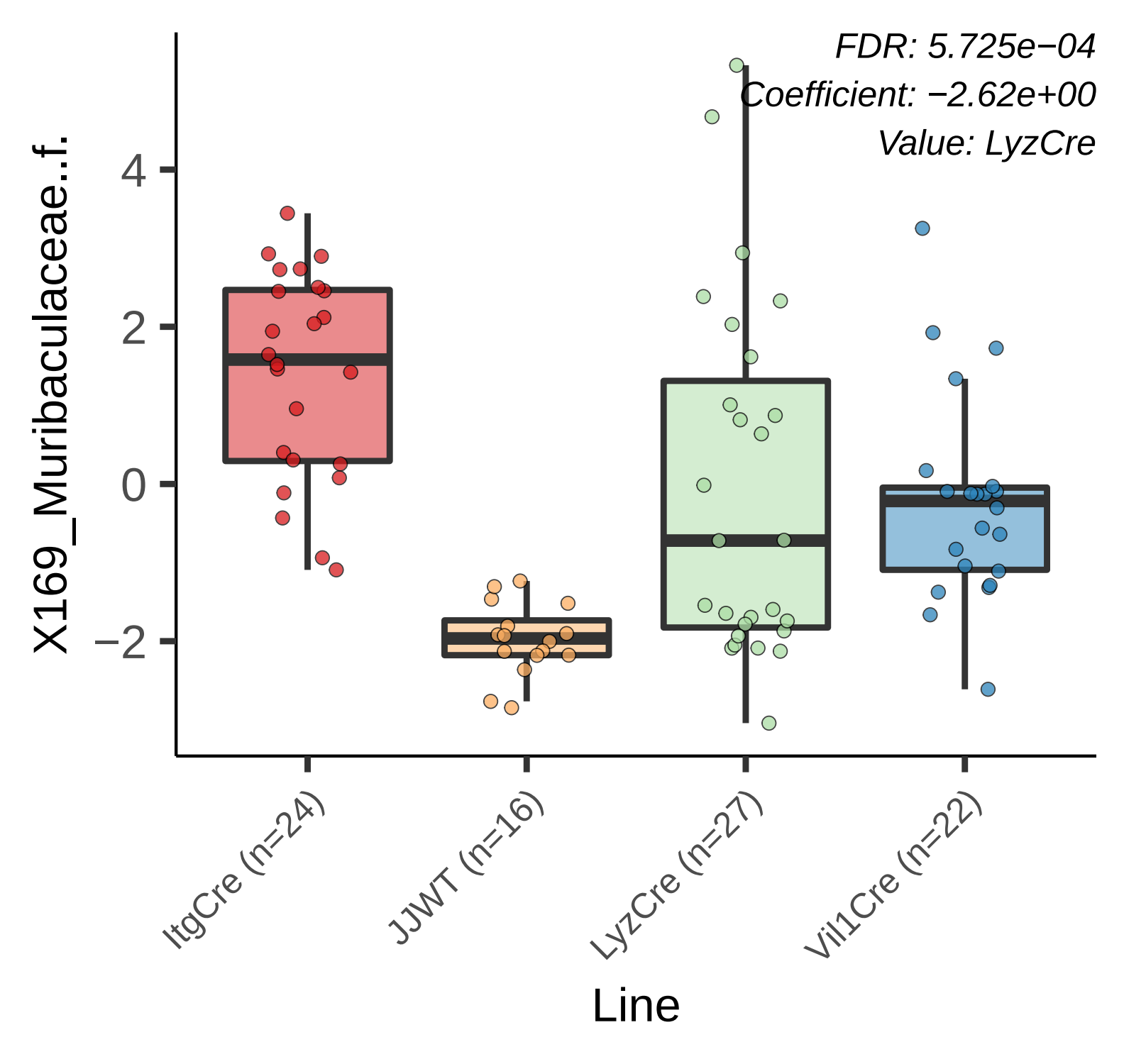

FDR: 3.956e−04
Coefficient: −1.73e+00
Value: LyzCre
X137_Defluviitaleaceae_UCG.011.
1
0
−1
−2
−3
ItgCre (n=24)
JJWT (n=16)
LyzCre (n=27)
Vil1Cre (n=22)
Line

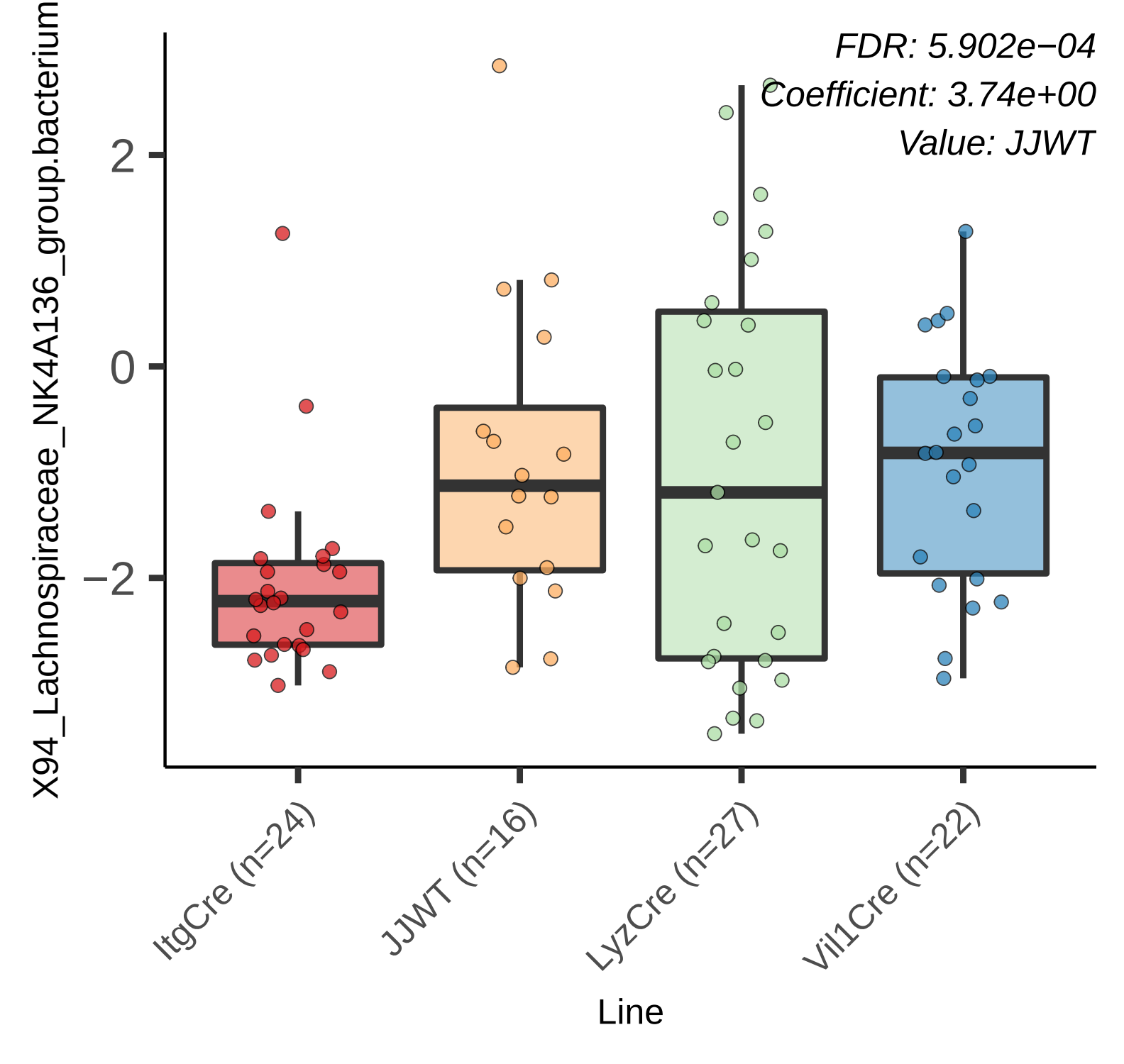

FDR: 4.116e−04
Coefficient: 2.94e+00
Value: LyzCre
X34_A2.
4
2
0
−2
ItgCre (n=24)
JJWT (n=16)
LyzCre (n=27)
Vil1Cre (n=22)
Line

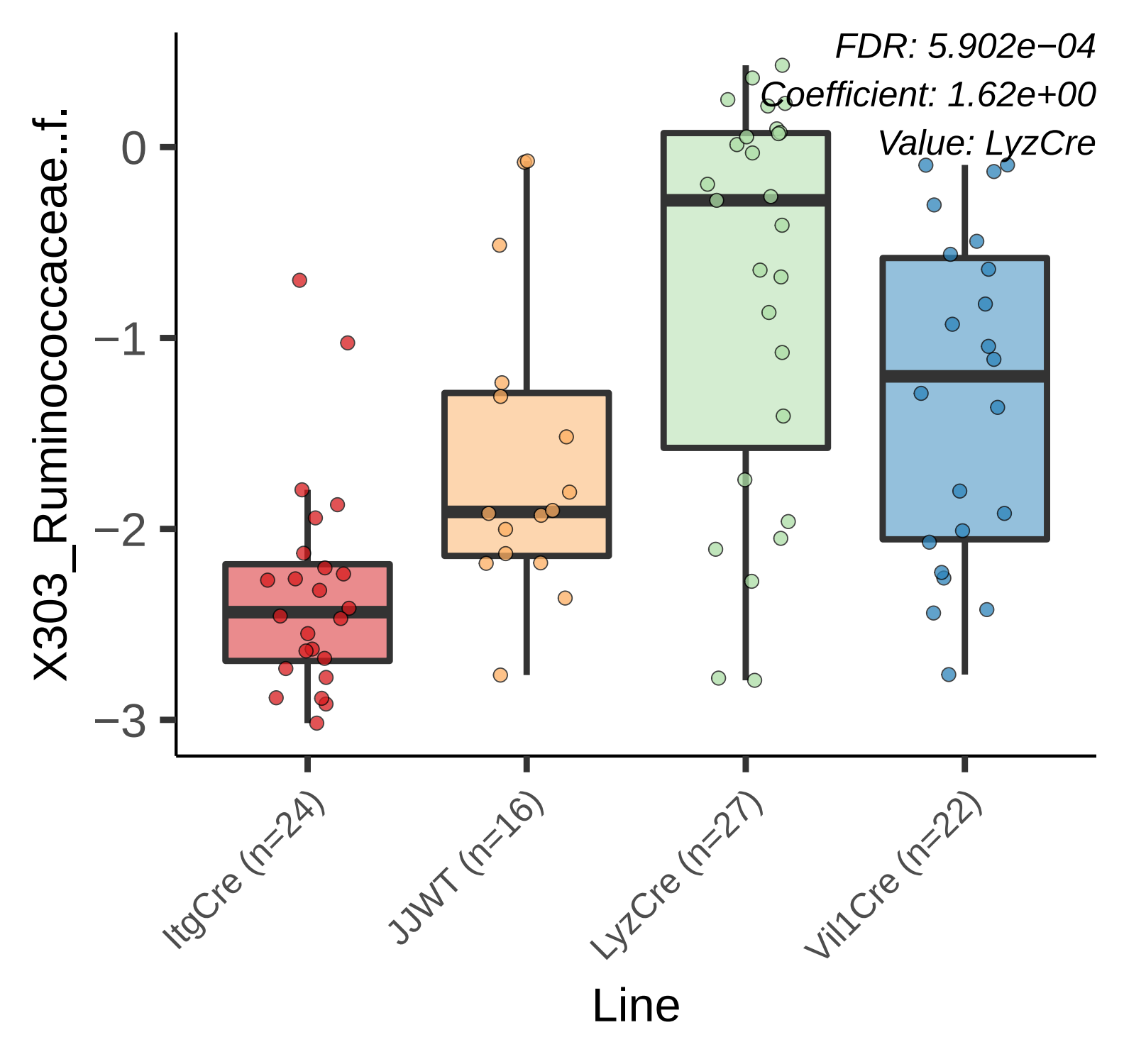

FDR: 4.277e−04
Coefficient: −2.70e+00
Value: LyzCre
X40_Lachnospiraceae..f.
2
0
−2
−4
ItgCre (n=24)
JJWT (n=16)
LyzCre (n=27)
Vil1Cre (n=22)
Line

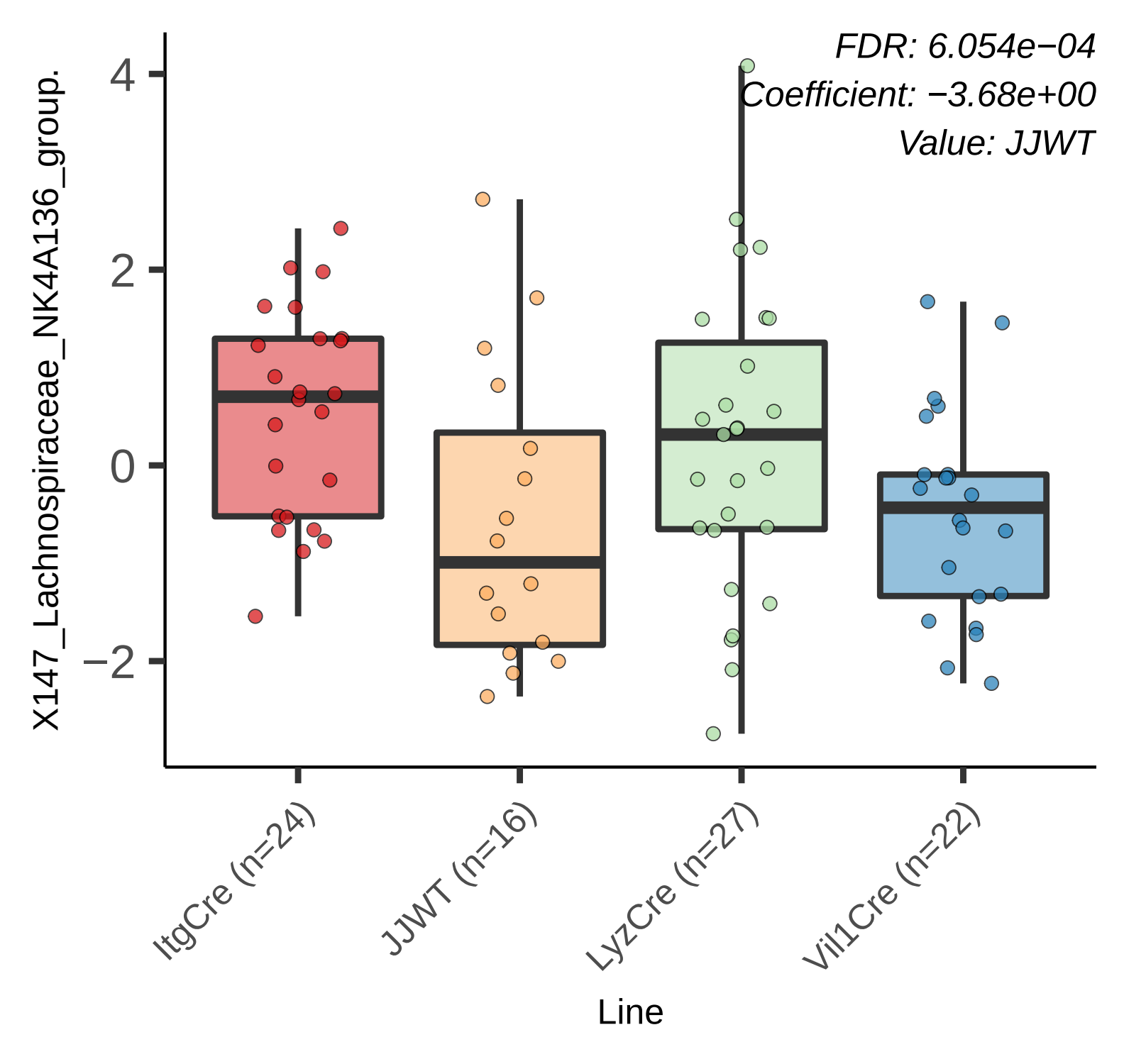

FDR: 5.661e−04
Coefficient: −4.25e+00
Value: JJWT
X156_Bacteroides.acidifaciens
6
4
2
0
−2
ItgCre (n=24)
JJWT (n=16)
LyzCre (n=27)
Vil1Cre (n=22)
Line

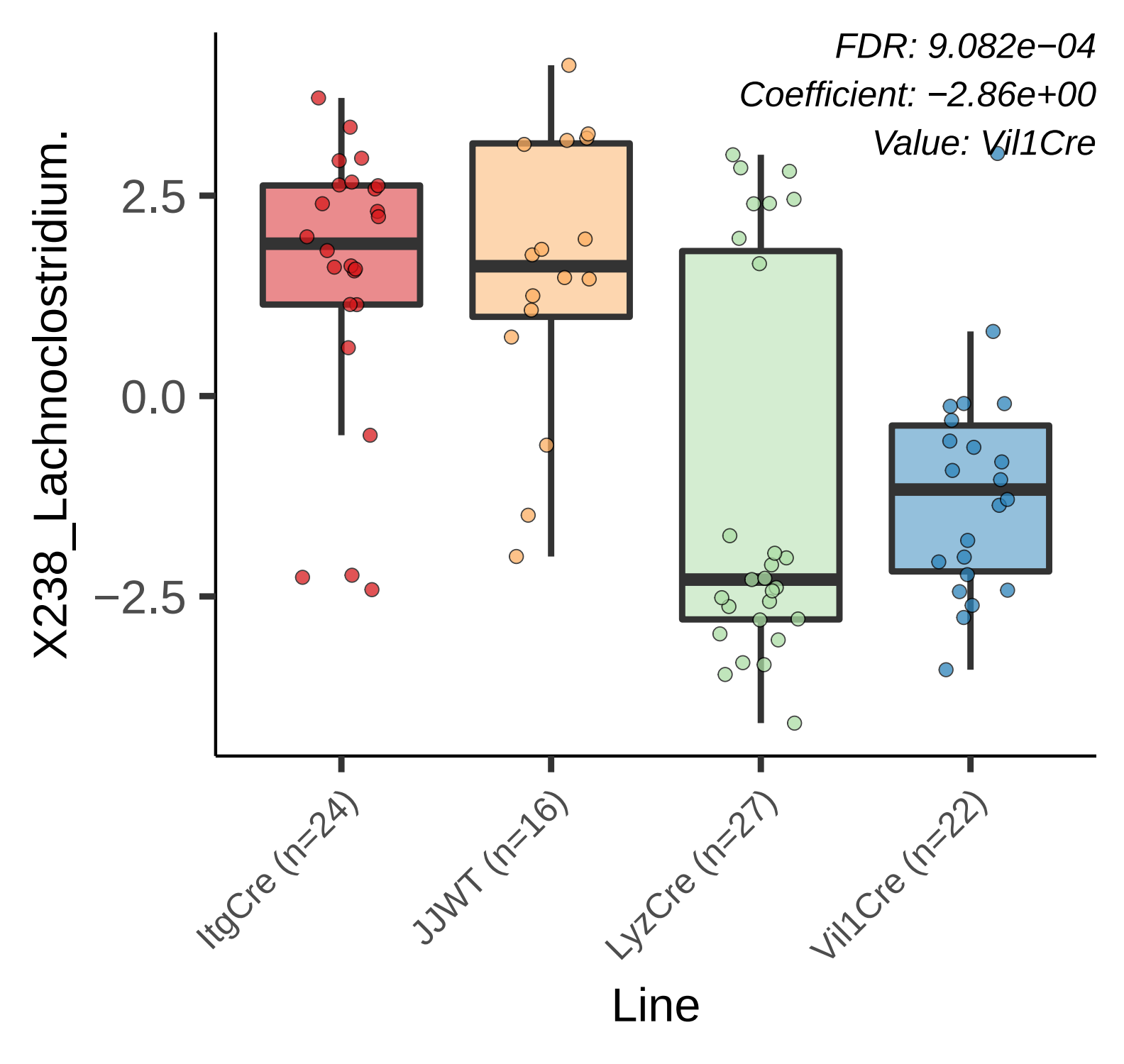

FDR: 5.661e−04
Coefficient: −1.83e+00
Value: LyzCre
X209_Lachnospiraceae.f.
1
0
−1
−2
−3
ItgCre (n=24)
JJWT (n=16)
LyzCre (n=27)
Vil1Cre (n=22)
Line

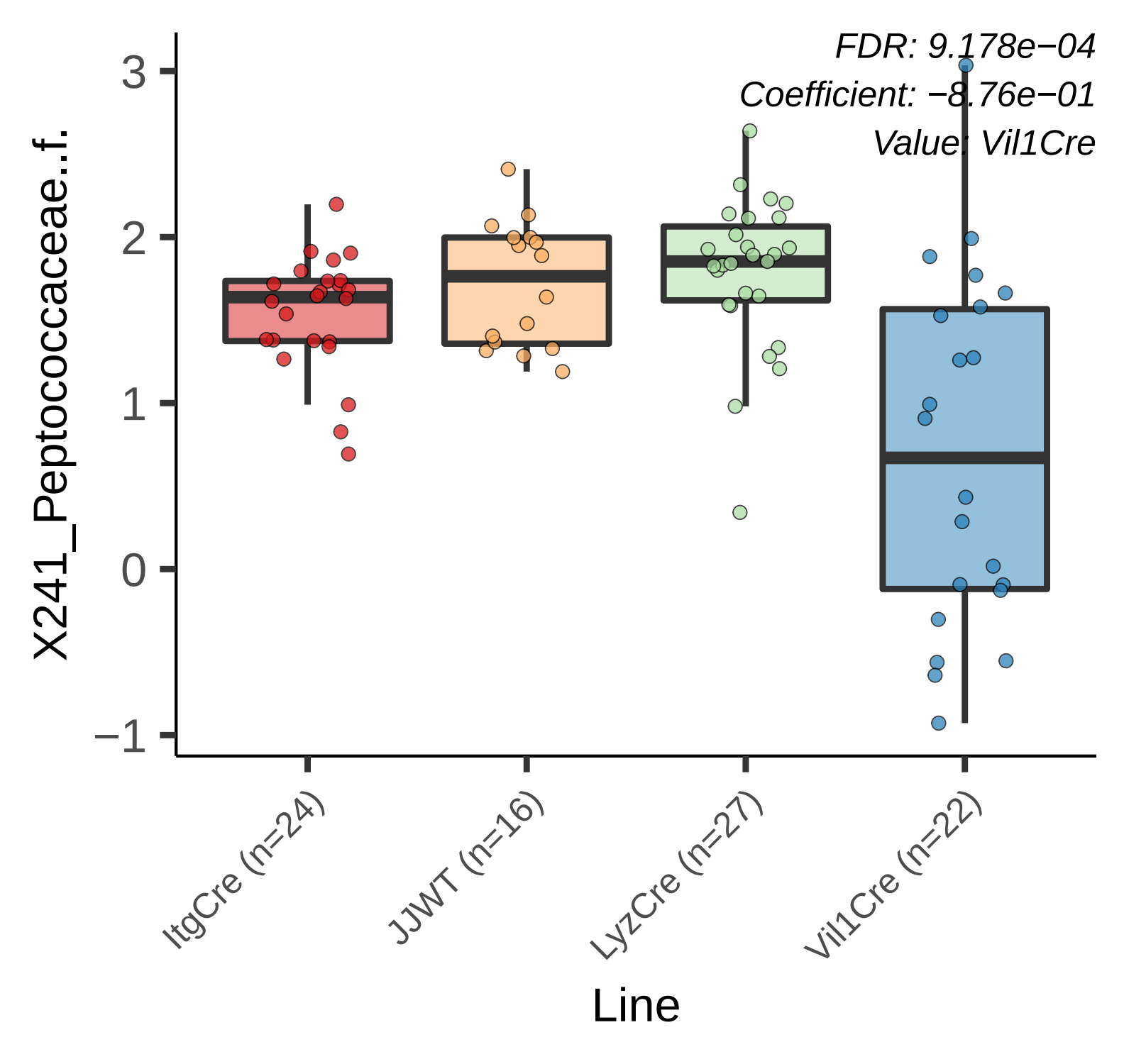

FDR: 5.725e−04
Coefficient: −2.62e+00
Value: LyzCre
X169_Muribaculaceae..f.
4
2
0
−2
ItgCre (n=24)
JJWT (n=16)
LyzCre (n=27)
Vil1Cre (n=22)
Line

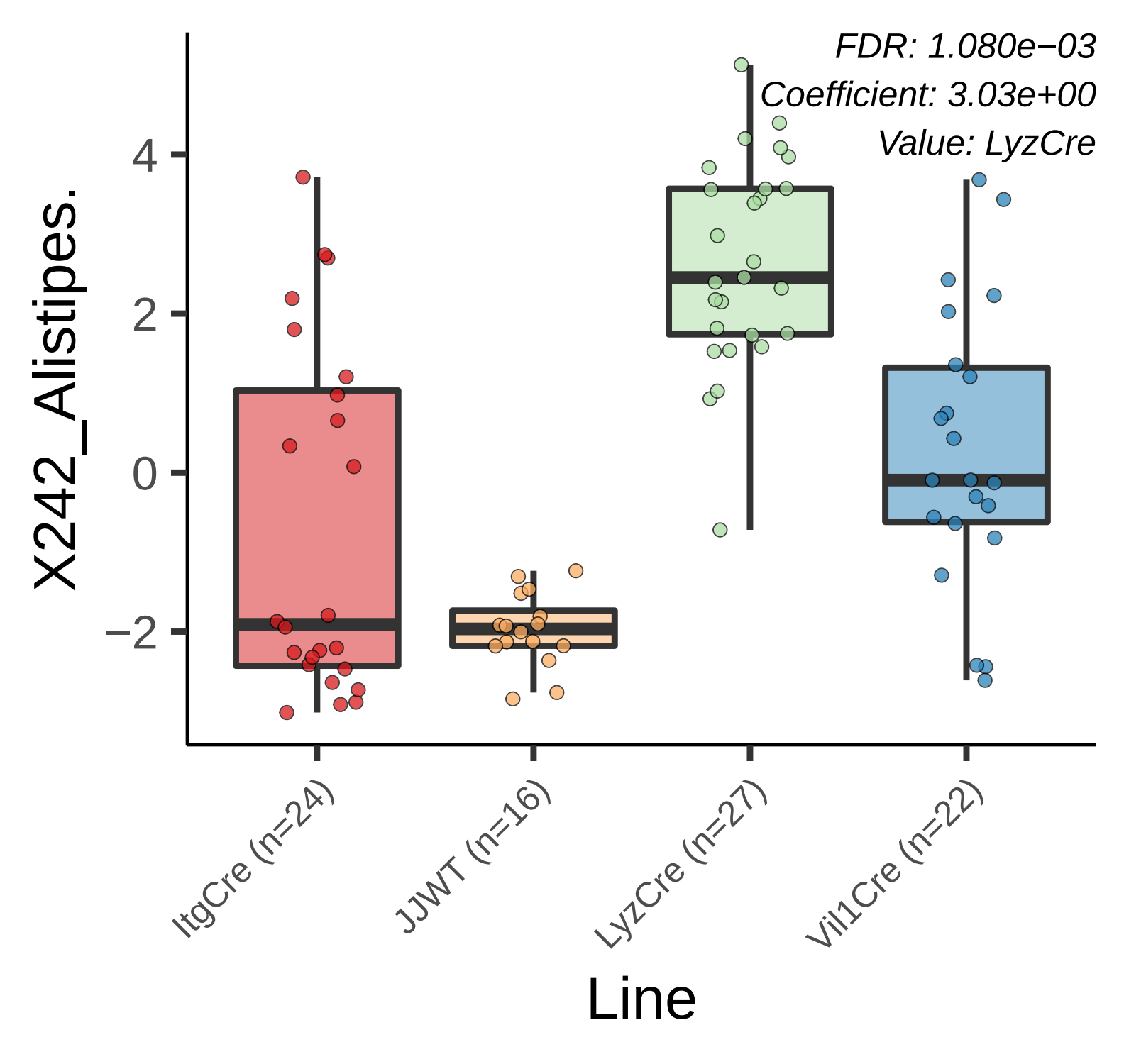

FDR: 5.902e−04
Coefficient: 3.74e+00
Value: JJWT
X94_Lachnospiraceae_NK4A136_group.bacterium
2
0
−2
ItgCre (n=24)
JJWT (n=16)
LyzCre (n=27)
Vil1Cre (n=22)
Line

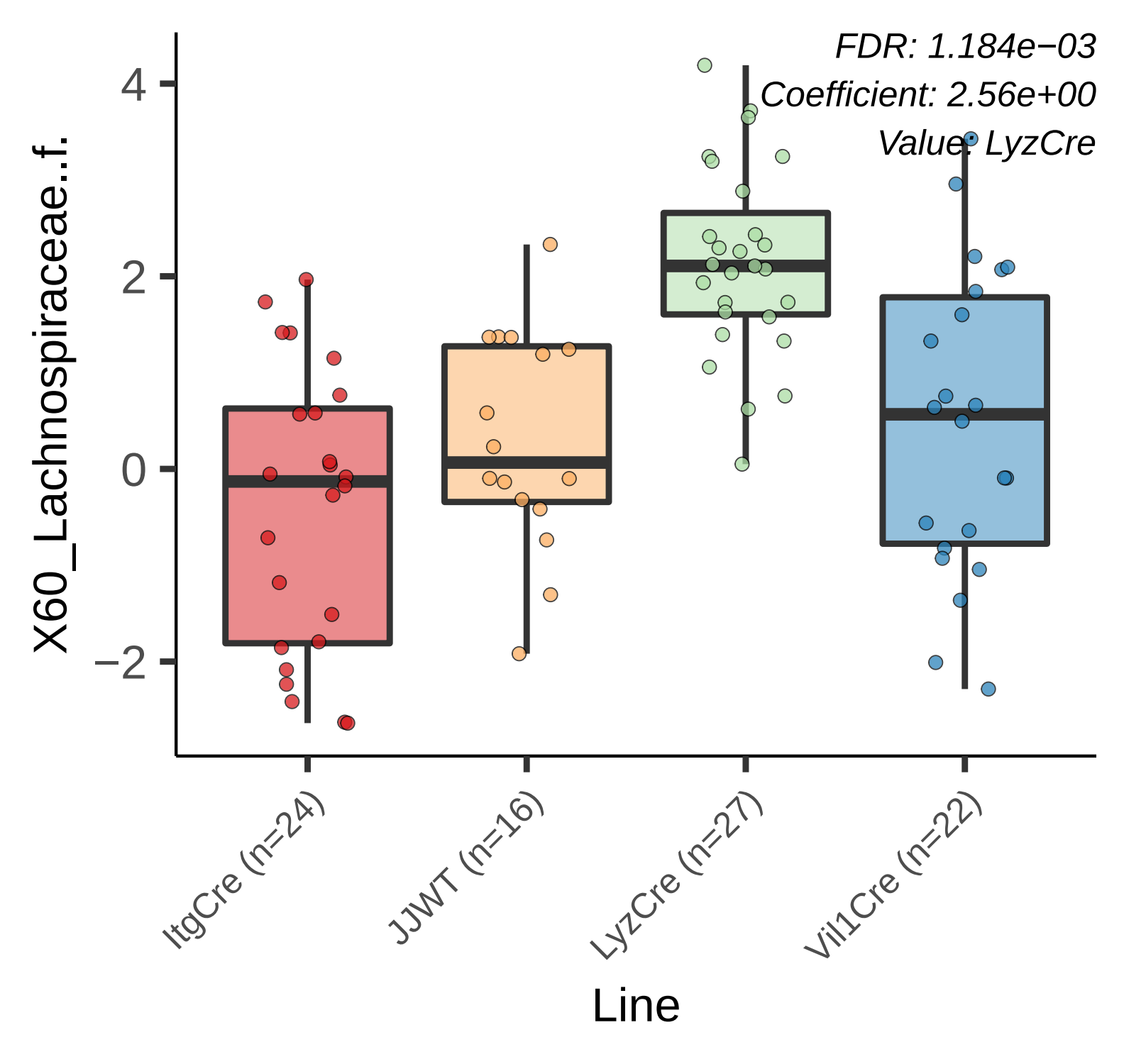

FDR: 5.902e−04
Coefficient: 1.62e+00
Value: LyzCre
X303_Ruminococcaceae..f.
0
−1
−2
−3
ItgCre (n=24)
JJWT (n=16)
LyzCre (n=27)
Vil1Cre (n=22)
Line

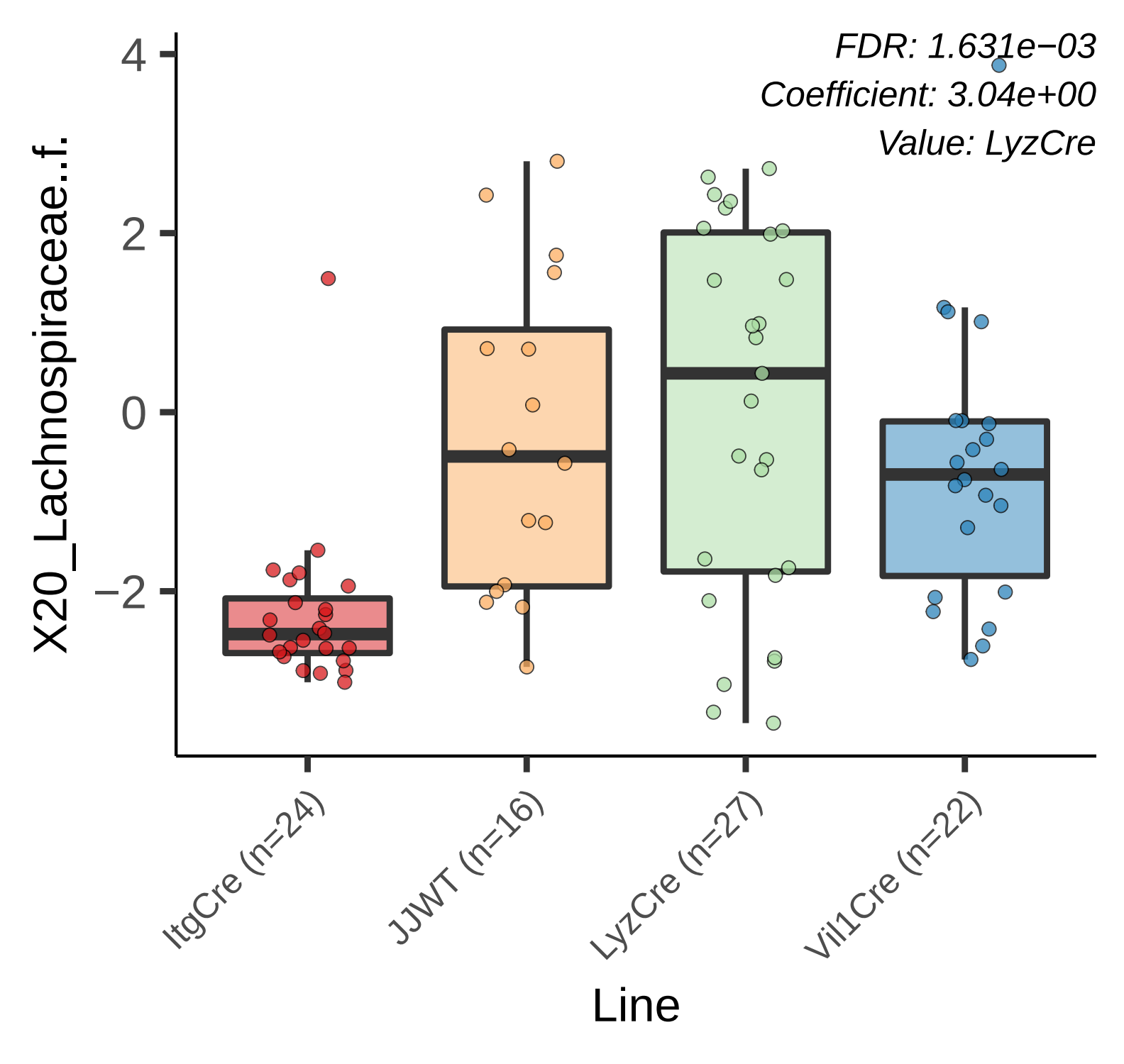

FDR: 6.054e−04
Coefficient: −3.68e+00
Value: JJWT
X147_Lachnospiraceae_NK4A136_group.
4
2
0
−2
ItgCre (n=24)
JJWT (n=16)
LyzCre (n=27)
Vil1Cre (n=22)
Line

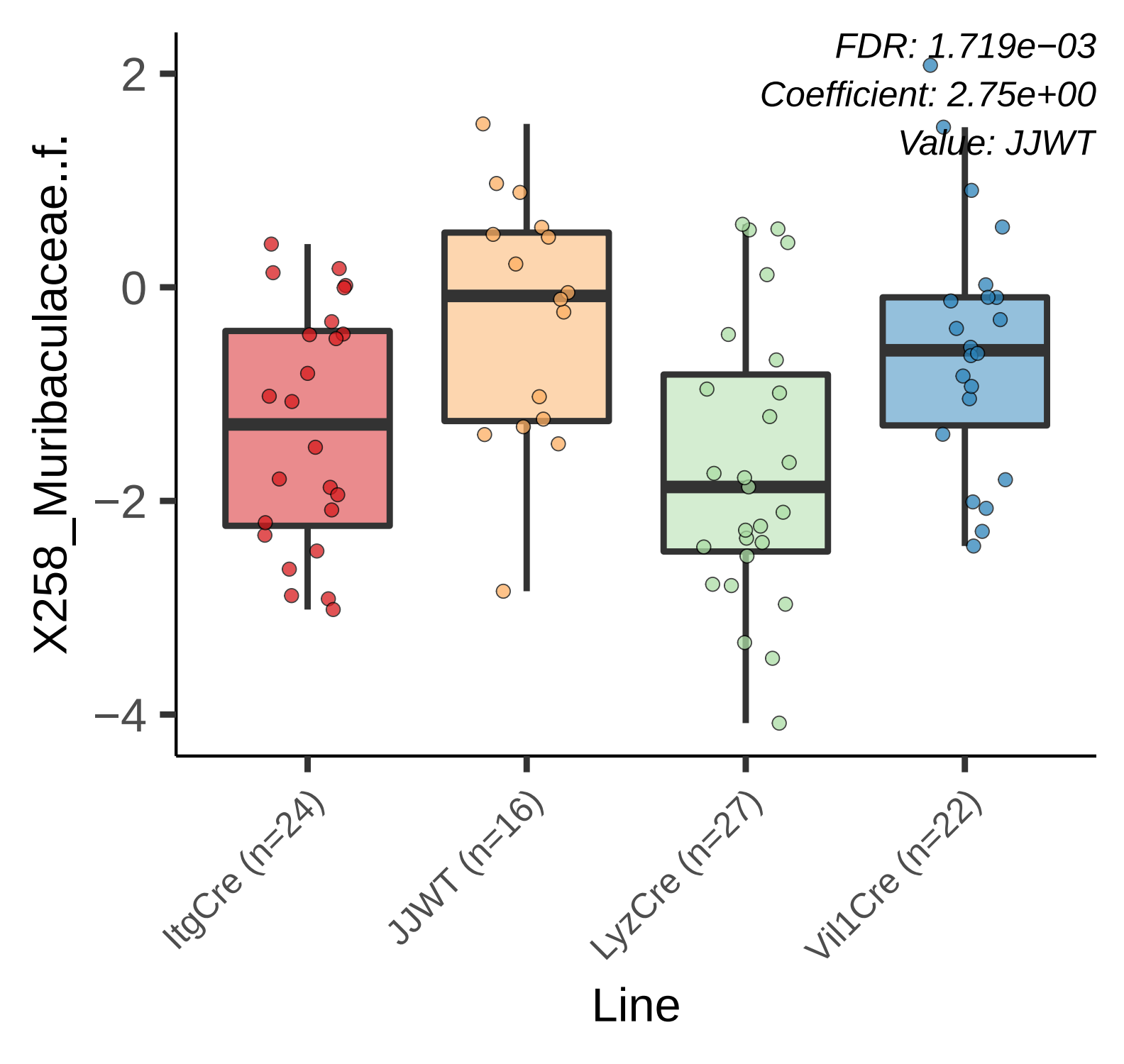

FDR: 9.082e−04
Coefficient: −2.86e+00
Value: Vil1Cre
X238_Lachnoclostridium.
2.5
0.0
−2.5
ItgCre (n=24)
JJWT (n=16)
LyzCre (n=27)
Vil1Cre (n=22)
Line

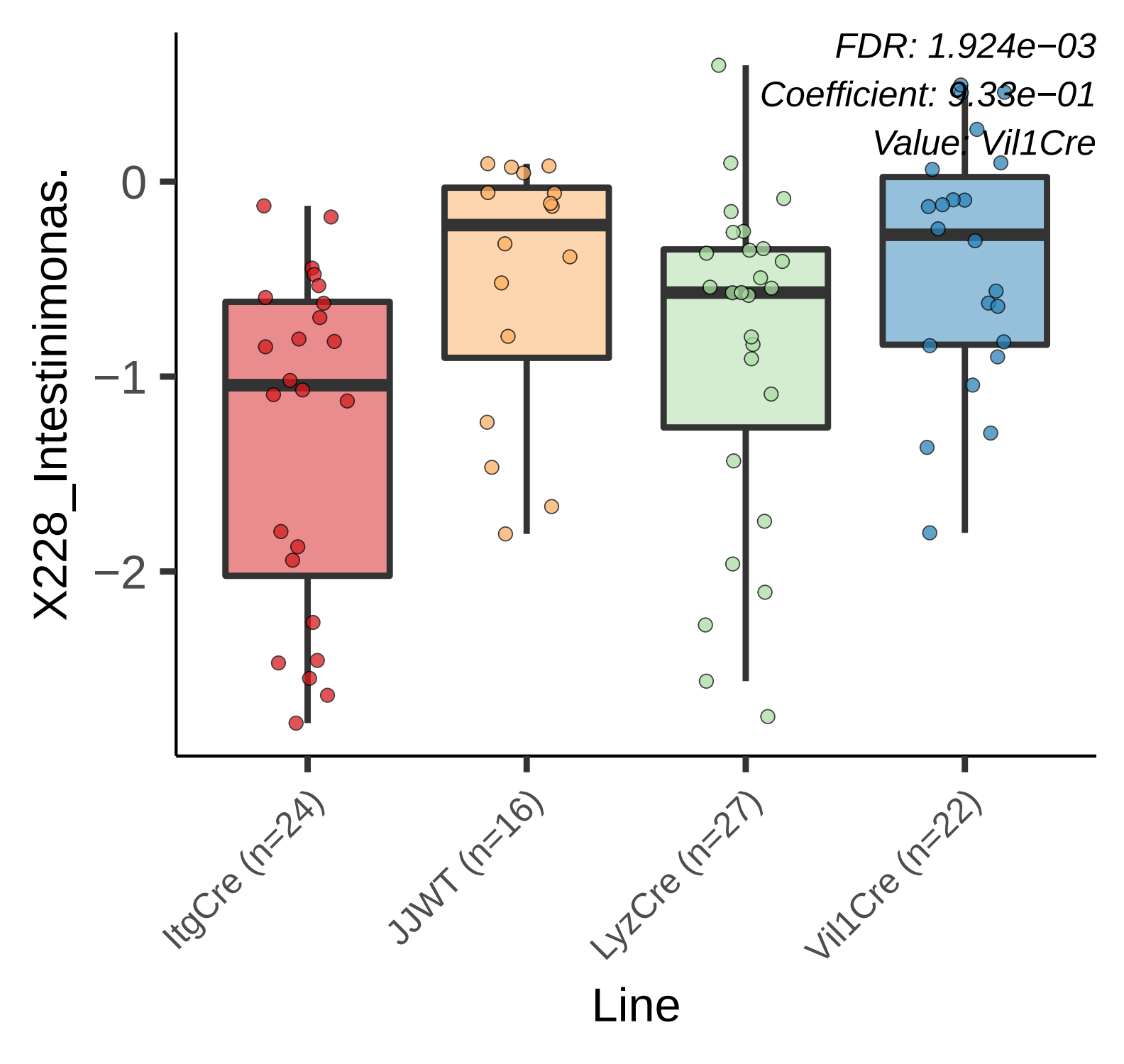

FDR: 9.178e−04
Coefficient: −8.76e−01
Value: Vil1Cre
X241_Peptococcaceae..f.
3
2
1
0
−1
ItgCre (n=24)
JJWT (n=16)
LyzCre (n=27)
Vil1Cre (n=22)
Line

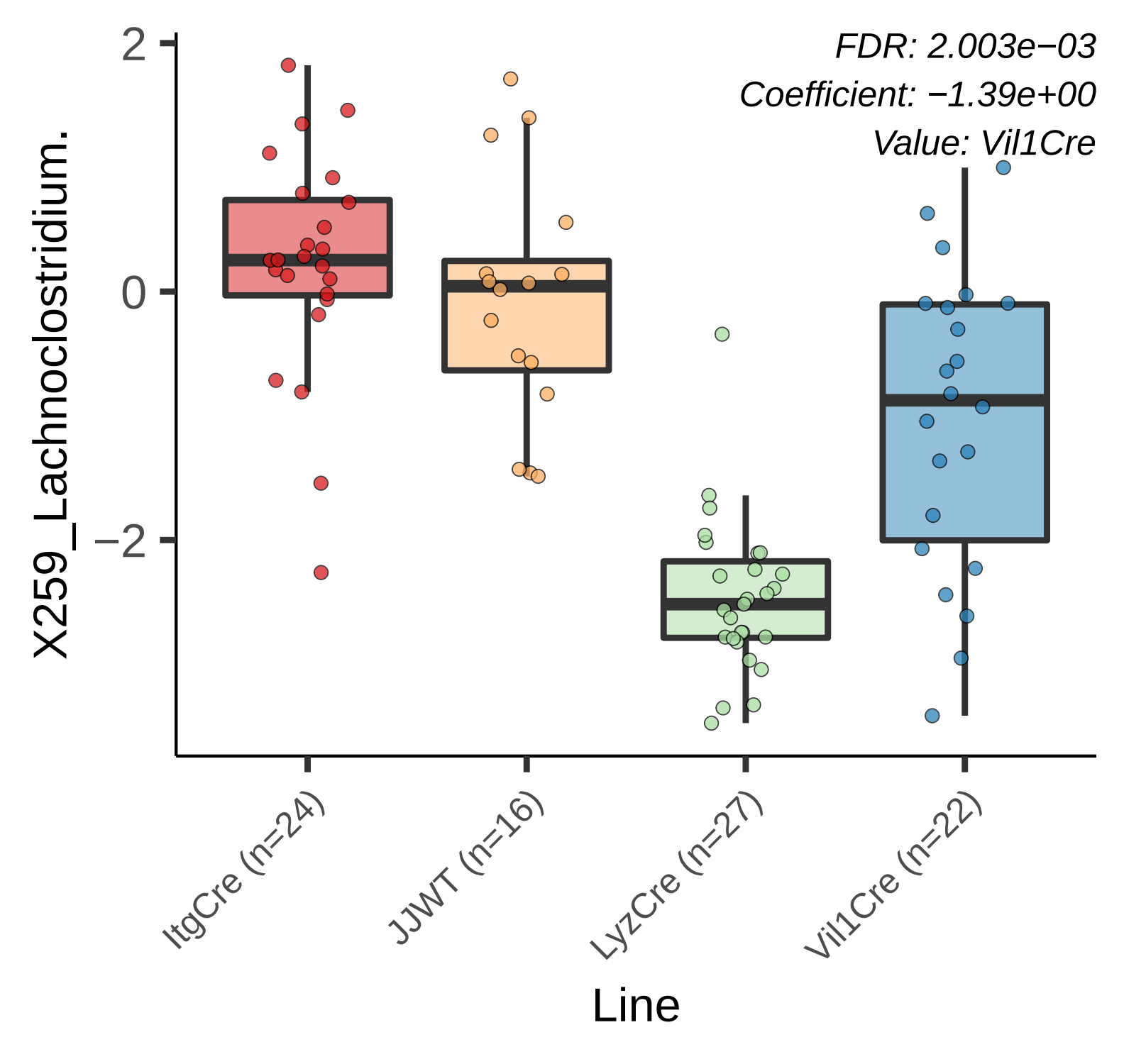

FDR: 1.080e−03
Coefficient: 3.03e+00
Value: LyzCre
X242_Alistipes.
4
2
0
−2
ItgCre (n=24)
JJWT (n=16)
LyzCre (n=27)
Vil1Cre (n=22)
Line

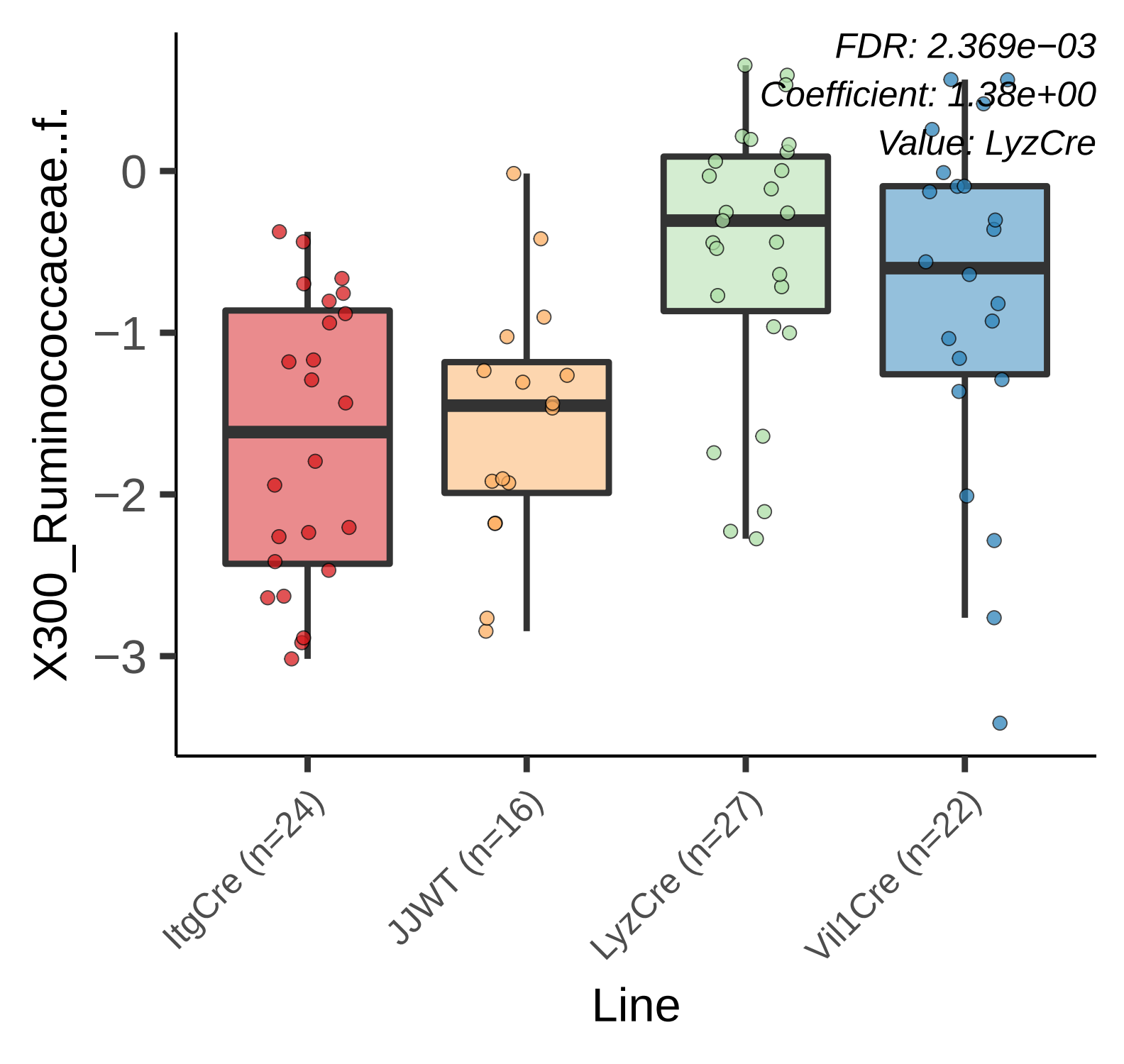

FDR: 1.184e−03
Coefficient: 2.56e+00
Value: LyzCre
X60_Lachnospiraceae..f.
4
2
0
−2
ItgCre (n=24)
JJWT (n=16)
LyzCre (n=27)
Vil1Cre (n=22)
Line

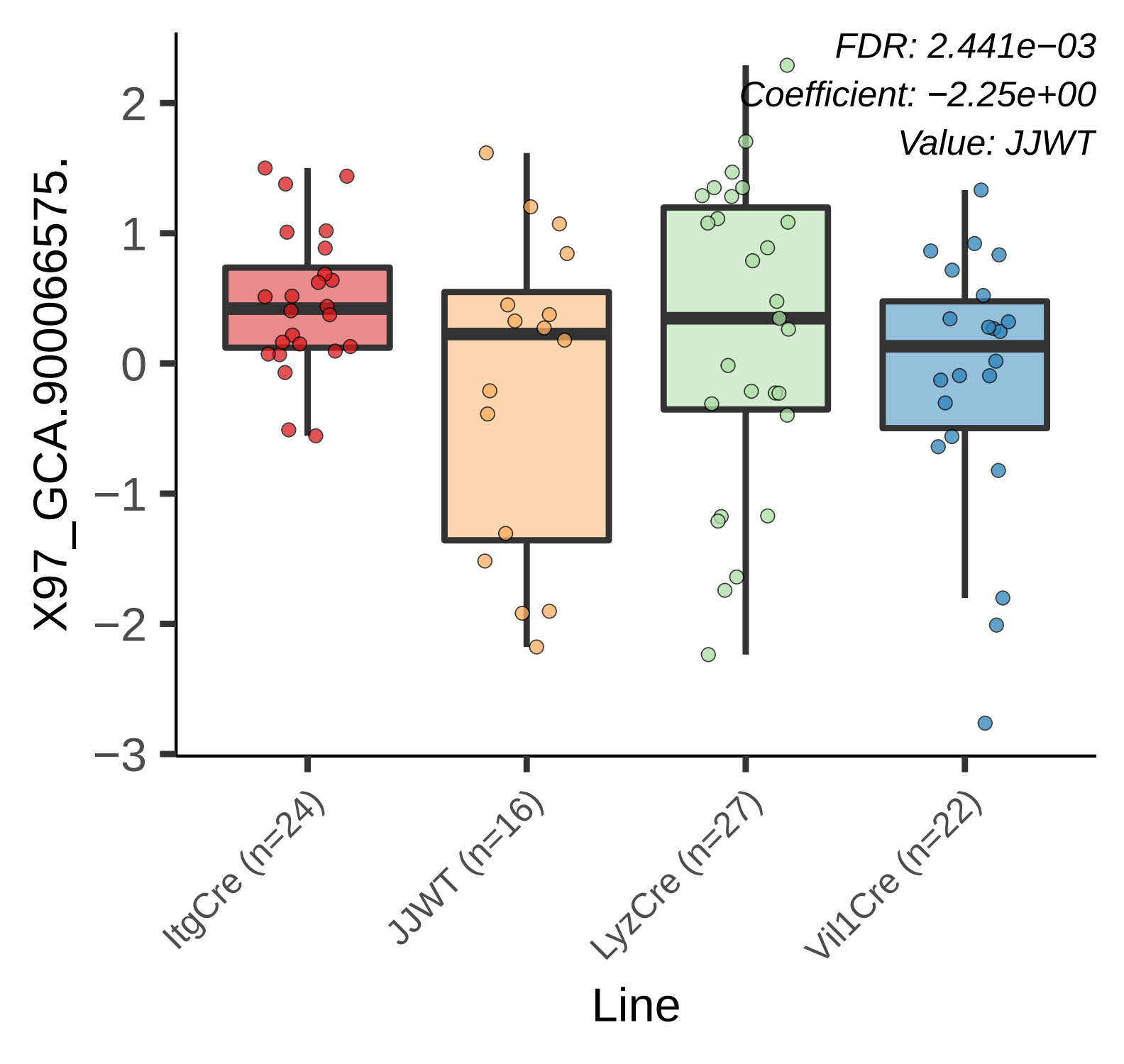

FDR: 1.631e−03
Coefficient: 3.04e+00
Value: LyzCre
X20_Lachnospiraceae..f.
4
2
0
−2
ItgCre (n=24)
JJWT (n=16)
LyzCre (n=27)
Vil1Cre (n=22)
Line

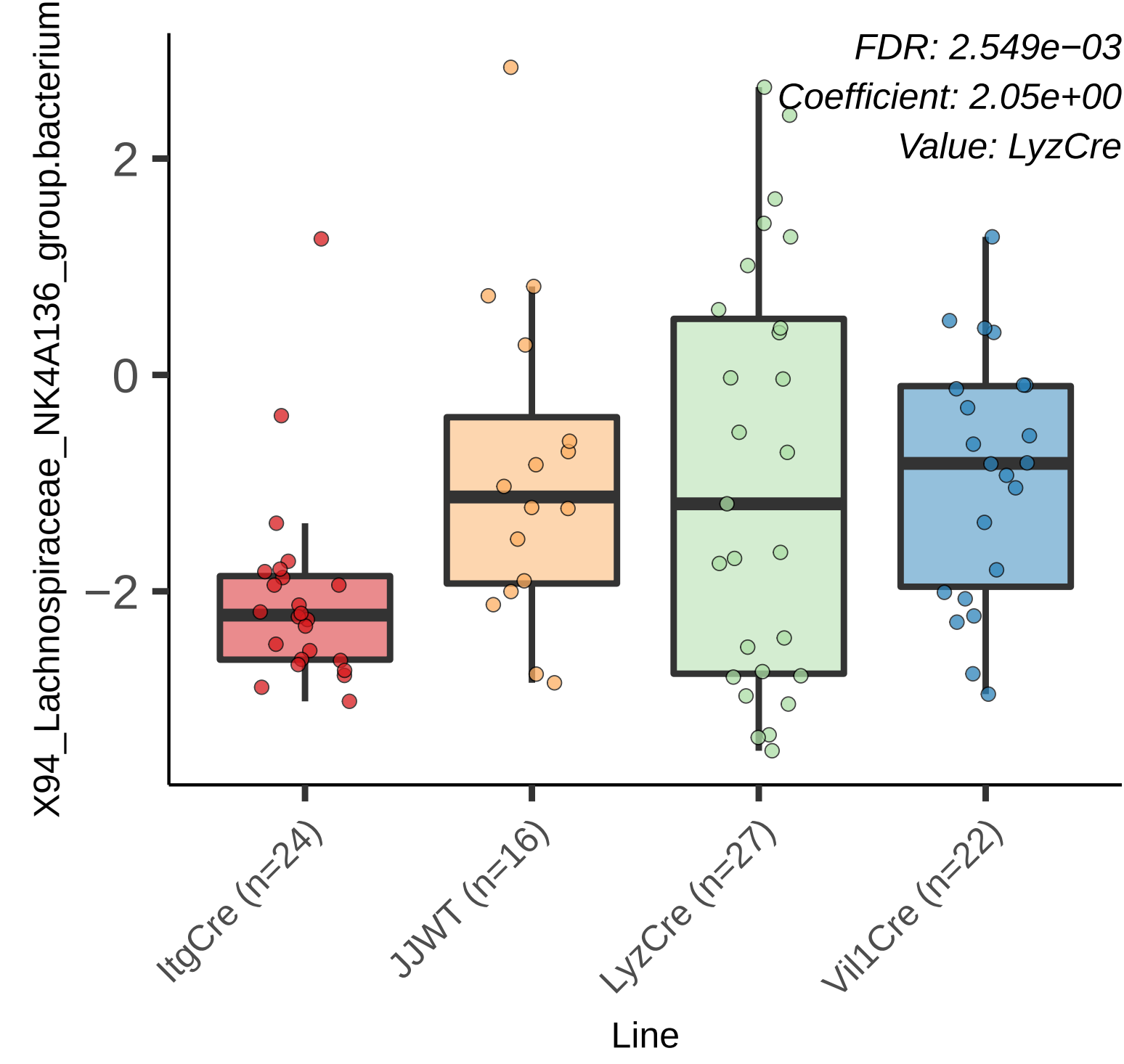

FDR: 1.719e−03
Coefficient: 2.75e+00
Value: JJWT
X258_Muribaculaceae..f.
2
0
−2
−4
ItgCre (n=24)
JJWT (n=16)
LyzCre (n=27)
Vil1Cre (n=22)
Line

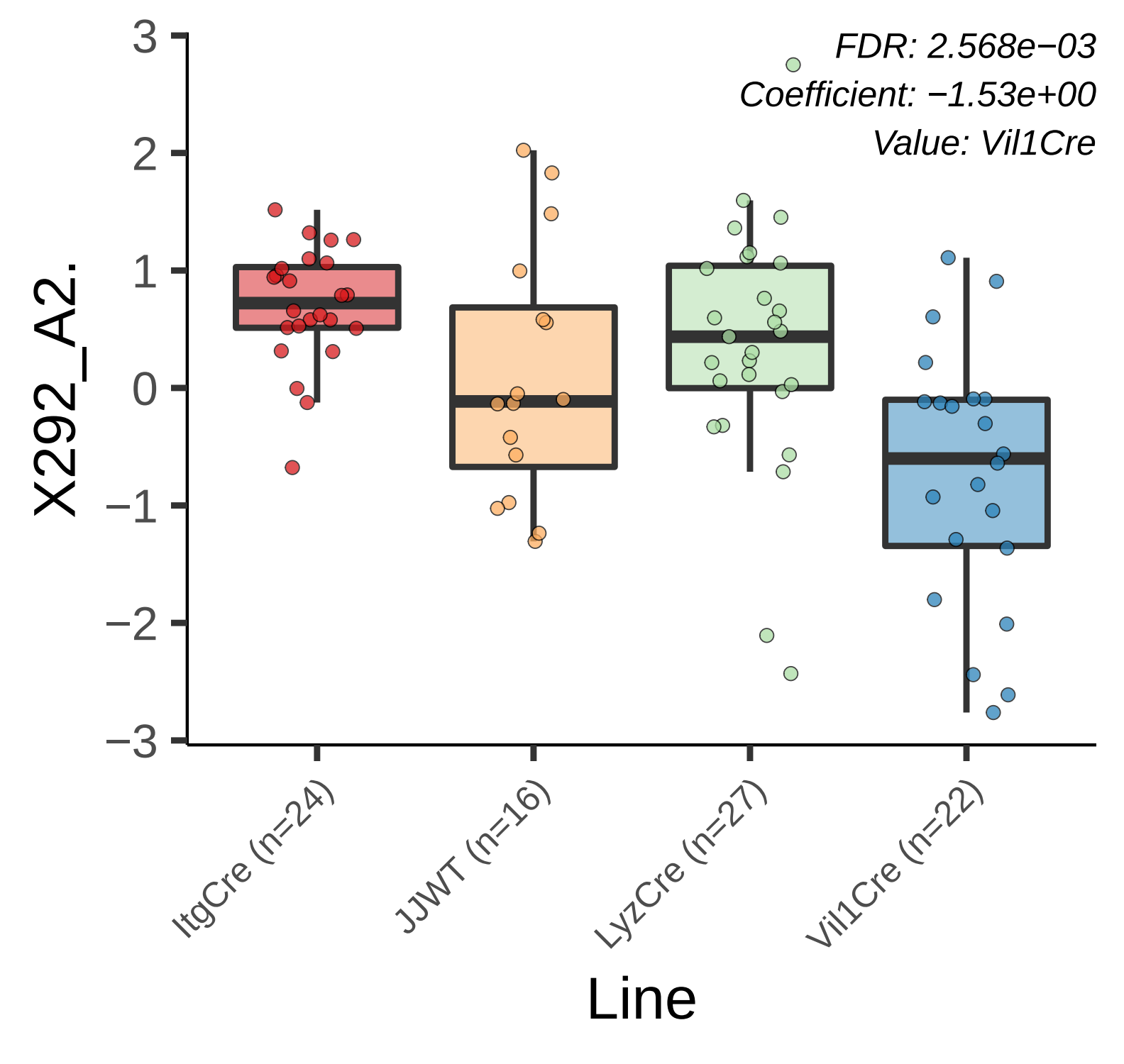

FDR: 1.924e−03
Coefficient: 9.33e−01
Value: Vil1Cre
X228_Intestinimonas.
0
−1
−2
ItgCre (n=24)
JJWT (n=16)
LyzCre (n=27)
Vil1Cre (n=22)
Line

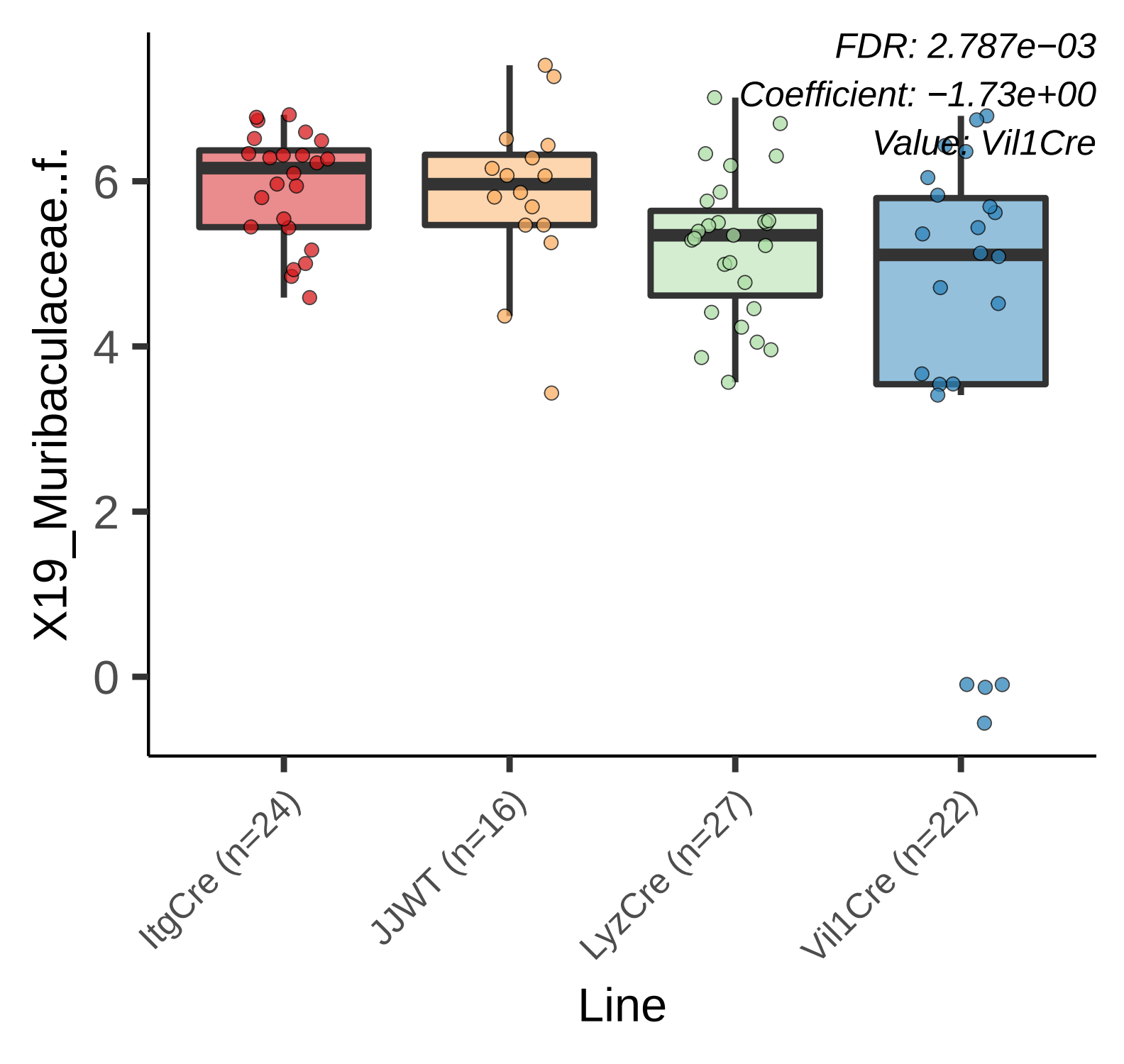

FDR: 2.003e−03
Coefficient: −1.39e+00
Value: Vil1Cre
X259_Lachnoclostridium.
2
0
−2
ItgCre (n=24)
JJWT (n=16)
LyzCre (n=27)
Vil1Cre (n=22)
Line

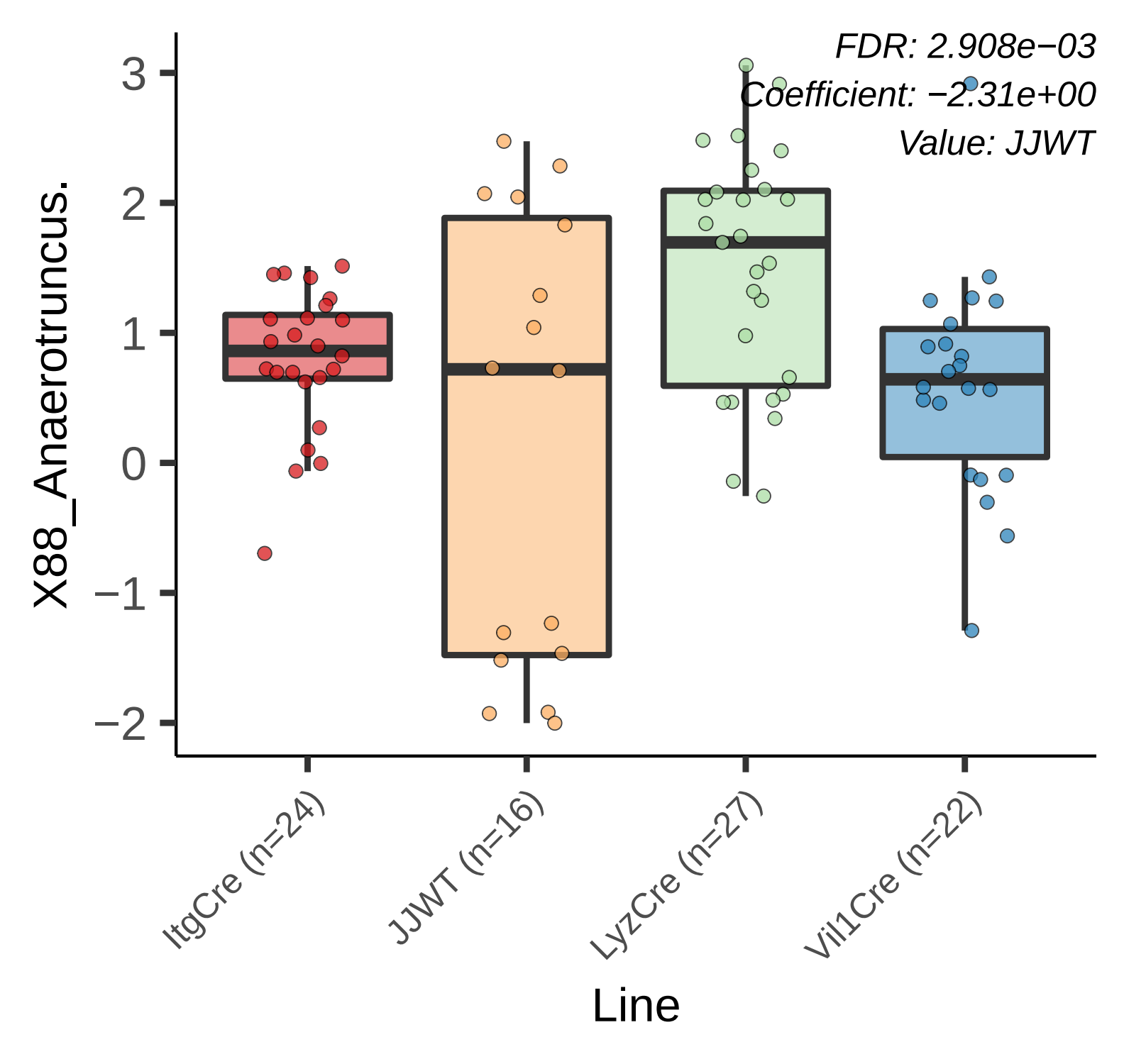

FDR: 2.369e−03
Coefficient: 1.38e+00
Value: LyzCre
X300_Ruminococcaceae..f.
0
−1
−2
−3
ItgCre (n=24)
JJWT (n=16)
LyzCre (n=27)
Vil1Cre (n=22)
Line

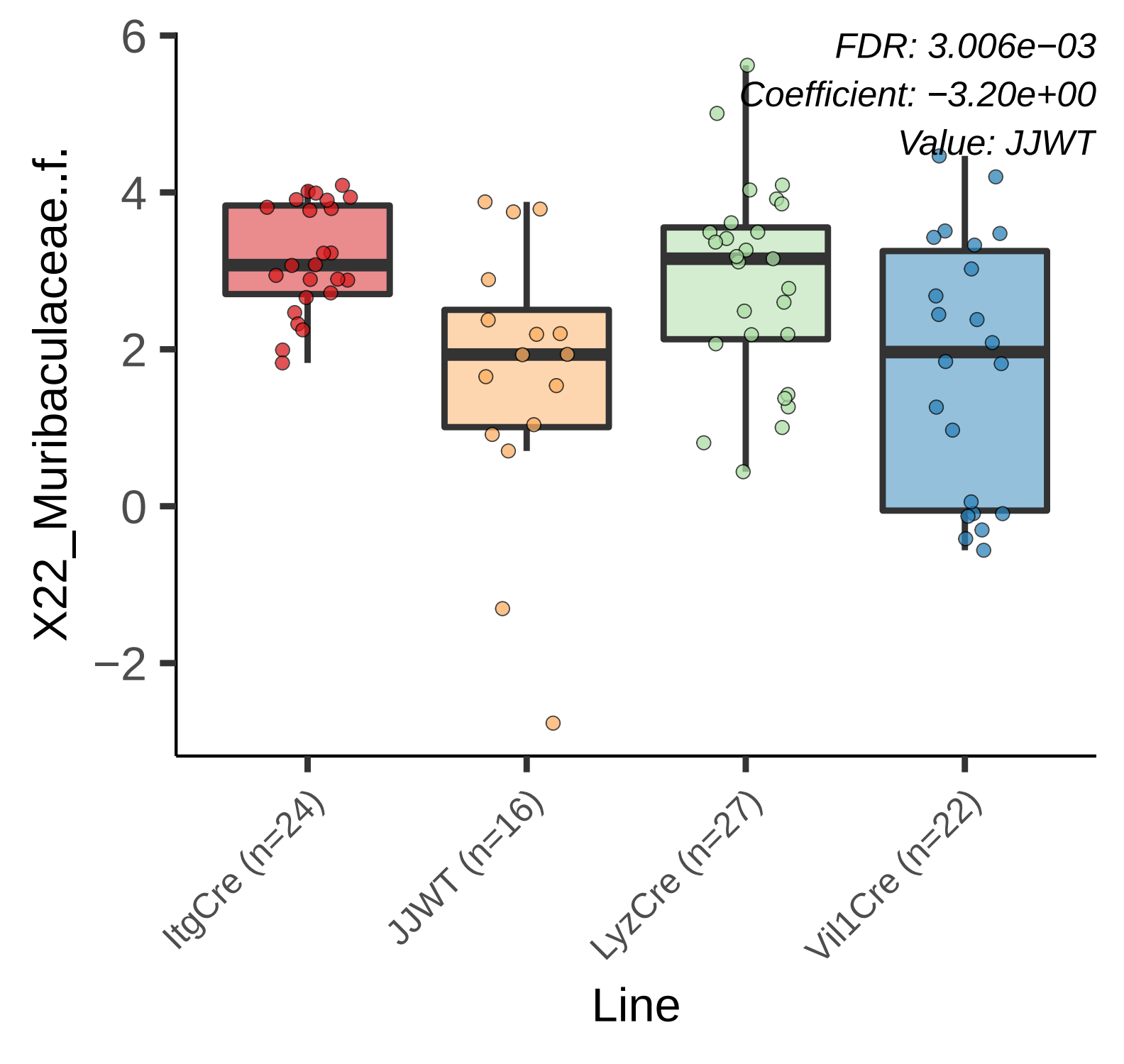

FDR: 2.441e−03
Coefficient: −2.25e+00
Value: JJWT
X97_GCA.900066575.
2
1
0
−1
−2
−3
ItgCre (n=24)
JJWT (n=16)
LyzCre (n=27)
Vil1Cre (n=22)
Line

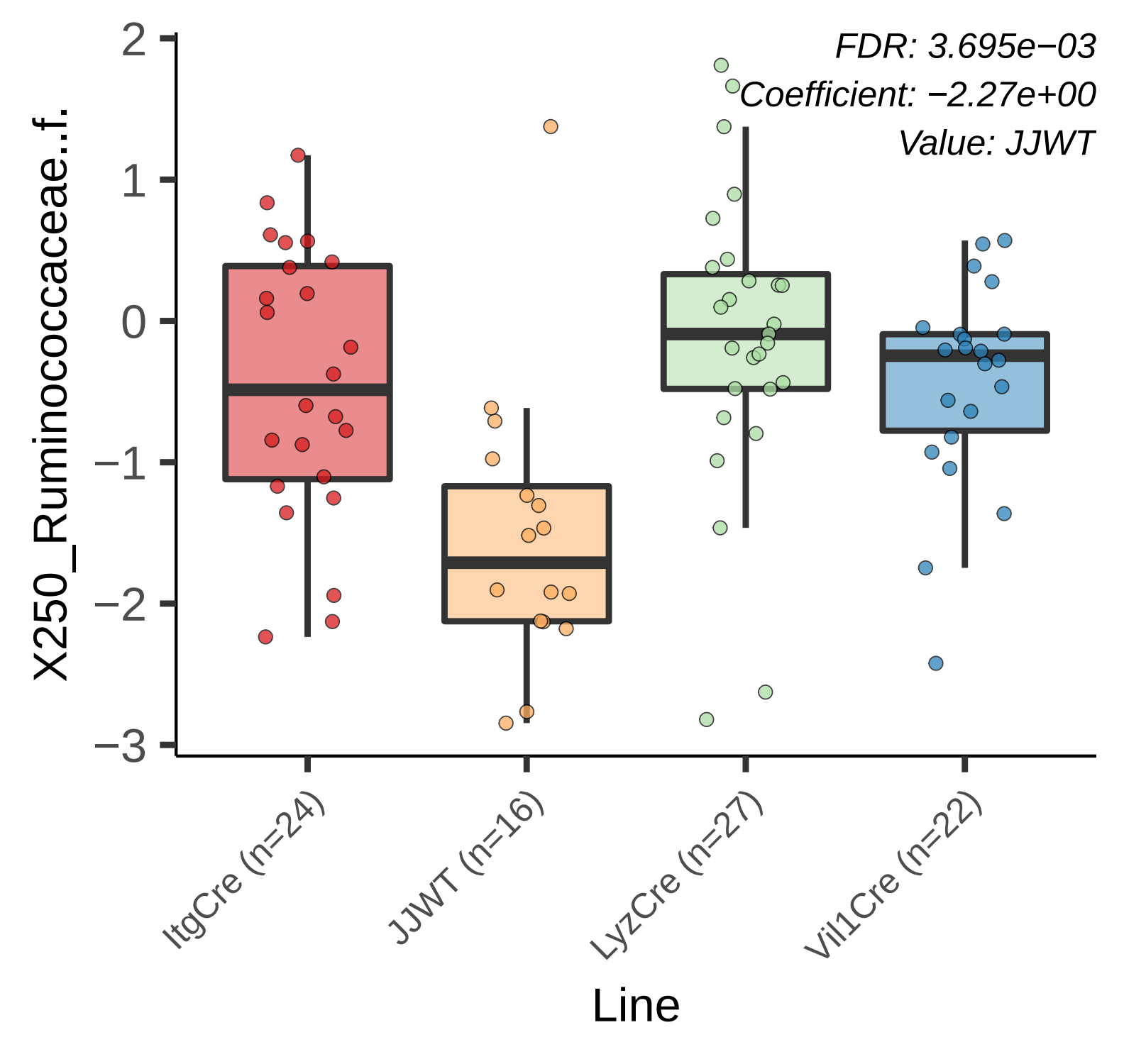

FDR: 2.549e−03
Coefficient: 2.05e+00
Value: LyzCre
X94_Lachnospiraceae_NK4A136_group.bacterium
2
0
−2
ItgCre (n=24)
JJWT (n=16)
LyzCre (n=27)
Vil1Cre (n=22)
Line

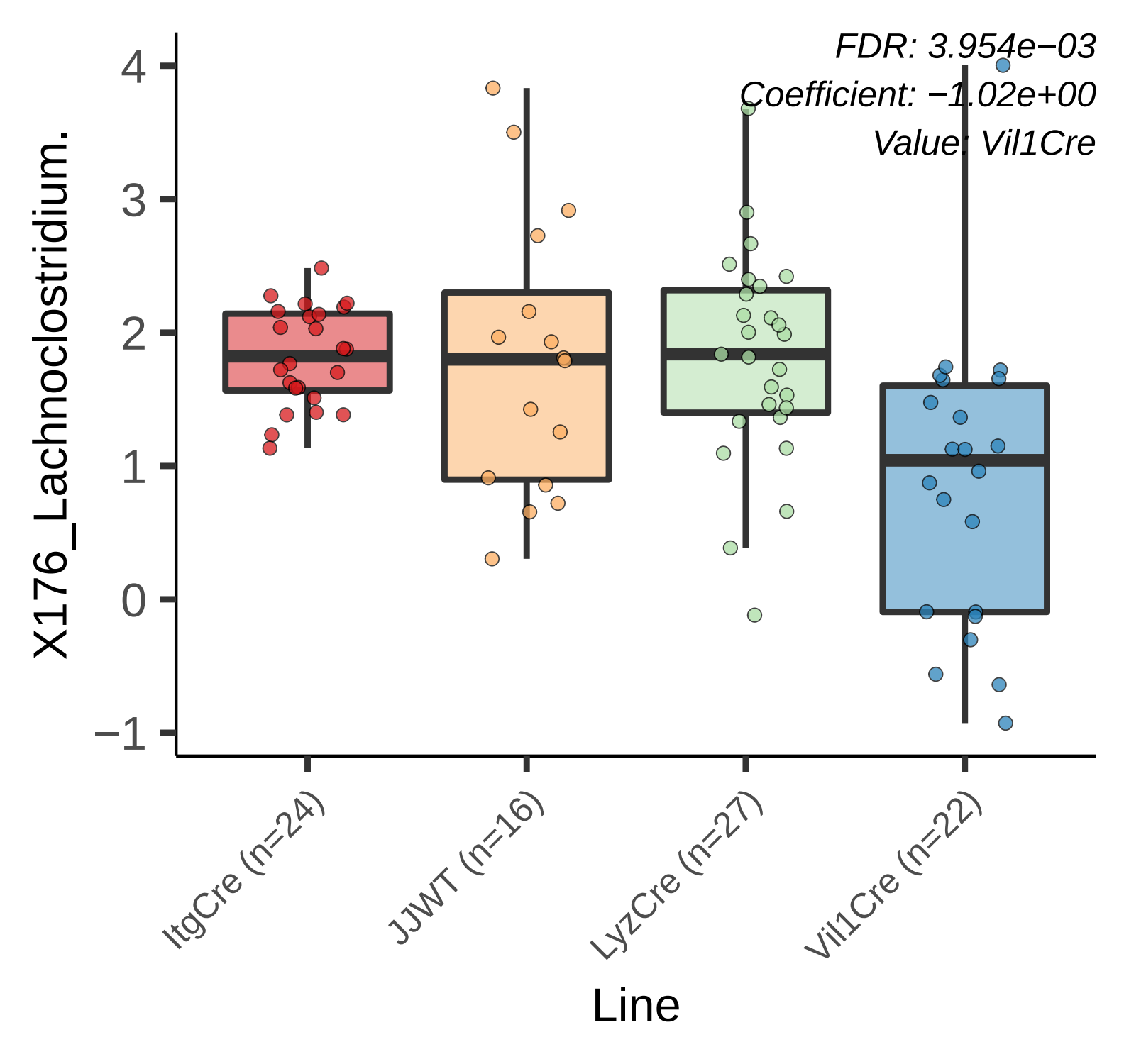

FDR: 2.568e−03
Coefficient: −1.53e+00
Value: Vil1Cre
X292_A2.
3
2
1
0
−1
−2
−3
ItgCre (n=24)
JJWT (n=16)
LyzCre (n=27)
Vil1Cre (n=22)
Line

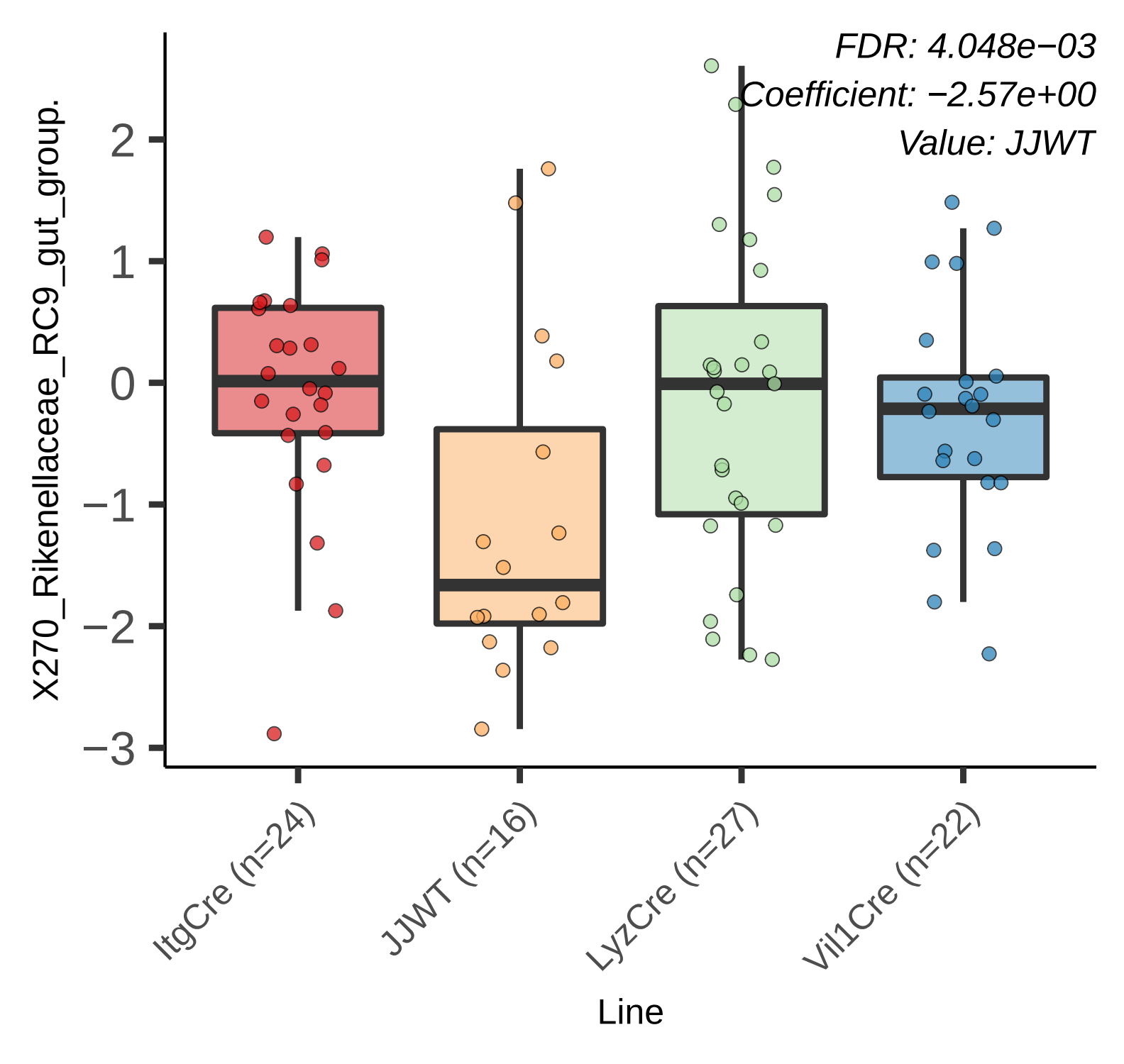

FDR: 2.787e−03
Coefficient: −1.73e+00
Value: Vil1Cre
X19_Muribaculaceae..f.
6
4
2
0
ItgCre (n=24)
JJWT (n=16)
LyzCre (n=27)
Vil1Cre (n=22)
Line

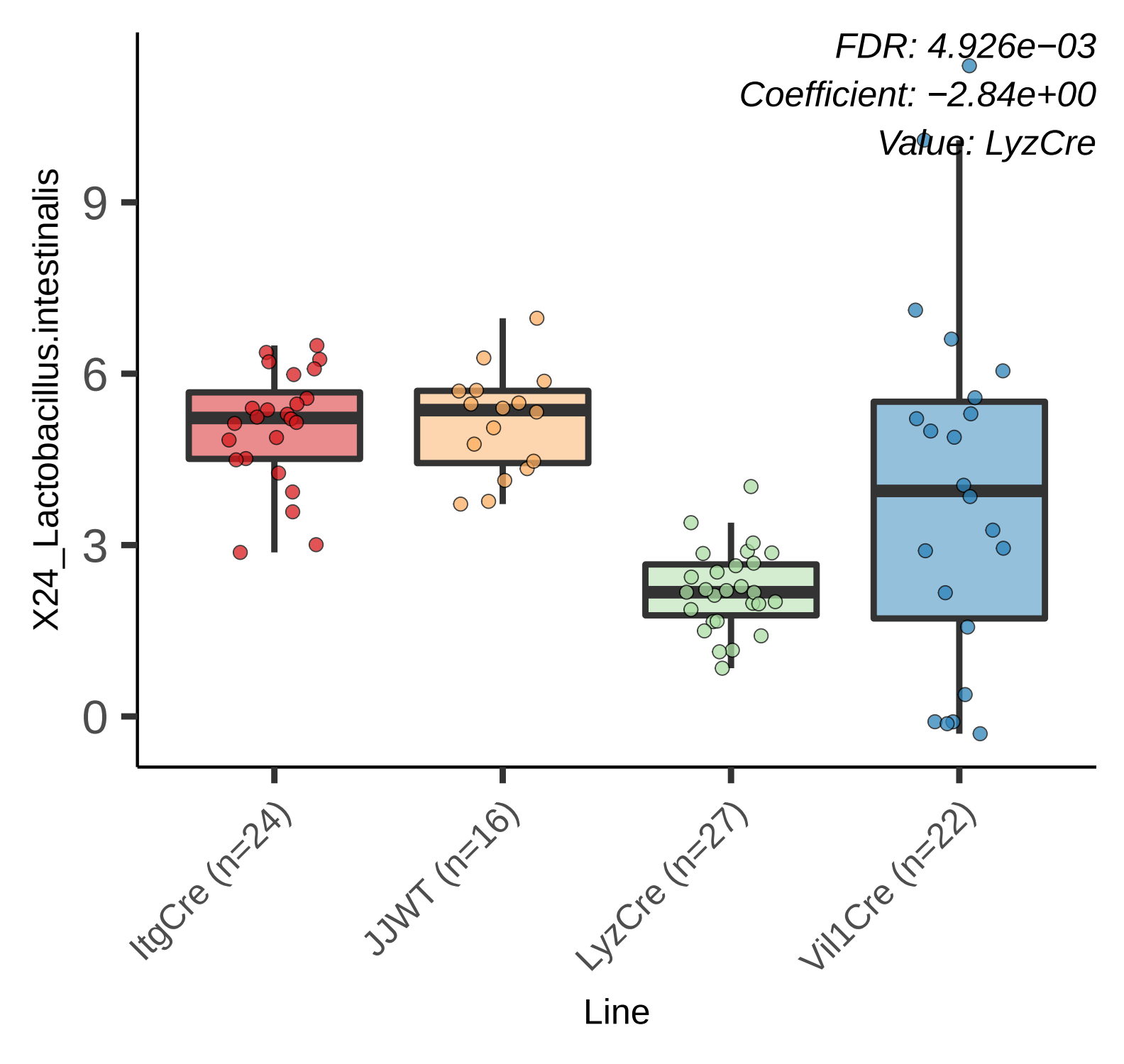

FDR: 2.908e−03
Coefficient: −2.31e+00
Value: JJWT
X88_Anaerotruncus.
3
2
1
0
−1
−2
ItgCre (n=24)
JJWT (n=16)
LyzCre (n=27)
Vil1Cre (n=22)
Line

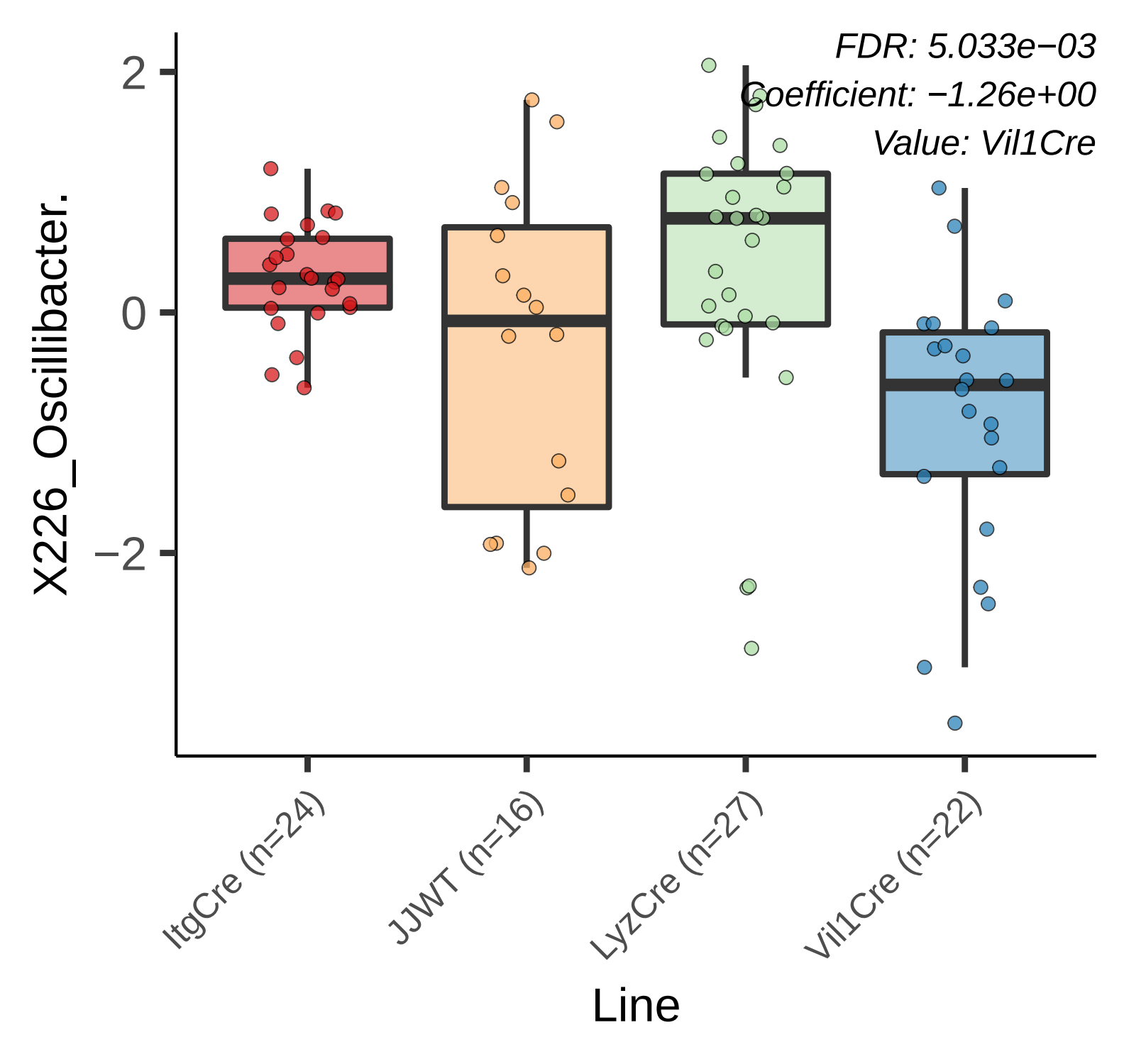

FDR: 3.006e−03
Coefficient: −3.20e+00
Value: JJWT
X22_Muribaculaceae..f.
6
4
2
0
−2
ItgCre (n=24)
JJWT (n=16)
LyzCre (n=27)
Vil1Cre (n=22)
Line

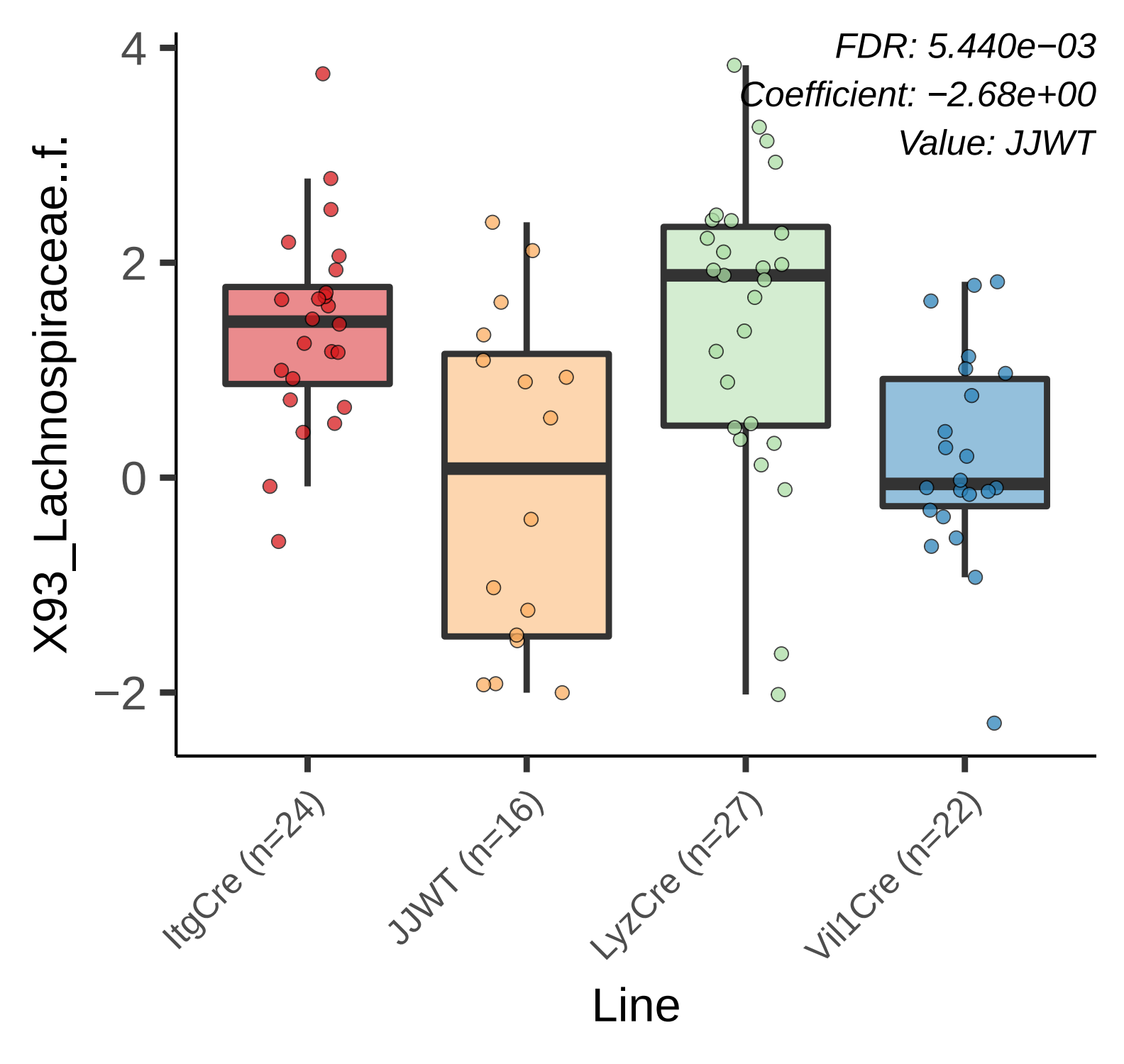

FDR: 3.695e−03
Coefficient: −2.27e+00
Value: JJWT
X250_Ruminococcaceae..f.
2
1
0
−1
−2
−3
ItgCre (n=24)
JJWT (n=16)
LyzCre (n=27)
Vil1Cre (n=22)
Line

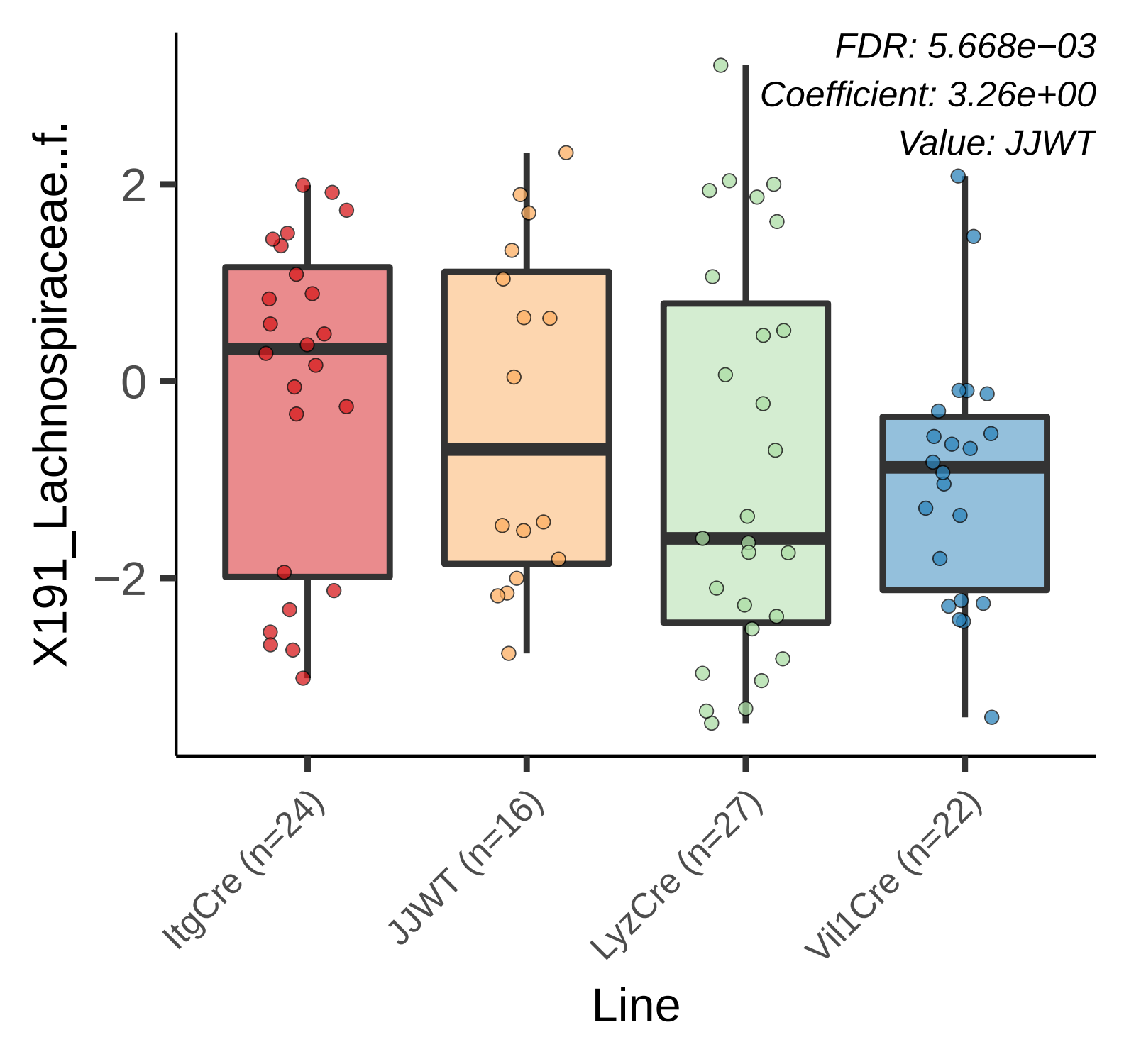

FDR: 3.954e−03
Coefficient: −1.02e+00
Value: Vil1Cre
X176_Lachnoclostridium.
4
3
2
1
0
−1
ItgCre (n=24)
JJWT (n=16)
LyzCre (n=27)
Vil1Cre (n=22)
Line

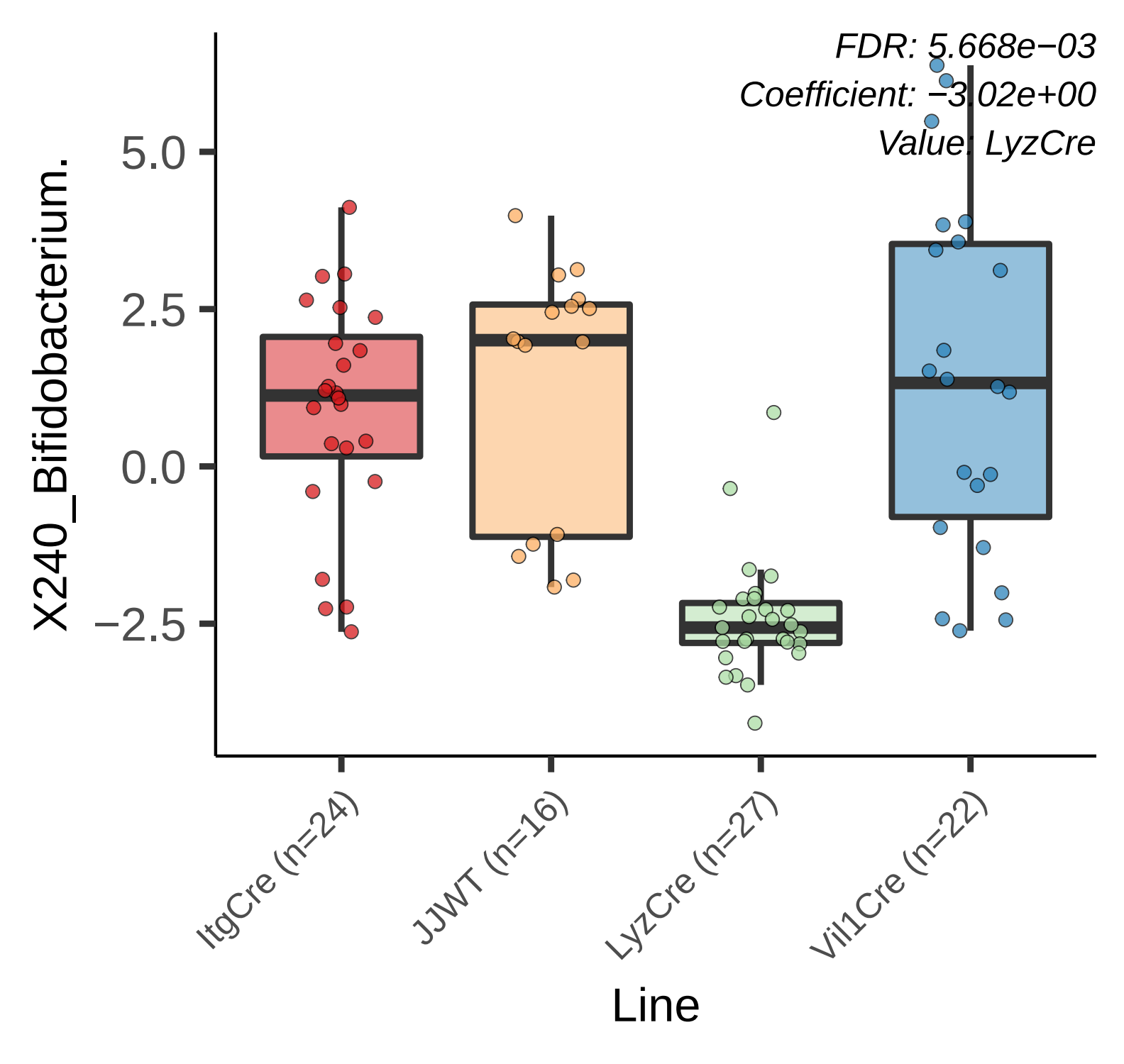

FDR: 4.048e−03
Coefficient: −2.57e+00
Value: JJWT
X270_Rikenellaceae_RC9_gut_group.
2
1
0
−1
−2
−3
ItgCre (n=24)
JJWT (n=16)
LyzCre (n=27)
Vil1Cre (n=22)
Line

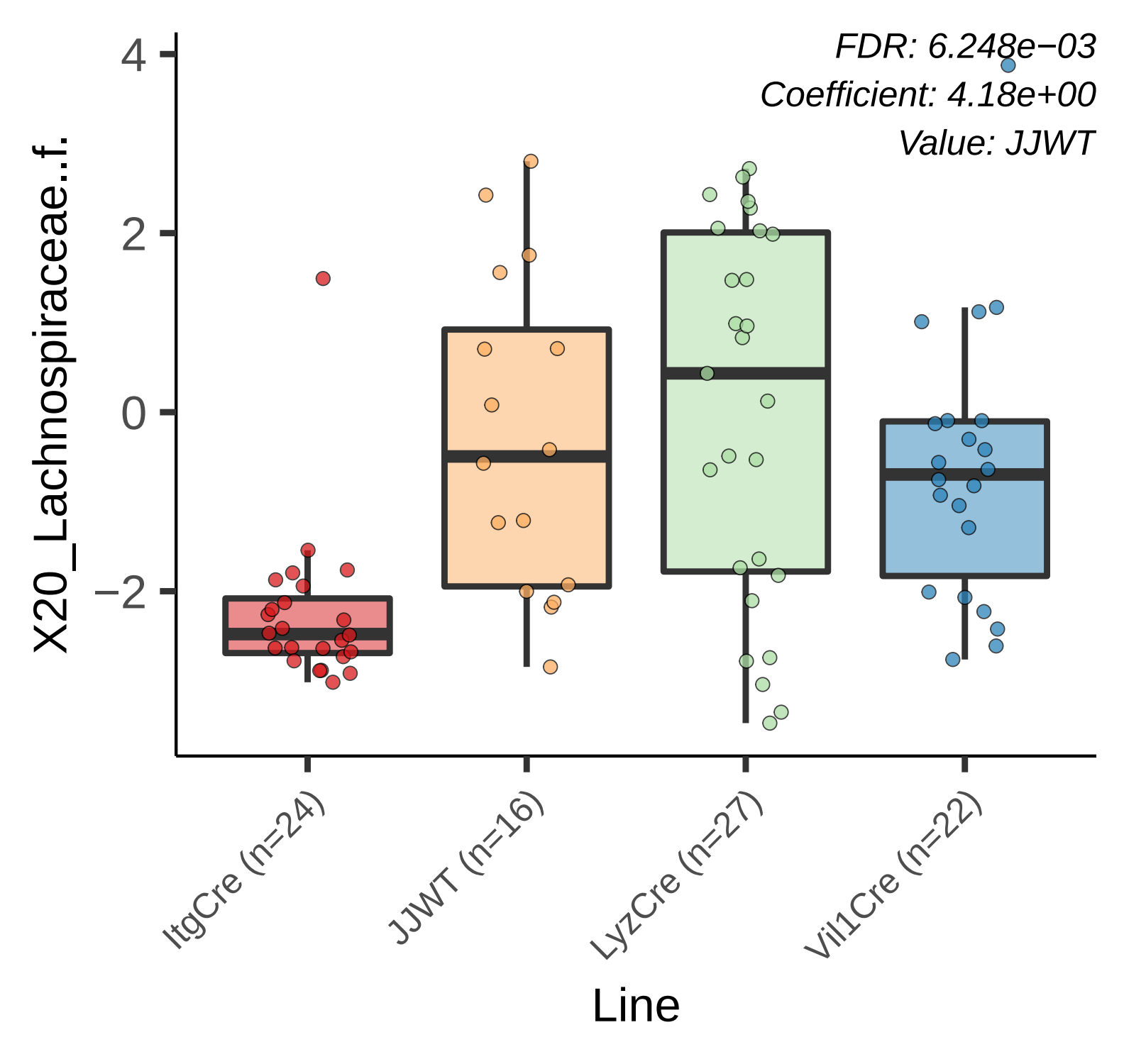

FDR: 4.926e−03
Coefficient: −2.84e+00
Value: LyzCre
X24_Lactobacillus.intestinalis
0
3
6
9
ItgCre (n=24)
JJWT (n=16)
LyzCre (n=27)
Vil1Cre (n=22)
Line

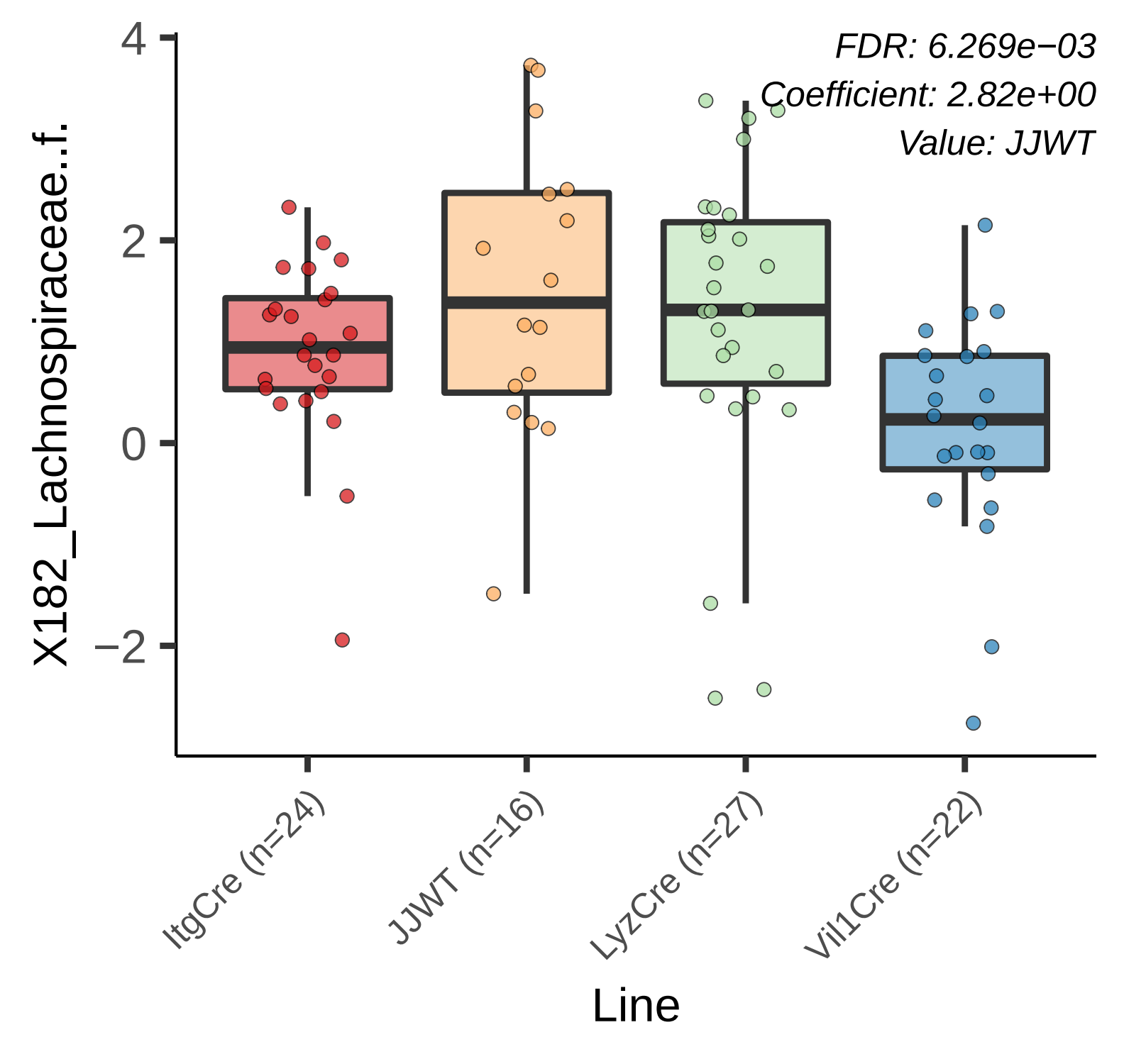

FDR: 5.033e−03
Coefficient: −1.26e+00
Value: Vil1Cre
X226_Oscillibacter.
2
0
−2
ItgCre (n=24)
JJWT (n=16)
LyzCre (n=27)
Vil1Cre (n=22)
Line

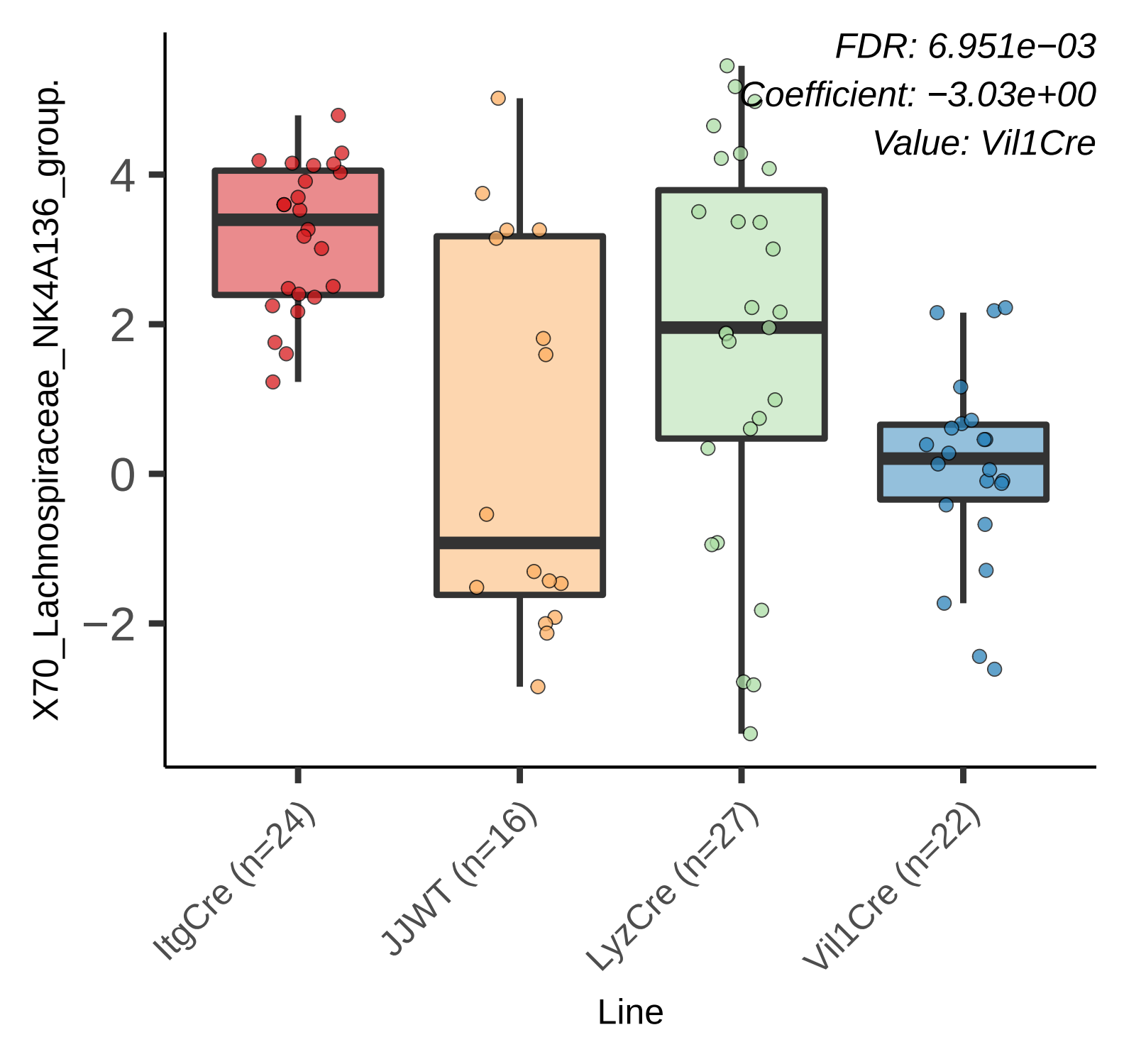

FDR: 5.440e−03
Coefficient: −2.68e+00
Value: JJWT
X93_Lachnospiraceae..f.
4
2
0
−2
ItgCre (n=24)
JJWT (n=16)
LyzCre (n=27)
Vil1Cre (n=22)
Line

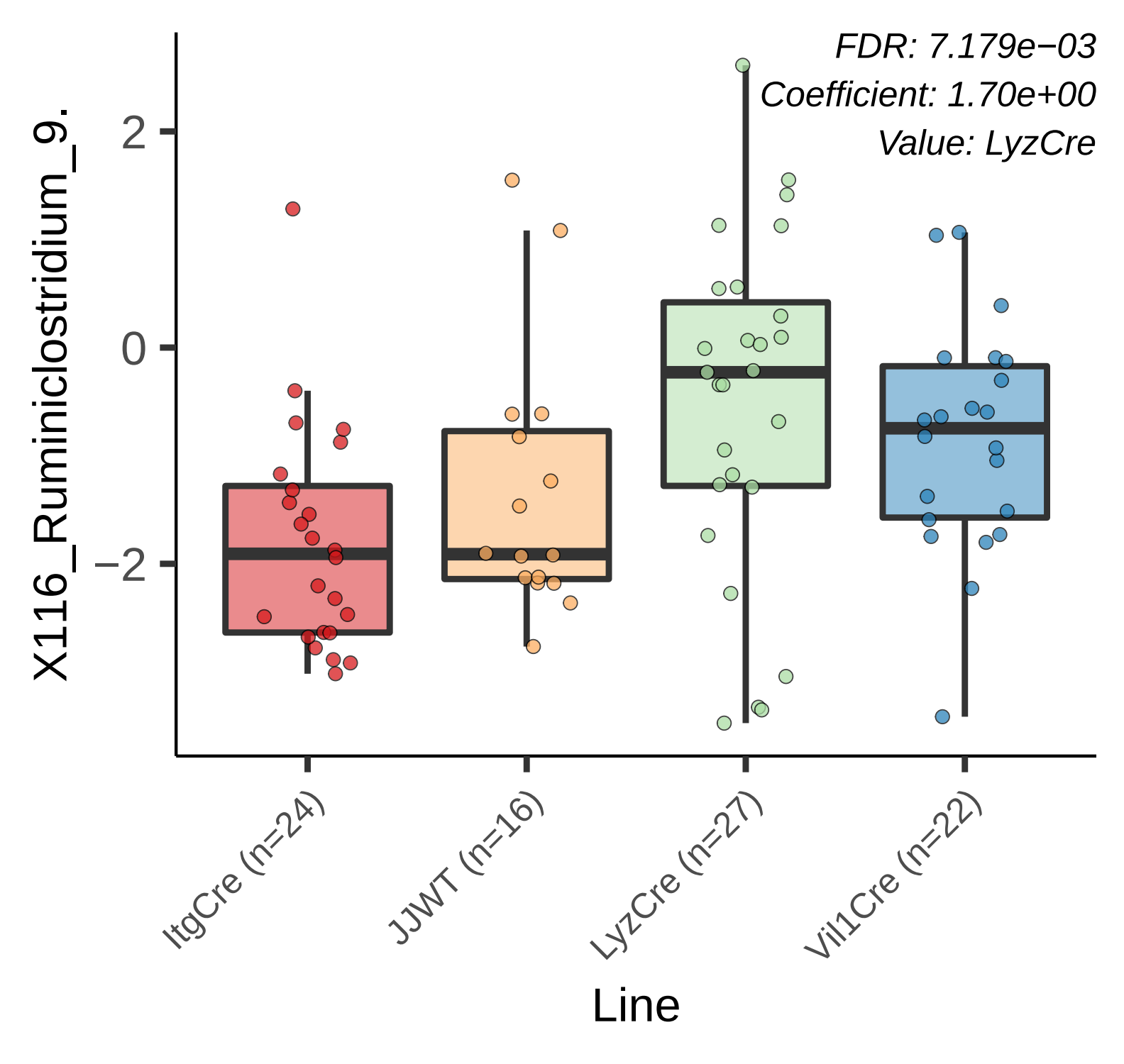

FDR: 5.668e−03
Coefficient: 3.26e+00
Value: JJWT
X191_Lachnospiraceae.f.
2
0
−2
ItgCre (n=24)
JJWT (n=16)
LyzCre (n=27)
Vil1Cre (n=22)
Line

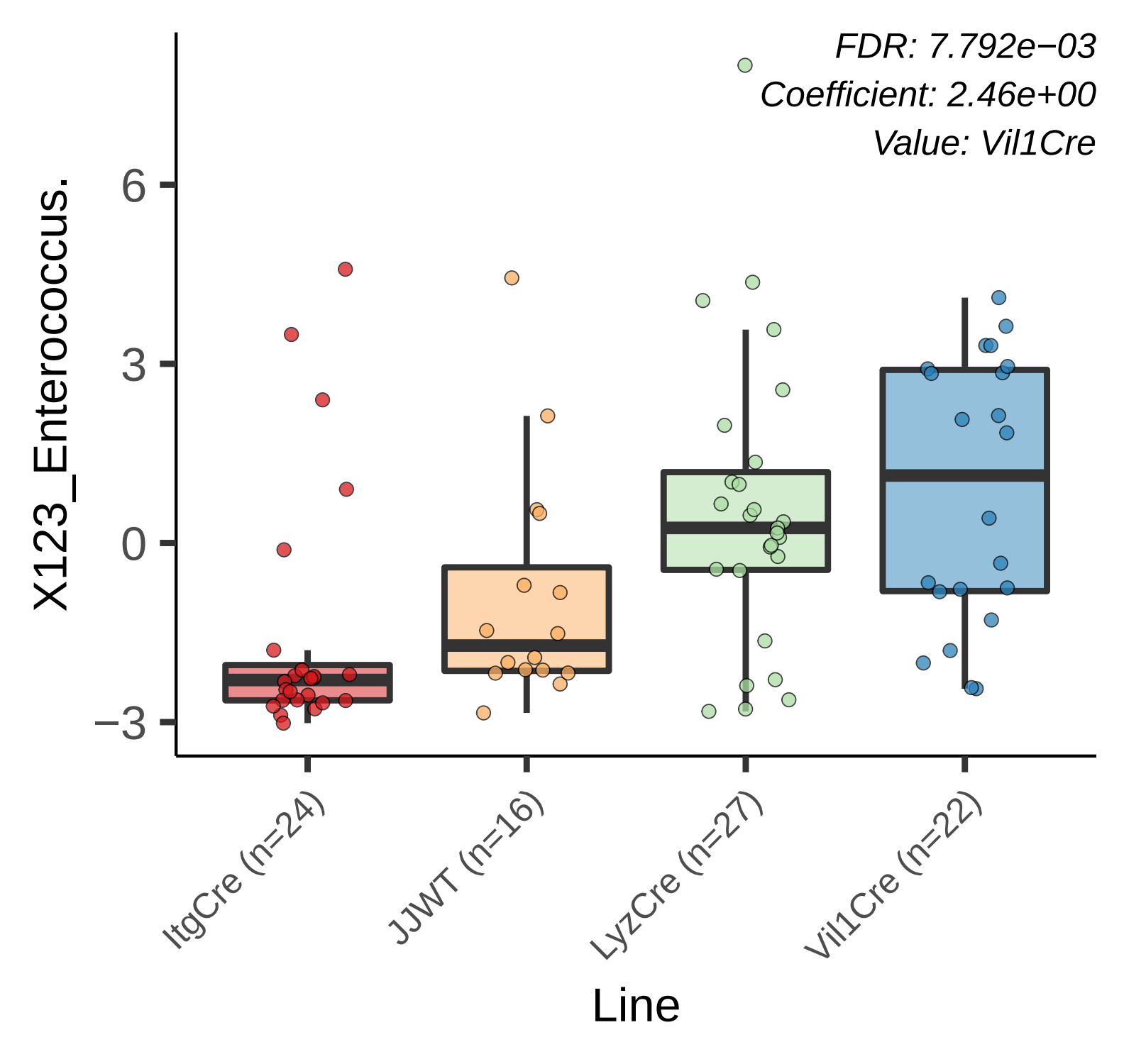

FDR: 5.668e−03
Coefficient: −3.02e+00
Value: LyzCre
X240_Bifidobacterium.
5.0
2.5
0.0
−2.5
ItgCre (n=24)
JJWT (n=16)
LyzCre (n=27)
Vil1Cre (n=22)
Line

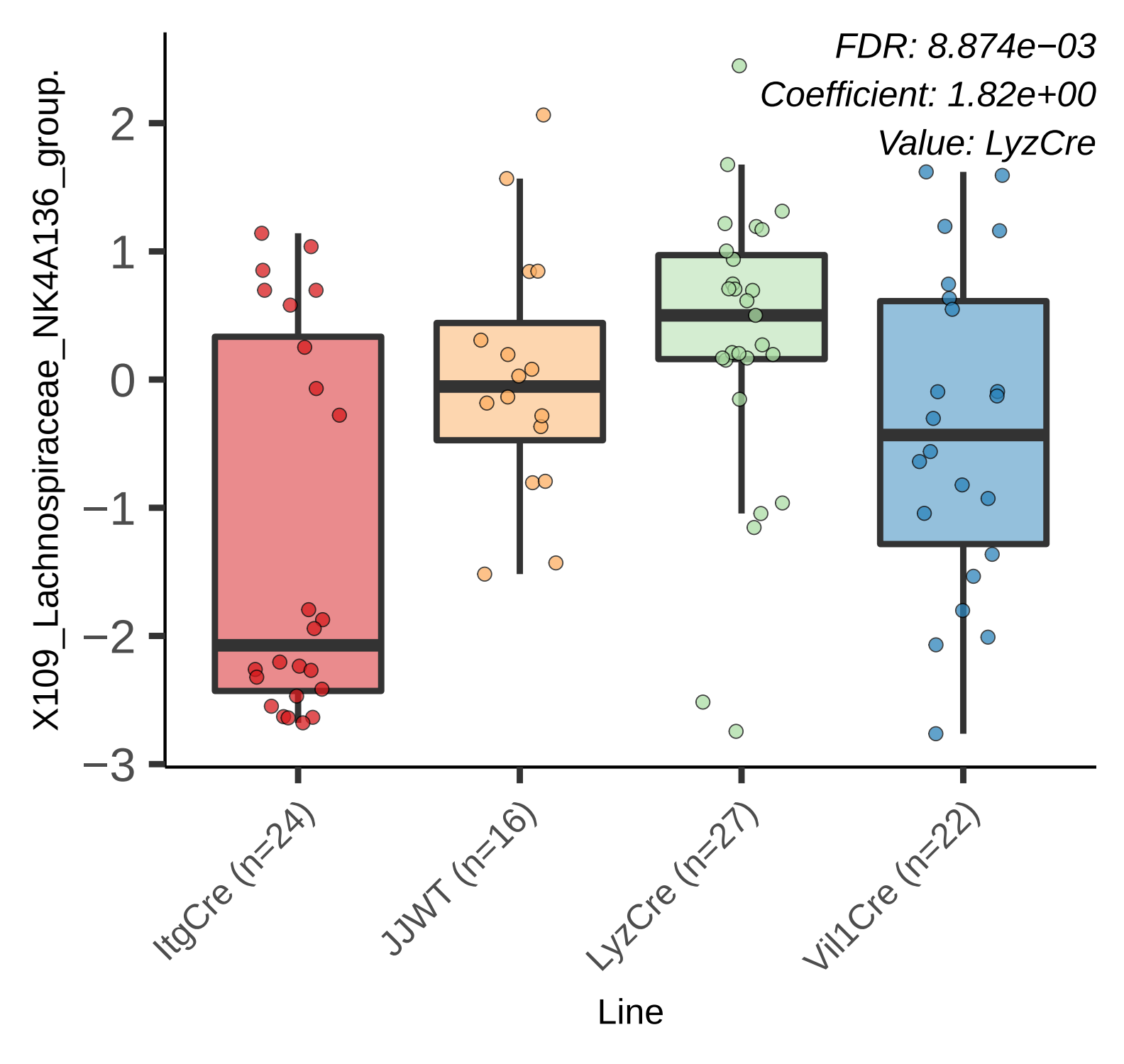

FDR: 6.248e−03
Coefficient: 4.18e+00
Value: JJWT
X20_Lachnospiraceae..f.
4
2
0
−2
ItgCre (n=24)
JJWT (n=16)
LyzCre (n=27)
Vil1Cre (n=22)
Line

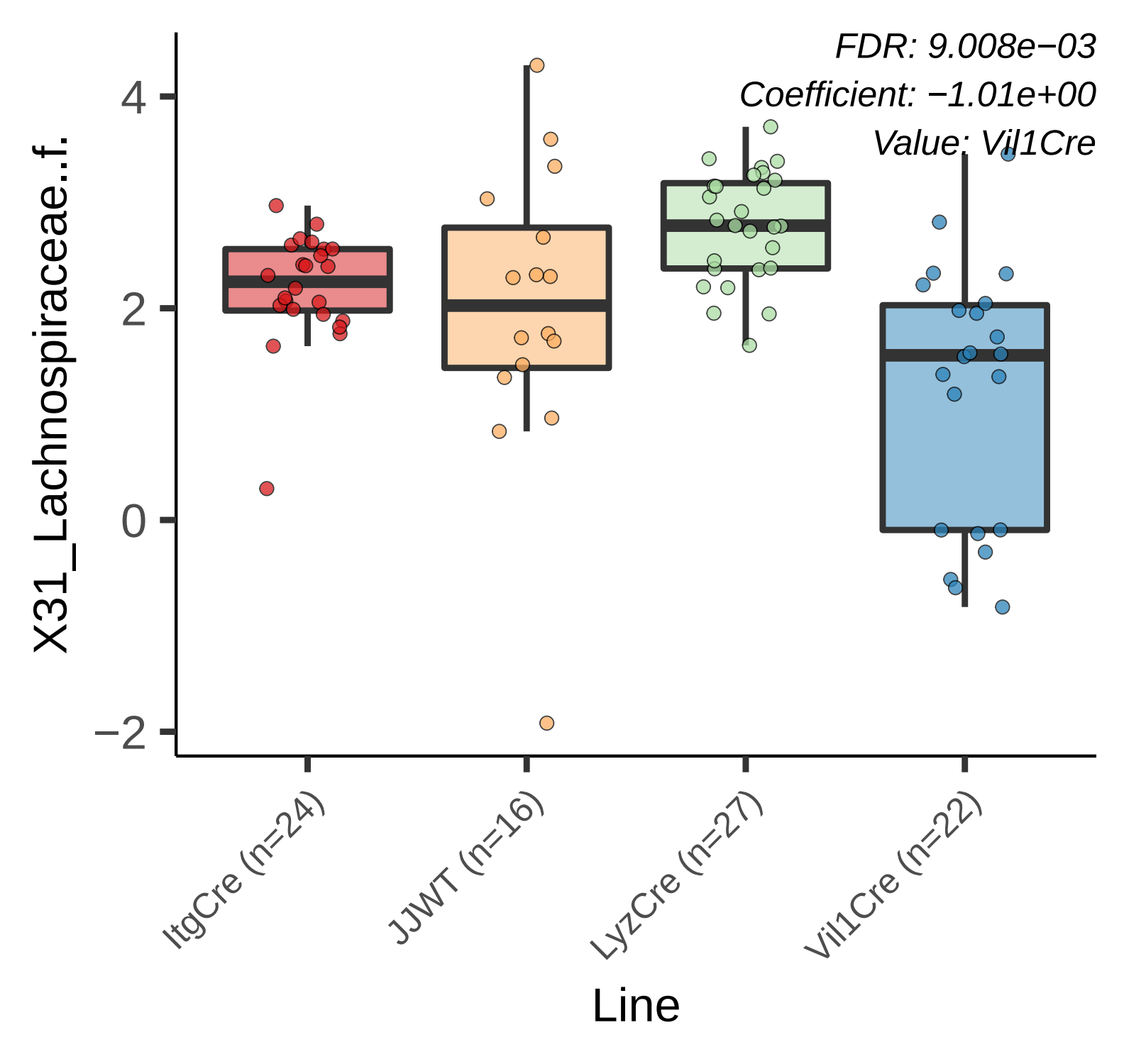

FDR: 6.269e−03
Coefficient: 2.82e+00
Value: JJWT
X182_Lachnospiraceae.f.
4
2
0
−2
ItgCre (n=24)
JJWT (n=16)
LyzCre (n=27)
Vil1Cre (n=22)
Line

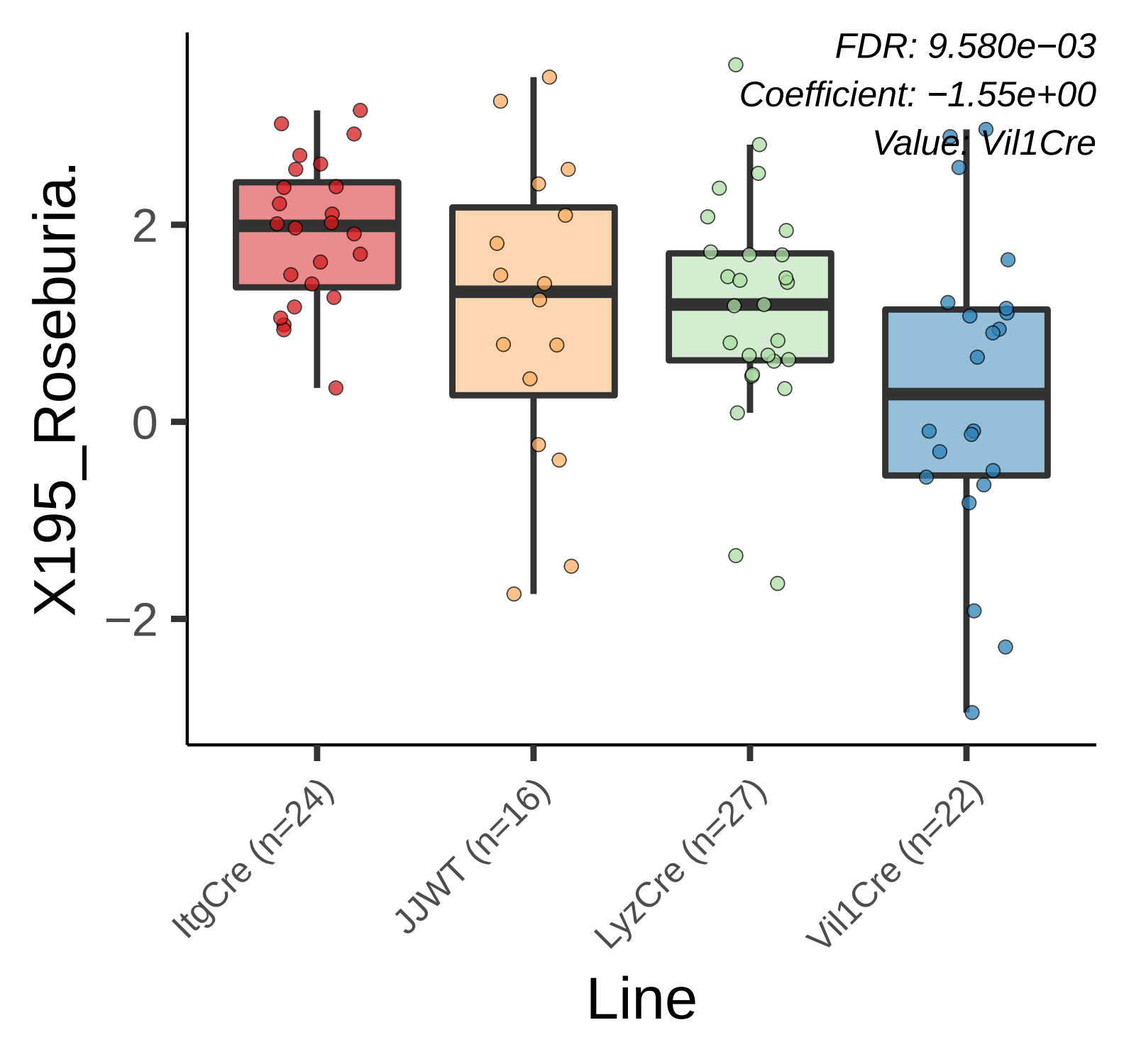

FDR: 6.951e−03
Coefficient: −3.03e+00
Value: Vil1Cre
X70_Lachnospiraceae_NK4A136_group.
4
2
0
−2
ItgCre (n=24)
JJWT (n=16)
LyzCre (n=27)
Vil1Cre (n=22)
Line

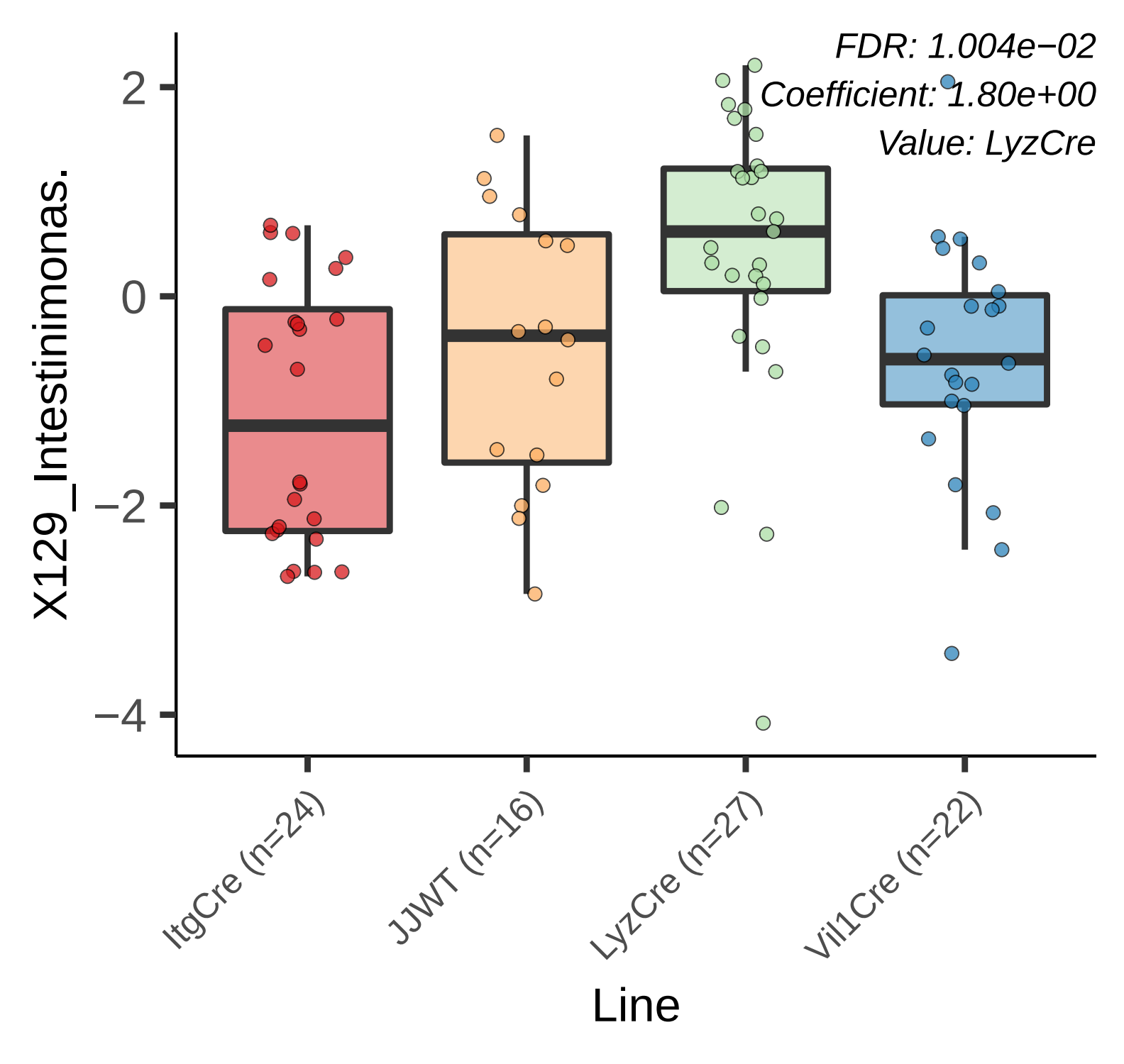

FDR: 7.179e−03
Coefficient: 1.70e+00
Value: LyzCre
X116_Ruminiclostridium_9.
2
0
−2
ItgCre (n=24)
JJWT (n=16)
LyzCre (n=27)
Vil1Cre (n=22)
Line

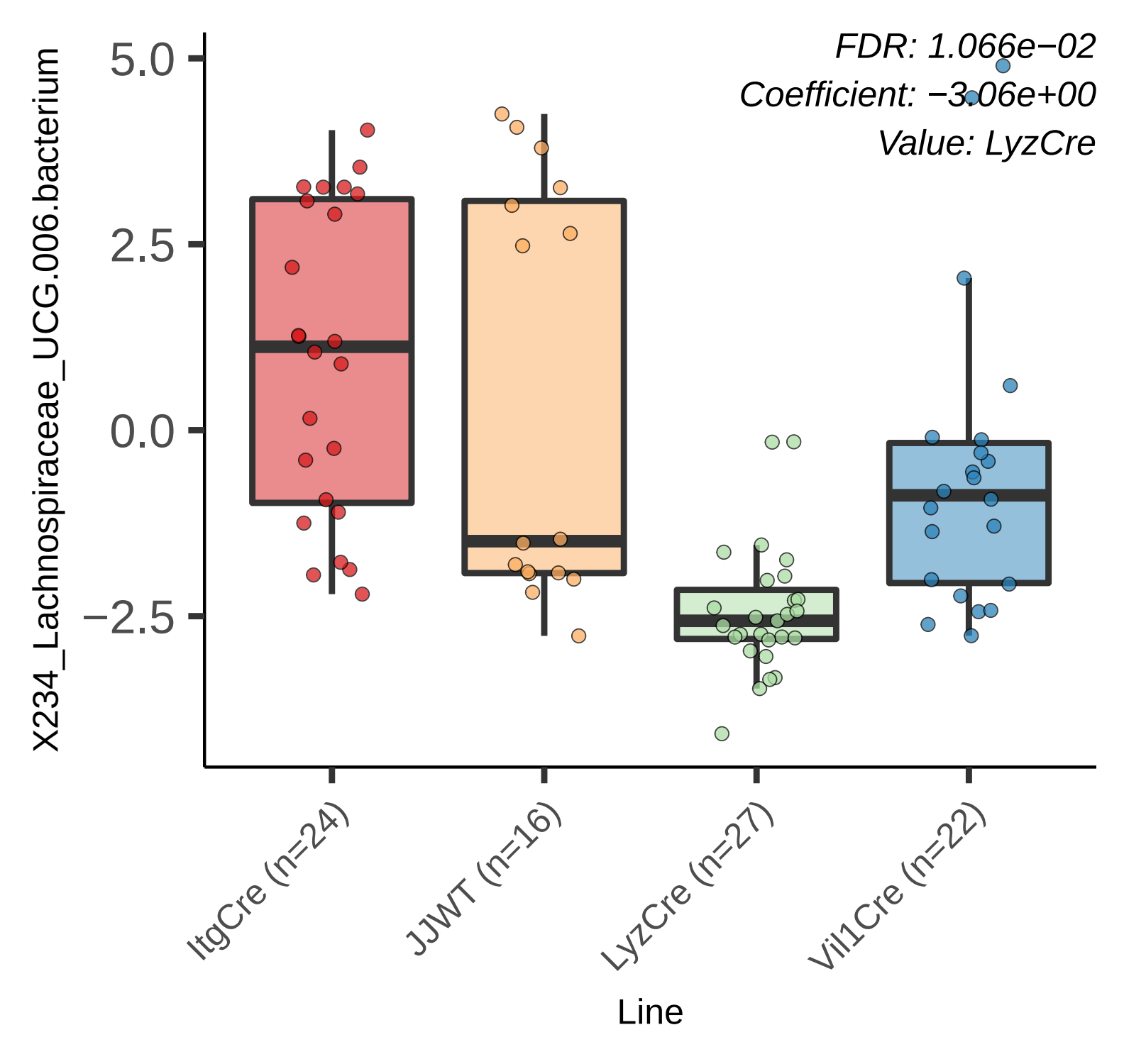

FDR: 7.792e−03
Coefficient: 2.46e+00
Value: Vil1Cre
X123_Enterococcus.
6
3
0
−3
ItgCre (n=24)
JJWT (n=16)
LyzCre (n=27)
Vil1Cre (n=22)
Line

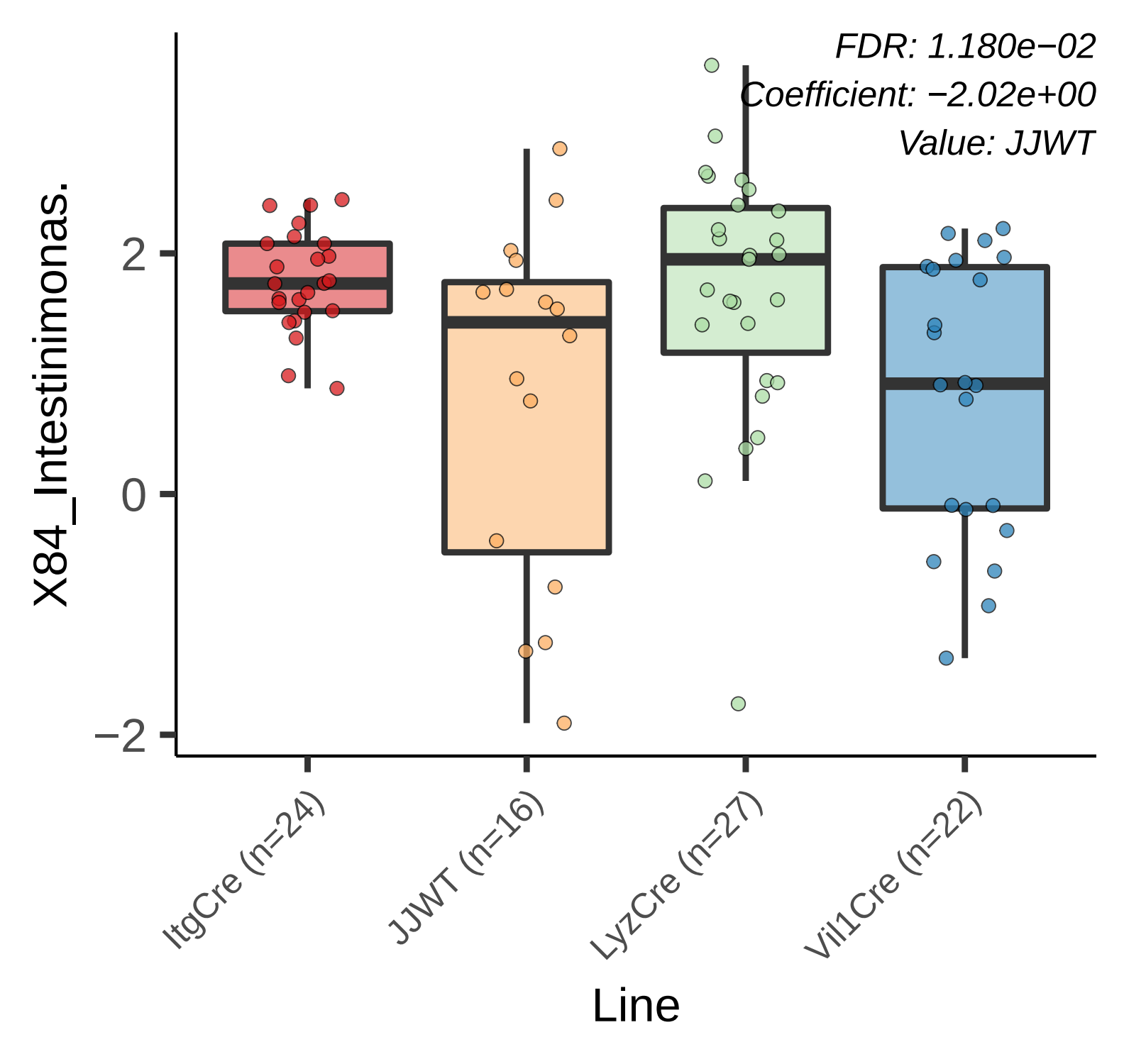

FDR: 8.874e−03
Coefficient: 1.82e+00
Value: LyzCre
X109_Lachnospiraceae_NK4A136_group.
2
1
0
−1
−2
−3
ItgCre (n=24)
JJWT (n=16)
LyzCre (n=27)
Vil1Cre (n=22)
Line

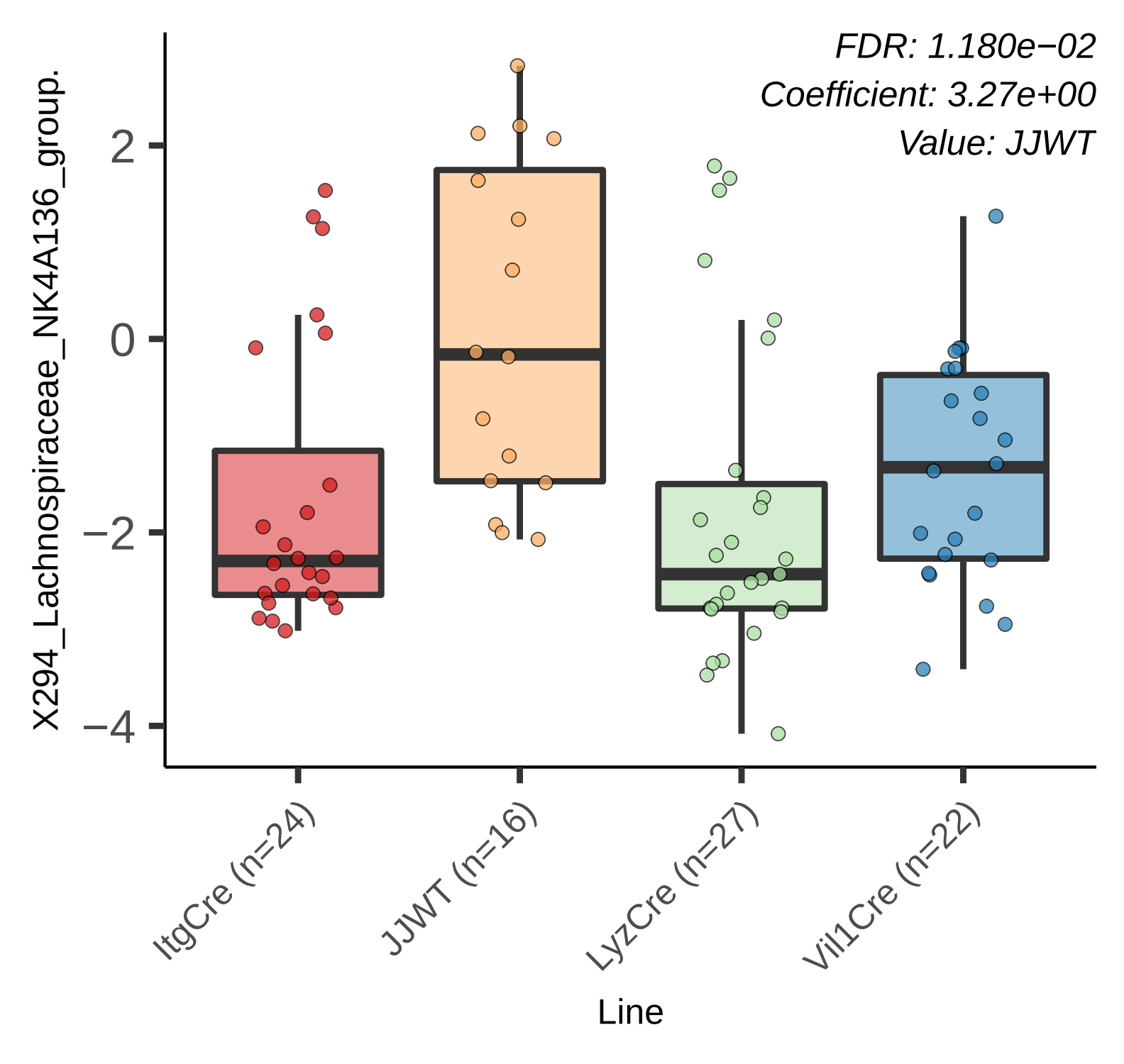

FDR: 9.008e−03
Coefficient: −1.01e+00
Value: Vil1Cre
X31_Lachnospiraceae..f.
4
2
0
−2
ItgCre (n=24)
JJWT (n=16)
LyzCre (n=27)
Vil1Cre (n=22)
Line

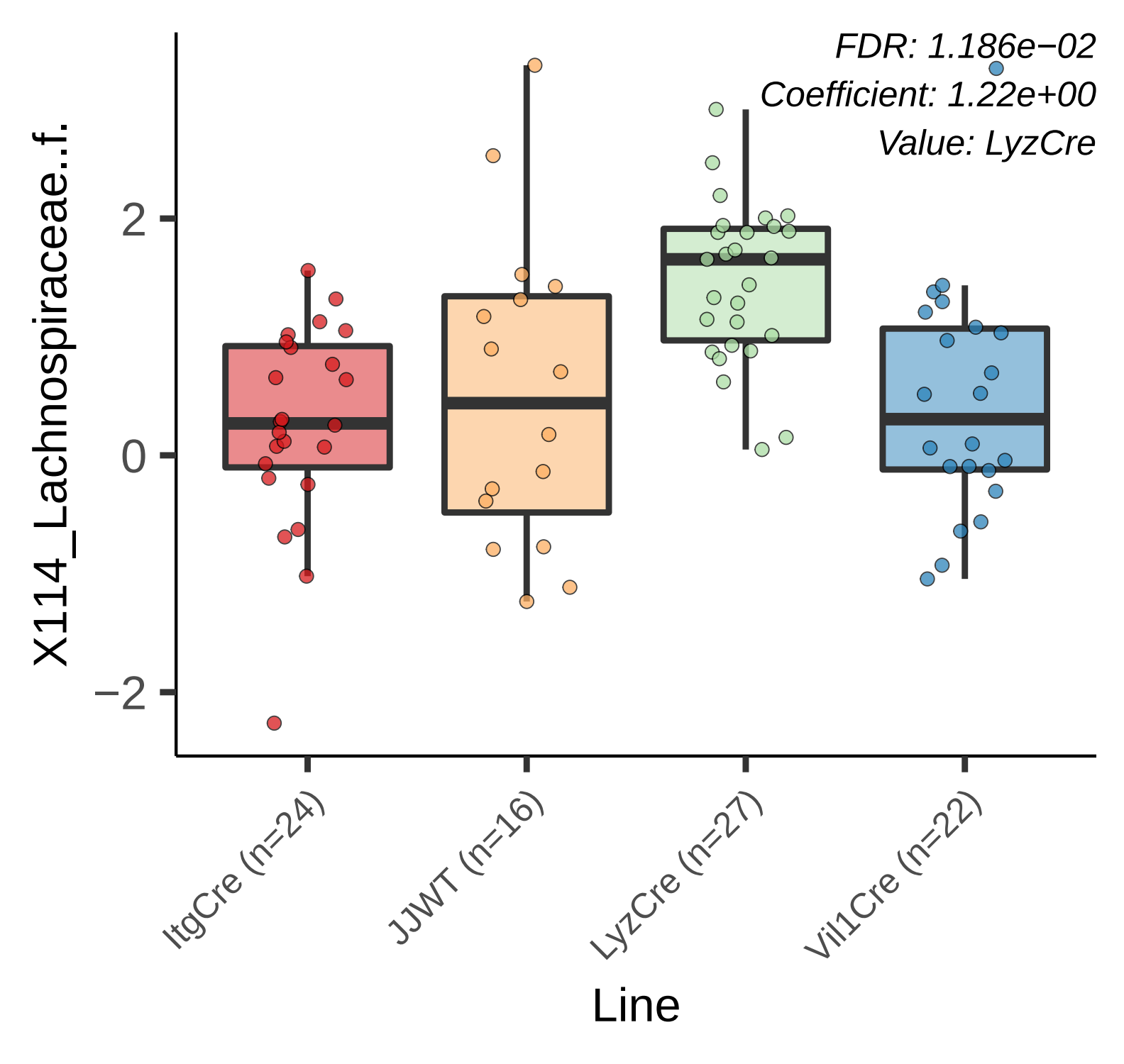

FDR: 9.580e−03
Coefficient: −1.55e+00
Value: Vil1Cre
X195_Roseburia.
2
0
−2
ItgCre (n=24)
JJWT (n=16)
LyzCre (n=27)
Vil1Cre (n=22)
Line

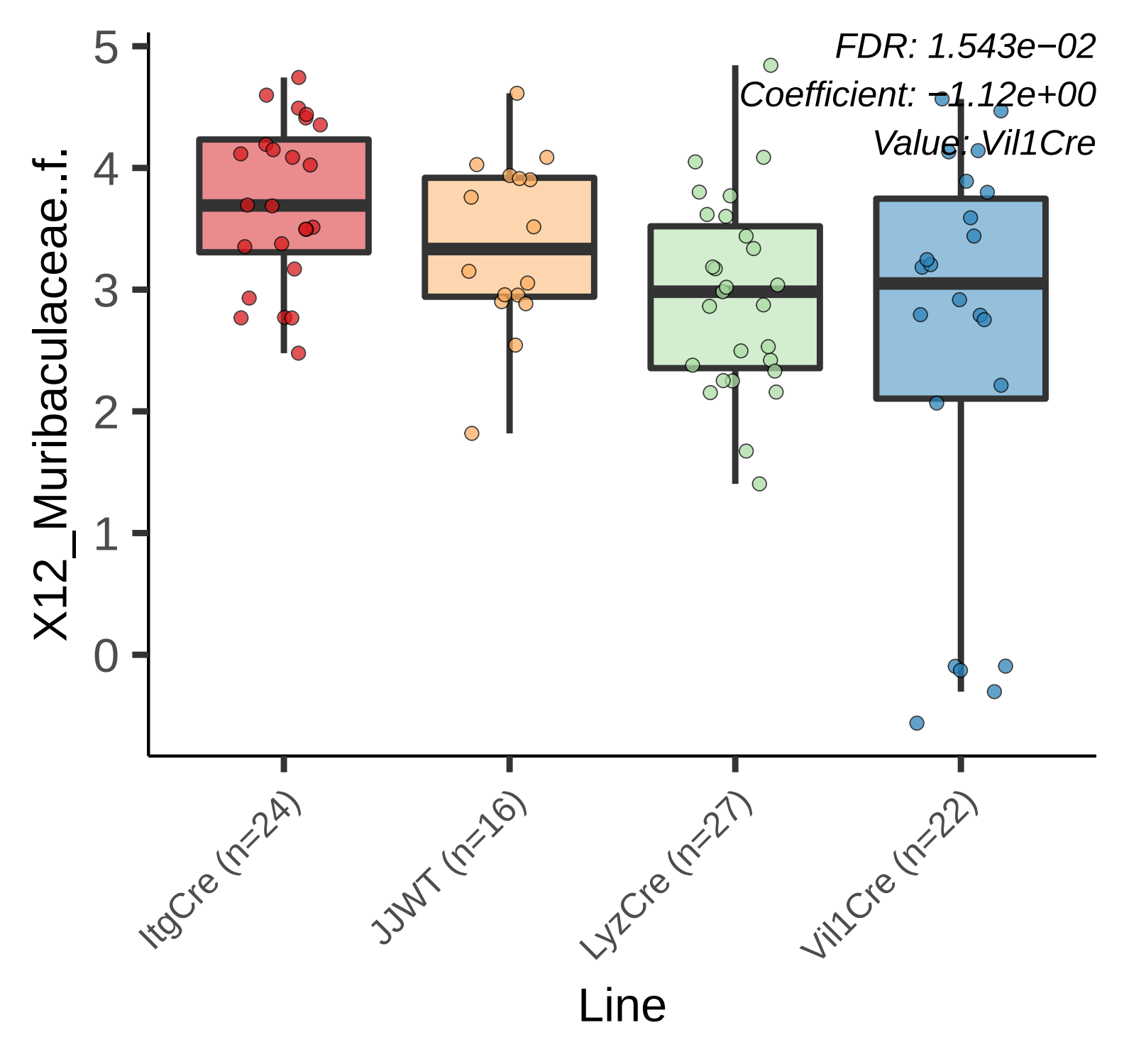

FDR: 1.004e−02
Coefficient: 1.80e+00
Value: LyzCre
X129_Intestinimonas.
2
0
−2
−4
ItgCre (n=24)
JJWT (n=16)
LyzCre (n=27)
Vil1Cre (n=22)
Line

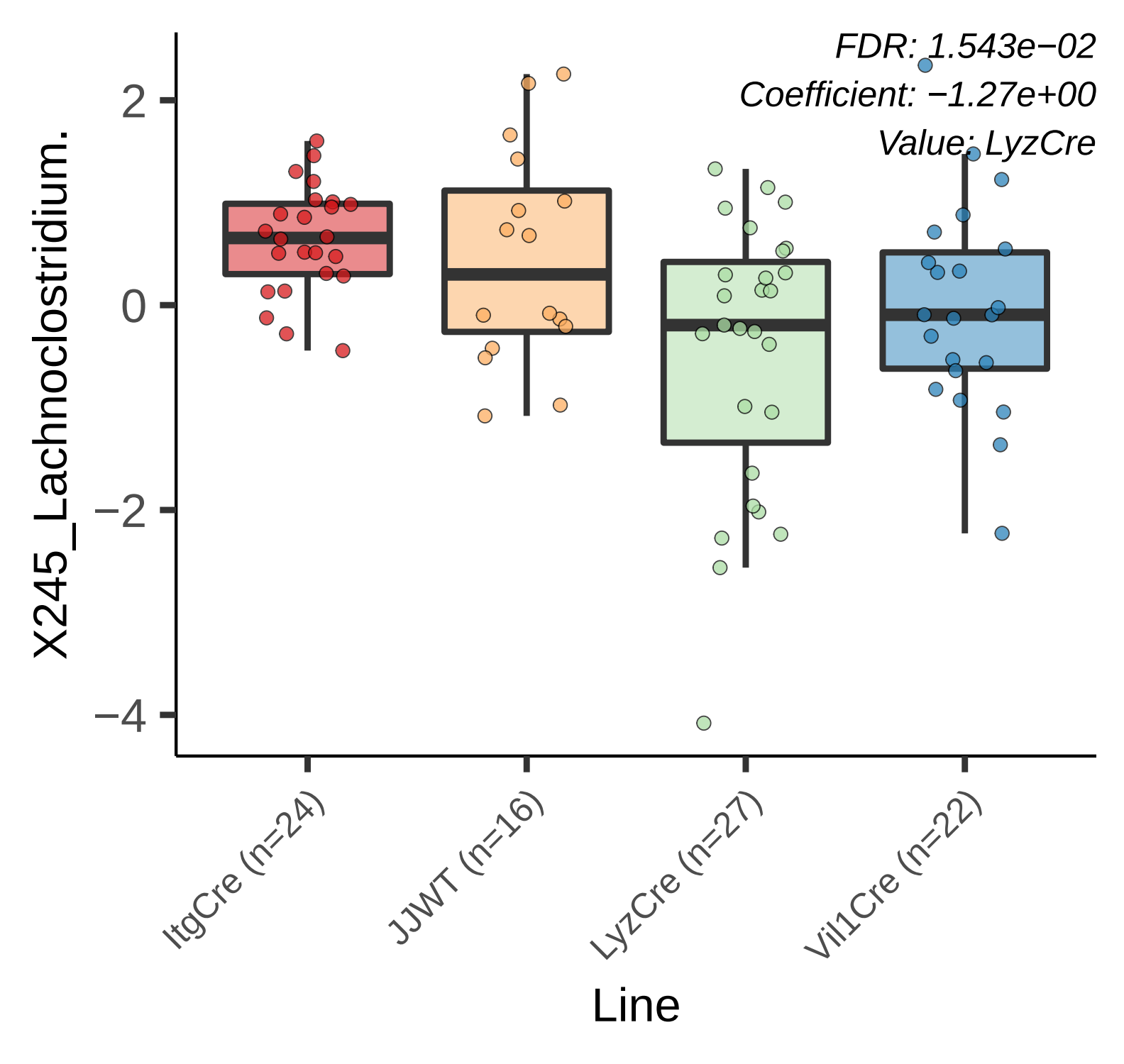

FDR: 1.066e−02
Coefficient: −3.06e+00
Value: LyzCre
X234_Lachnospiraceae_UCG.006.bacterium
5.0
2.5
0.0
−2.5
ItgCre (n=24)
JJWT (n=16)
LyzCre (n=27)
Vil1Cre (n=22)
Line

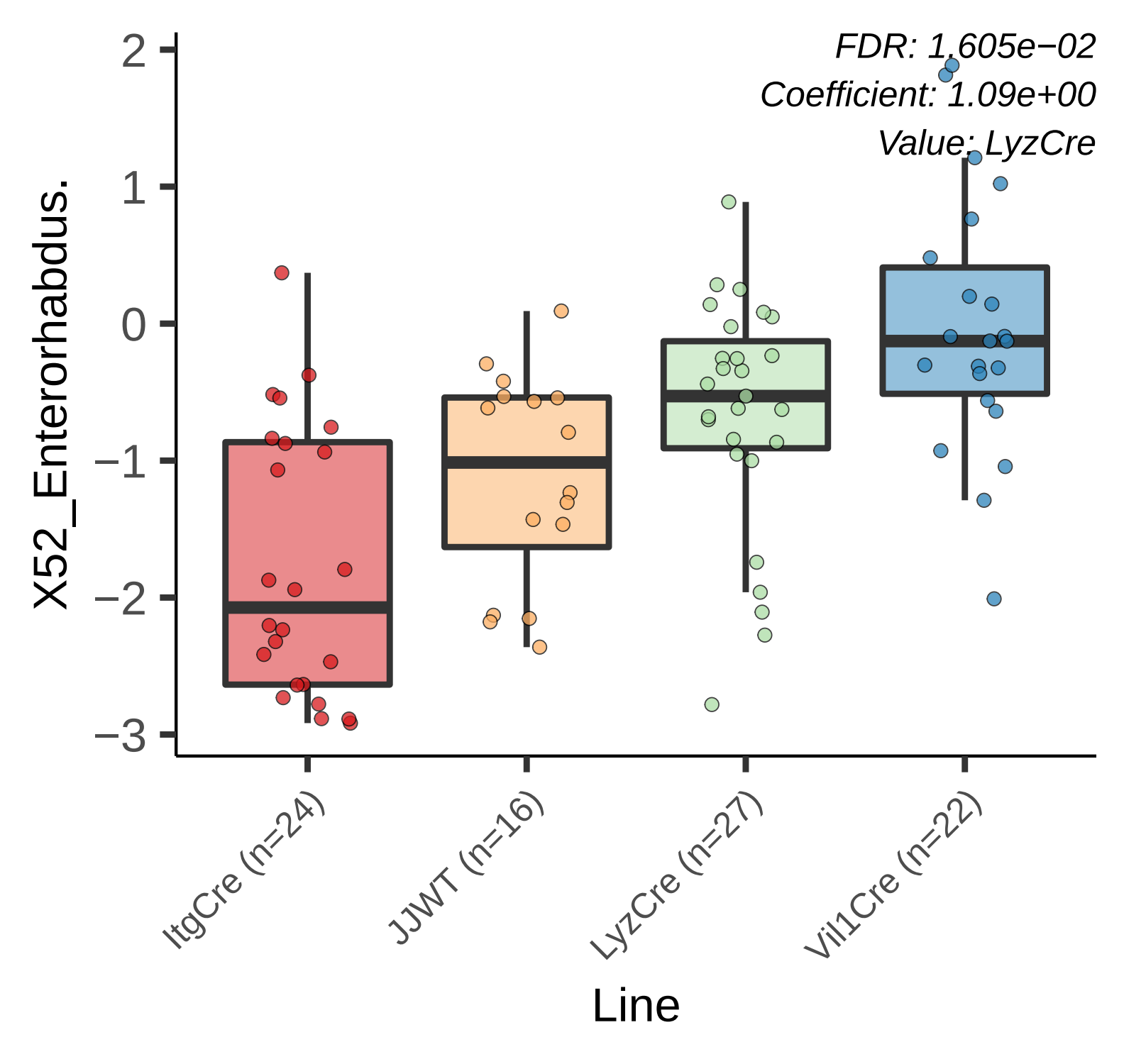

FDR: 1.180e−02
Coefficient: −2.02e+00
Value: JJWT
X84_Intestinimonas.
2
0
−2
ItgCre (n=24)
JJWT (n=16)
LyzCre (n=27)
Vil1Cre (n=22)
Line

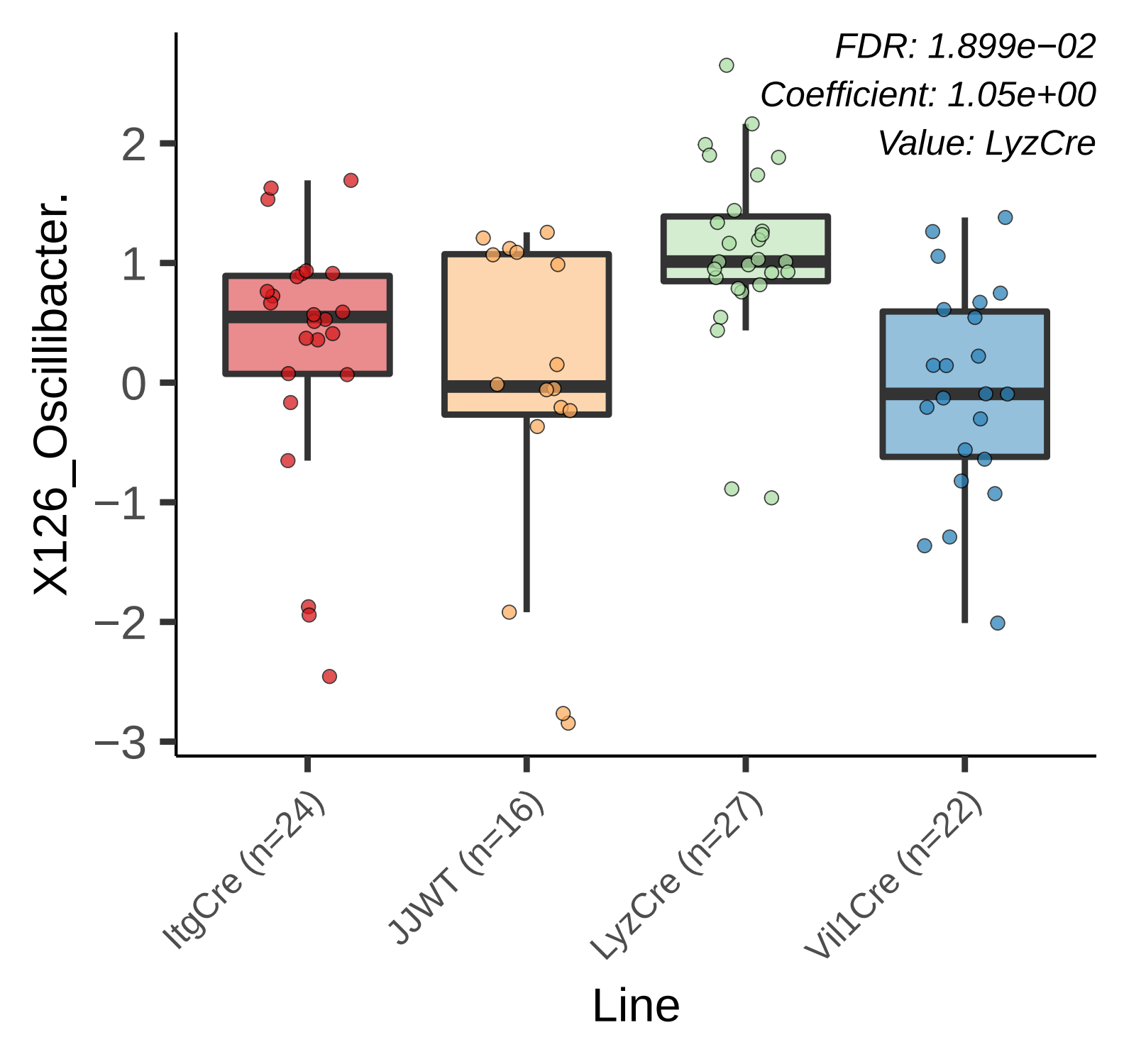

FDR: 1.180e−02
Coefficient: 3.27e+00
Value: JJWT
X294_Lachnospiraceae_NK4A136_group.
2
0
−2
−4
ItgCre (n=24)
JJWT (n=16)
LyzCre (n=27)
Vil1Cre (n=22)
Line

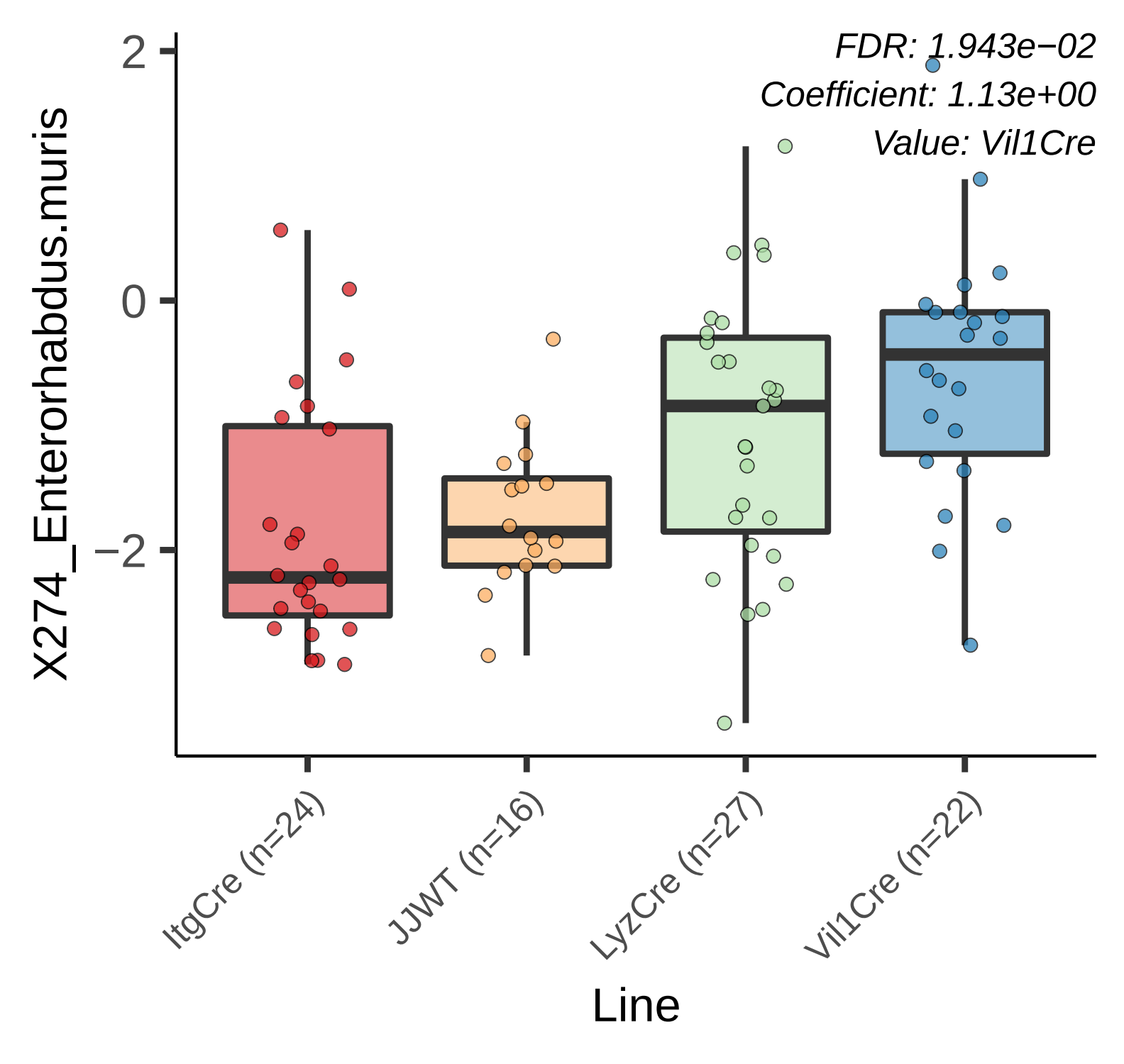

FDR: 1.186e−02
Coefficient: 1.22e+00
Value: LyzCre
X114_Lachnospiraceae.f.
2
0
−2
ItgCre (n=24)
JJWT (n=16)
LyzCre (n=27)
Vil1Cre (n=22)
Line

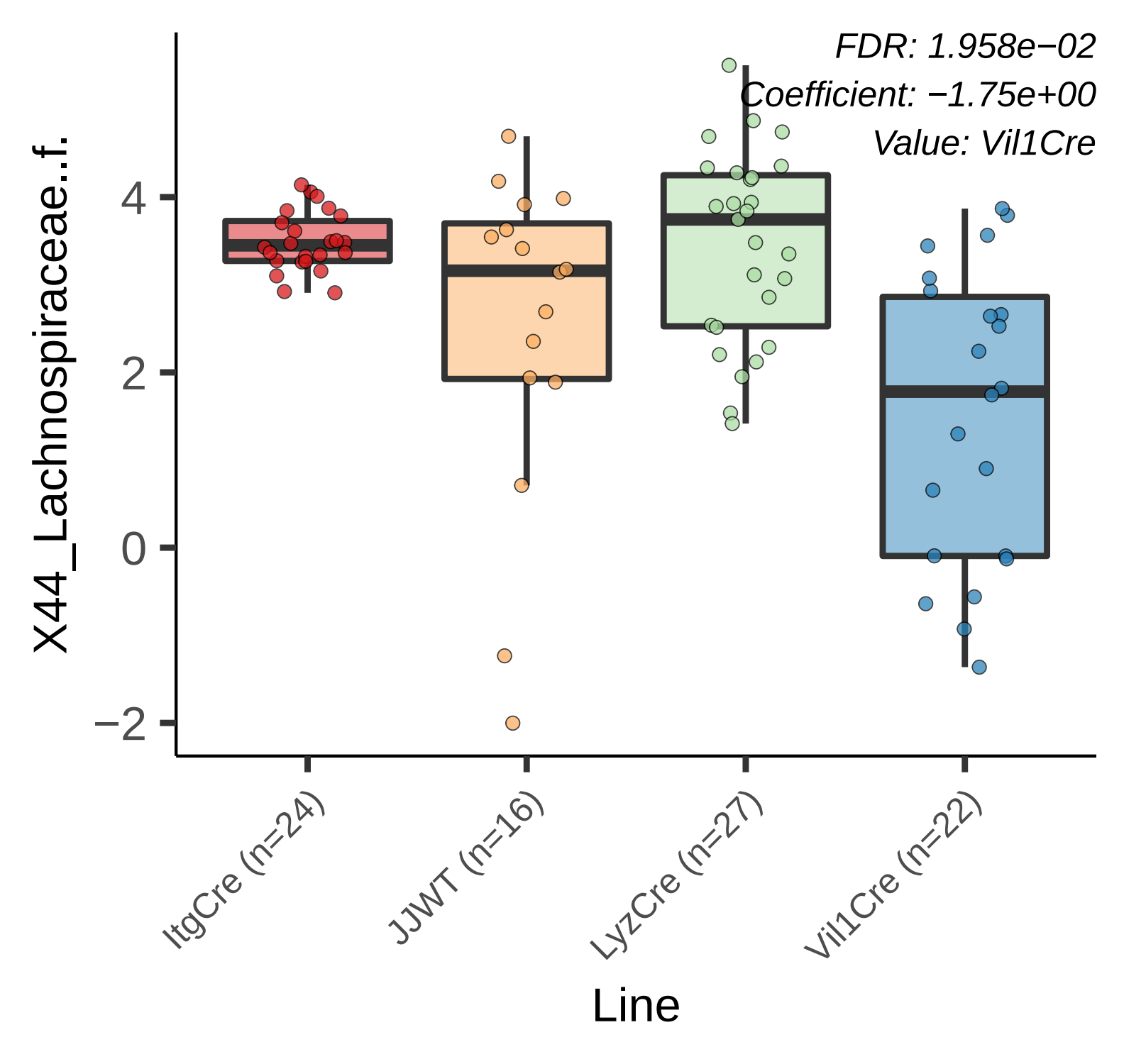

FDR: 1.543e−02
Coefficient: −1.12e+00
Value: Vil1Cre
X12_Muribaculaceae..f.
5
4
3
2
1
0
ItgCre (n=24)
JJWT (n=16)
LyzCre (n=27)
Vil1Cre (n=22)
Line

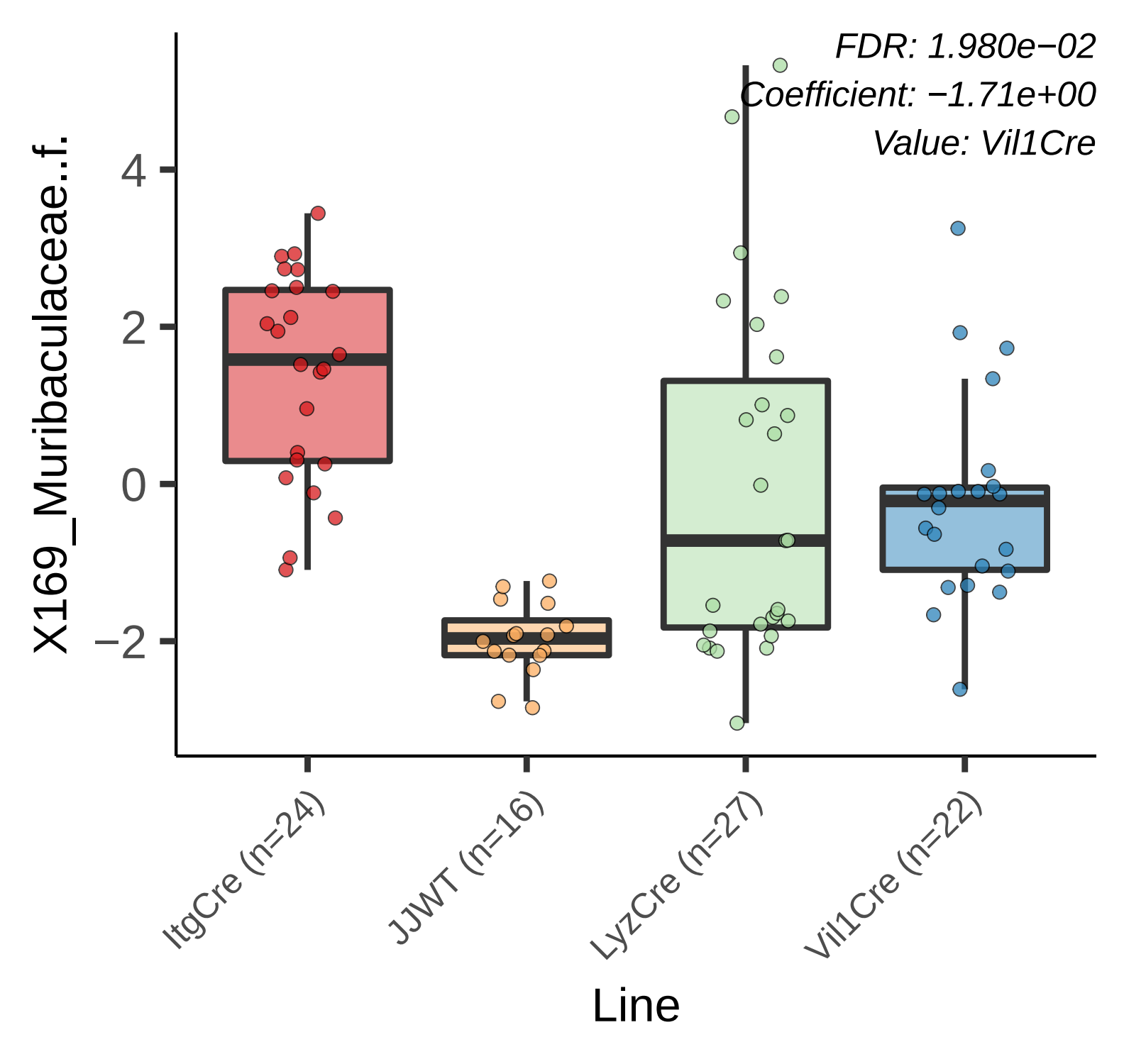

FDR: 1.543e−02
Coefficient: −1.27e+00
Value: LyzCre
X245_Lachnoclostridium.
2
0
−2
−4
ItgCre (n=24)
JJWT (n=16)
LyzCre (n=27)
Vil1Cre (n=22)
Line

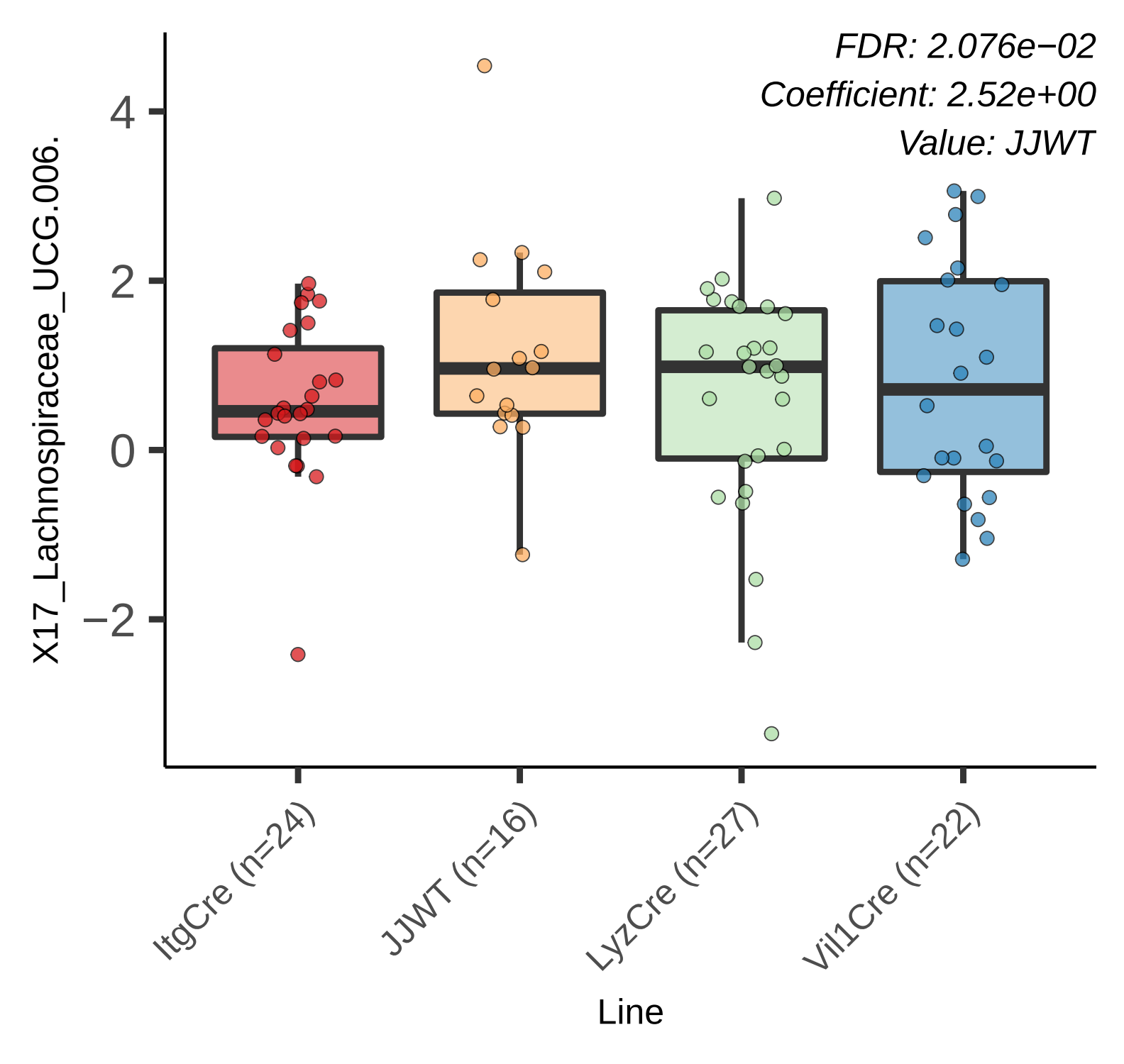

FDR: 1.605e−02
Coefficient: 1.09e+00
Value: LyzCre
X52_Enterorhabdus.
2
1
0
−1
−2
−3
ItgCre (n=24)
JJWT (n=16)
LyzCre (n=27)
Vil1Cre (n=22)
Line

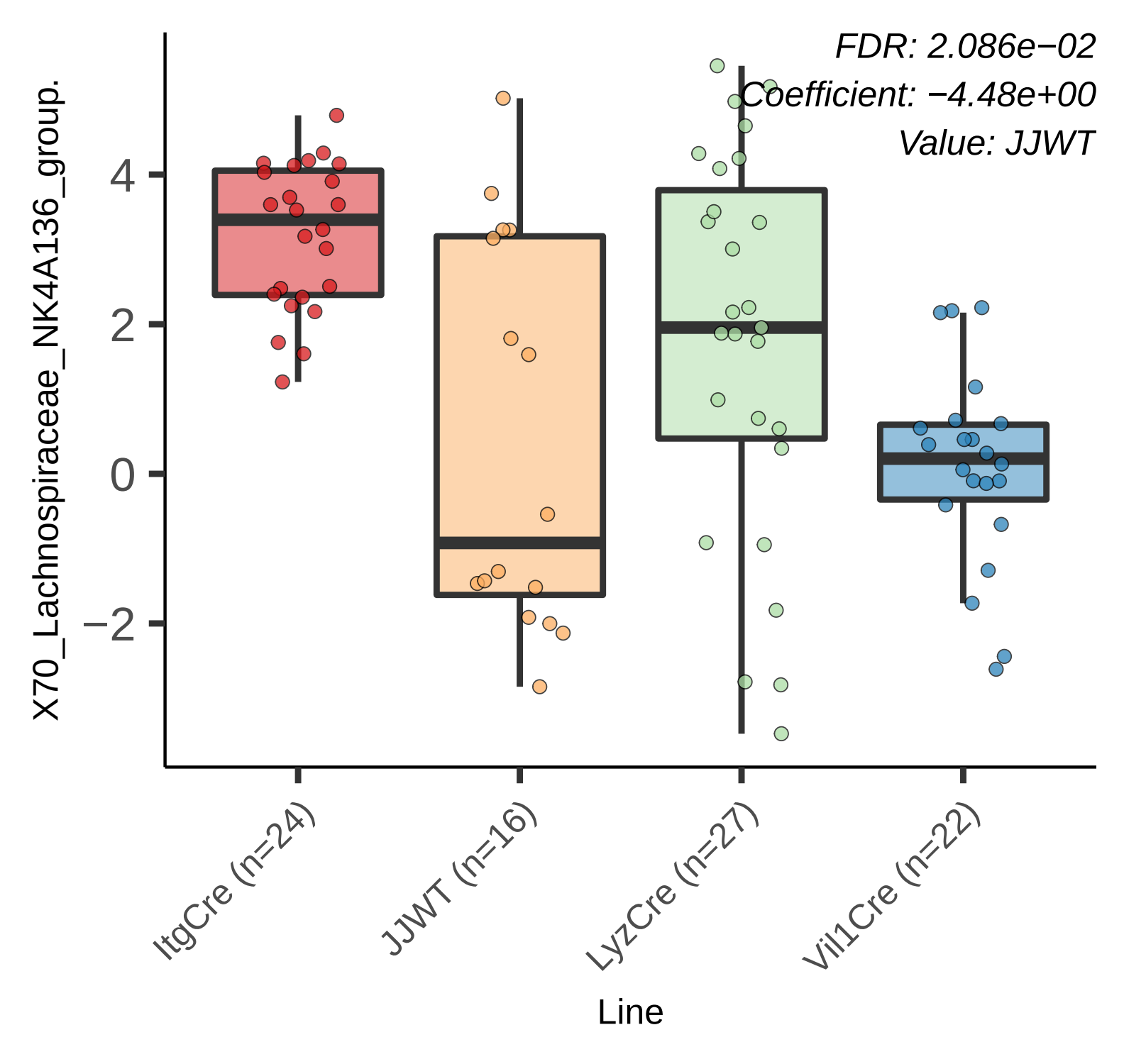

FDR: 1.899e−02
Coefficient: 1.05e+00
Value: LyzCre
X126_Oscillibacter.
2
1
0
−1
−2
−3
ItgCre (n=24)
JJWT (n=16)
LyzCre (n=27)
Vil1Cre (n=22)
Line

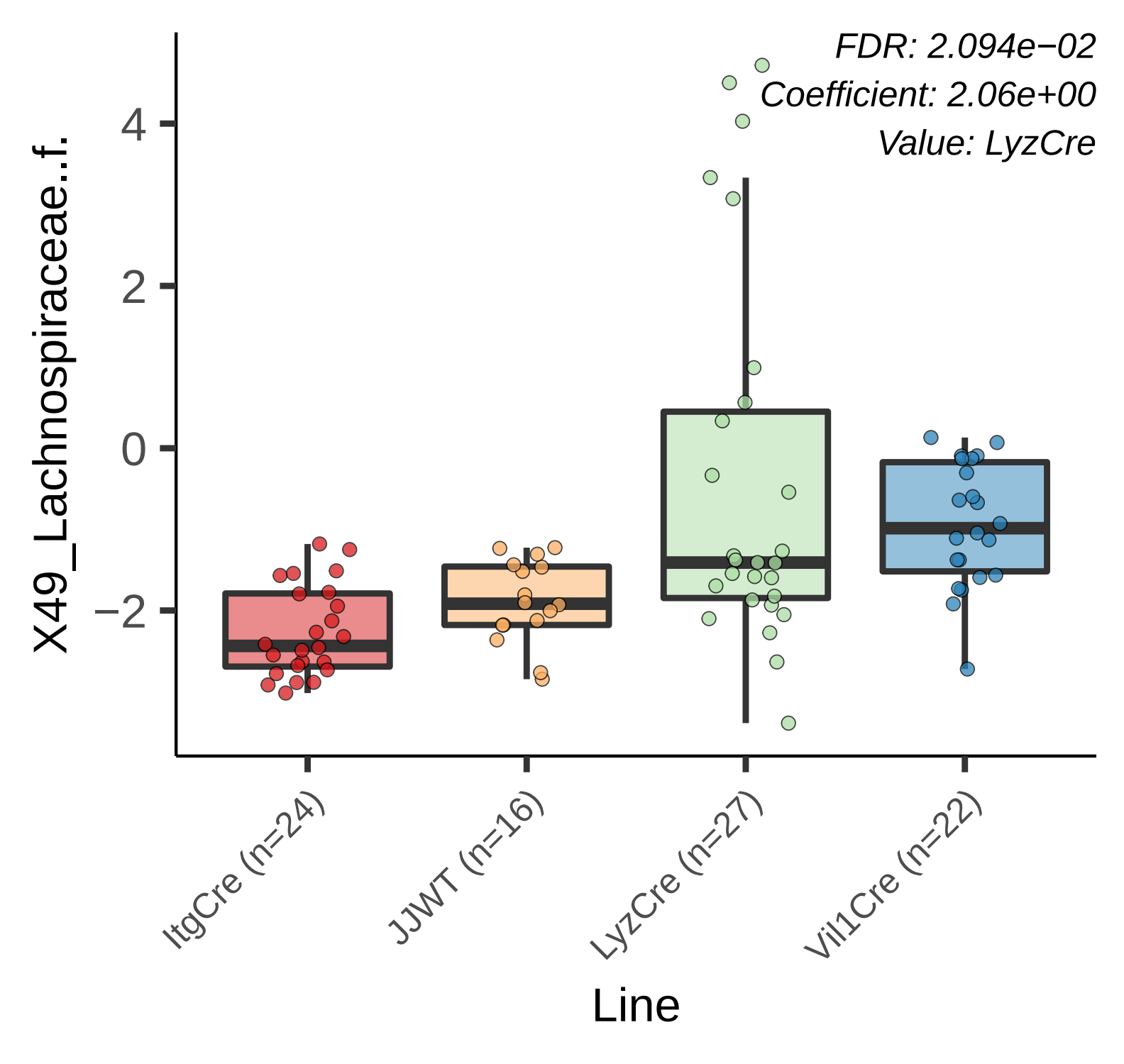

FDR: 1.943e−02
Coefficient: 1.13e+00
Value: Vil1Cre
X274_Enterorhabdus.muris
2
0
−2
ItgCre (n=24)
JJWT (n=16)
LyzCre (n=27)
Vil1Cre (n=22)
Line

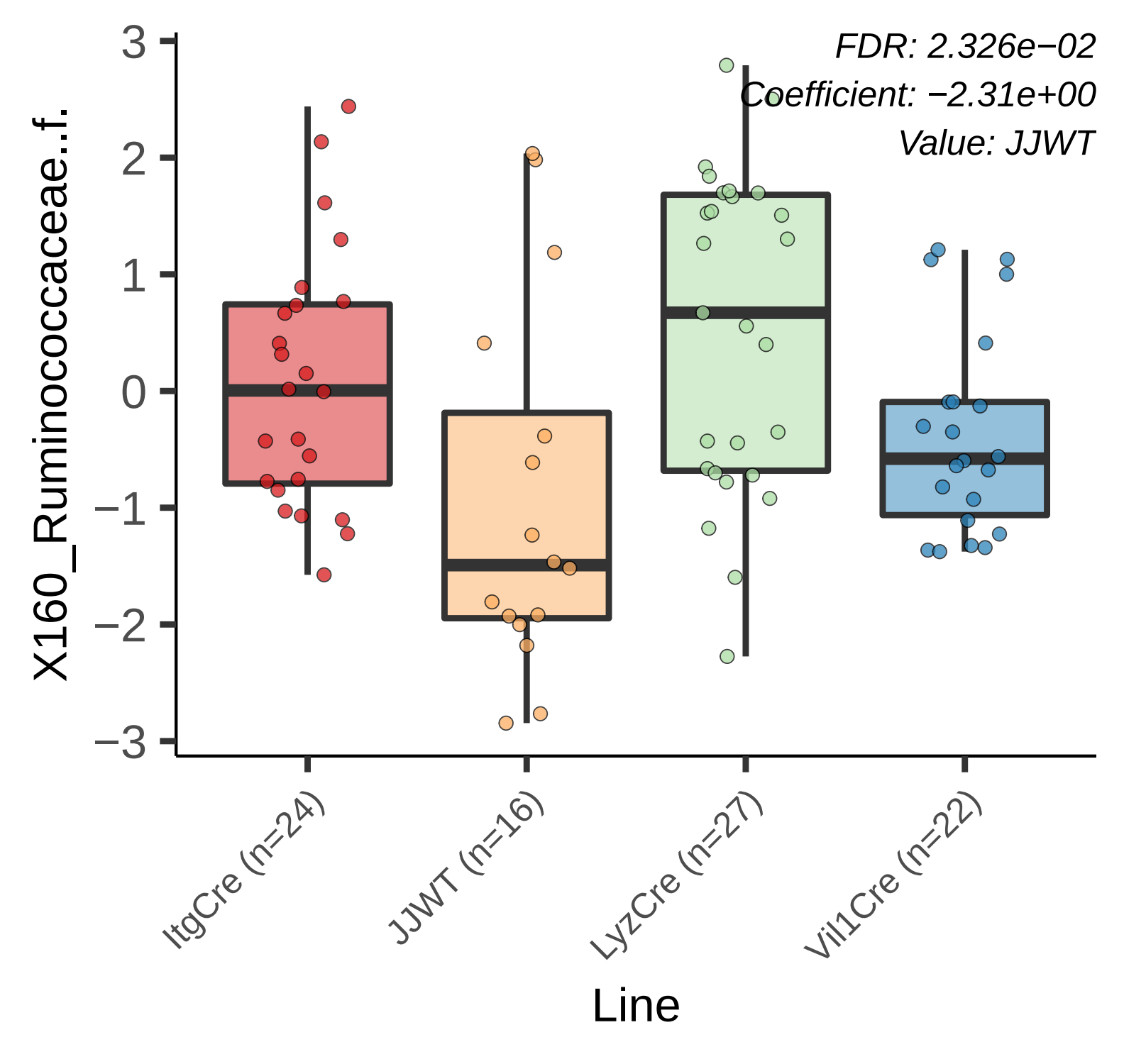

FDR: 1.958e−02
Coefficient: −1.75e+00
Value: Vil1Cre
X44_Lachnospiraceae..f.
4
2
0
−2
ItgCre (n=24)
JJWT (n=16)
LyzCre (n=27)
Vil1Cre (n=22)
Line

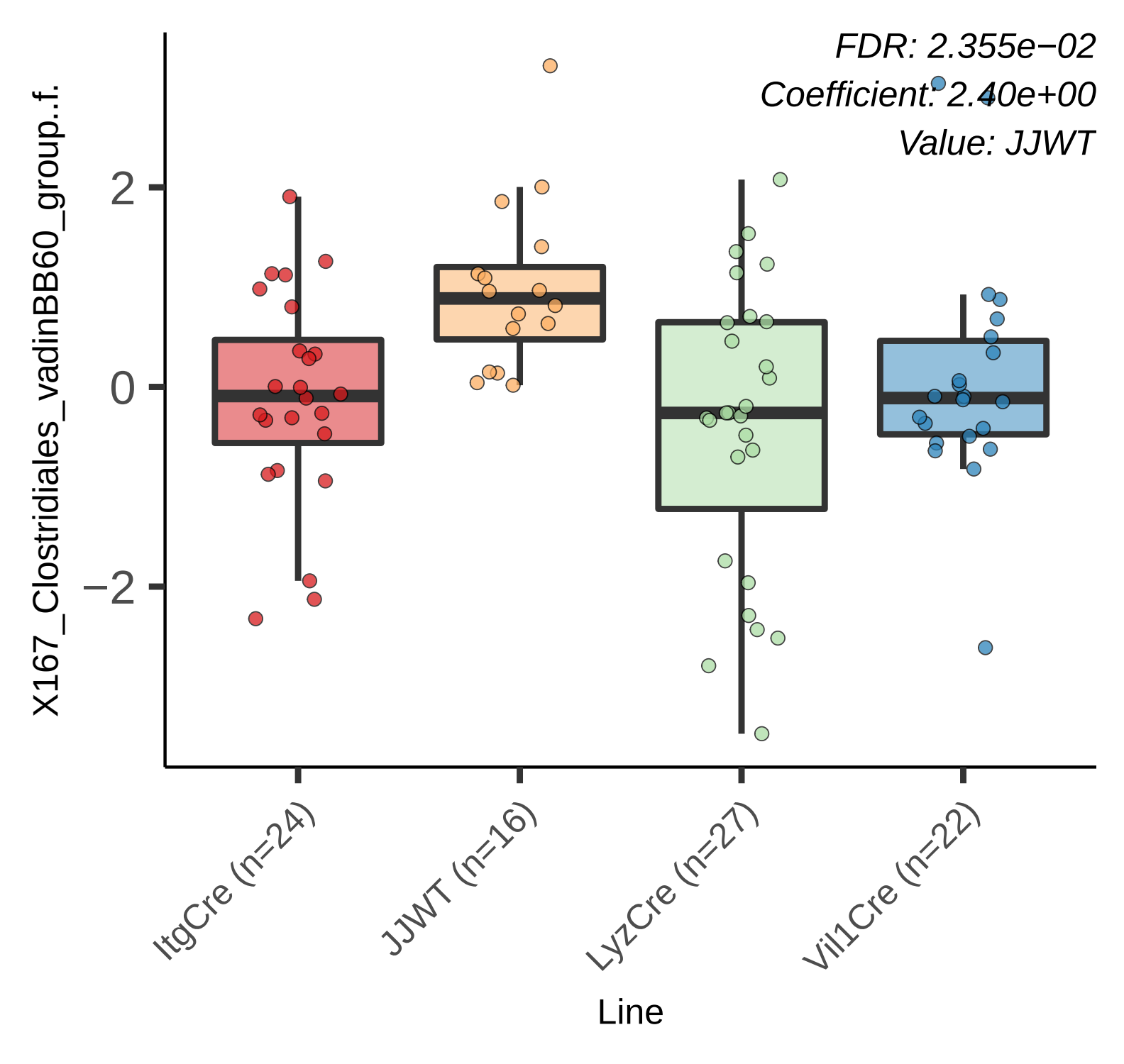

FDR: 1.980e−02
Coefficient: −1.71e+00
Value: Vil1Cre
X169_Muribaculaceae..f.
4
2
0
−2
ItgCre (n=24)
JJWT (n=16)
LyzCre (n=27)
Vil1Cre (n=22)
Line

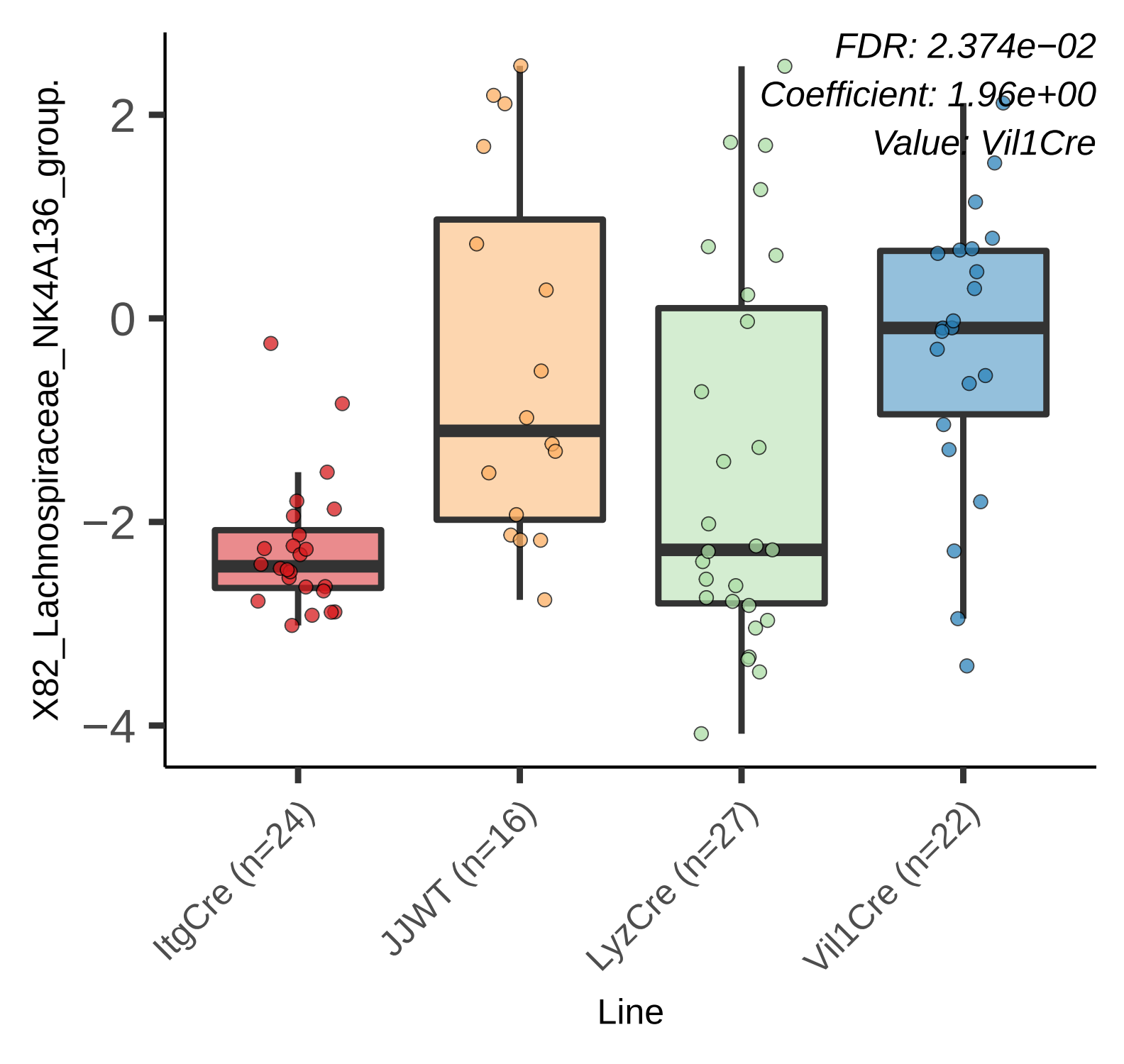

FDR: 2.076e−02
Coefficient: 2.52e+00
Value: JJWT
X17_Lachnospiraceae_UCG.006.
4
2
0
−2
ItgCre (n=24)
JJWT (n=16)
LyzCre (n=27)
Vil1Cre (n=22)
Line

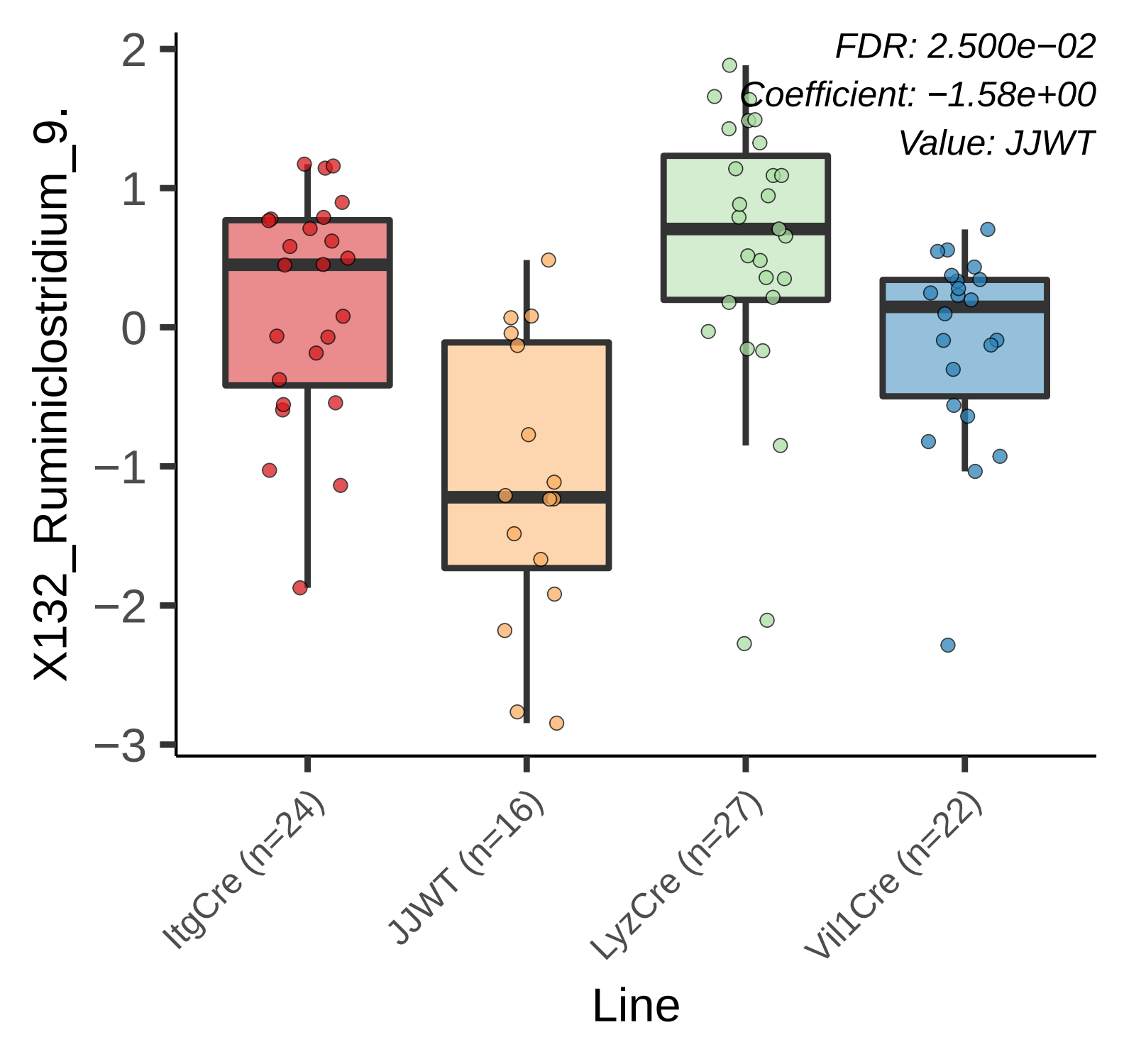

FDR: 2.086e−02
Coefficient: −4.48e+00
Value: JJWT
X70_Lachnospiraceae_NK4A136_group.
4
2
0
−2
ItgCre (n=24)
JJWT (n=16)
LyzCre (n=27)
Vil1Cre (n=22)
Line

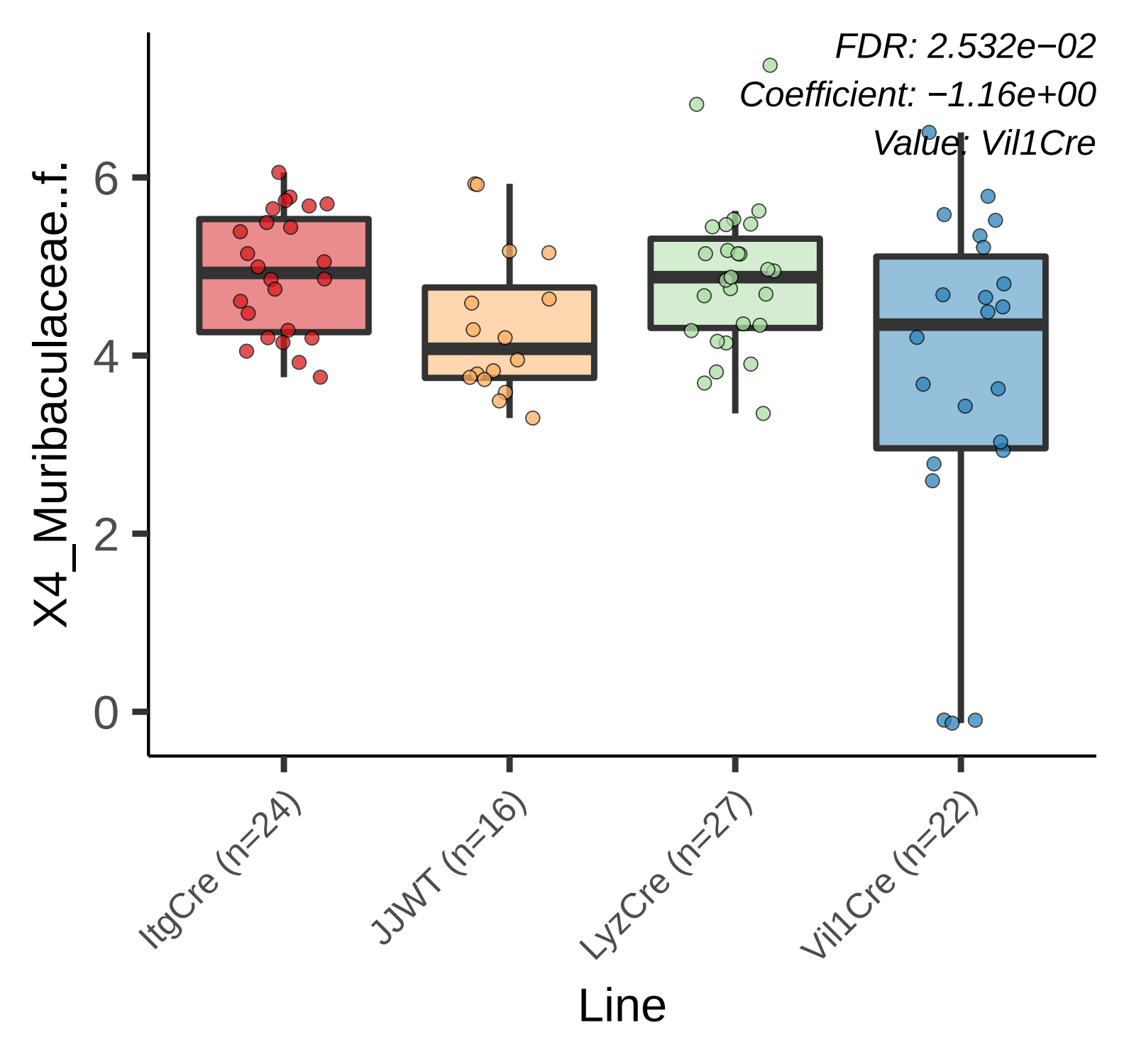

FDR: 2.094e−02
Coefficient: 2.06e+00
Value: LyzCre
X49_Lachnospiraceae..f.
4
2
0
−2
ItgCre (n=24)
JJWT (n=16)
LyzCre (n=27)
Vil1Cre (n=22)
Line

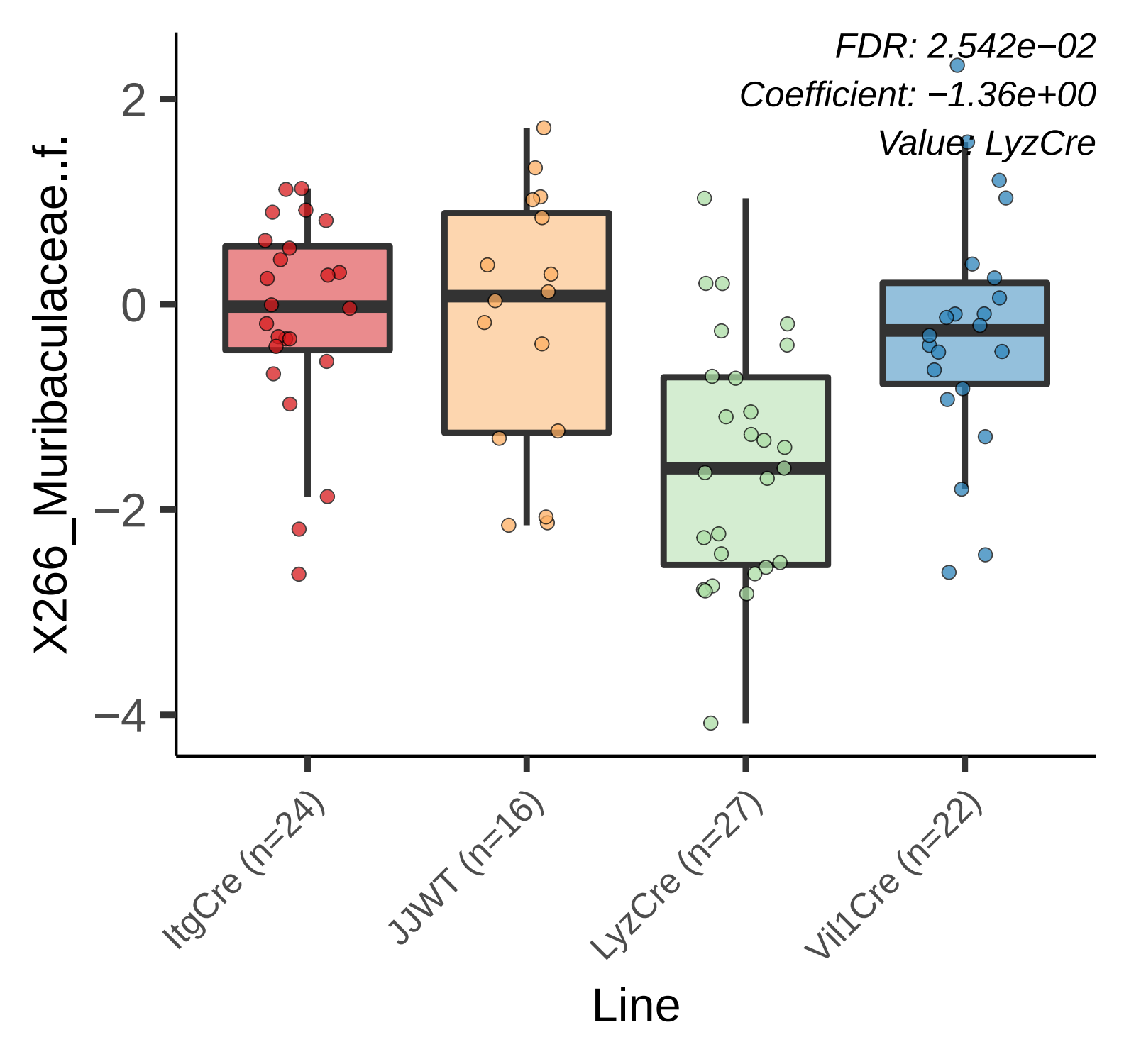

FDR: 2.326e−02
Coefficient: −2.31e+00
Value: JJWT
X160_Ruminococcaceae..f.
3
2
1
0
−1
−2
−3
ItgCre (n=24)
JJWT (n=16)
LyzCre (n=27)
Vil1Cre (n=22)
Line

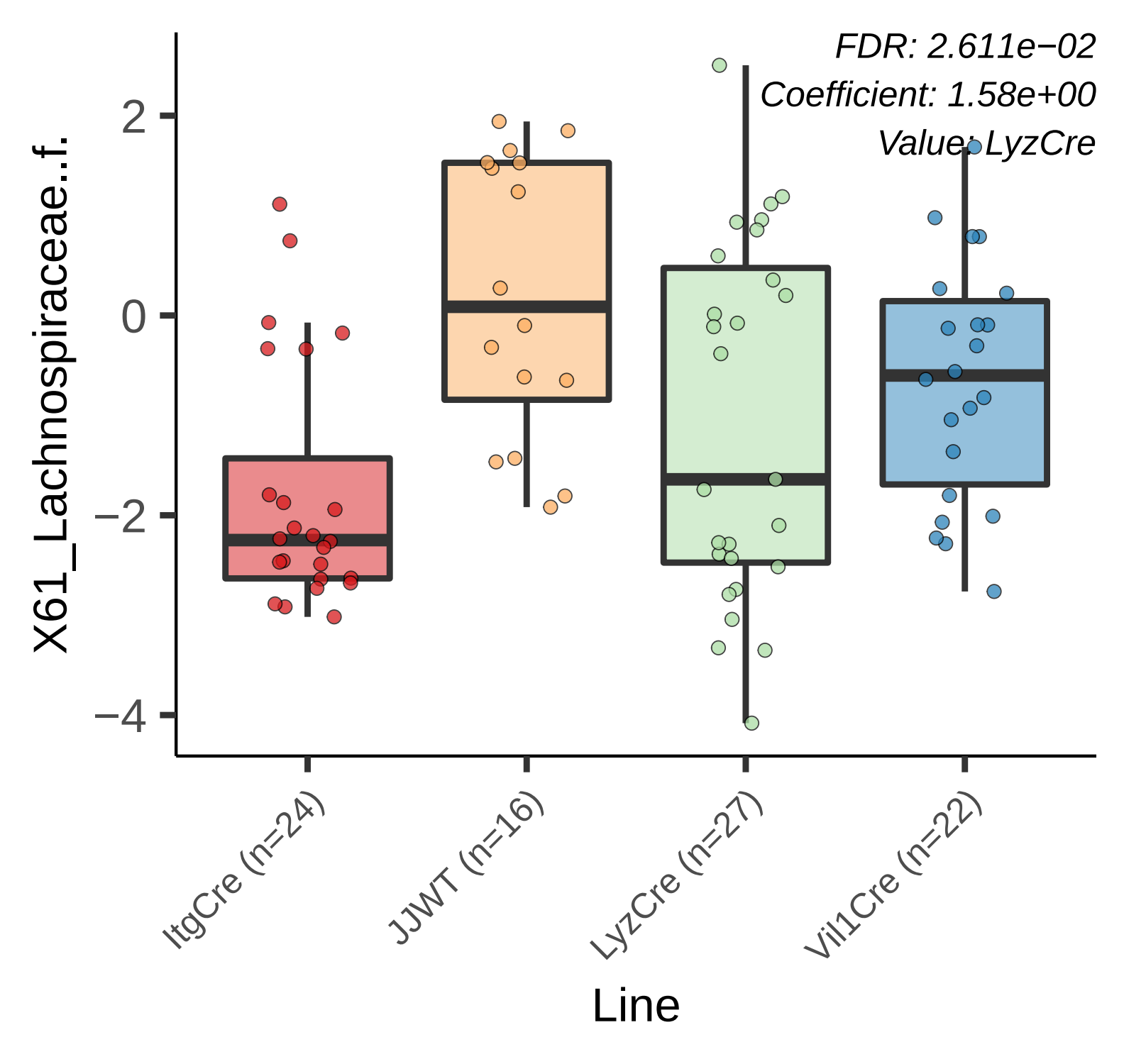

FDR: 2.355e−02
Coefficient: 2.40e+00
Value: JJWT
X167_Clostridiales_vadinBB60_group..f.
2
0
−2
ItgCre (n=24)
JJWT (n=16)
LyzCre (n=27)
Vil1Cre (n=22)
Line

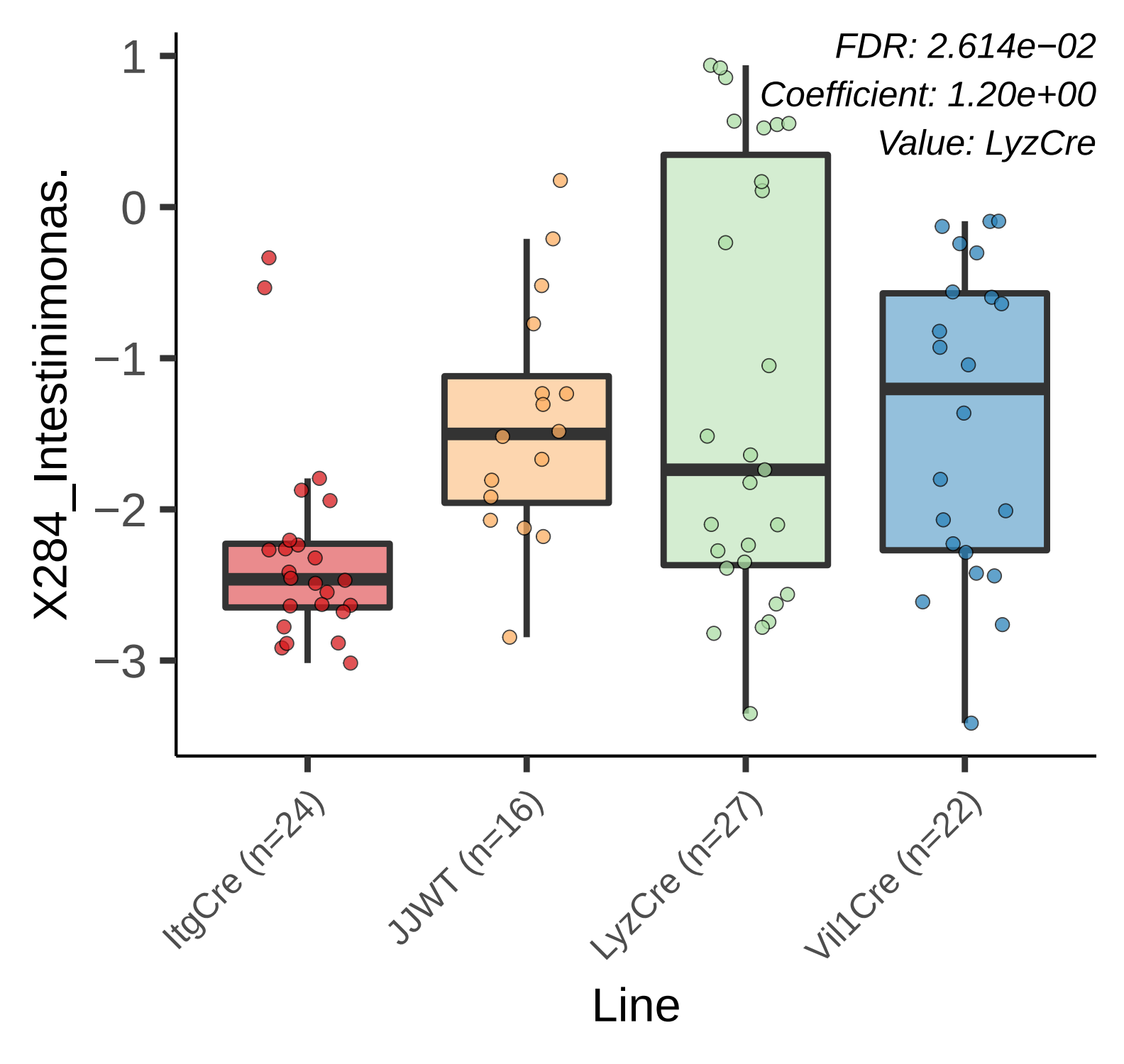

FDR: 2.374e−02
Coefficient: 1.96e+00
Value: Vil1Cre
X82_Lachnospiraceae_NK4A136_group.
2
0
−2
−4
ItgCre (n=24)
JJWT (n=16)
LyzCre (n=27)
Vil1Cre (n=22)
Line

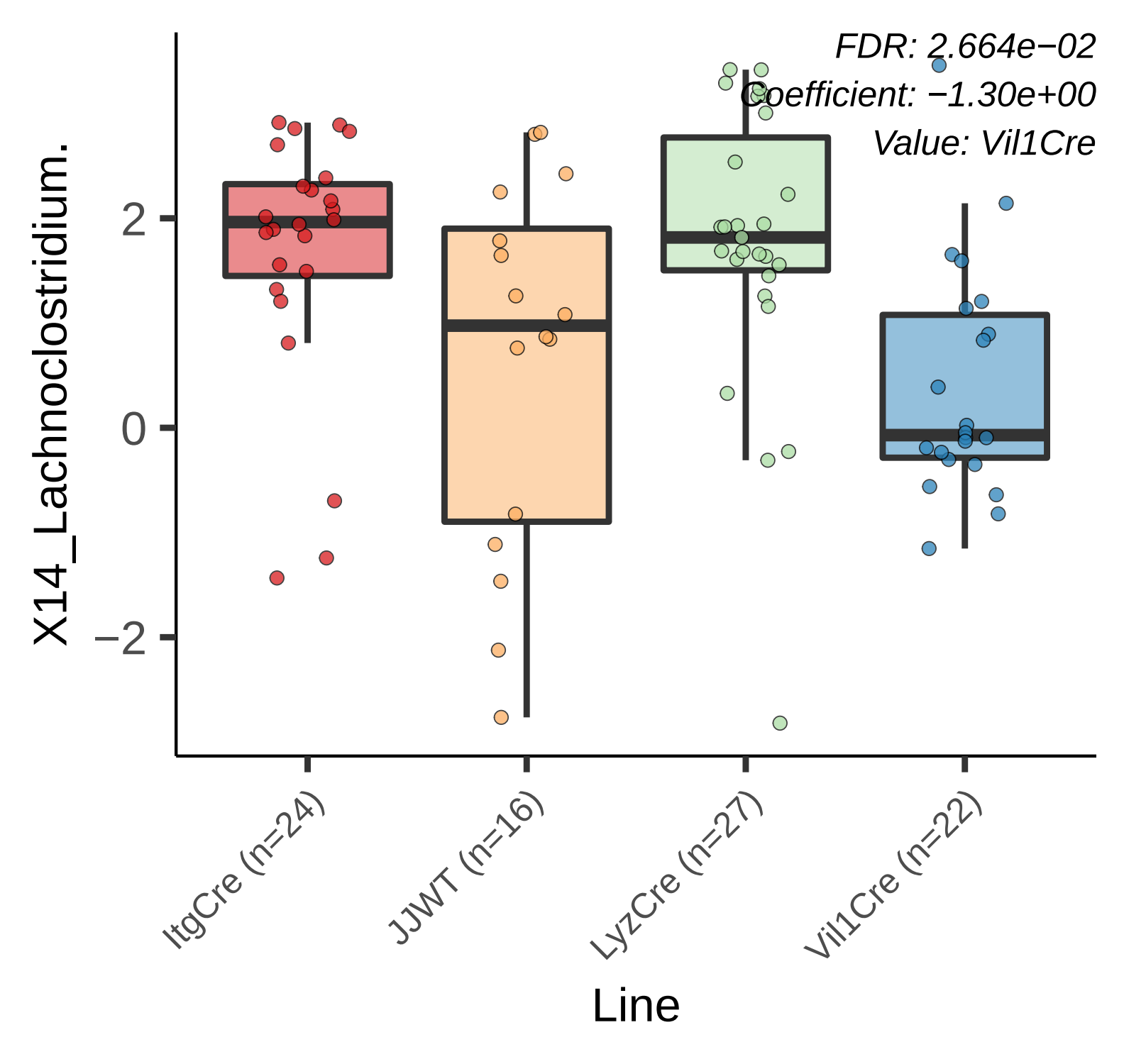

FDR: 2.500e−02
Coefficient: −1.58e+00
Value: JJWT
X132_Ruminiclostridium_9.
2
1
0
−1
−2
−3
ItgCre (n=24)
JJWT (n=16)
LyzCre (n=27)
Vil1Cre (n=22)
Line

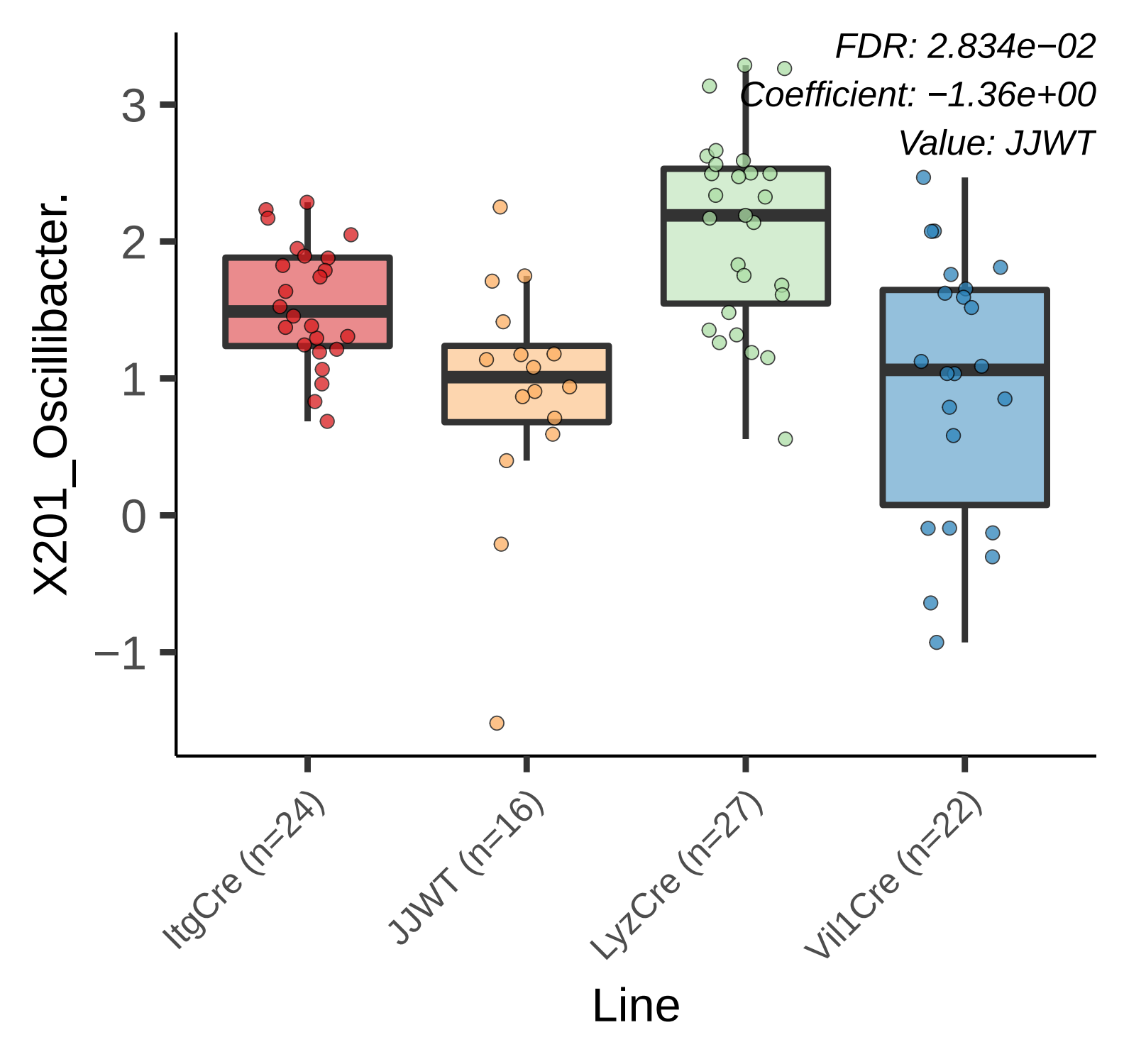

FDR: 2.532e−02
Coefficient: −1.16e+00
Value: Vil1Cre
X4_Muribaculaceae..f.
6
4
2
0
ItgCre (n=24)
JJWT (n=16)
LyzCre (n=27)
Vil1Cre (n=22)
Line

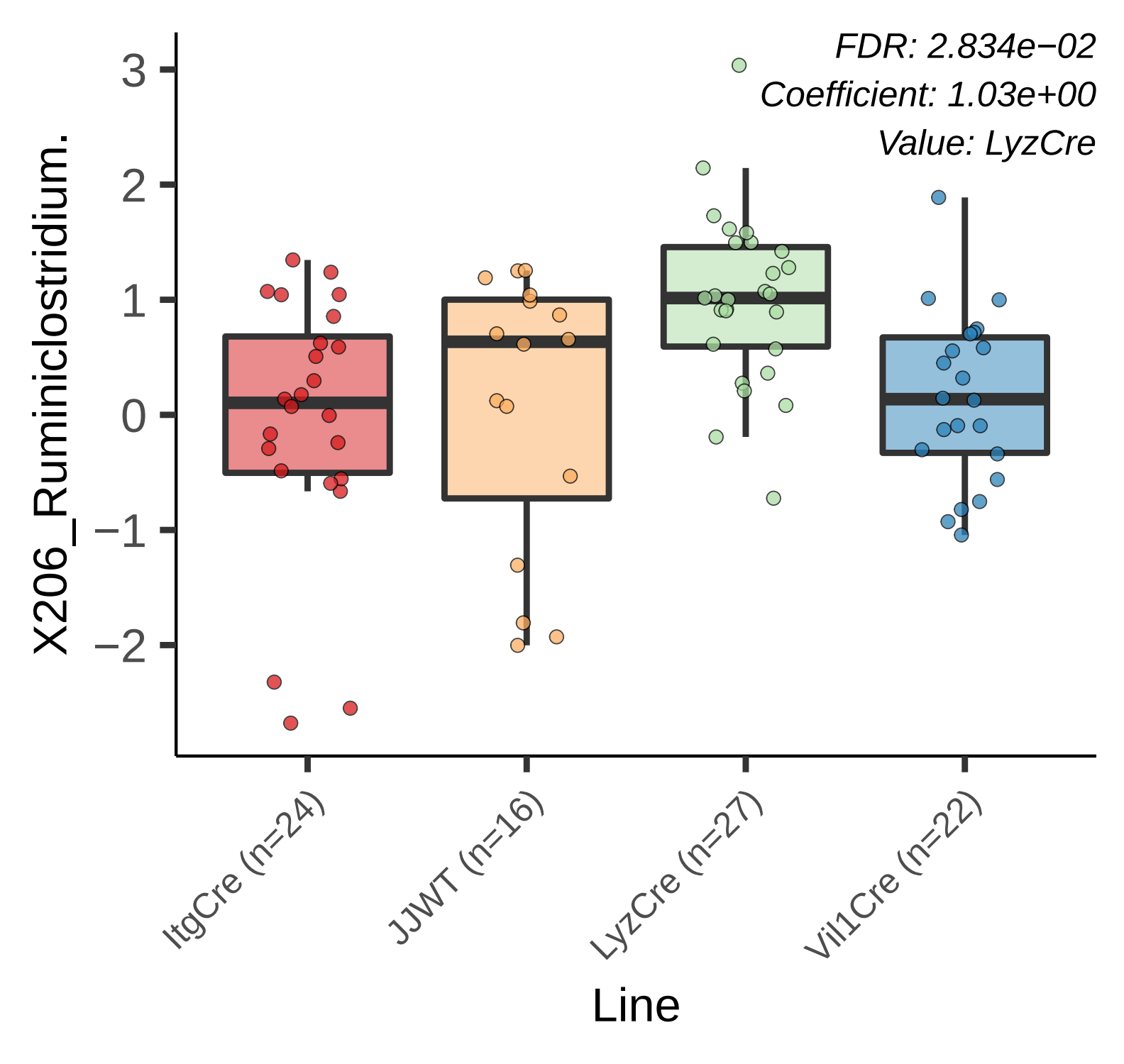

FDR: 2.542e−02
Coefficient: −1.36e+00
Value: LyzCre
X266_Muribaculaceae..f.
2
0
−2
−4
ItgCre (n=24)
JJWT (n=16)
LyzCre (n=27)
Vil1Cre (n=22)
Line

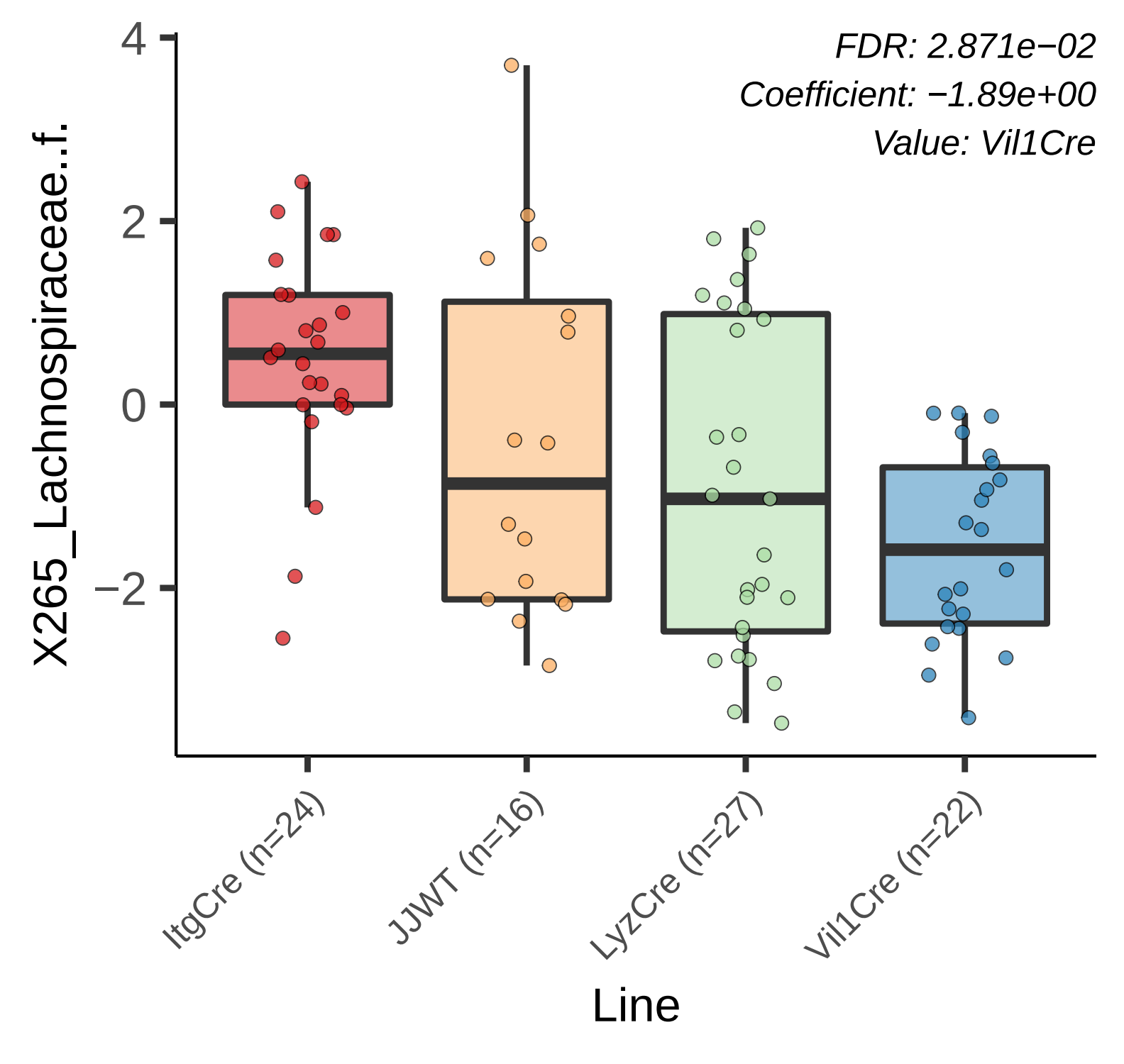

FDR: 2.611e−02
Coefficient: 1.58e+00
Value: LyzCre
X61_Lachnospiraceae..f.
2
0
−2
−4
ItgCre (n=24)
JJWT (n=16)
LyzCre (n=27)
Vil1Cre (n=22)
Line

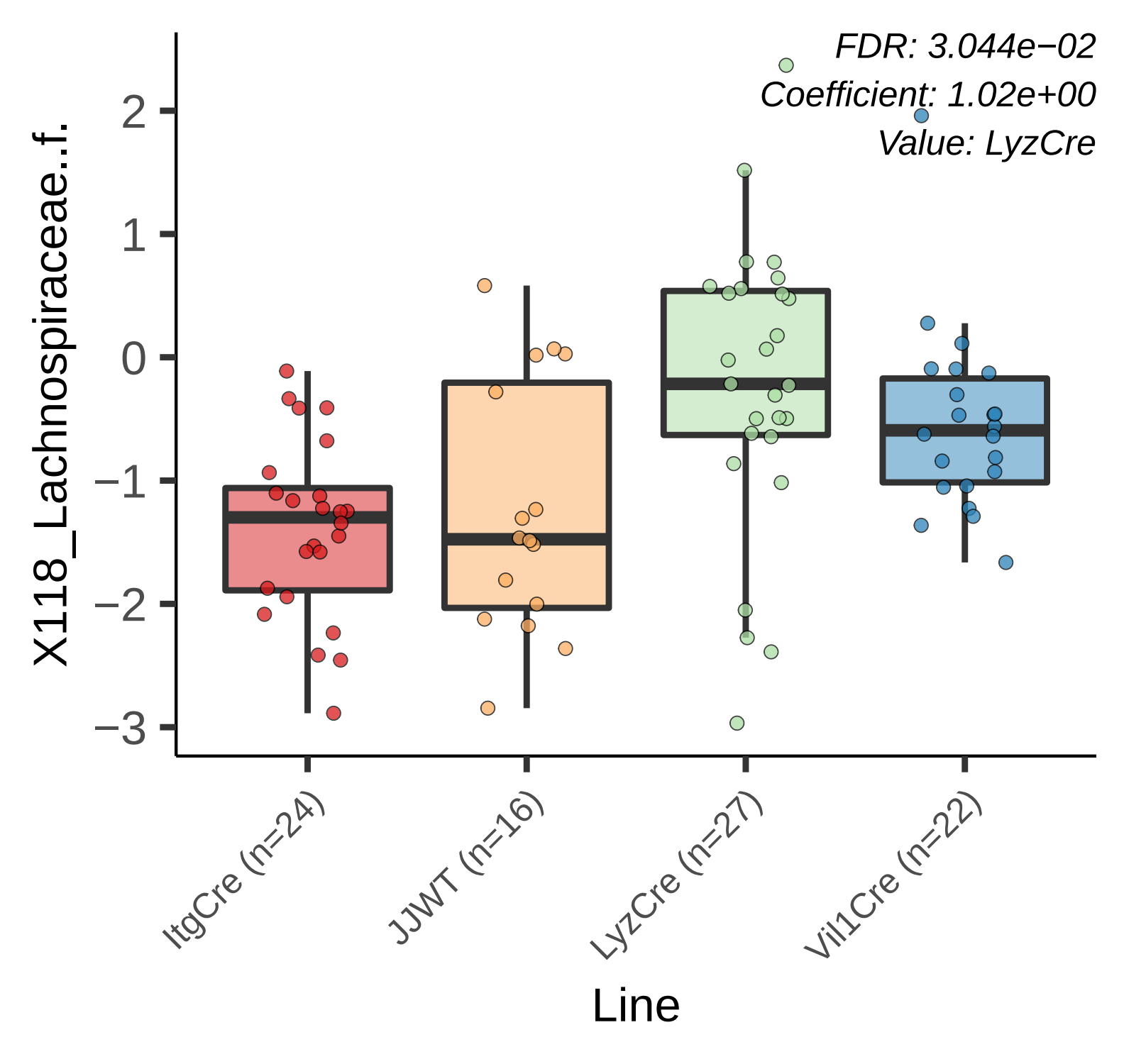

FDR: 2.614e−02
Coefficient: 1.20e+00
Value: LyzCre
X284_Intestinimonas.
1
0
−1
−2
−3
ItgCre (n=24)
JJWT (n=16)
LyzCre (n=27)
Vil1Cre (n=22)
Line

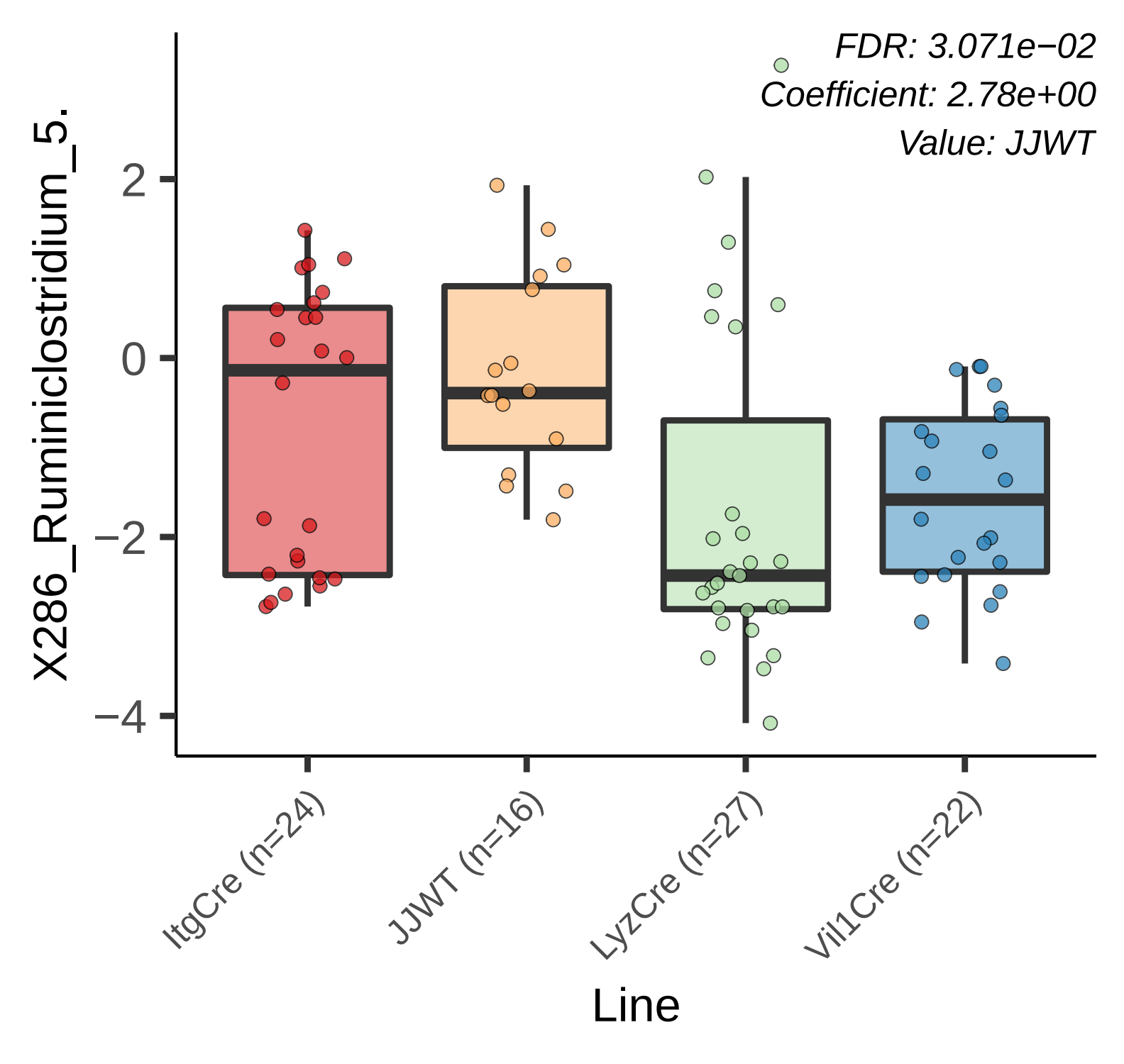

FDR: 2.664e−02
Coefficient: −1.30e+00
Value: Vil1Cre
X14_Lachnoclostridium.
2
0
−2
ItgCre (n=24)
JJWT (n=16)
LyzCre (n=27)
Vil1Cre (n=22)
Line

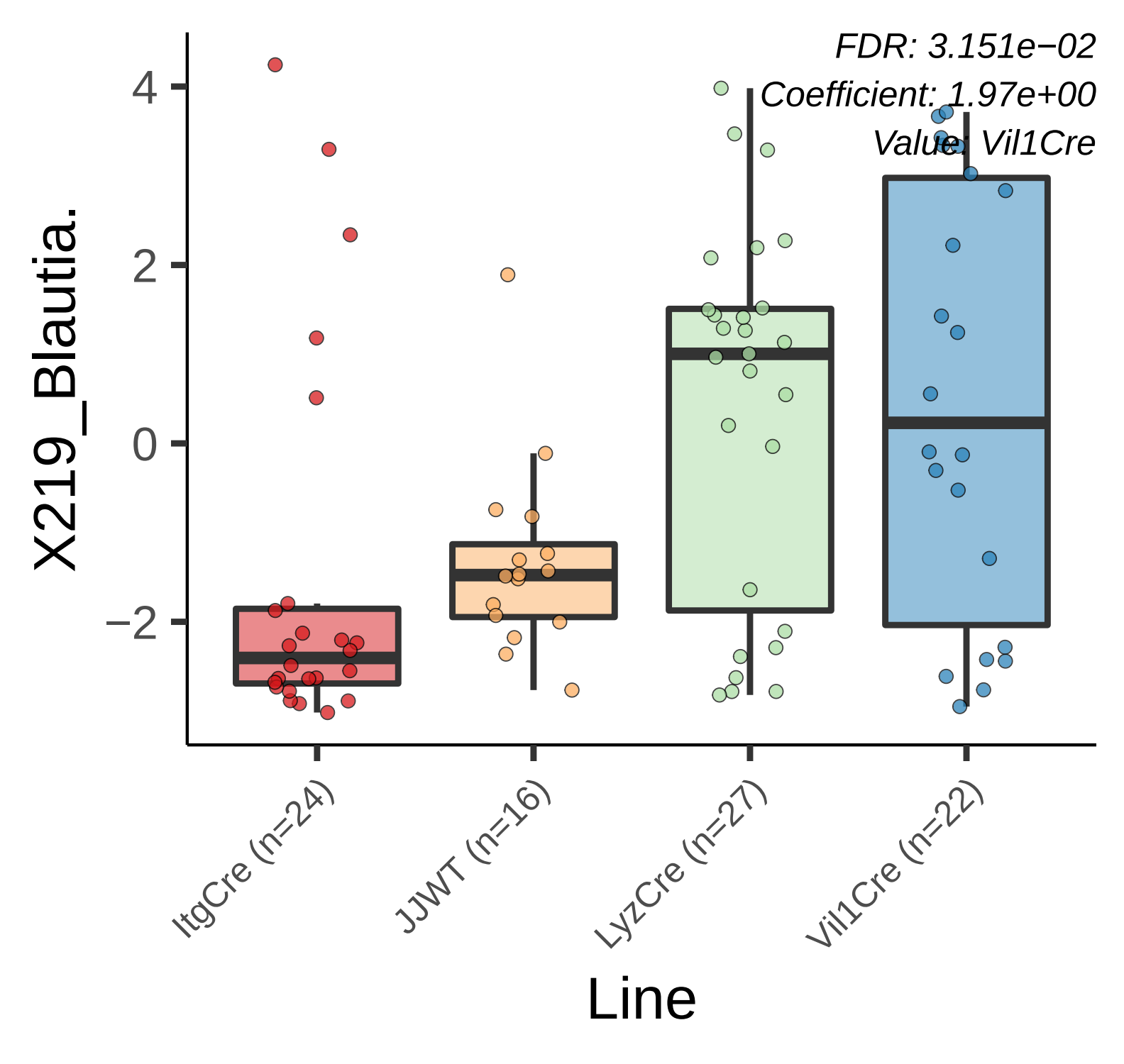

FDR: 2.834e−02
Coefficient: −1.36e+00
Value: JJWT
X201_Oscillibacter.
3
2
1
0
−1
ItgCre (n=24)
JJWT (n=16)
LyzCre (n=27)
Vil1Cre (n=22)
Line

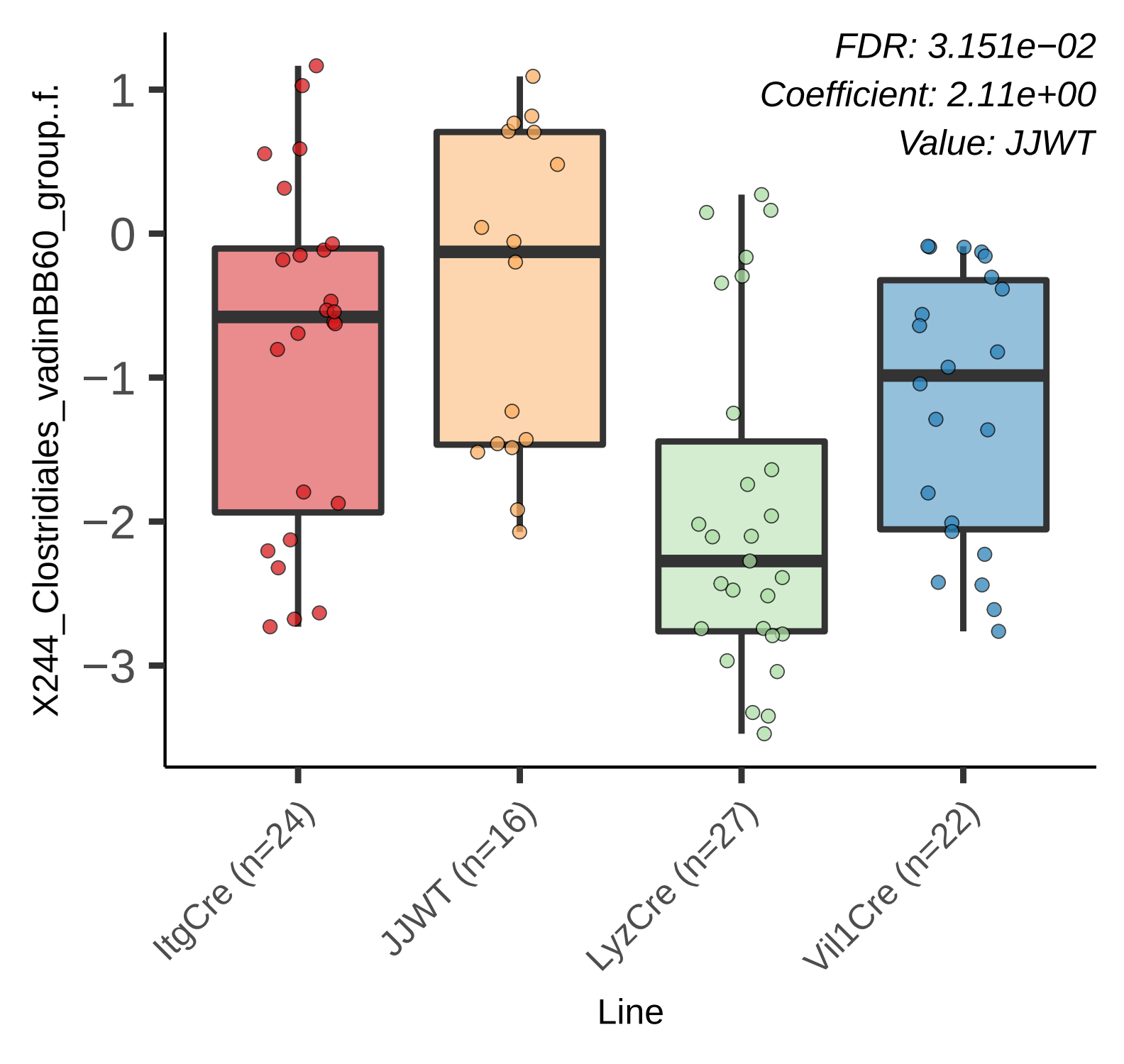

FDR: 2.834e−02
Coefficient: 1.03e+00
Value: LyzCre
X206_Ruminiclostridium.
3
2
1
0
−1
−2
ItgCre (n=24)
JJWT (n=16)
LyzCre (n=27)
Vil1Cre (n=22)
Line

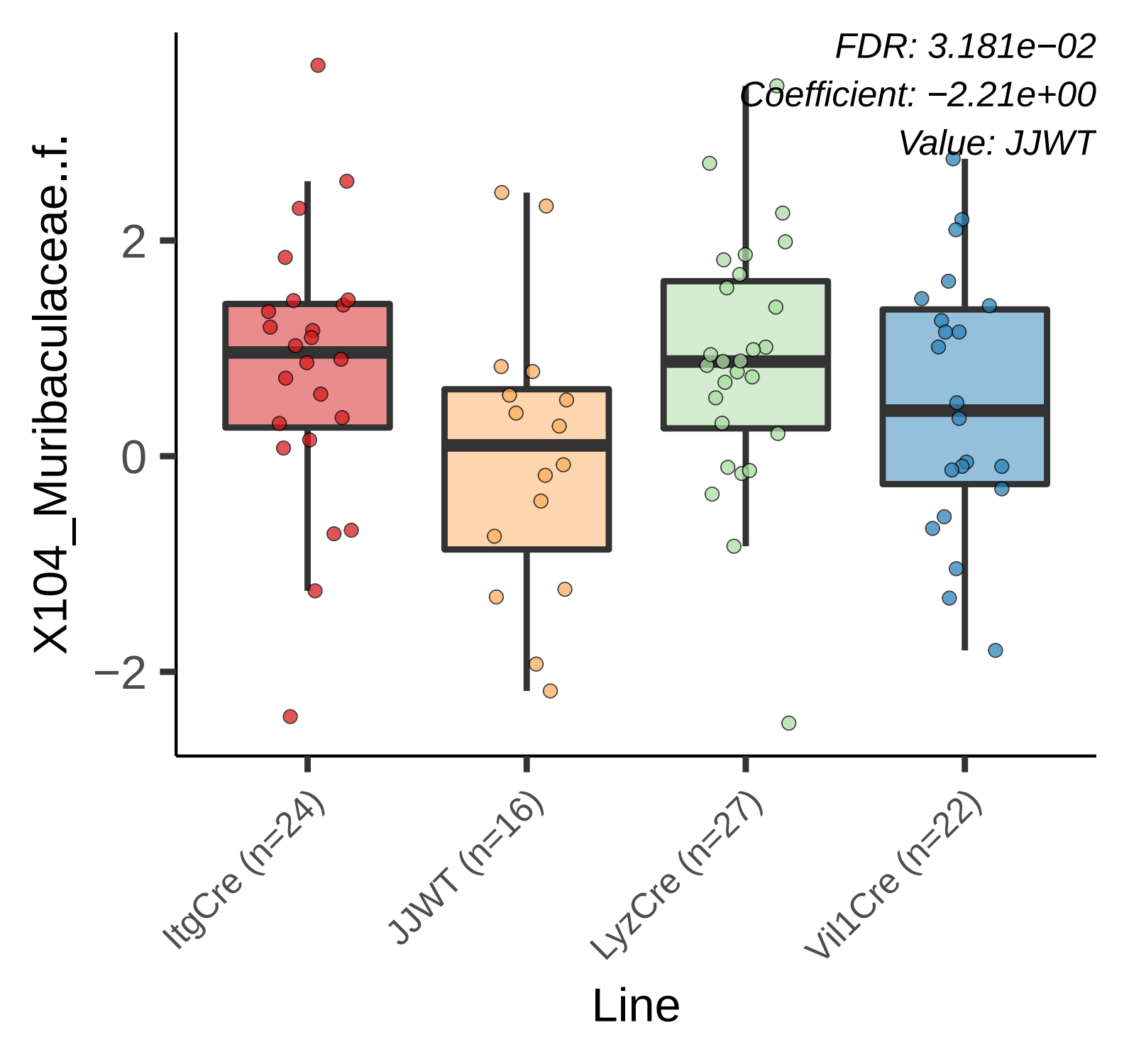

FDR: 2.871e−02
Coefficient: −1.89e+00
Value: Vil1Cre
X265_Lachnospiraceae.f.
4
2
0
−2
ItgCre (n=24)
JJWT (n=16)
LyzCre (n=27)
Vil1Cre (n=22)
Line

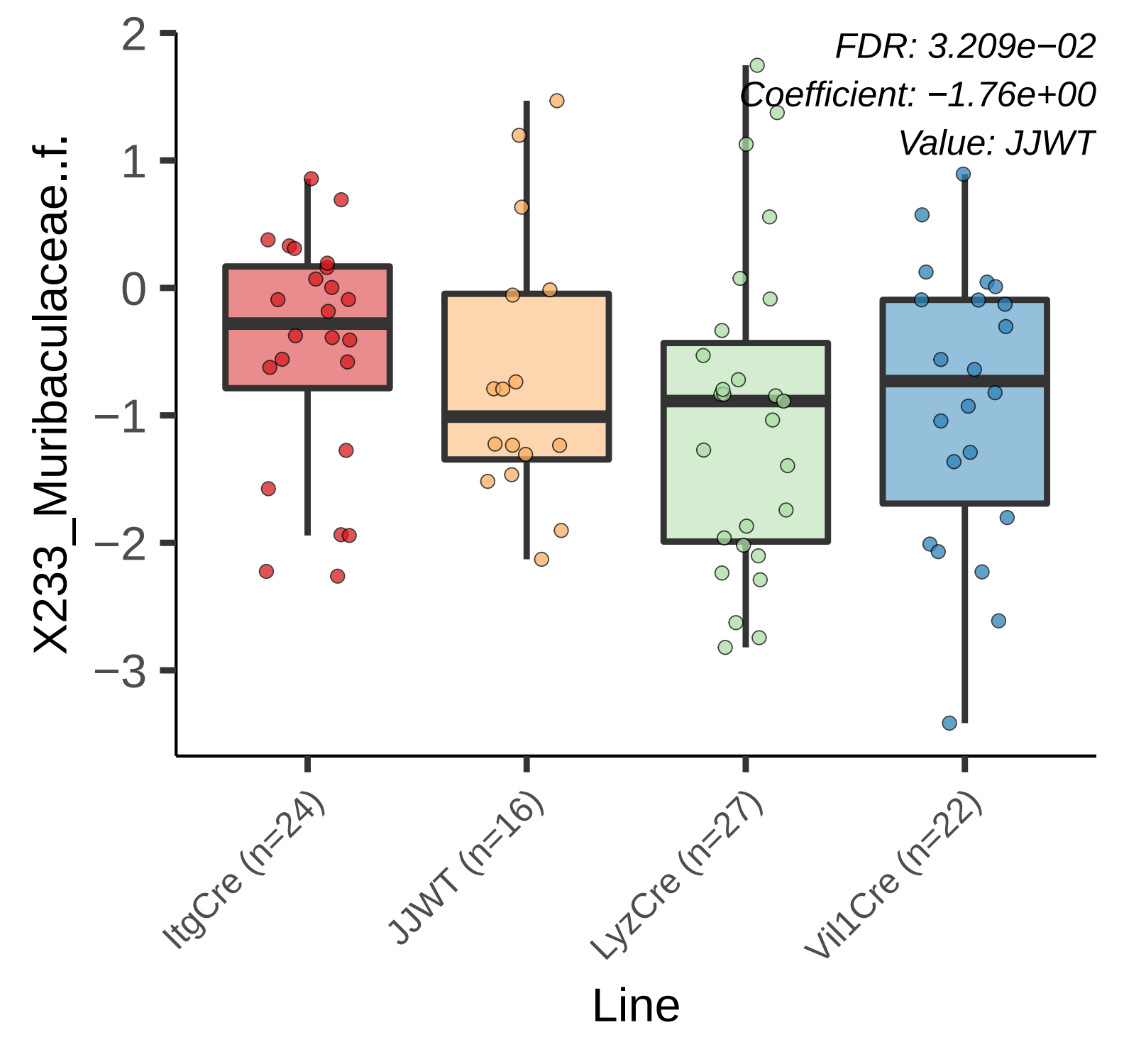

FDR: 3.044e−02
Coefficient: 1.02e+00
Value: LyzCre
X118_Lachnospiraceae.f.
2
1
0
−1
−2
−3
ItgCre (n=24)
JJWT (n=16)
LyzCre (n=27)
Vil1Cre (n=22)
Line

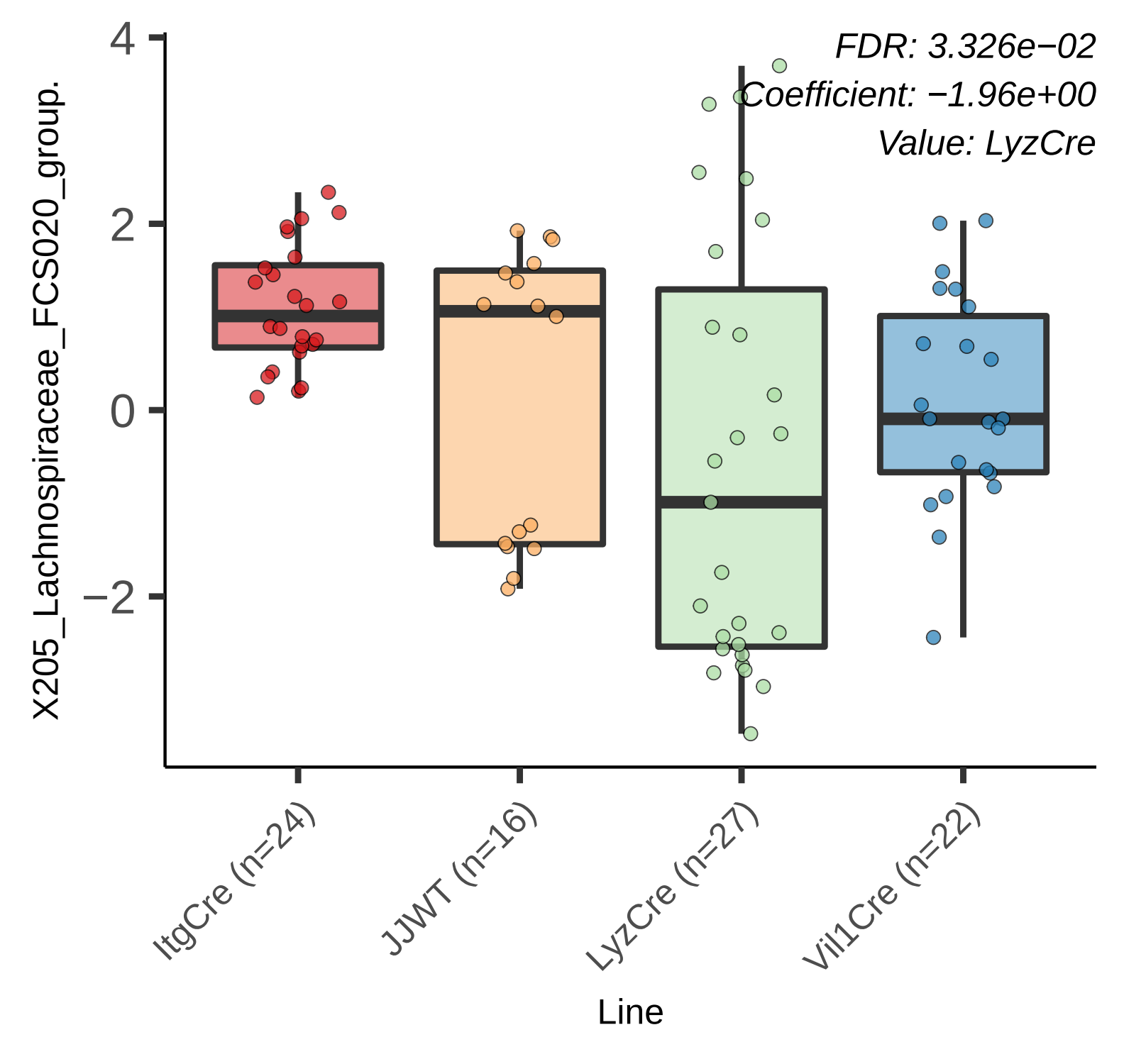

FDR: 3.071e−02
Coefficient: 2.78e+00
Value: JJWT
X286_Ruminiclostridium_5.
2
0
−2
−4
ItgCre (n=24)
JJWT (n=16)
LyzCre (n=27)
Vil1Cre (n=22)
Line

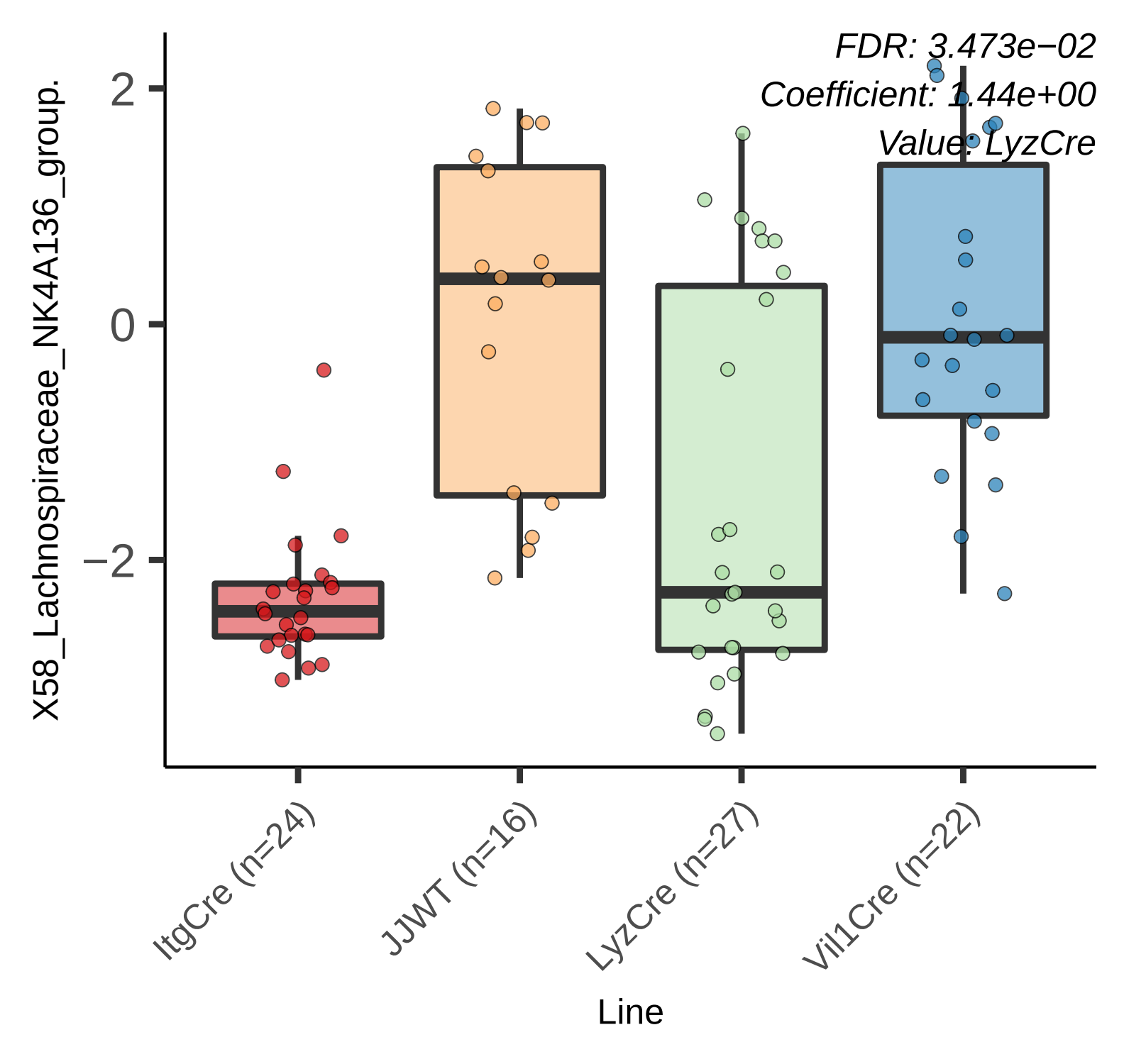

FDR: 3.151e−02
Coefficient: 1.97e+00
Value: Vil1Cre
X219_Blautia.
4
2
0
−2
ItgCre (n=24)
JJWT (n=16)
LyzCre (n=27)
Vil1Cre (n=22)
Line

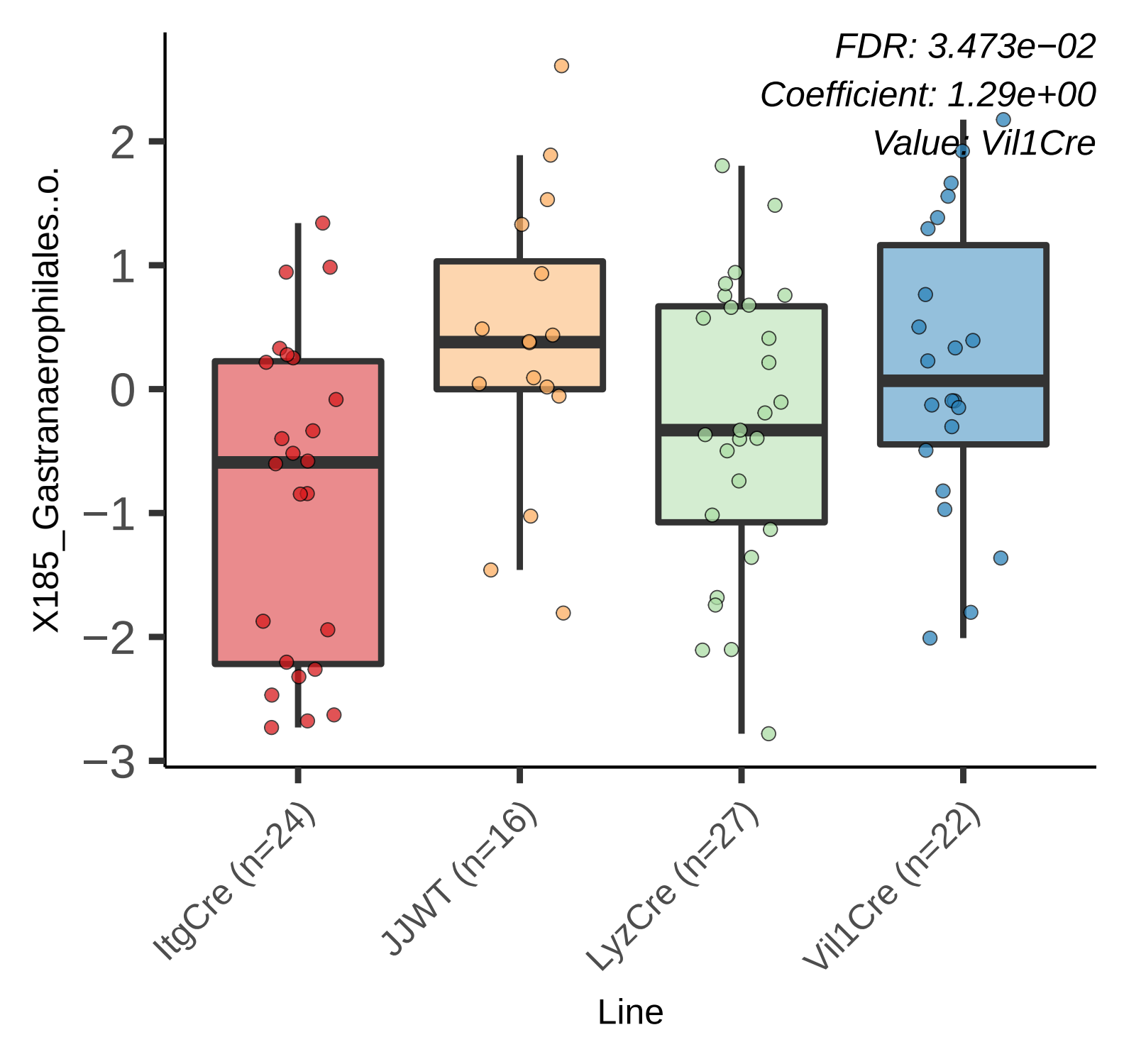

FDR: 3.151e−02
Coefficient: 2.11e+00
Value: JJWT
X244_Clostridiales_vadinBB60_group..f.
1
0
−1
−2
−3
ItgCre (n=24)
JJWT (n=16)
LyzCre (n=27)
Vil1Cre (n=22)
Line

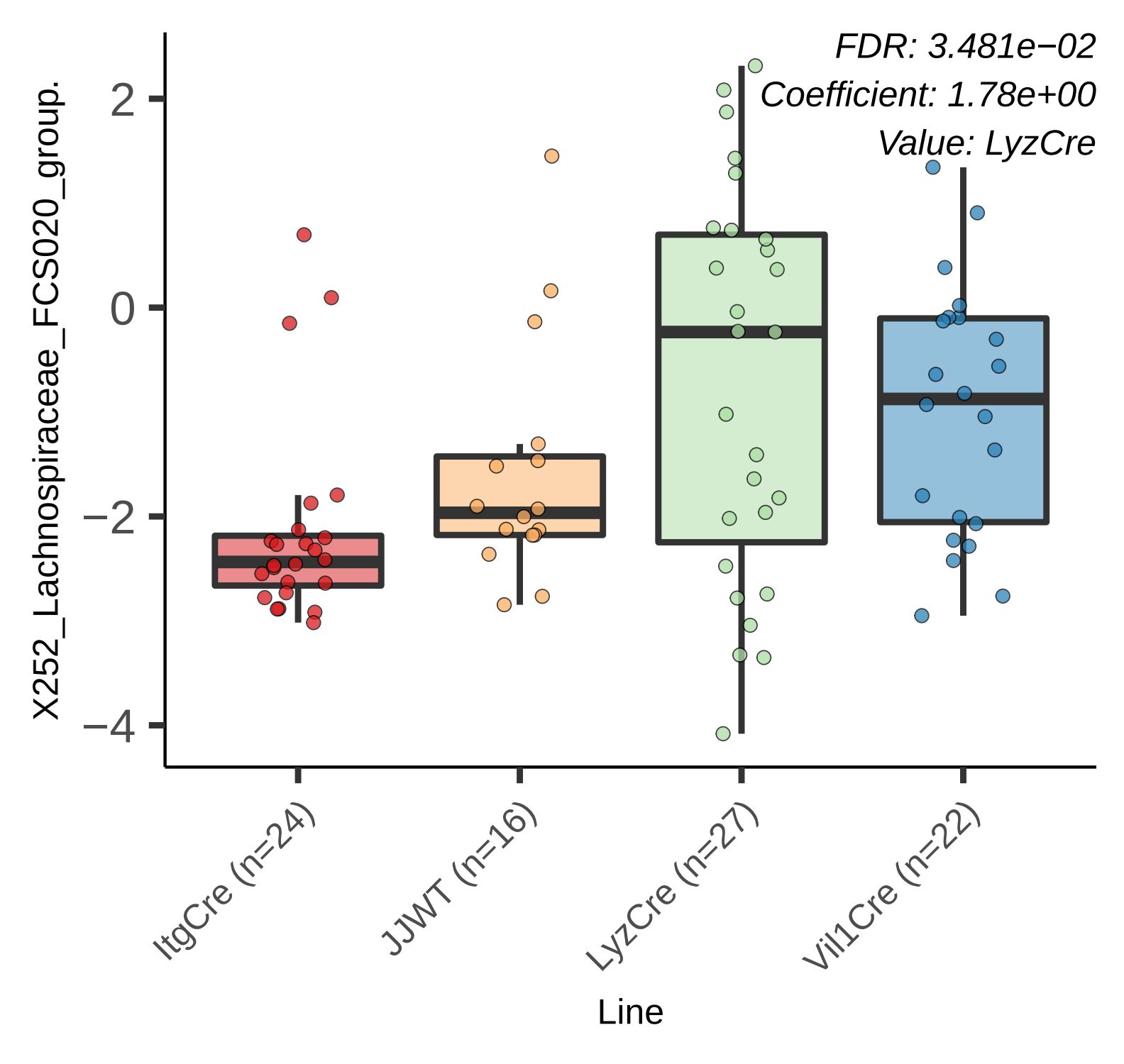

FDR: 3.181e−02
Coefficient: −2.21e+00
Value: JJWT
X104_Muribaculaceae..f.
2
0
−2
ItgCre (n=24)
JJWT (n=16)
LyzCre (n=27)
Vil1Cre (n=22)
Line

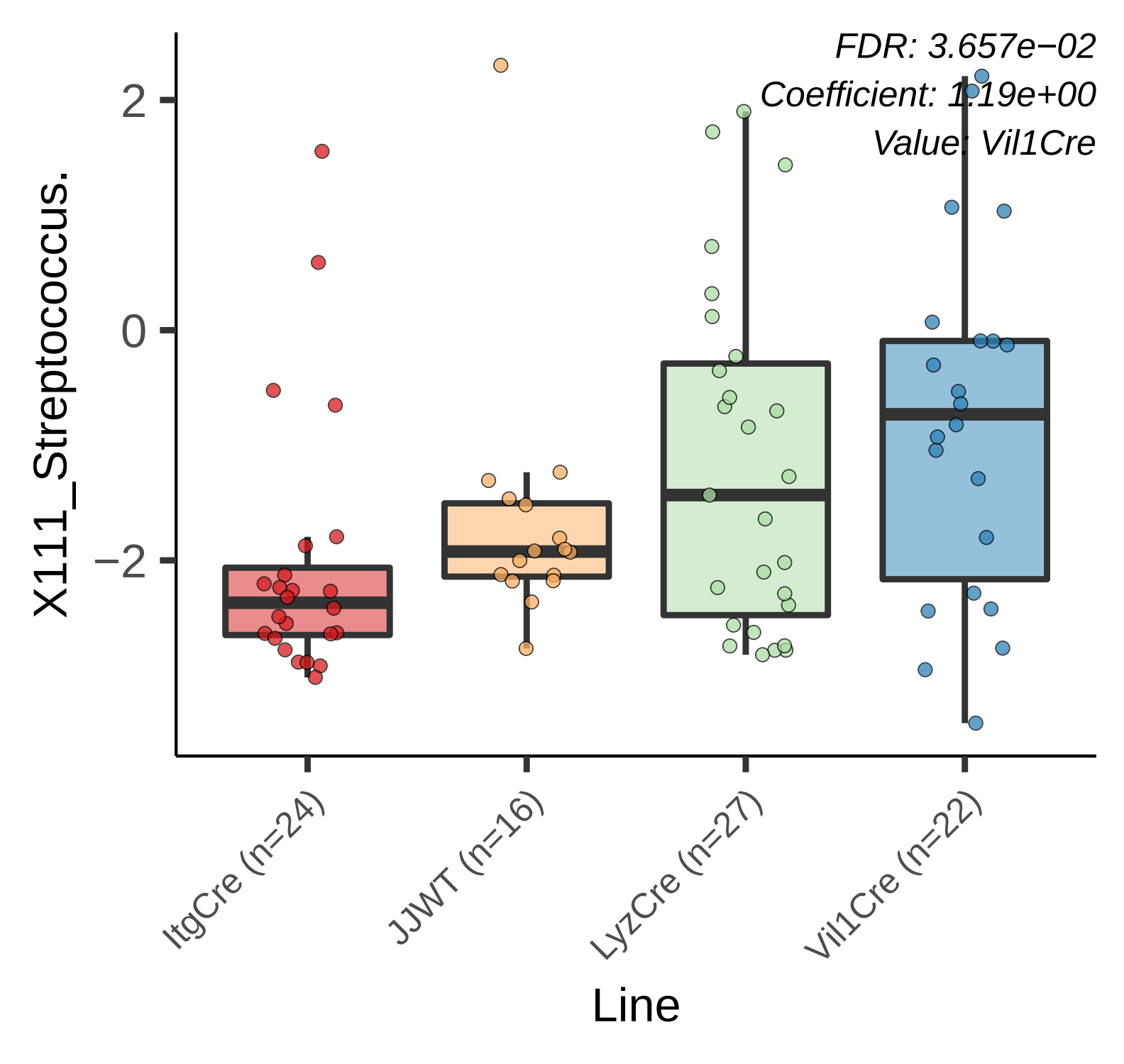

FDR: 3.209e−02
Coefficient: −1.76e+00
Value: JJWT
X233_Muribaculaceae..f.
2
1
0
−1
−2
−3
ItgCre (n=24)
JJWT (n=16)
LyzCre (n=27)
Vil1Cre (n=22)
Line

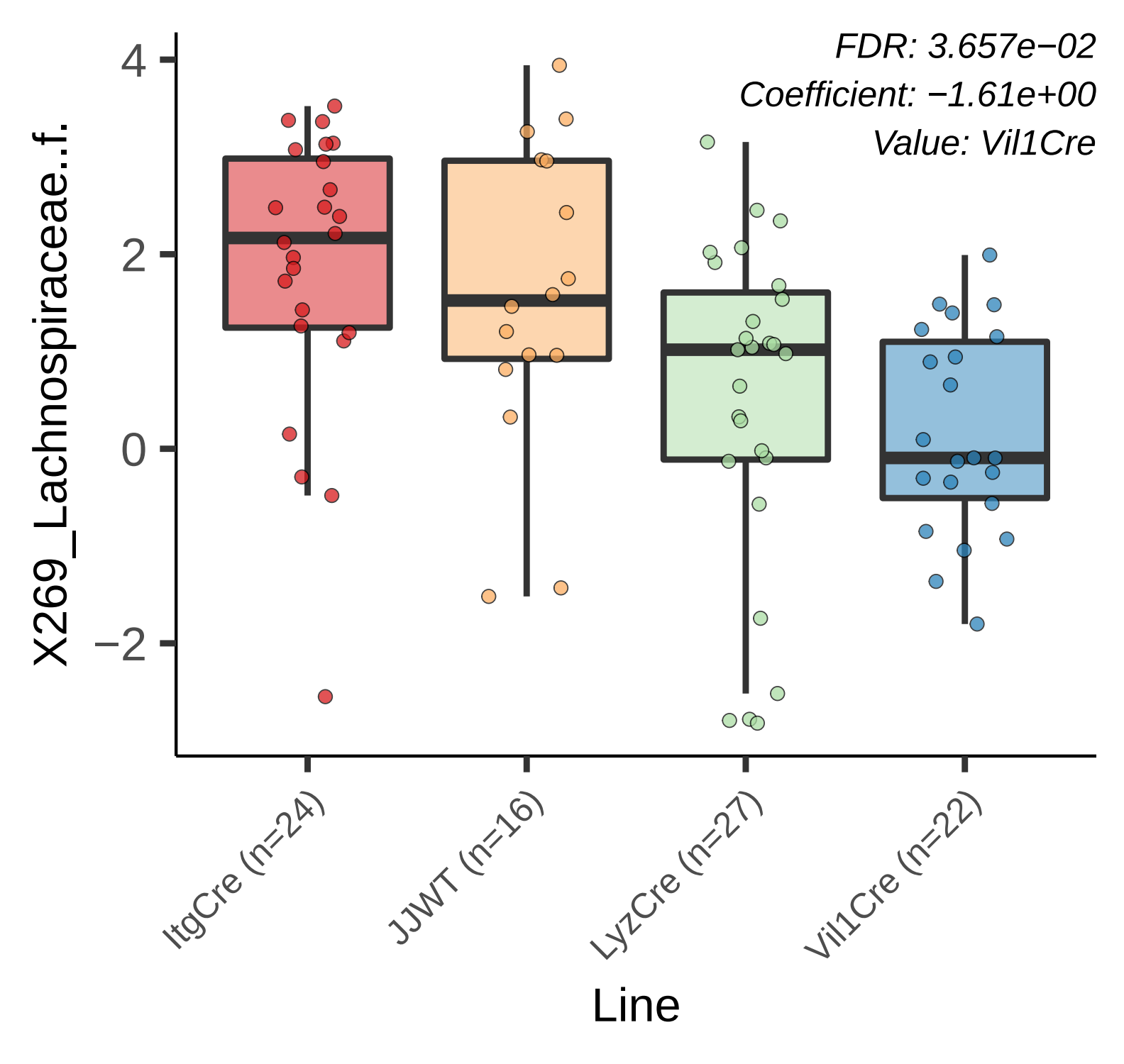

FDR: 3.326e−02
Coefficient: −1.96e+00
Value: LyzCre
X205_Lachnospiraceae_FCS020_group.
4
2
0
−2
ItgCre (n=24)
JJWT (n=16)
LyzCre (n=27)
Vil1Cre (n=22)
Line

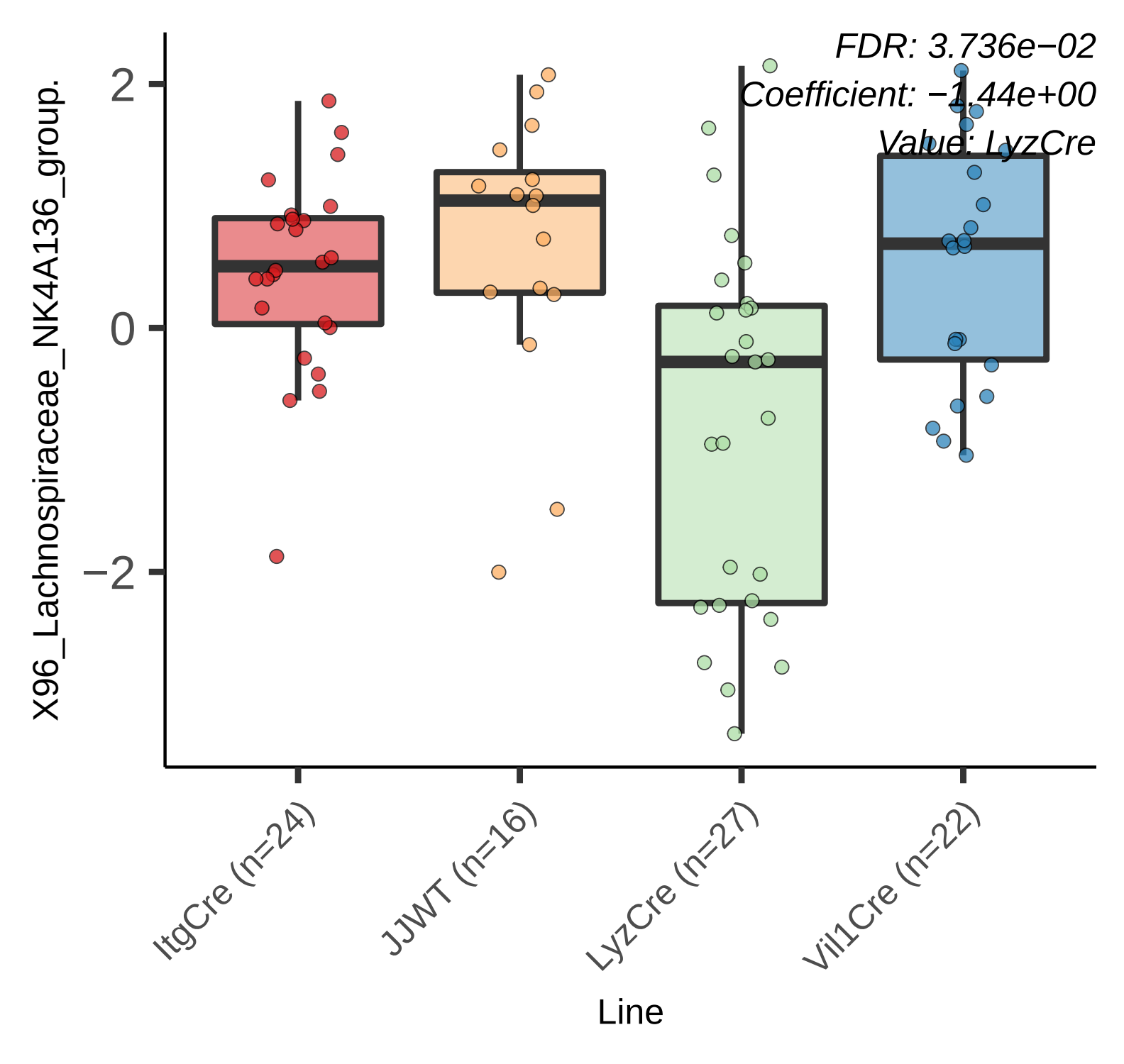

FDR: 3.473e−02
Coefficient: 1.44e+00
Value: LyzCre
X58_Lachnospiraceae_NK4A136_group.
2
0
−2
ItgCre (n=24)
JJWT (n=16)
LyzCre (n=27)
Vil1Cre (n=22)
Line

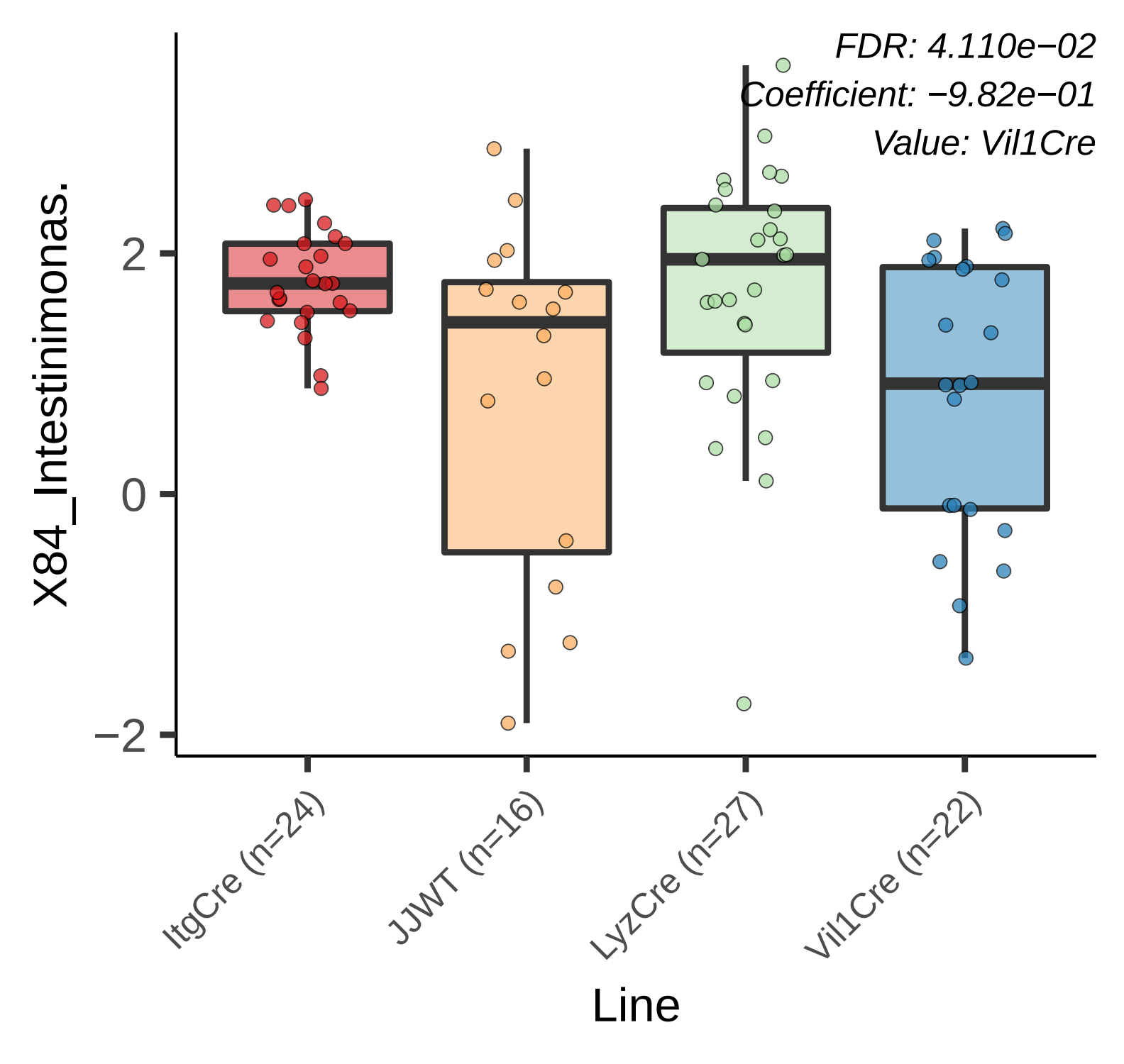

FDR: 3.473e−02
Coefficient: 1.29e+00
Value: Vil1Cre
X185_Gastranaerophilales..o.
2
1
0
−1
−2
−3
ItgCre (n=24)
JJWT (n=16)
LyzCre (n=27)
Vil1Cre (n=22)
Line

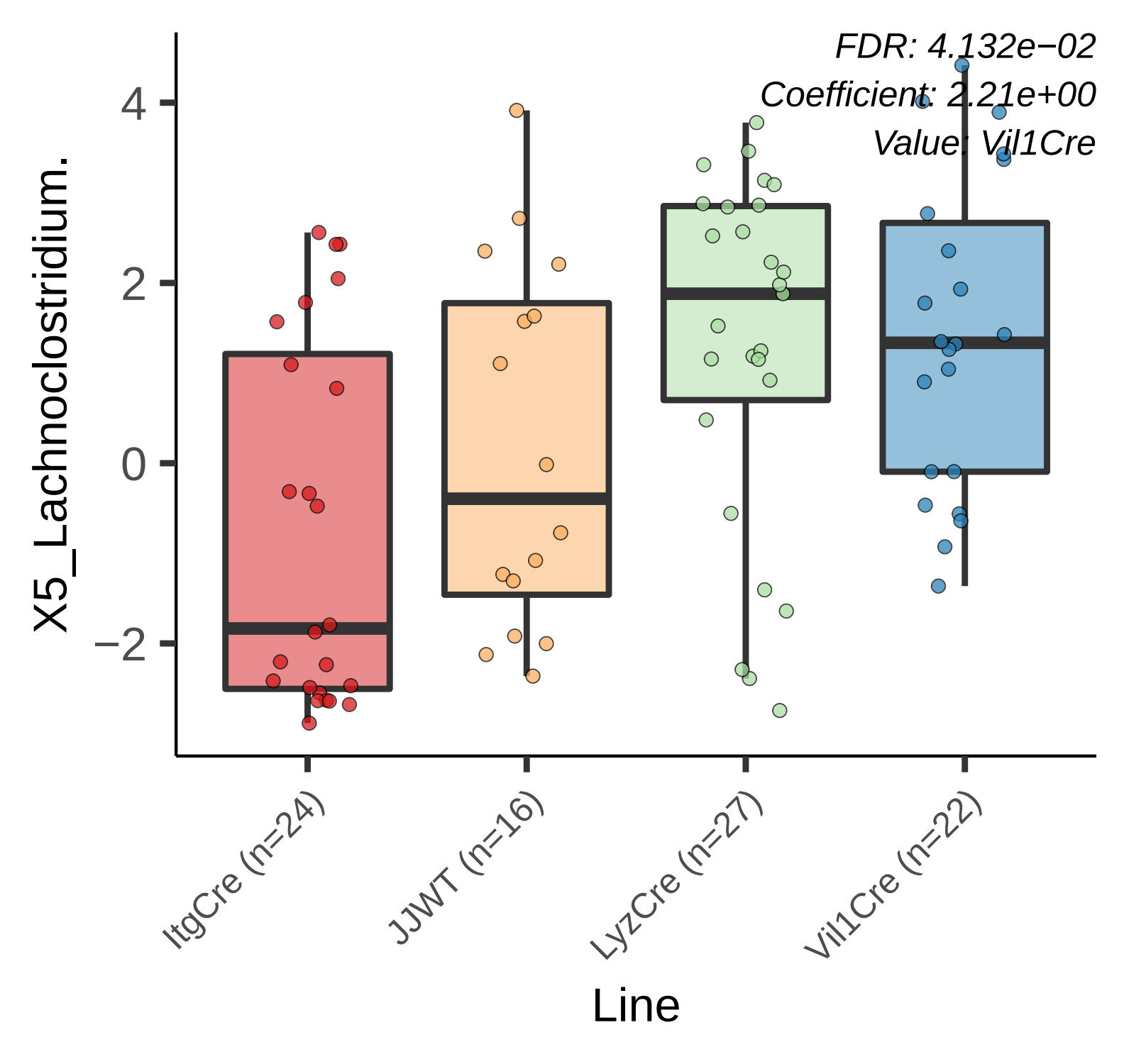

FDR: 3.481e−02
Coefficient: 1.78e+00
Value: LyzCre
X252_Lachnospiraceae_FCS020_group.
2
0
−2
−4
ItgCre (n=24)
JJWT (n=16)
LyzCre (n=27)
Vil1Cre (n=22)
Line

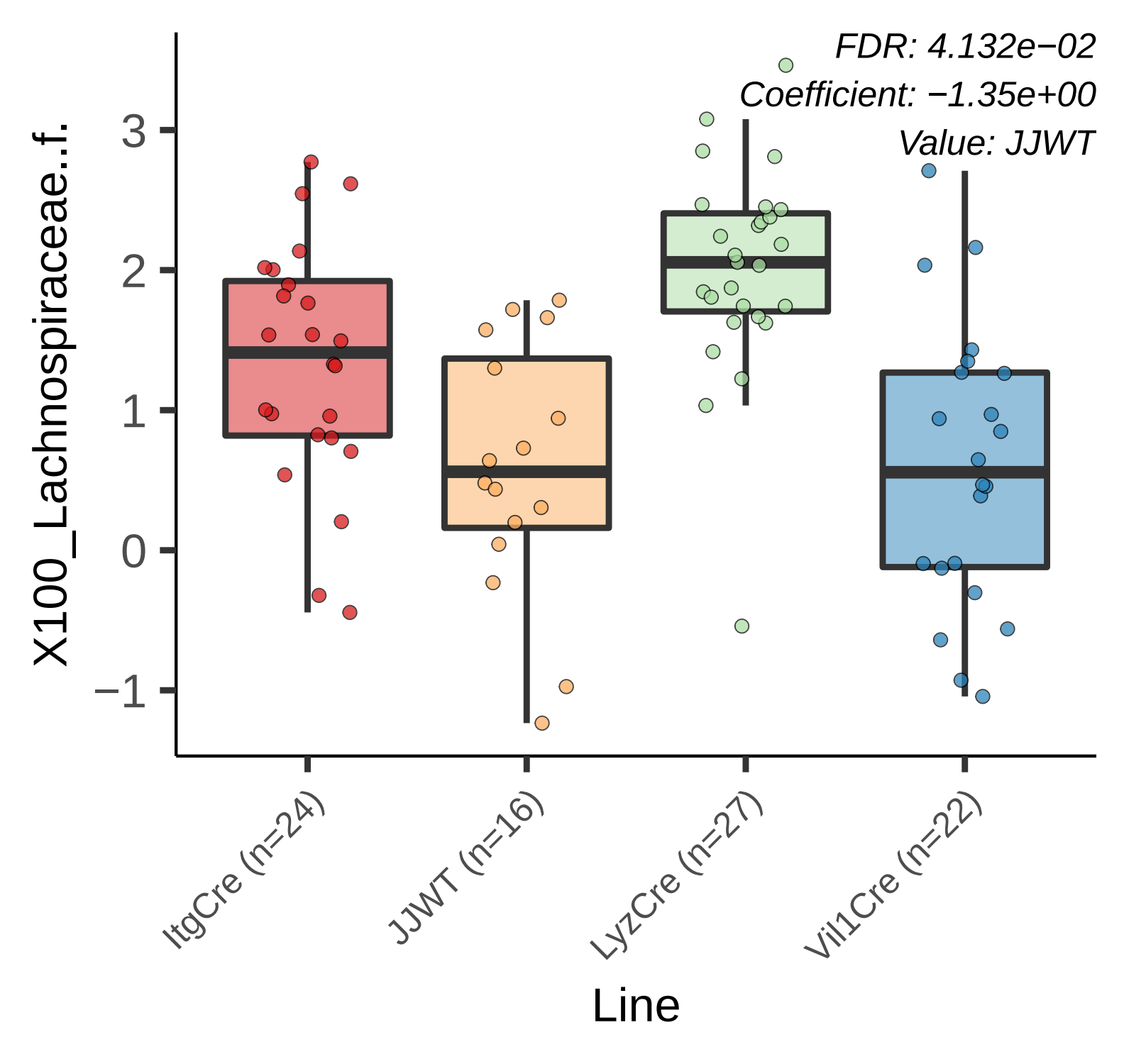

FDR: 3.657e−02
Coefficient: 1.19e+00
Value: Vil1Cre
X111_Streptococcus.
2
0
−2
ItgCre (n=24)
JJWT (n=16)
LyzCre (n=27)
Vil1Cre (n=22)
Line

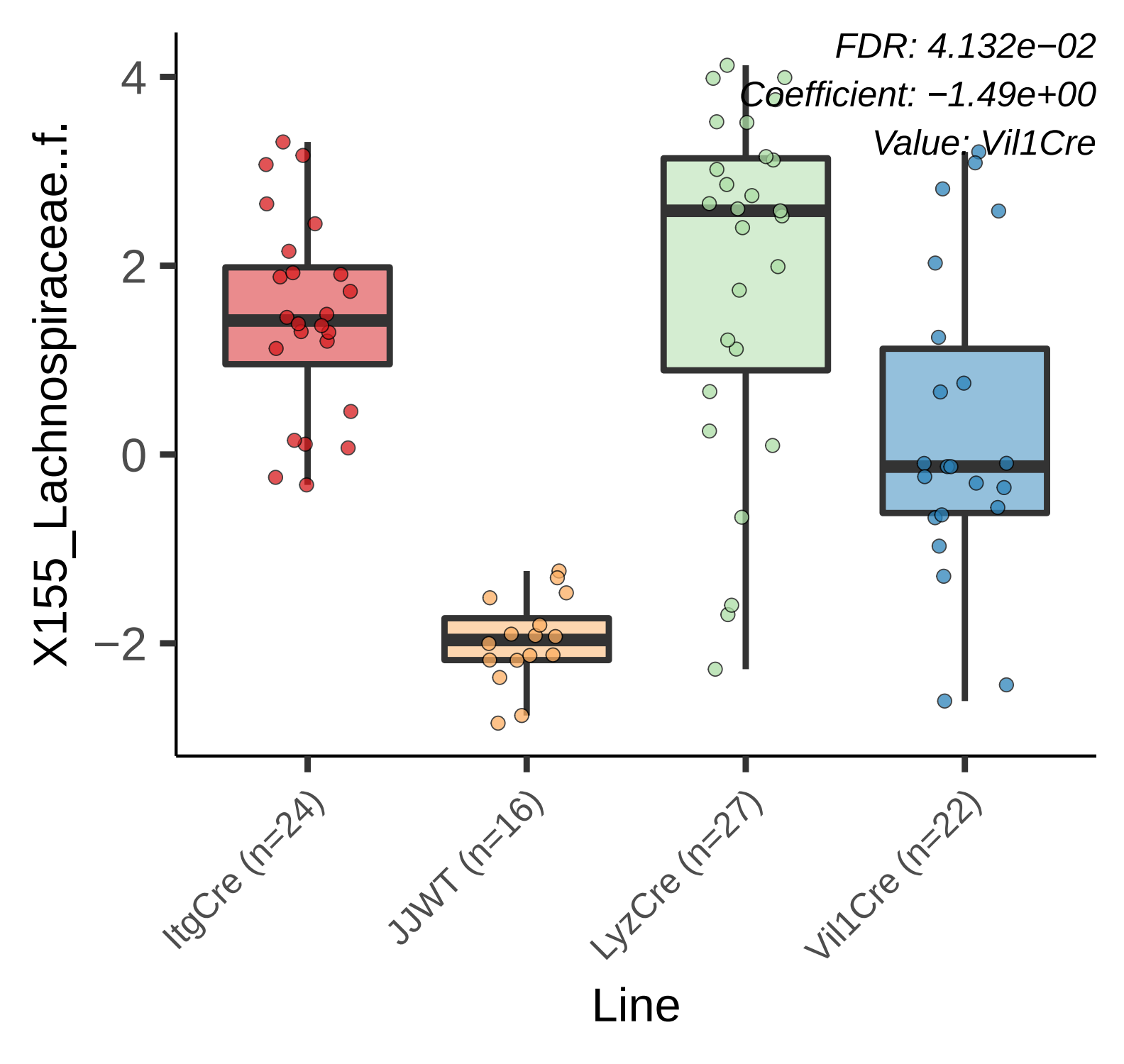

FDR: 3.657e−02
Coefficient: −1.61e+00
Value: Vil1Cre
X269_Lachnospiraceae.f.
4
2
0
−2
ItgCre (n=24)
JJWT (n=16)
LyzCre (n=27)
Vil1Cre (n=22)
Line

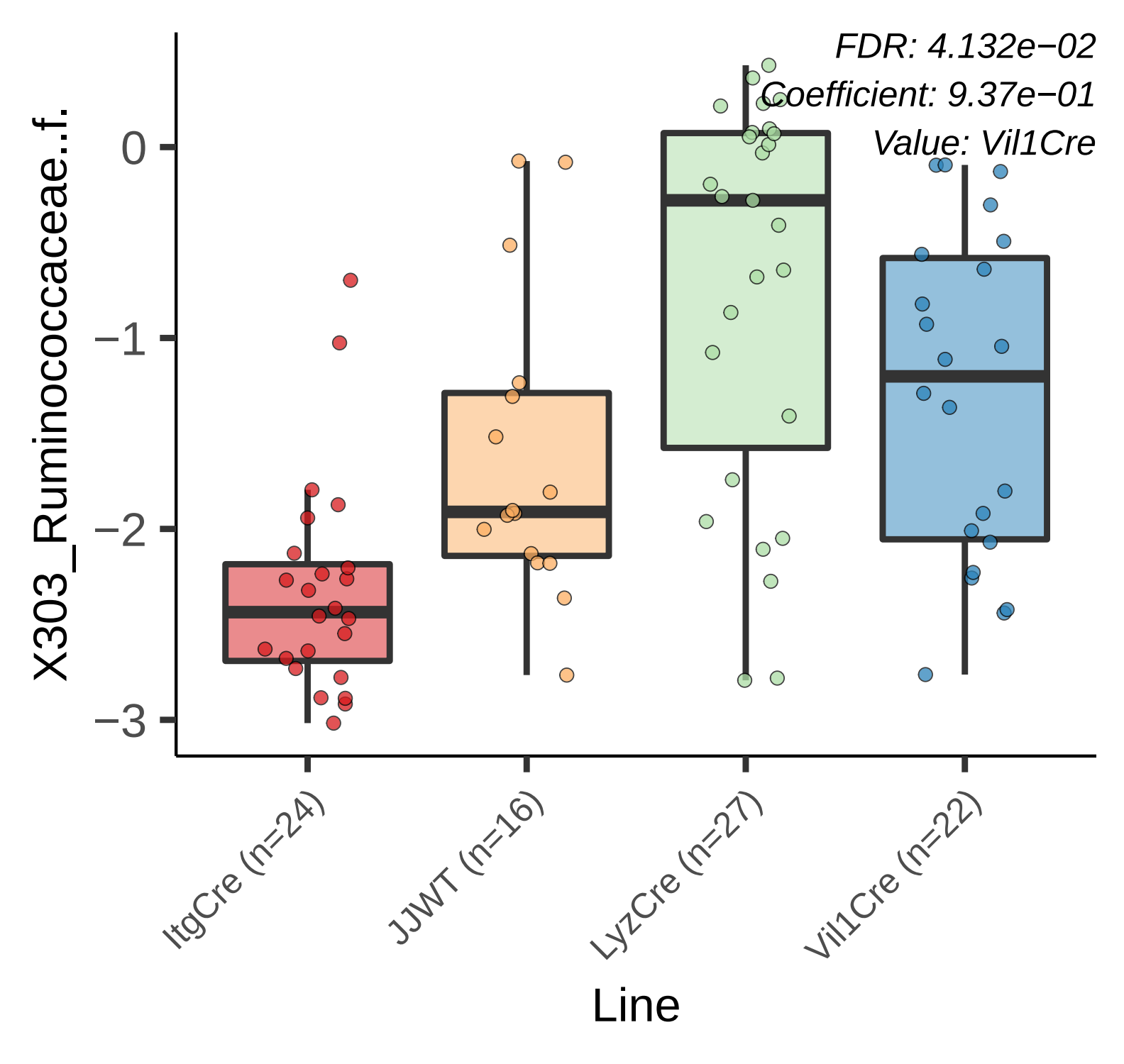

FDR: 3.736e−02
Coefficient: −1.44e+00
Value: LyzCre
X96_Lachnospiraceae_NK4A136_group.
2
0
−2
ItgCre (n=24)
JJWT (n=16)
LyzCre (n=27)
Vil1Cre (n=22)
Line

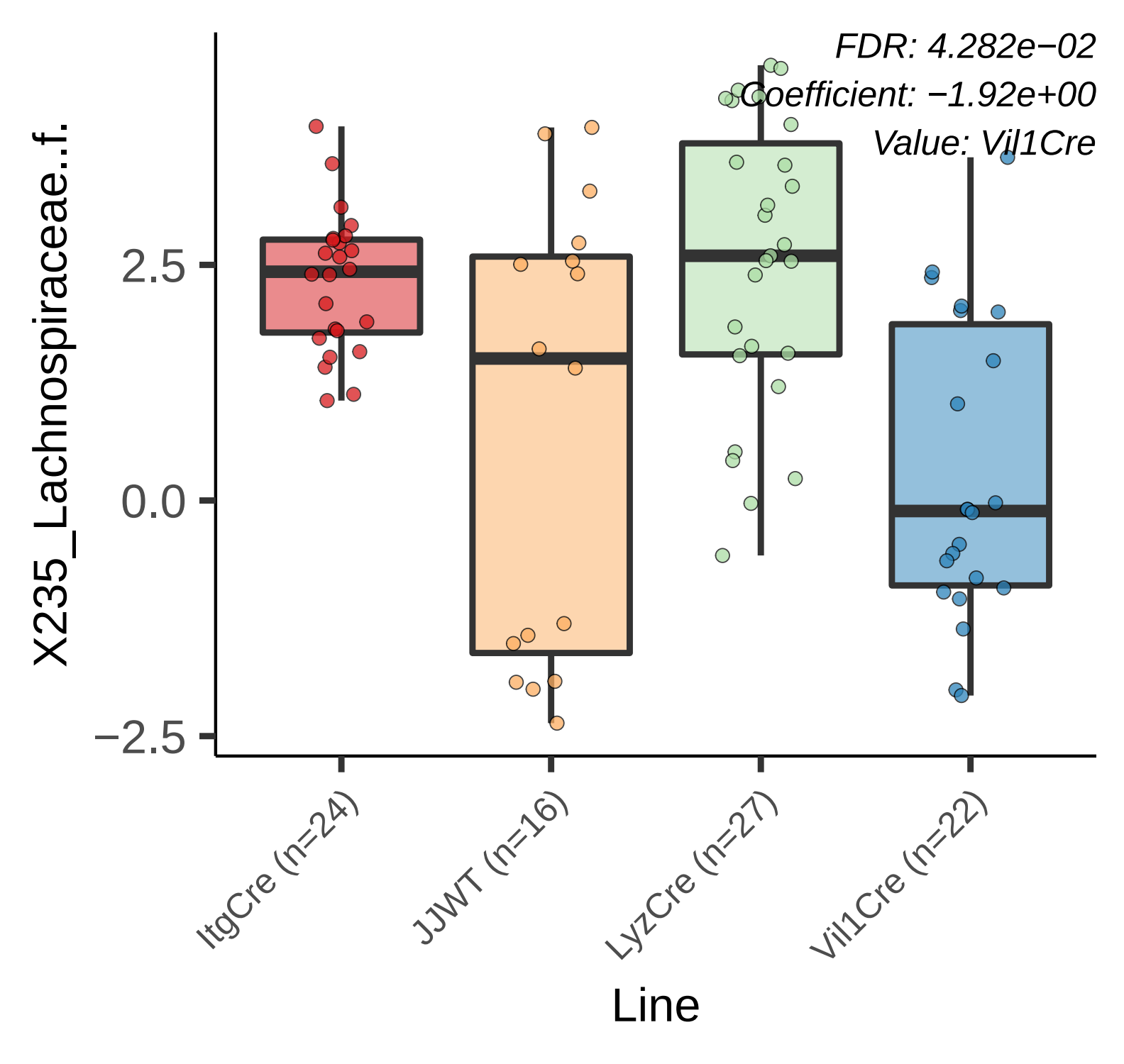

FDR: 4.110e−02
Coefficient: −9.82e−01
Value: Vil1Cre
X84_Intestinimonas.
2
0
−2
ItgCre (n=24)
JJWT (n=16)
LyzCre (n=27)
Vil1Cre (n=22)
Line

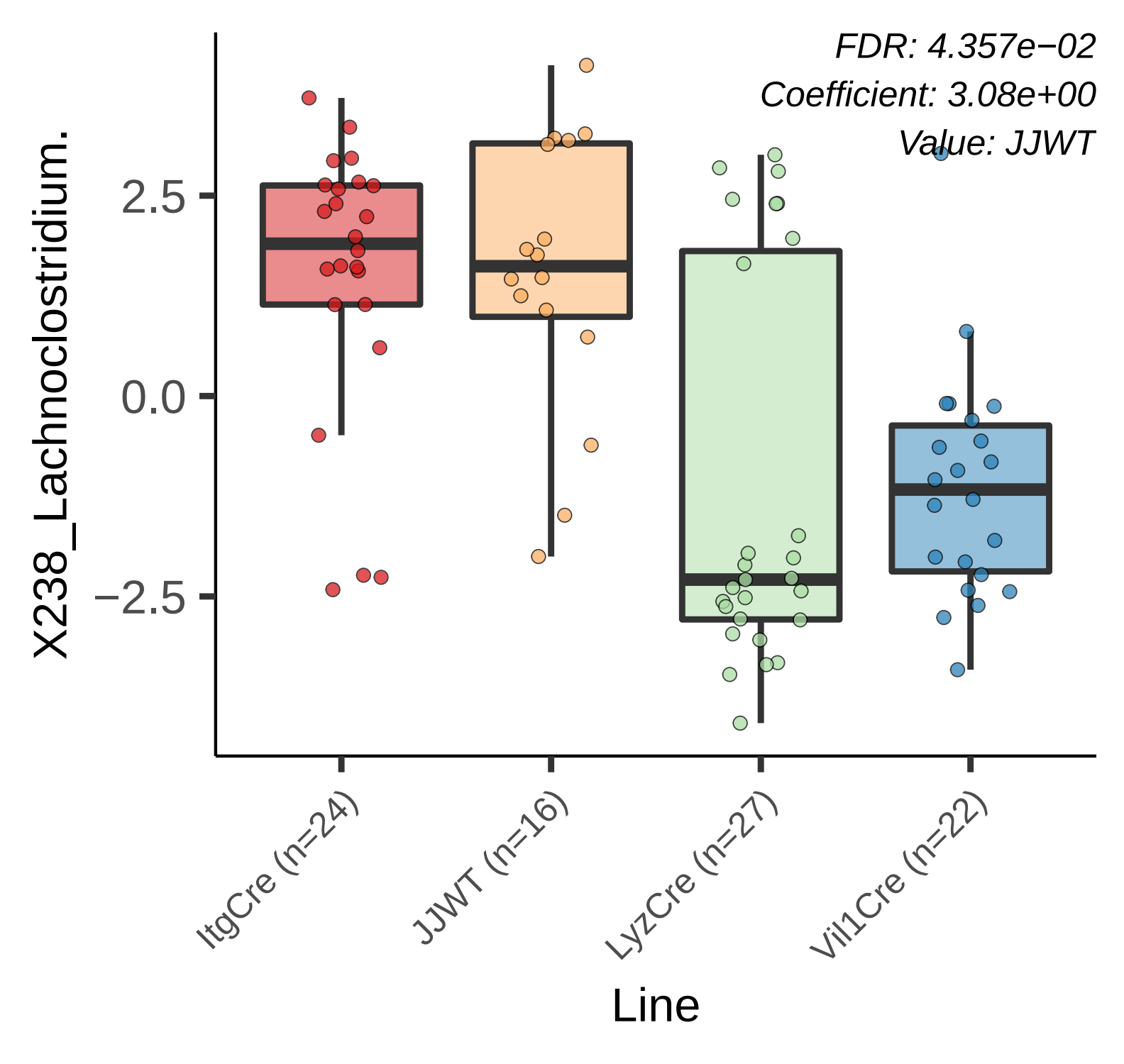

FDR: 4.132e−02
Coefficient: 2.21e+00
Value: Vil1Cre
X5_Lachnoclostridium.
4
2
0
−2
ItgCre (n=24)
JJWT (n=16)
LyzCre (n=27)
Vil1Cre (n=22)
Line

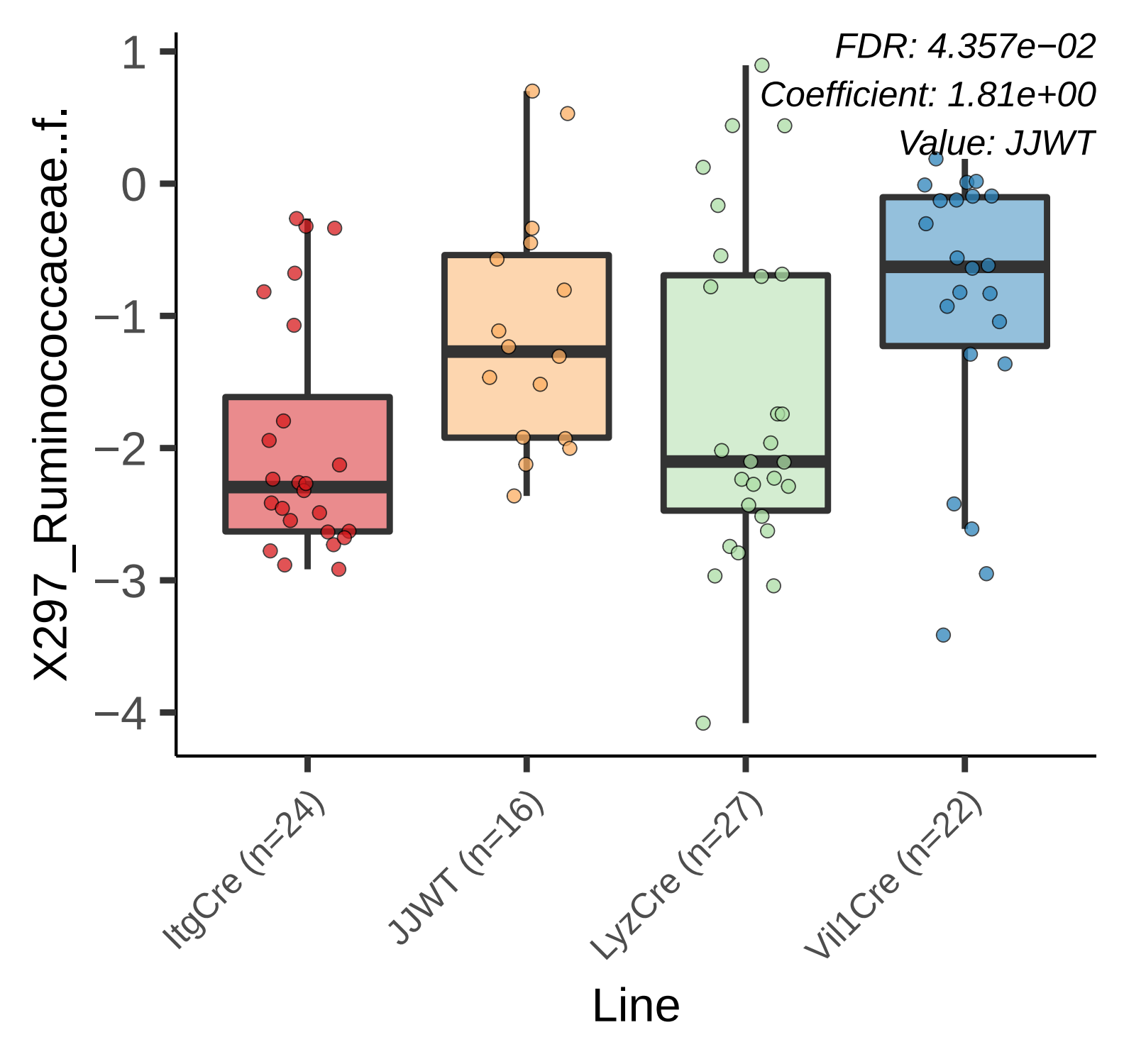

FDR: 4.132e−02
Coefficient: −1.35e+00
Value: JJWT
X100_Lachnospiraceae.f.
3
2
1
0
−1
ItgCre (n=24)
JJWT (n=16)
LyzCre (n=27)
Vil1Cre (n=22)
Line

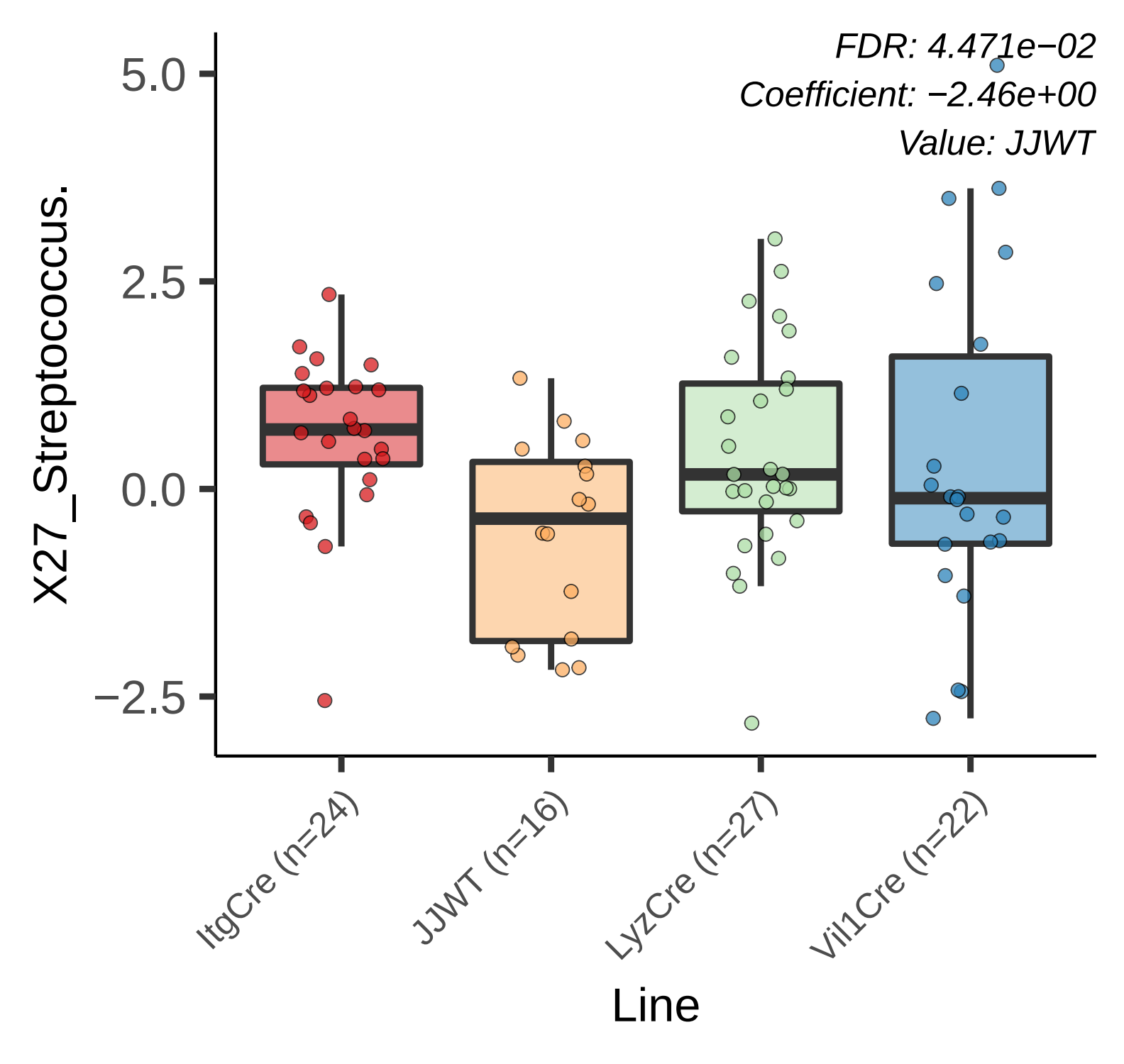

FDR: 4.132e−02
Coefficient: −1.49e+00
Value: Vil1Cre
X155_Lachnospiraceae.f.
4
2
0
−2
ItgCre (n=24)
JJWT (n=16)
LyzCre (n=27)
Vil1Cre (n=22)
Line

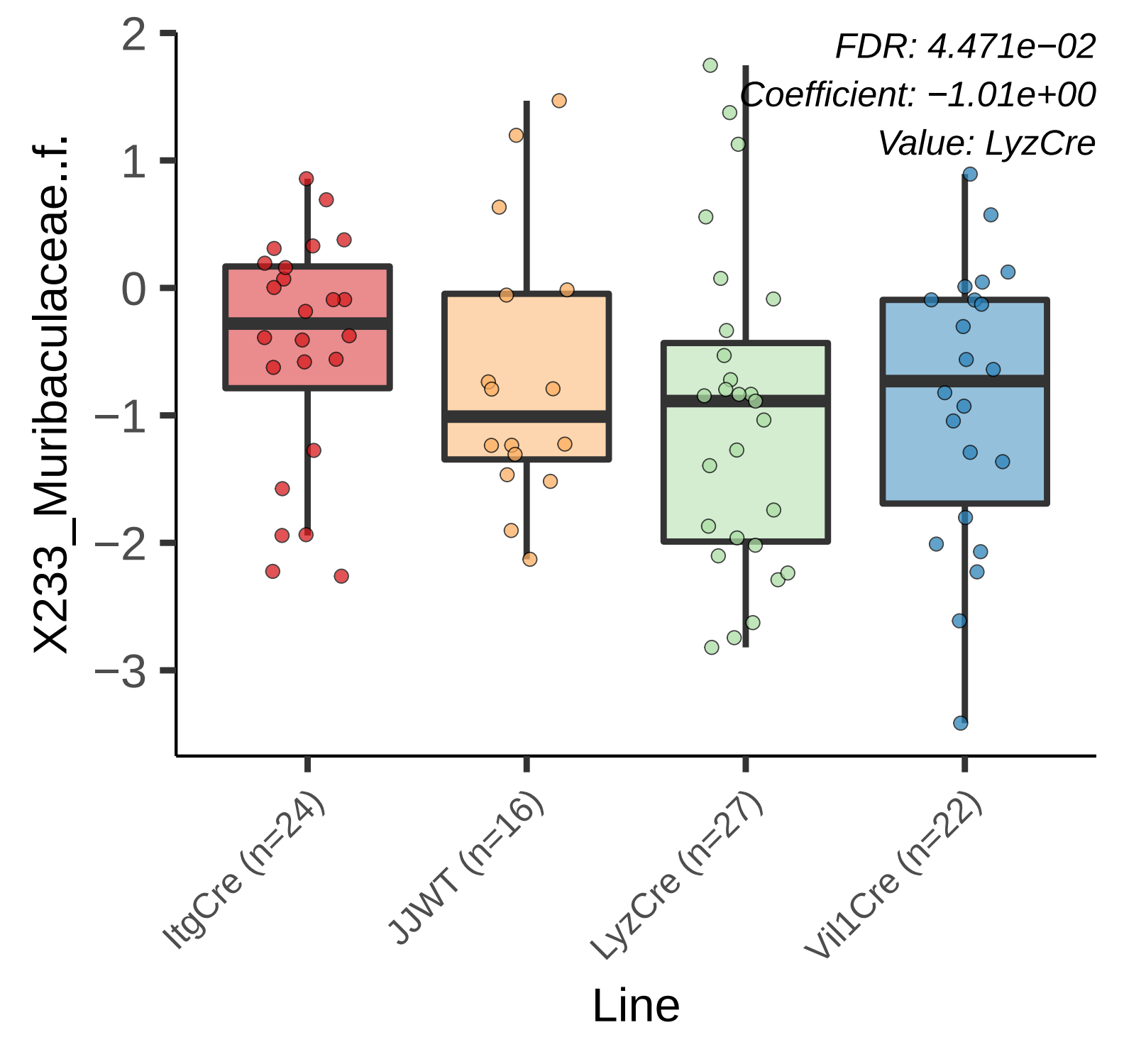

FDR: 4.132e−02
Coefficient: 9.37e−01
Value: Vil1Cre
X303_Ruminococcaceae..f.
0
−1
−2
−3
ItgCre (n=24)
JJWT (n=16)
LyzCre (n=27)
Vil1Cre (n=22)
Line

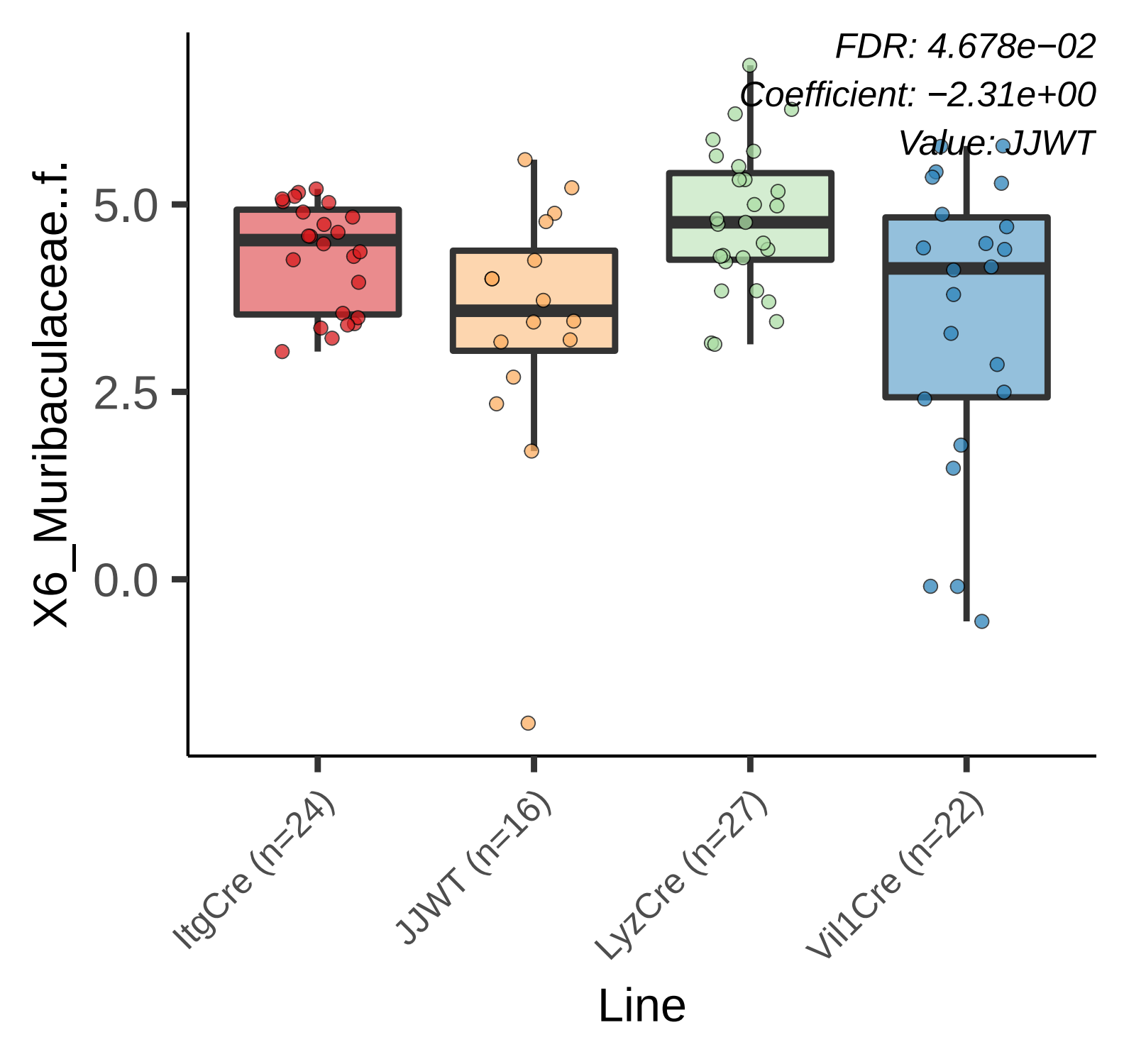

FDR: 4.282e−02
Coefficient: −1.92e+00
Value: Vil1Cre
X235_Lachnospiraceae.f.
2.5
0.0
−2.5
ItgCre (n=24)
JJWT (n=16)
LyzCre (n=27)
Vil1Cre (n=22)
Line

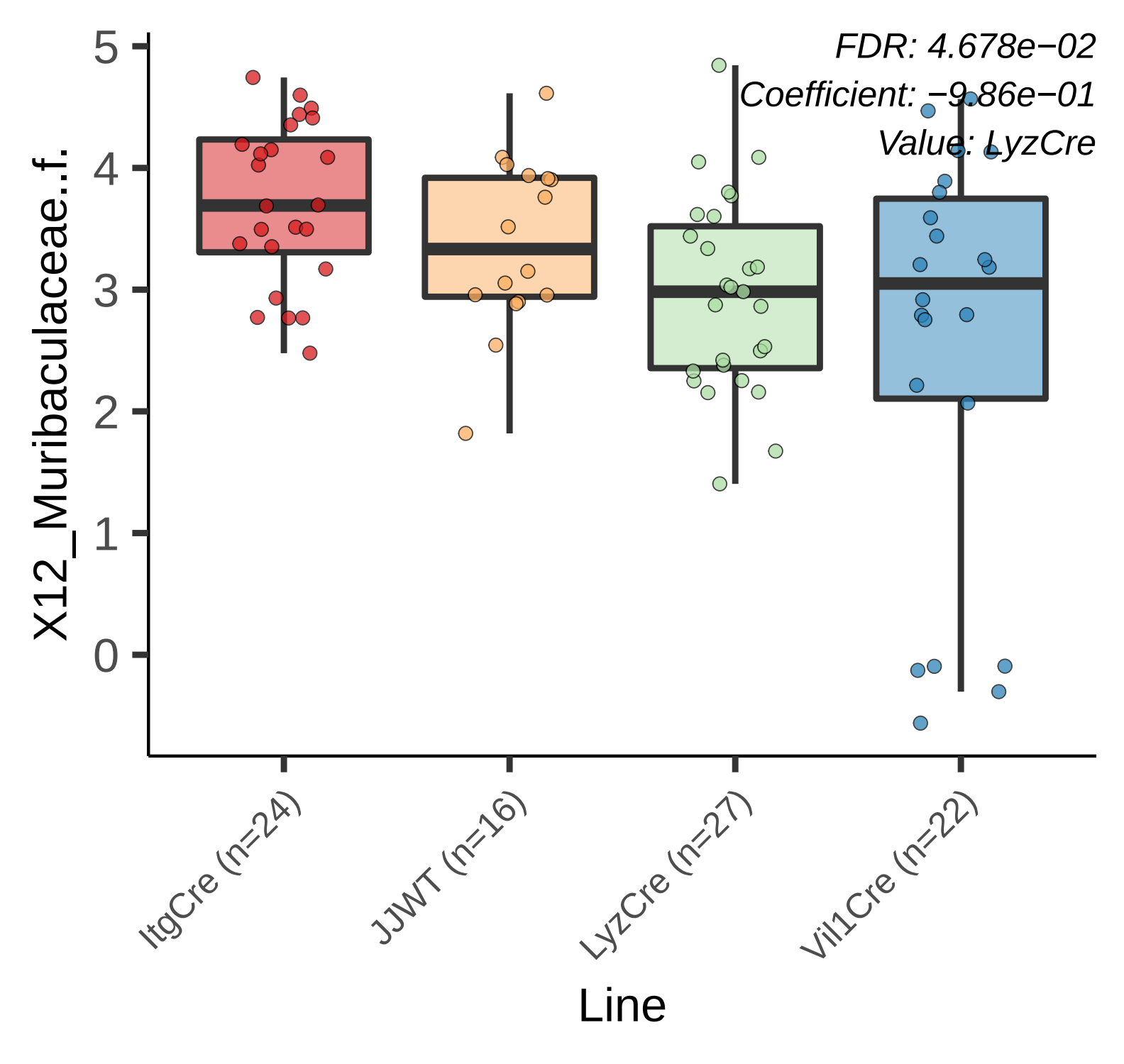

FDR: 4.357e−02
Coefficient: 3.08e+00
Value: JJWT
X238_Lachnoclostridium.
2.5
0.0
−2.5
ItgCre (n=24)
JJWT (n=16)
LyzCre (n=27)
Vil1Cre (n=22)
Line

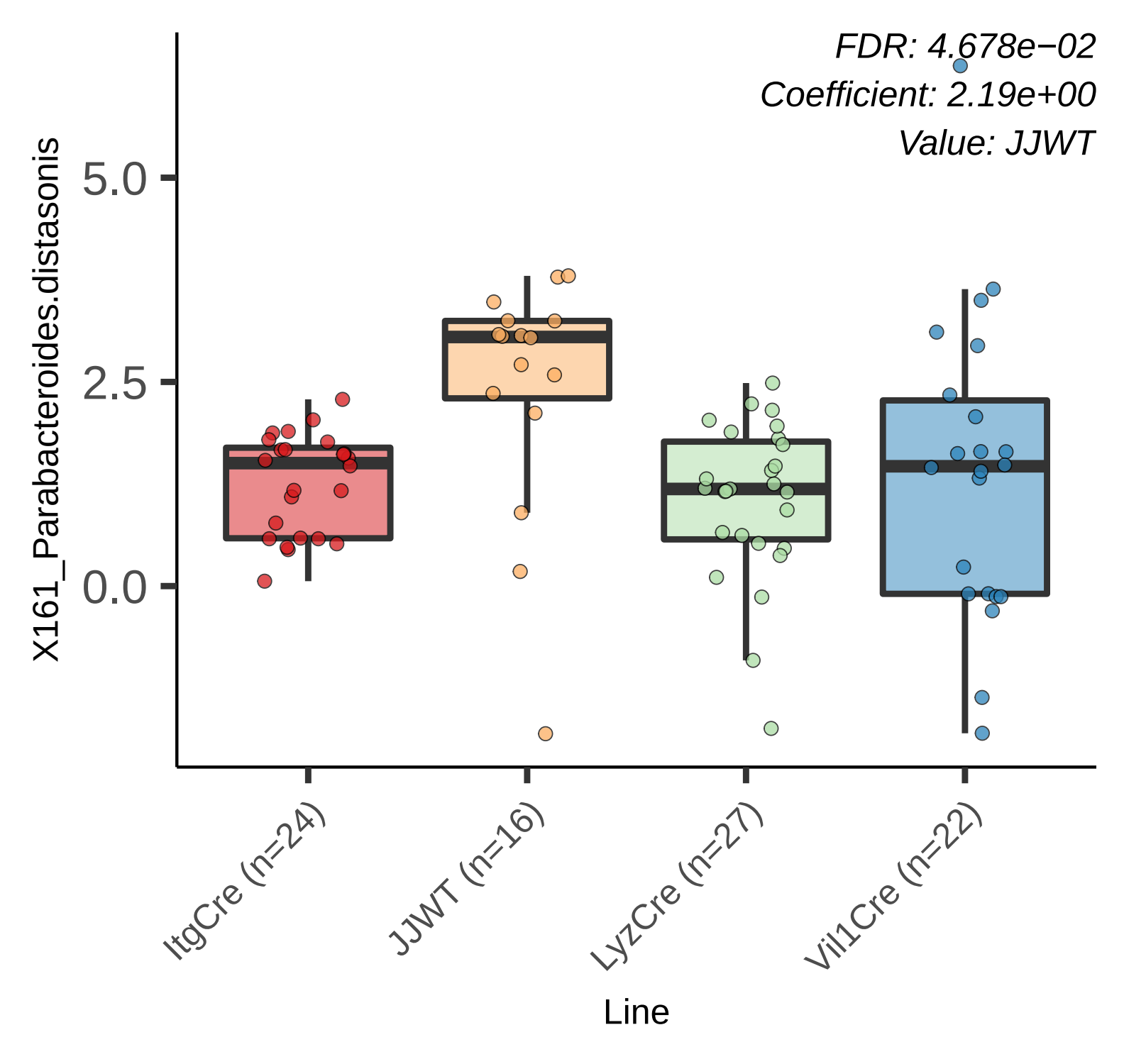

FDR: 4.357e−02
Coefficient: 1.81e+00
Value: JJWT
X297_Ruminococcaceae..f.
1
0
−1
−2
−3
−4
ItgCre (n=24)
JJWT (n=16)
LyzCre (n=27)
Vil1Cre (n=22)
Line

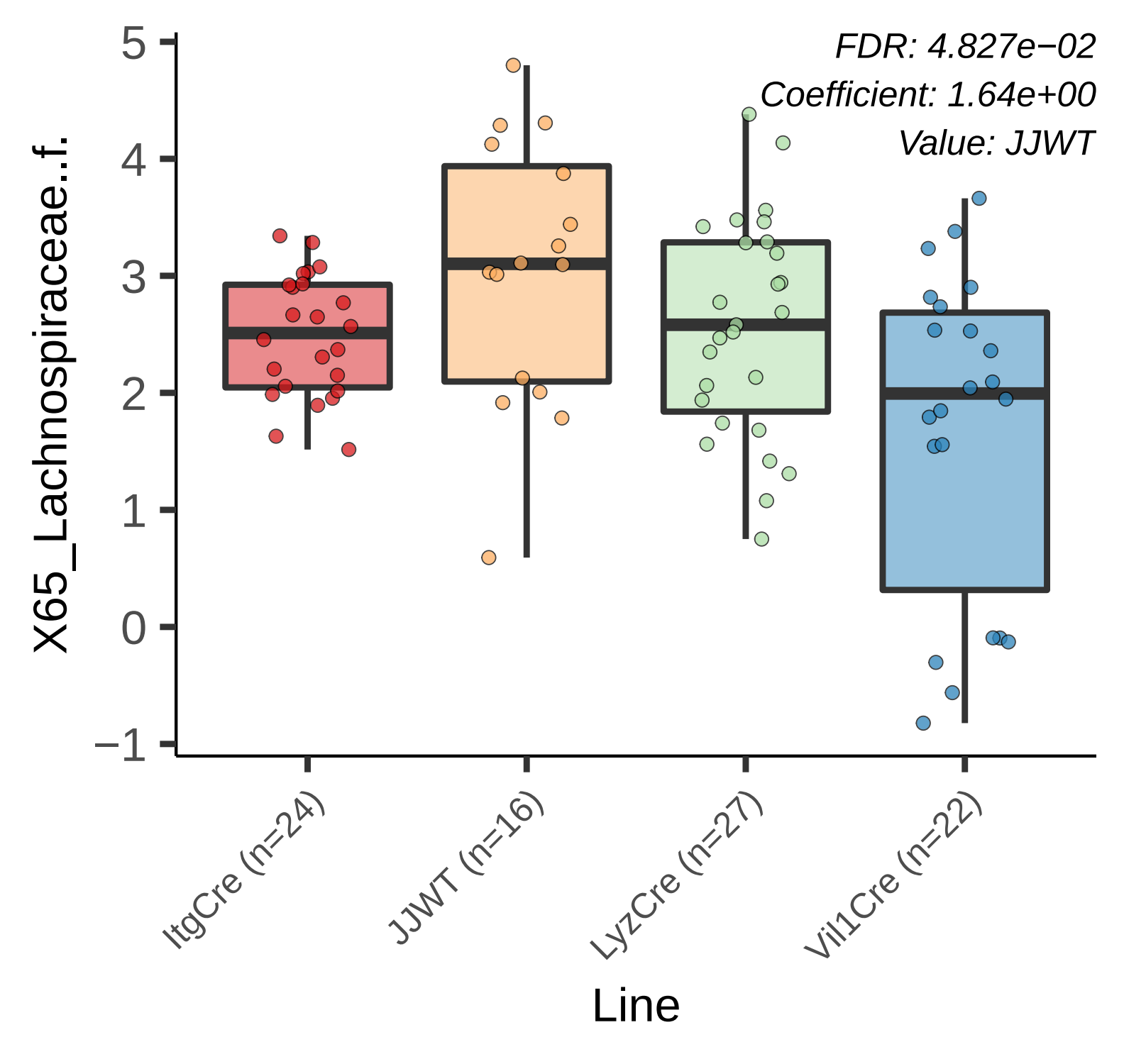

FDR: 4.471e−02
Coefficient: −2.46e+00
Value: JJWT
X27_Streptococcus.
5.0
2.5
0.0
−2.5
ItgCre (n=24)
JJWT (n=16)
LyzCre (n=27)
Vil1Cre (n=22)
Line

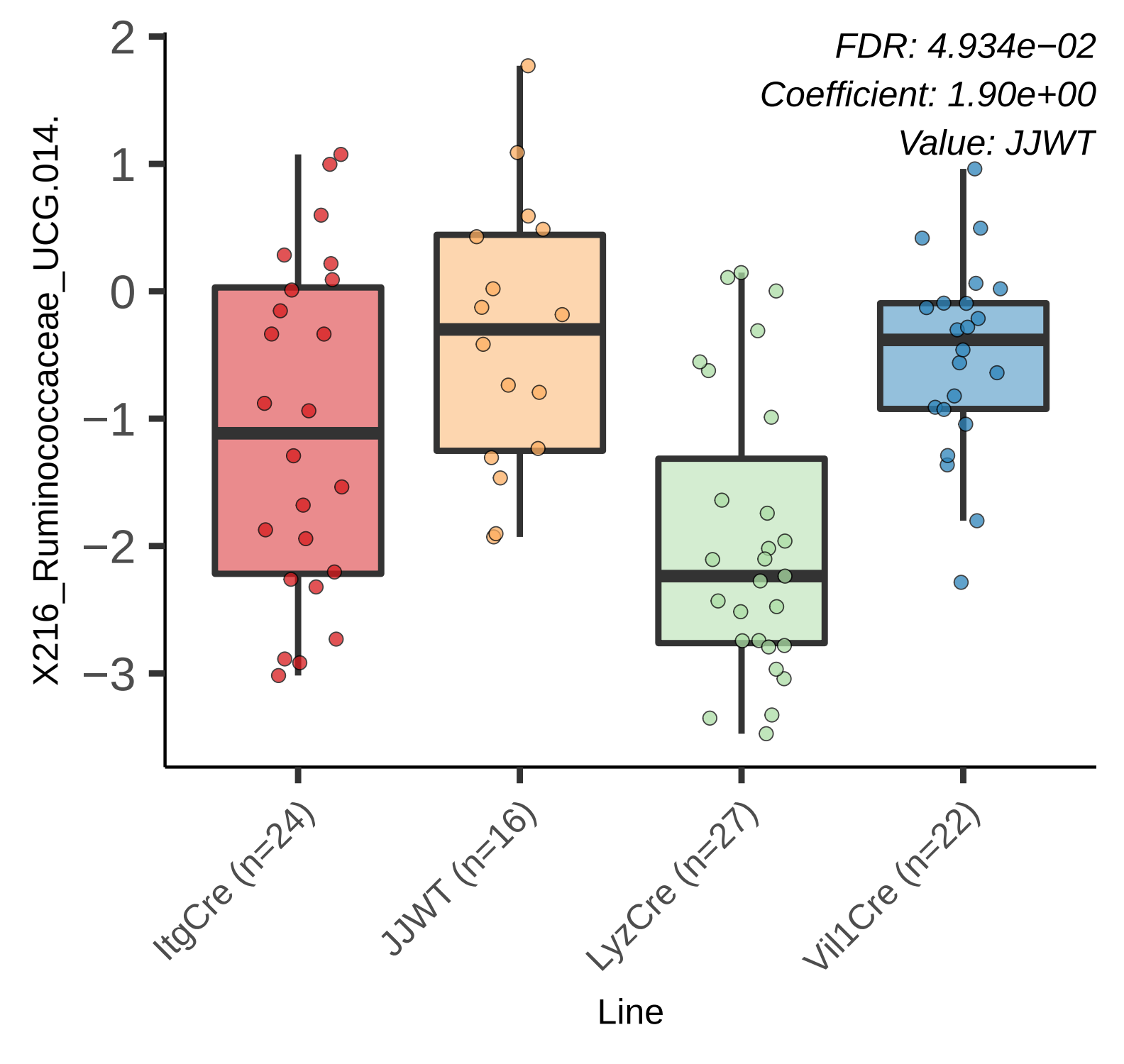

FDR: 4.471e−02
Coefficient: −1.01e+00
Value: LyzCre
X233_Muribaculaceae..f.
2
1
0
−1
−2
−3
ItgCre (n=24)
JJWT (n=16)
LyzCre (n=27)
Vil1Cre (n=22)
Line

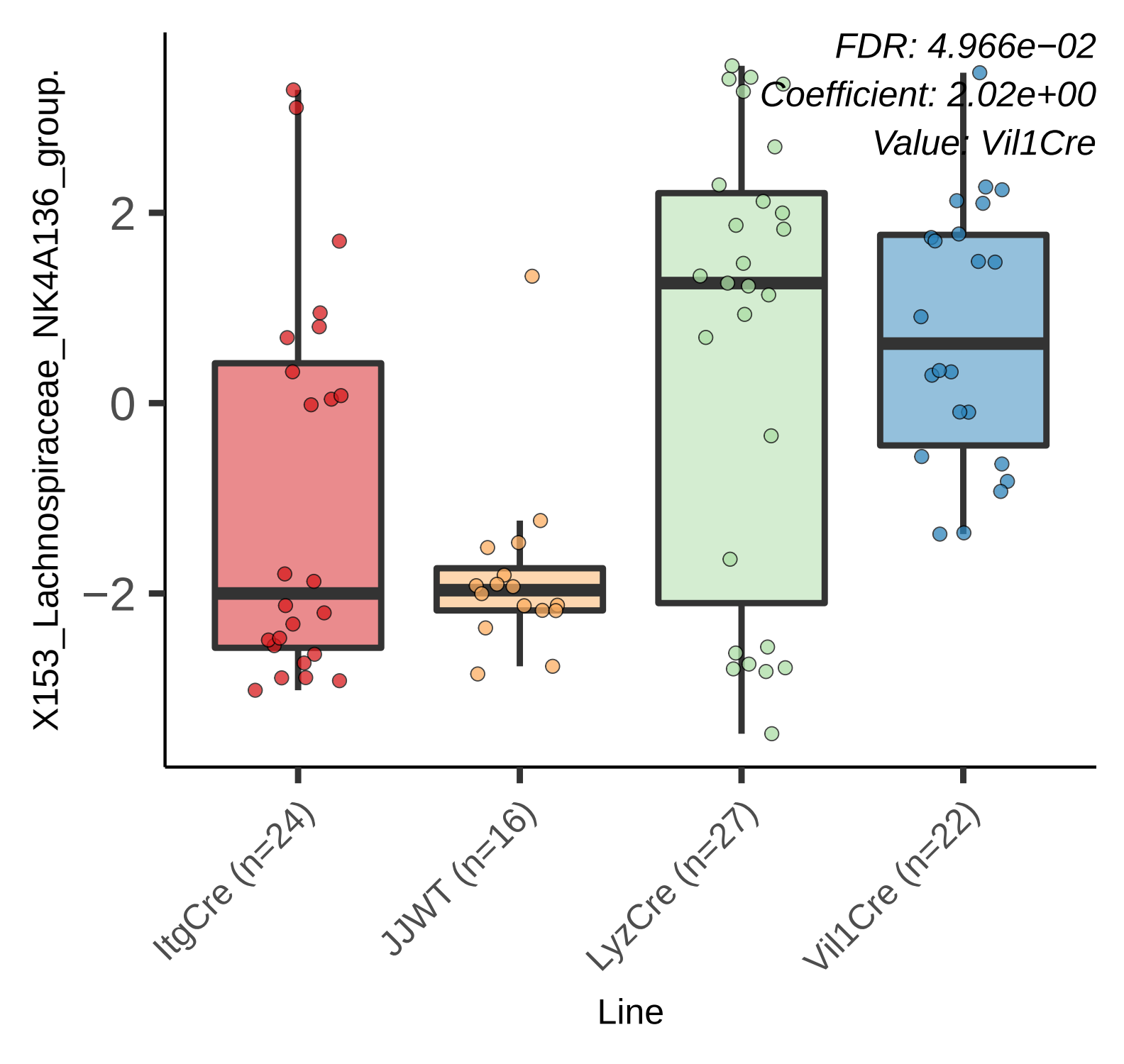

FDR: 4.678e−02
Coefficient: −2.31e+00
Value: JJWT
X6_Muribaculaceae..f.
5.0
2.5
0.0
ItgCre (n=24)
JJWT (n=16)
LyzCre (n=27)
Vil1Cre (n=22)
Line

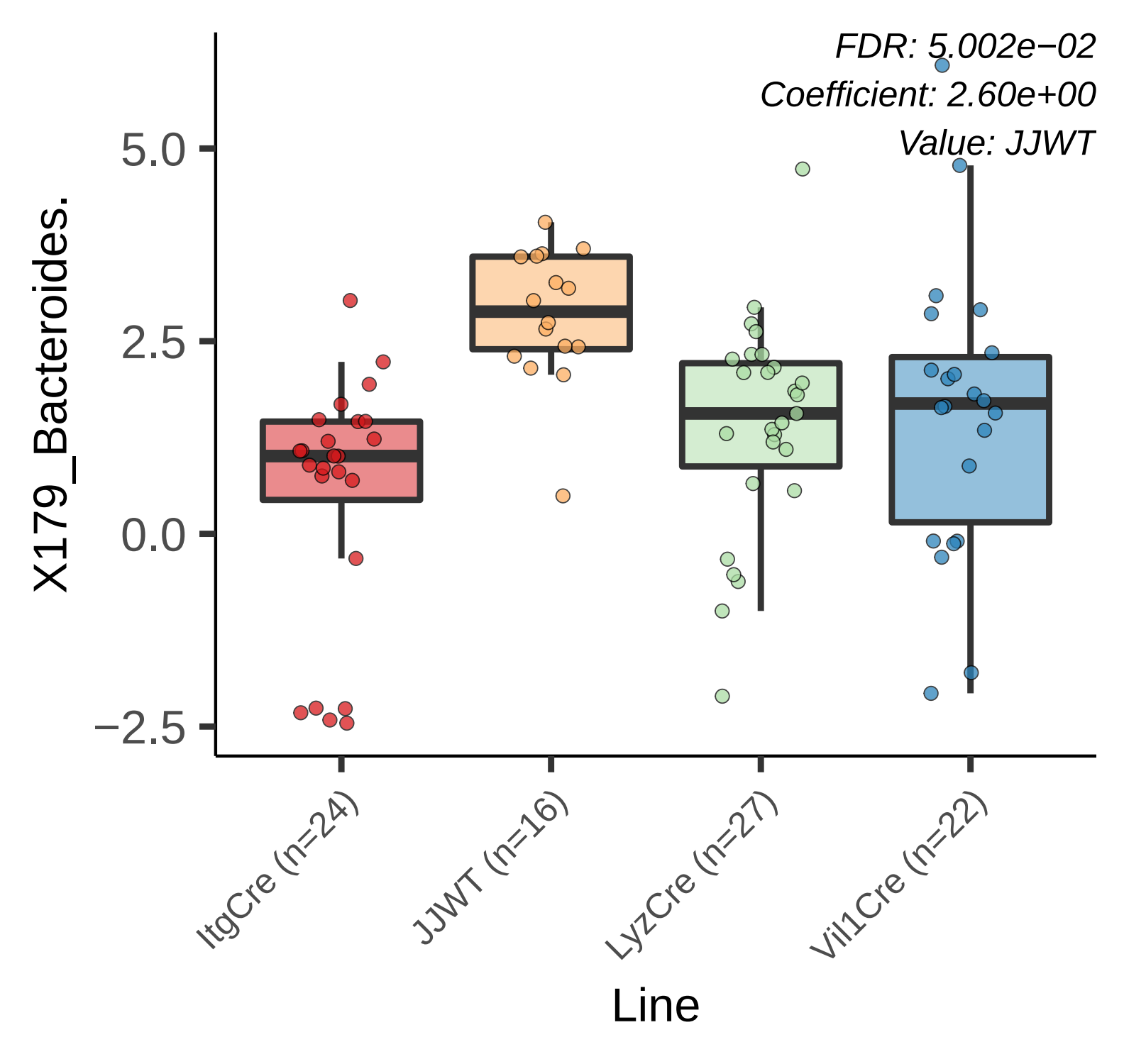

FDR: 4.678e−02
Coefficient: −9.86e−01
Value: LyzCre
X12_Muribaculaceae..f.
5
4
3
2
1
0
ItgCre (n=24)
JJWT (n=16)
LyzCre (n=27)
Vil1Cre (n=22)
Line

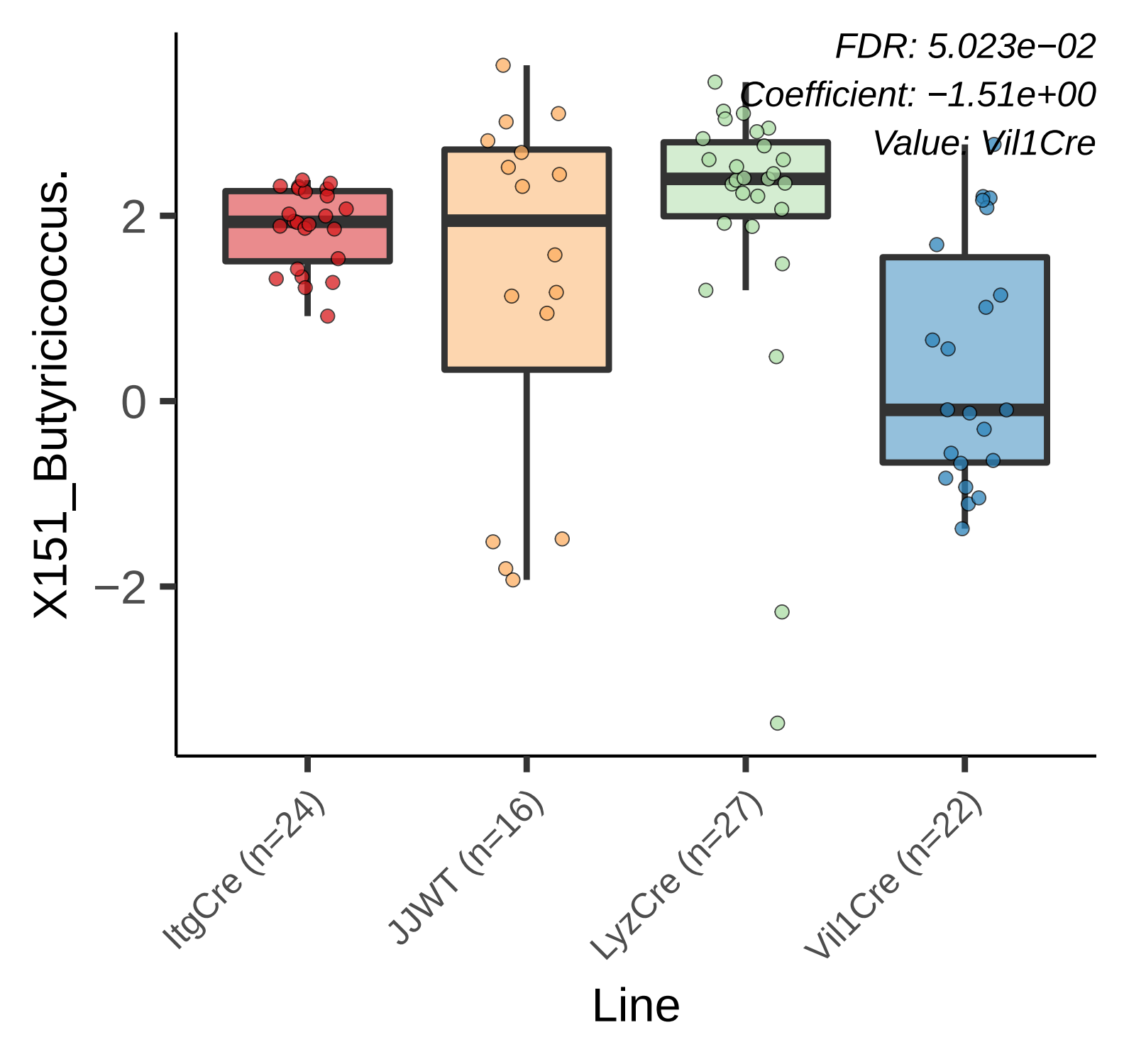

FDR: 4.678e−02
Coefficient: 2.19e+00
Value: JJWT
X161_Parabacteroides.distasonis
5.0
2.5
0.0
ItgCre (n=24)
JJWT (n=16)
LyzCre (n=27)
Vil1Cre (n=22)
Line

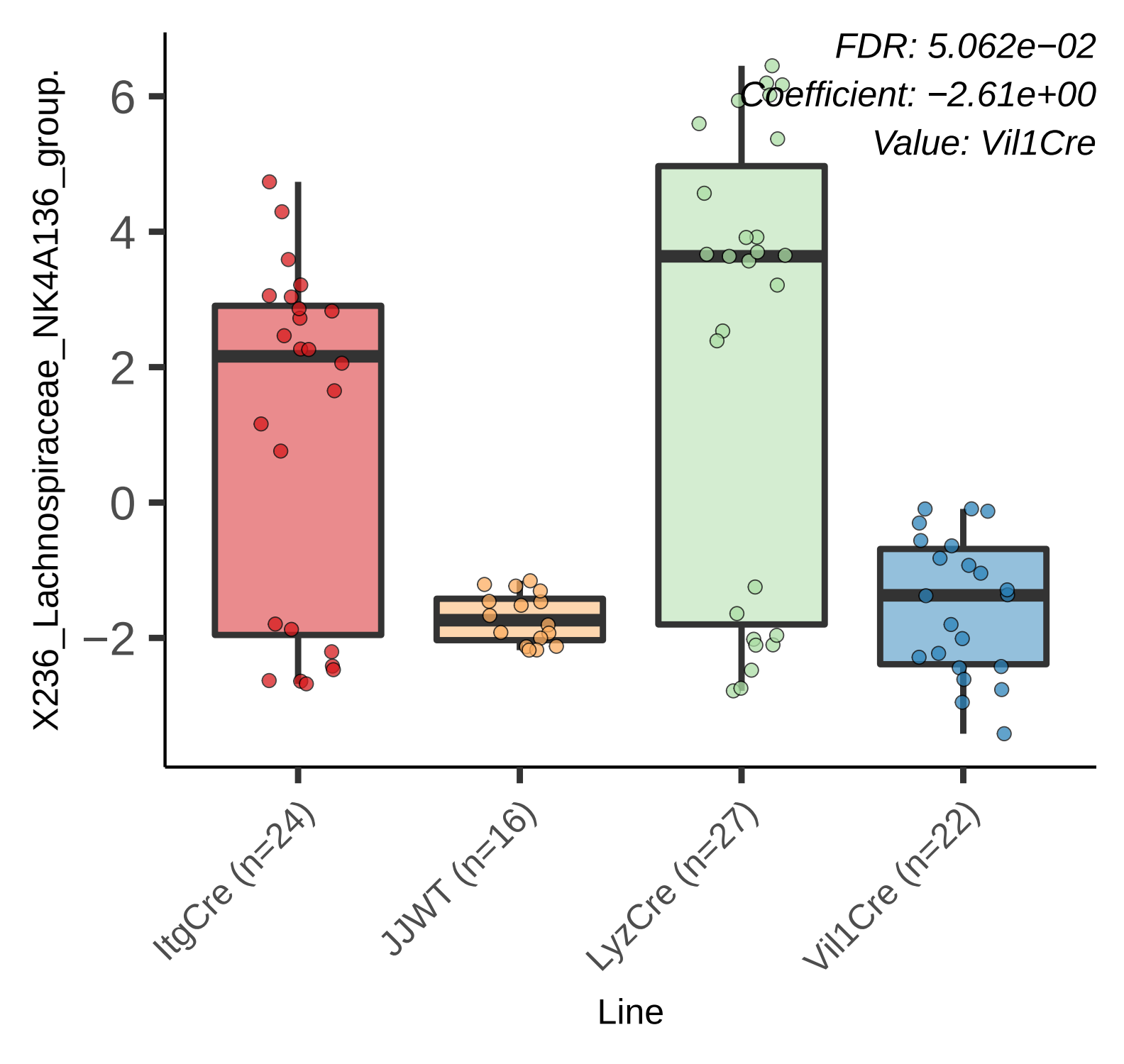

FDR: 4.827e−02
Coefficient: 1.64e+00
Value: JJWT
X65_Lachnospiraceae..f.
5
4
3
2
1
0
−1
ItgCre (n=24)
JJWT (n=16)
LyzCre (n=27)
Vil1Cre (n=22)
Line

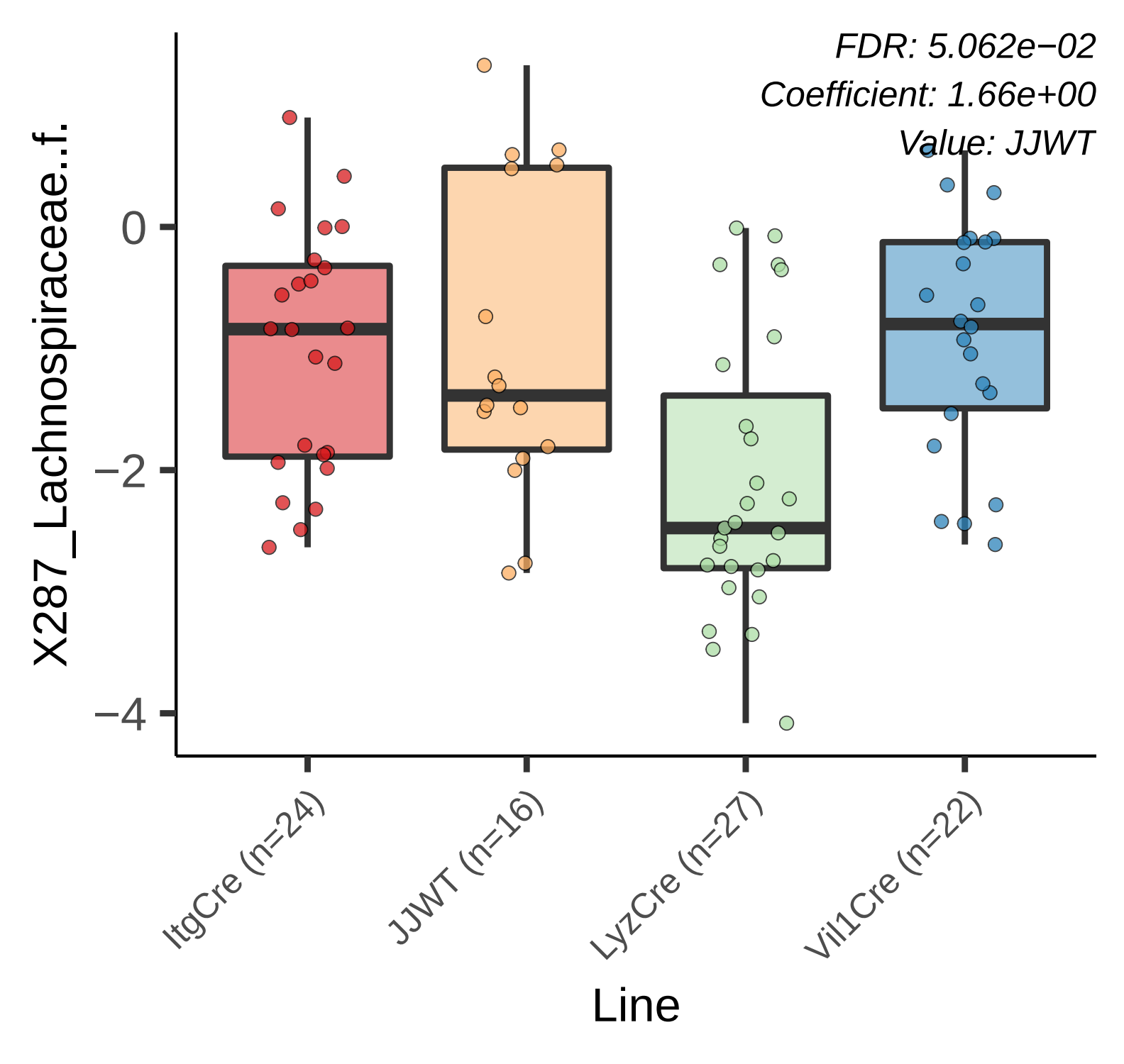

FDR: 4.934e−02
Coefficient: 1.90e+00
Value: JJWT
X216_Ruminococcaceae_UCG.014.
2
1
0
−1
−2
−3
ItgCre (n=24)
JJWT (n=16)
LyzCre (n=27)
Vil1Cre (n=22)
Line

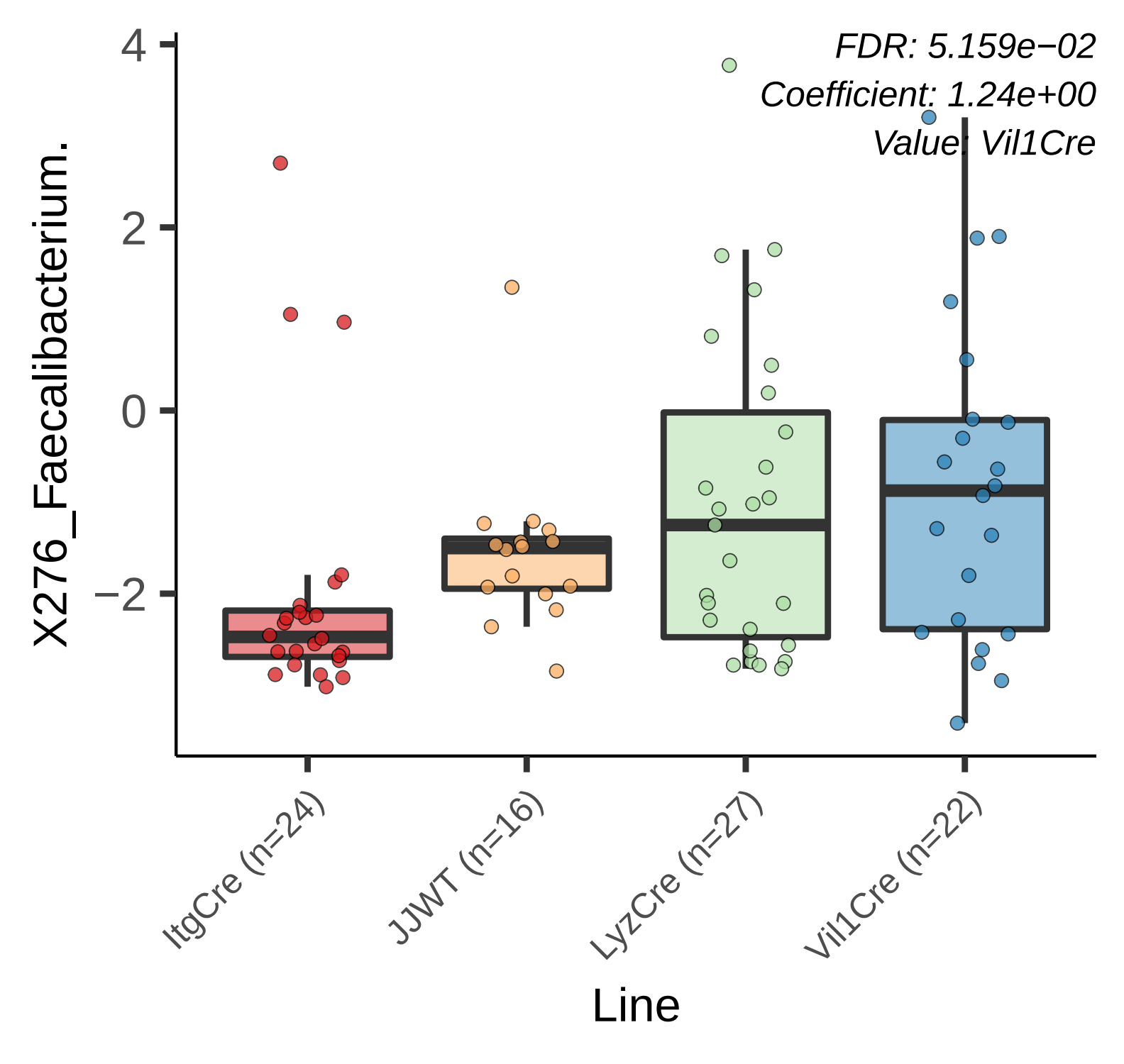

FDR: 4.966e−02
Coefficient: 2.02e+00
Value: Vil1Cre
X153_Lachnospiraceae_NK4A136_group.
2
0
−2
ItgCre (n=24)
JJWT (n=16)
LyzCre (n=27)
Vil1Cre (n=22)
Line

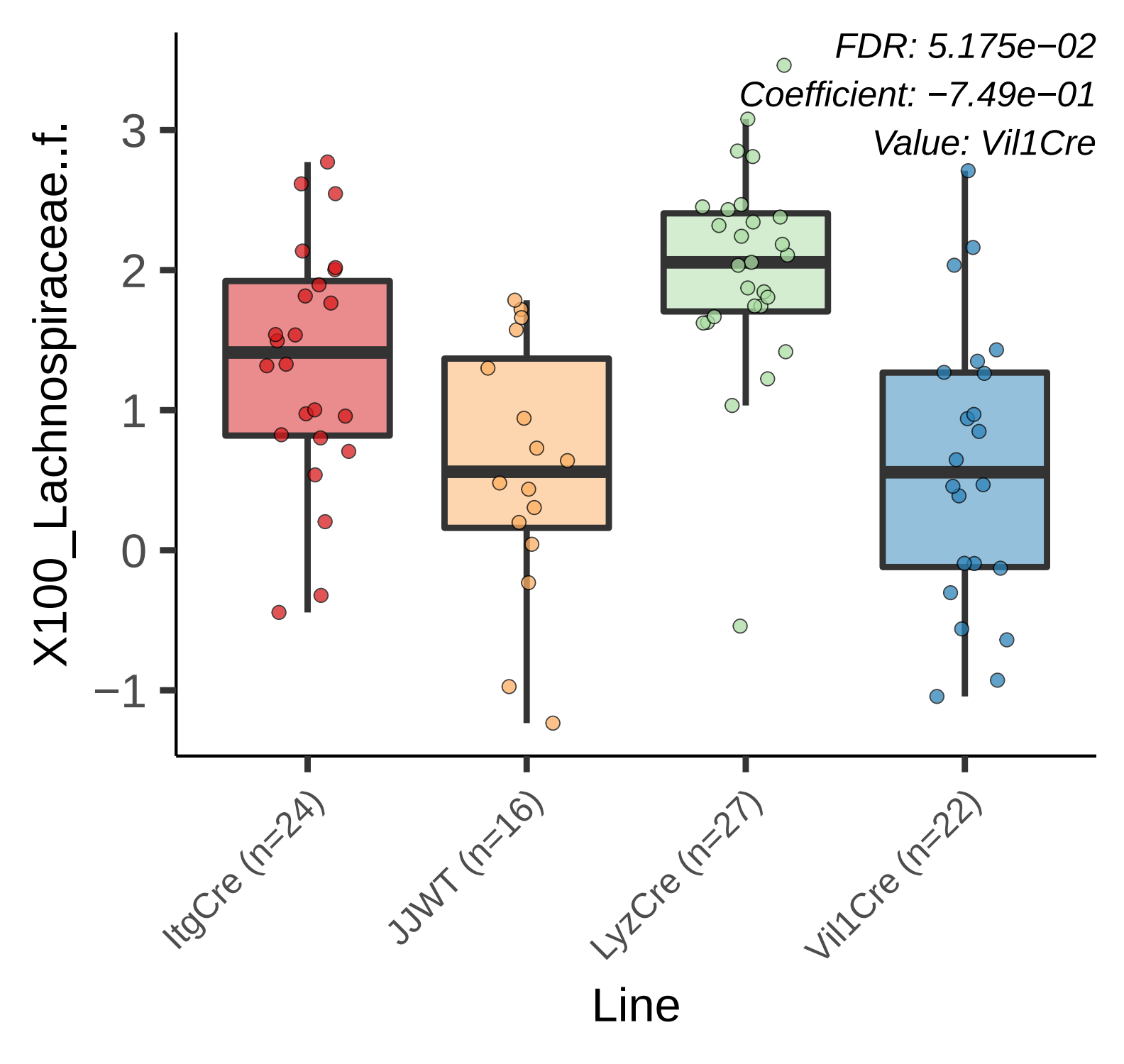

FDR: 5.002e−02
Coefficient: 2.60e+00
Value: JJWT
X179_Bacteroides.
5.0
2.5
0.0
−2.5
ItgCre (n=24)
JJWT (n=16)
LyzCre (n=27)
Vil1Cre (n=22)
Line

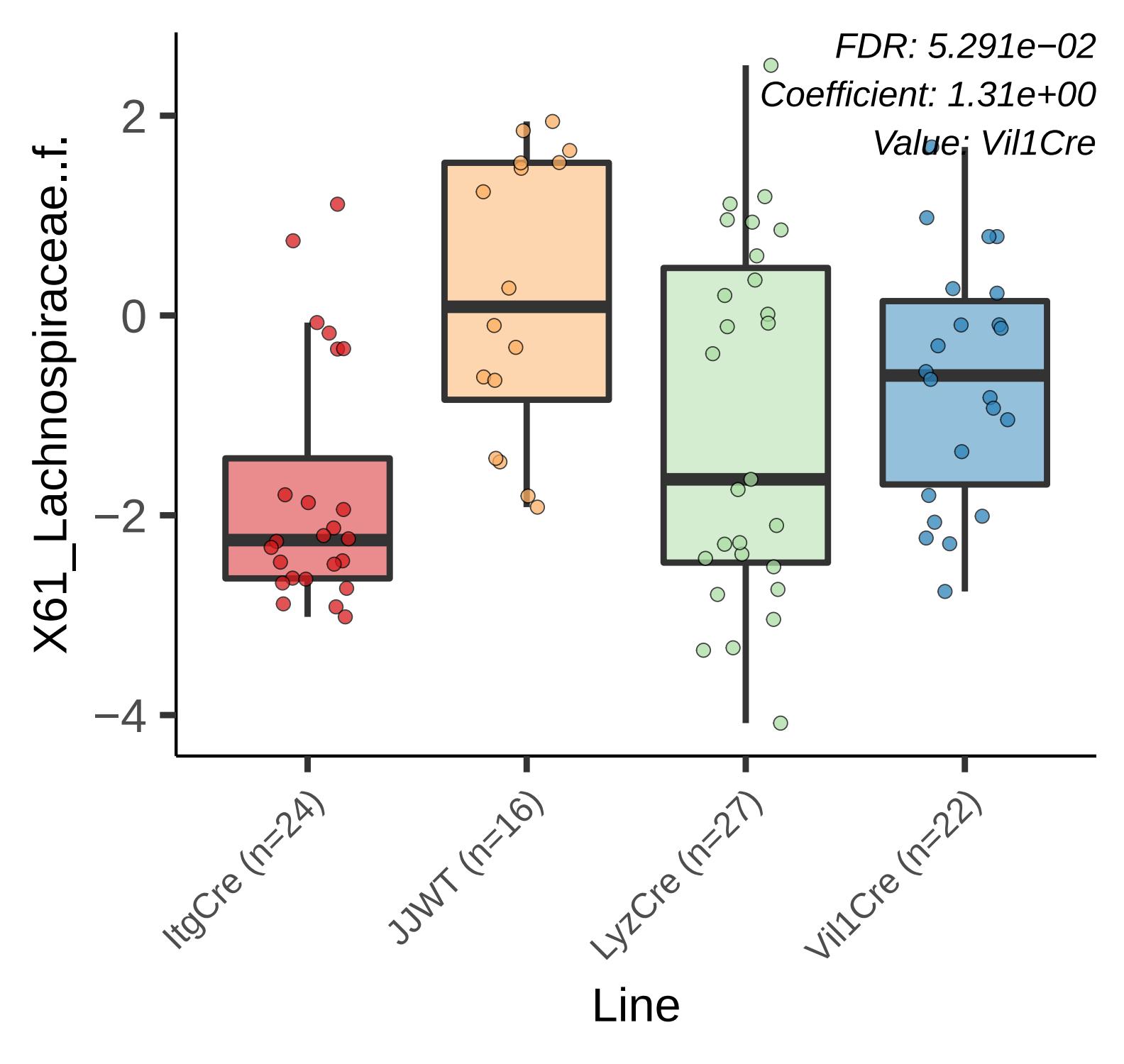

FDR: 5.023e−02
Coefficient: −1.51e+00
Value: Vil1Cre
X151_Butyricicoccus.
2
0
−2
ItgCre (n=24)
JJWT (n=16)
LyzCre (n=27)
Vil1Cre (n=22)
Line

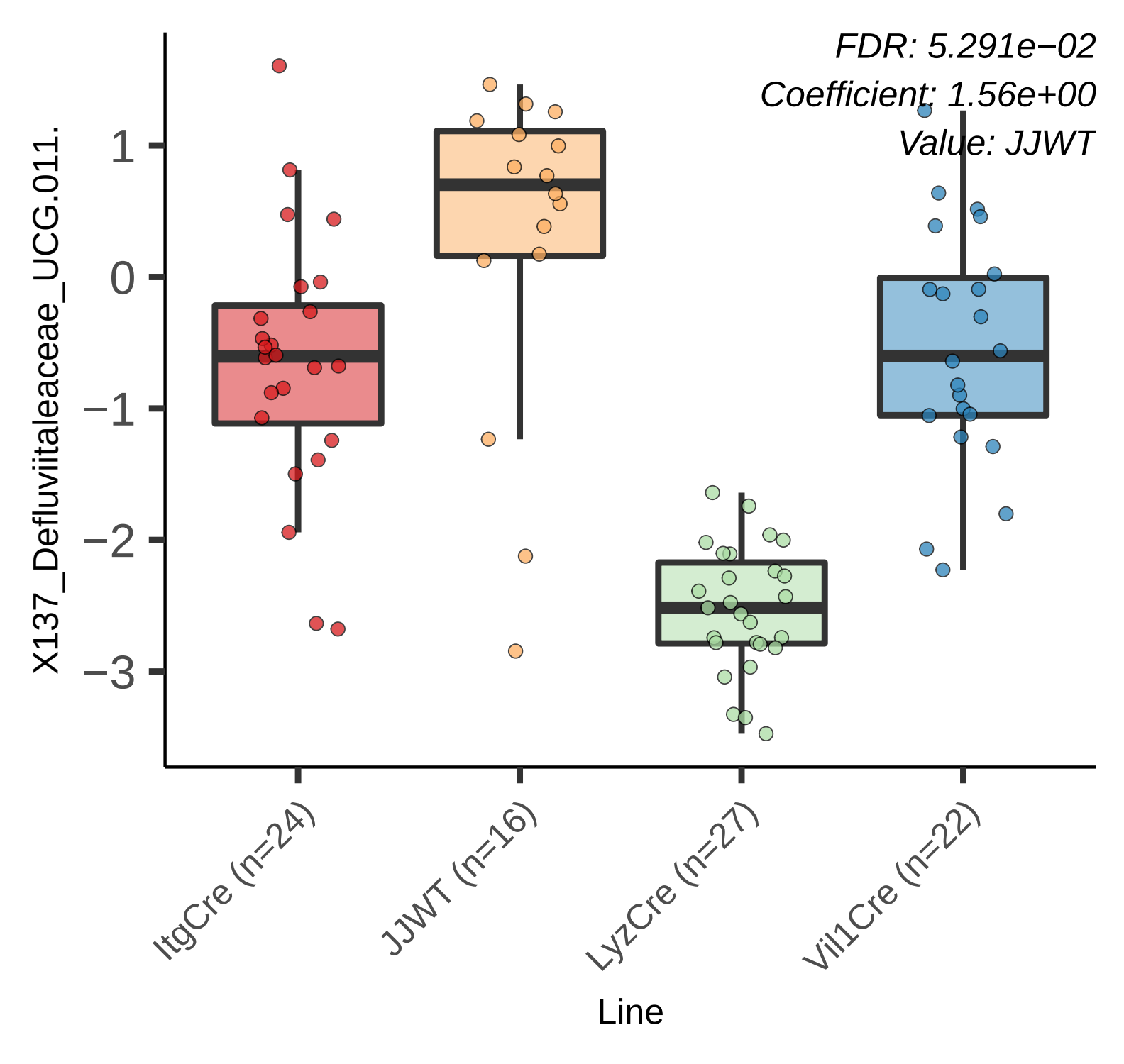

FDR: 5.062e−02
Coefficient: −2.61e+00
Value: Vil1Cre
X236_Lachnospiraceae_NK4A136_group.
6
4
2
0
−2
ItgCre (n=24)
JJWT (n=16)
LyzCre (n=27)
Vil1Cre (n=22)
Line

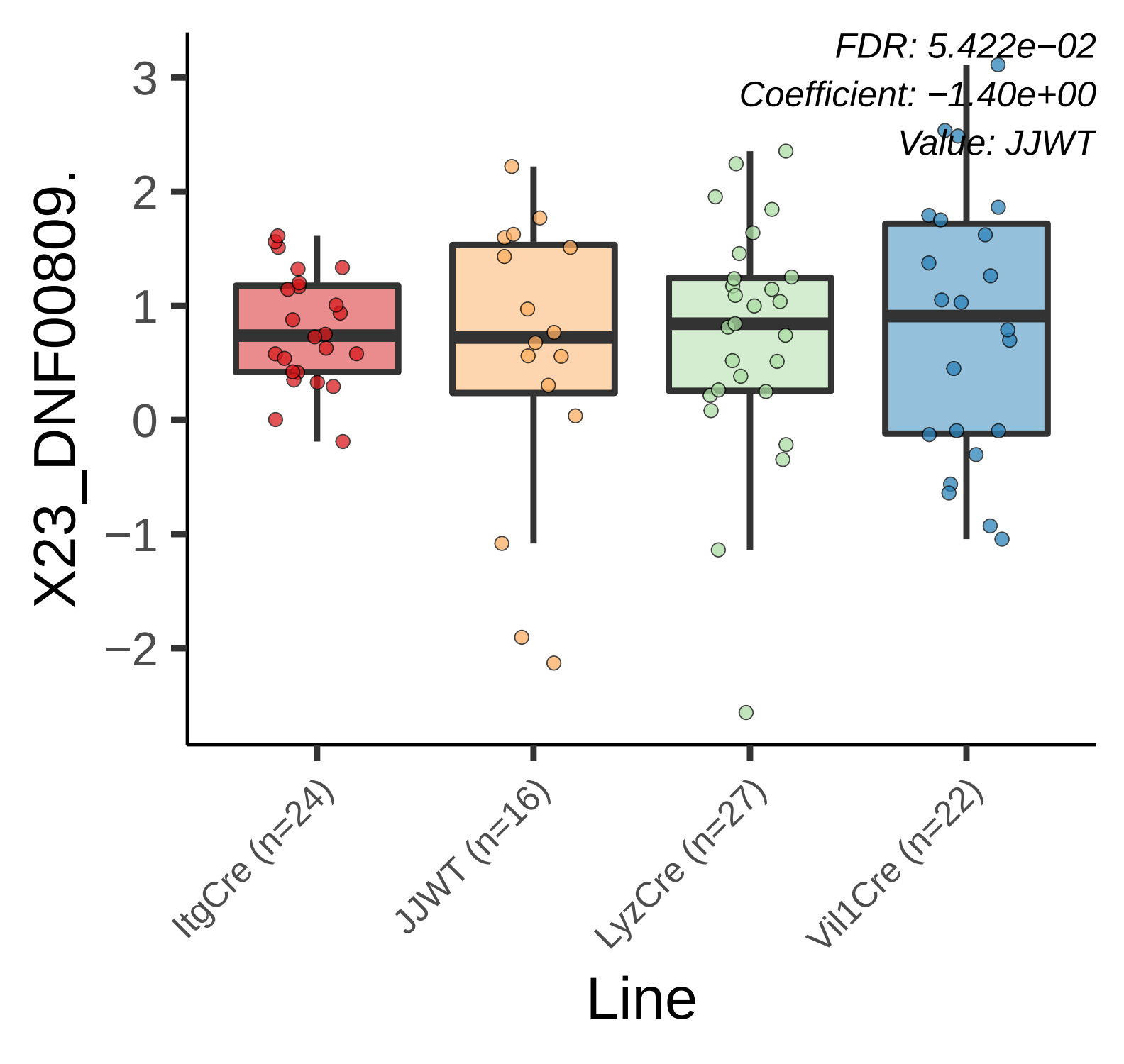

FDR: 5.062e−02
Coefficient: 1.66e+00
Value: JJWT
X287_Lachnospiraceae.f.
0
−2
−4
ItgCre (n=24)
JJWT (n=16)
LyzCre (n=27)
Vil1Cre (n=22)
Line

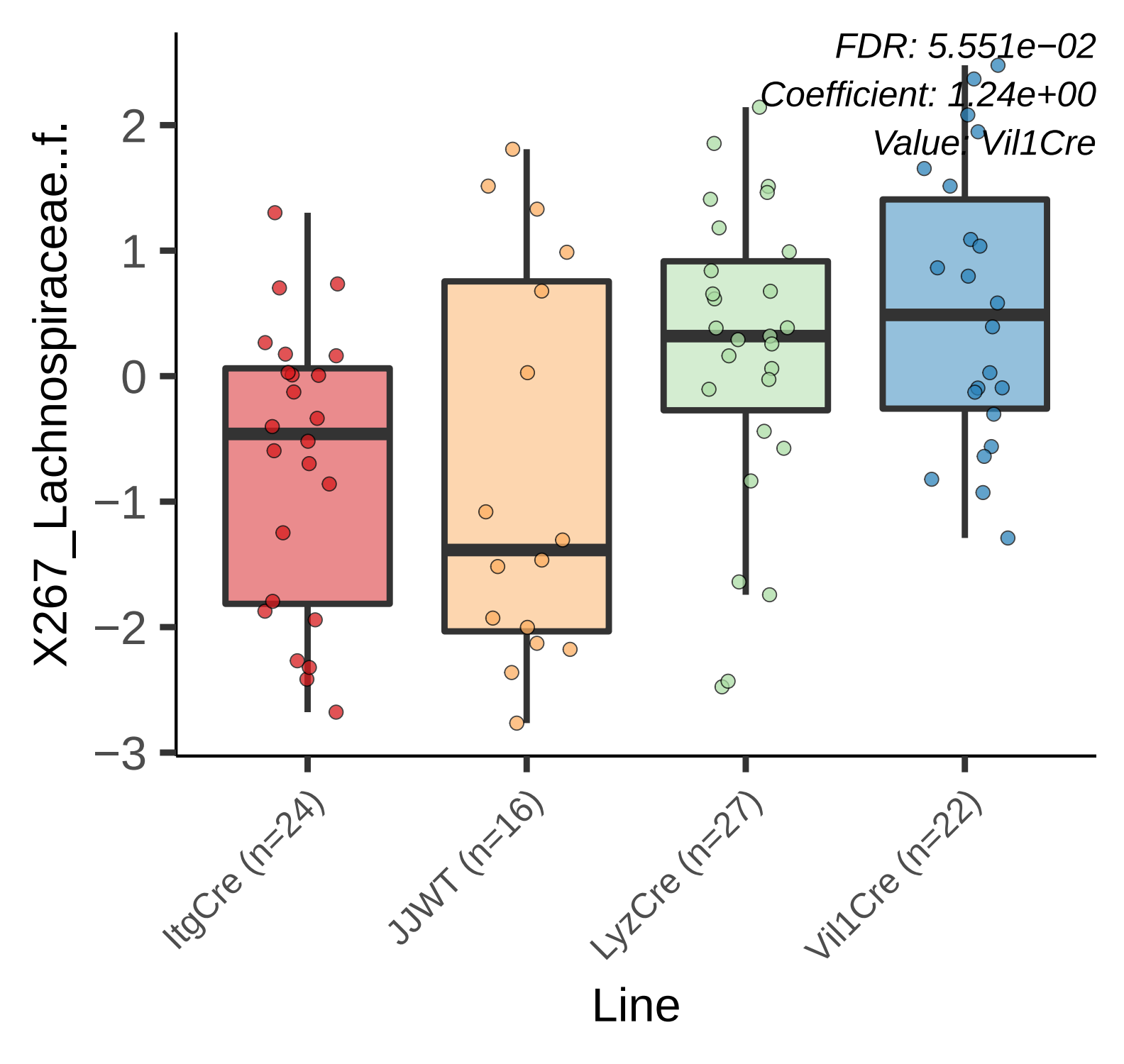

FDR: 5.159e−02
Coefficient: 1.24e+00
Value: Vil1Cre
X276_Faecalibacterium.
4
2
0
−2
ItgCre (n=24)
JJWT (n=16)
LyzCre (n=27)
Vil1Cre (n=22)
Line

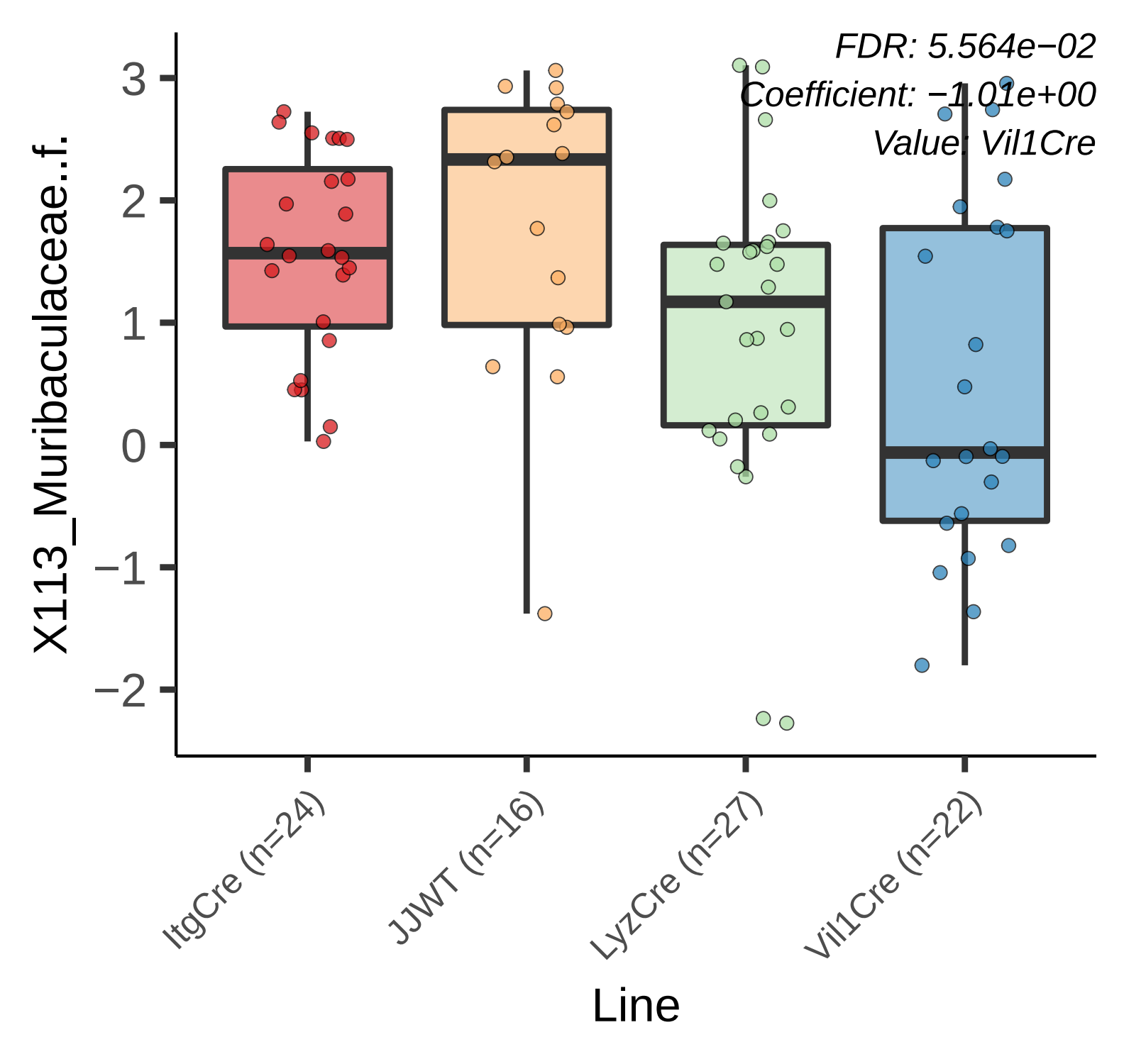

FDR: 5.175e−02
Coefficient: −7.49e−01
Value: Vil1Cre
X100_Lachnospiraceae.f.
3
2
1
0
−1
ItgCre (n=24)
JJWT (n=16)
LyzCre (n=27)
Vil1Cre (n=22)
Line

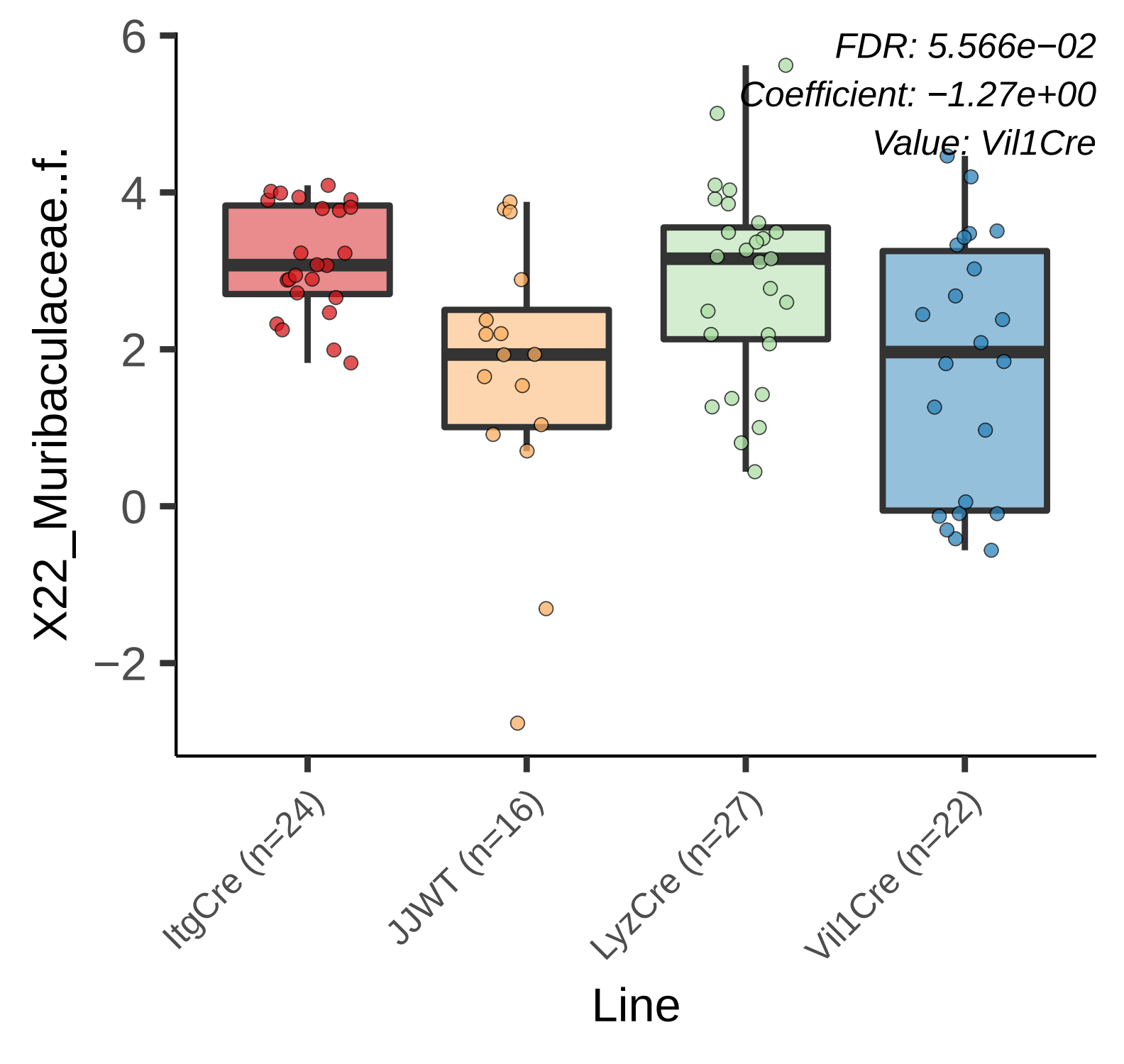

FDR: 5.291e−02
Coefficient: 1.31e+00
Value: Vil1Cre
X61_Lachnospiraceae..f.
2
0
−2
−4
ItgCre (n=24)
JJWT (n=16)
LyzCre (n=27)
Vil1Cre (n=22)
Line

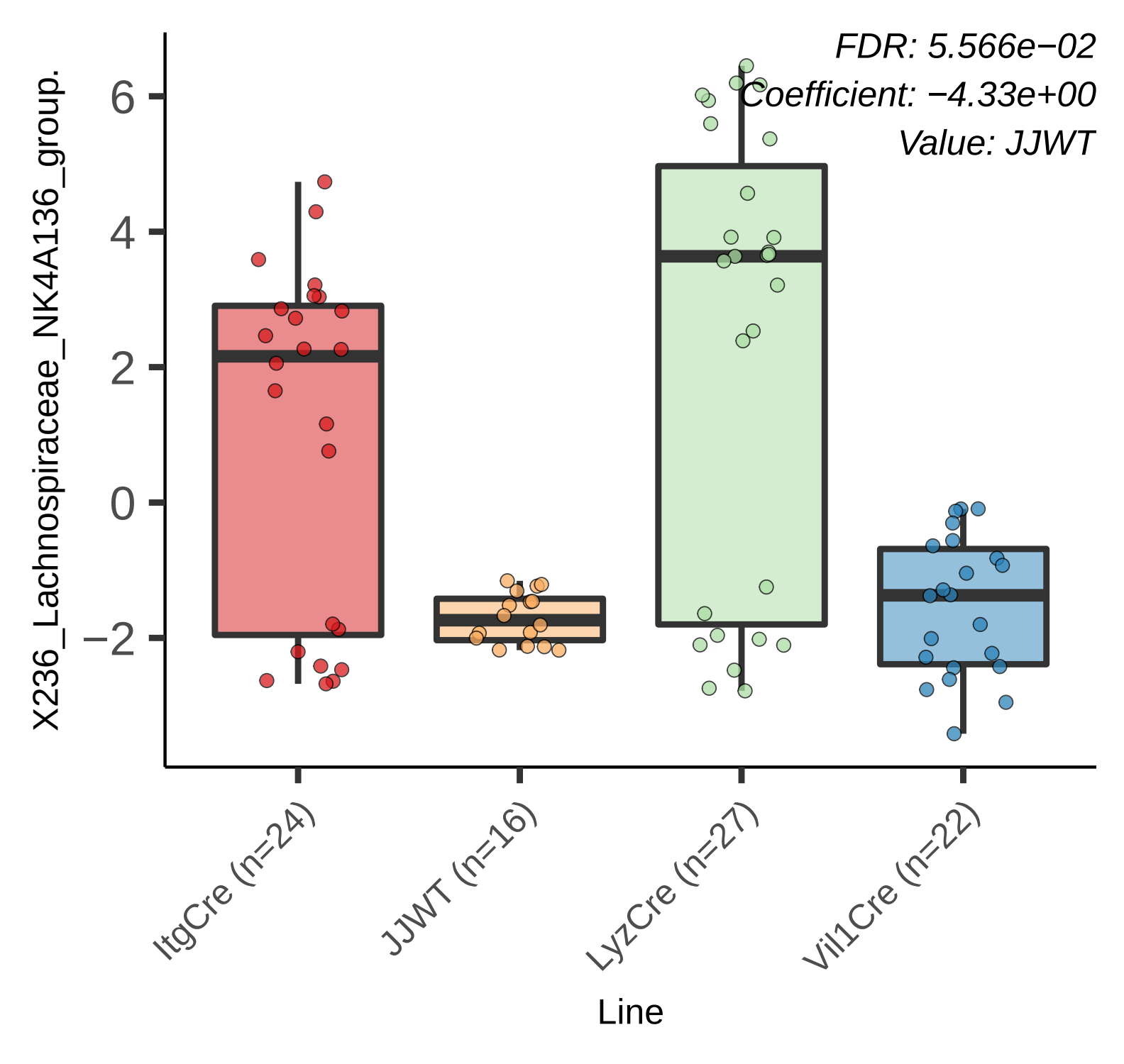

FDR: 5.291e−02
Coefficient: 1.56e+00
Value: JJWT
X137_Defluviitaleaceae_UCG.011.
1
0
−1
−2
−3
ItgCre (n=24)
JJWT (n=16)
LyzCre (n=27)
Vil1Cre (n=22)
Line

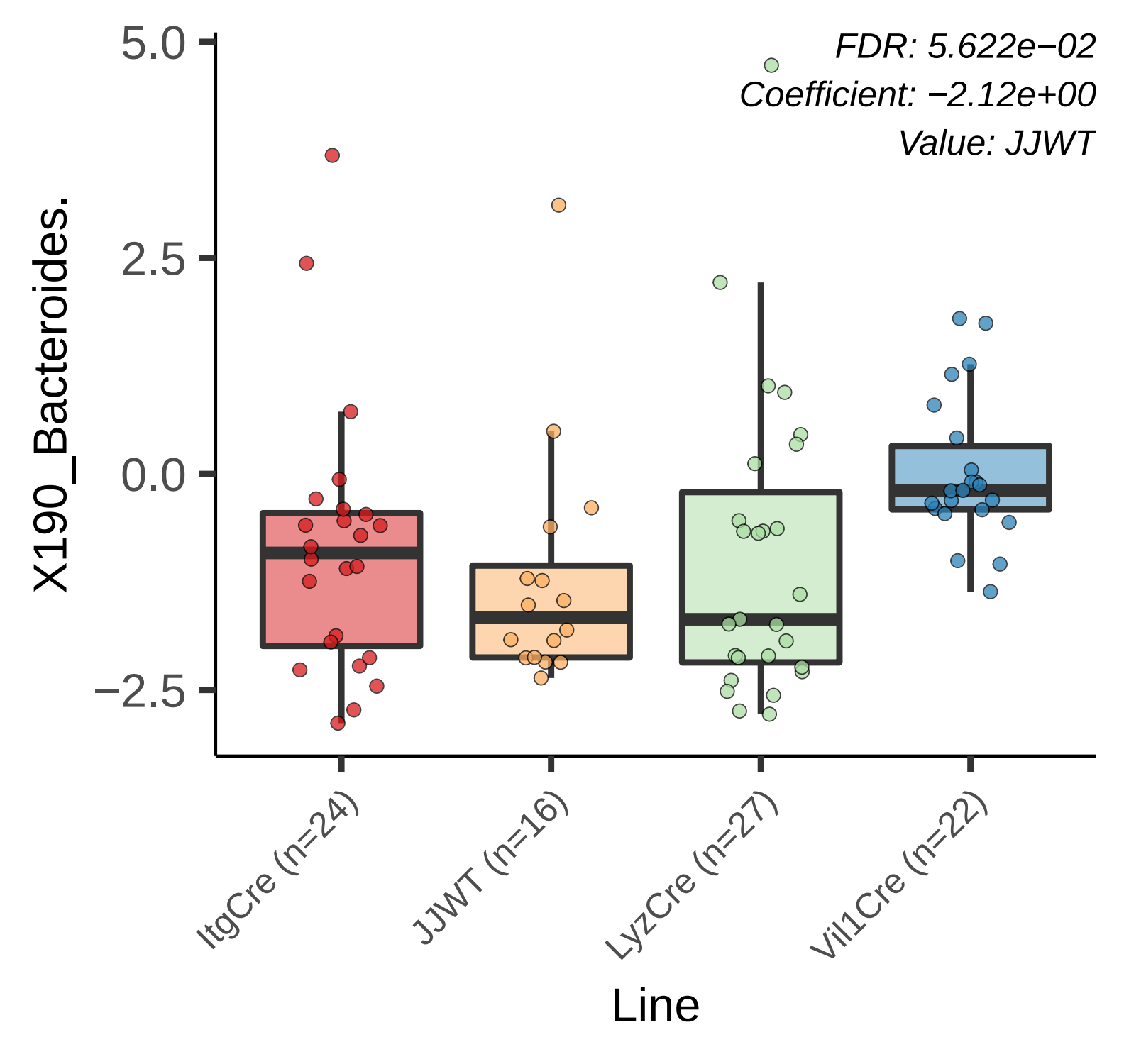

FDR: 5.422e−02
Coefficient: −1.40e+00
Value: JJWT
X23_DNF00809.
3
2
1
0
−1
−2
ItgCre (n=24)
JJWT (n=16)
LyzCre (n=27)
Vil1Cre (n=22)
Line

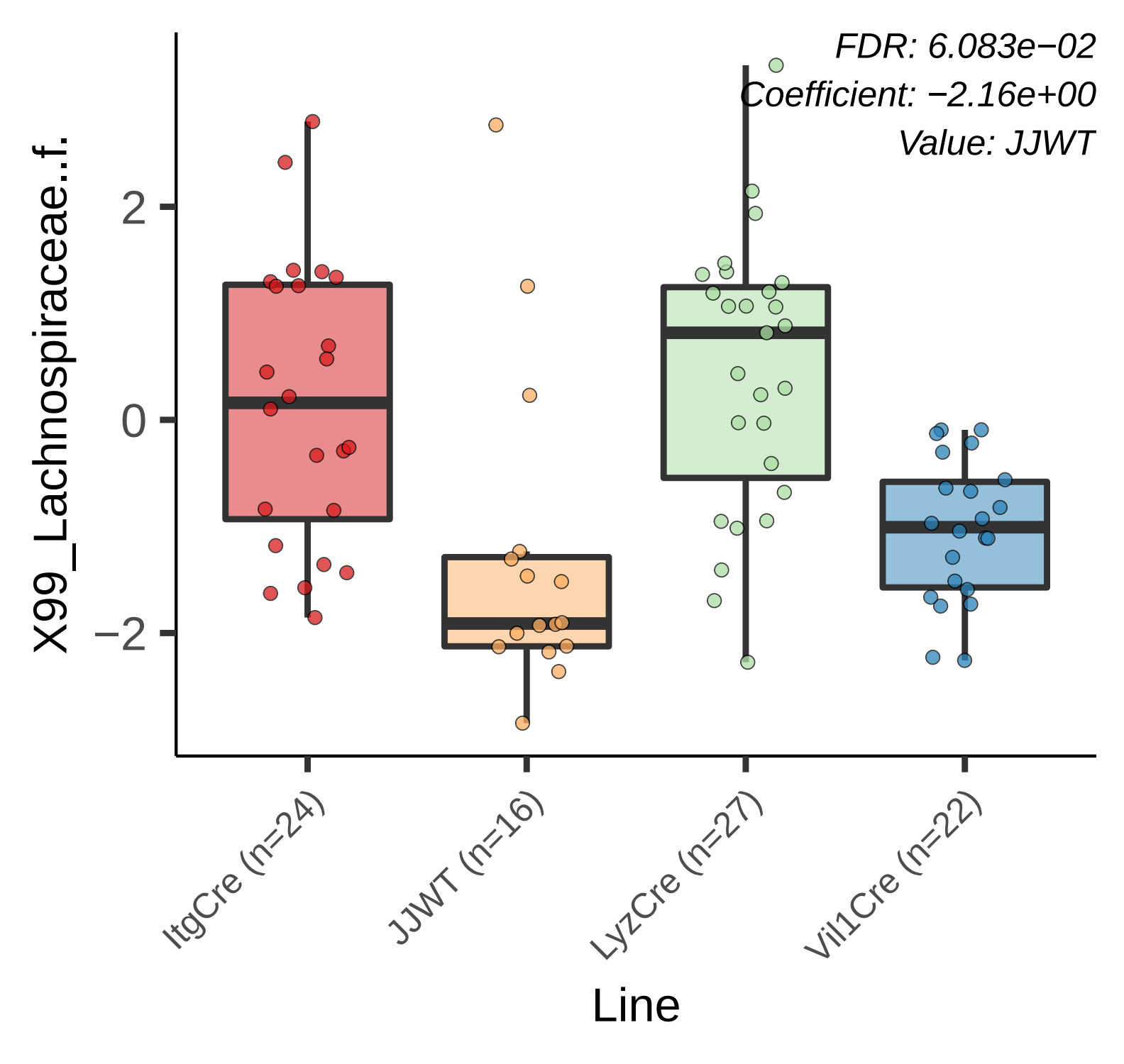

FDR: 5.551e−02
Coefficient: 1.24e+00
Value: Vil1Cre
X267_Lachnospiraceae.f.
2
1
0
−1
−2
−3
ItgCre (n=24)
JJWT (n=16)
LyzCre (n=27)
Vil1Cre (n=22)
Line

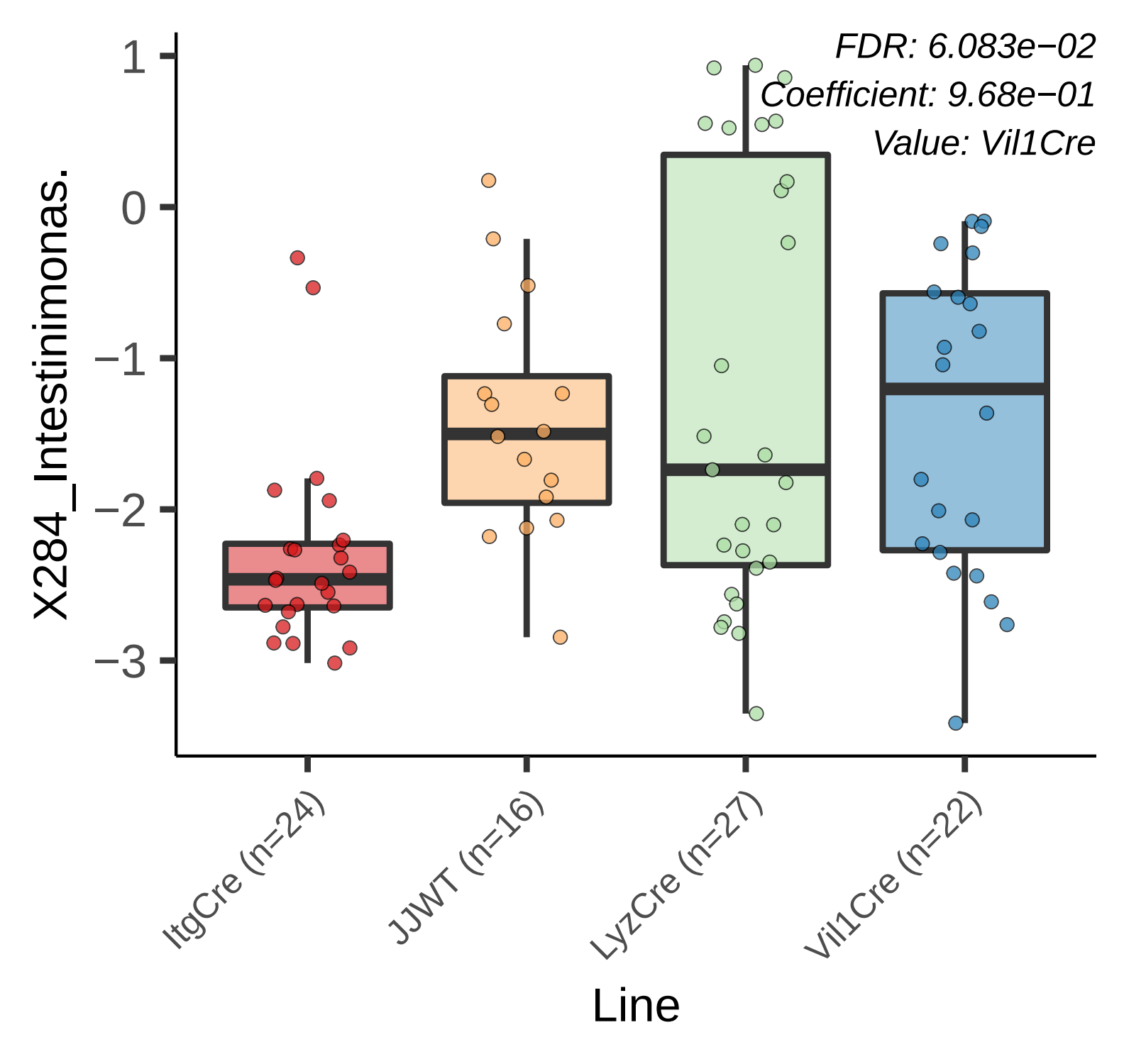

FDR: 5.564e−02
Coefficient: −1.01e+00
Value: Vil1Cre
X113_Muribaculaceae..f.
3
2
1
0
−1
−2
ItgCre (n=24)
JJWT (n=16)
LyzCre (n=27)
Vil1Cre (n=22)
Line

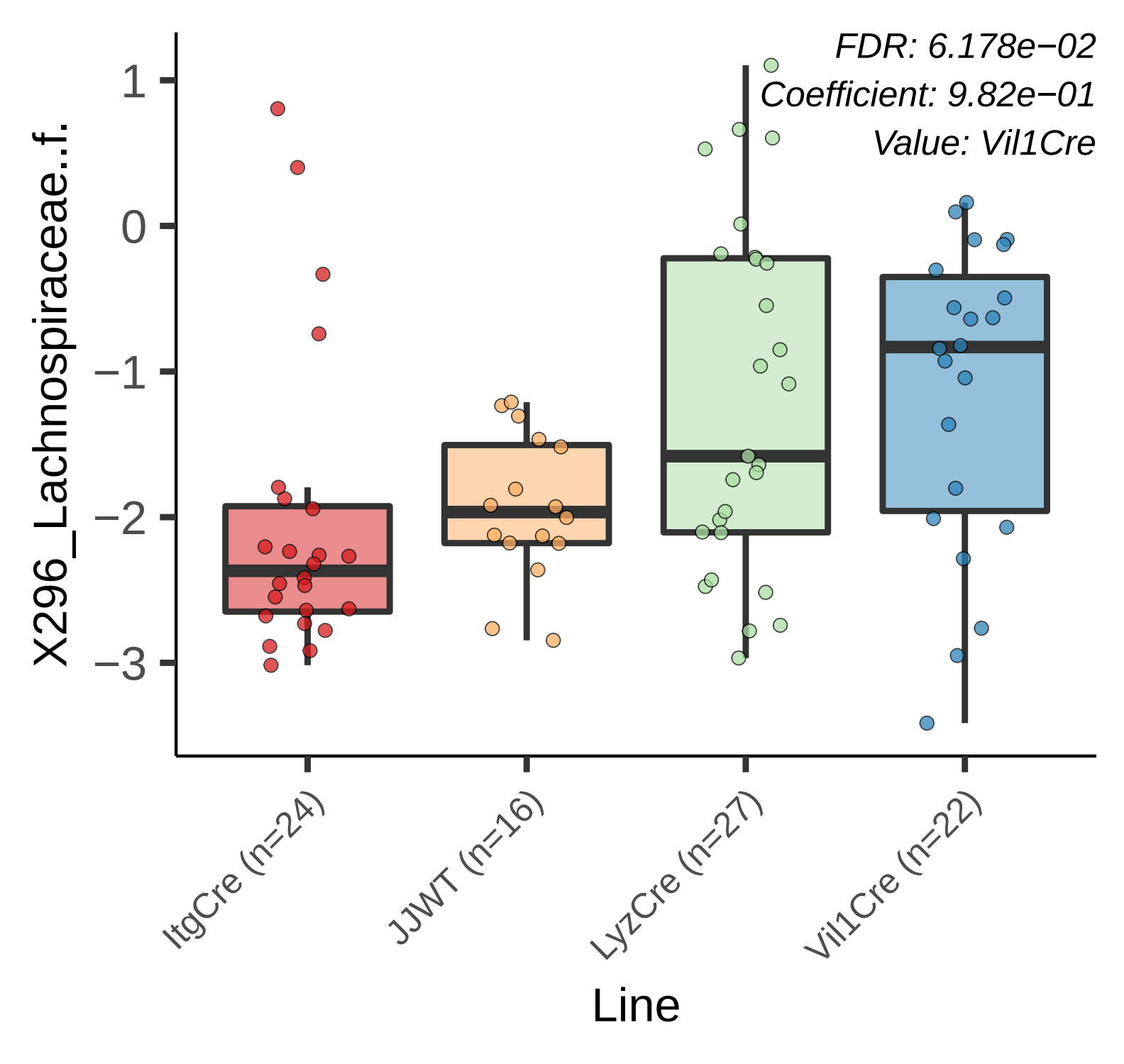

FDR: 5.566e−02
Coefficient: −1.27e+00
Value: Vil1Cre
X22_Muribaculaceae..f.
6
4
2
0
−2
ItgCre (n=24)
JJWT (n=16)
LyzCre (n=27)
Vil1Cre (n=22)
Line

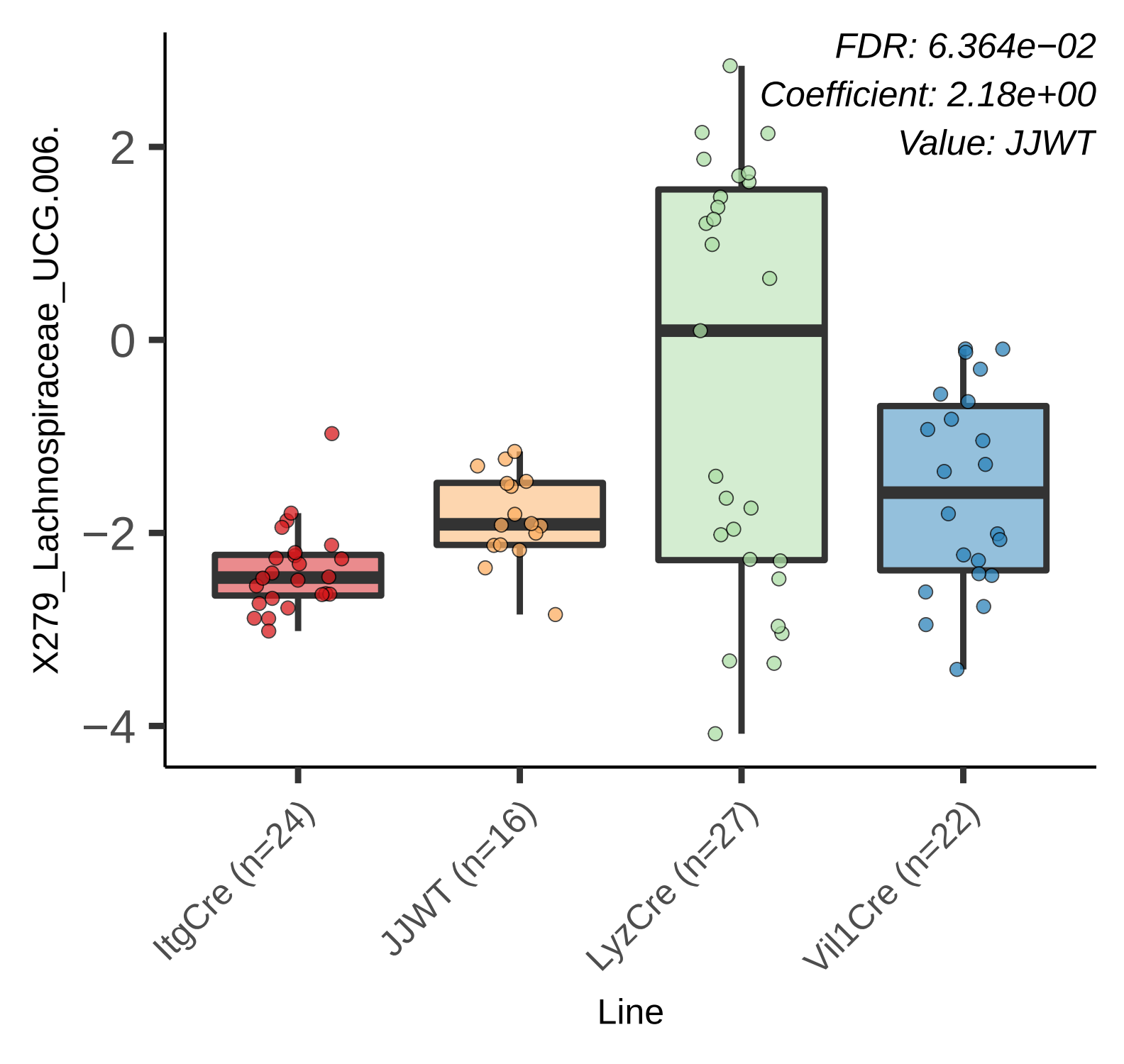

FDR: 5.566e−02
Coefficient: −4.33e+00
Value: JJWT
X236_Lachnospiraceae_NK4A136_group.
6
4
2
0
−2
ItgCre (n=24)
JJWT (n=16)
LyzCre (n=27)
Vil1Cre (n=22)
Line

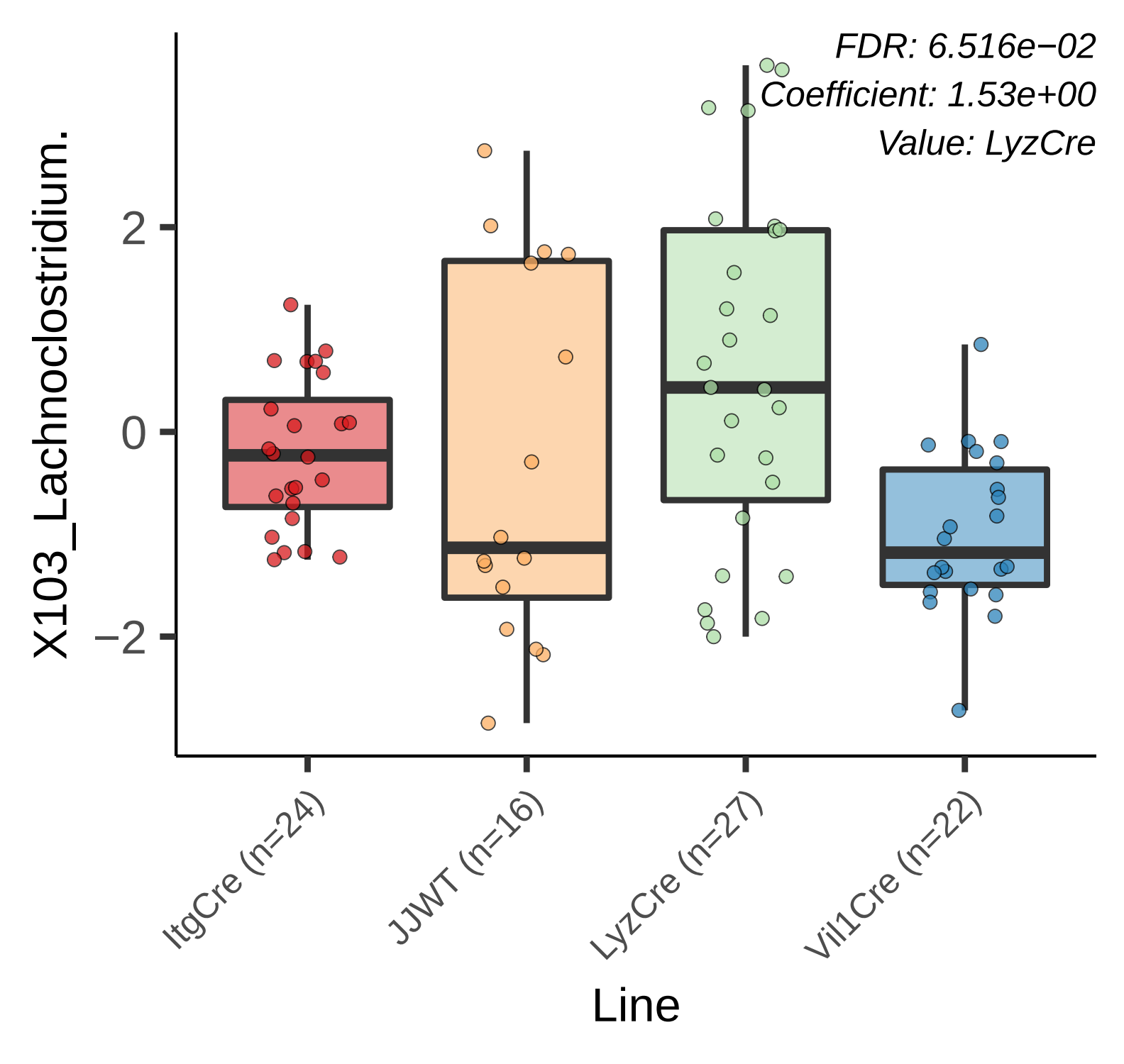

FDR: 5.622e−02
Coefficient: −2.12e+00
Value: JJWT
X190_Bacteroides.
5.0
2.5
0.0
−2.5
ItgCre (n=24)
JJWT (n=16)
LyzCre (n=27)
Vil1Cre (n=22)
Line

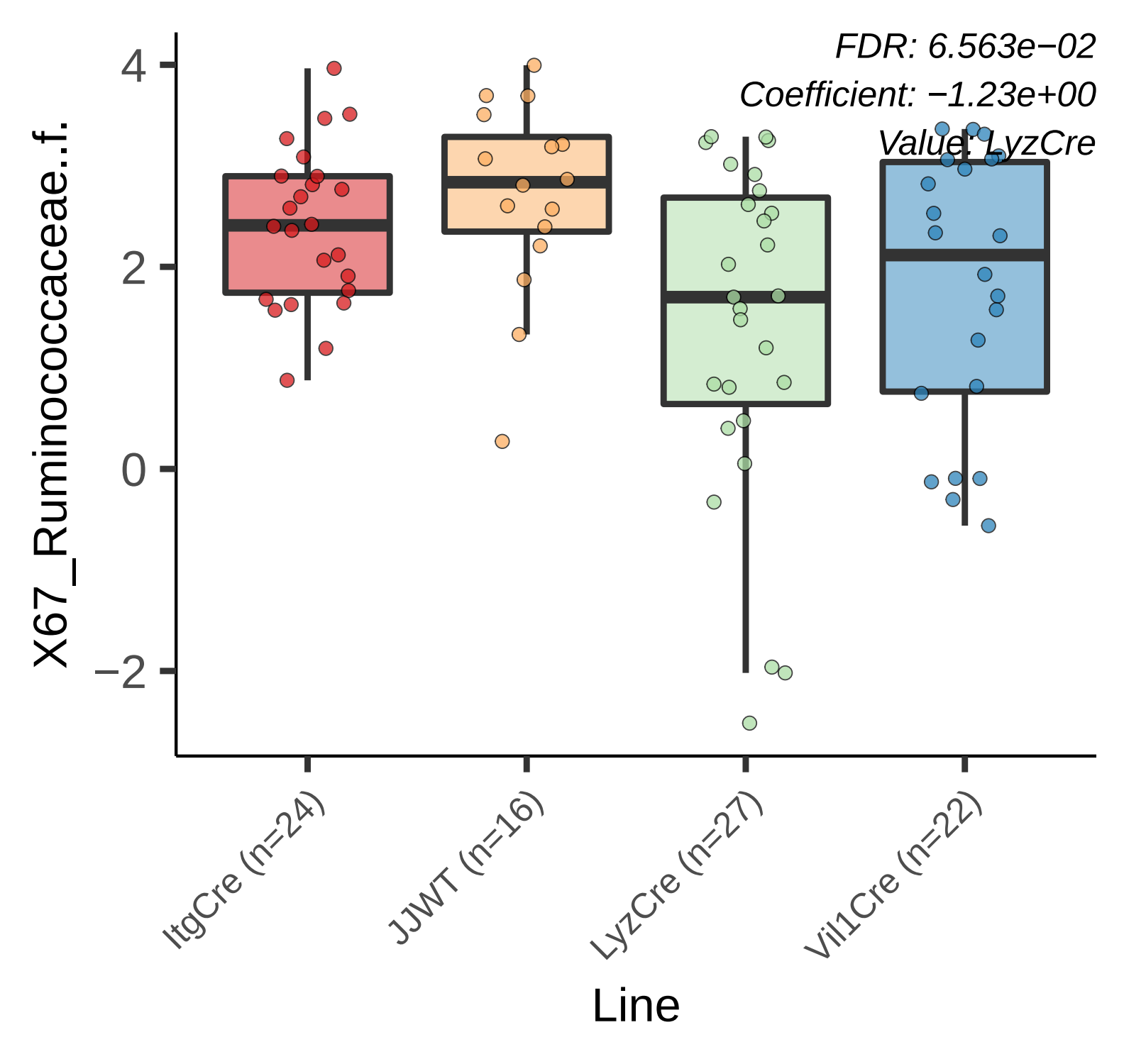

FDR: 6.083e−02
Coefficient: −2.16e+00
Value: JJWT
X99_Lachnospiraceae..f.
2
0
−2
ItgCre (n=24)
JJWT (n=16)
LyzCre (n=27)
Vil1Cre (n=22)
Line

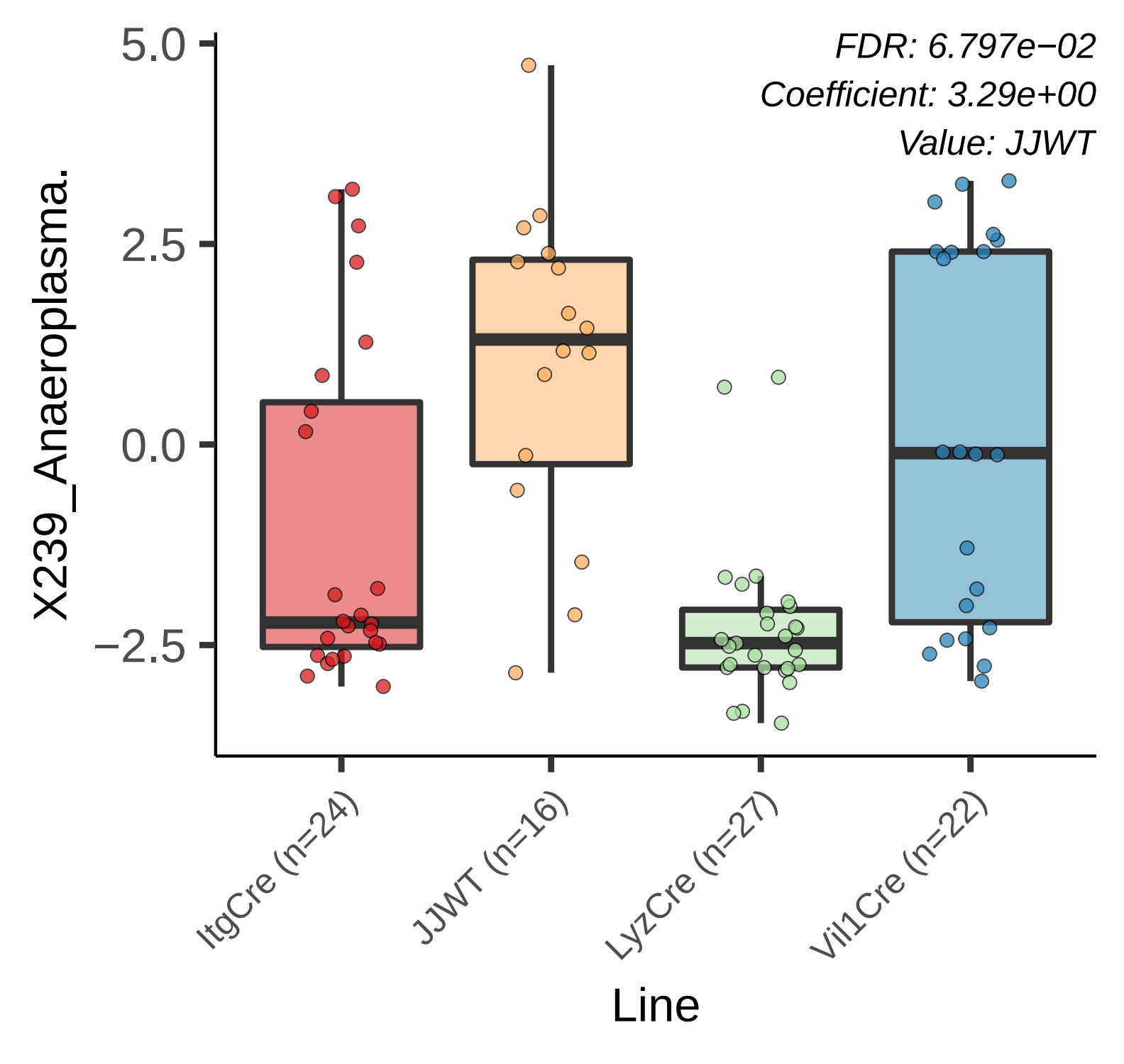

FDR: 6.083e−02
Coefficient: 9.68e−01
Value: Vil1Cre
X284_Intestinimonas.
1
0
−1
−2
−3
ItgCre (n=24)
JJWT (n=16)
LyzCre (n=27)
Vil1Cre (n=22)
Line

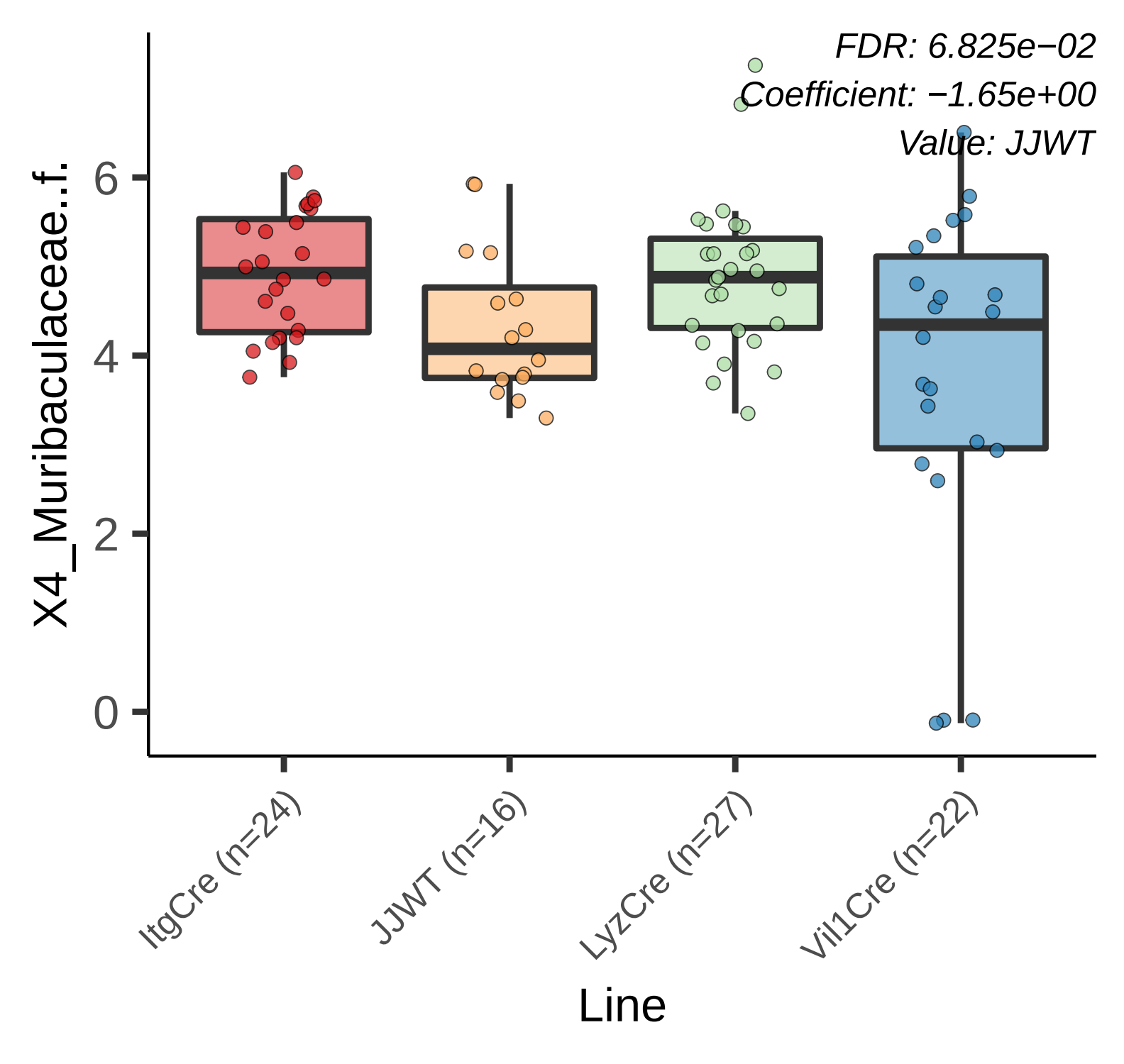

FDR: 6.178e−02
Coefficient: 9.82e−01
Value: Vil1Cre
X296_Lachnospiraceae.f.
1
0
−1
−2
−3
ItgCre (n=24)
JJWT (n=16)
LyzCre (n=27)
Vil1Cre (n=22)
Line

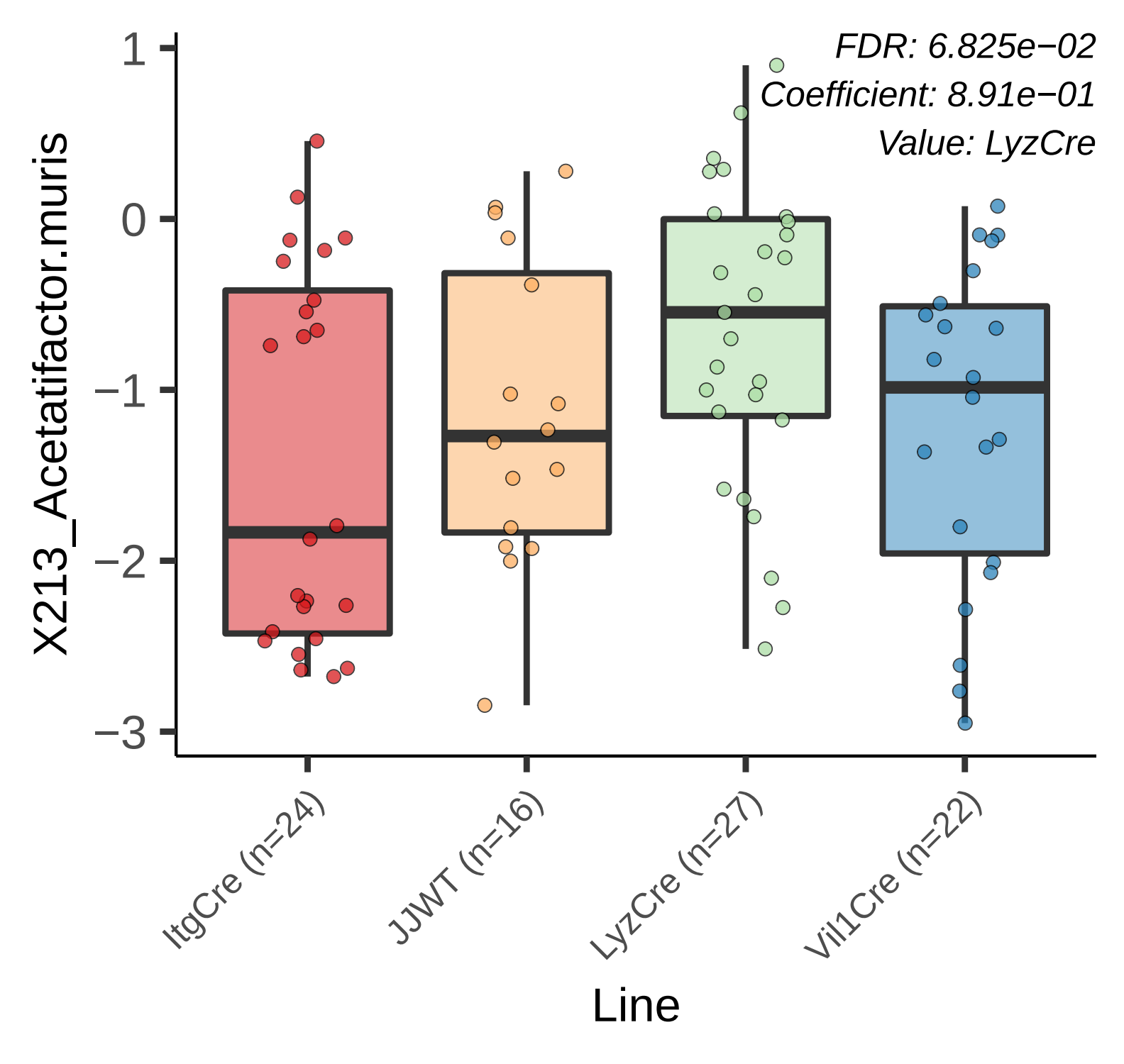

FDR: 6.364e−02
Coefficient: 2.18e+00
Value: JJWT
X279_Lachnospiraceae_UCG.006.
2
0
−2
−4
ItgCre (n=24)
JJWT (n=16)
LyzCre (n=27)
Vil1Cre (n=22)
Line

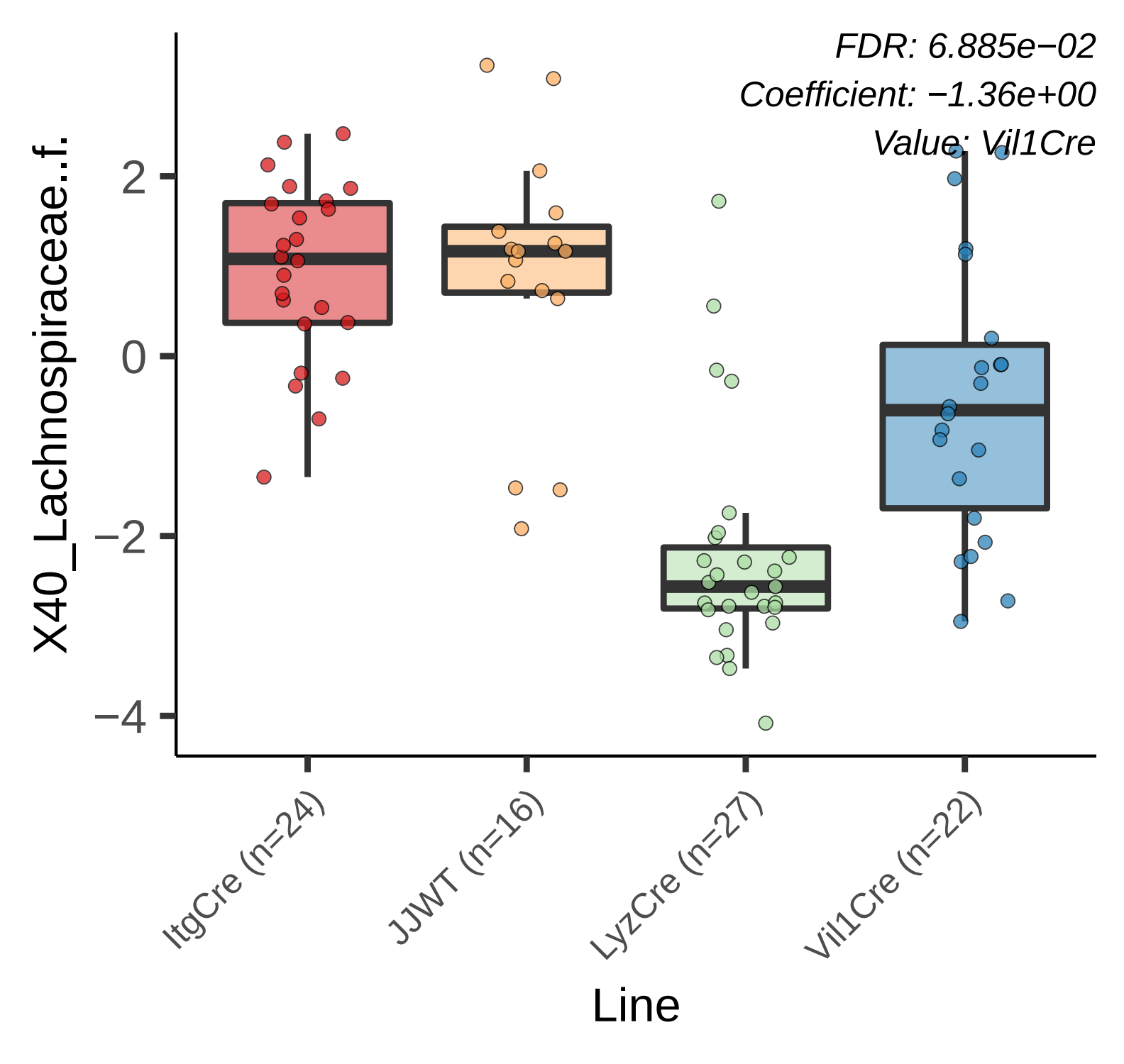

FDR: 6.516e−02
Coefficient: 1.53e+00
Value: LyzCre
X103_Lachnoclostridium.
2
0
−2
ItgCre (n=24)
JJWT (n=16)
LyzCre (n=27)
Vil1Cre (n=22)
Line

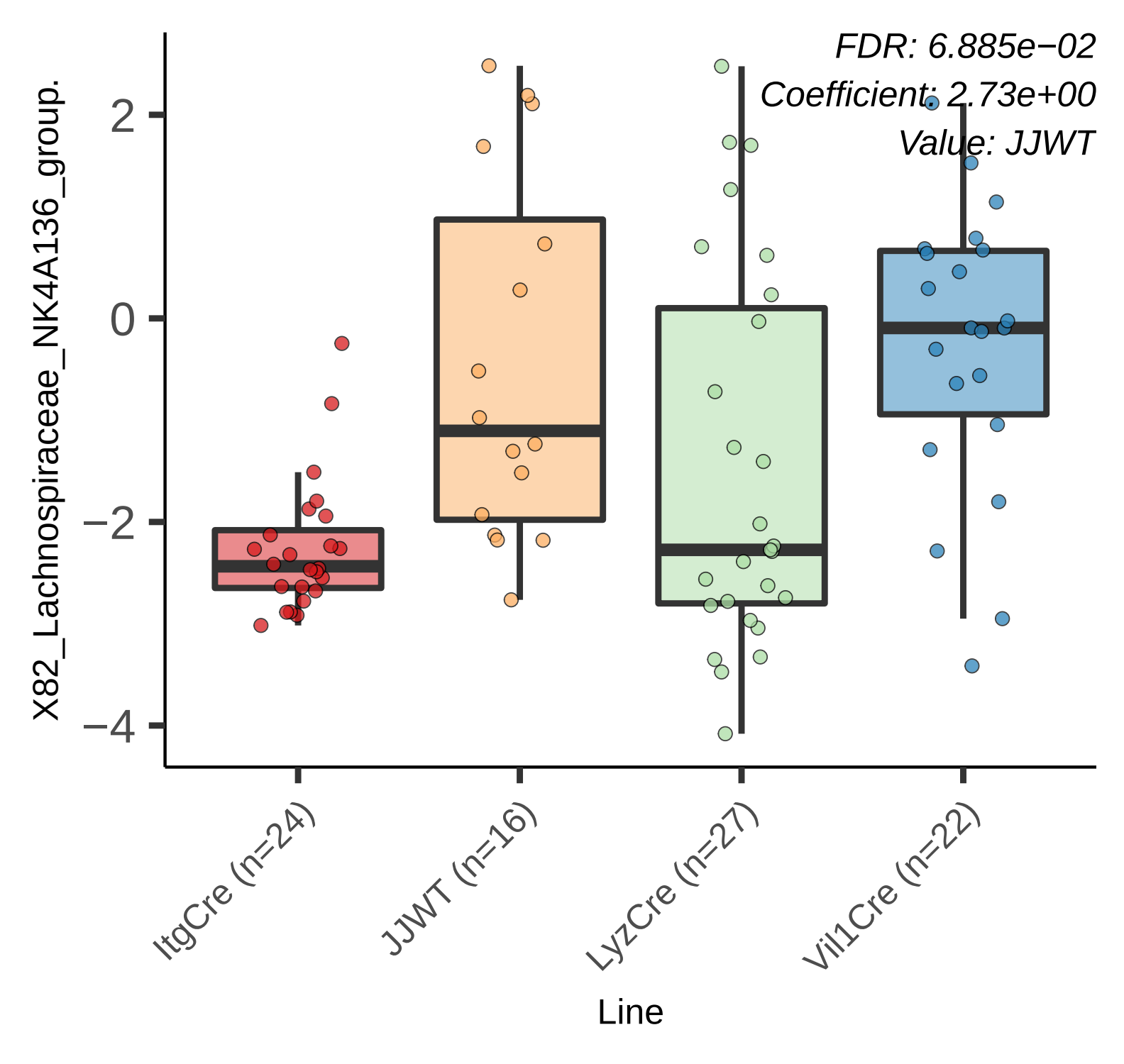

FDR: 6.563e−02
Coefficient: −1.23e+00
Value: LyzCre
X67_Ruminococcaceae..f.
4
2
0
−2
ItgCre (n=24)
JJWT (n=16)
LyzCre (n=27)
Vil1Cre (n=22)
Line

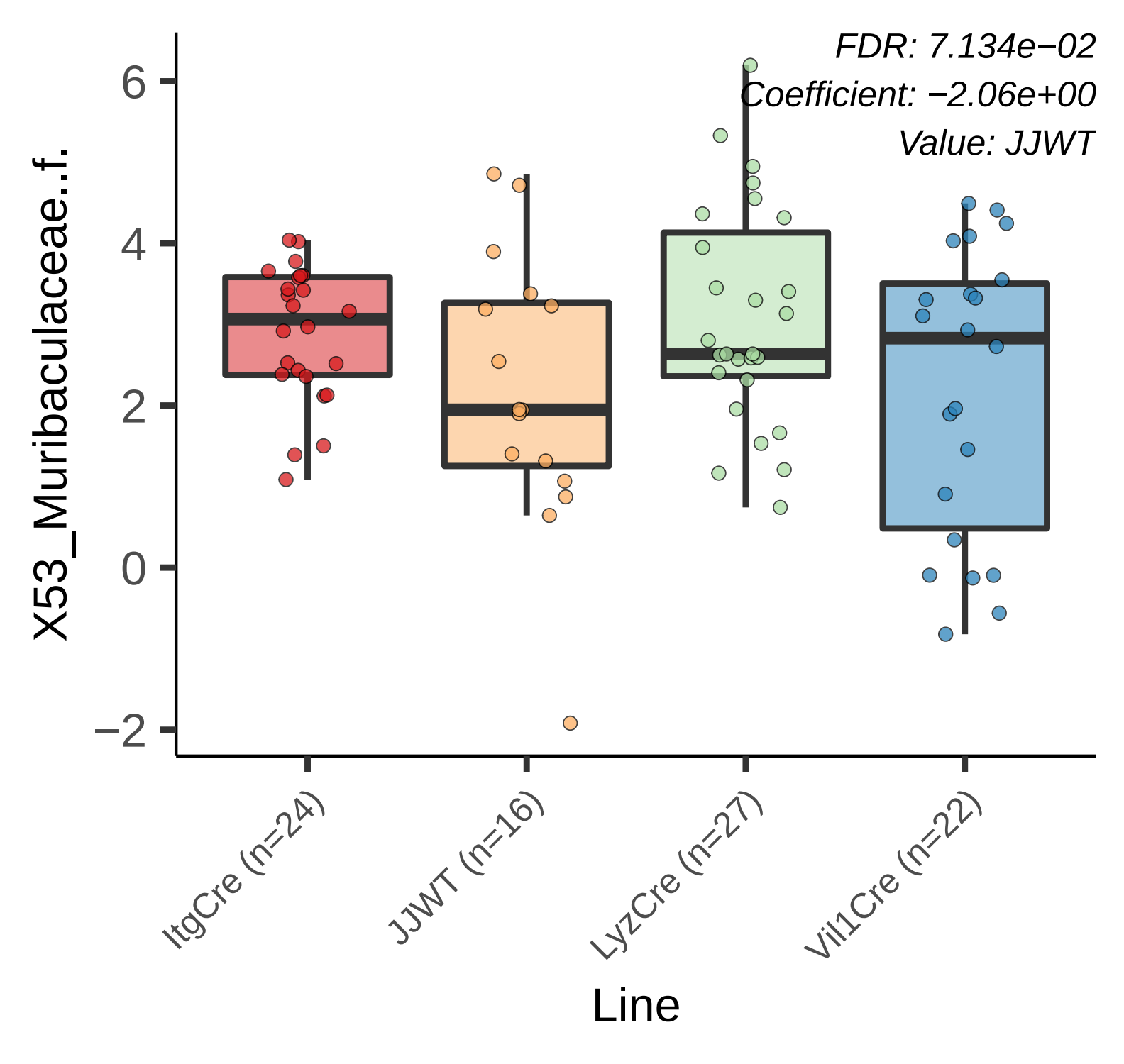

FDR: 6.797e−02
Coefficient: 3.29e+00
Value: JJWT
X239_Anaeroplasma.
5.0
2.5
0.0
−2.5
ItgCre (n=24)
JJWT (n=16)
LyzCre (n=27)
Vil1Cre (n=22)
Line

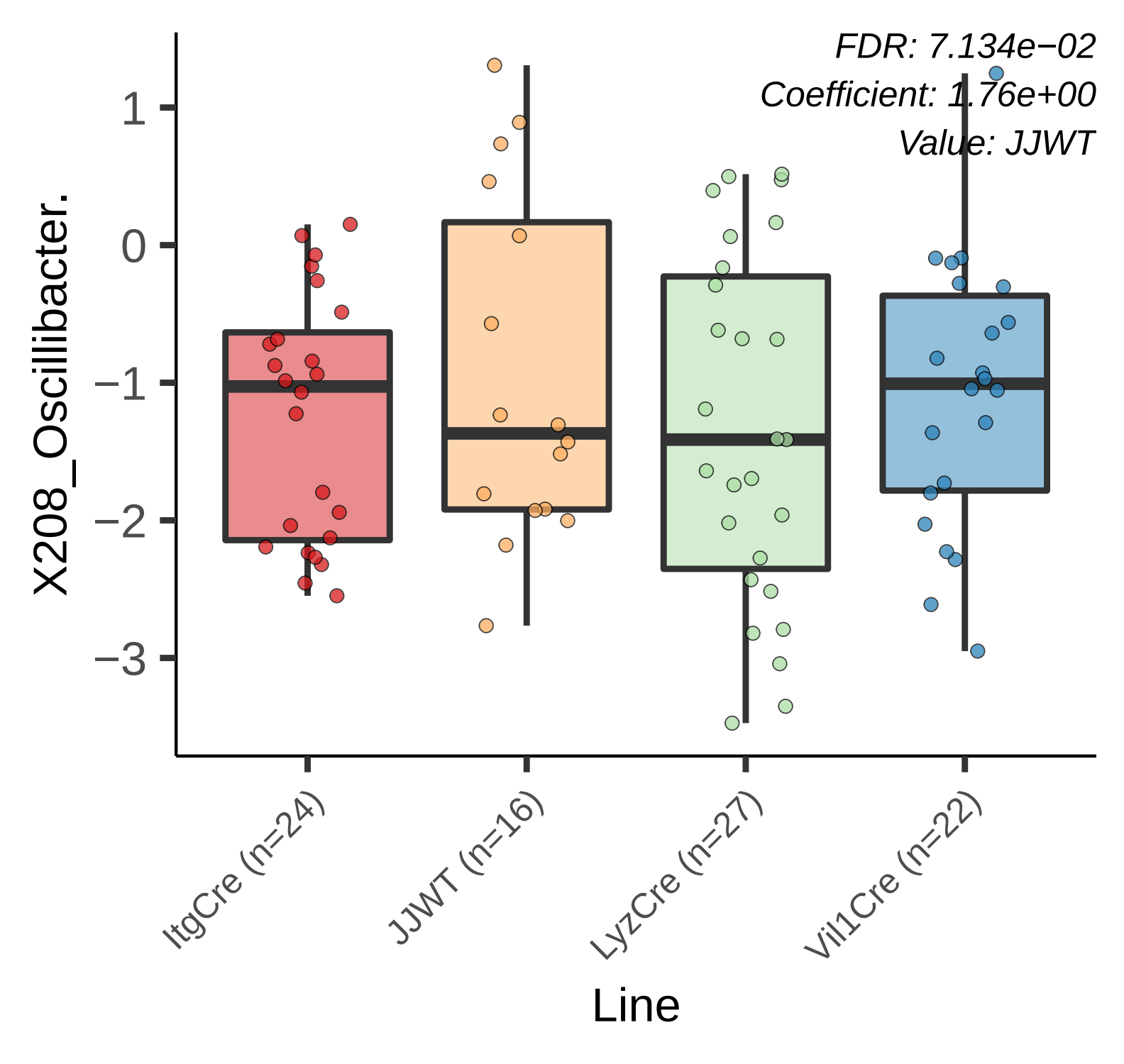

FDR: 6.825e−02
Coefficient: −1.65e+00
Value: JJWT
X4_Muribaculaceae..f.
0
2
4
6
ItgCre (n=24)
JJWT (n=16)
LyzCre (n=27)
Vil1Cre (n=22)
Line

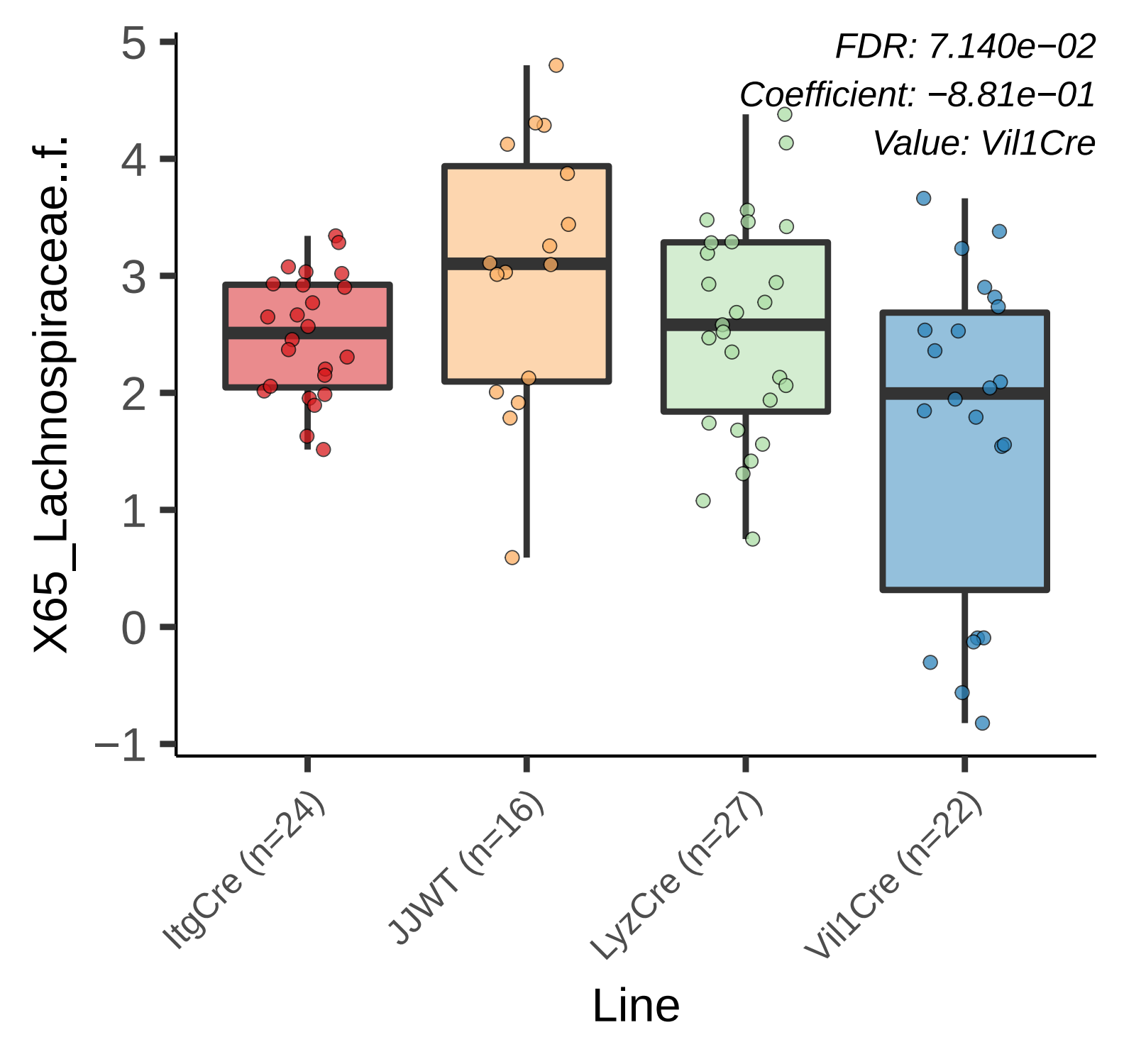

FDR: 6.825e−02
Coefficient: 8.91e−01
Value: LyzCre
X213_Acetatifactor.muris
1
0
−1
−2
−3
ItgCre (n=24)
JJWT (n=16)
LyzCre (n=27)
Vil1Cre (n=22)
Line

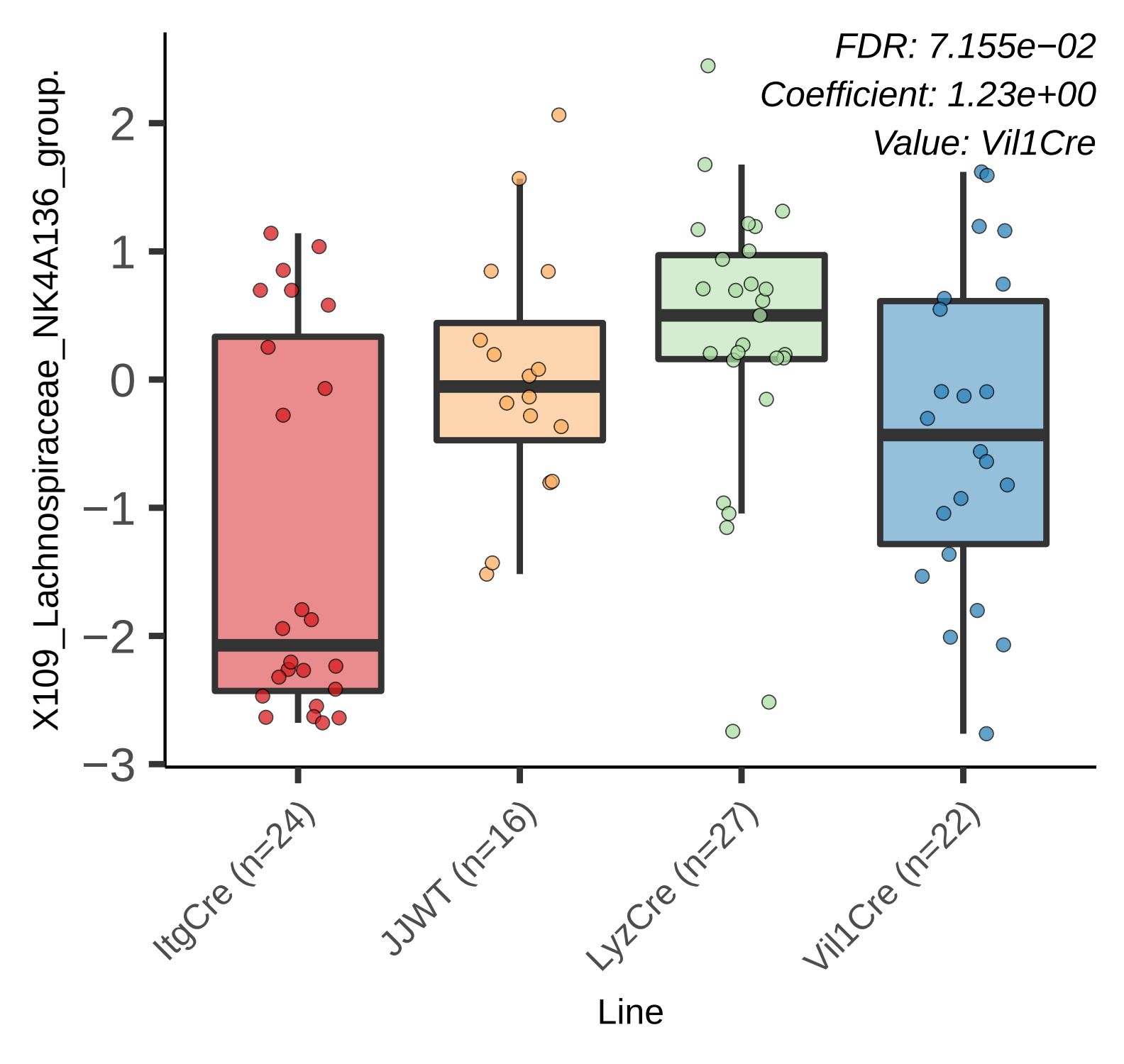

FDR: 6.885e−02
Coefficient: −1.36e+00
Value: Vil1Cre
X40_Lachnospiraceae..f.
2
0
−2
−4
ItgCre (n=24)
JJWT (n=16)
LyzCre (n=27)
Vil1Cre (n=22)
Line

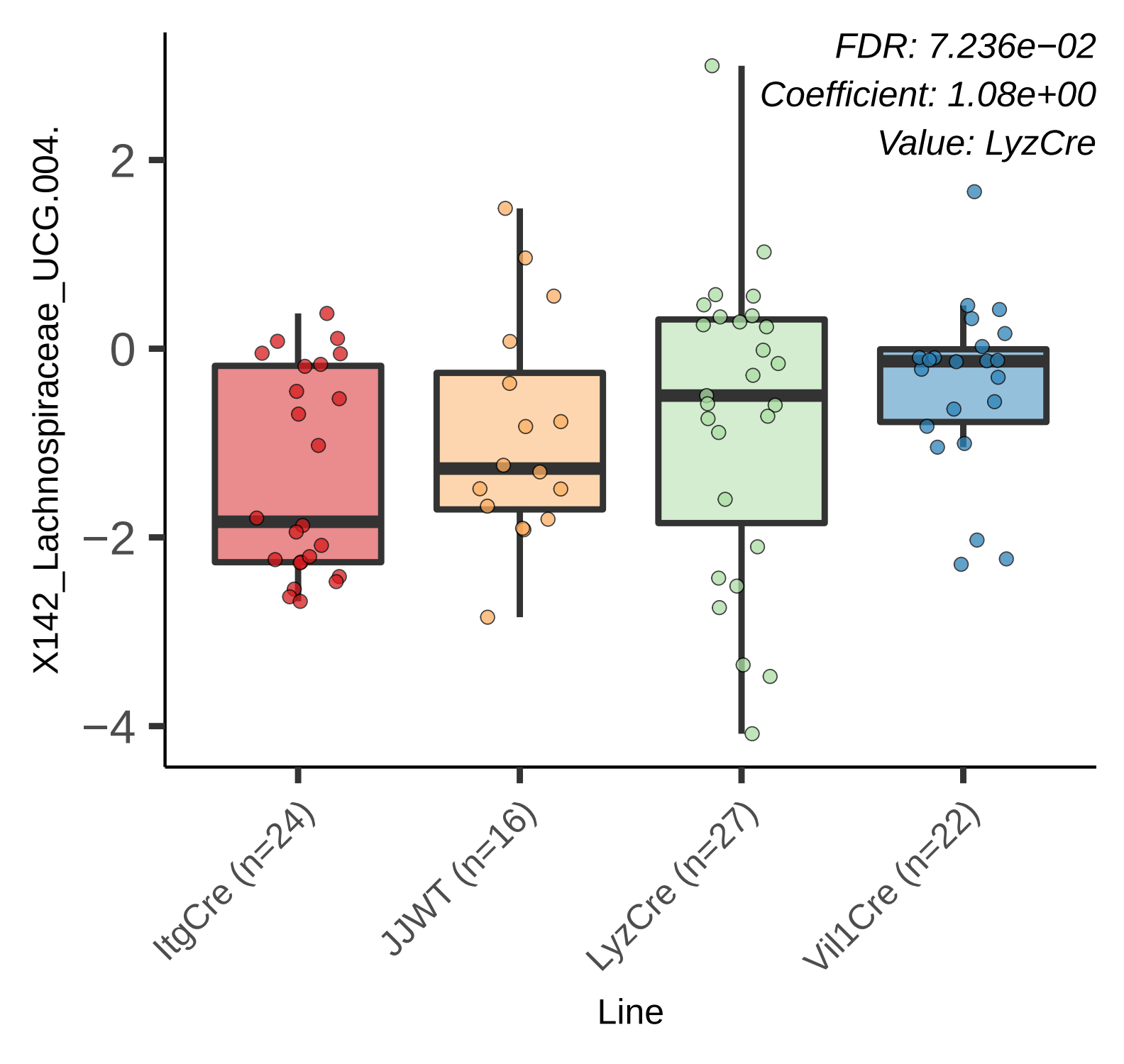

FDR: 6.885e−02
Coefficient: 2.73e+00
Value: JJWT
X82_Lachnospiraceae_NK4A136_group.
2
0
−2
−4
ItgCre (n=24)
JJWT (n=16)
LyzCre (n=27)
Vil1Cre (n=22)
Line

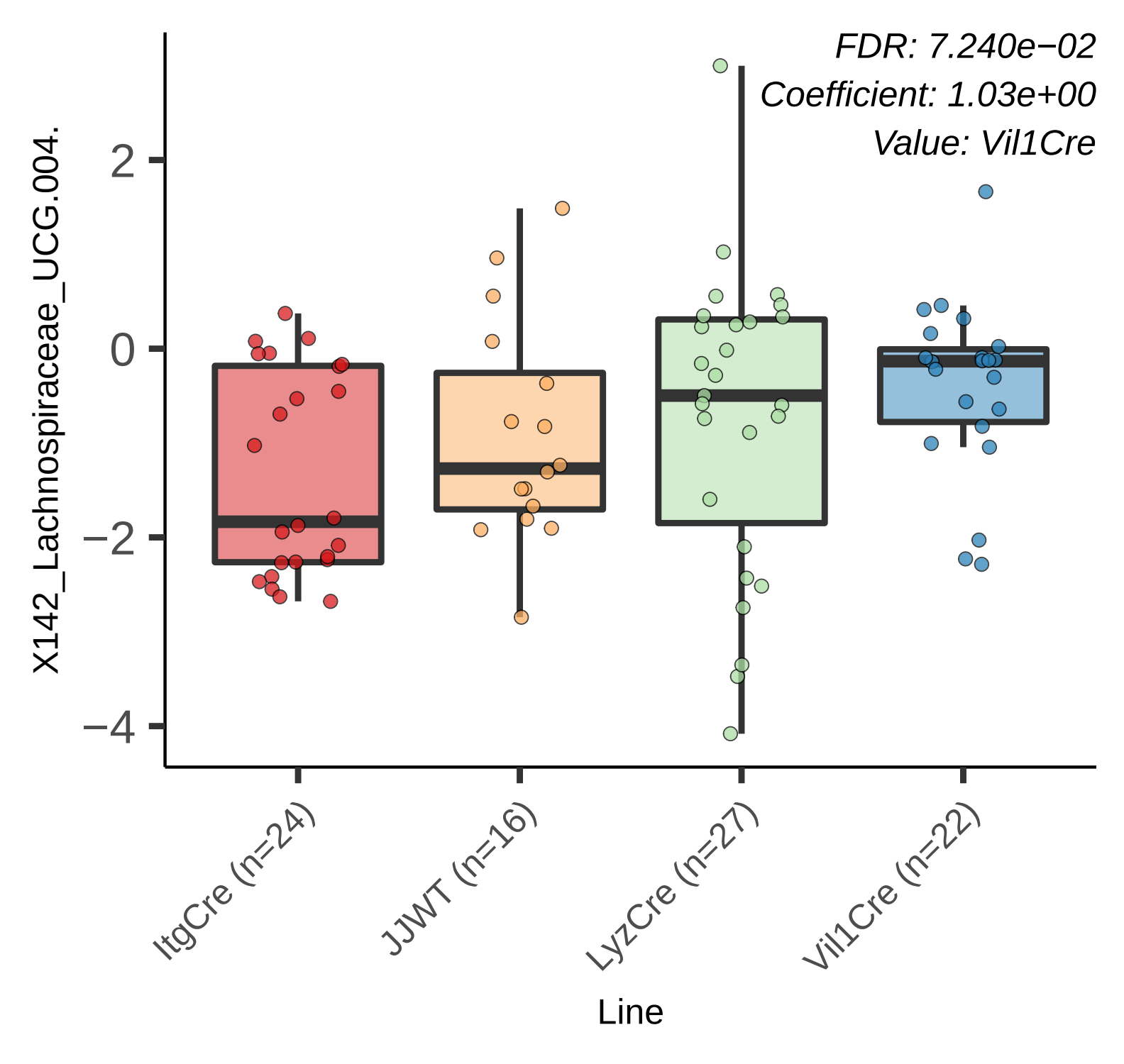

FDR: 7.134e−02
Coefficient: −2.06e+00
Value: JJWT
X53_Muribaculaceae..f.
6
4
2
0
−2
ItgCre (n=24)
JJWT (n=16)
LyzCre (n=27)
Vil1Cre (n=22)
Line

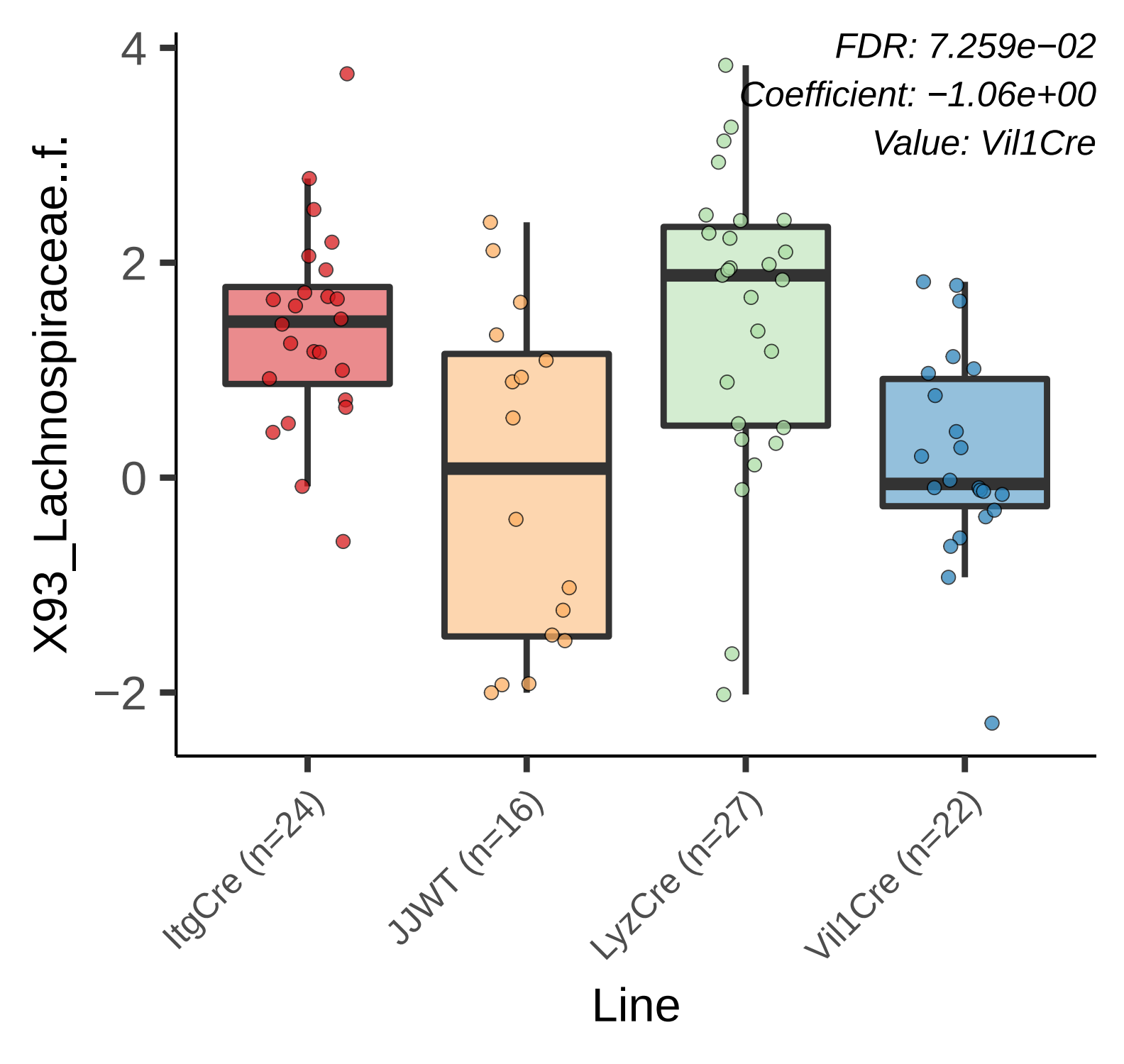

FDR: 7.134e−02
Coefficient: 1.76e+00
Value: JJWT
X208_Oscillibacter.
1
0
−1
−2
−3
ItgCre (n=24)
JJWT (n=16)
LyzCre (n=27)
Vil1Cre (n=22)
Line

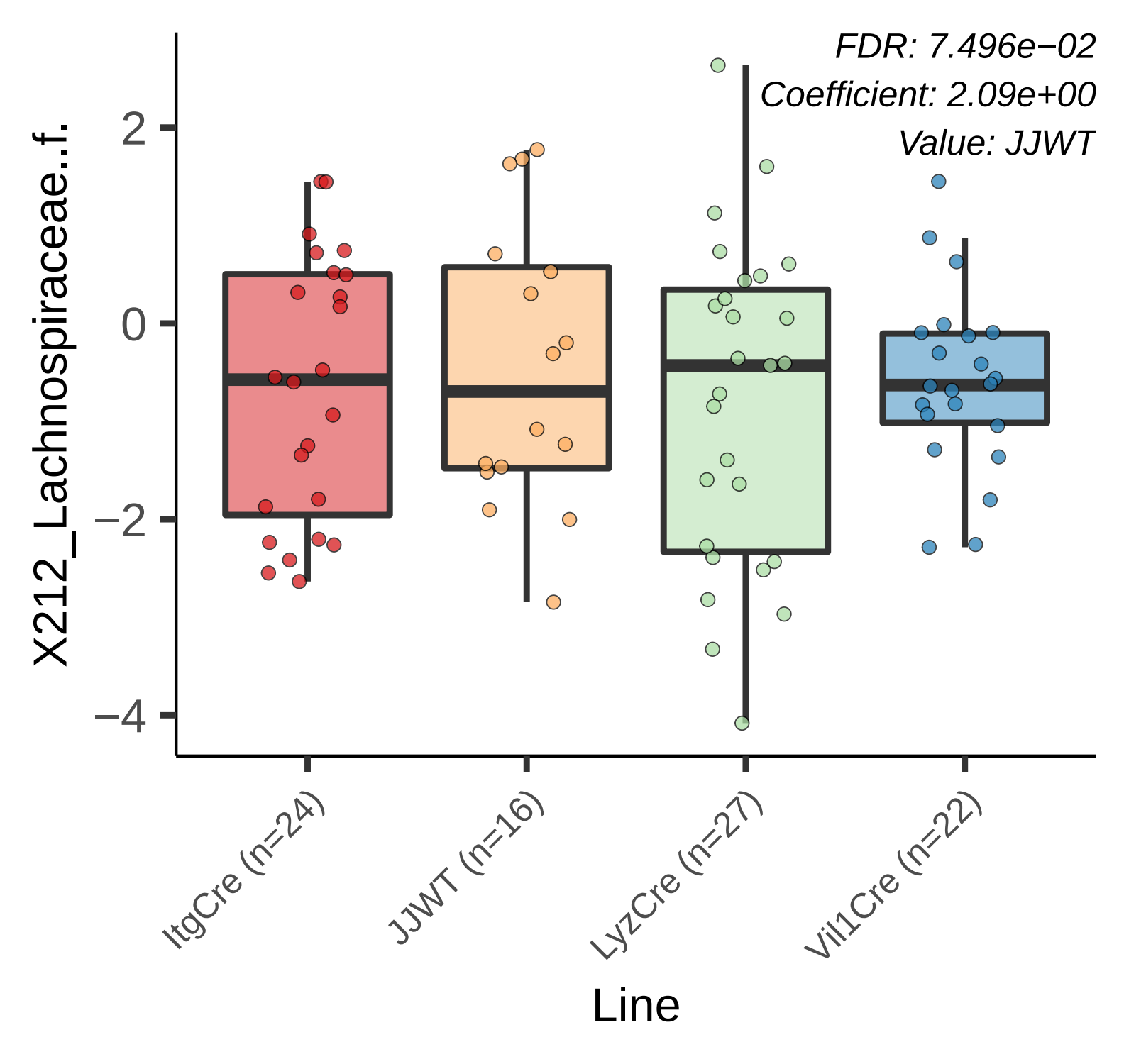

FDR: 7.140e−02
Coefficient: −8.81e−01
Value: Vil1Cre
X65_Lachnospiraceae..f.
5
4
3
2
1
0
−1
ItgCre (n=24)
JJWT (n=16)
LyzCre (n=27)
Vil1Cre (n=22)
Line

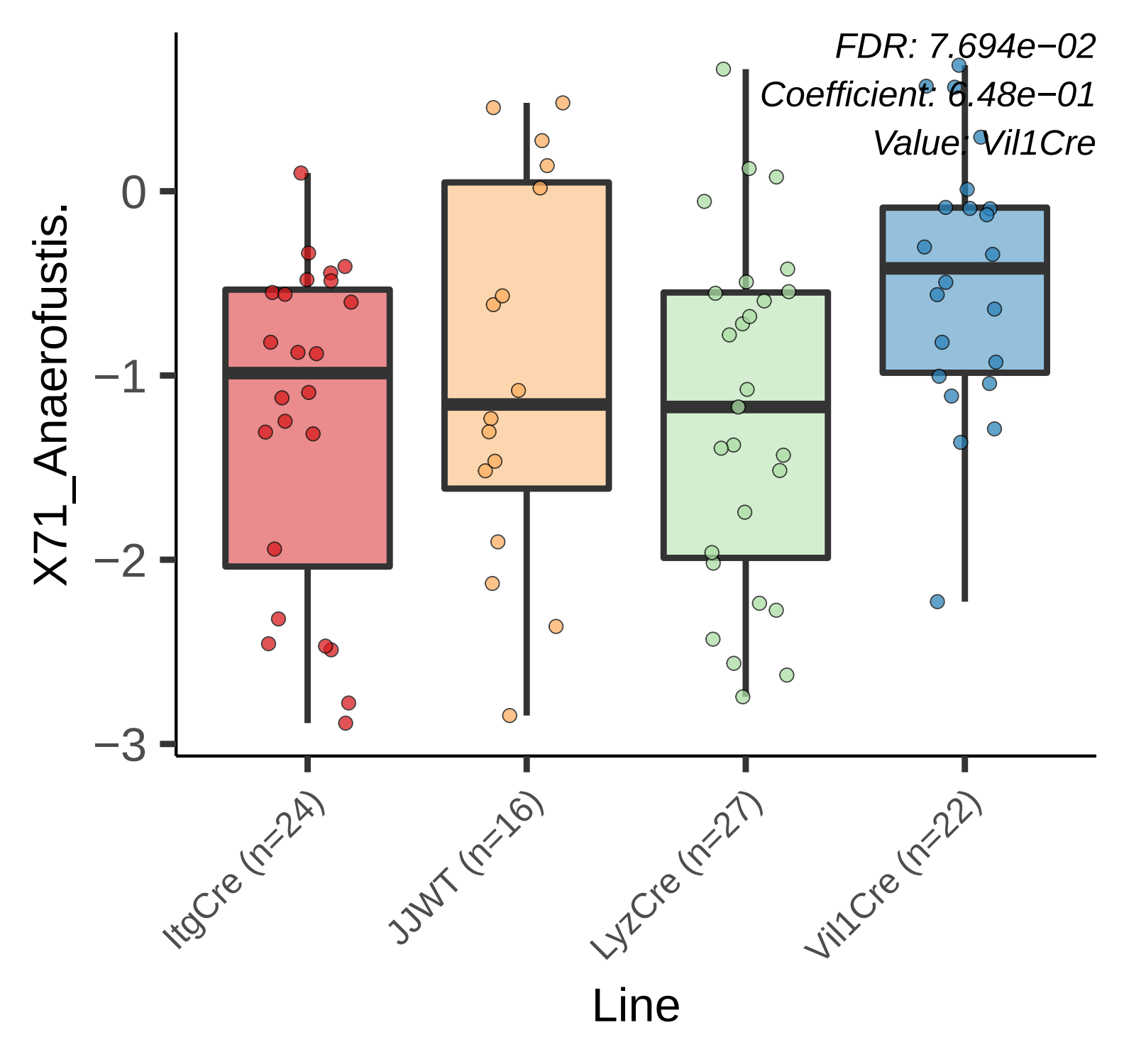

FDR: 7.155e−02
Coefficient: 1.23e+00
Value: Vil1Cre
X109_Lachnospiraceae_NK4A136_group.
2
1
0
−1
−2
−3
ItgCre (n=24)
JJWT (n=16)
LyzCre (n=27)
Vil1Cre (n=22)
Line

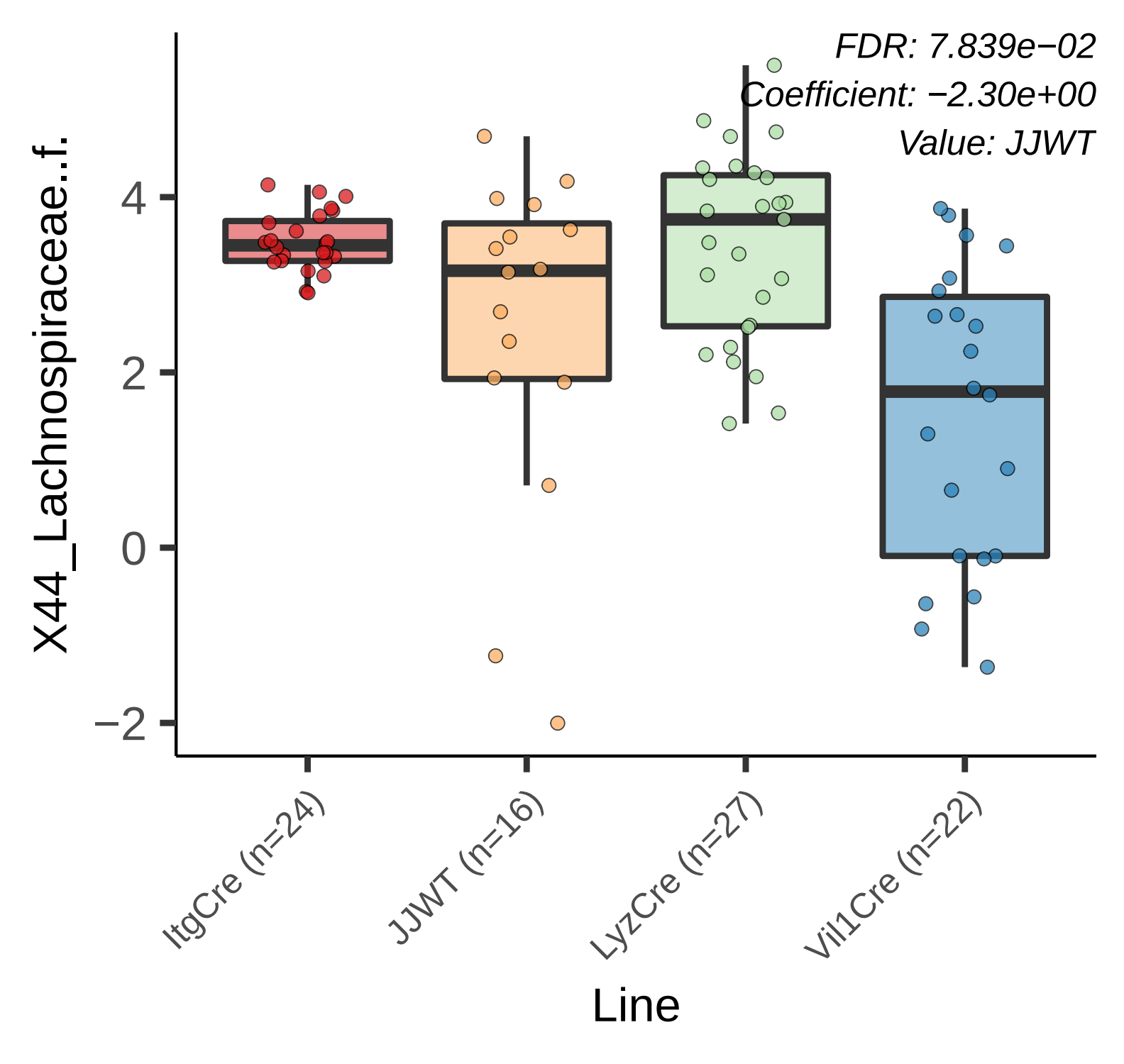

FDR: 7.236e−02
Coefficient: 1.08e+00
Value: LyzCre
X142_Lachnospiraceae_UCG.004.
2
0
−2
−4
ItgCre (n=24)
JJWT (n=16)
LyzCre (n=27)
Vil1Cre (n=22)
Line

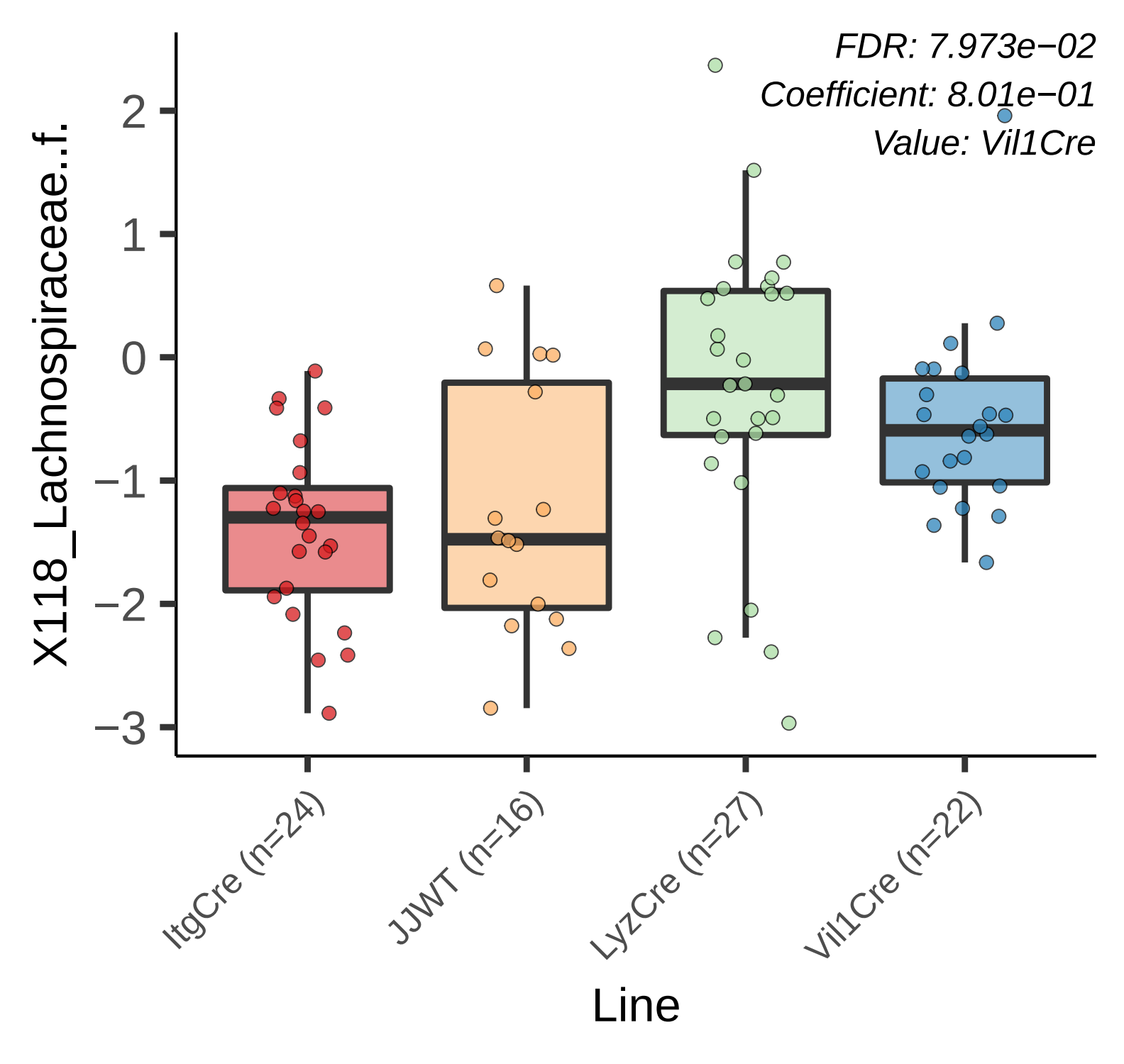

FDR: 7.240e−02
Coefficient: 1.03e+00
Value: Vil1Cre
X142_Lachnospiraceae_UCG.004.
2
0
−2
−4
ItgCre (n=24)
JJWT (n=16)
LyzCre (n=27)
Vil1Cre (n=22)
Line

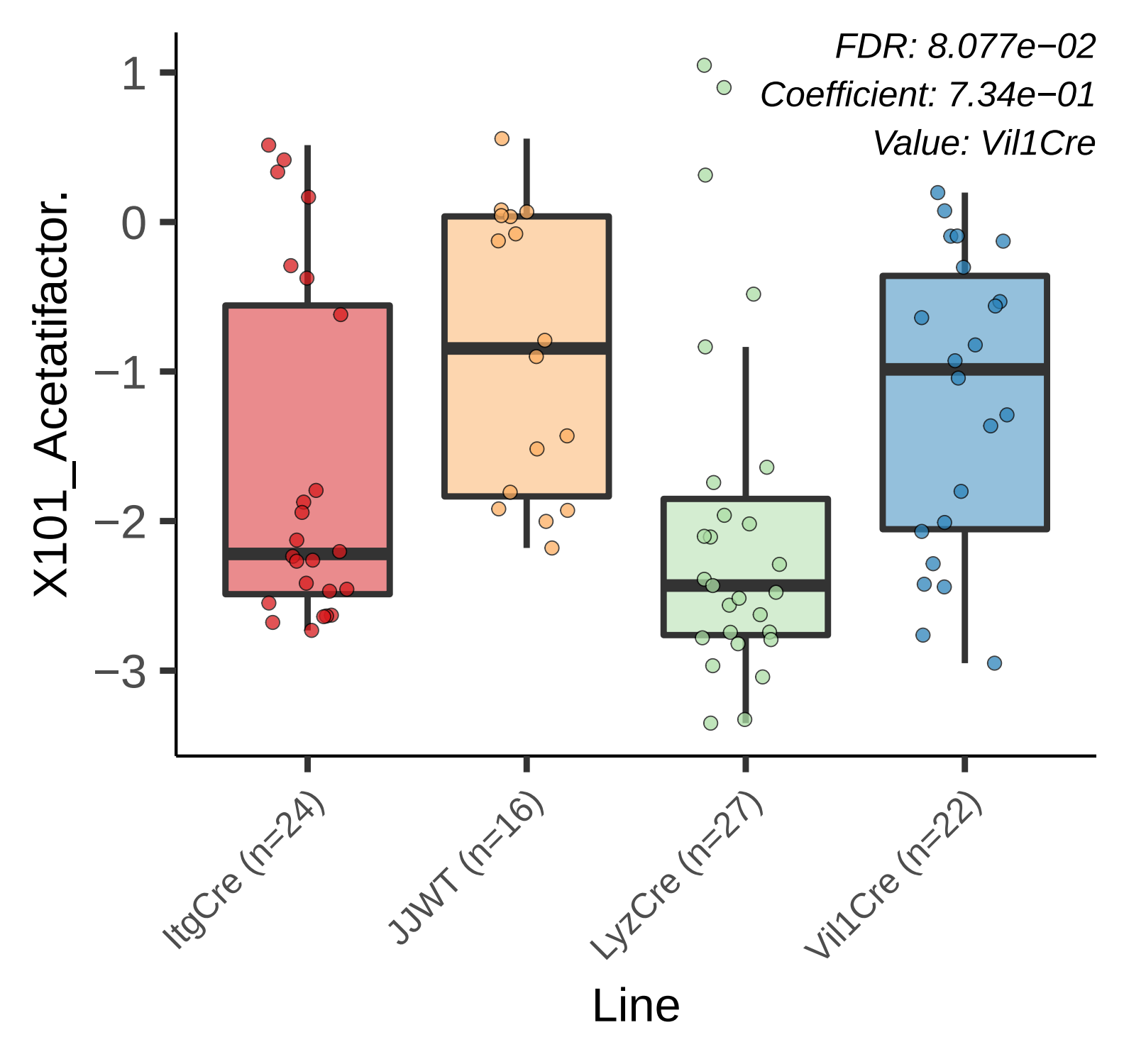

FDR: 7.259e−02
Coefficient: −1.06e+00
Value: Vil1Cre
X93_Lachnospiraceae..f.
4
2
0
−2
ItgCre (n=24)
JJWT (n=16)
LyzCre (n=27)
Vil1Cre (n=22)
Line

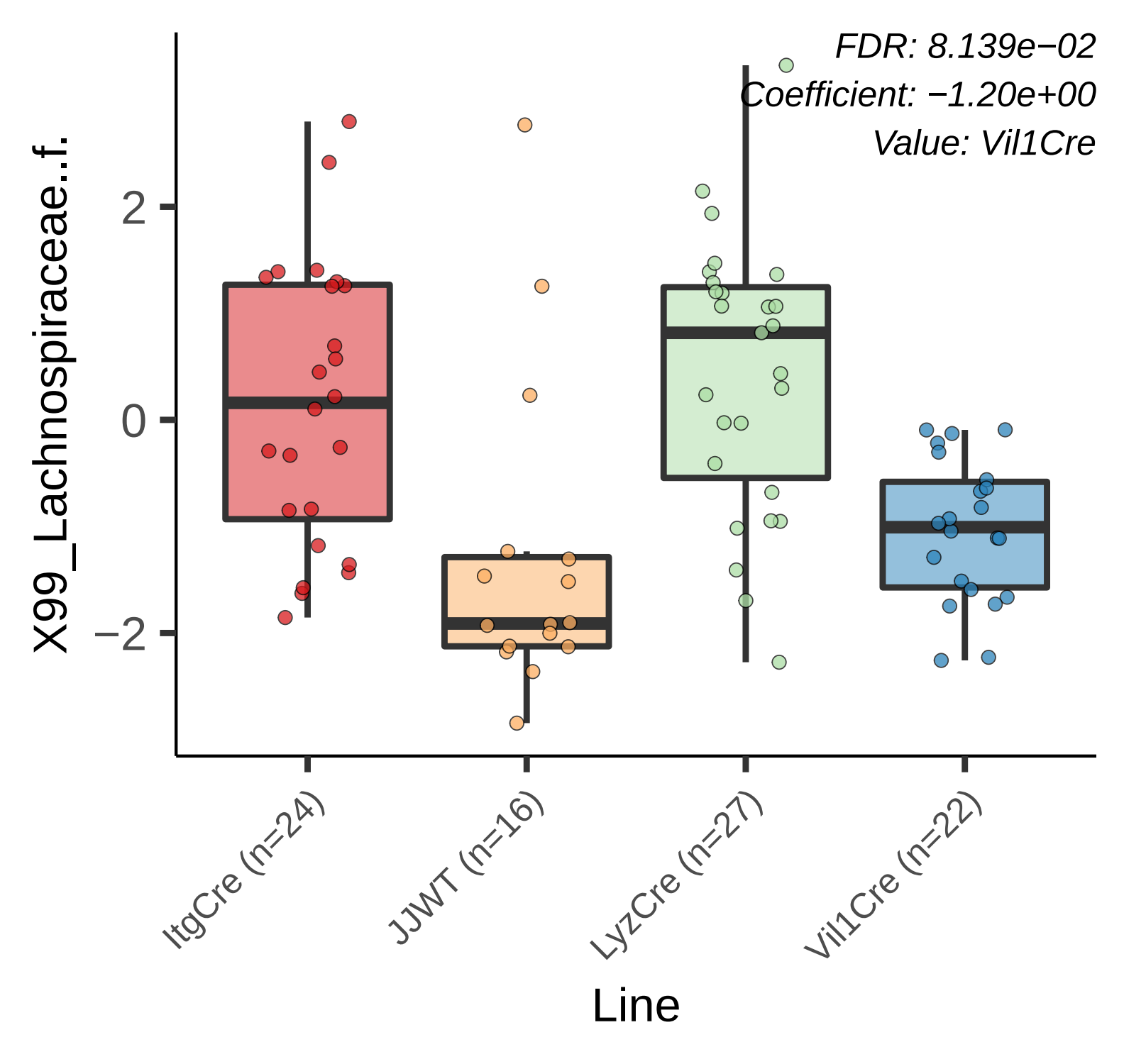

FDR: 7.496e−02
Coefficient: 2.09e+00
Value: JJWT
X212_Lachnospiraceae.f.
2
0
−2
−4
ItgCre (n=24)
JJWT (n=16)
LyzCre (n=27)
Vil1Cre (n=22)
Line

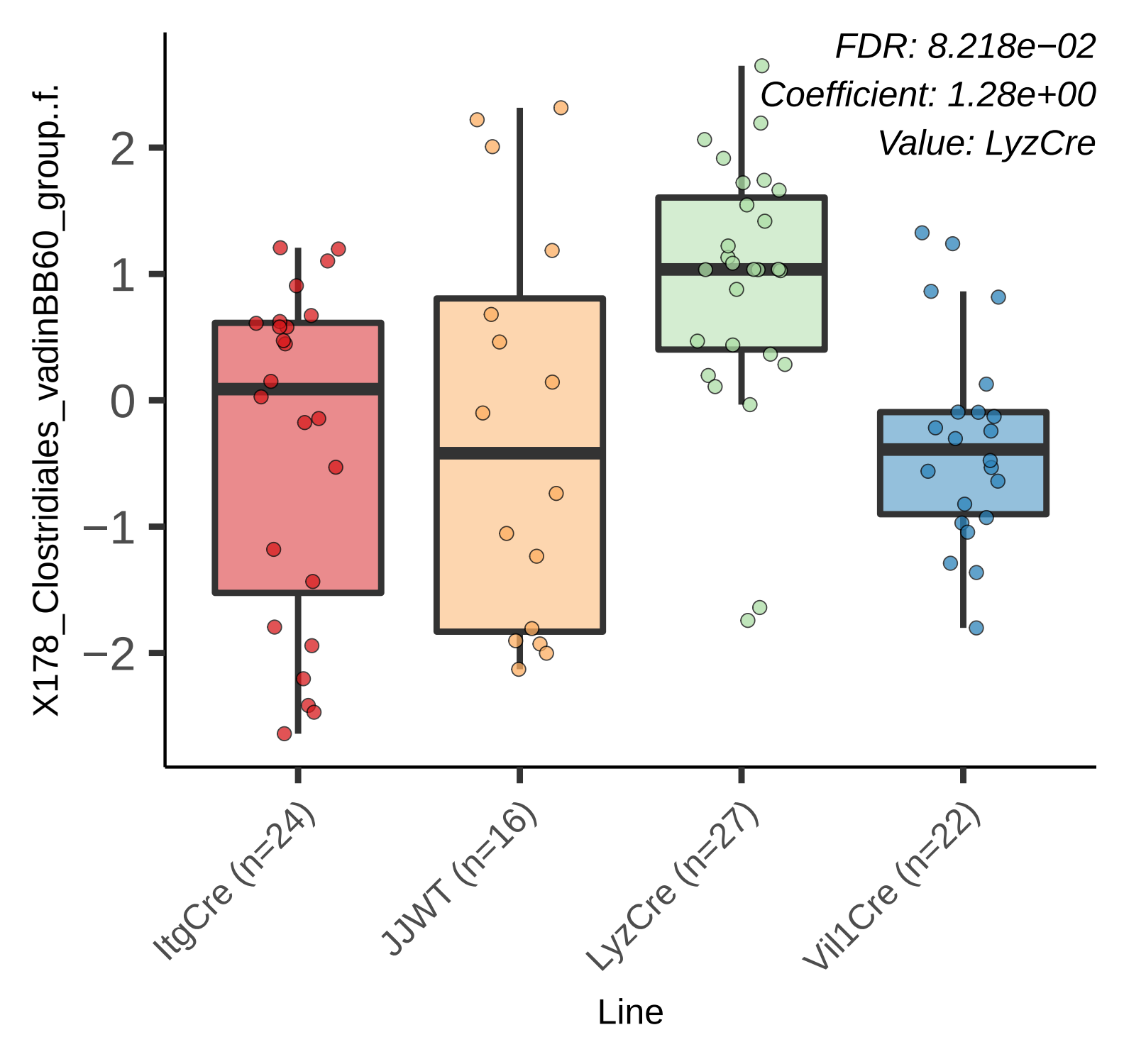

FDR: 7.694e−02
Coefficient: 6.48e−01
Value: Vil1Cre
X71_Anaerofustis.
0
−1
−2
−3
ItgCre (n=24)
JJWT (n=16)
LyzCre (n=27)
Vil1Cre (n=22)
Line

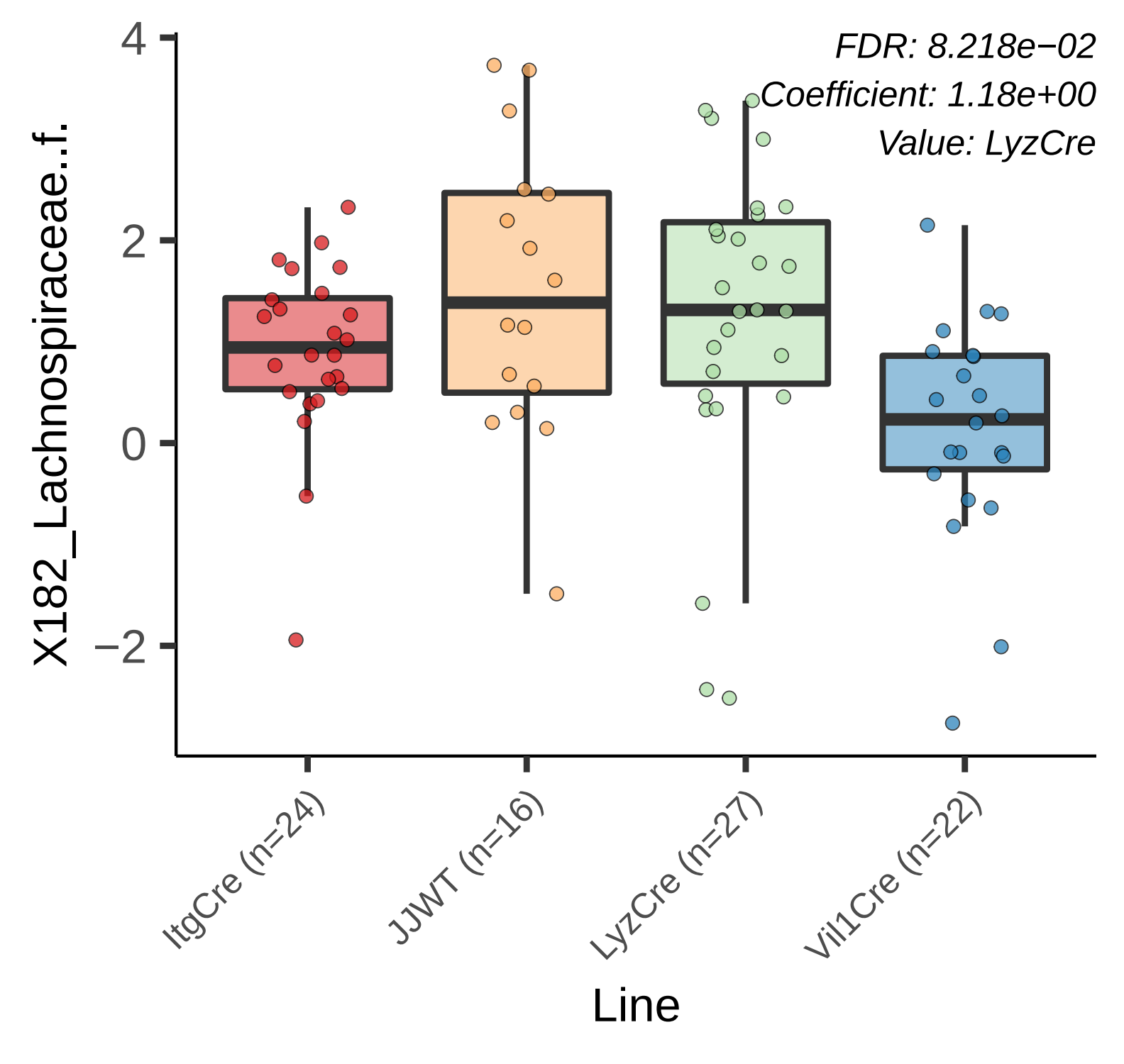

FDR: 7.839e−02
Coefficient: −2.30e+00
Value: JJWT
X44_Lachnospiraceae..f.
4
2
0
−2
ItgCre (n=24)
JJWT (n=16)
LyzCre (n=27)
Vil1Cre (n=22)
Line

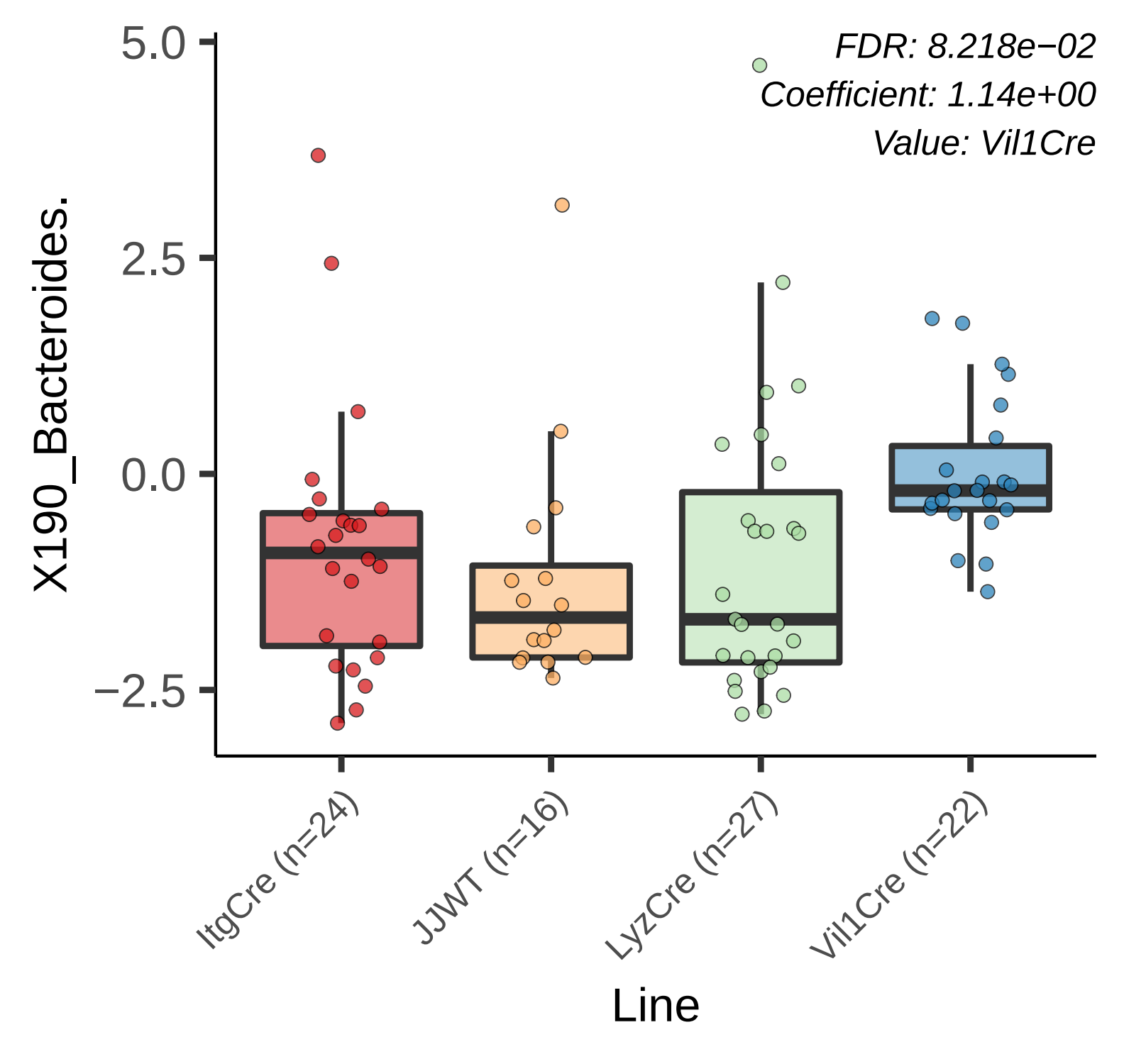

FDR: 7.973e−02
Coefficient: 8.01e−01
Value: Vil1Cre
X118_Lachnospiraceae.f.
2
1
0
−1
−2
−3
ItgCre (n=24)
JJWT (n=16)
LyzCre (n=27)
Vil1Cre (n=22)
Line

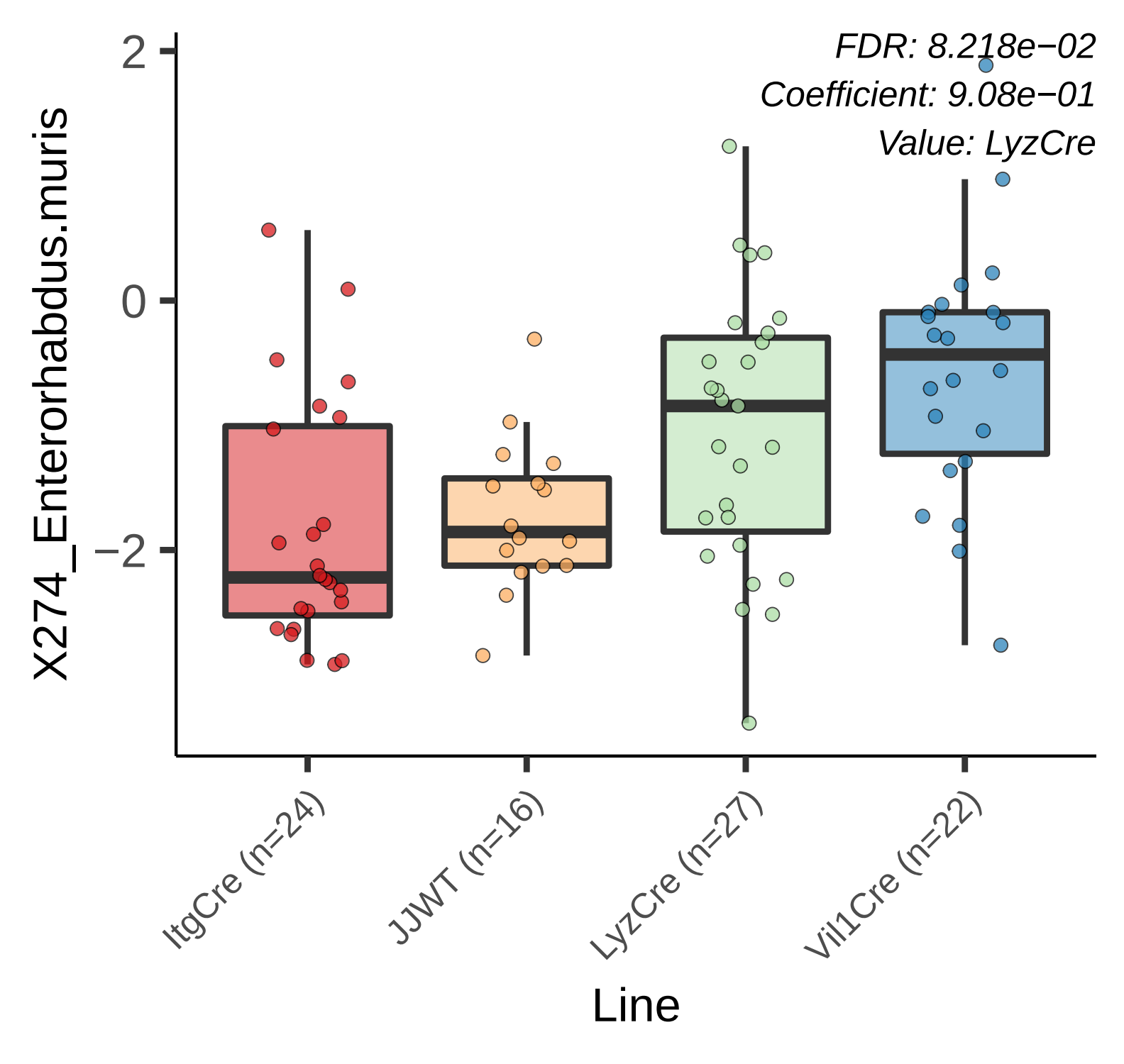

FDR: 8.077e−02
Coefficient: 7.34e−01
Value: Vil1Cre
X101_Acetatifactor.
1
0
−1
−2
−3
ItgCre (n=24)
JJWT (n=16)
LyzCre (n=27)
Vil1Cre (n=22)
Line

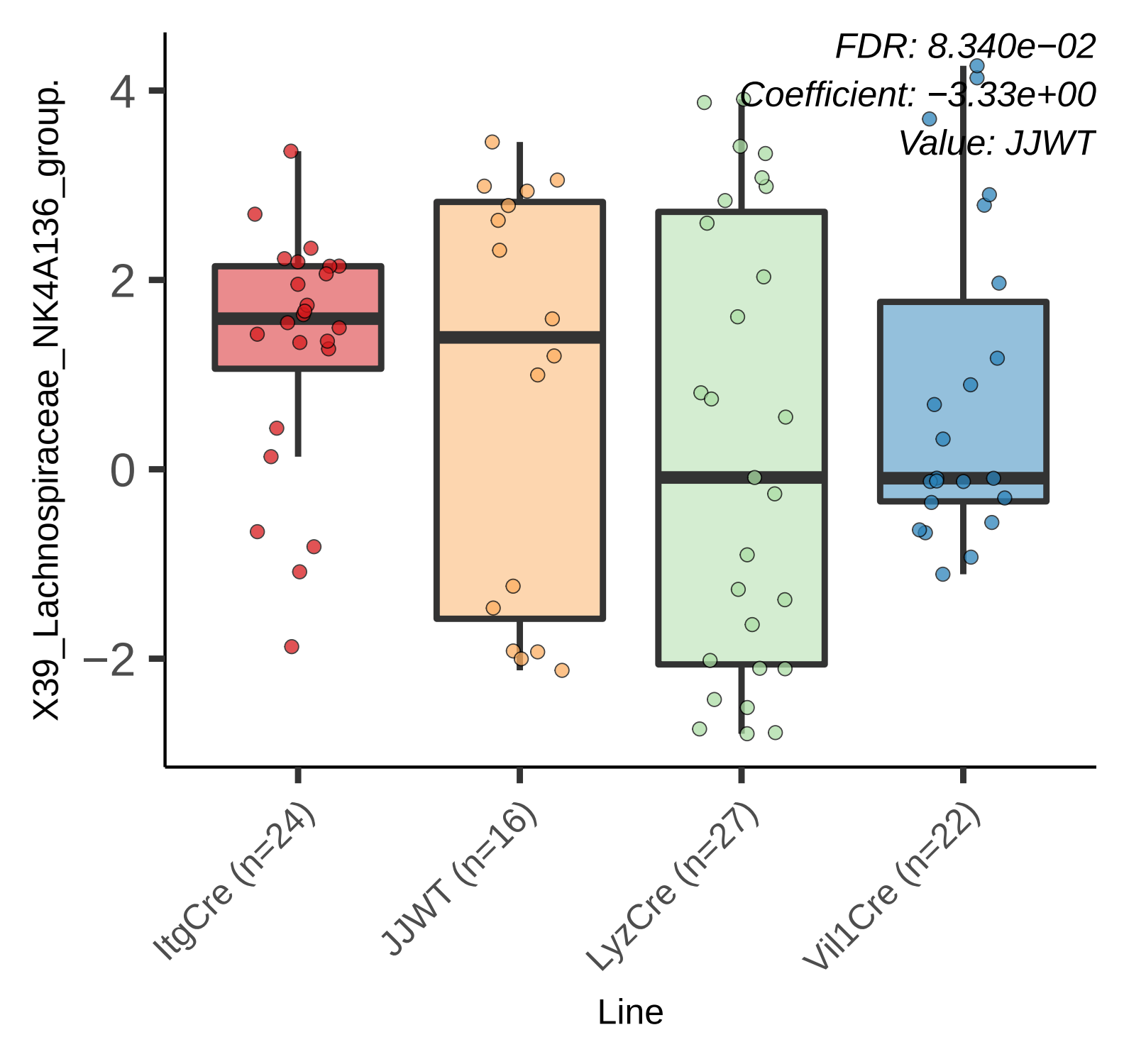

FDR: 8.139e−02
Coefficient: −1.20e+00
Value: Vil1Cre
X99_Lachnospiraceae..f.
2
0
−2
ItgCre (n=24)
JJWT (n=16)
LyzCre (n=27)
Vil1Cre (n=22)
Line

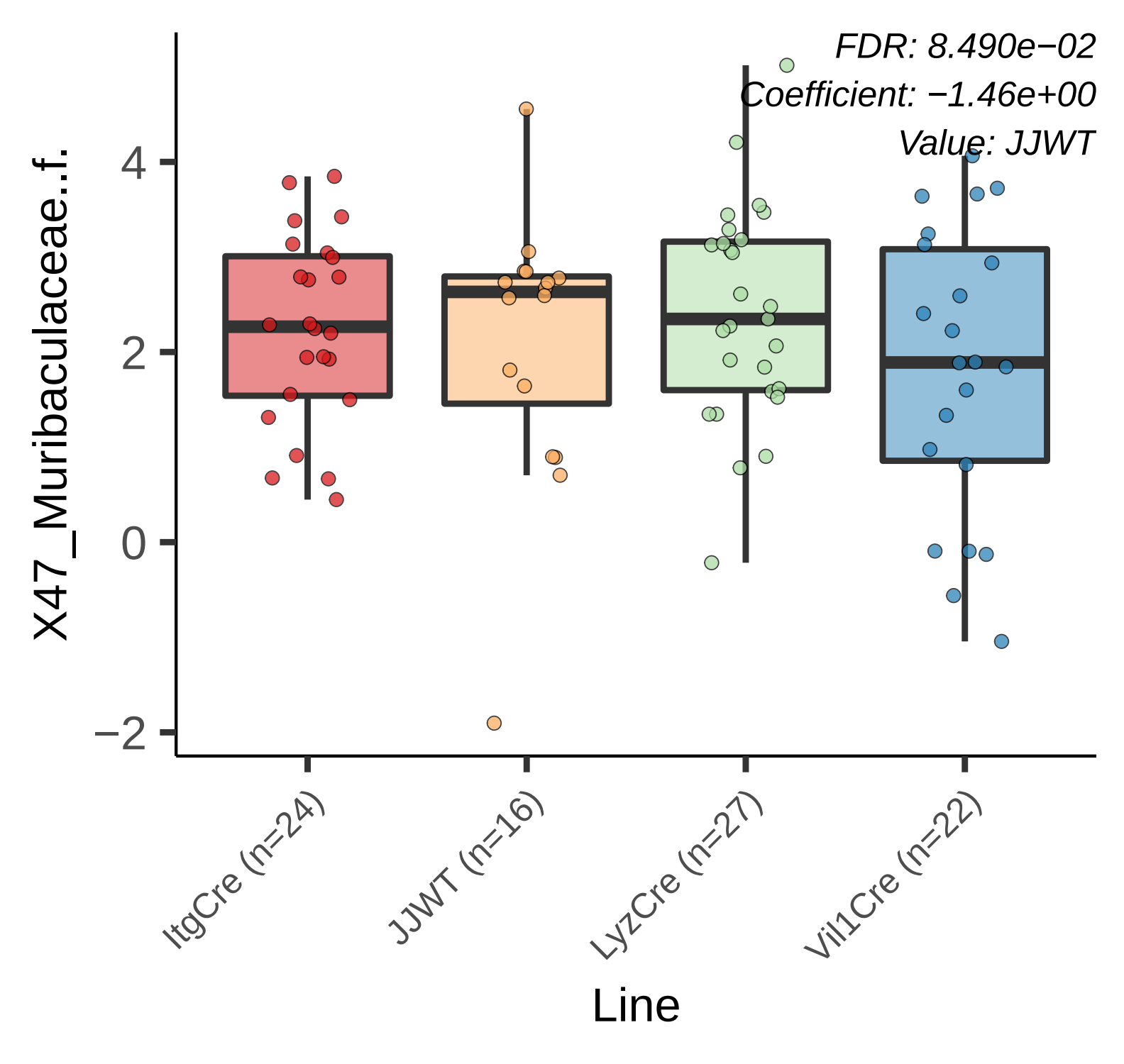

FDR: 8.218e−02
Coefficient: 1.28e+00
Value: LyzCre
X178_Clostridiales_vadinBB60_group..f.
2
1
0
−1
−2
ItgCre (n=24)
JJWT (n=16)
LyzCre (n=27)
Vil1Cre (n=22)
Line

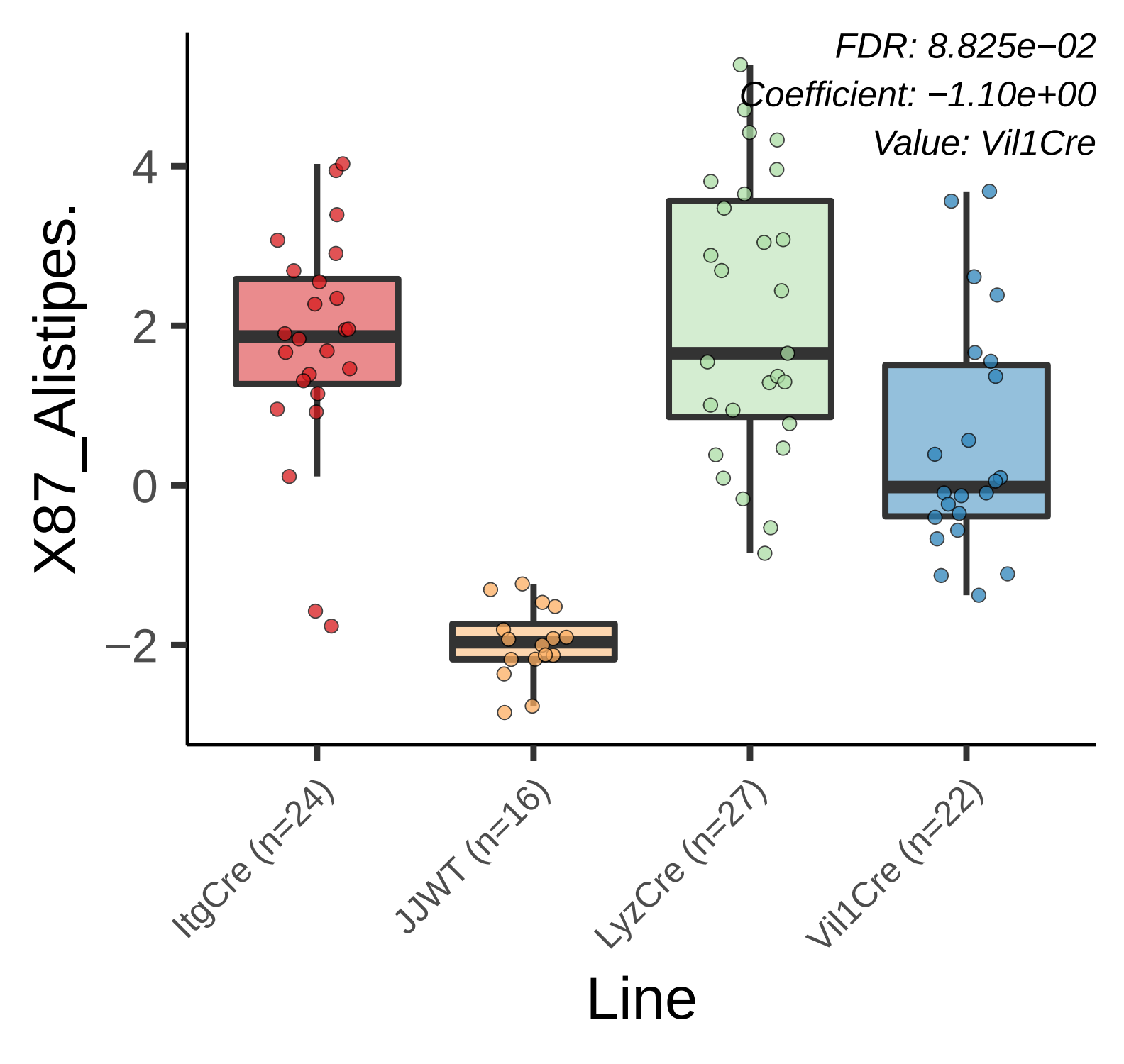

FDR: 8.218e−02
Coefficient: 1.18e+00
Value: LyzCre
X182_Lachnospiraceae.f.
4
2
0
−2
ItgCre (n=24)
JJWT (n=16)
LyzCre (n=27)
Vil1Cre (n=22)
Line

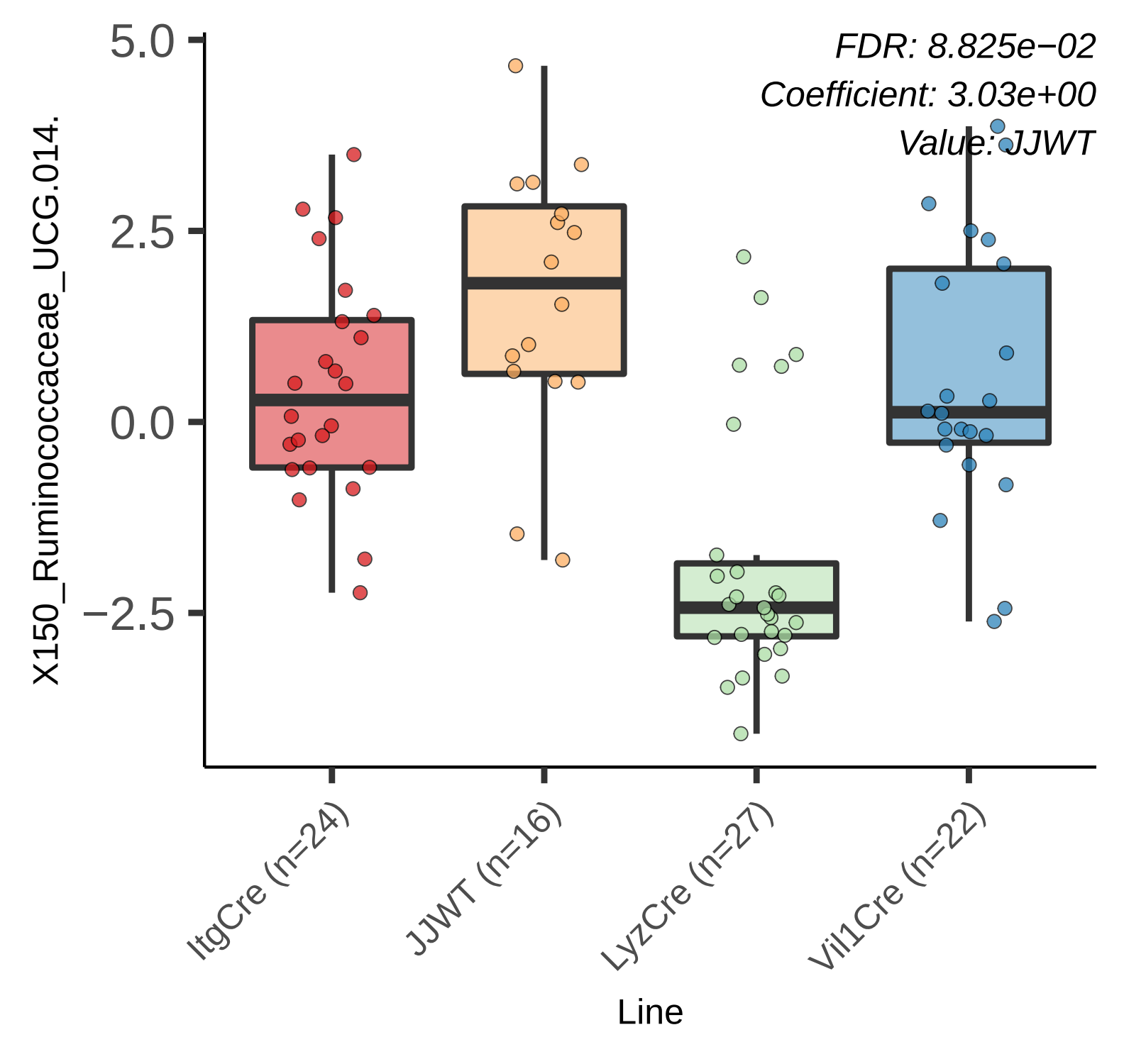

FDR: 8.218e−02
Coefficient: 1.14e+00
Value: Vil1Cre
X190_Bacteroides.
5.0
2.5
0.0
−2.5
ItgCre (n=24)
JJWT (n=16)
LyzCre (n=27)
Vil1Cre (n=22)
Line

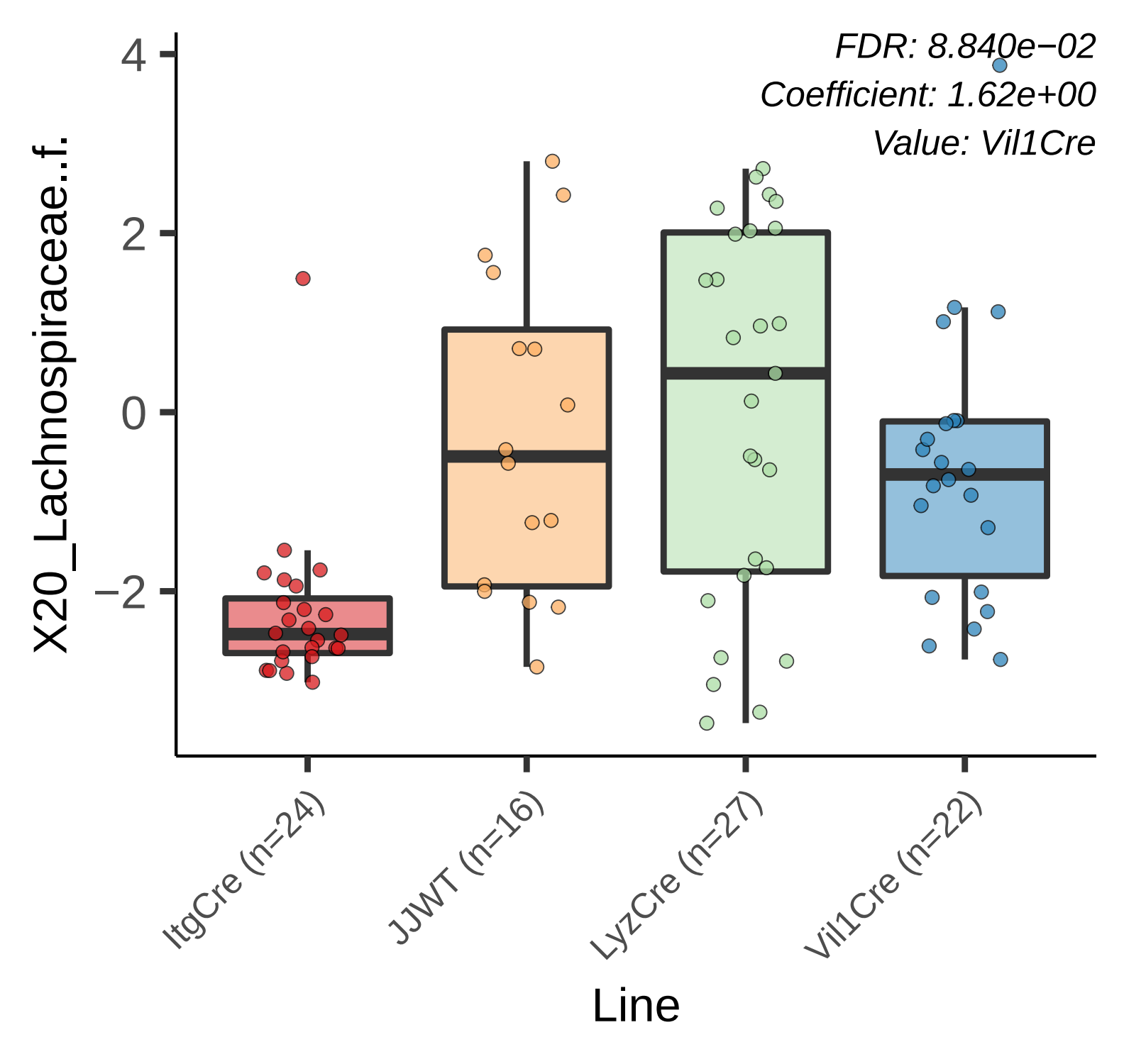

FDR: 8.218e−02
Coefficient: 9.08e−01
Value: LyzCre
X274_Enterorhabdus.muris
2
0
−2
ItgCre (n=24)
JJWT (n=16)
LyzCre (n=27)
Vil1Cre (n=22)
Line

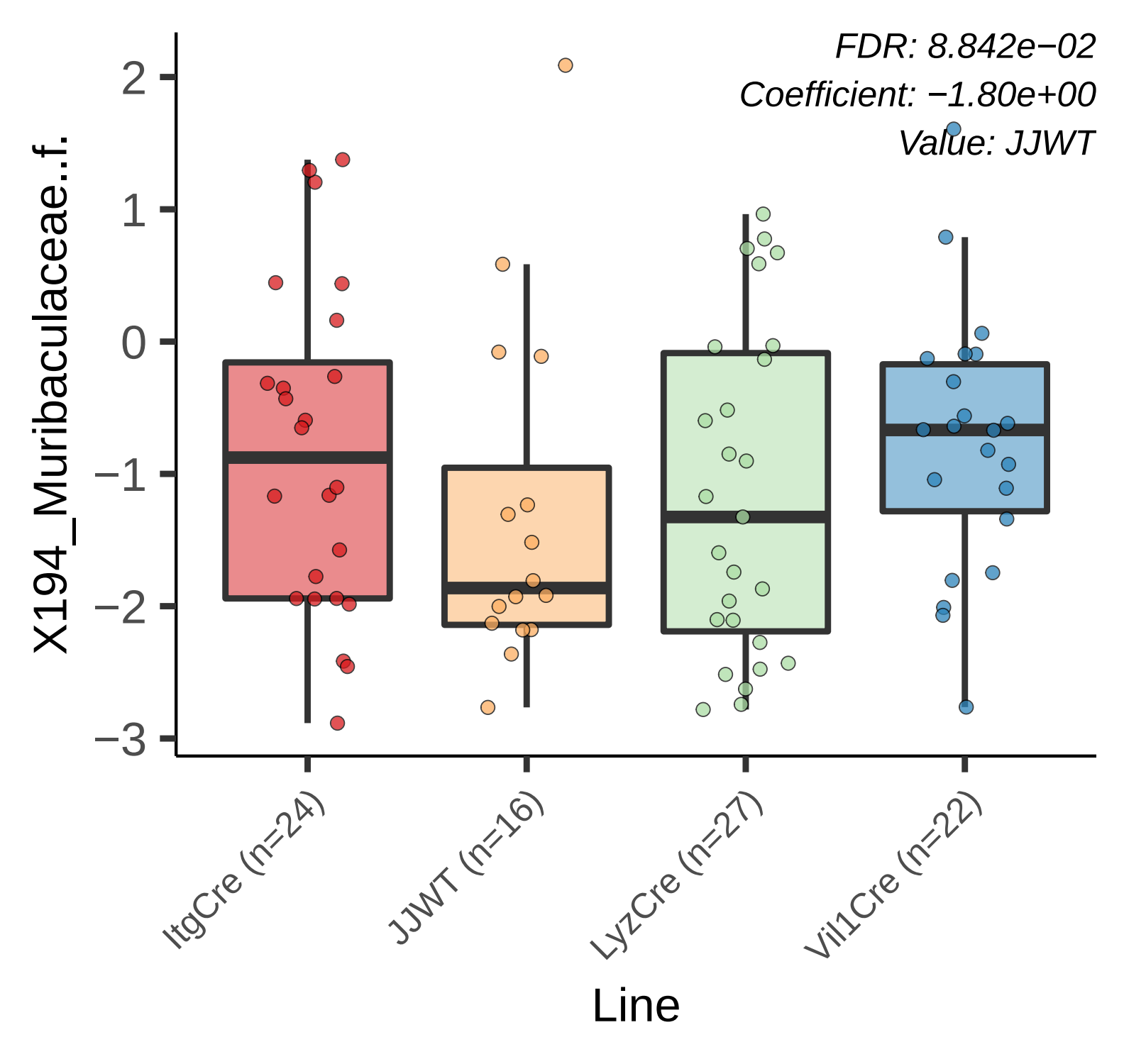

FDR: 8.340e−02
Coefficient: −3.33e+00
Value: JJWT
X39_Lachnospiraceae_NK4A136_group.
4
2
0
−2
ItgCre (n=24)
JJWT (n=16)
LyzCre (n=27)
Vil1Cre (n=22)
Line

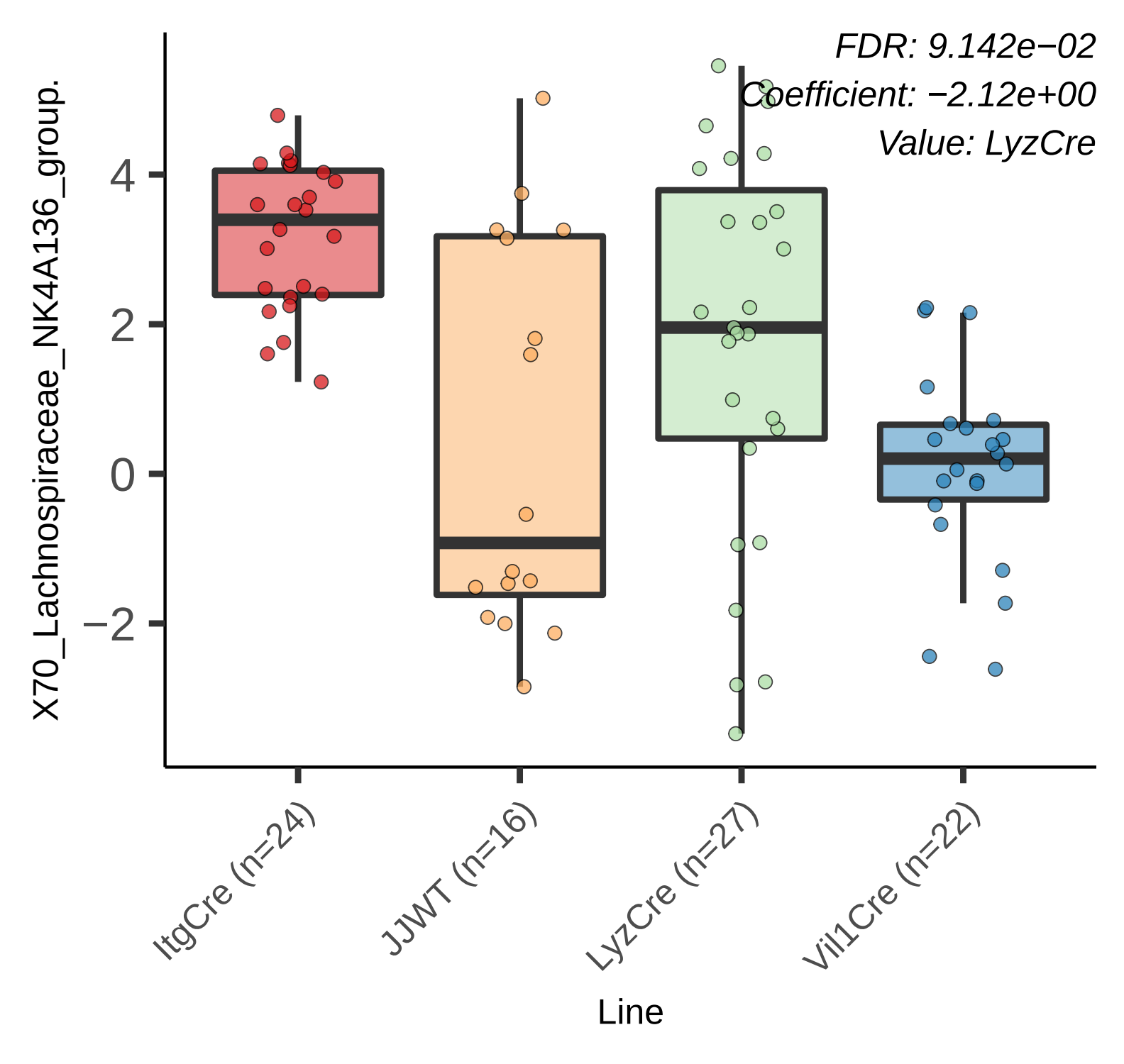

FDR: 8.490e−02
Coefficient: −1.46e+00
Value: JJWT
X47_Muribaculaceae..f.
4
2
0
−2
ItgCre (n=24)
JJWT (n=16)
LyzCre (n=27)
Vil1Cre (n=22)
Line

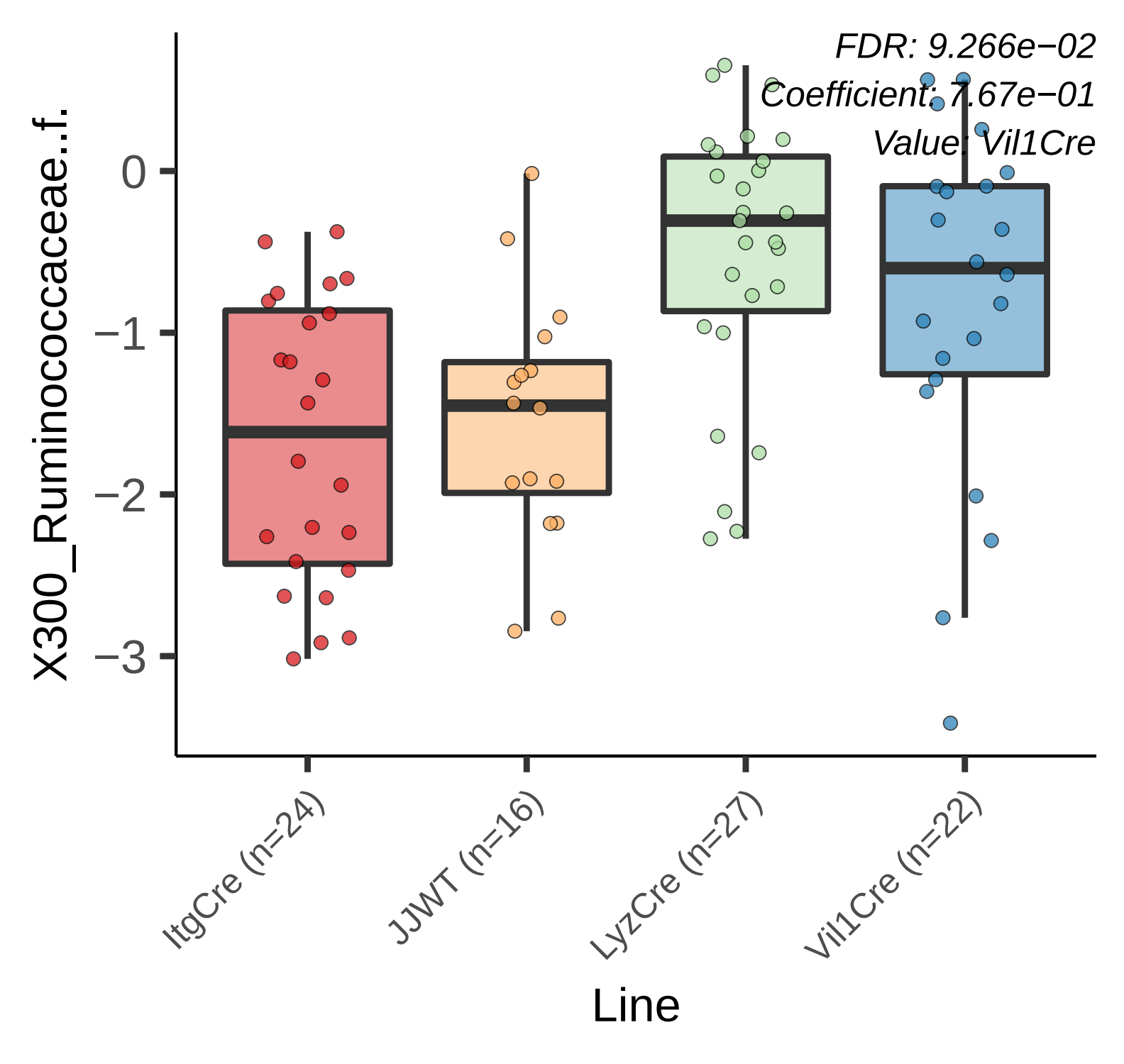

FDR: 8.825e−02
Coefficient: −1.10e+00
Value: Vil1Cre
X87_Alistipes.
4
2
0
−2
ItgCre (n=24)
JJWT (n=16)
LyzCre (n=27)
Vil1Cre (n=22)
Line

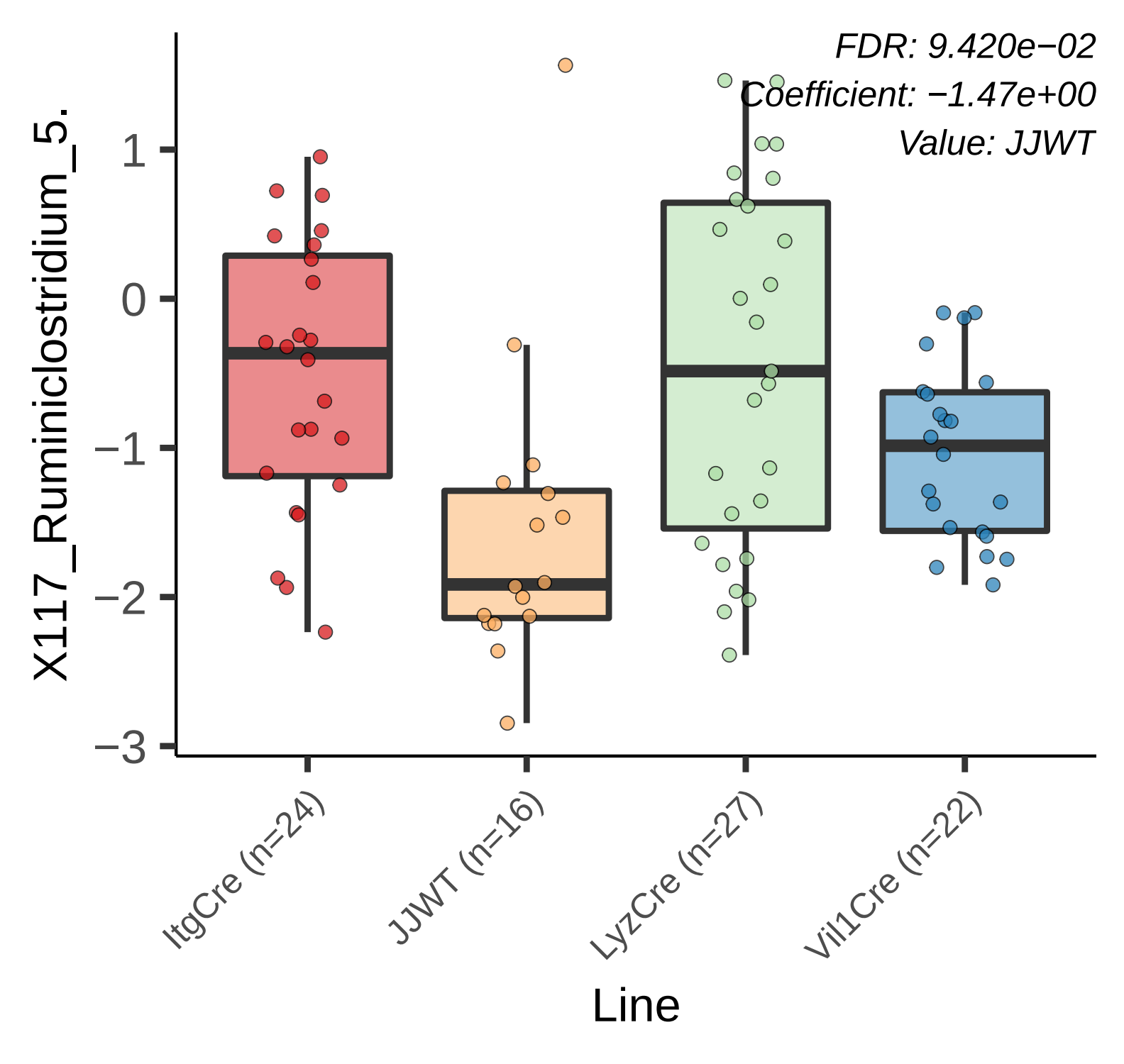

FDR: 8.825e−02
Coefficient: 3.03e+00
Value: JJWT
X150_Ruminococcaceae_UCG.014.
5.0
2.5
0.0
−2.5
ItgCre (n=24)
JJWT (n=16)
LyzCre (n=27)
Vil1Cre (n=22)
Line

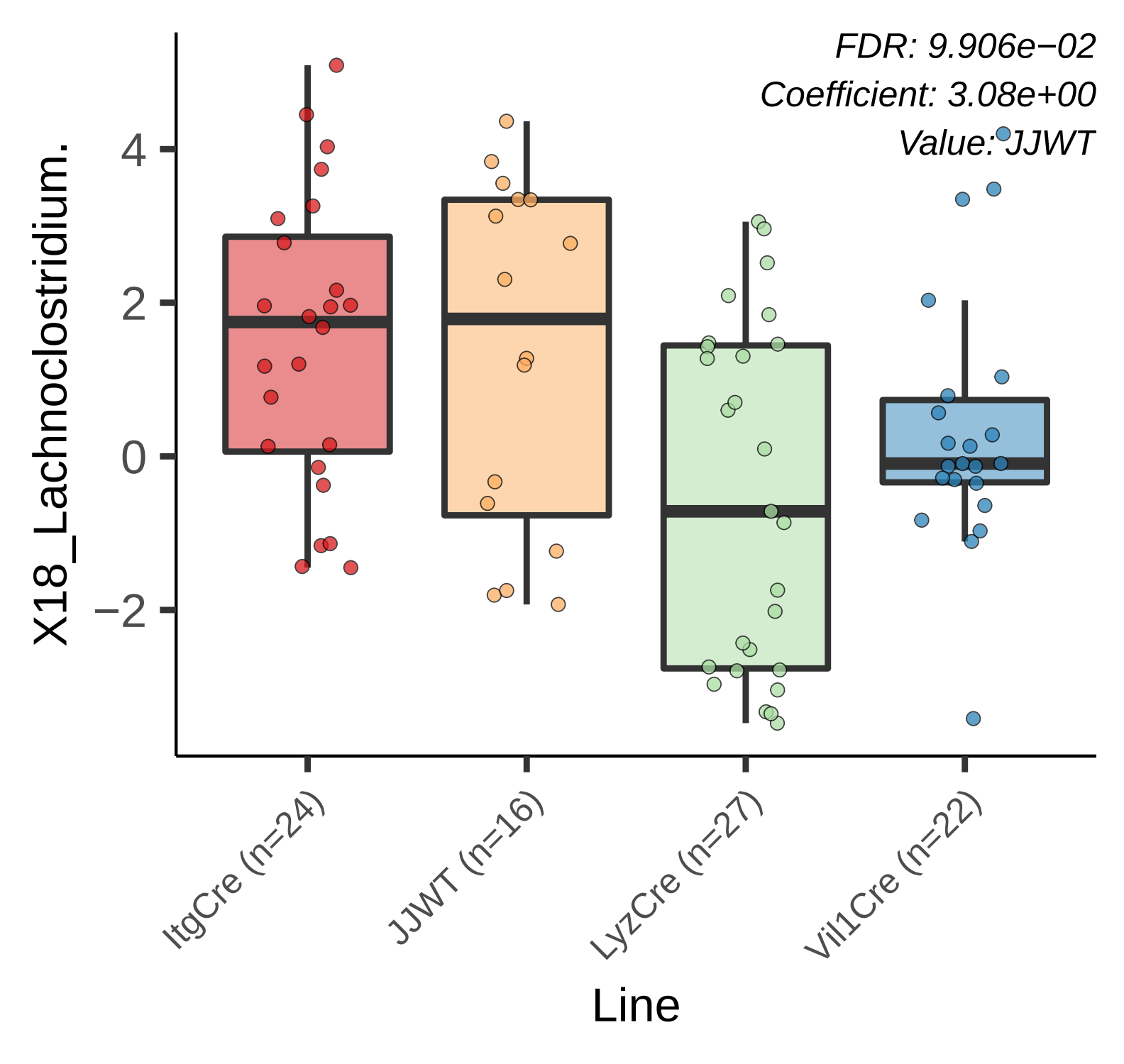

FDR: 8.840e−02
Coefficient: 1.62e+00
Value: Vil1Cre
X20_Lachnospiraceae..f.
4
2
0
−2
ItgCre (n=24)
JJWT (n=16)
LyzCre (n=27)
Vil1Cre (n=22)
Line

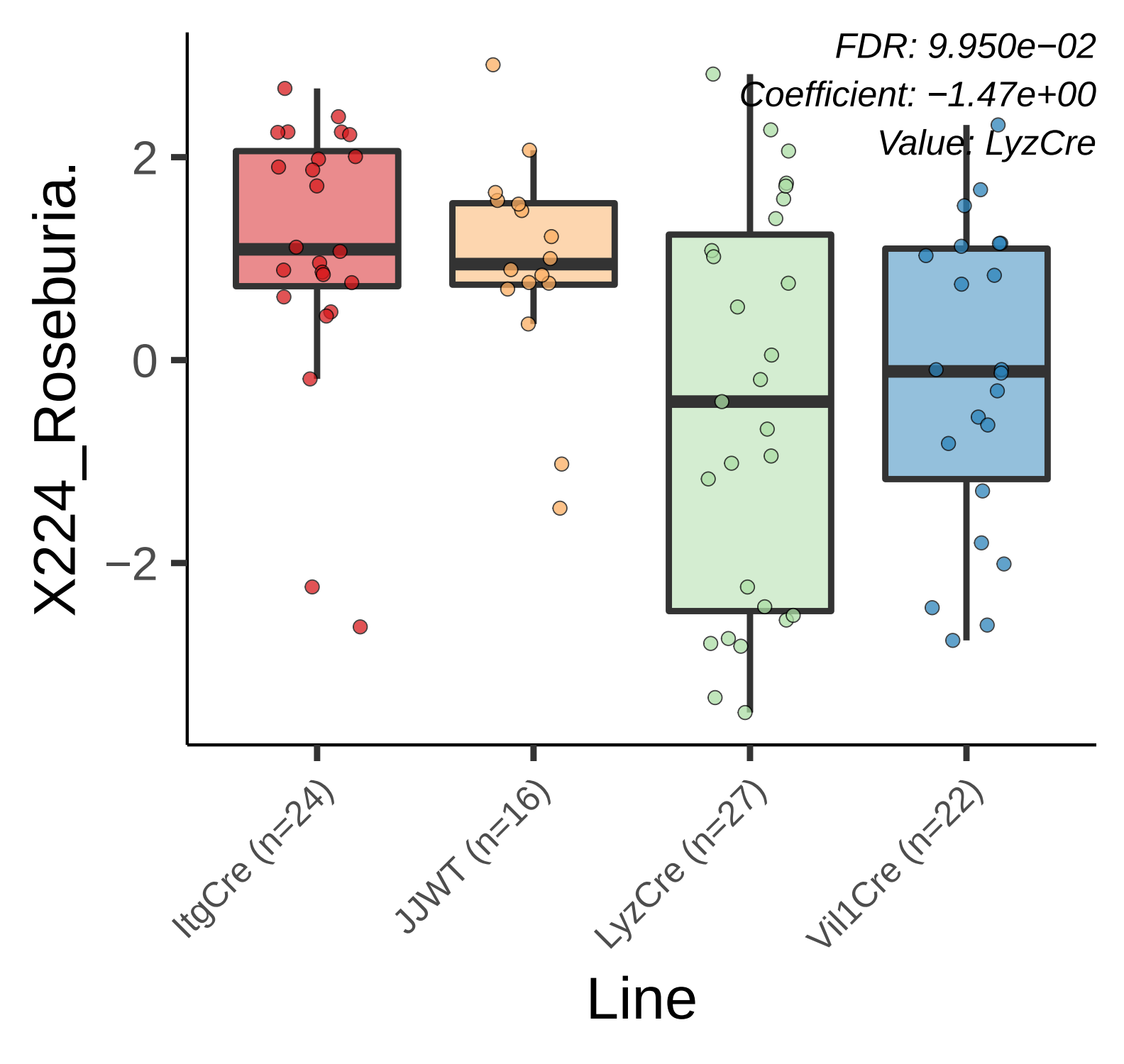

FDR: 8.842e−02
Coefficient: −1.80e+00
Value: JJWT
X194_Muribaculaceae..f.
2
1
0
−1
−2
−3
ItgCre (n=24)
JJWT (n=16)
LyzCre (n=27)
Vil1Cre (n=22)
Line

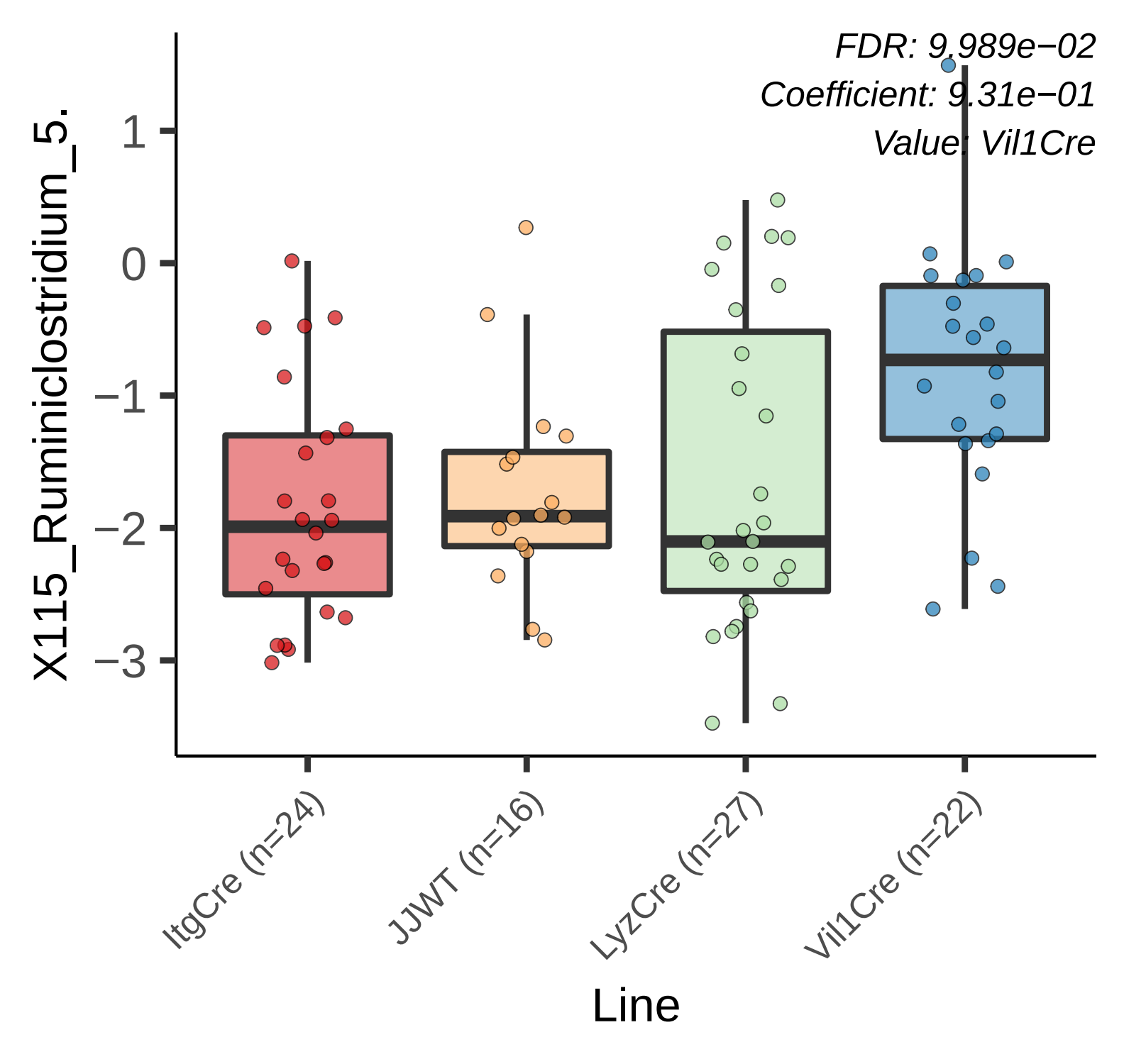

FDR: 9.142e−02
Coefficient: −2.12e+00
Value: LyzCre
X70_Lachnospiraceae_NK4A136_group.
4
2
0
−2
ItgCre (n=24)
JJWT (n=16)
LyzCre (n=27)
Vil1Cre (n=22)
Line

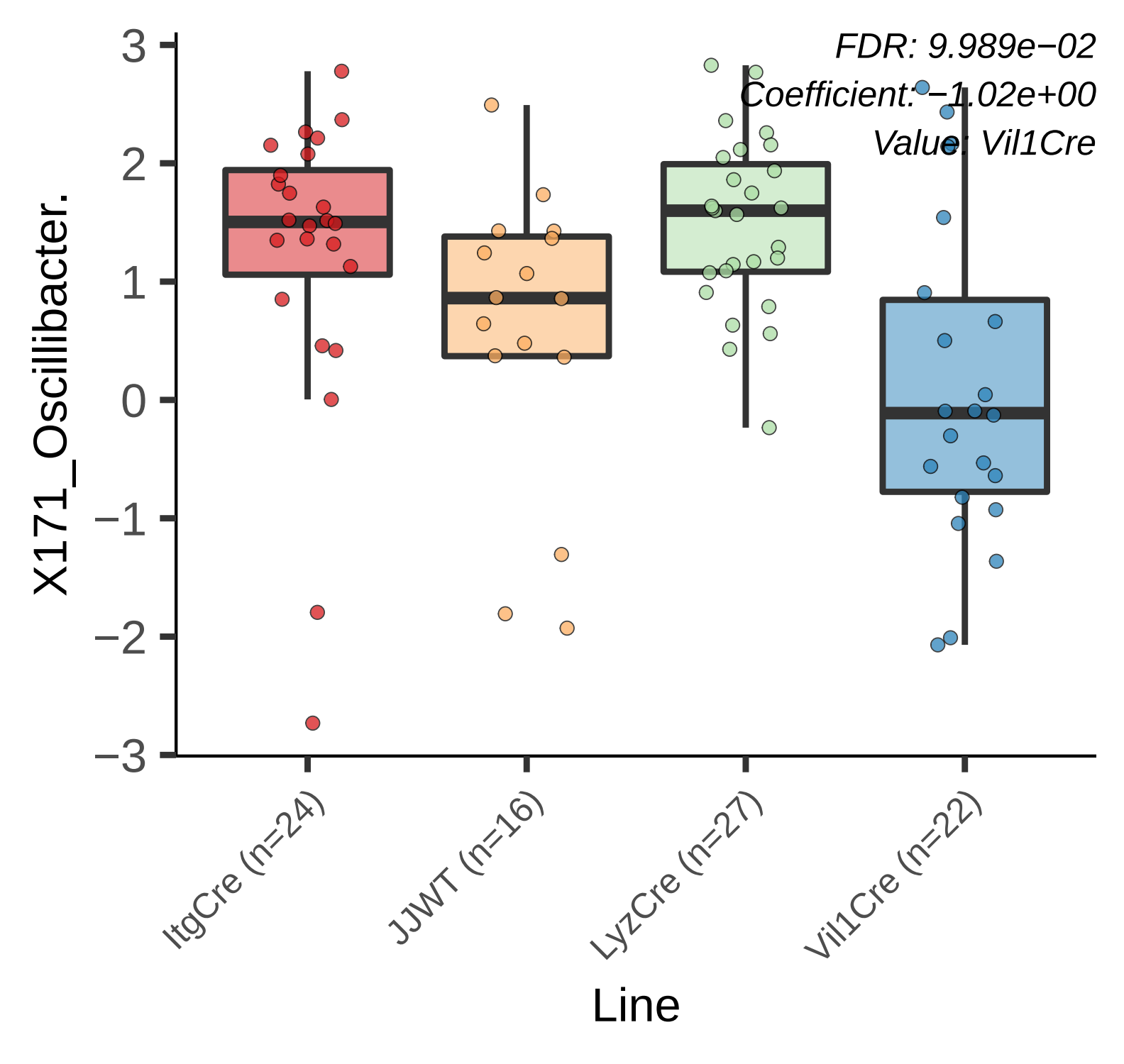

FDR: 9.266e−02
Coefficient: 7.67e−01
Value: Vil1Cre
X300_Ruminococcaceae..f.
0
−1
−2
−3
ItgCre (n=24)
JJWT (n=16)
LyzCre (n=27)
Vil1Cre (n=22)
Line

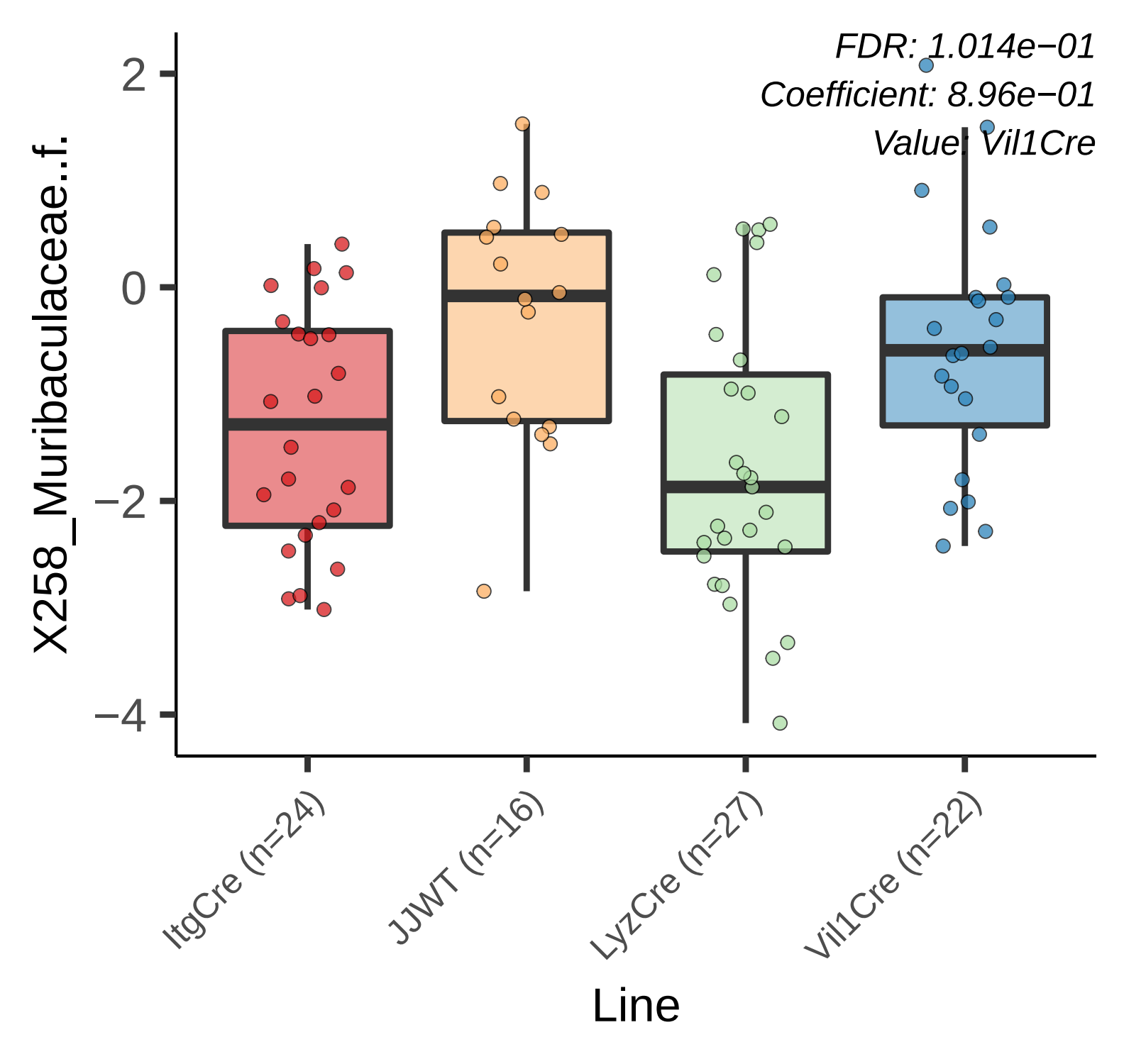

FDR: 9.420e−02
Coefficient: −1.47e+00
Value: JJWT
X117_Ruminiclostridium_5.
1
0
−1
−2
−3
ItgCre (n=24)
JJWT (n=16)
LyzCre (n=27)
Vil1Cre (n=22)
Line

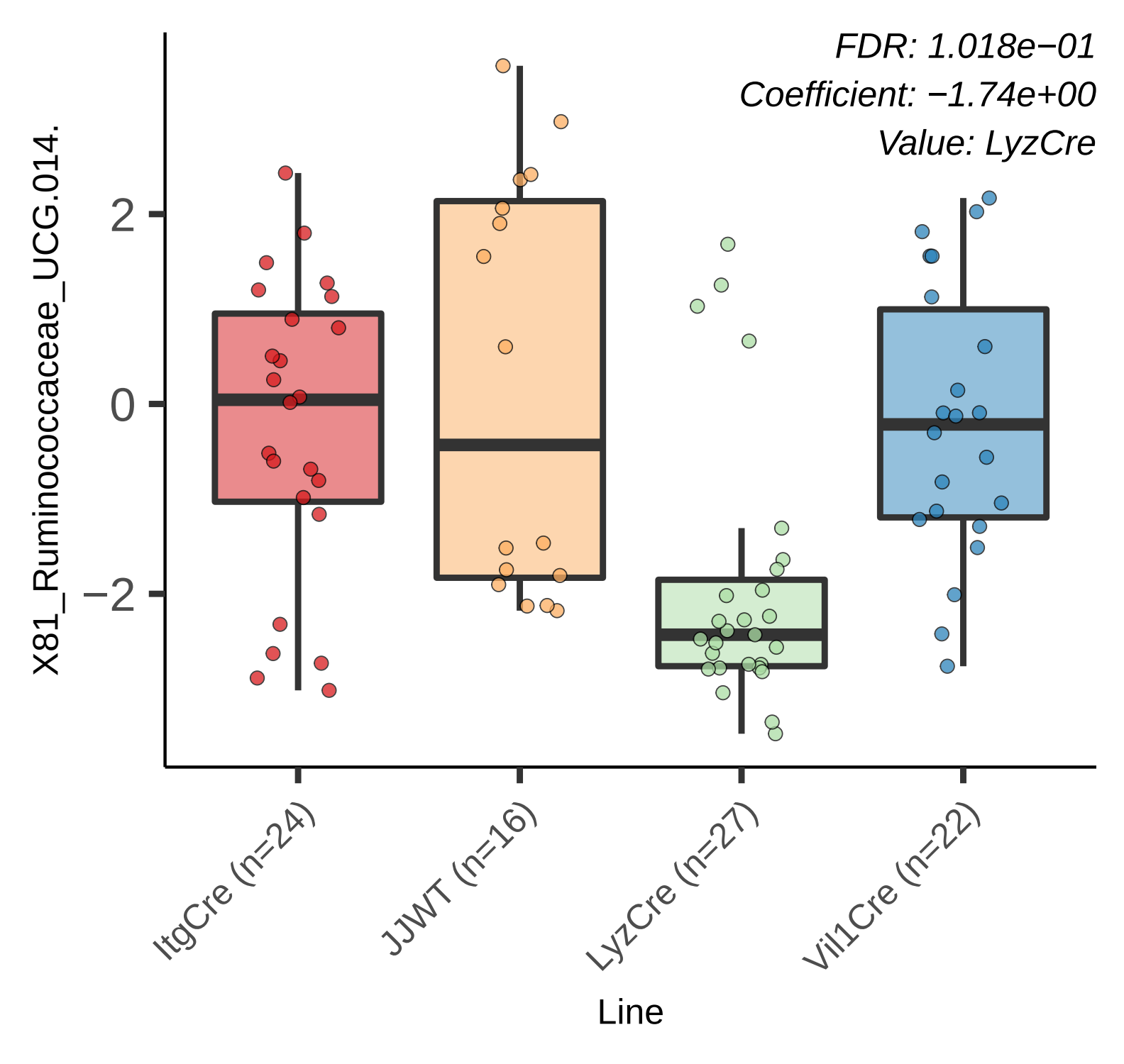

FDR: 9.906e−02
Coefficient: 3.08e+00
Value: JJWT
X18_Lachnoclostridium.
4
2
0
−2
ItgCre (n=24)
JJWT (n=16)
LyzCre (n=27)
Vil1Cre (n=22)
Line

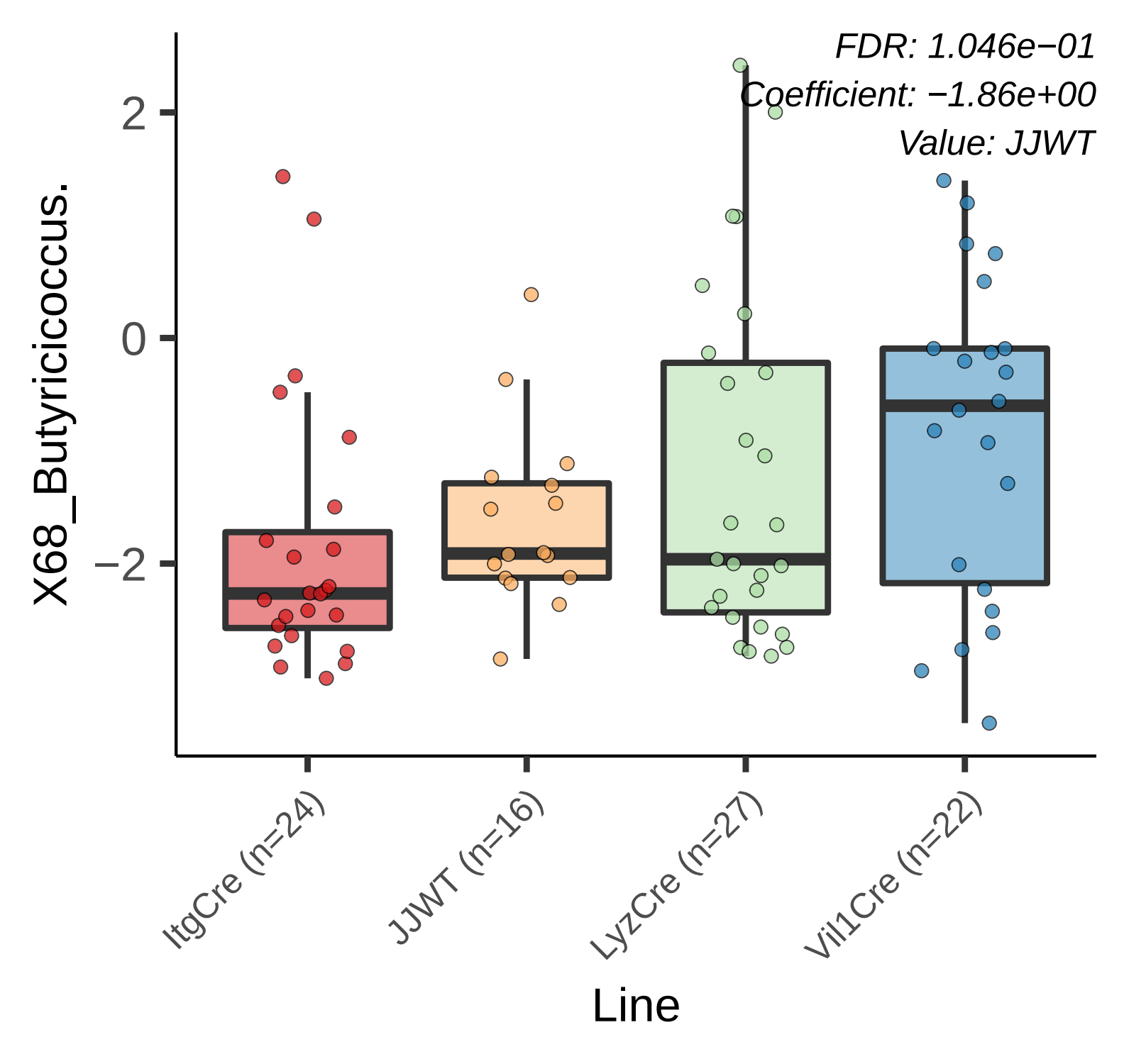

FDR: 9.950e−02
Coefficient: −1.47e+00
Value: LyzCre
X224_Roseburia.
2
0
−2
ItgCre (n=24)
JJWT (n=16)
LyzCre (n=27)
Vil1Cre (n=22)
Line

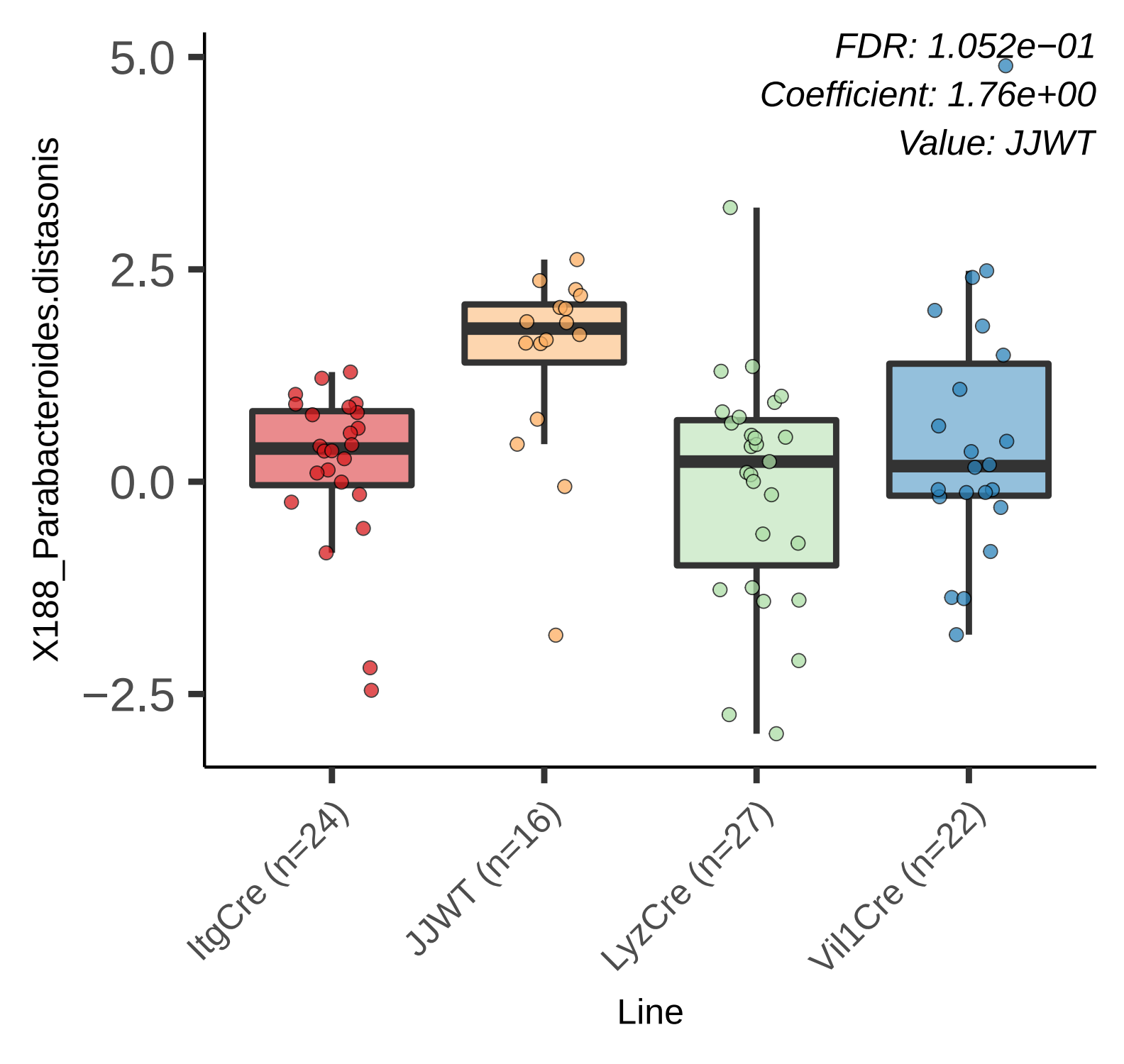

FDR: 9.989e−02
Coefficient: 9.31e−01
Value: Vil1Cre
X115_Ruminiclostridium_5.
1
0
−1
−2
−3
ItgCre (n=24)
JJWT (n=16)
LyzCre (n=27)
Vil1Cre (n=22)
Line

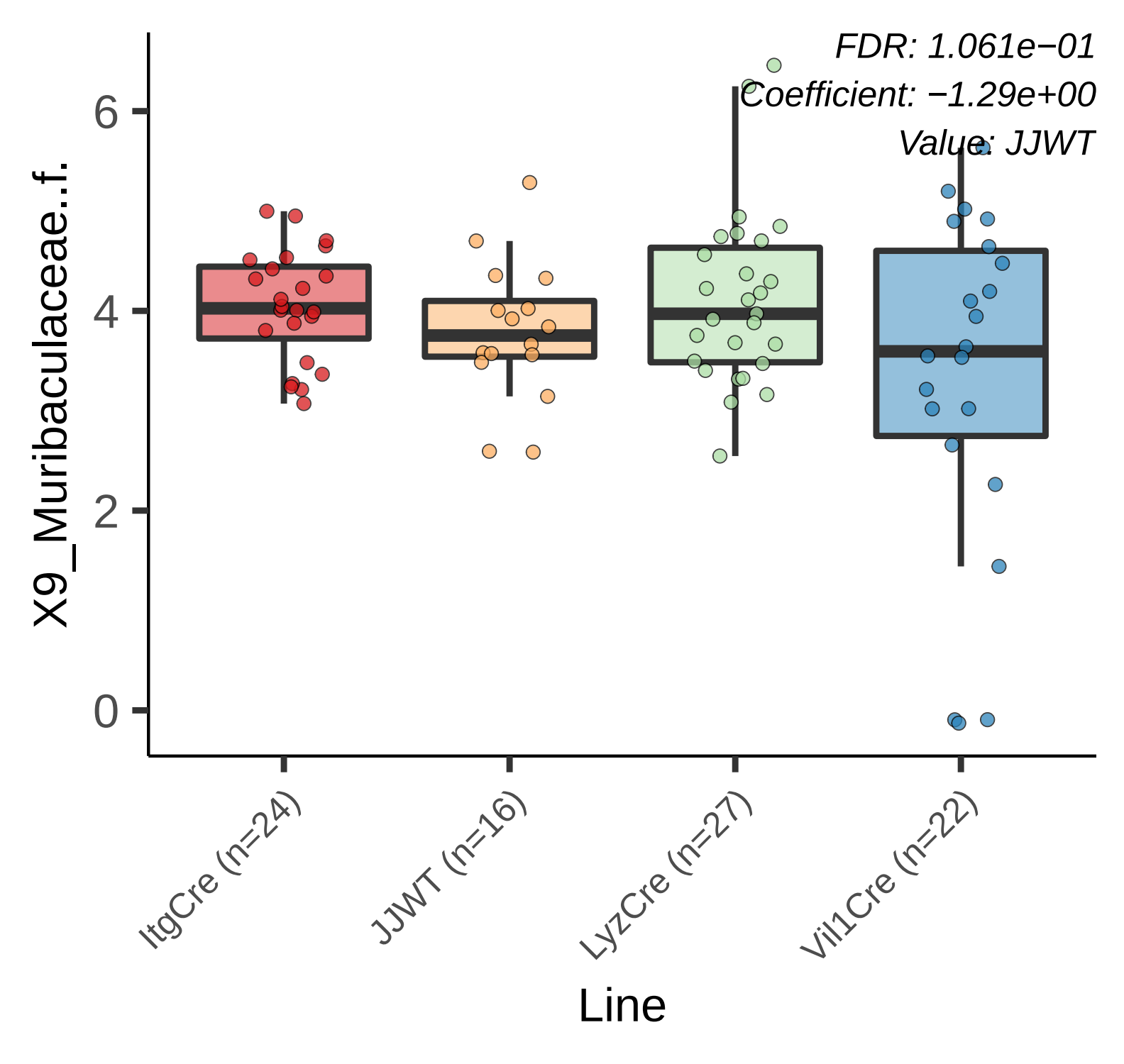

FDR: 9.989e−02
Coefficient: −1.02e+00
Value: Vil1Cre
X171_Oscillibacter.
3
2
1
0
−1
−2
−3
ItgCre (n=24)
JJWT (n=16)
LyzCre (n=27)
Vil1Cre (n=22)
Line

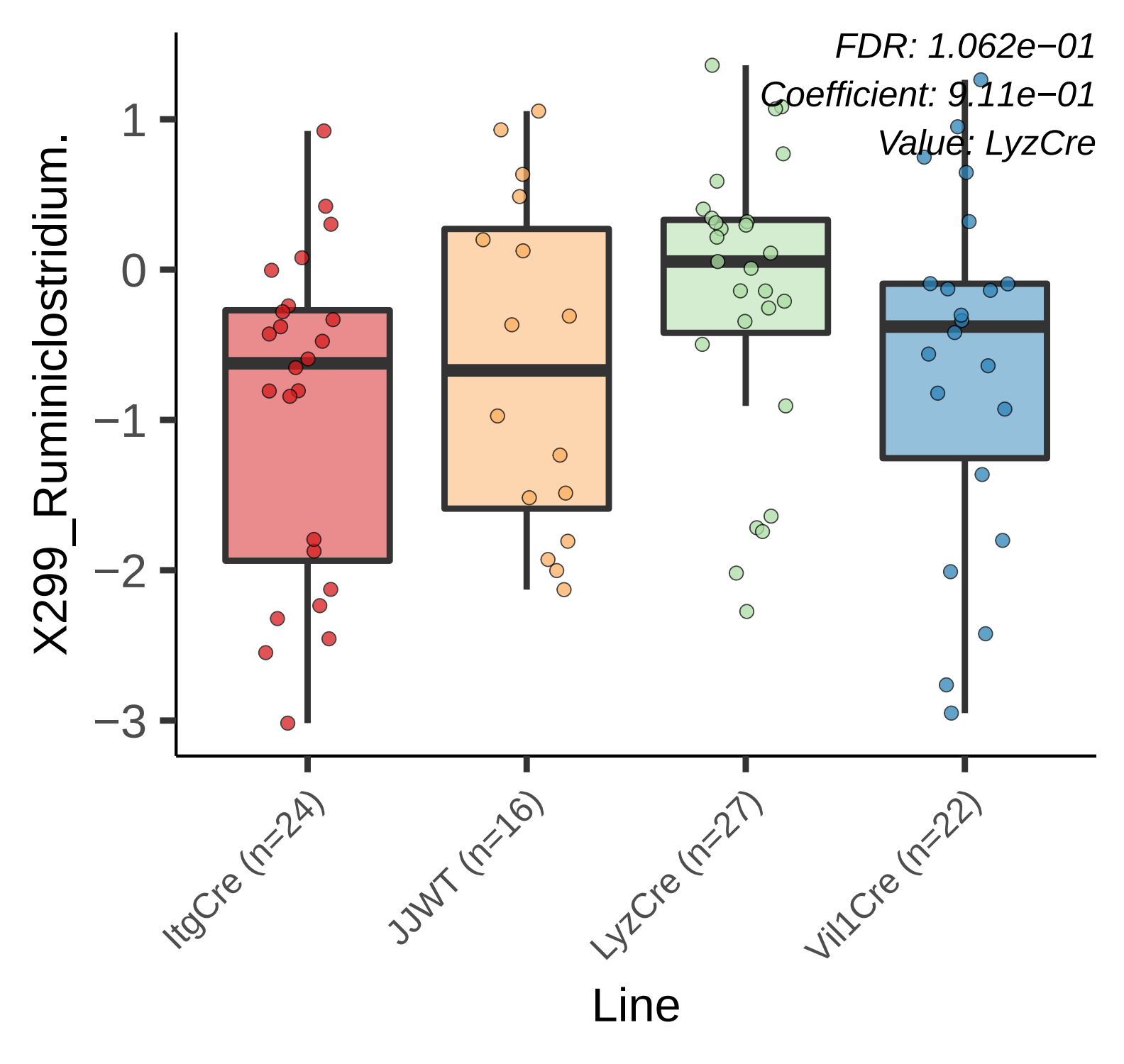

FDR: 1.014e−01
Coefficient: 8.96e−01
Value: Vil1Cre
X258_Muribaculaceae..f.
2
0
−2
−4
ItgCre (n=24)
JJWT (n=16)
LyzCre (n=27)
Vil1Cre (n=22)
Line

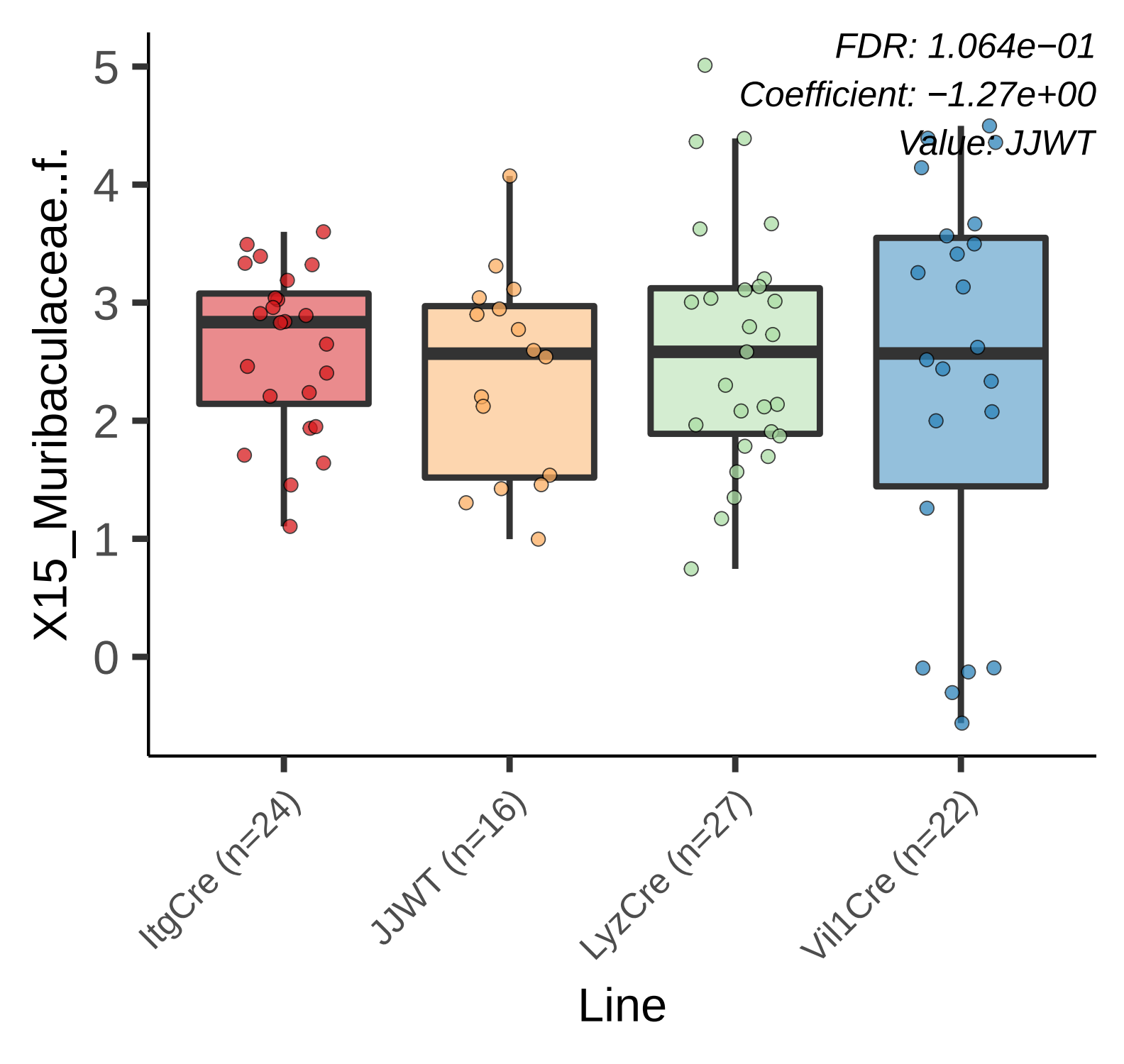

FDR: 1.018e−01
Coefficient: −1.74e+00
Value: LyzCre
X81_Ruminococcaceae_UCG.014.
2
0
−2
ItgCre (n=24)
JJWT (n=16)
LyzCre (n=27)
Vil1Cre (n=22)
Line

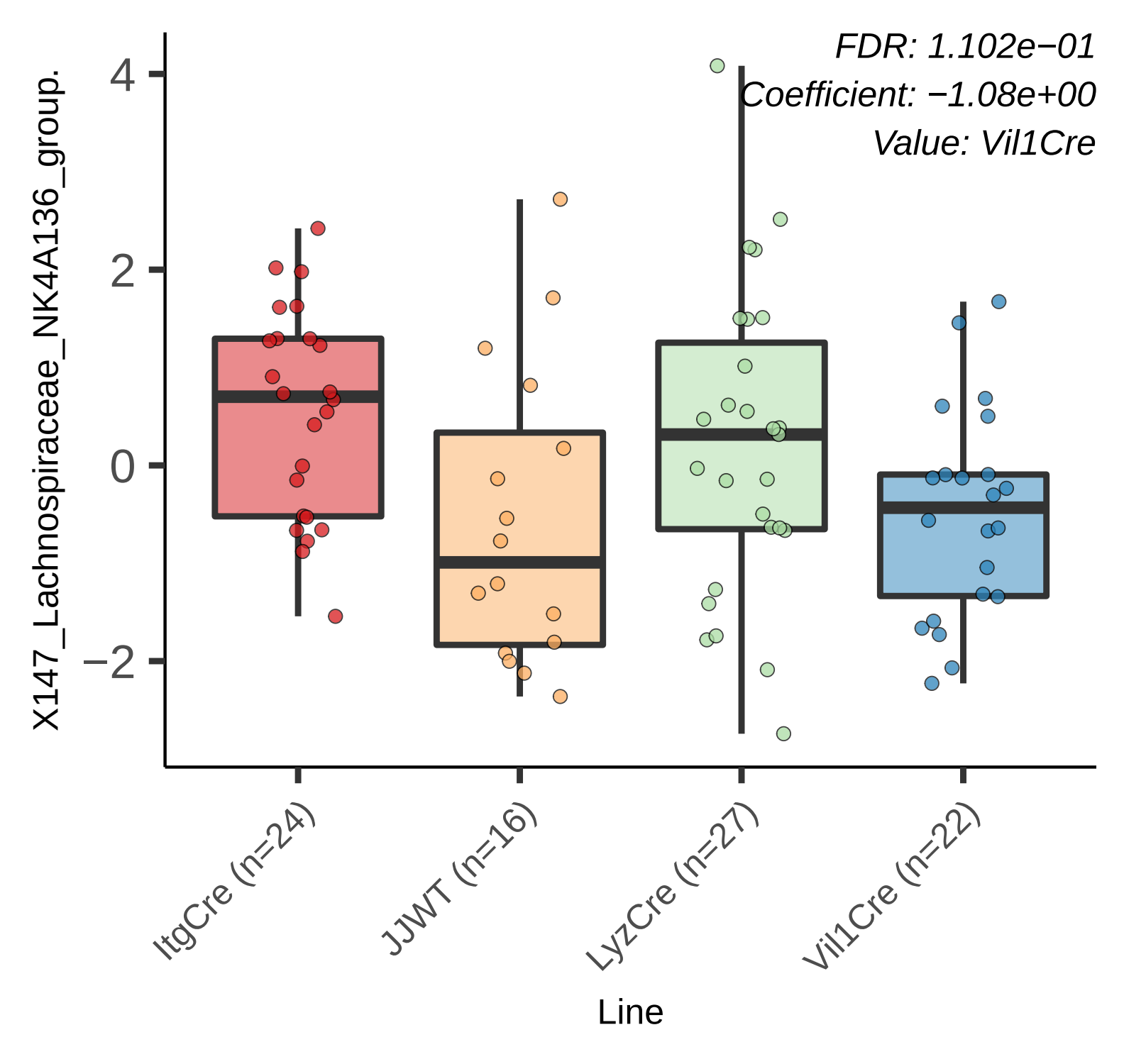

FDR: 1.046e−01
Coefficient: −1.86e+00
Value: JJWT
X68_Butyricicoccus.
2
0
−2
ItgCre (n=24)
JJWT (n=16)
LyzCre (n=27)
Vil1Cre (n=22)
Line

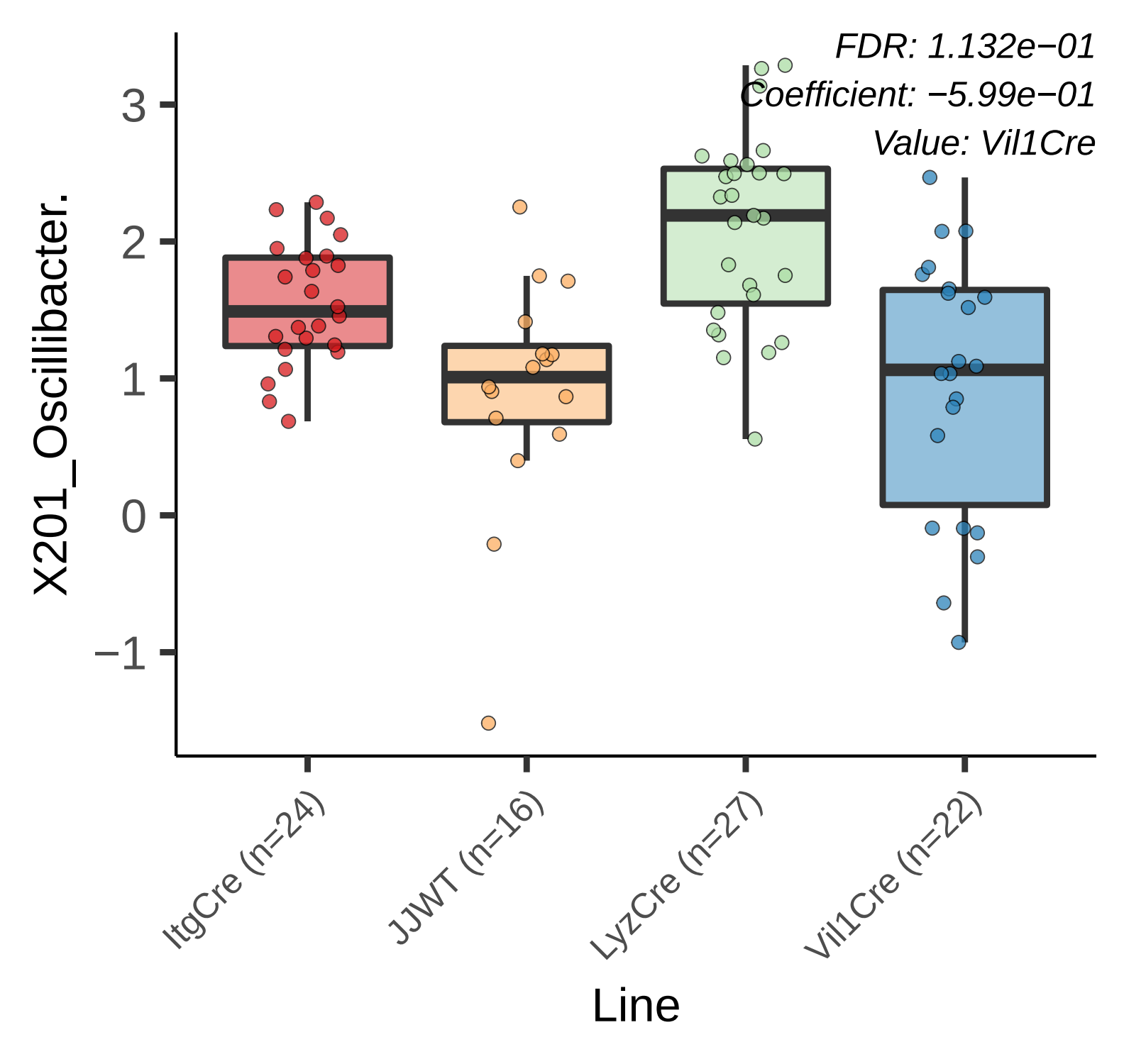

FDR: 1.052e−01
Coefficient: 1.76e+00
Value: JJWT
X188_Parabacteroides.distasonis
5.0
2.5
0.0
−2.5
ItgCre (n=24)
JJWT (n=16)
LyzCre (n=27)
Vil1Cre (n=22)
Line

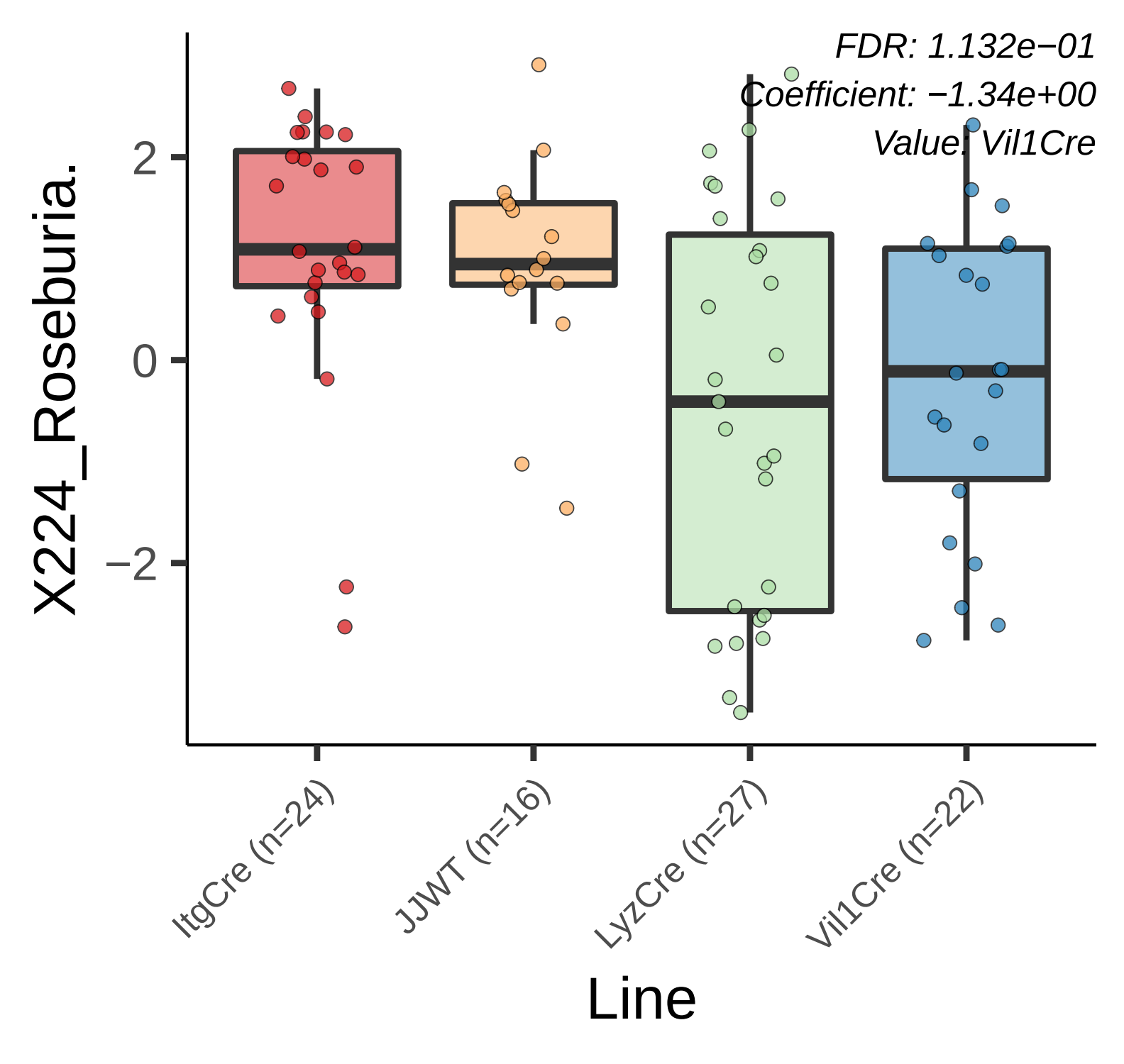

FDR: 1.061e−01
Coefficient: −1.29e+00
Value: JJWT
X9_Muribaculaceae..f.
6
4
2
0
ItgCre (n=24)
JJWT (n=16)
LyzCre (n=27)
Vil1Cre (n=22)
Line

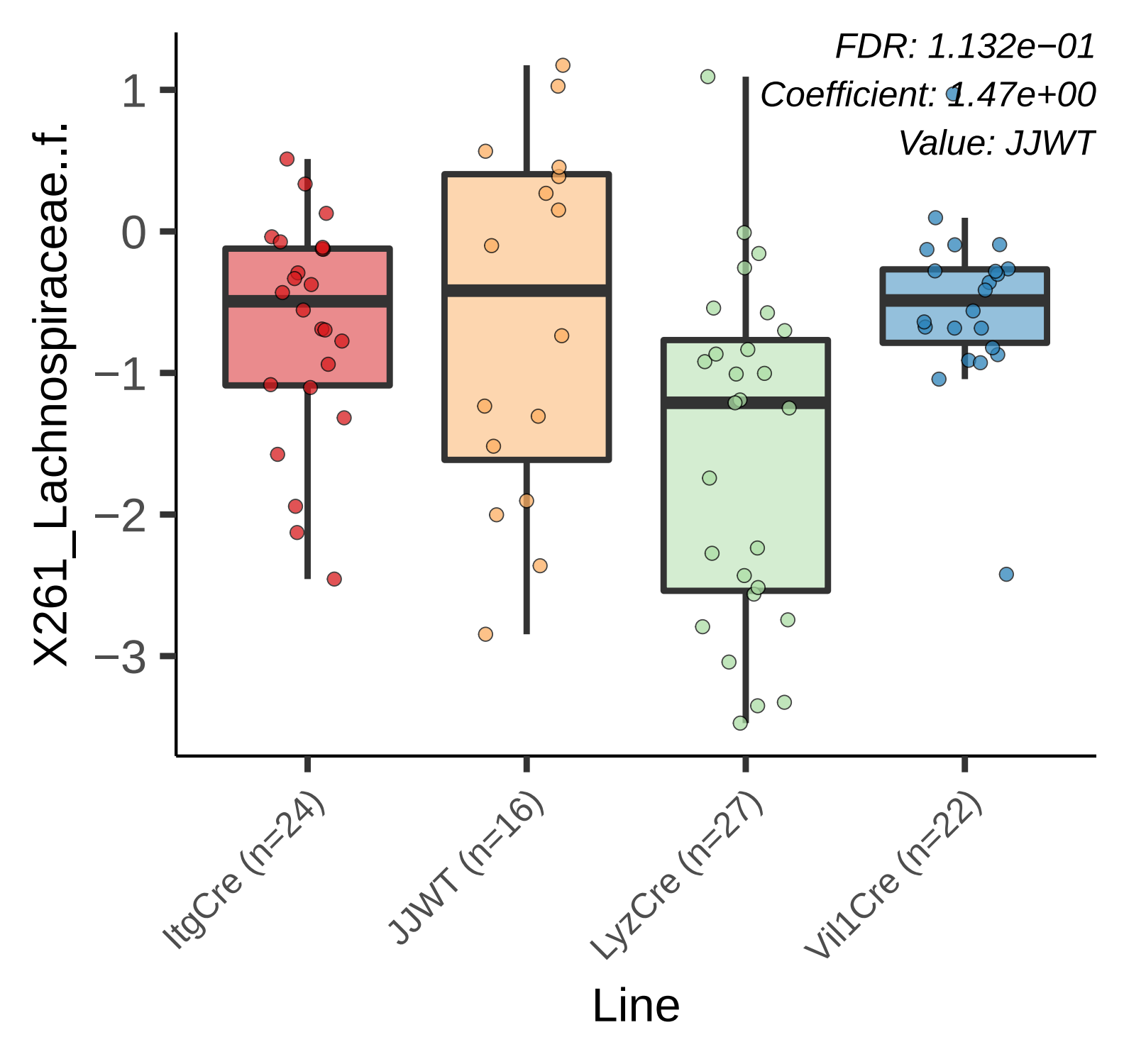

FDR: 1.062e−01
Coefficient: 9.11e−01
Value: LyzCre
X299_Ruminiclostridium.
1
0
−1
−2
−3
ItgCre (n=24)
JJWT (n=16)
LyzCre (n=27)
Vil1Cre (n=22)
Line

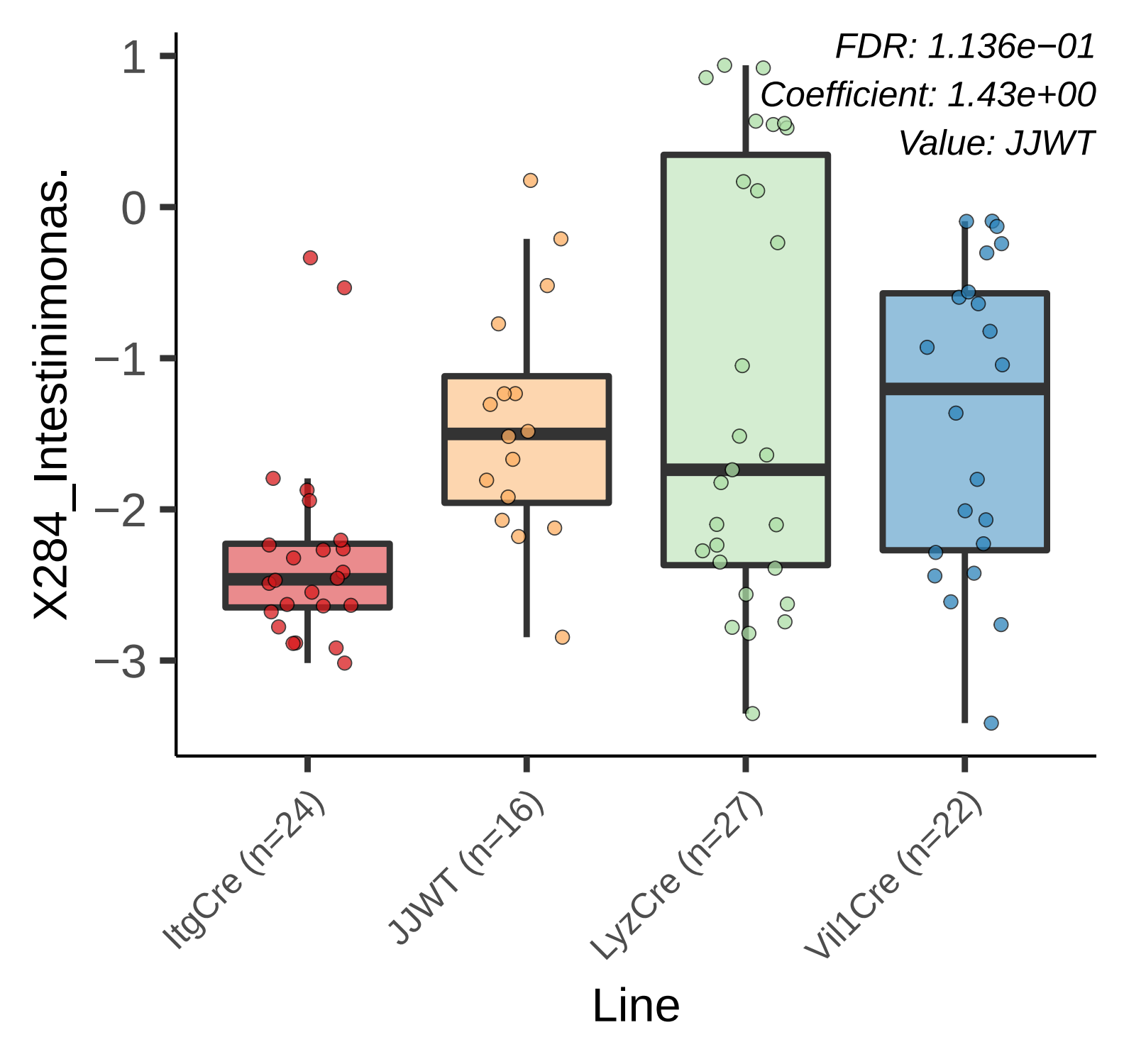

FDR: 1.064e−01
Coefficient: −1.27e+00
Value: JJWT
X15_Muribaculaceae..f.
5
4
3
2
1
0
ItgCre (n=24)
JJWT (n=16)
LyzCre (n=27)
Vil1Cre (n=22)
Line

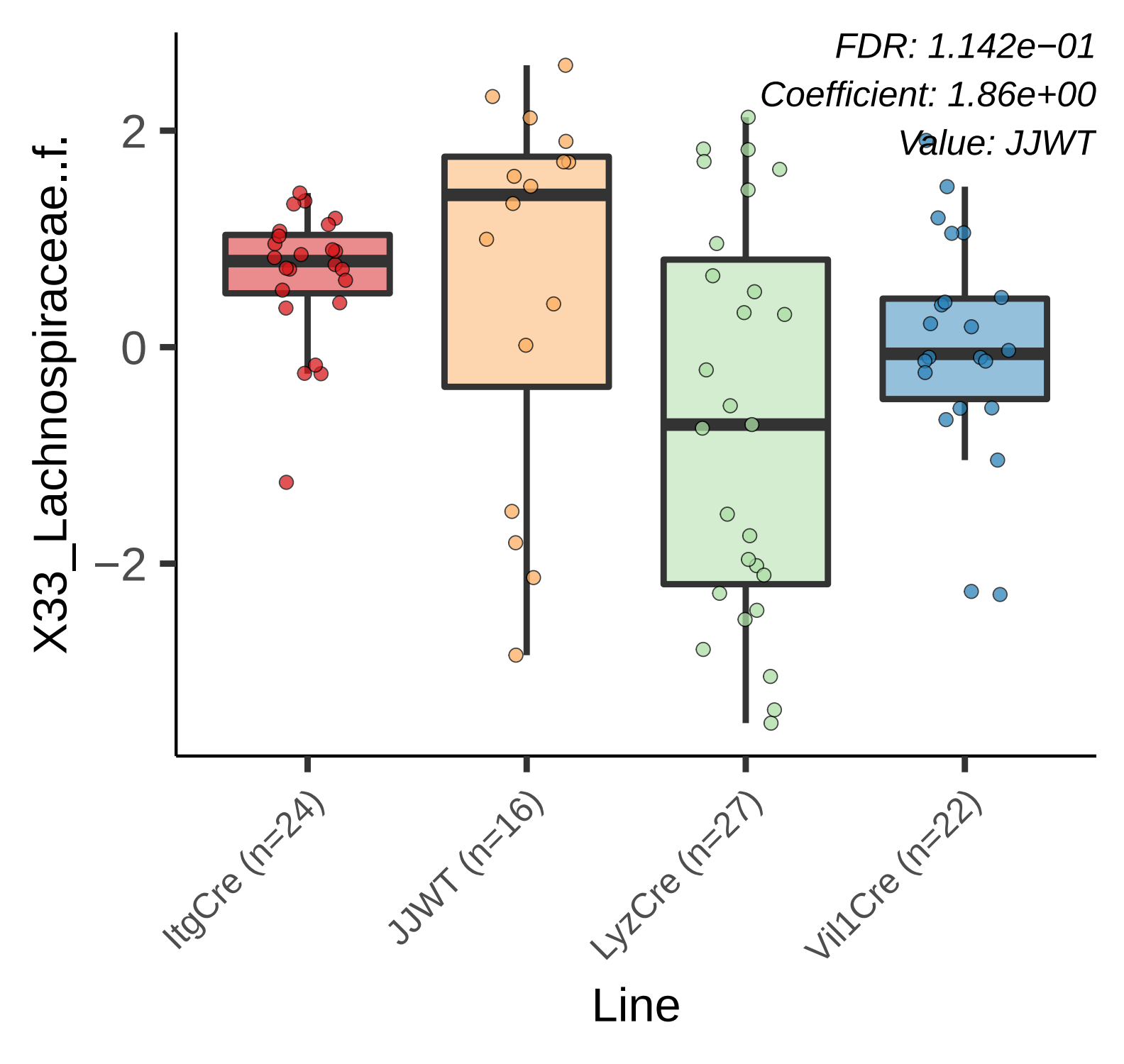

FDR: 1.102e−01
Coefficient: −1.08e+00
Value: Vil1Cre
X147_Lachnospiraceae_NK4A136_group.
4
2
0
−2
ItgCre (n=24)
JJWT (n=16)
LyzCre (n=27)
Vil1Cre (n=22)
Line

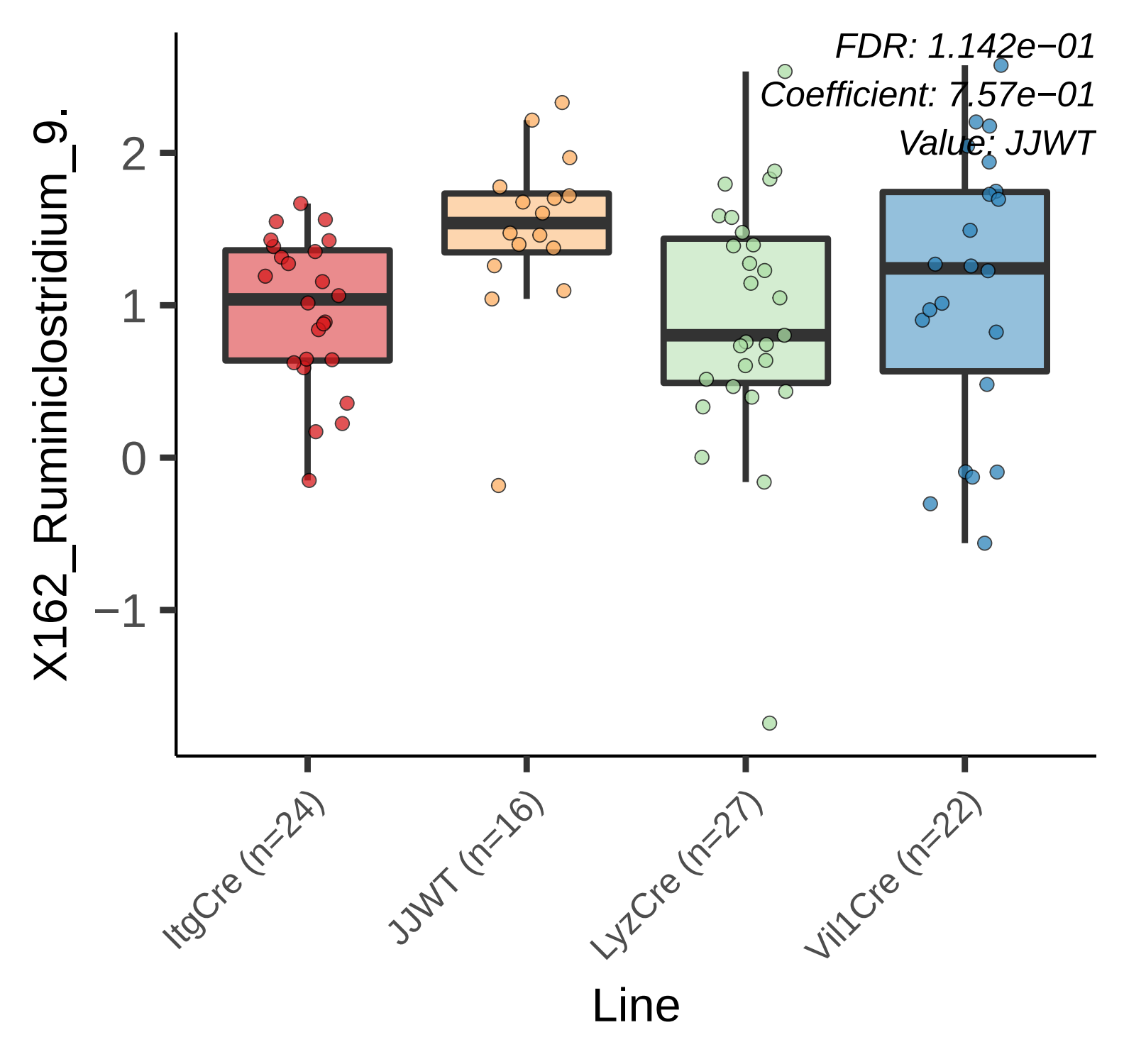

FDR: 1.132e−01
Coefficient: −5.99e−01
Value: Vil1Cre
X201_Oscillibacter.
3
2
1
0
−1
ItgCre (n=24)
JJWT (n=16)
LyzCre (n=27)
Vil1Cre (n=22)
Line

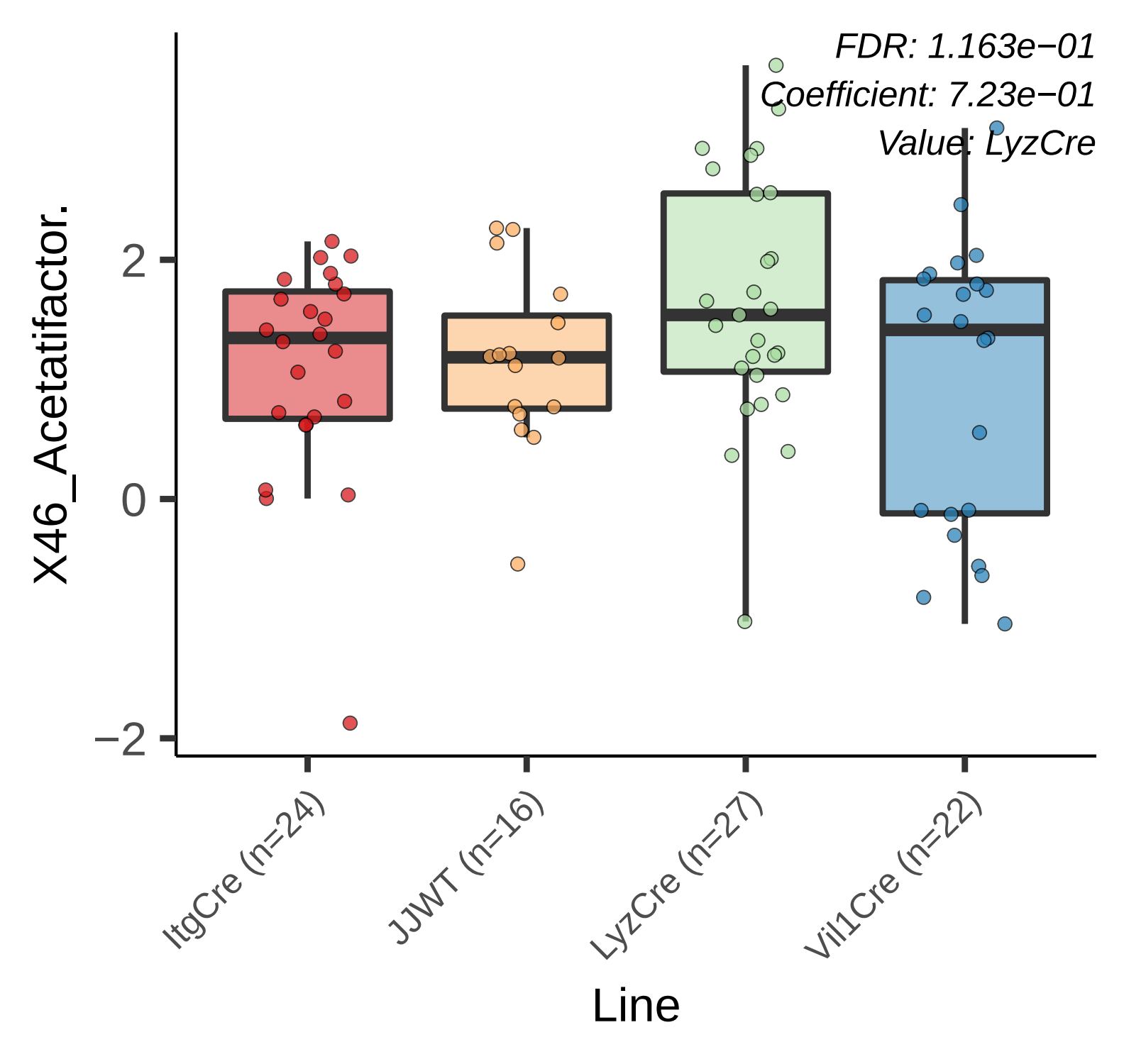

FDR: 1.132e−01
Coefficient: −1.34e+00
Value: Vil1Cre
X224_Roseburia.
2
0
−2
ItgCre (n=24)
JJWT (n=16)
LyzCre (n=27)
Vil1Cre (n=22)
Line

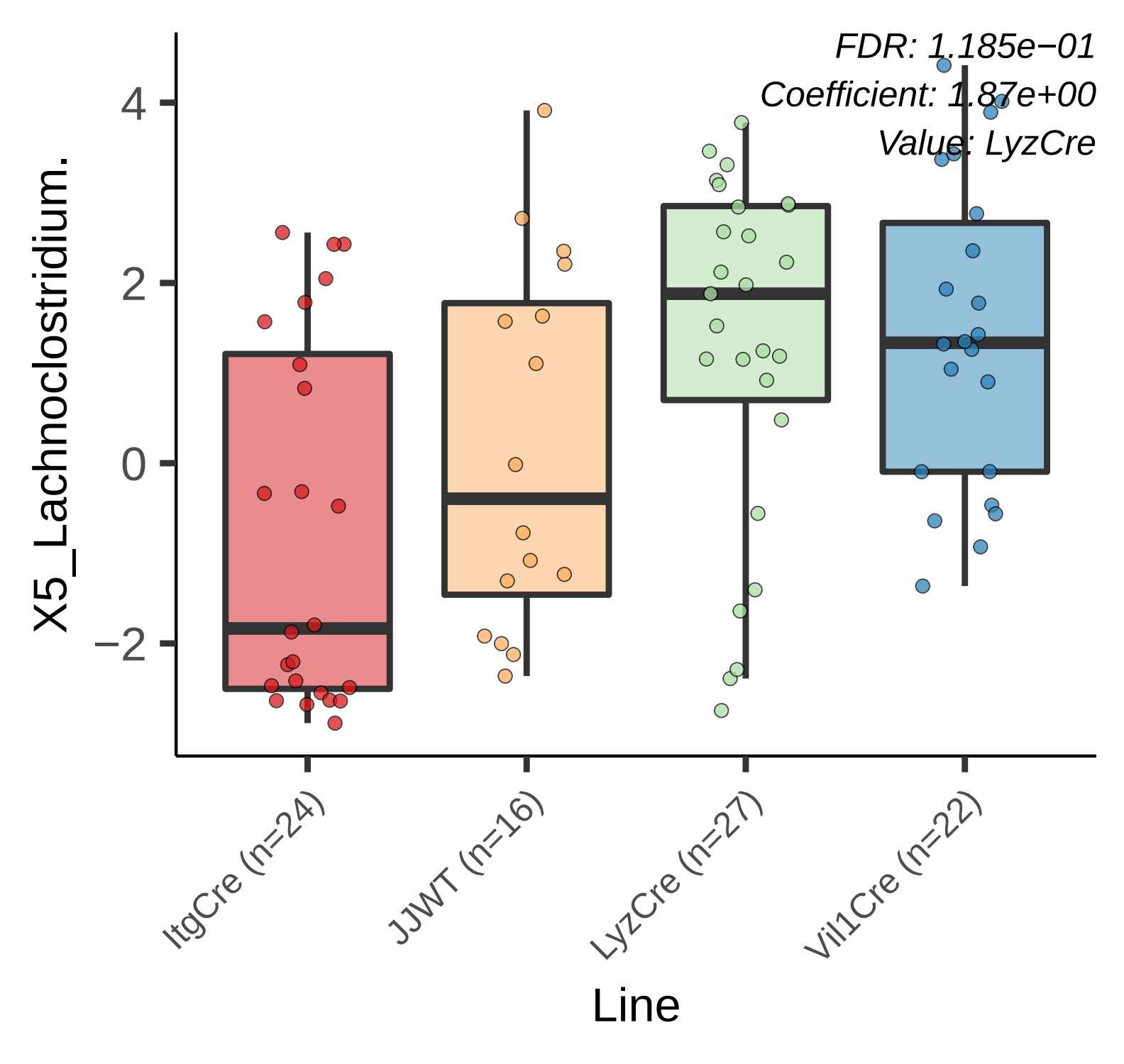

FDR: 1.132e−01
Coefficient: 1.47e+00
Value: JJWT
X261_Lachnospiraceae.f.
1
0
−1
−2
−3
ItgCre (n=24)
JJWT (n=16)
LyzCre (n=27)
Vil1Cre (n=22)
Line

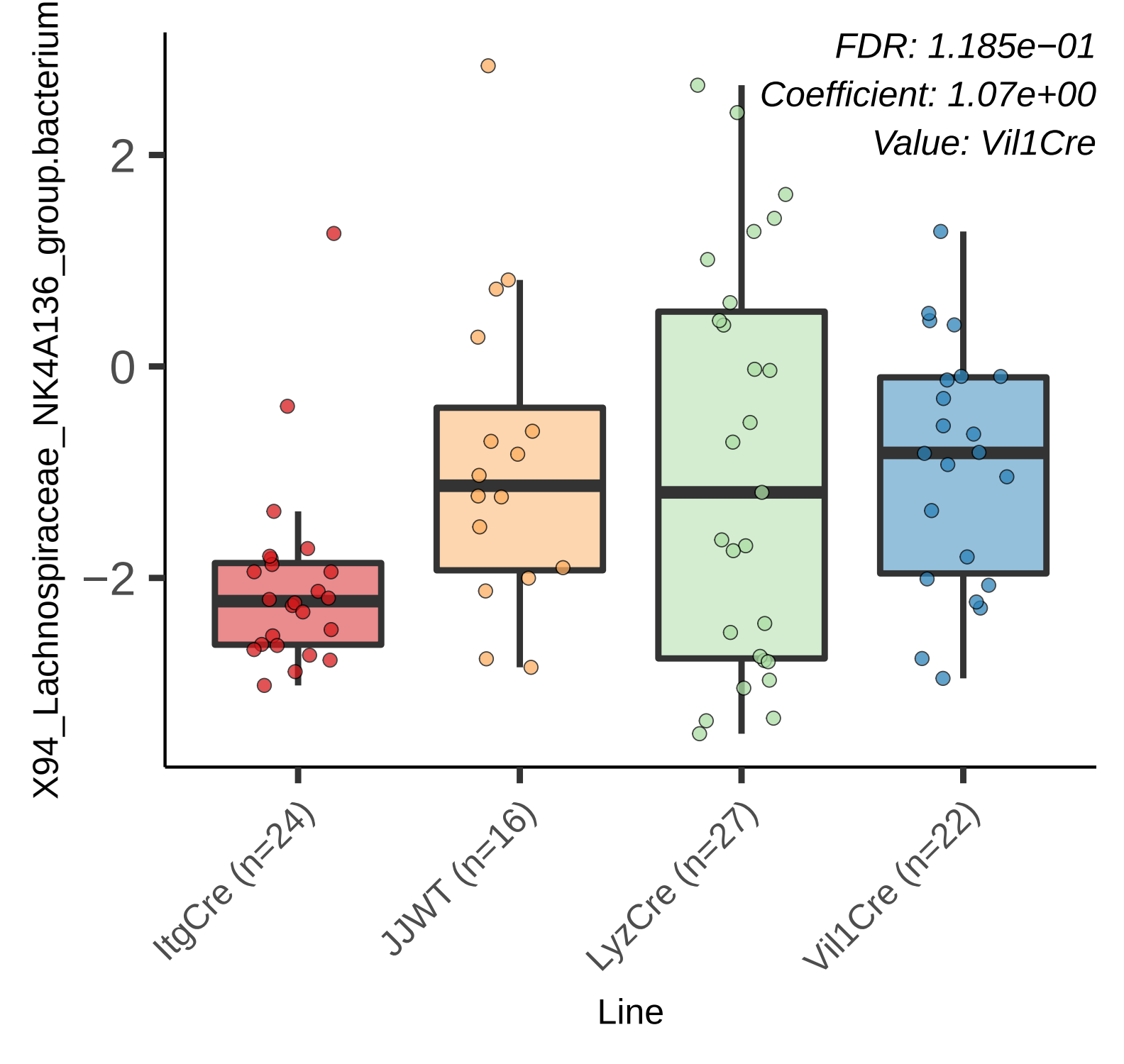

FDR: 1.136e−01
Coefficient: 1.43e+00
Value: JJWT
X284_Intestinimonas.
1
0
−1
−2
−3
ItgCre (n=24)
JJWT (n=16)
LyzCre (n=27)
Vil1Cre (n=22)
Line

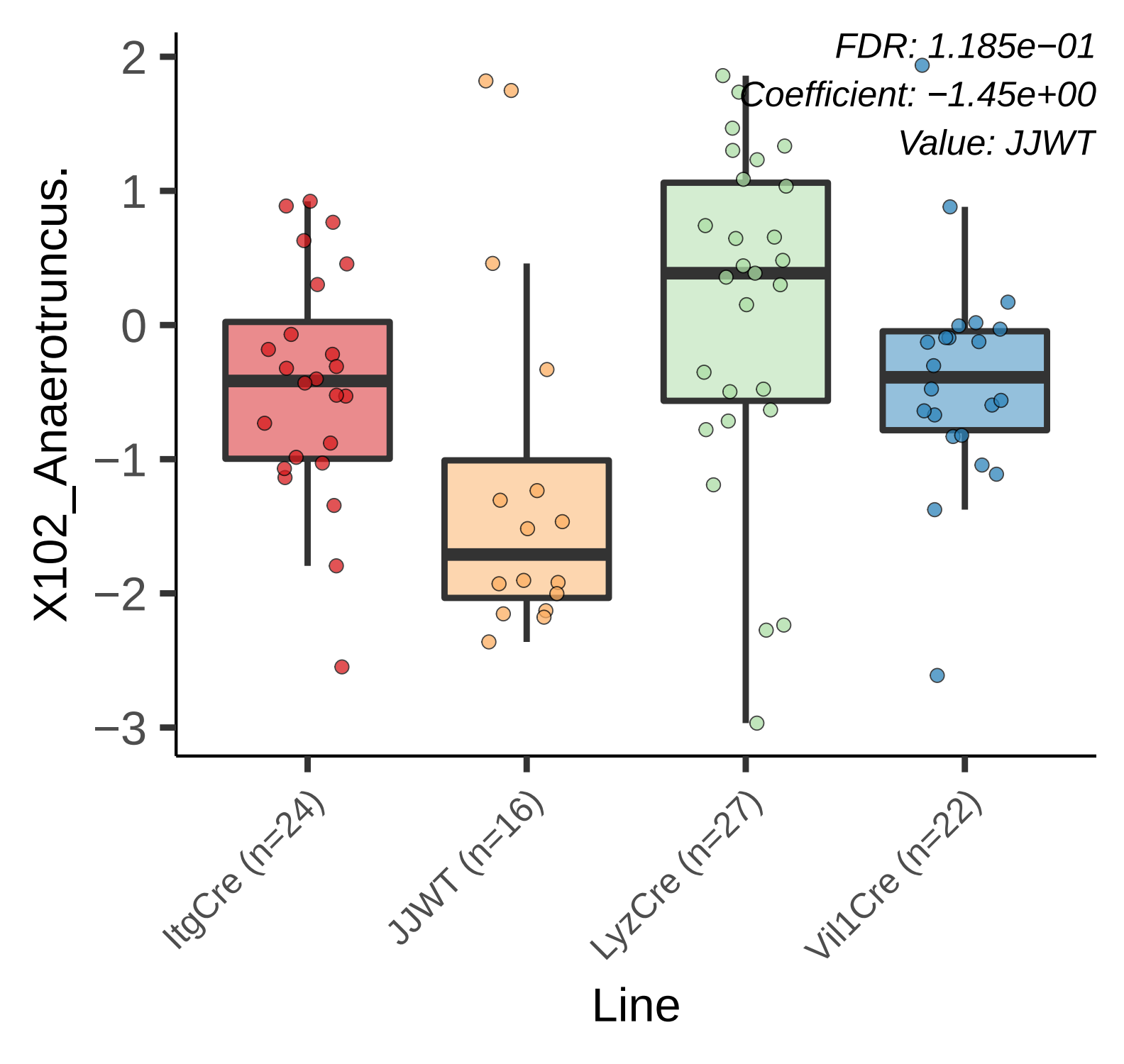

FDR: 1.142e−01
Coefficient: 1.86e+00
Value: JJWT
X33_Lachnospiraceae..f.
2
0
−2
ItgCre (n=24)
JJWT (n=16)
LyzCre (n=27)
Vil1Cre (n=22)
Line

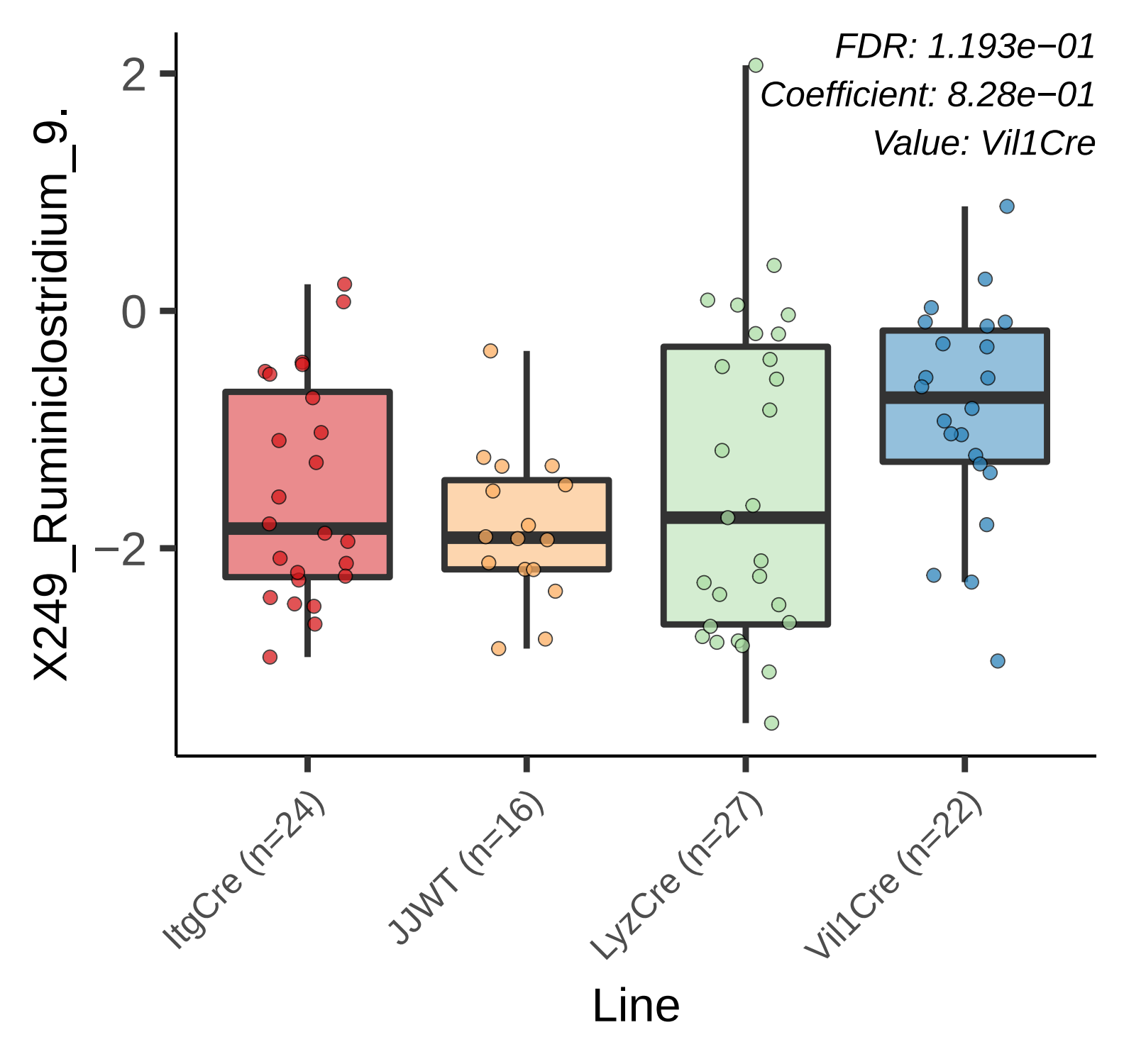

FDR: 1.142e−01
Coefficient: 7.57e−01
Value: JJWT
X162_Ruminiclostridium_9.
2
1
0
−1
ItgCre (n=24)
JJWT (n=16)
LyzCre (n=27)
Vil1Cre (n=22)
Line

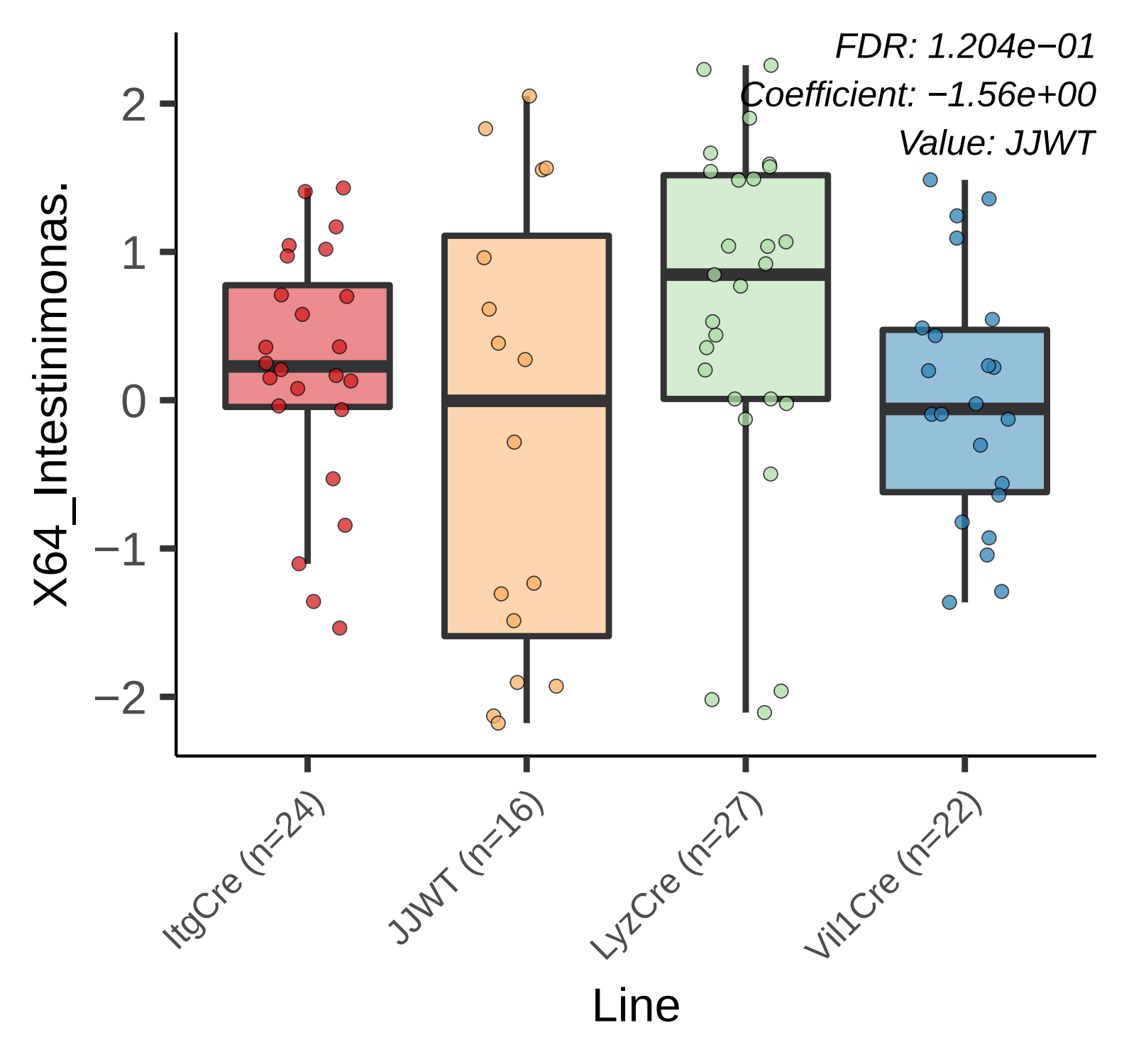

FDR: 1.163e−01
Coefficient: 7.23e−01
Value: LyzCre
X46_Acetatifactor.
2
0
−2
ItgCre (n=24)
JJWT (n=16)
LyzCre (n=27)
Vil1Cre (n=22)
Line

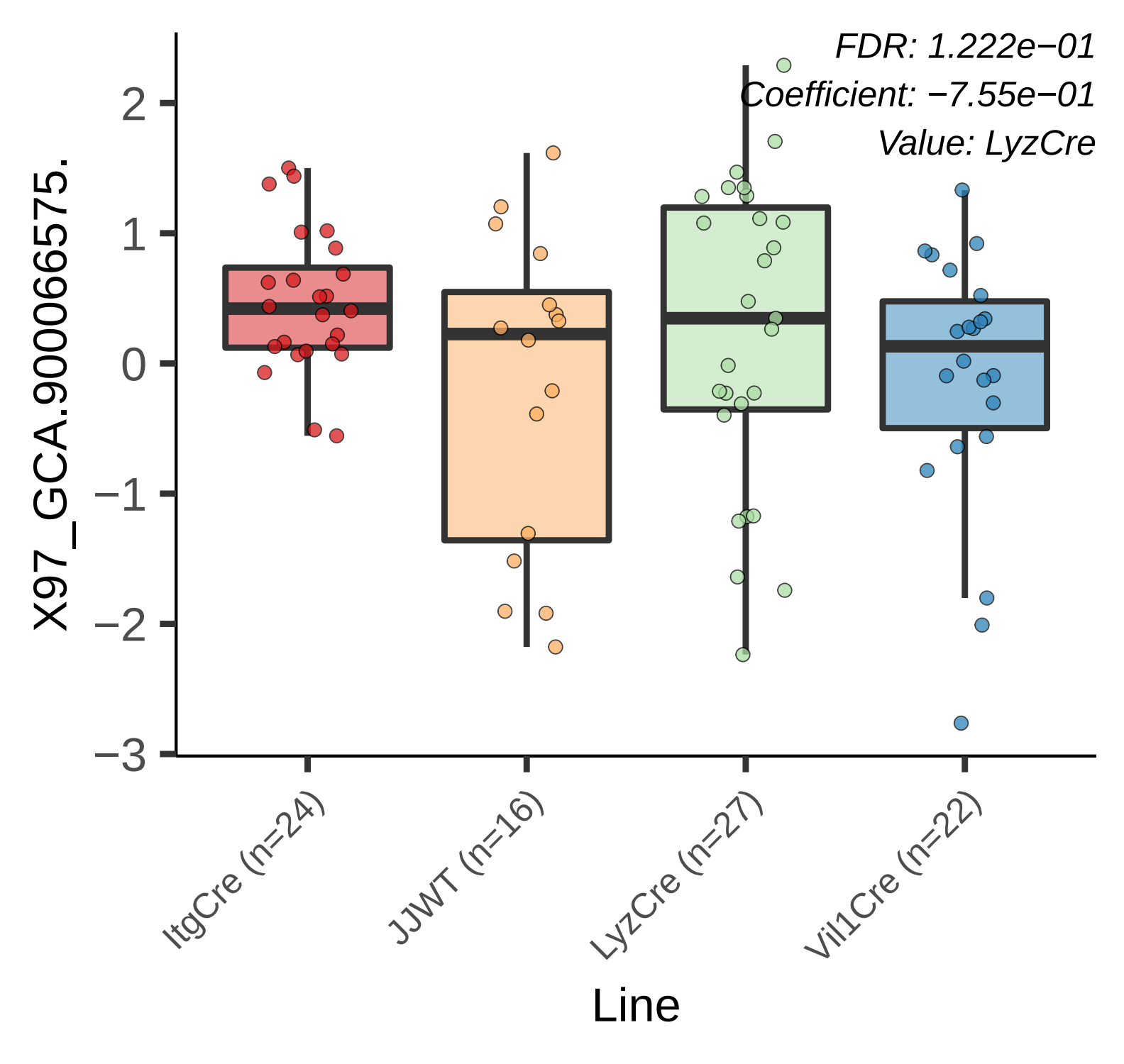

FDR: 1.185e−01
Coefficient: 1.87e+00
Value: LyzCre
X5_Lachnoclostridium.
4
2
0
−2
ItgCre (n=24)
JJWT (n=16)
LyzCre (n=27)
Vil1Cre (n=22)
Line

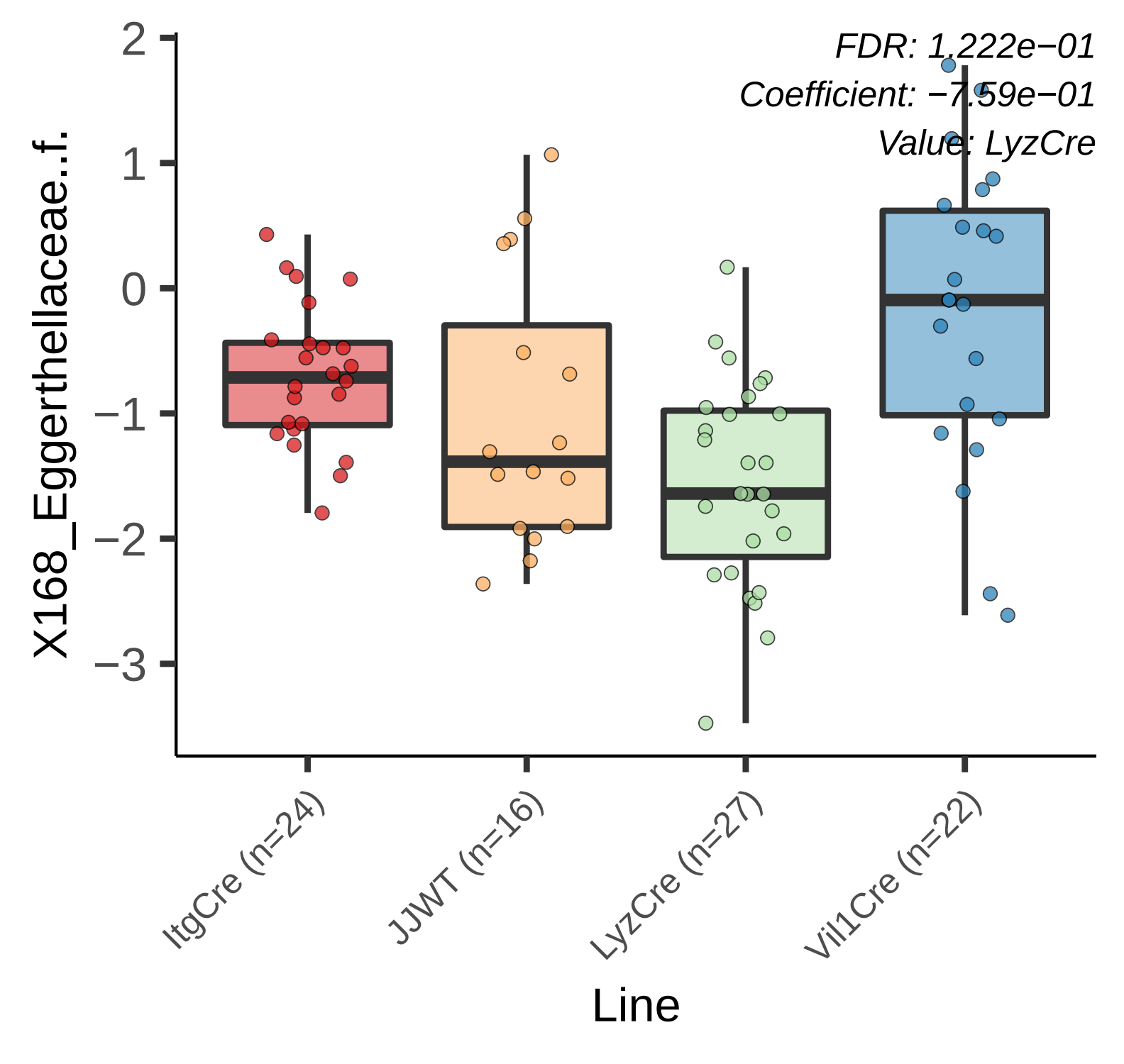

FDR: 1.185e−01
Coefficient: 1.07e+00
Value: Vil1Cre
X94_Lachnospiraceae_NK4A136_group.bacterium
2
0
−2
ItgCre (n=24)
JJWT (n=16)
LyzCre (n=27)
Vil1Cre (n=22)
Line

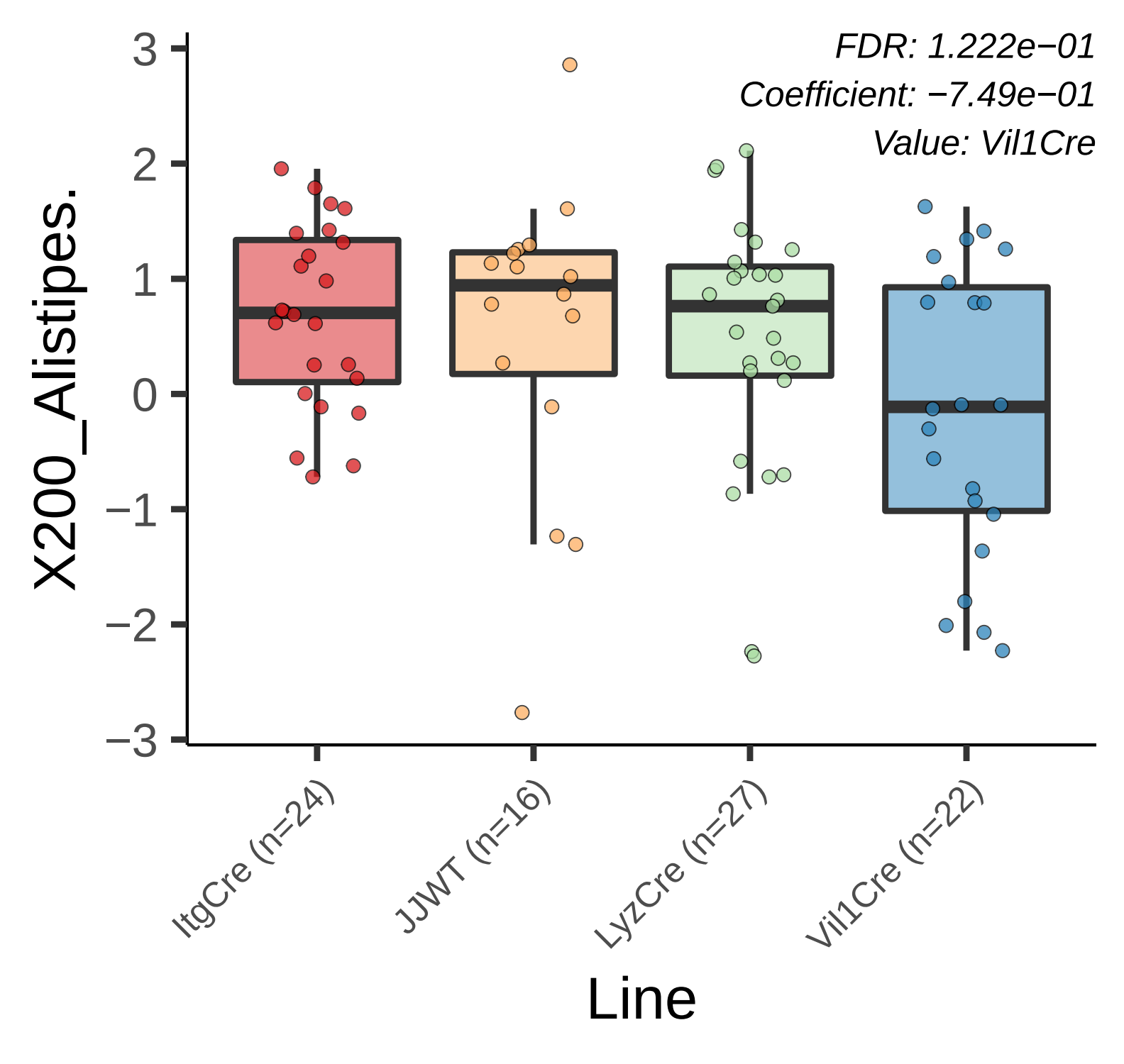

FDR: 1.185e−01
Coefficient: −1.45e+00
Value: JJWT
X102_Anaerotruncus.
2
1
0
−1
−2
−3
ItgCre (n=24)
JJWT (n=16)
LyzCre (n=27)
Vil1Cre (n=22)
Line

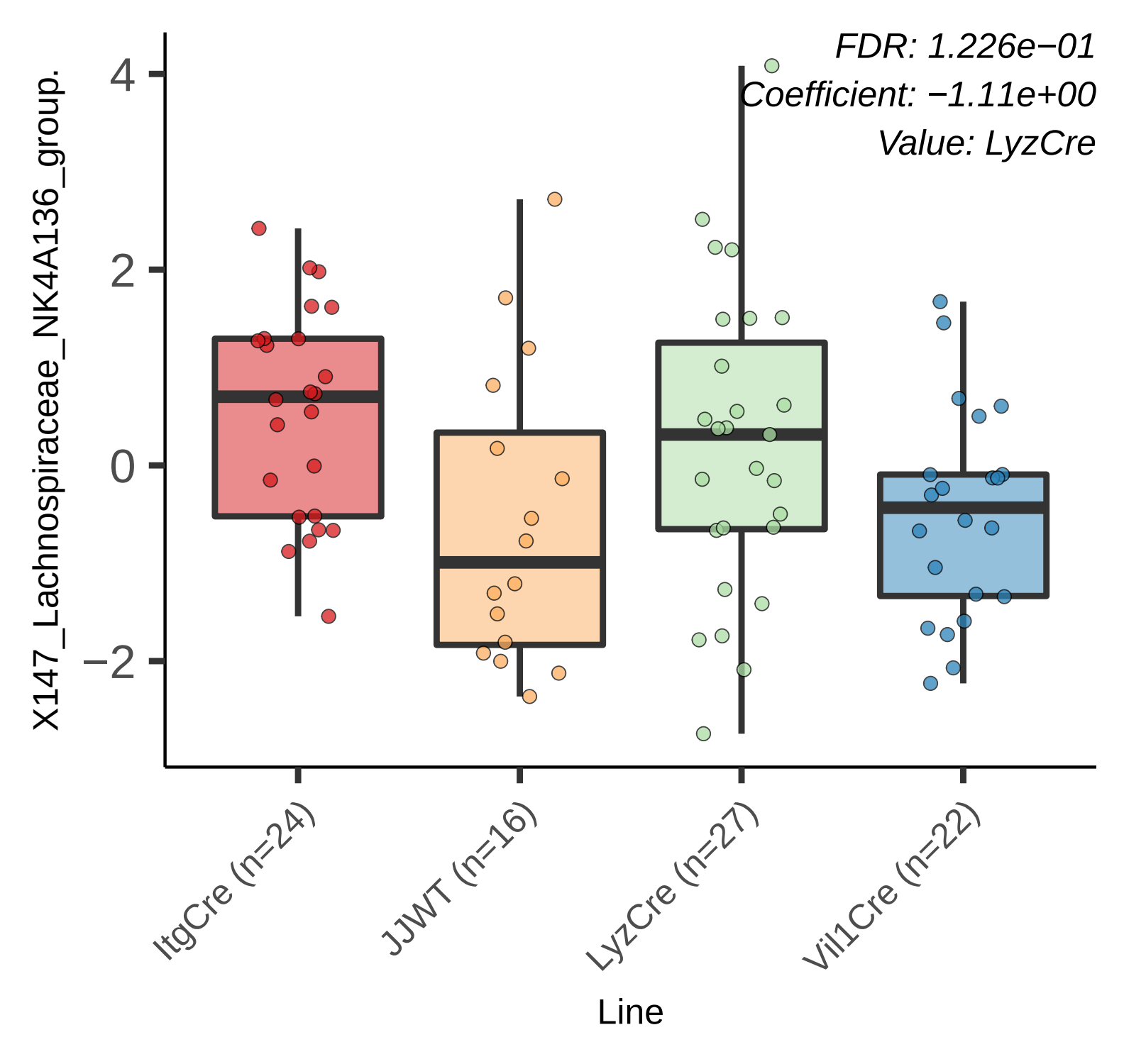

FDR: 1.193e−01
Coefficient: 8.28e−01
Value: Vil1Cre
X249_Ruminiclostridium_9.
2
0
−2
ItgCre (n=24)
JJWT (n=16)
LyzCre (n=27)
Vil1Cre (n=22)
Line

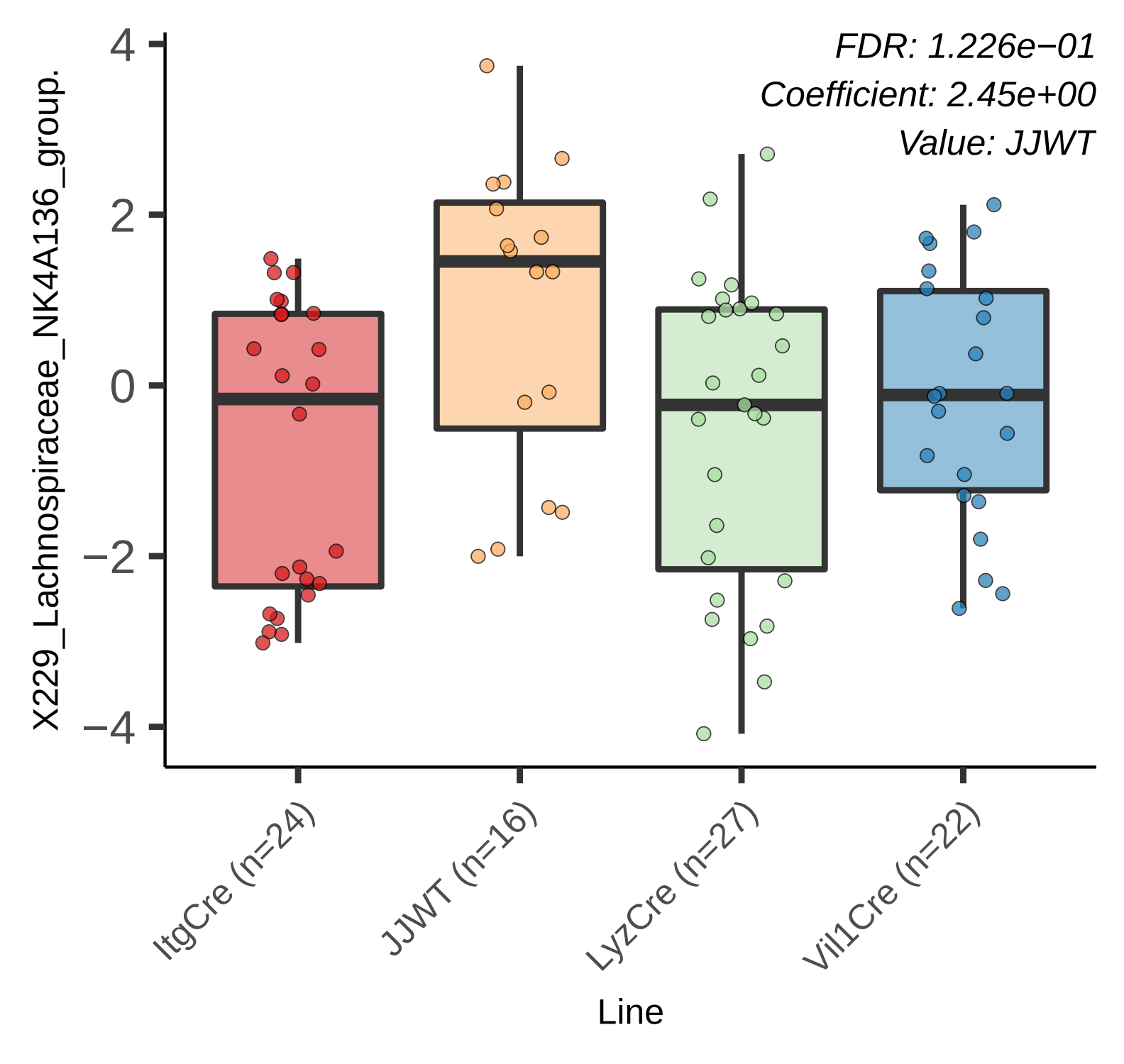

FDR: 1.204e−01
Coefficient: −1.56e+00
Value: JJWT
X64_Intestinimonas.
2
1
0
−1
−2
ItgCre (n=24)
JJWT (n=16)
LyzCre (n=27)
Vil1Cre (n=22)
Line

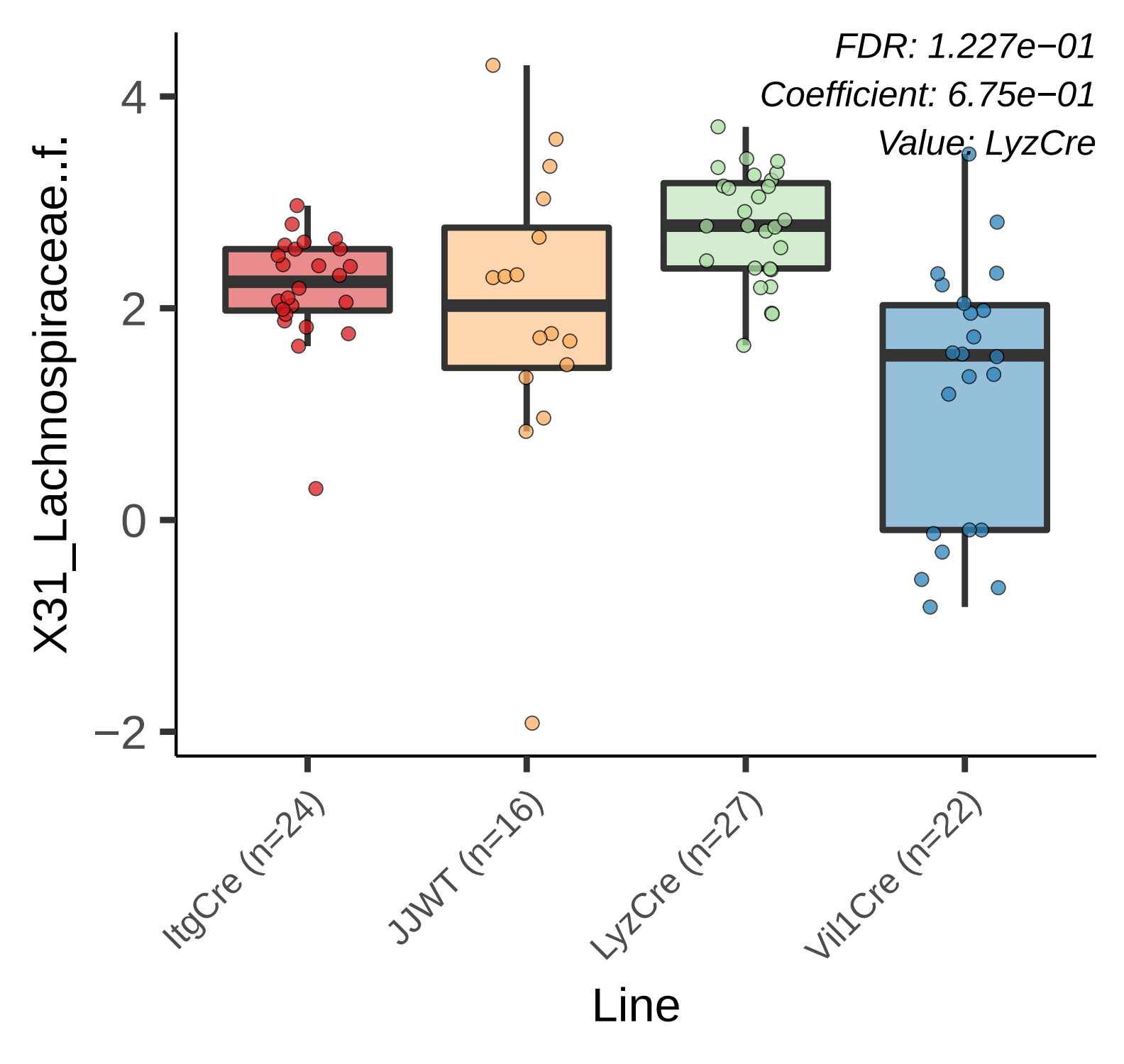

FDR: 1.222e−01
Coefficient: −7.55e−01
Value: LyzCre
X97_GCA.900066575.
2
1
0
−1
−2
−3
ItgCre (n=24)
JJWT (n=16)
LyzCre (n=27)
Vil1Cre (n=22)
Line

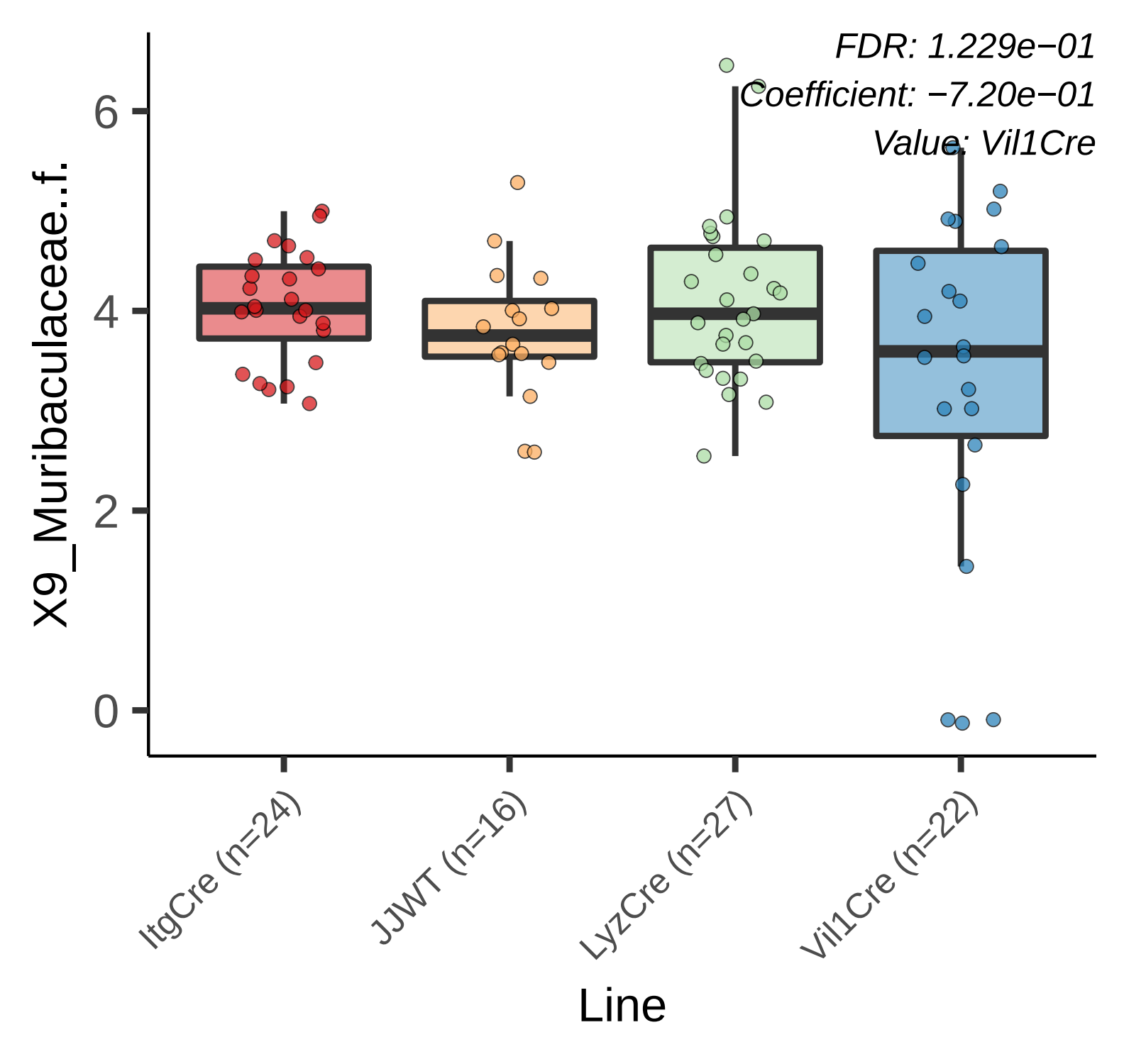

FDR: 1.222e−01
Coefficient: −7.59e−01
Value: LyzCre
X168_Eggerthellaceae..f.
2
1
0
−1
−2
−3
ItgCre (n=24)
JJWT (n=16)
LyzCre (n=27)
Vil1Cre (n=22)
Line

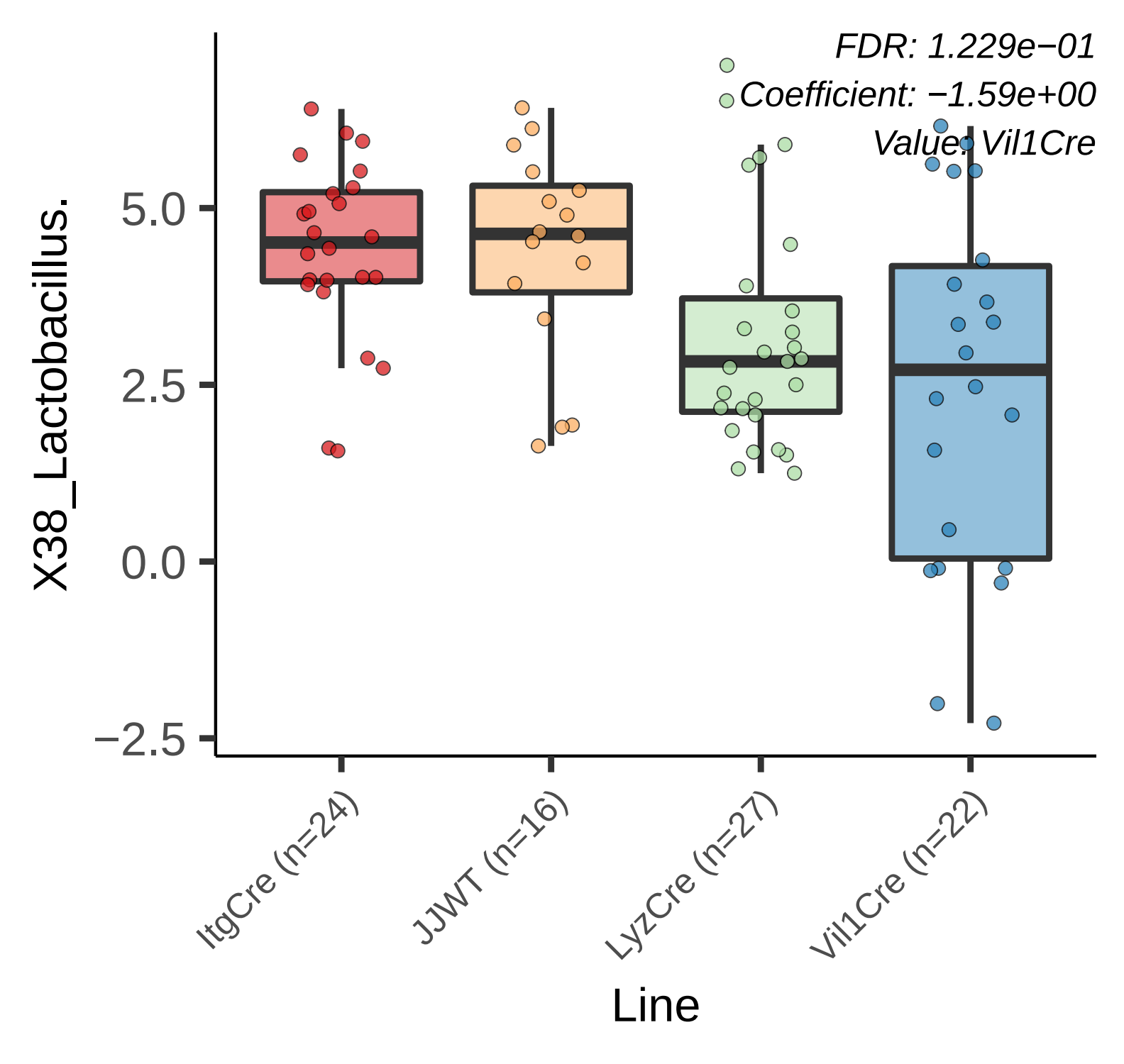

FDR: 1.222e−01
Coefficient: −7.49e−01
Value: Vil1Cre
X200_Alistipes.
3
2
1
0
−1
−2
−3
ItgCre (n=24)
JJWT (n=16)
LyzCre (n=27)
Vil1Cre (n=22)
Line

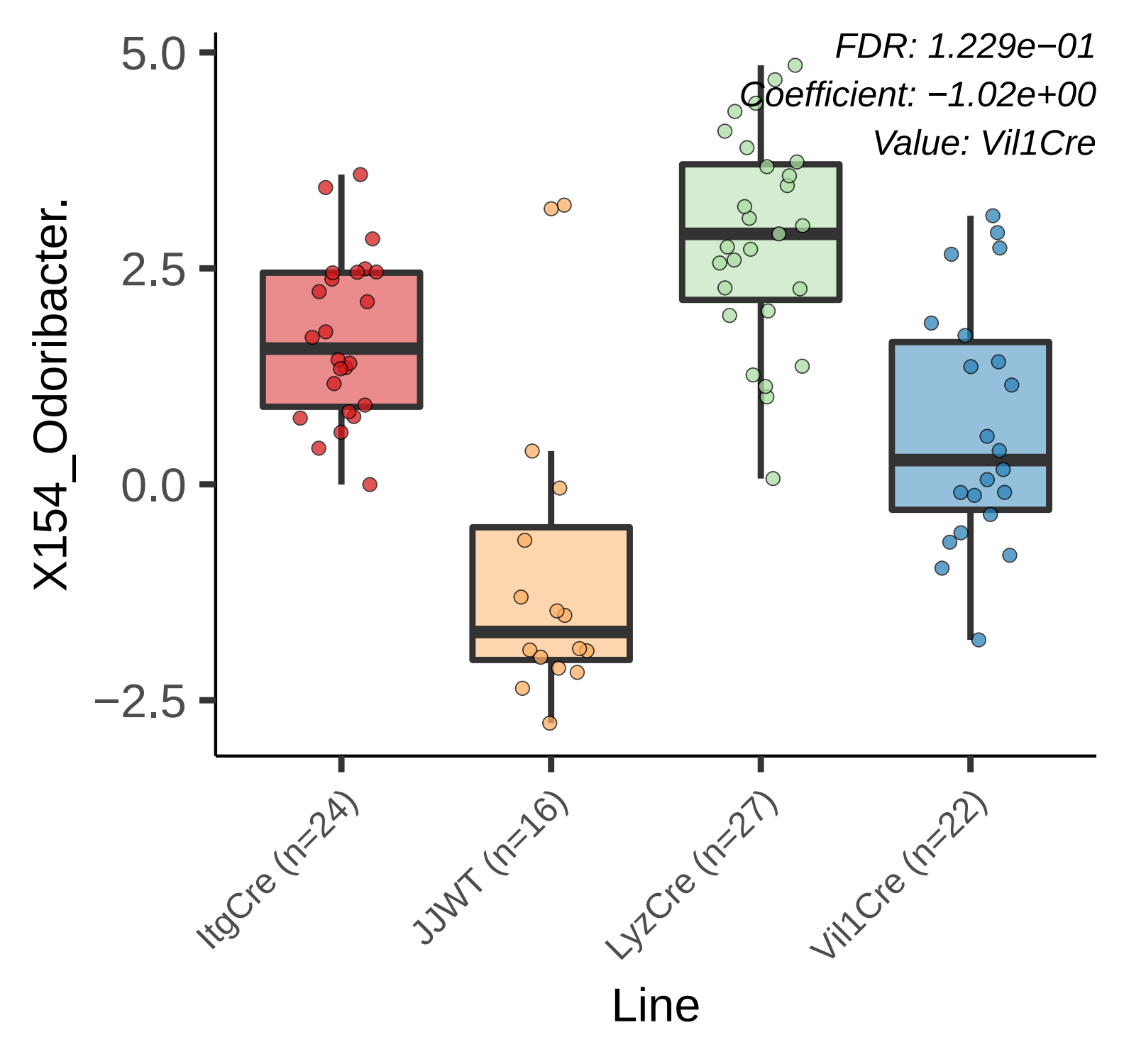

FDR: 1.226e−01
Coefficient: −1.11e+00
Value: LyzCre
X147_Lachnospiraceae_NK4A136_group.
4
2
0
−2
ItgCre (n=24)
JJWT (n=16)
LyzCre (n=27)
Vil1Cre (n=22)
Line

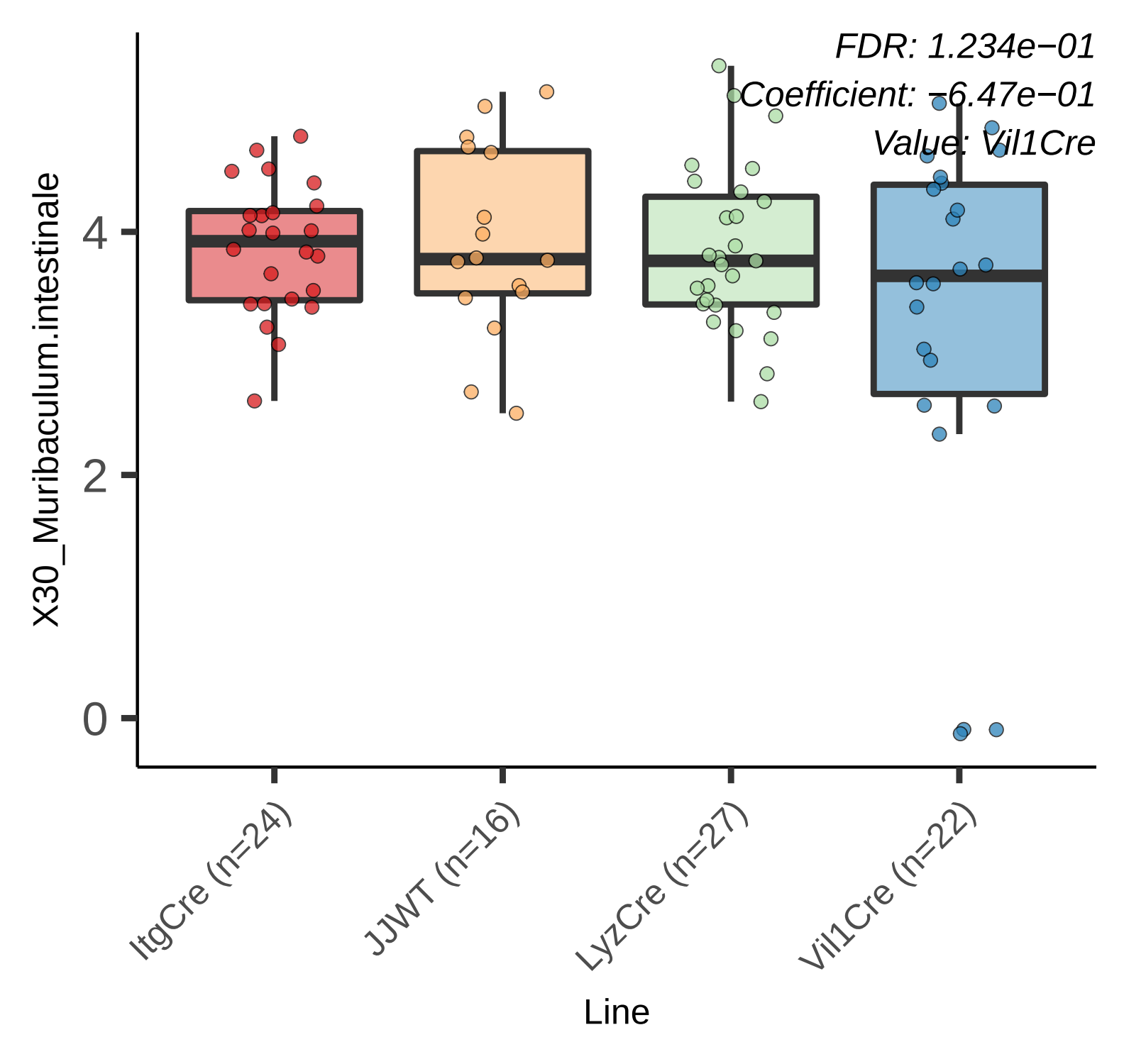

FDR: 1.226e−01
Coefficient: 2.45e+00
Value: JJWT
X229_Lachnospiraceae_NK4A136_group.
4
2
0
−2
−4
ItgCre (n=24)
JJWT (n=16)
LyzCre (n=27)
Vil1Cre (n=22)
Line

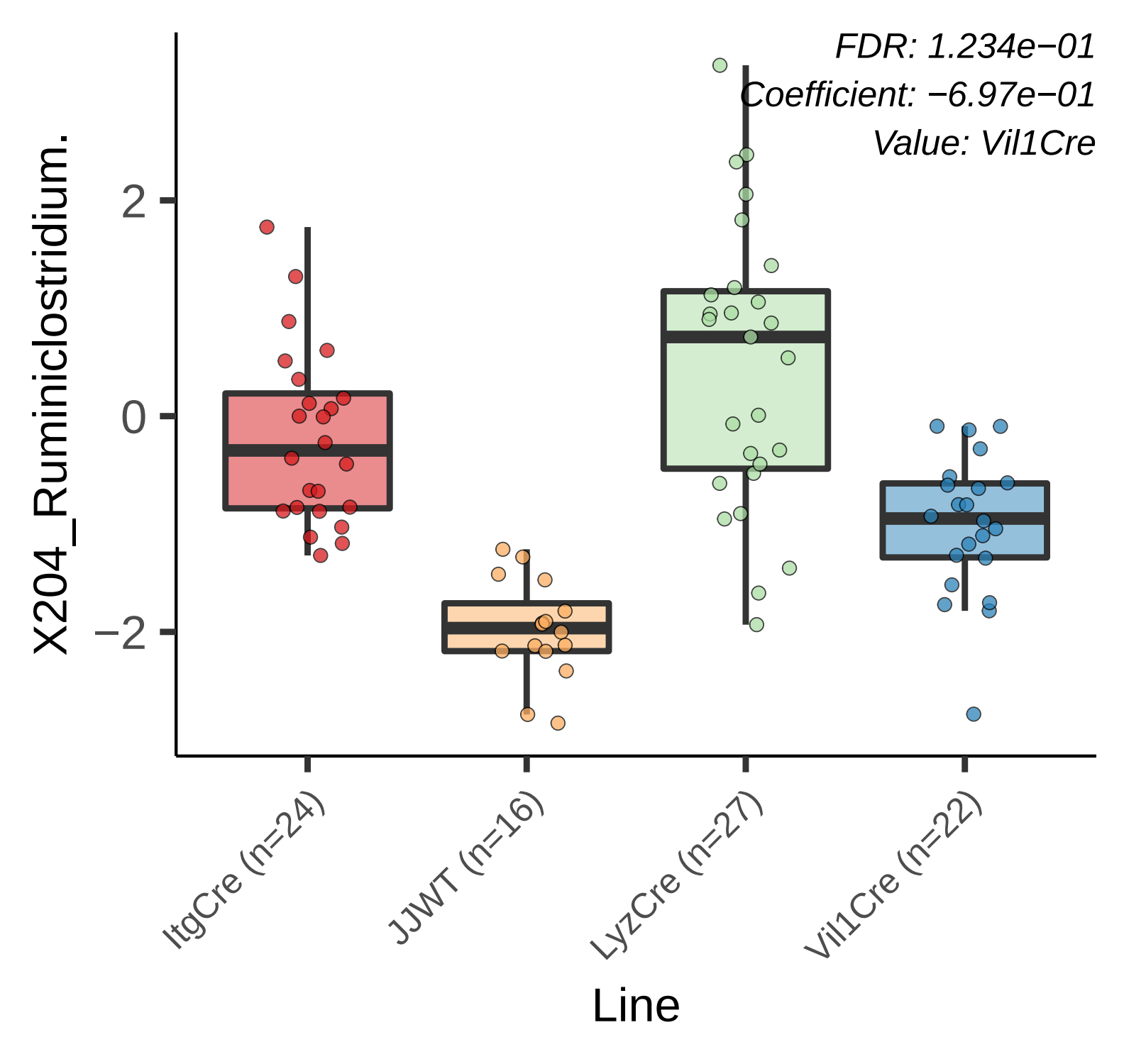

FDR: 1.227e−01
Coefficient: 6.75e−01
Value: LyzCre
X31_Lachnospiraceae..f.
4
2
0
−2
ItgCre (n=24)
JJWT (n=16)
LyzCre (n=27)
Vil1Cre (n=22)
Line

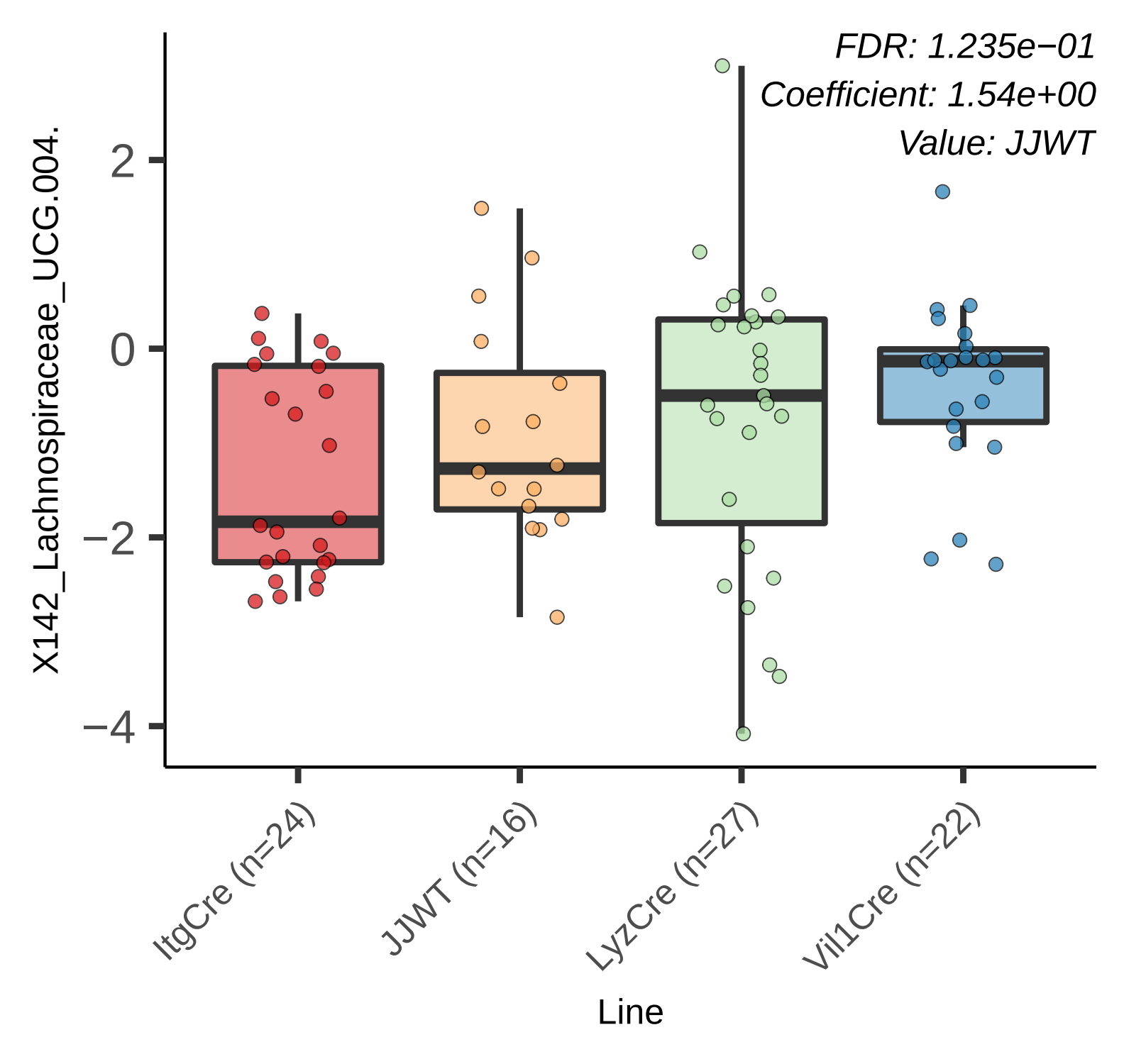

FDR: 1.229e−01
Coefficient: −7.20e−01
Value: Vil1Cre
X9_Muribaculaceae..f.
6
4
2
0
ItgCre (n=24)
JJWT (n=16)
LyzCre (n=27)
Vil1Cre (n=22)
Line

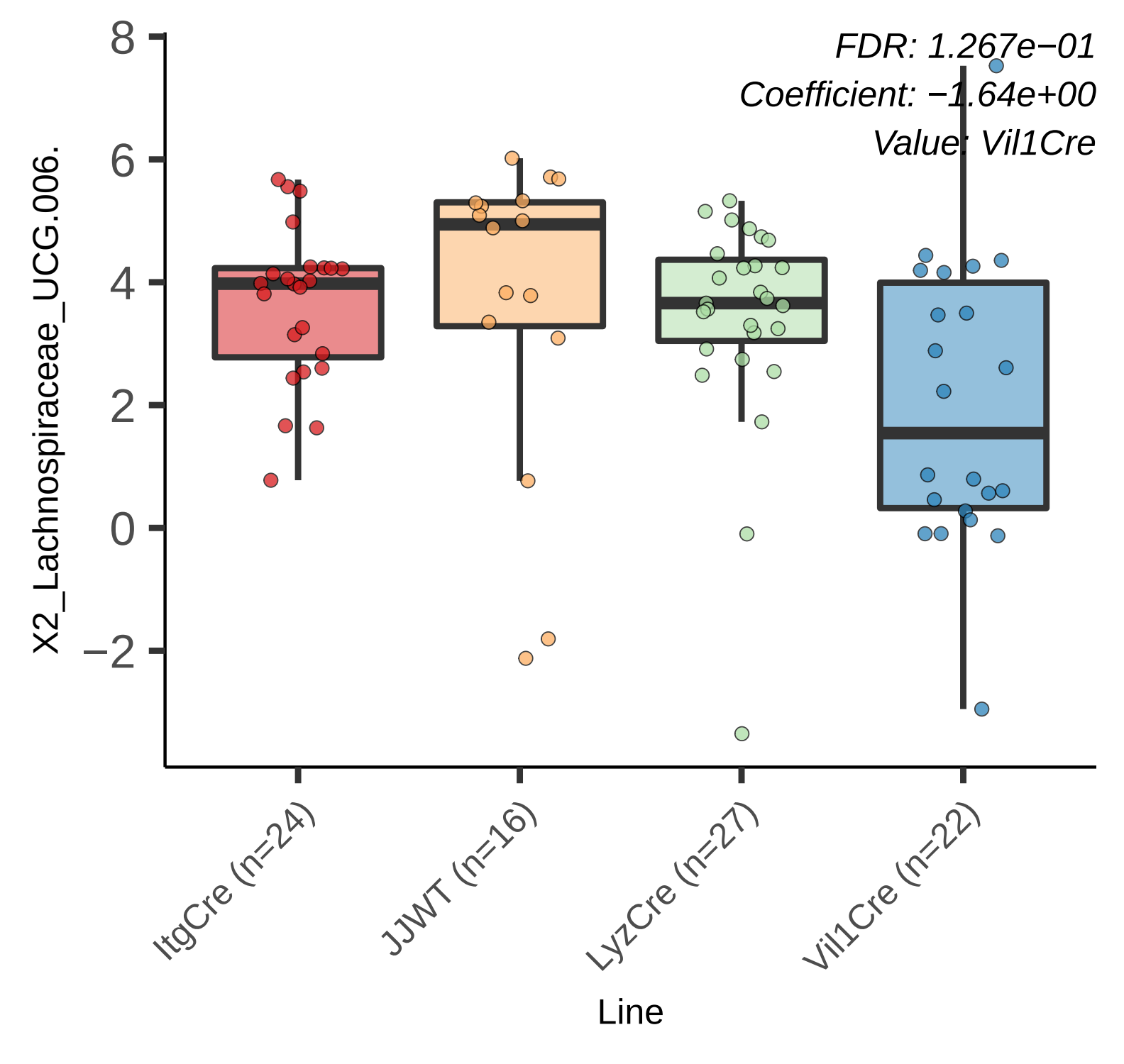

FDR: 1.229e−01
Coefficient: −1.59e+00
Value: Vil1Cre
X38_Lactobacillus.
5.0
2.5
0.0
−2.5
ItgCre (n=24)
JJWT (n=16)
LyzCre (n=27)
Vil1Cre (n=22)
Line

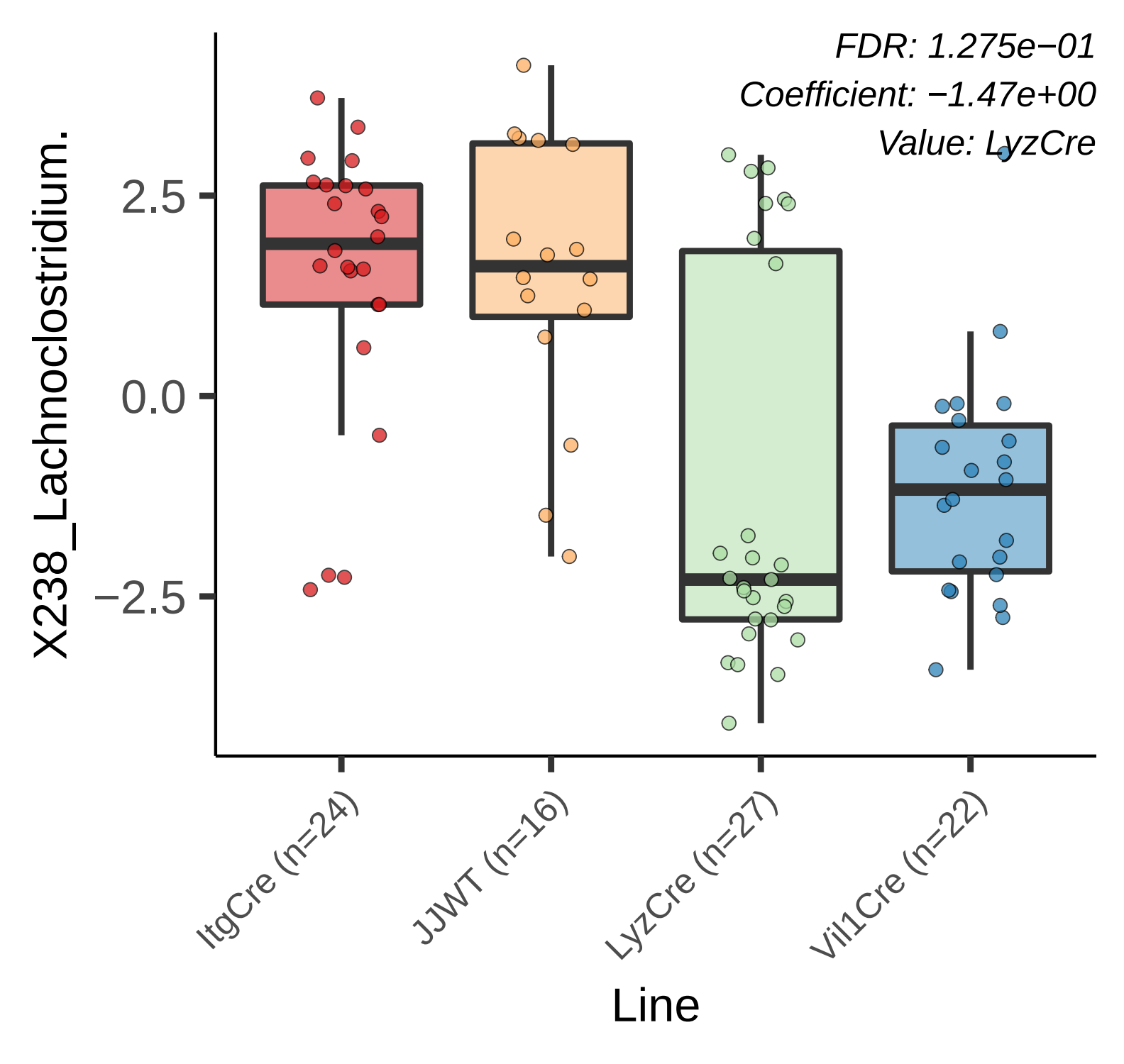

FDR: 1.229e−01
Coefficient: −1.02e+00
Value: Vil1Cre
X154_Odoribacter.
5.0
2.5
0.0
−2.5
ItgCre (n=24)
JJWT (n=16)
LyzCre (n=27)
Vil1Cre (n=22)
Line

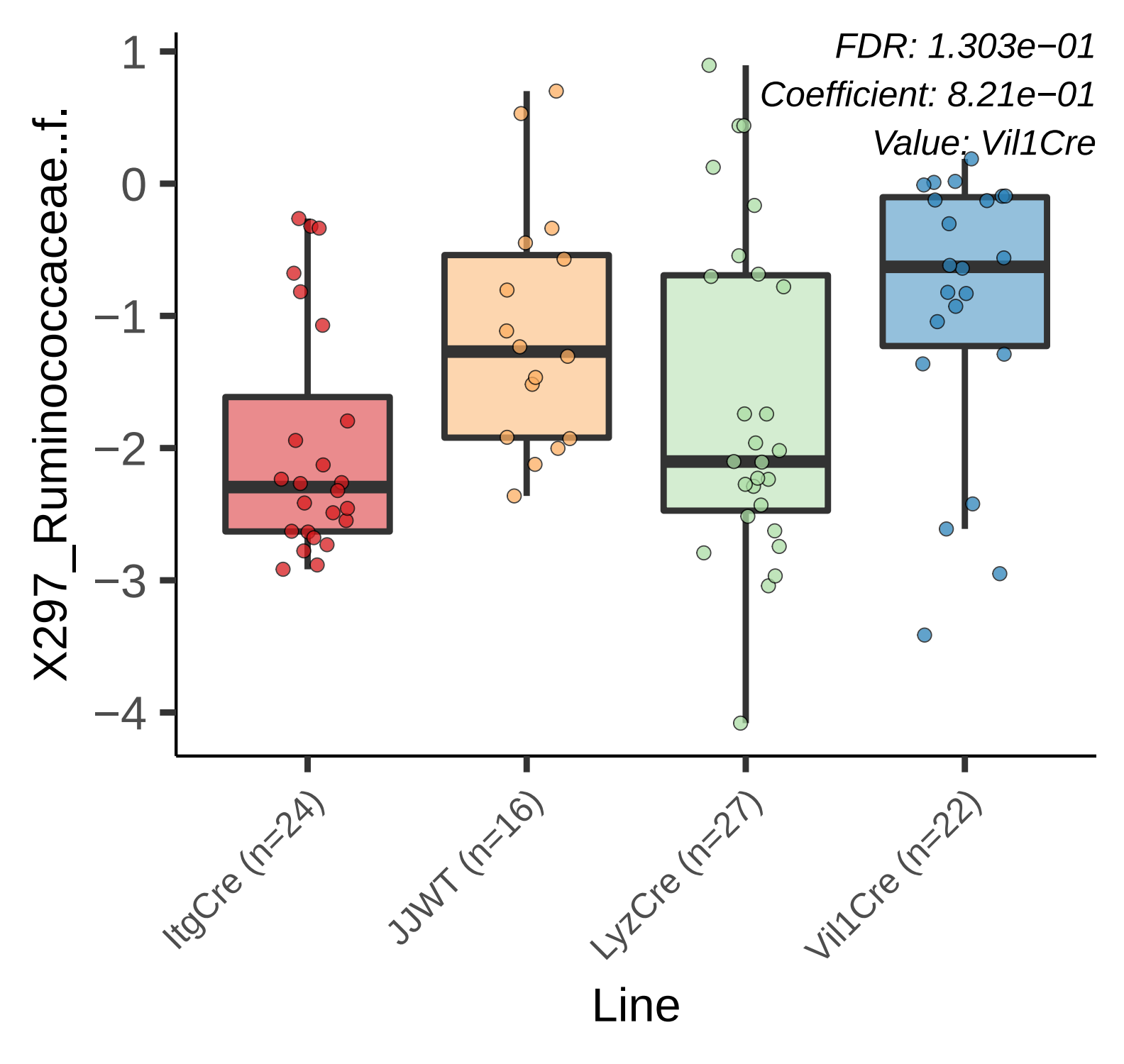

FDR: 1.234e−01
Coefficient: −6.47e−01
Value: Vil1Cre
X30_Muribaculum.intestinale
4
2
0
ItgCre (n=24)
JJWT (n=16)
LyzCre (n=27)
Vil1Cre (n=22)
Line

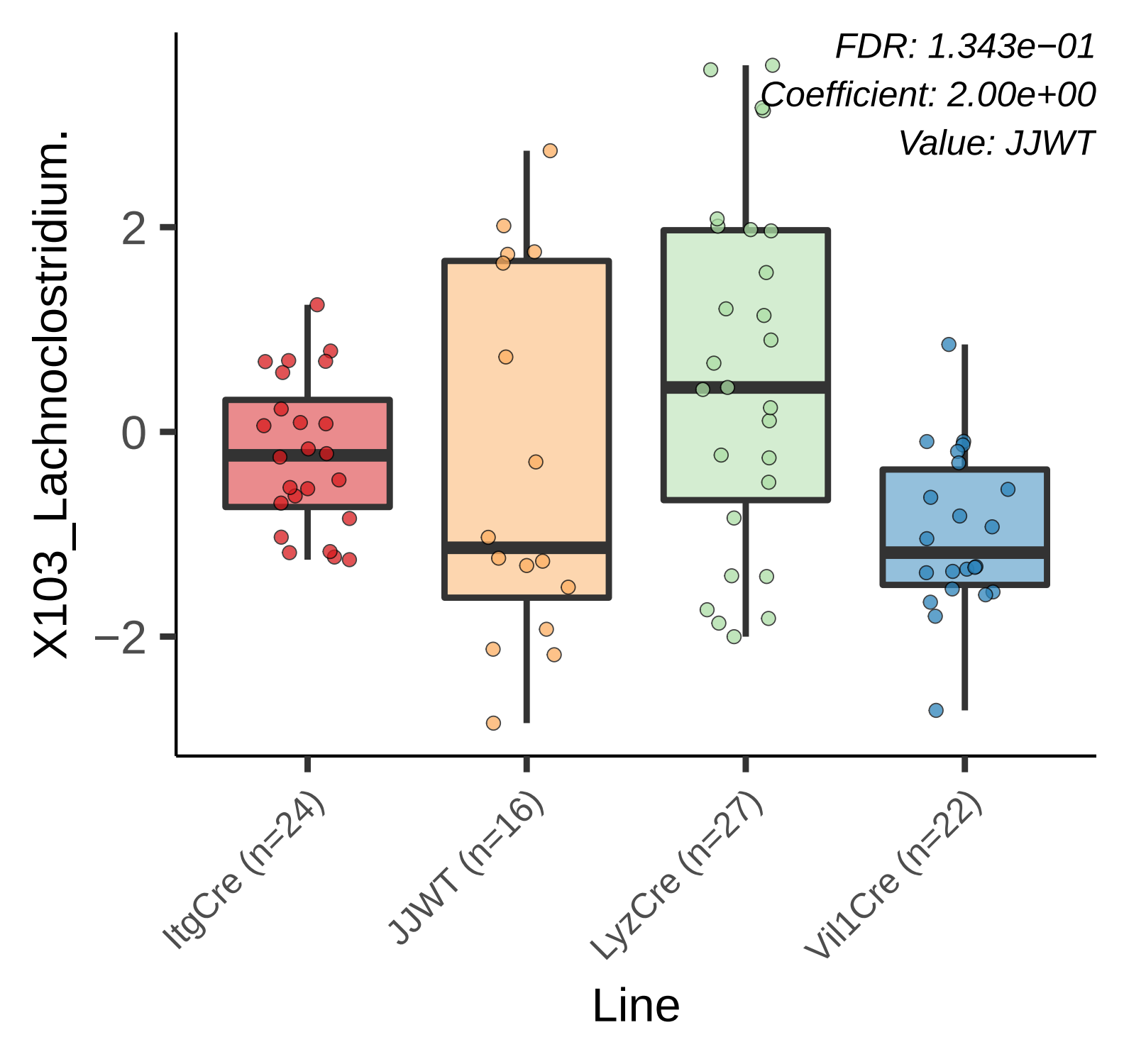

FDR: 1.234e−01
Coefficient: −6.97e−01
Value: Vil1Cre
X204_Ruminiclostridium.
2
0
−2
ItgCre (n=24)
JJWT (n=16)
LyzCre (n=27)
Vil1Cre (n=22)
Line

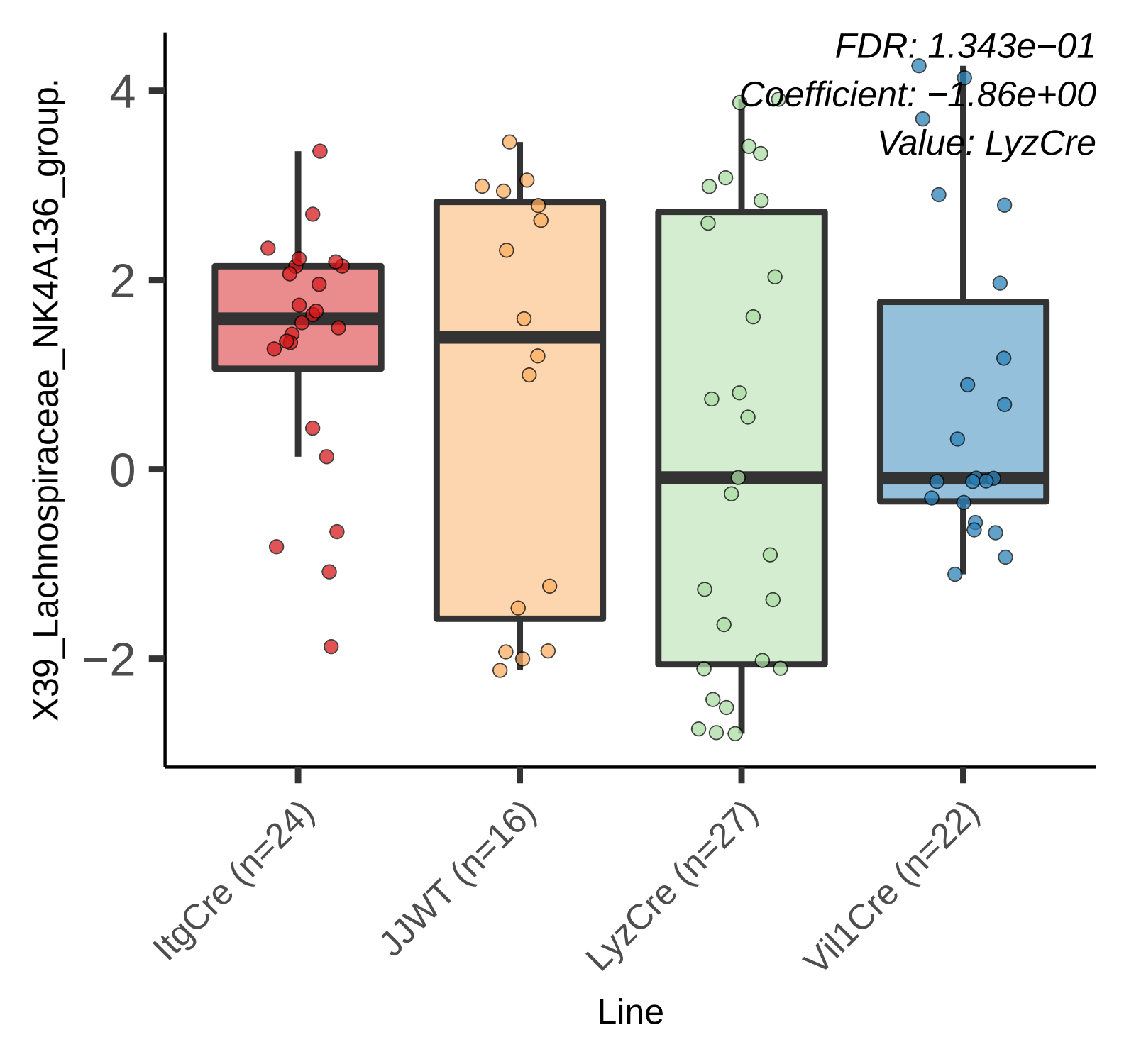

FDR: 1.235e−01
Coefficient: 1.54e+00
Value: JJWT
X142_Lachnospiraceae_UCG.004.
2
0
−2
−4
ItgCre (n=24)
JJWT (n=16)
LyzCre (n=27)
Vil1Cre (n=22)
Line

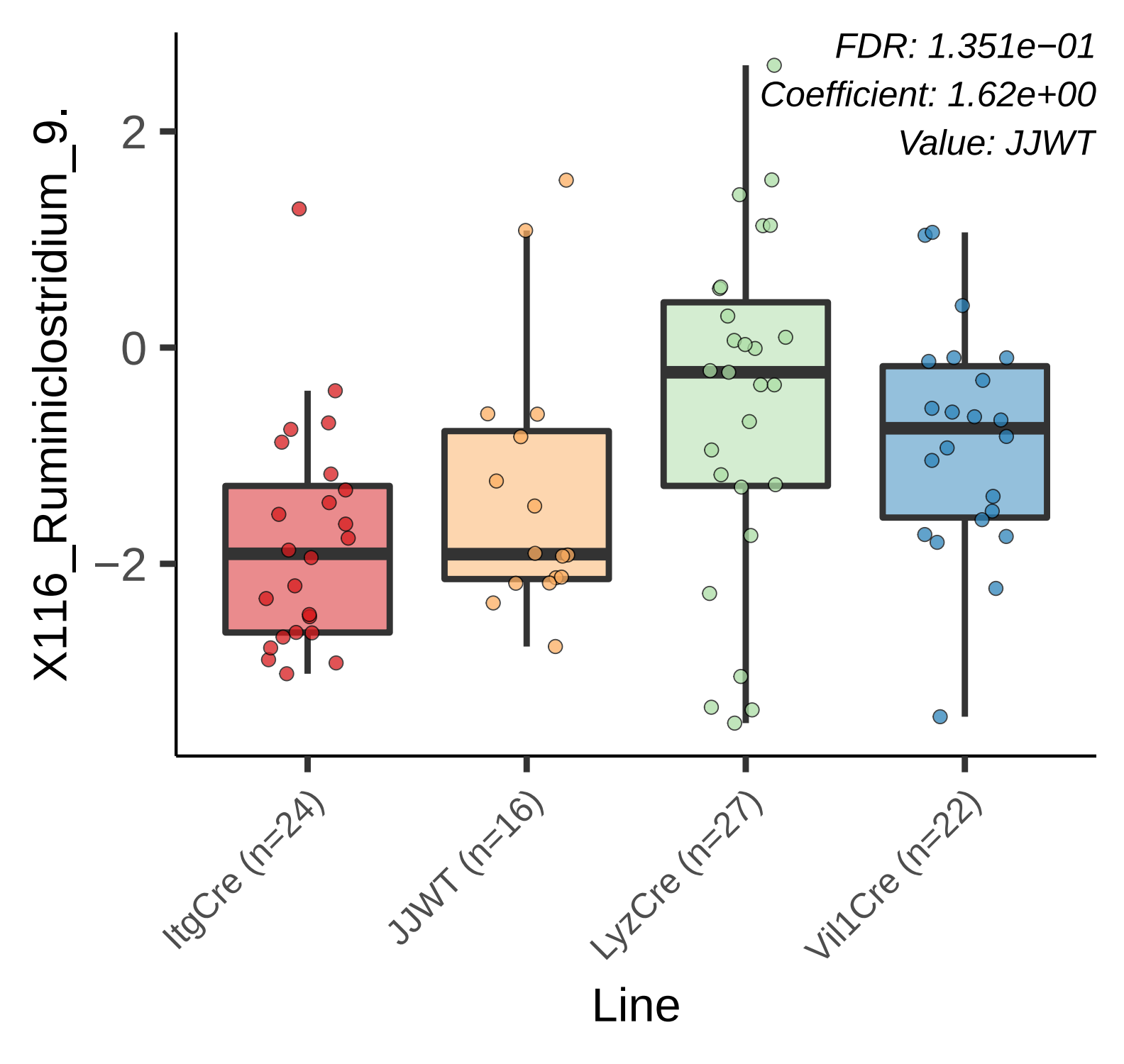

FDR: 1.267e−01
Coefficient: −1.64e+00
Value: Vil1Cre
X2_Lachnospiraceae_UCG.006.
8
6
4
2
0
−2
ItgCre (n=24)
JJWT (n=16)
LyzCre (n=27)
Vil1Cre (n=22)
Line

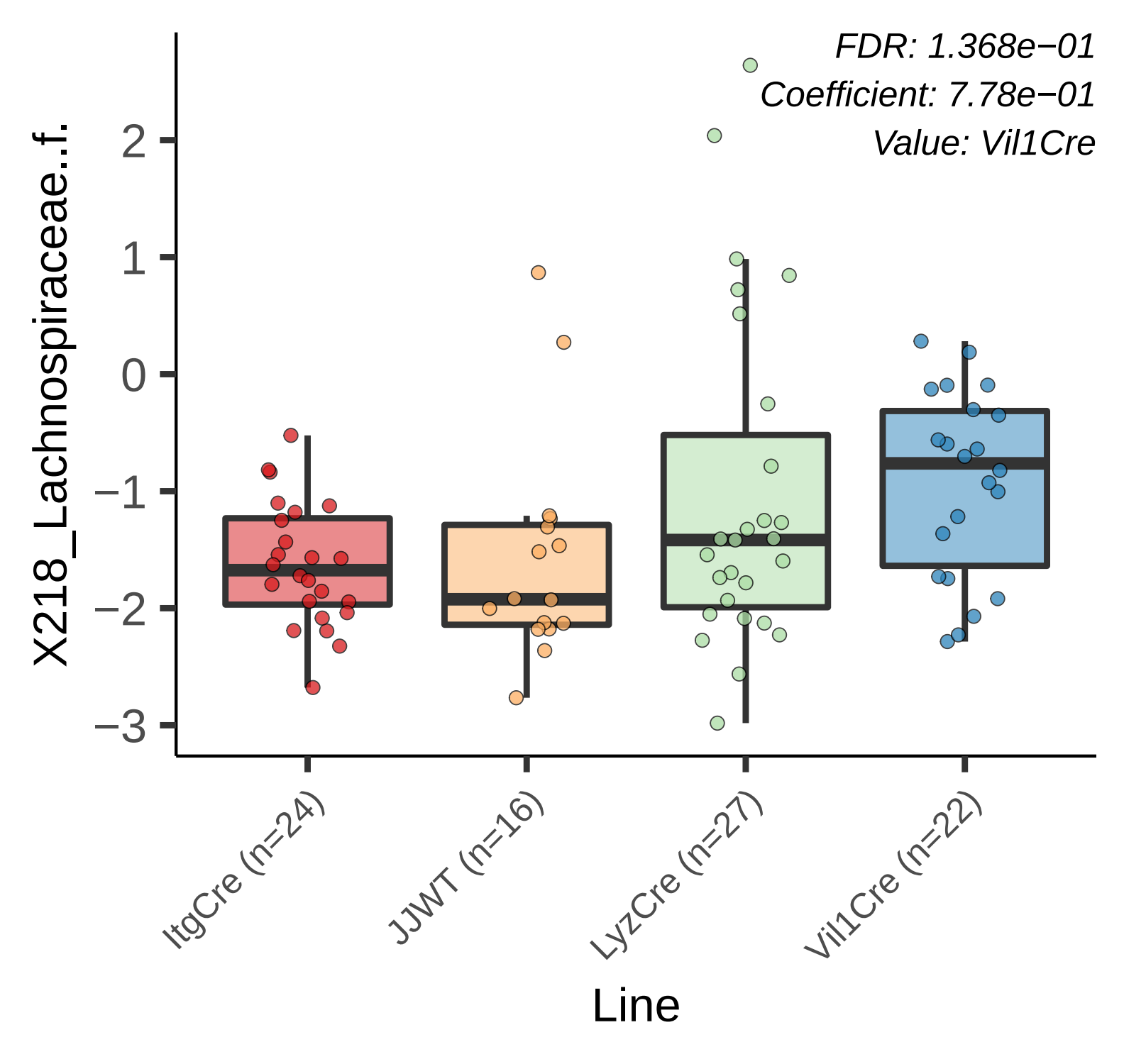

FDR: 1.275e−01
Coefficient: −1.47e+00
Value: LyzCre
X238_Lachnoclostridium.
2.5
0.0
−2.5
ItgCre (n=24)
JJWT (n=16)
LyzCre (n=27)
Vil1Cre (n=22)
Line

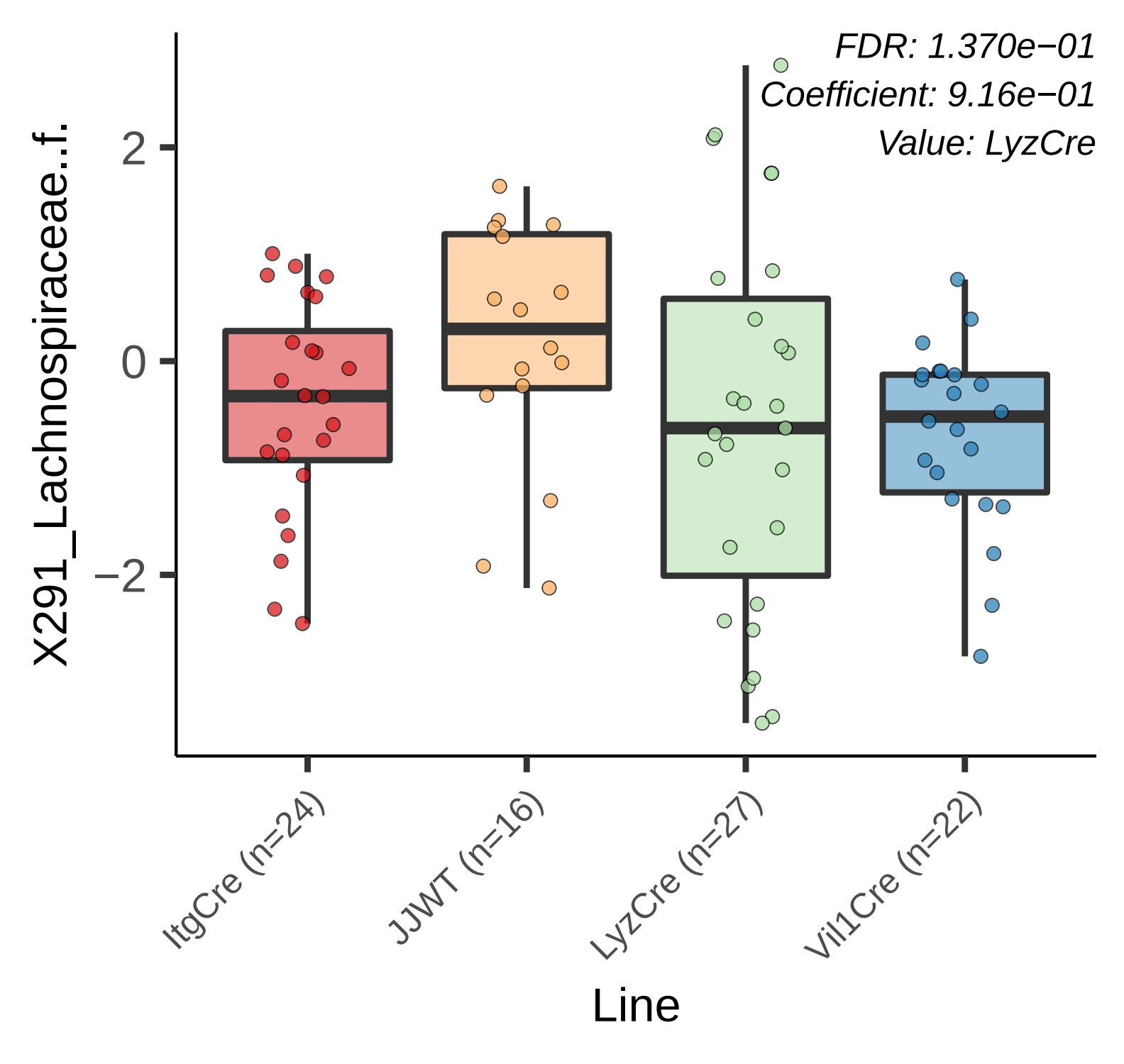

FDR: 1.303e−01
Coefficient: 8.21e−01
Value: Vil1Cre
X297_Ruminococcaceae..f.
1
0
−1
−2
−3
−4
ItgCre (n=24)
JJWT (n=16)
LyzCre (n=27)
Vil1Cre (n=22)
Line

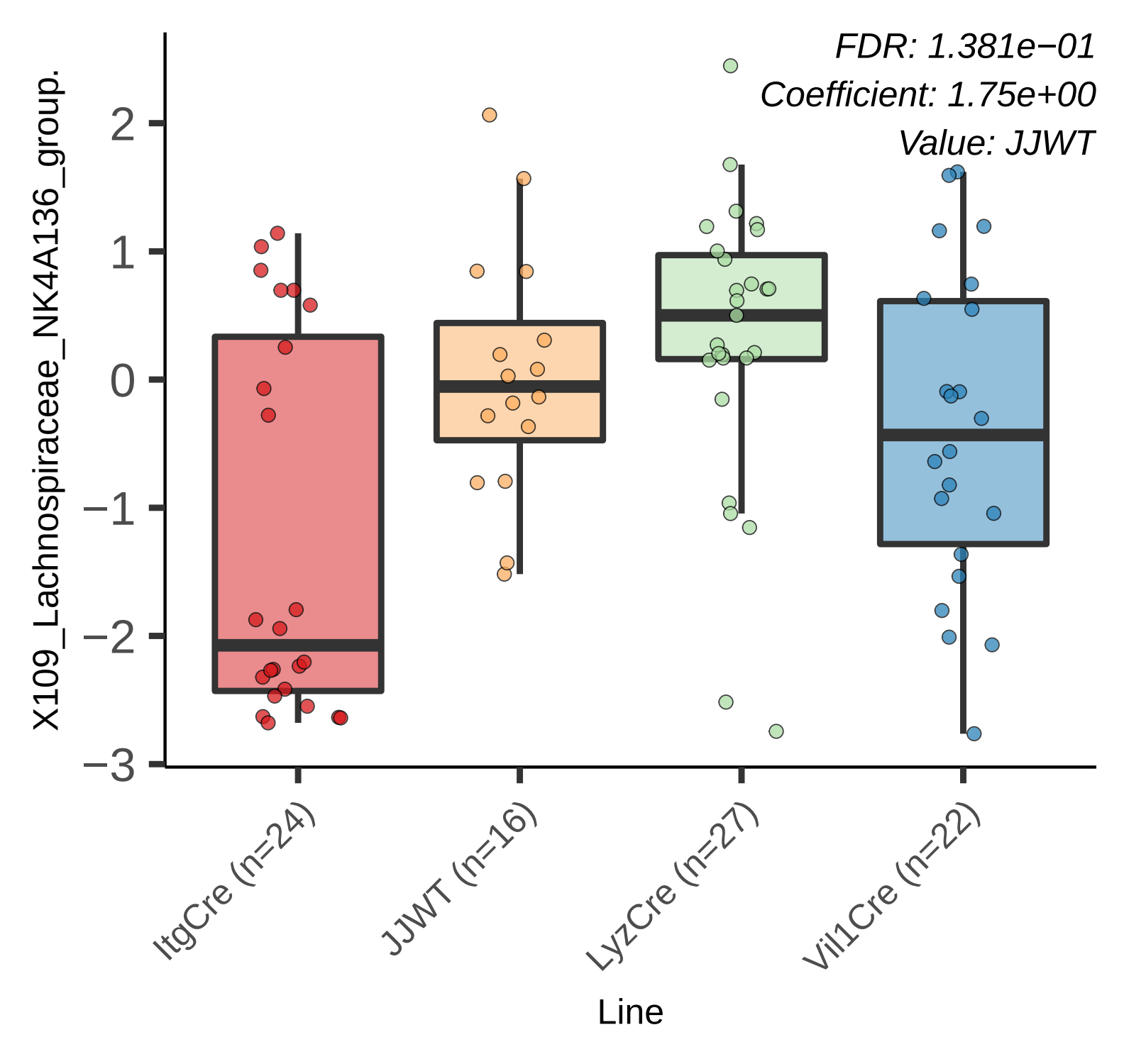

FDR: 1.343e−01
Coefficient: 2.00e+00
Value: JJWT
X103_Lachnoclostridium.
2
0
−2
ItgCre (n=24)
JJWT (n=16)
LyzCre (n=27)
Vil1Cre (n=22)
Line

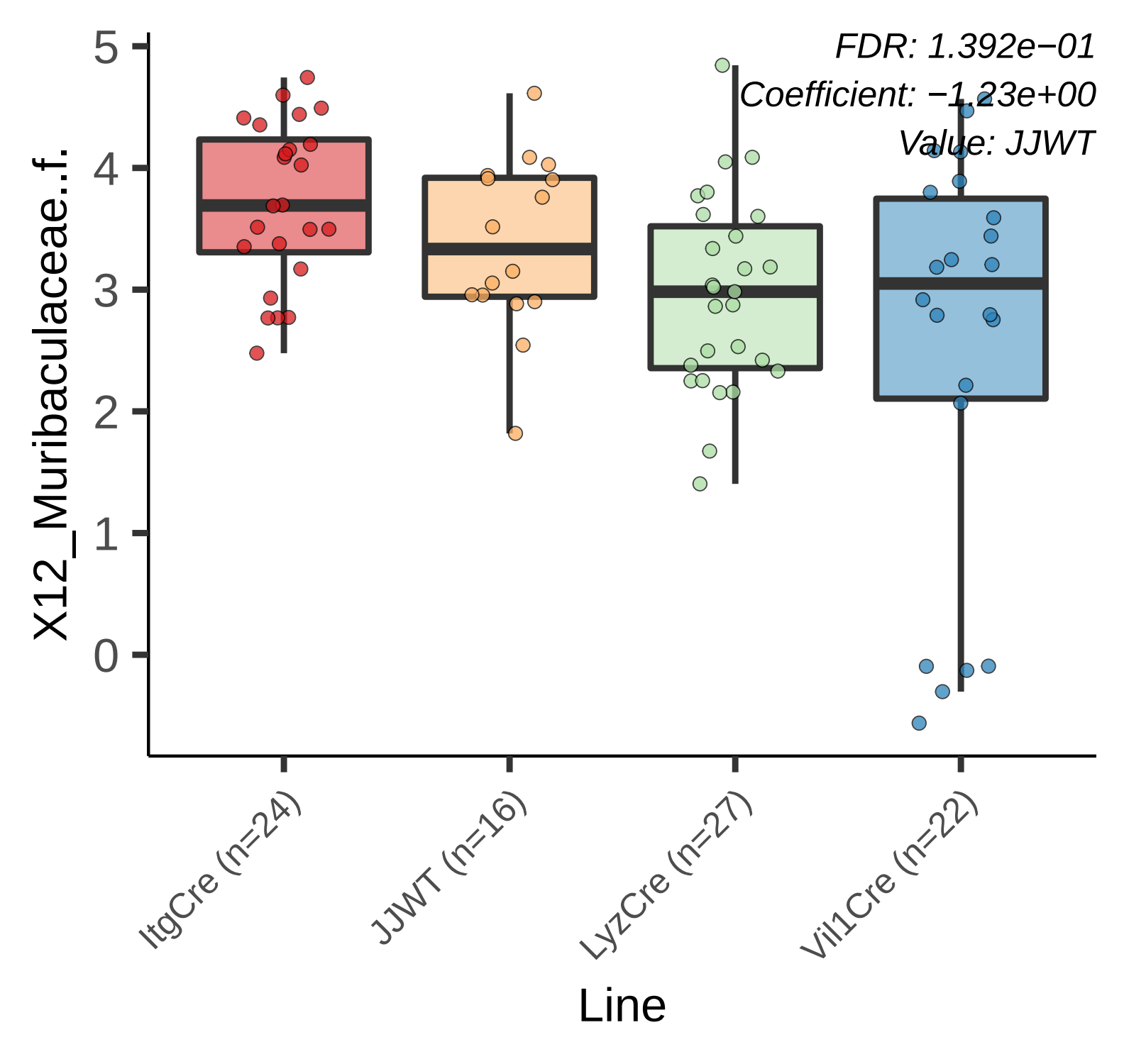

FDR: 1.343e−01
Coefficient: −1.86e+00
Value: LyzCre
X39_Lachnospiraceae_NK4A136_group.
4
2
0
−2
ItgCre (n=24)
JJWT (n=16)
LyzCre (n=27)
Vil1Cre (n=22)
Line

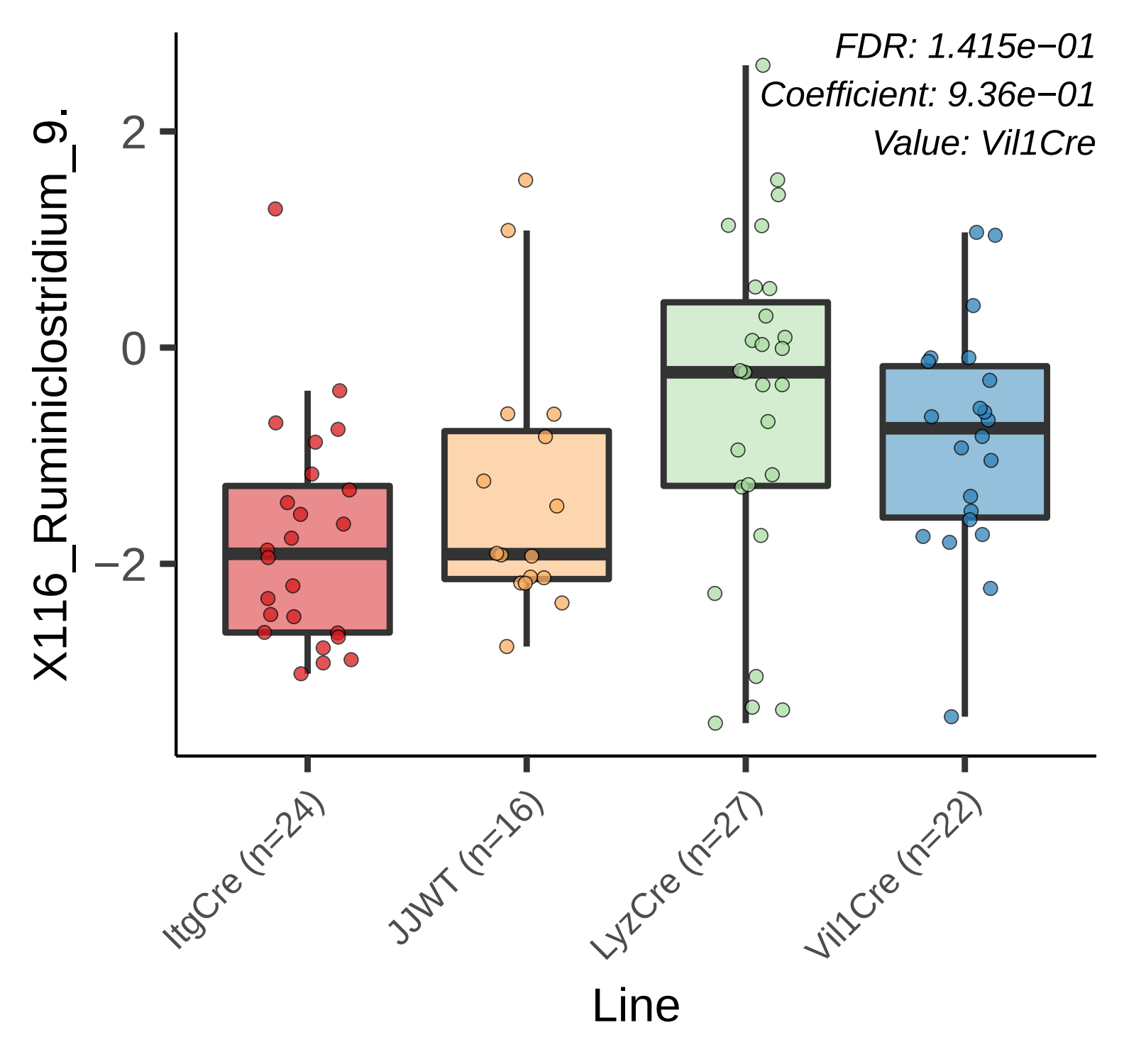

FDR: 1.351e−01
Coefficient: 1.62e+00
Value: JJWT
X116_Ruminiclostridium_9.
2
0
−2
ItgCre (n=24)
JJWT (n=16)
LyzCre (n=27)
Vil1Cre (n=22)
Line

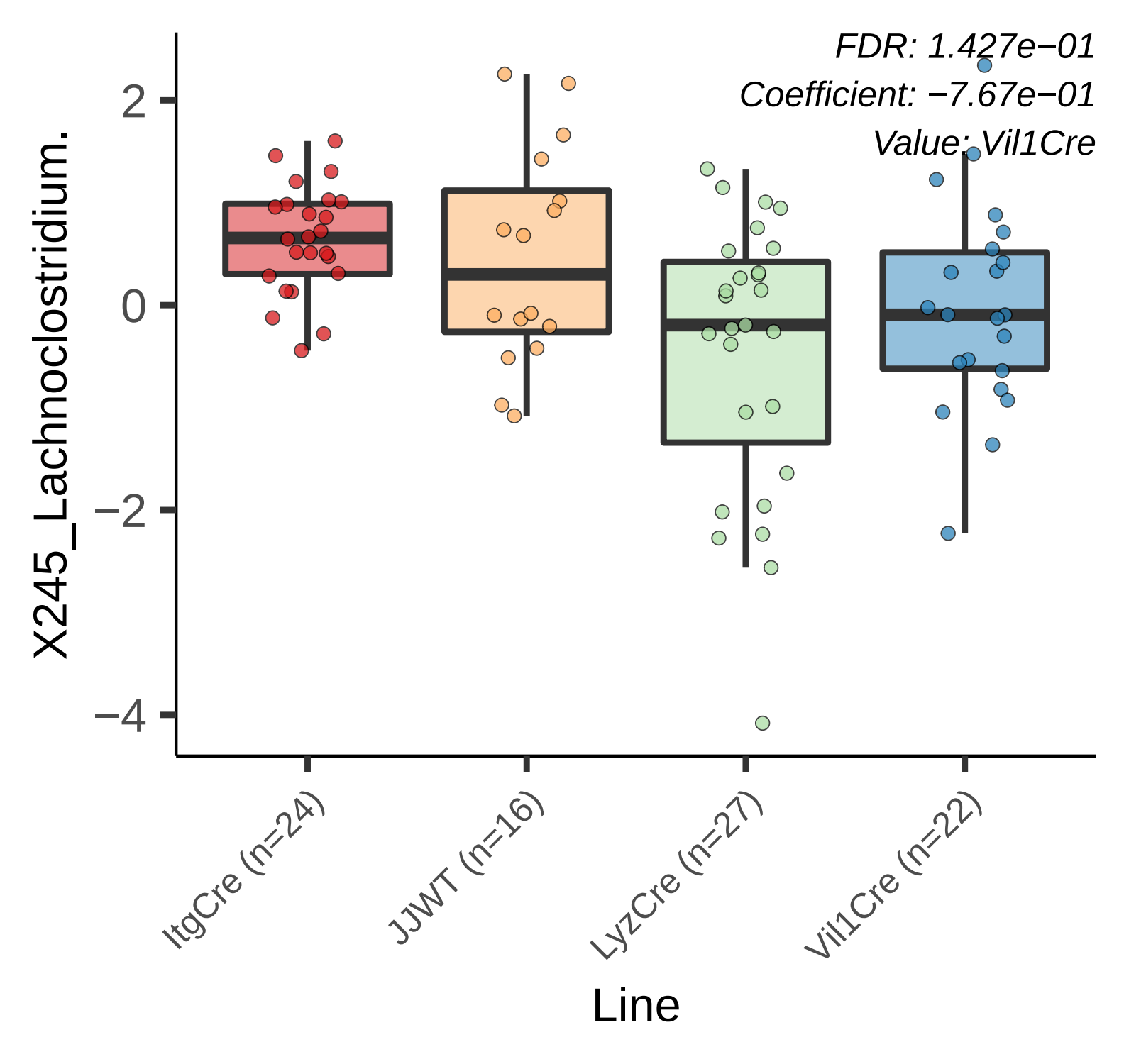

FDR: 1.368e−01
Coefficient: 7.78e−01
Value: Vil1Cre
X218_Lachnospiraceae.f.
2
1
0
−1
−2
−3
ItgCre (n=24)
JJWT (n=16)
LyzCre (n=27)
Vil1Cre (n=22)
Line

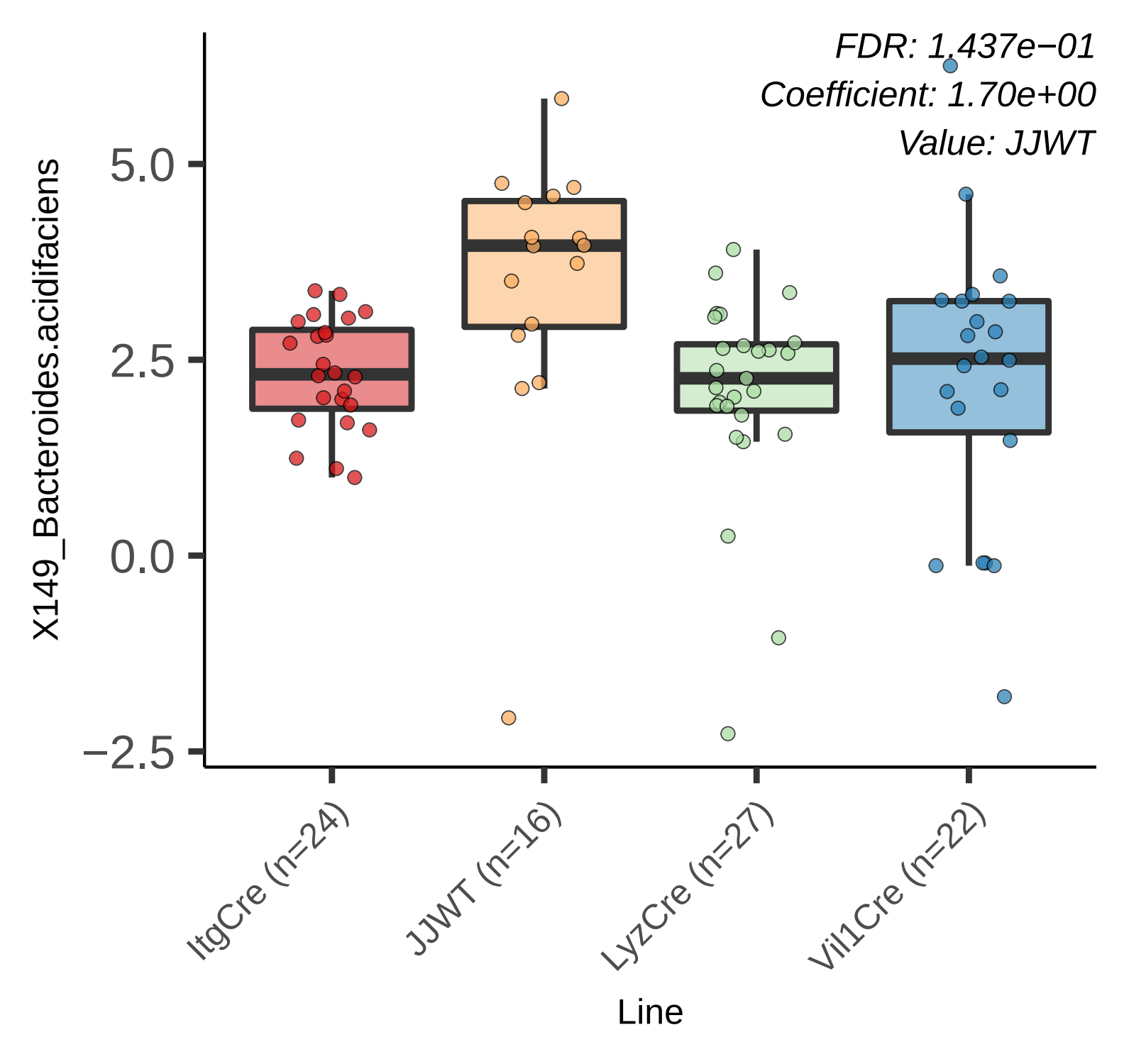

FDR: 1.370e−01
Coefficient: 9.16e−01
Value: LyzCre
X291_Lachnospiraceae.f.
2
0
−2
ItgCre (n=24)
JJWT (n=16)
LyzCre (n=27)
Vil1Cre (n=22)
Line

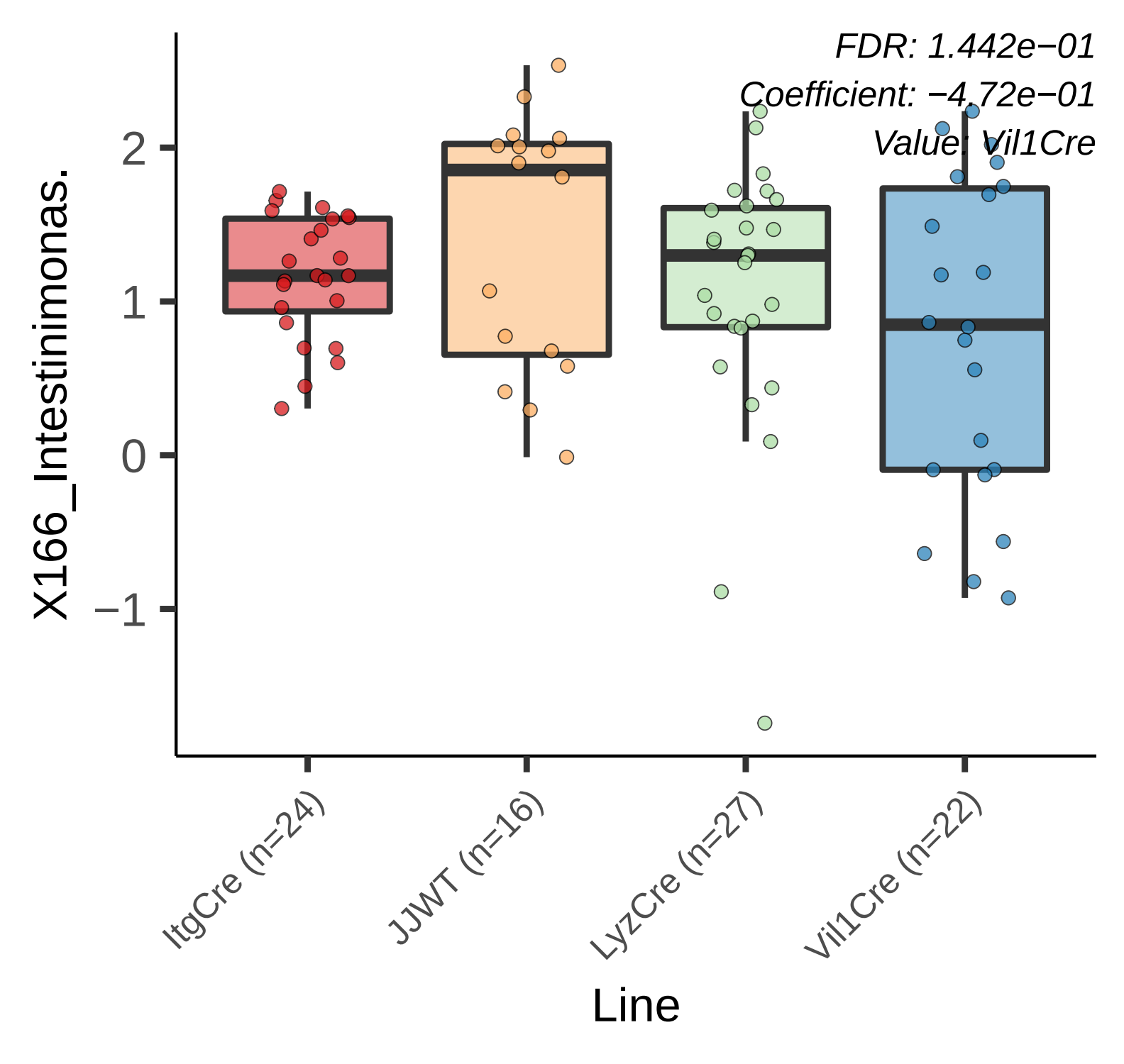

FDR: 1.381e−01
Coefficient: 1.75e+00
Value: JJWT
X109_Lachnospiraceae_NK4A136_group.
2
1
0
−1
−2
−3
ItgCre (n=24)
JJWT (n=16)
LyzCre (n=27)
Vil1Cre (n=22)
Line

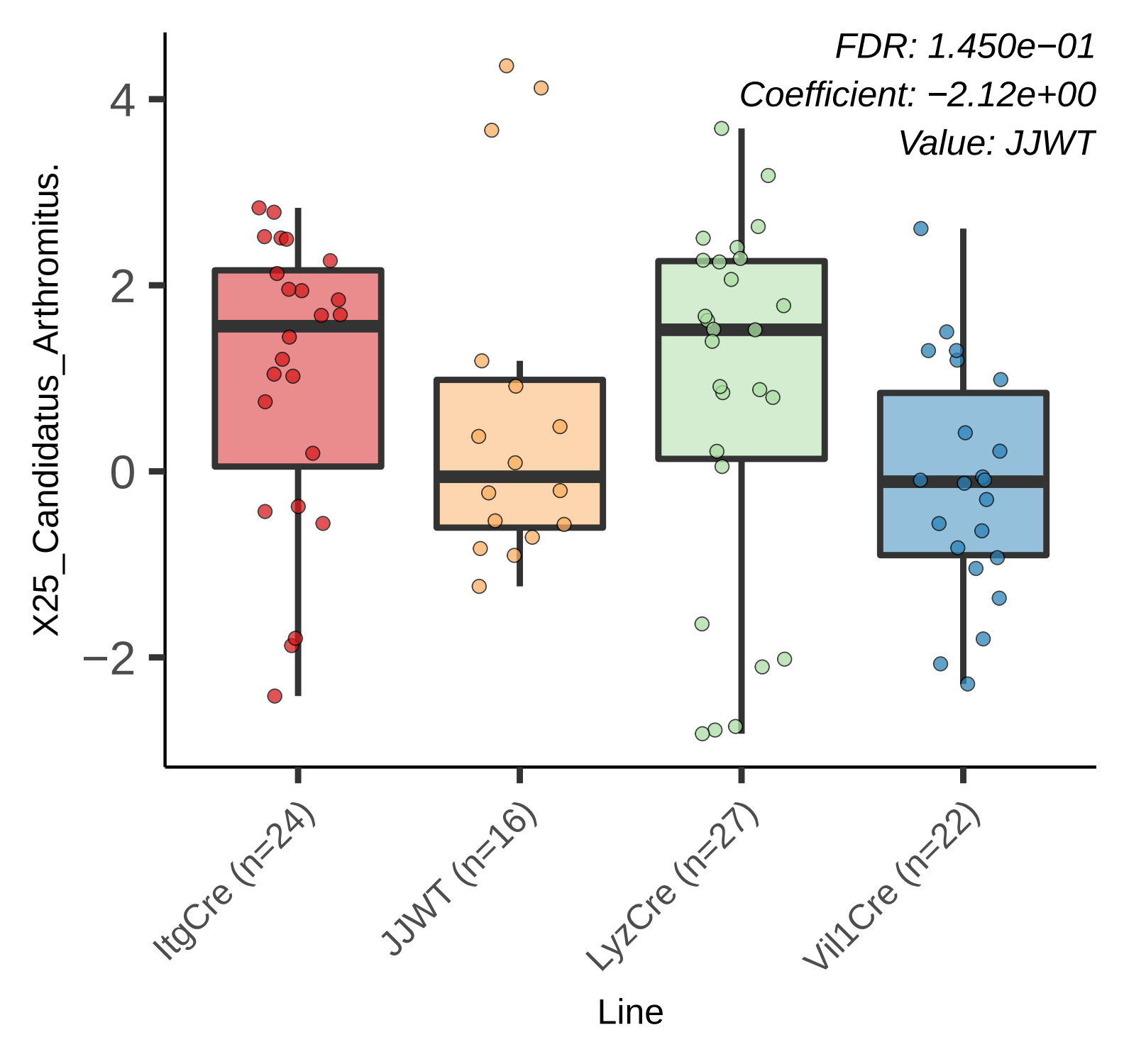

FDR: 1.392e−01
Coefficient: −1.23e+00
Value: JJWT
X12_Muribaculaceae..f.
5
4
3
2
1
0
ItgCre (n=24)
JJWT (n=16)
LyzCre (n=27)
Vil1Cre (n=22)
Line

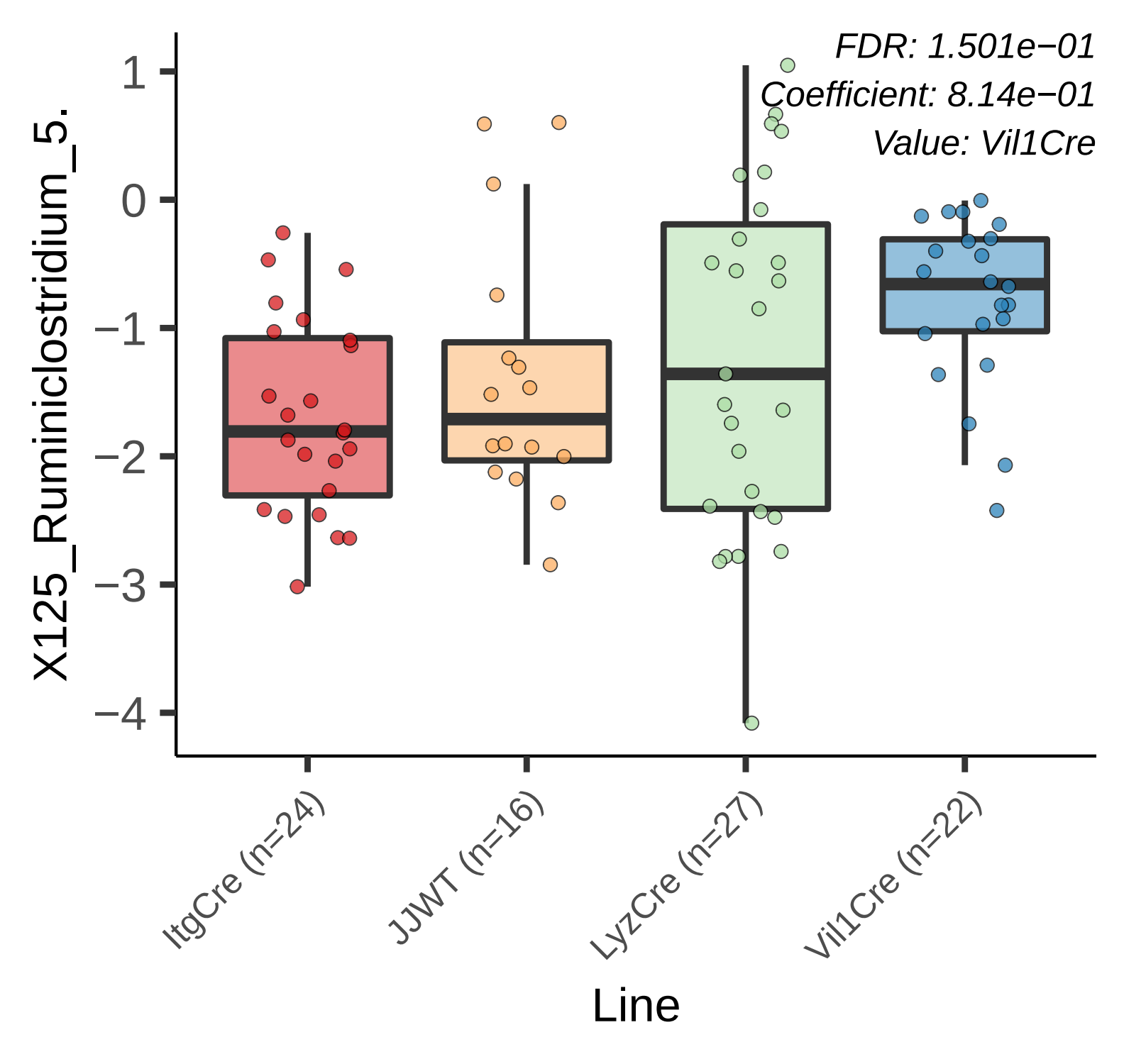

FDR: 1.415e−01
Coefficient: 9.36e−01
Value: Vil1Cre
X116_Ruminiclostridium_9.
2
0
−2
ItgCre (n=24)
JJWT (n=16)
LyzCre (n=27)
Vil1Cre (n=22)
Line

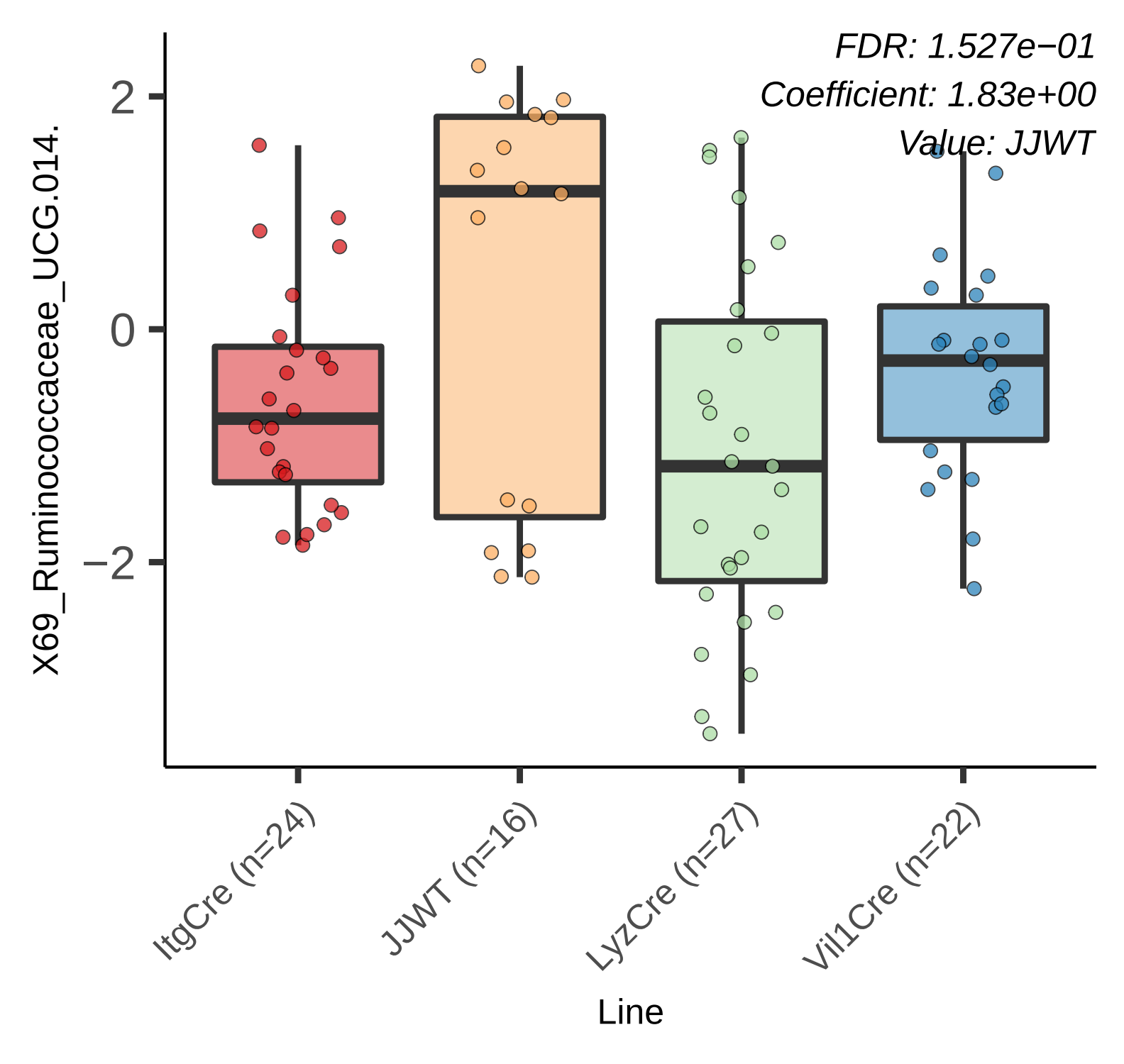

FDR: 1.427e−01
Coefficient: −7.67e−01
Value: Vil1Cre
X245_Lachnoclostridium.
2
0
−2
−4
ItgCre (n=24)
JJWT (n=16)
LyzCre (n=27)
Vil1Cre (n=22)
Line

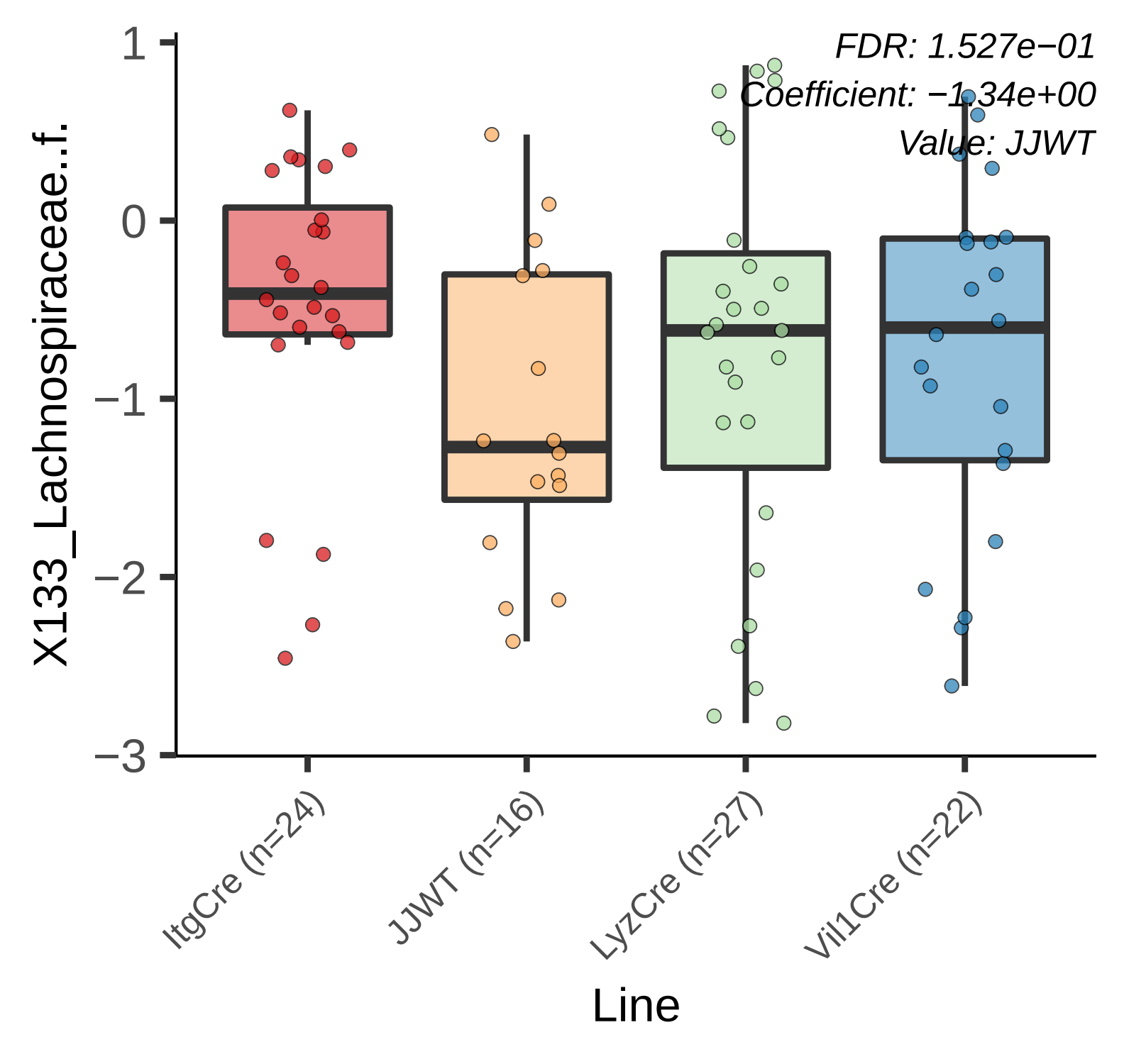

FDR: 1.437e−01
Coefficient: 1.70e+00
Value: JJWT
X149_Bacteroides.acidifaciens
5.0
2.5
0.0
−2.5
ItgCre (n=24)
JJWT (n=16)
LyzCre (n=27)
Vil1Cre (n=22)
Line

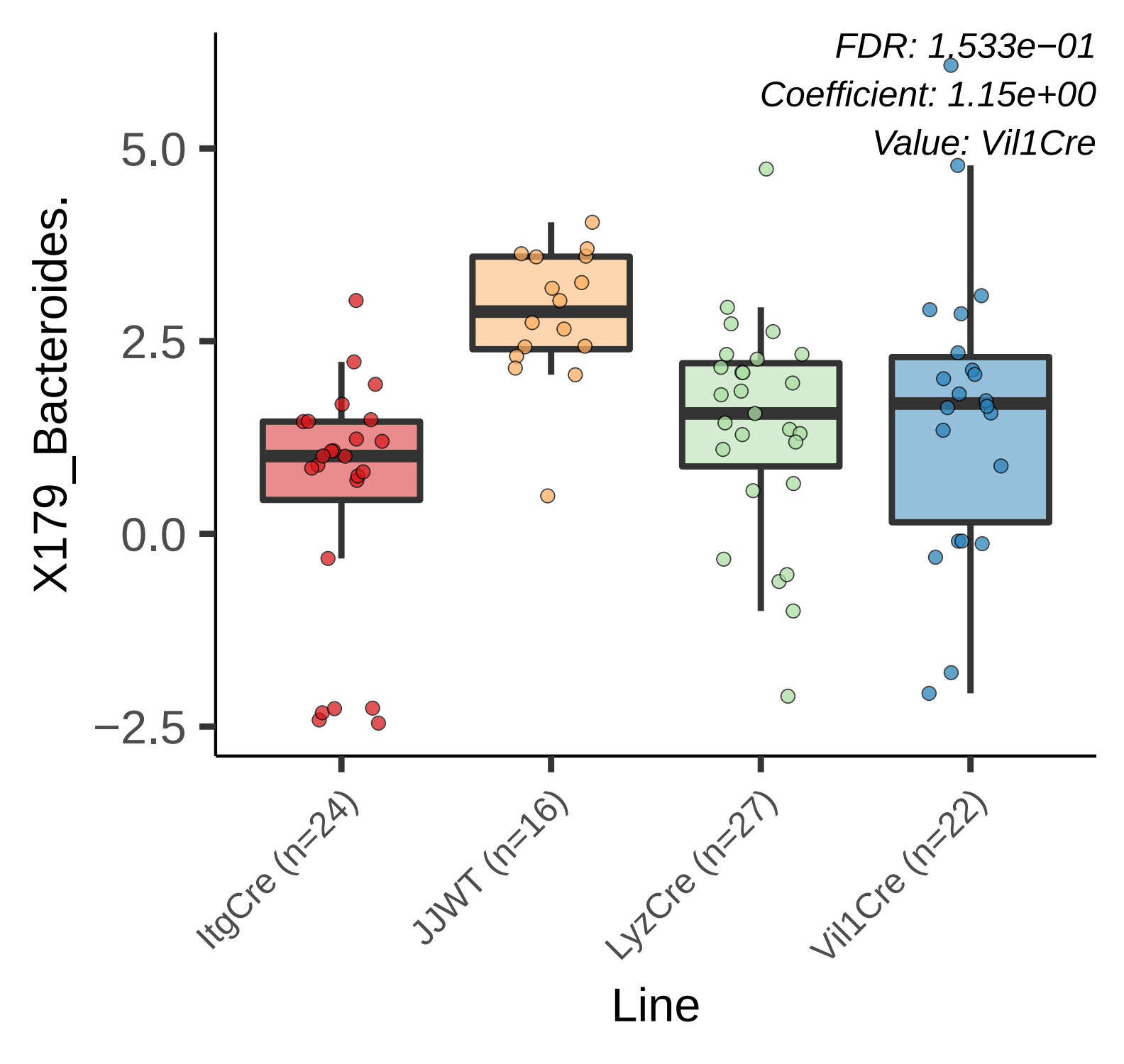

FDR: 1.442e−01
Coefficient: −4.72e−01
Value: Vil1Cre
X166_Intestinimonas.
2
1
0
−1
ItgCre (n=24)
JJWT (n=16)
LyzCre (n=27)
Vil1Cre (n=22)
Line

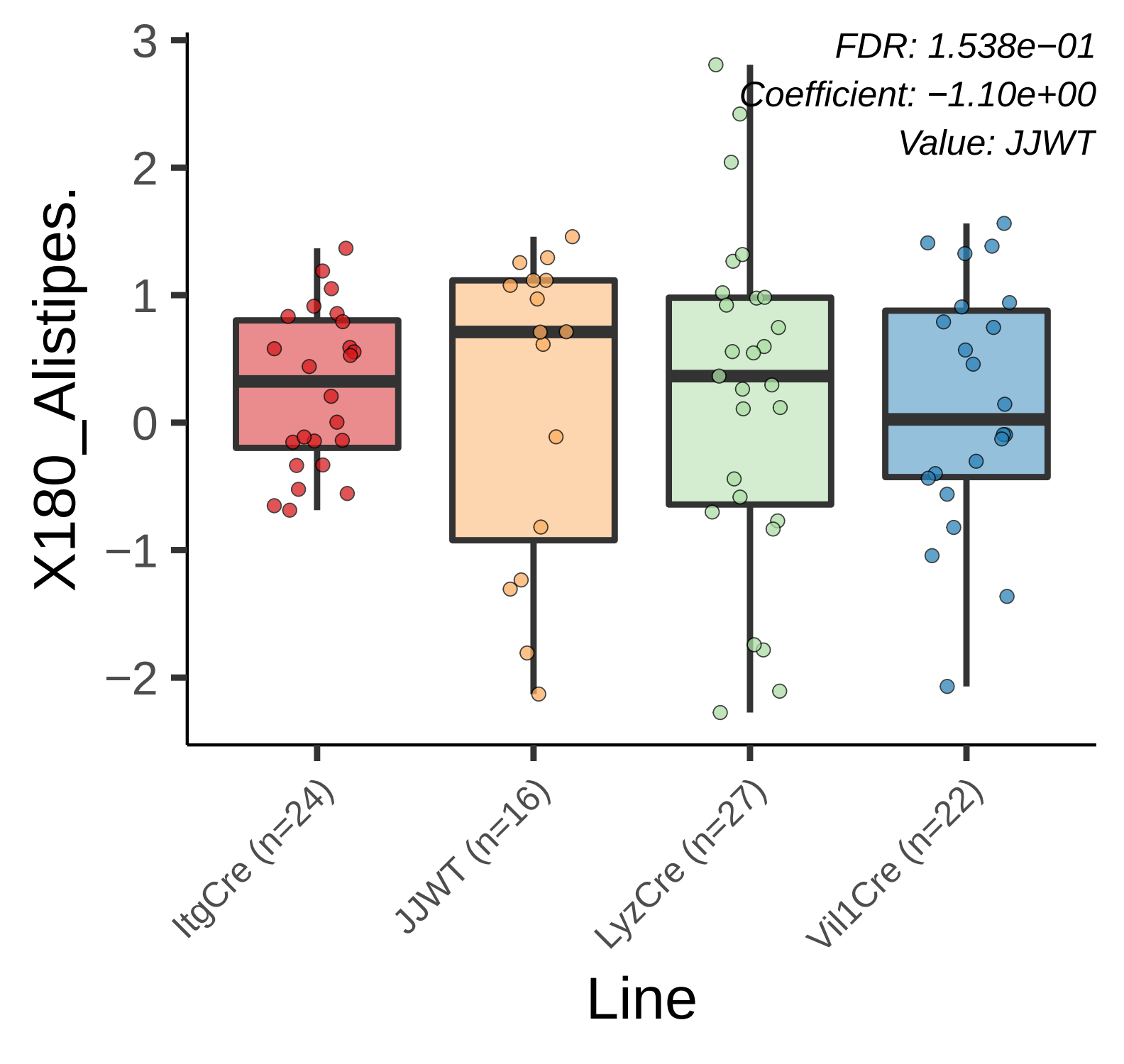

FDR: 1.450e−01
Coefficient: −2.12e+00
Value: JJWT
X25_Candidatus_Arthromitus.
4
2
0
−2
ItgCre (n=24)
JJWT (n=16)
LyzCre (n=27)
Vil1Cre (n=22)
Line

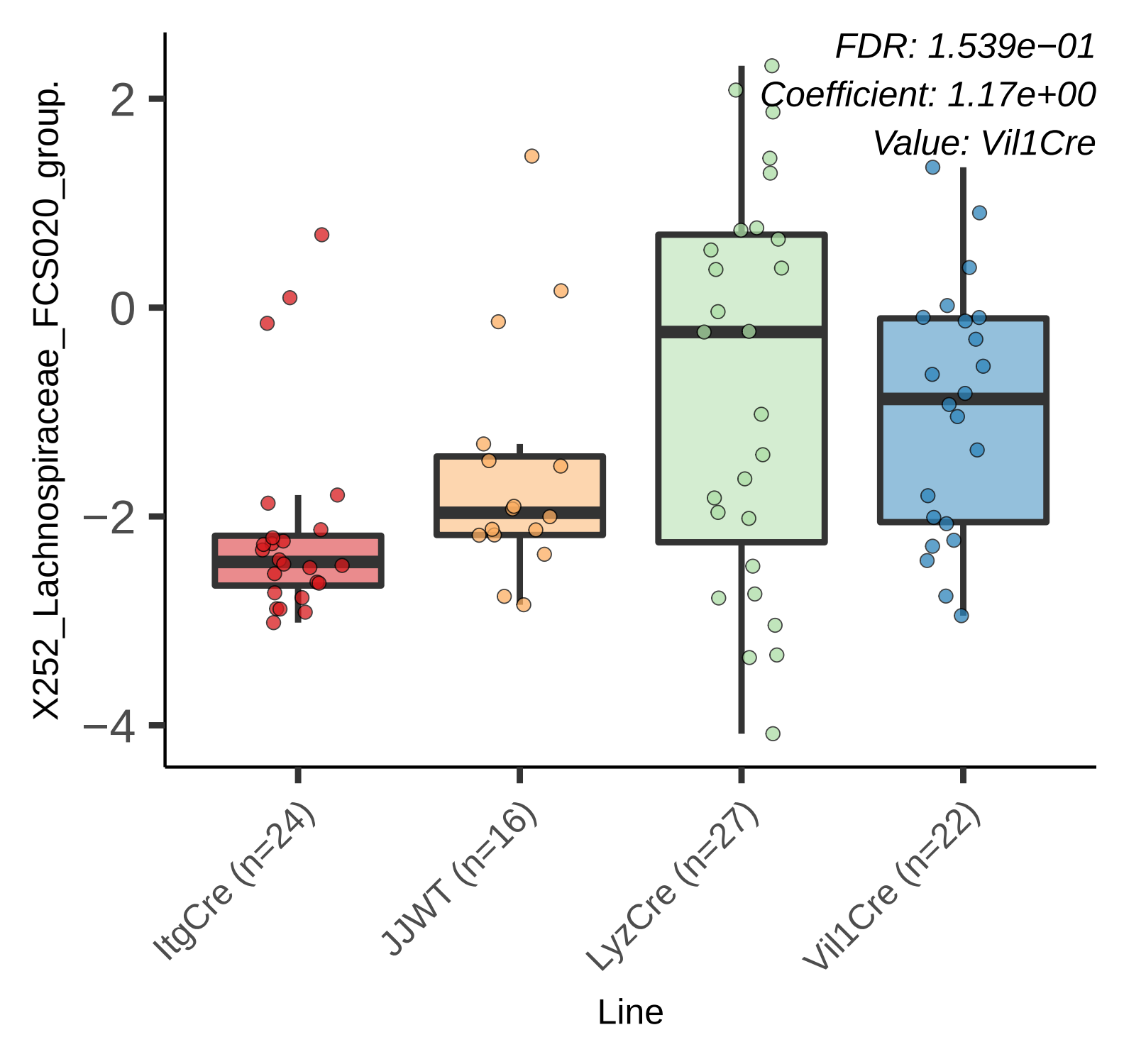

FDR: 1.501e−01
Coefficient: 8.14e−01
Value: Vil1Cre
X125_Ruminiclostridium_5.
1
0
−1
−2
−3
−4
ItgCre (n=24)
JJWT (n=16)
LyzCre (n=27)
Vil1Cre (n=22)
Line

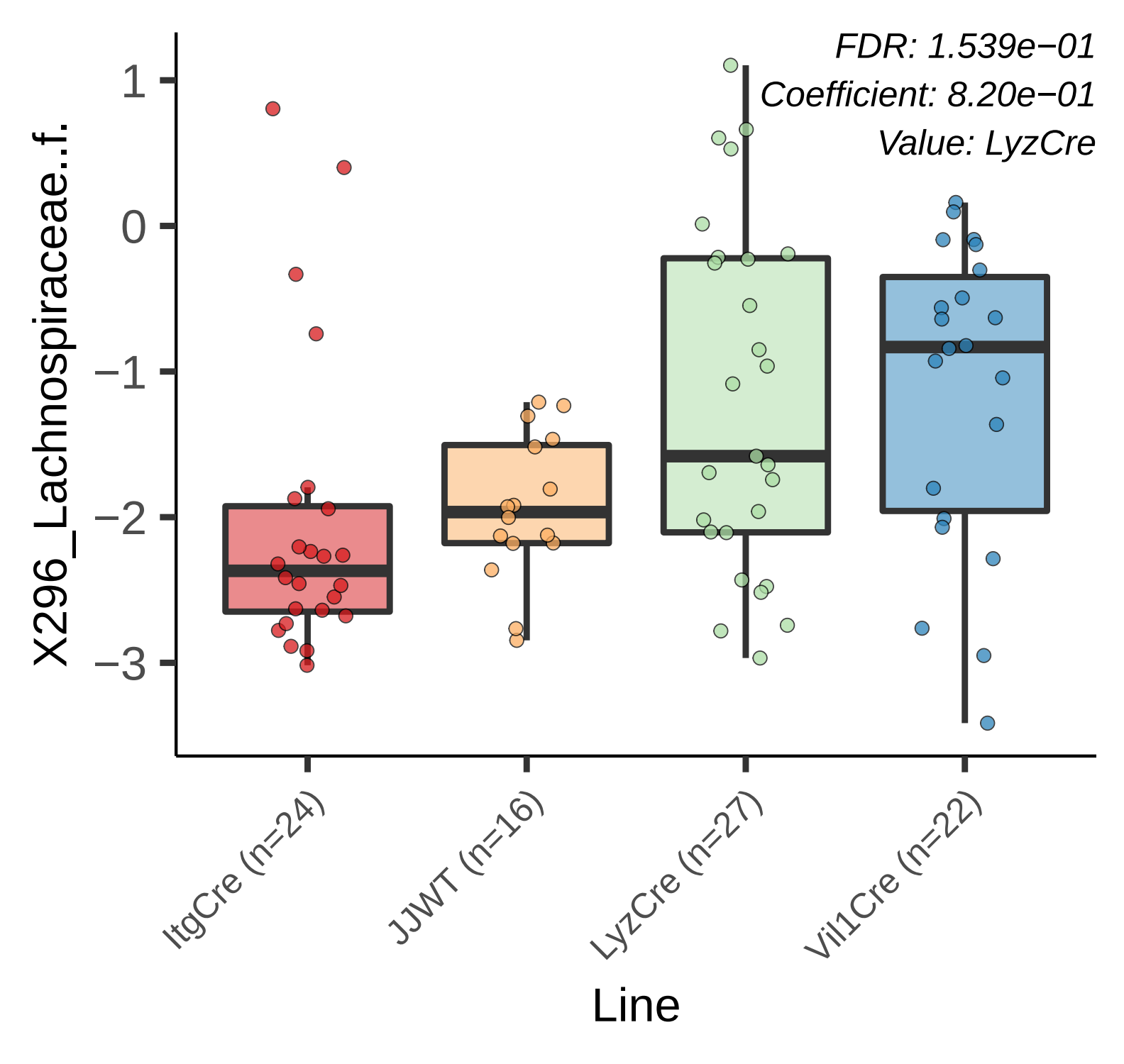

FDR: 1.527e−01
Coefficient: 1.83e+00
Value: JJWT
X69_Ruminococcaceae_UCG.014.
2
0
−2
ItgCre (n=24)
JJWT (n=16)
LyzCre (n=27)
Vil1Cre (n=22)
Line

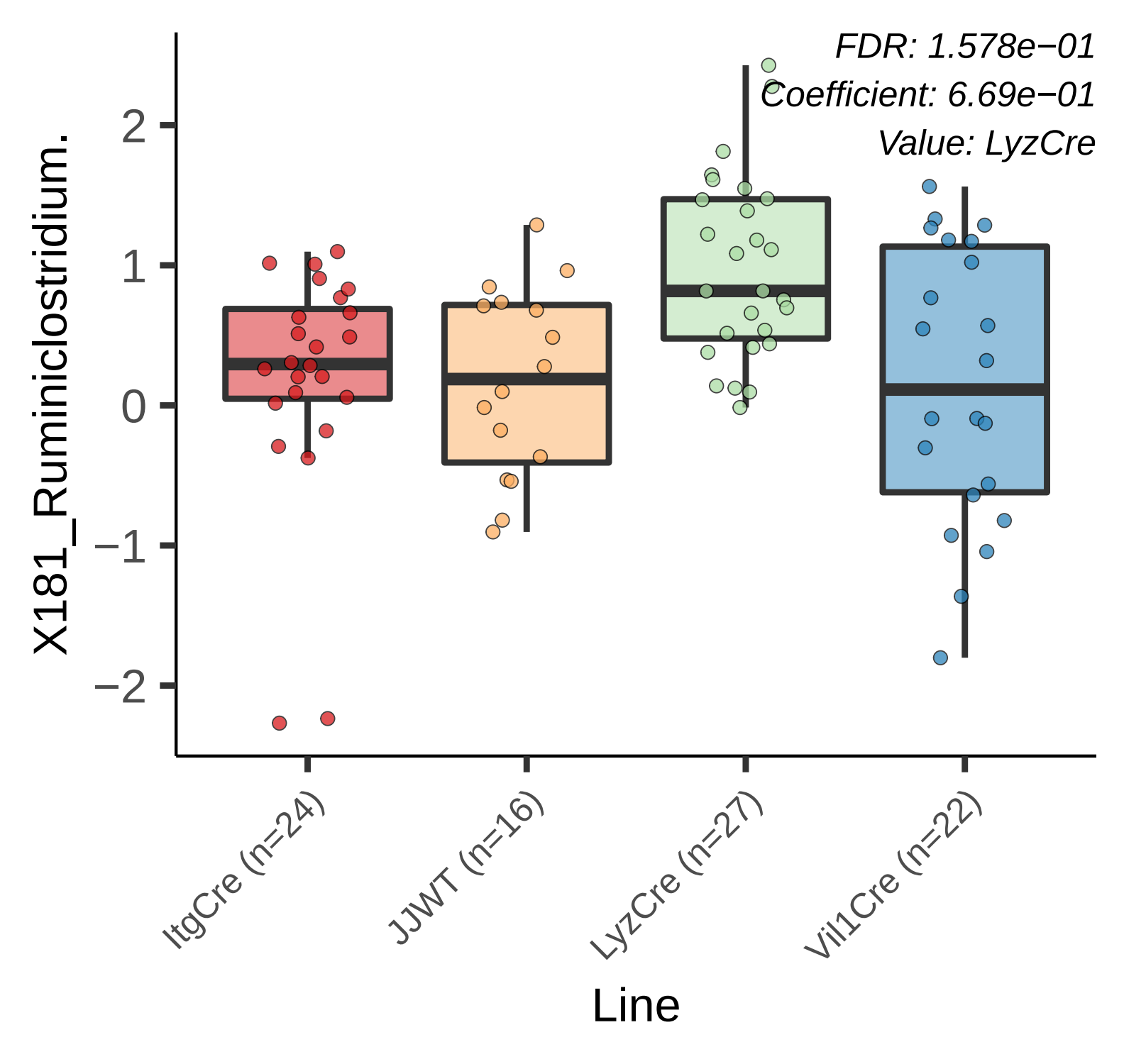

FDR: 1.527e−01
Coefficient: −1.34e+00
Value: JJWT
X133_Lachnospiraceae.f.
1
0
−1
−2
−3
ItgCre (n=24)
JJWT (n=16)
LyzCre (n=27)
Vil1Cre (n=22)
Line

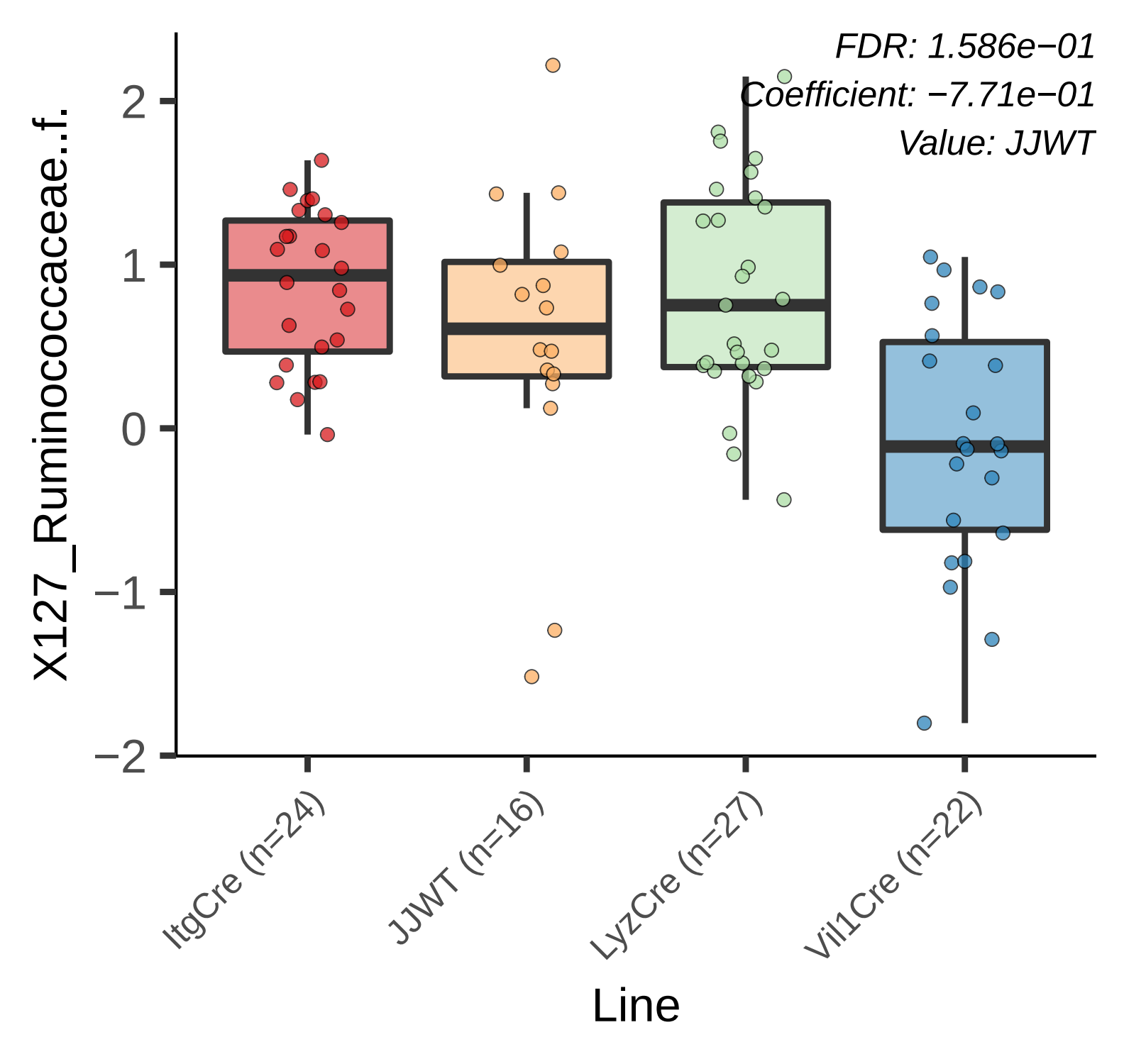

FDR: 1.533e−01
Coefficient: 1.15e+00
Value: Vil1Cre
X179_Bacteroides.
5.0
2.5
0.0
−2.5
ItgCre (n=24)
JJWT (n=16)
LyzCre (n=27)
Vil1Cre (n=22)
Line

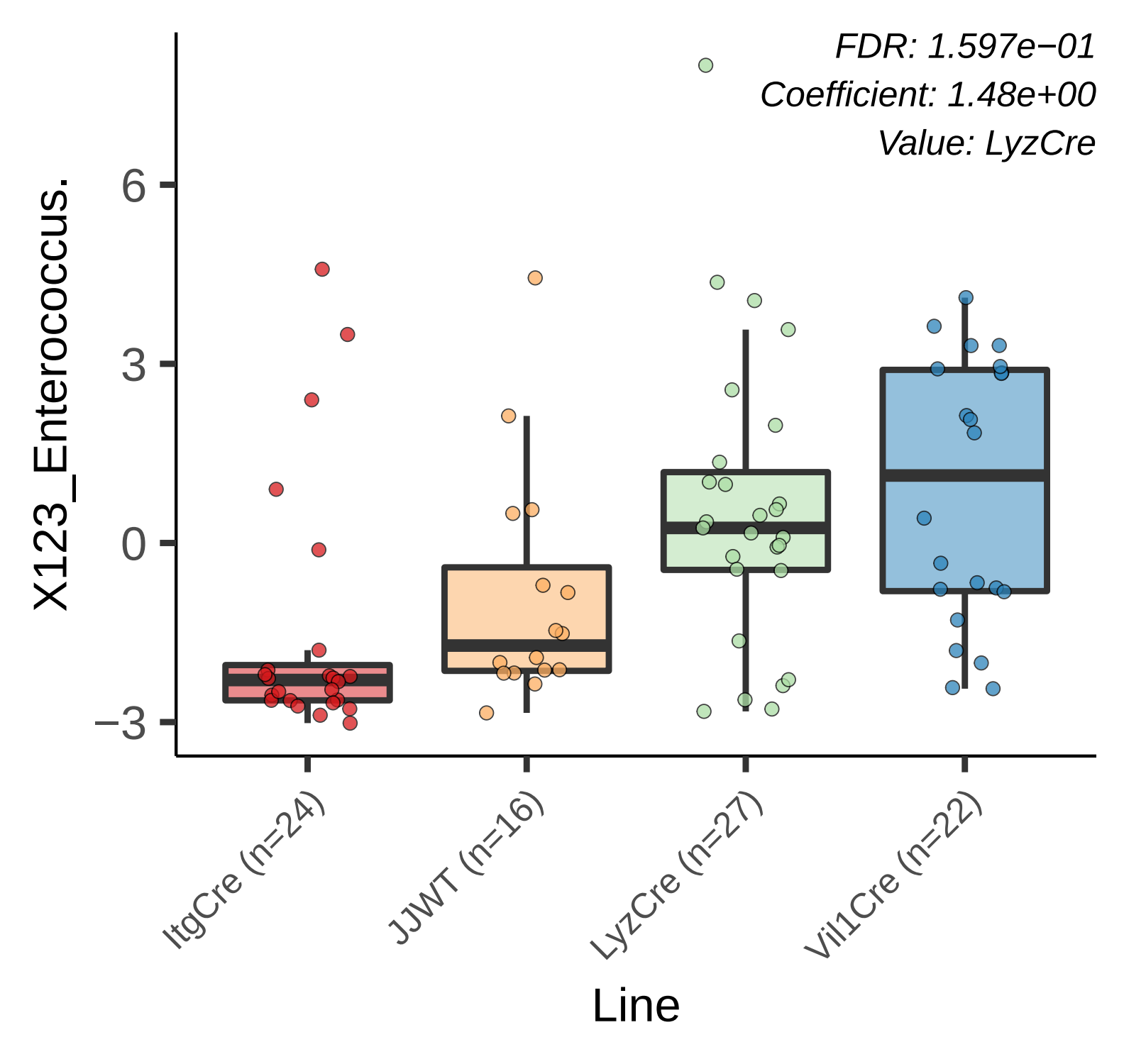

FDR: 1.538e−01
Coefficient: −1.10e+00
Value: JJWT
X180_Alistipes.
3
2
1
0
−1
−2
ItgCre (n=24)
JJWT (n=16)
LyzCre (n=27)
Vil1Cre (n=22)
Line

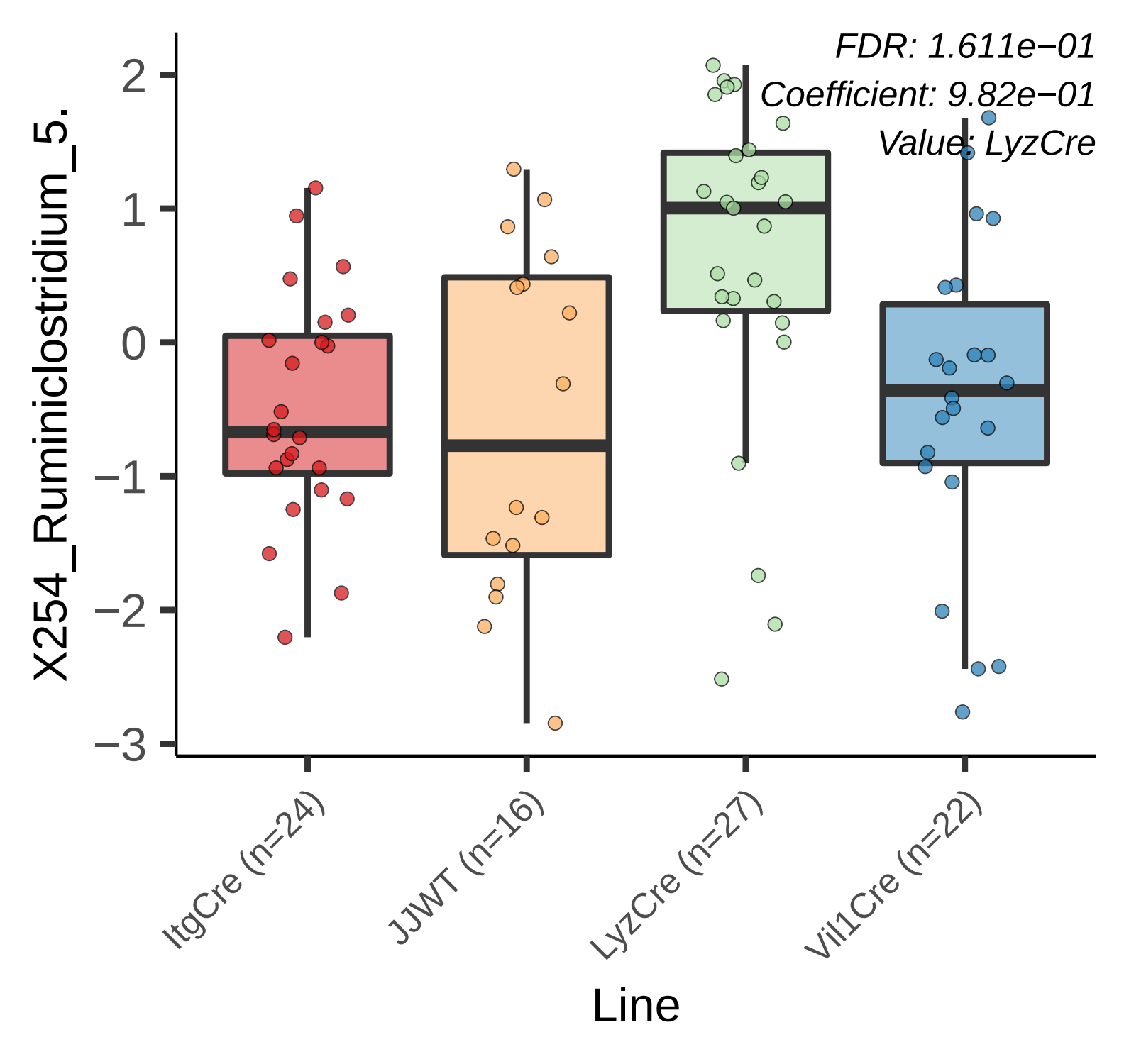

FDR: 1.539e−01
Coefficient: 1.17e+00
Value: Vil1Cre
X252_Lachnospiraceae_FCS020_group.
2
0
−2
−4
ItgCre (n=24)
JJWT (n=16)
LyzCre (n=27)
Vil1Cre (n=22)
Line

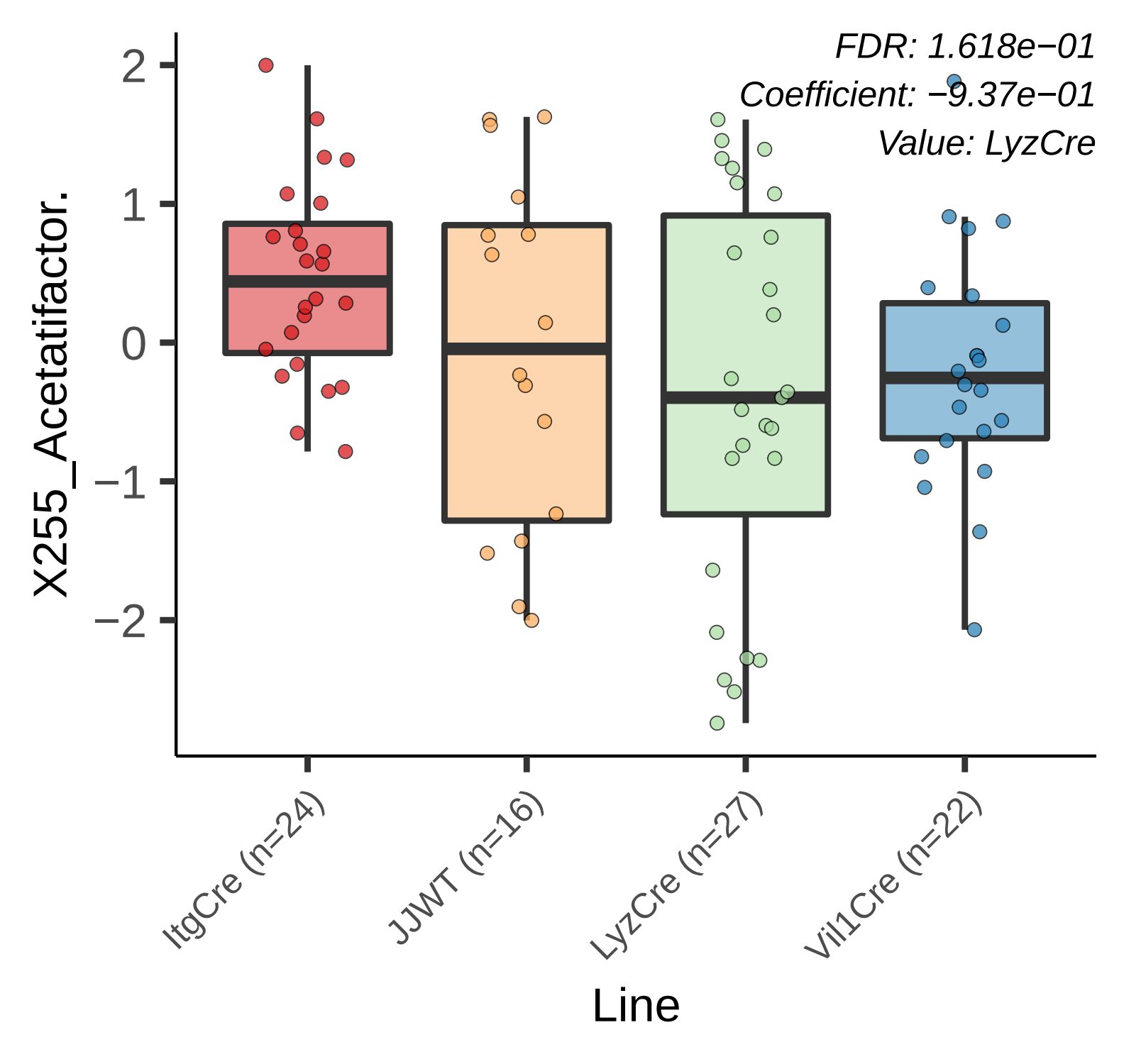

FDR: 1.539e−01
Coefficient: 8.20e−01
Value: LyzCre
X296_Lachnospiraceae.f.
1
0
−1
−2
−3
ItgCre (n=24)
JJWT (n=16)
LyzCre (n=27)
Vil1Cre (n=22)
Line

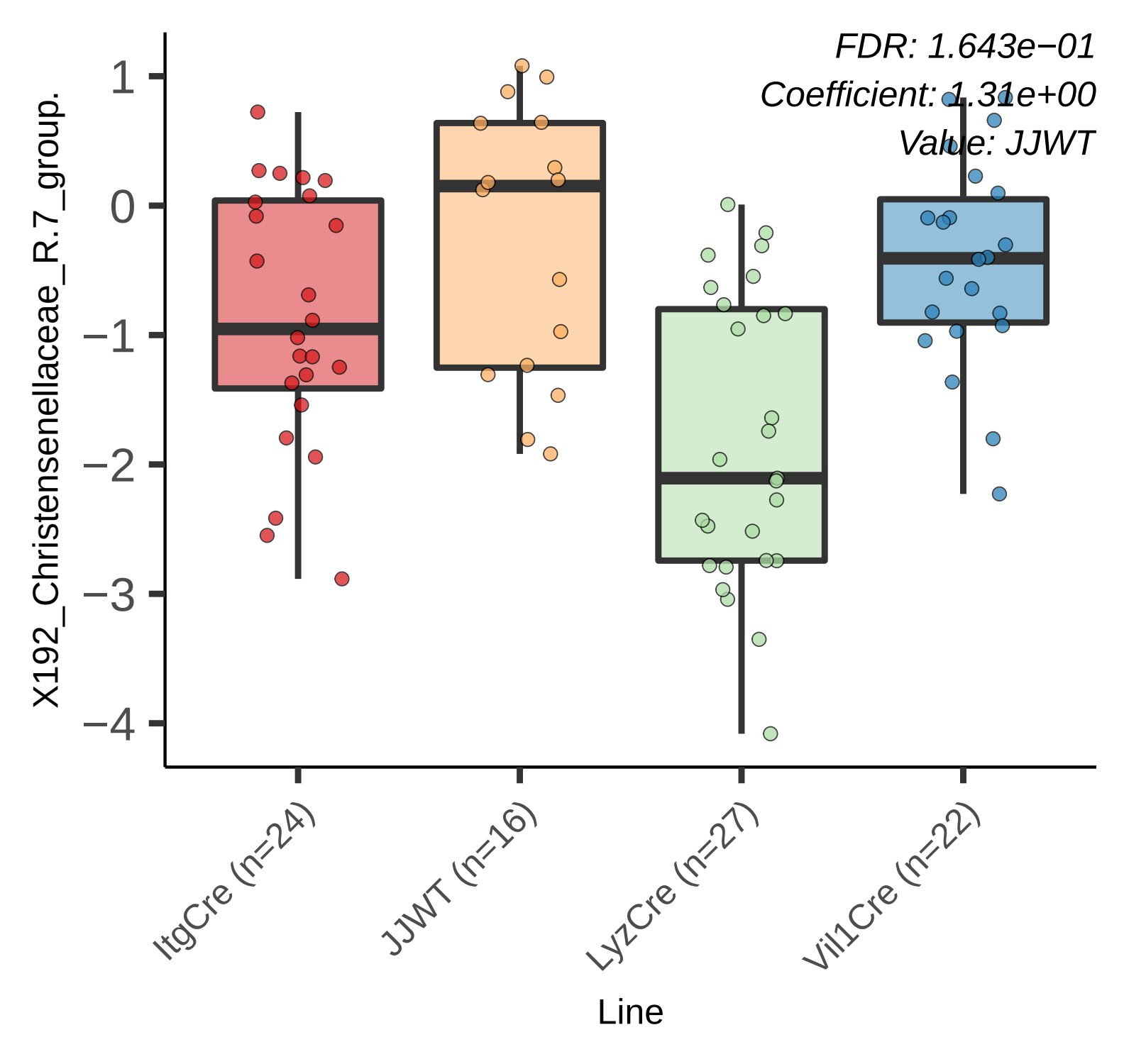

FDR: 1.578e−01
Coefficient: 6.69e−01
Value: LyzCre
X181_Ruminiclostridium.
2
1
0
−1
−2
ItgCre (n=24)
JJWT (n=16)
LyzCre (n=27)
Vil1Cre (n=22)
Line

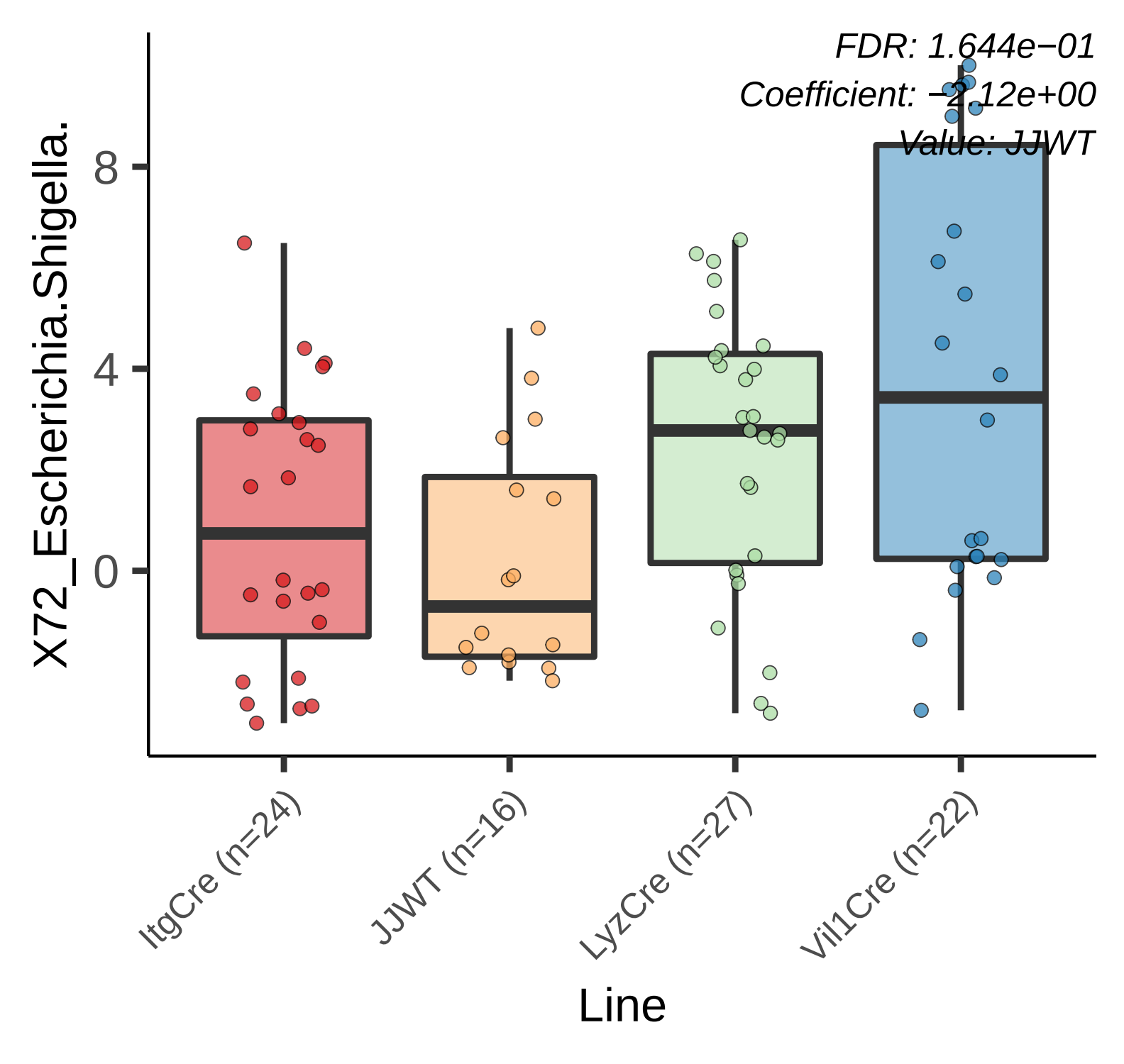

FDR: 1.586e−01
Coefficient: −7.71e−01
Value: JJWT
X127_Ruminococcaceae..f.
2
1
0
−1
−2
ItgCre (n=24)
JJWT (n=16)
LyzCre (n=27)
Vil1Cre (n=22)
Line

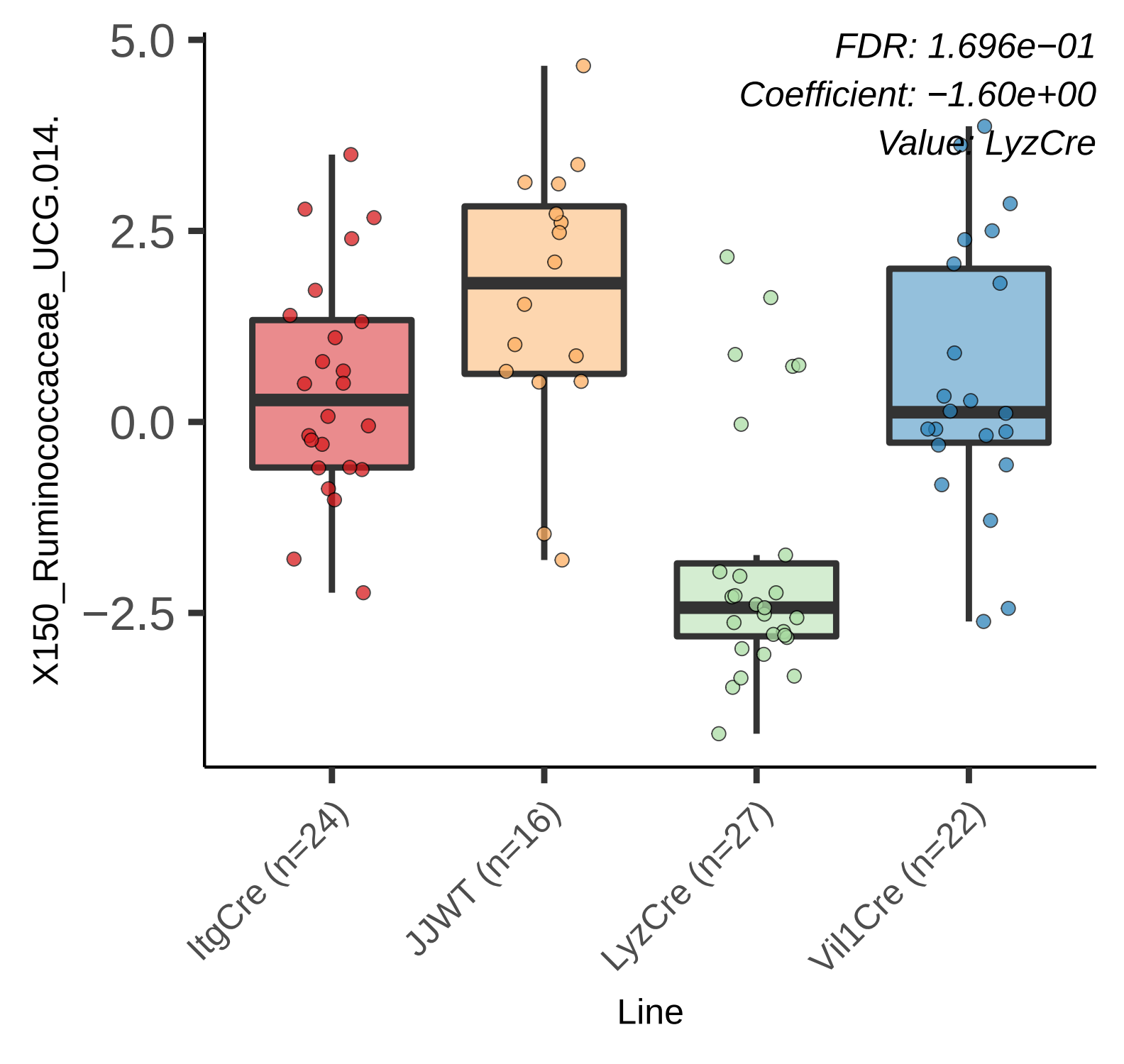

FDR: 1.597e−01
Coefficient: 1.48e+00
Value: LyzCre
X123_Enterococcus.
6
3
0
−3
ItgCre (n=24)
JJWT (n=16)
LyzCre (n=27)
Vil1Cre (n=22)
Line

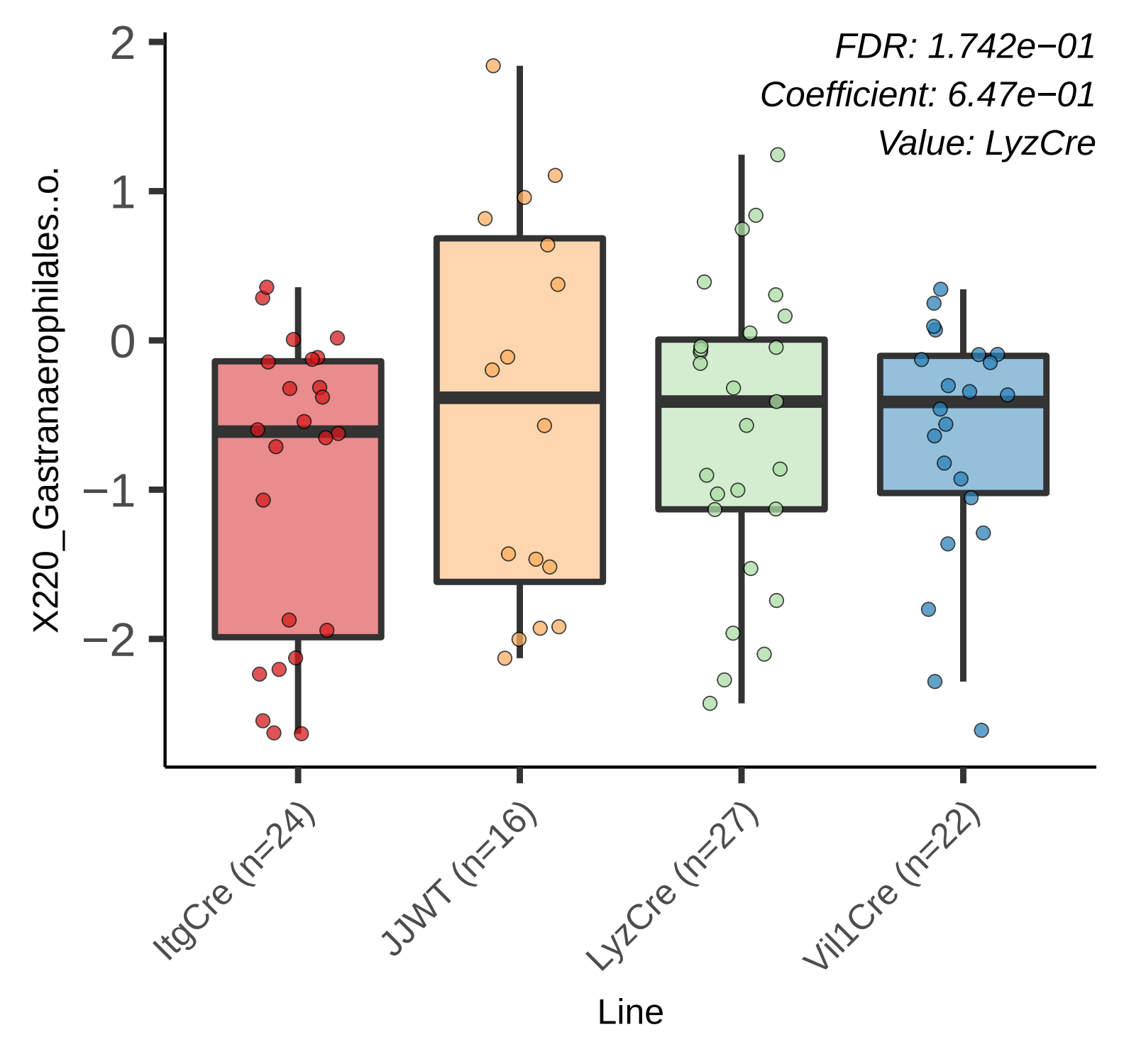

FDR: 1.611e−01
Coefficient: 9.82e−01
Value: LyzCre
X254_Ruminiclostridium_5.
2
1
0
−1
−2
−3
ItgCre (n=24)
JJWT (n=16)
LyzCre (n=27)
Vil1Cre (n=22)
Line

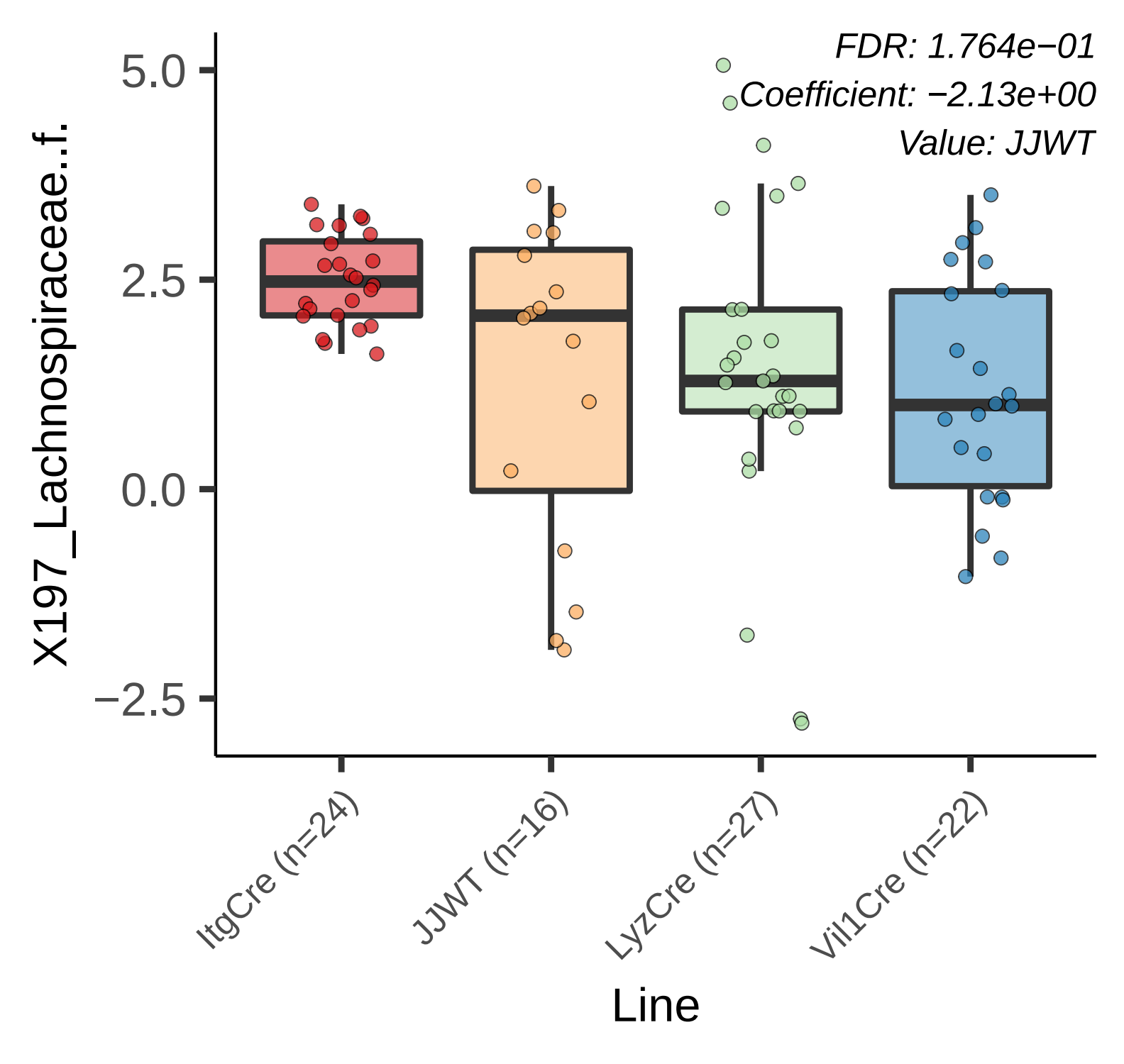

FDR: 1.618e−01
Coefficient: −9.37e−01
Value: LyzCre
X255_Acetatifactor.
2
1
0
−1
−2
ItgCre (n=24)
JJWT (n=16)
LyzCre (n=27)
Vil1Cre (n=22)
Line

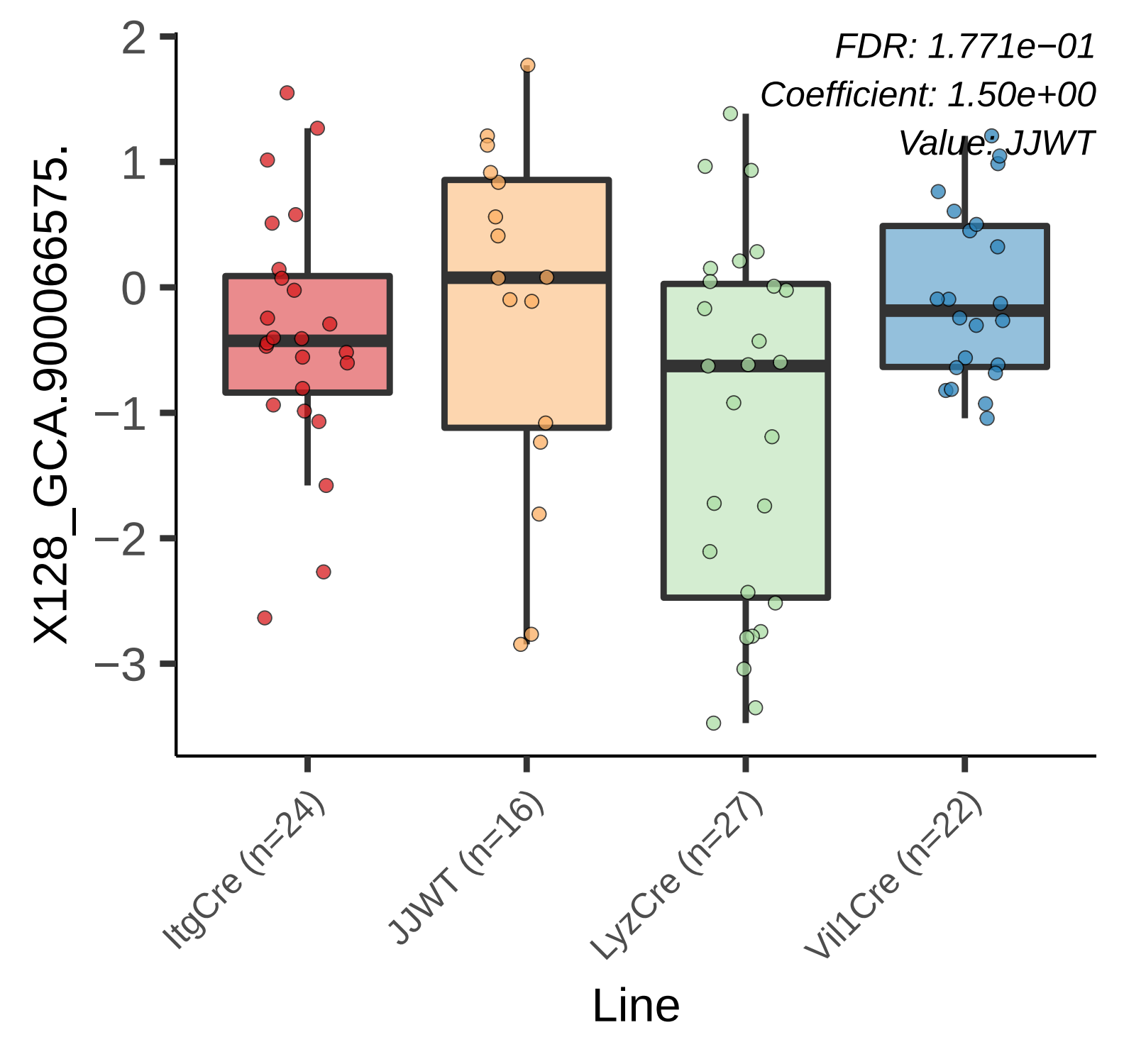

FDR: 1.643e−01
Coefficient: 1.31e+00
Value: JJWT
X192_Christensenellaceae_R.7_group.
1
0
−1
−2
−3
−4
ItgCre (n=24)
JJWT (n=16)
LyzCre (n=27)
Vil1Cre (n=22)
Line

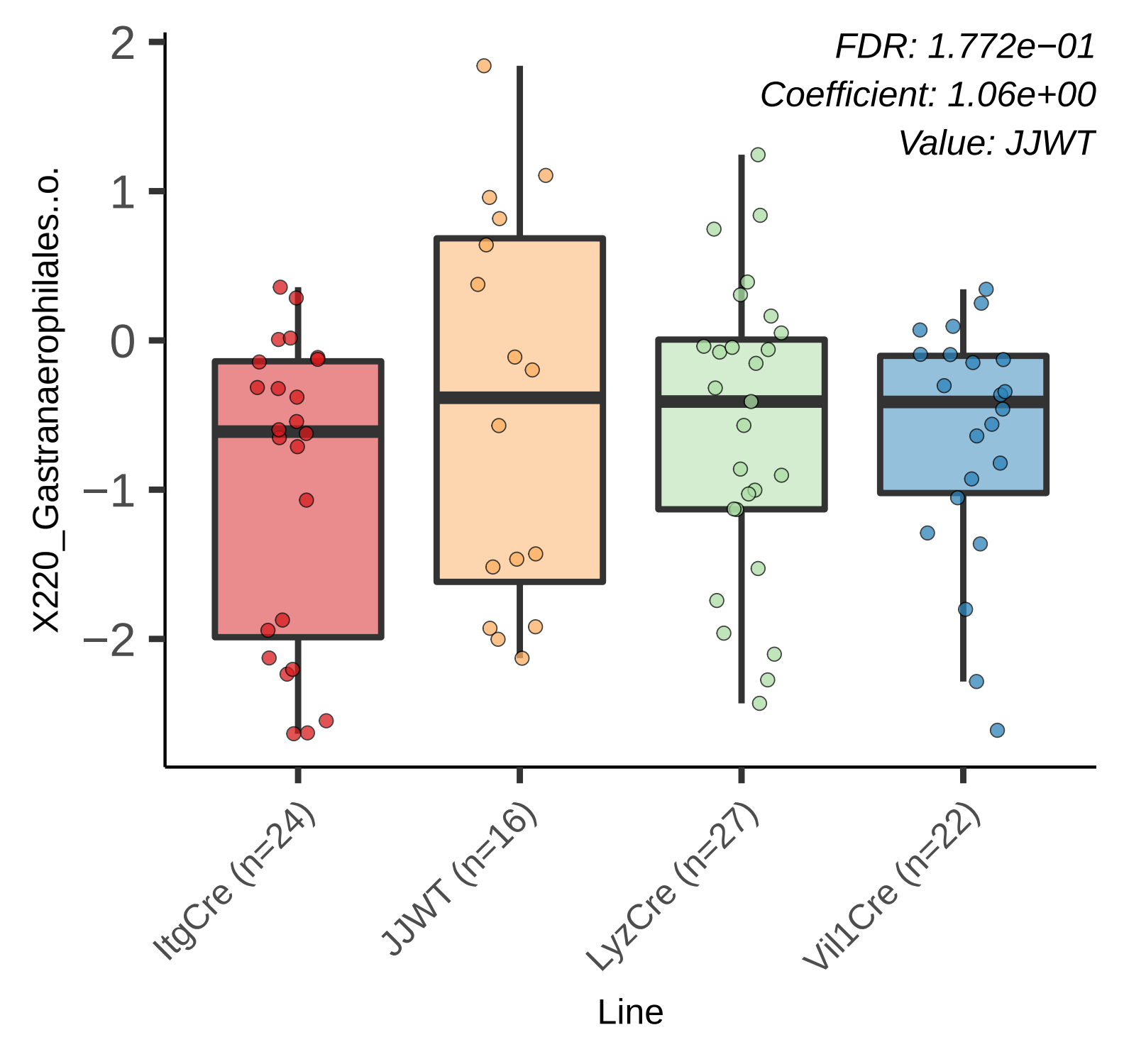

FDR: 1.644e−01
Coefficient: −2.12e+00
Value: JJWT
X72_Escherichia.Shigella.
8
4
0
ItgCre (n=24)
JJWT (n=16)
LyzCre (n=27)
Vil1Cre (n=22)
Line

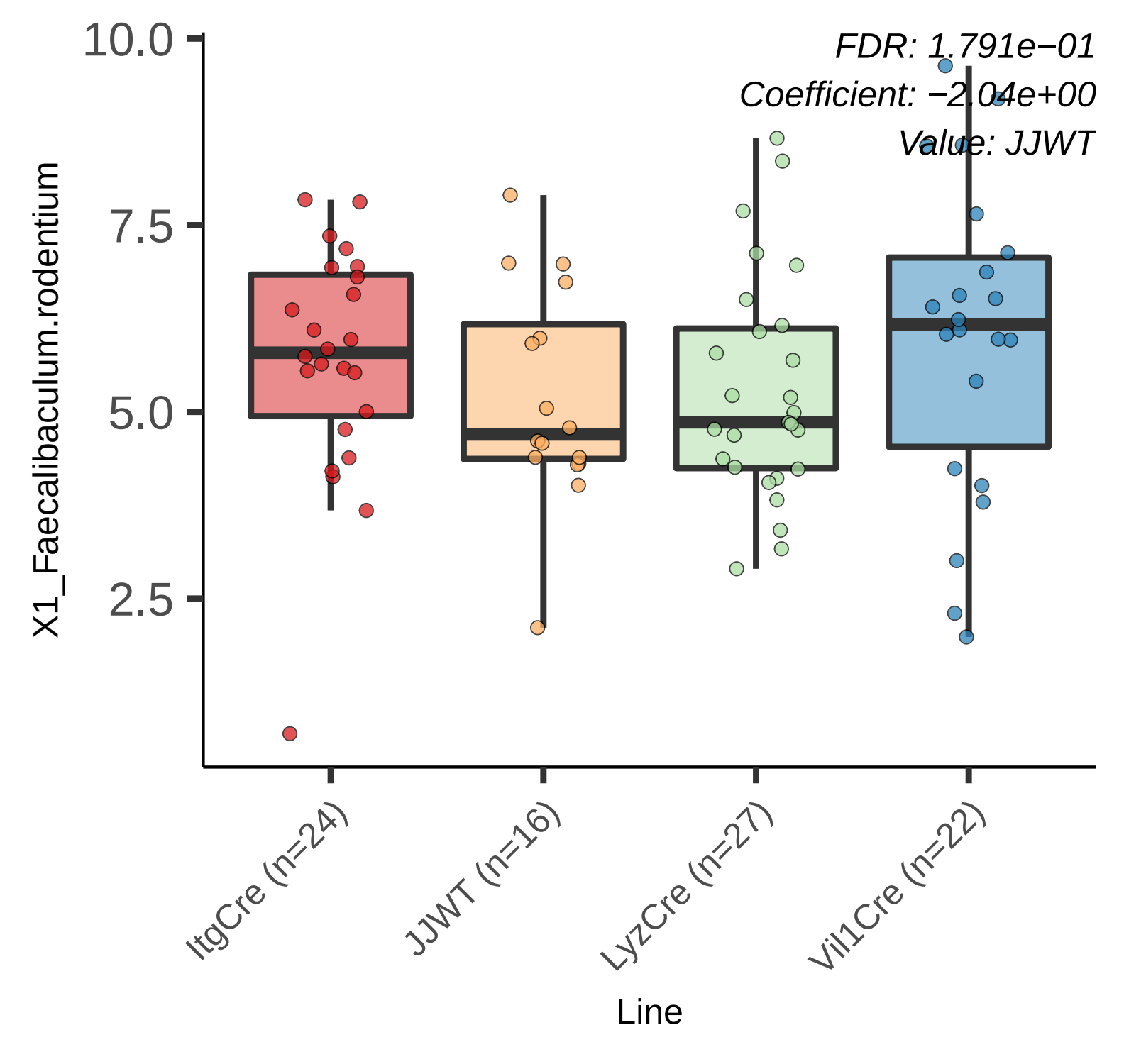

FDR: 1.696e−01
Coefficient: −1.60e+00
Value: LyzCre
X150_Ruminococcaceae_UCG.014.
5.0
2.5
0.0
−2.5
ItgCre (n=24)
JJWT (n=16)
LyzCre (n=27)
Vil1Cre (n=22)
Line

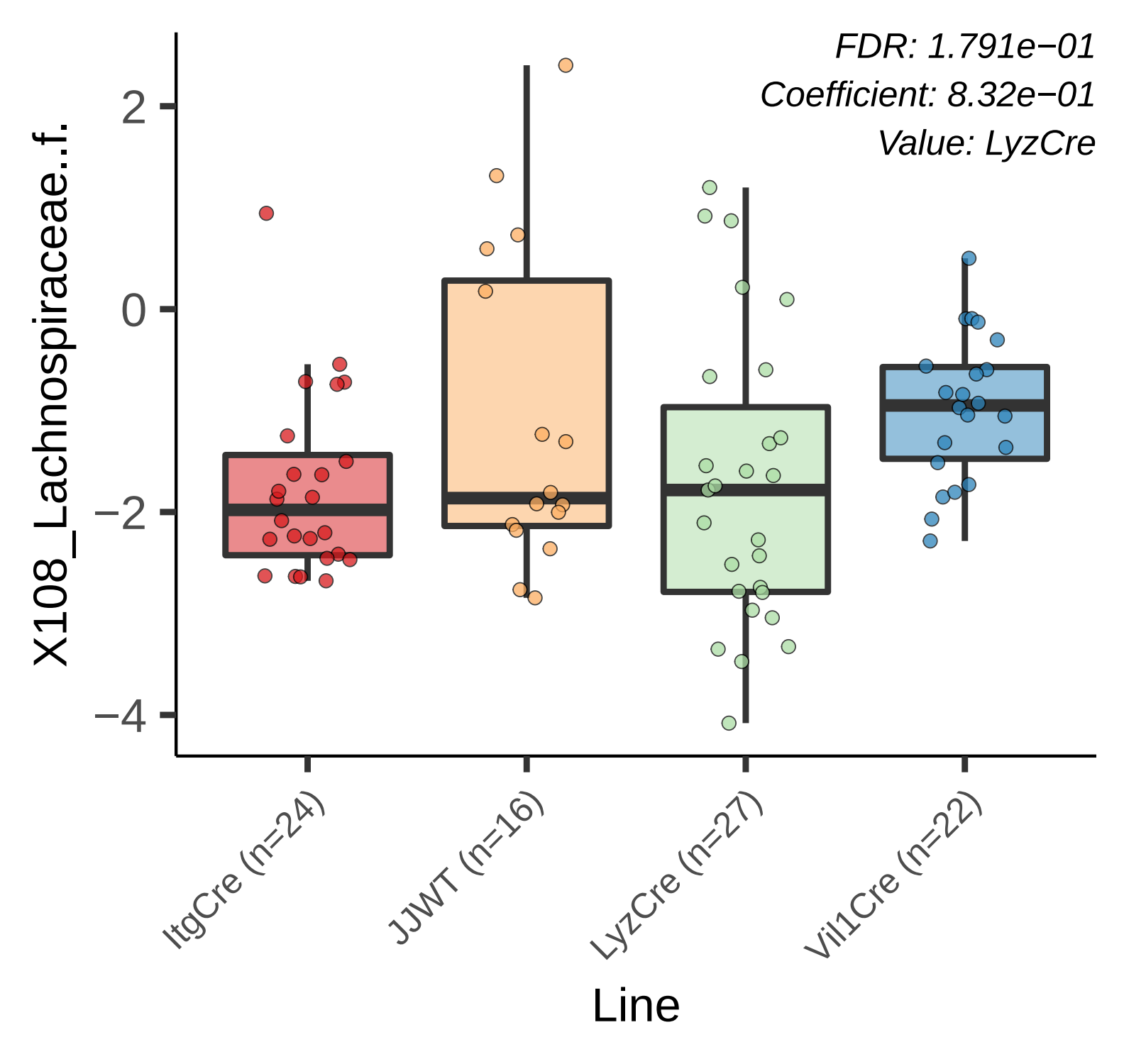

FDR: 1.742e−01
Coefficient: 6.47e−01
Value: LyzCre
X220_Gastranaerophilales..o.
2
1
0
−1
−2
ItgCre (n=24)
JJWT (n=16)
LyzCre (n=27)
Vil1Cre (n=22)
Line

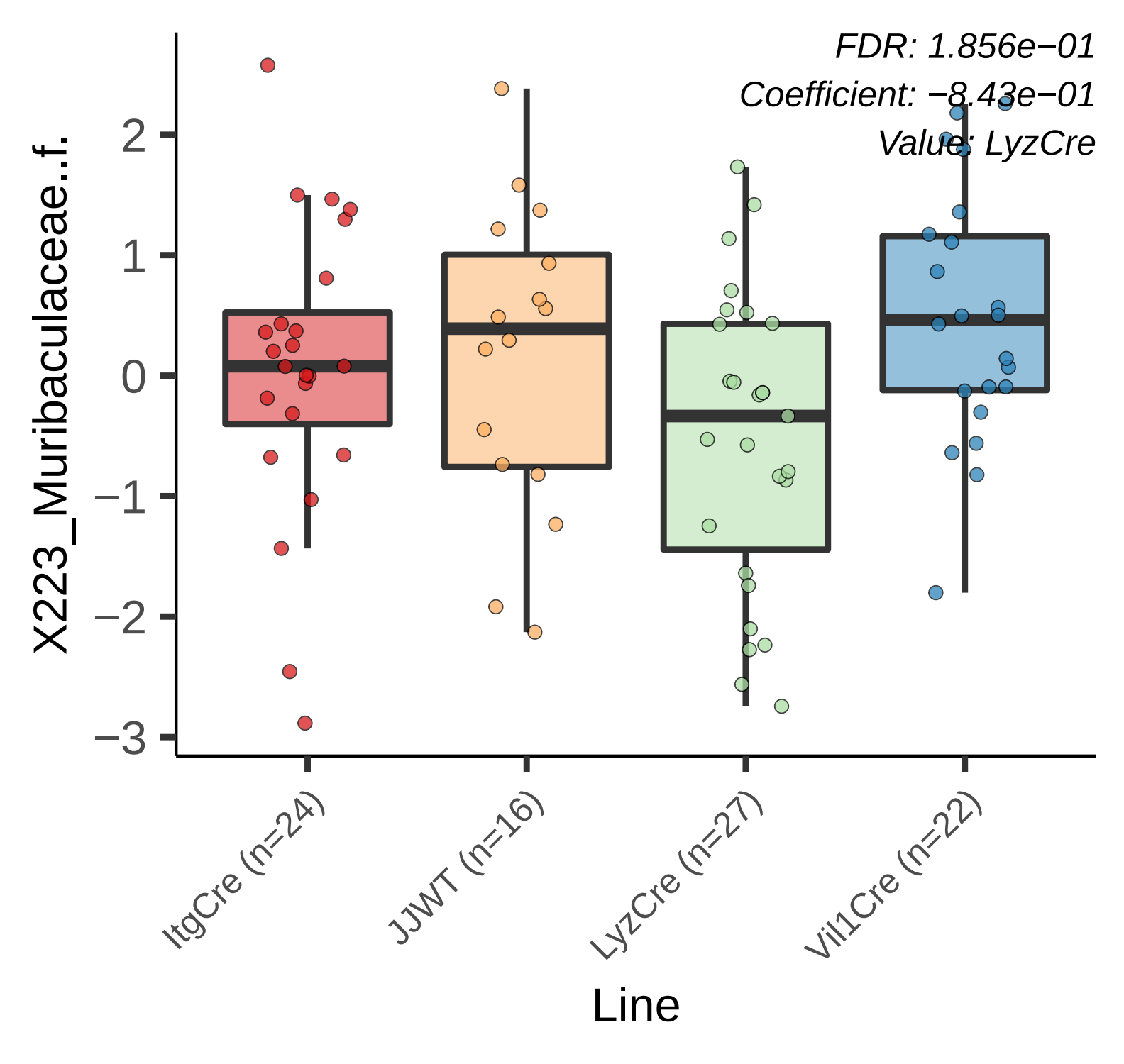

FDR: 1.764e−01
Coefficient: −2.13e+00
Value: JJWT
X197_Lachnospiraceae.f.
5.0
2.5
0.0
−2.5
ItgCre (n=24)
JJWT (n=16)
LyzCre (n=27)
Vil1Cre (n=22)
Line

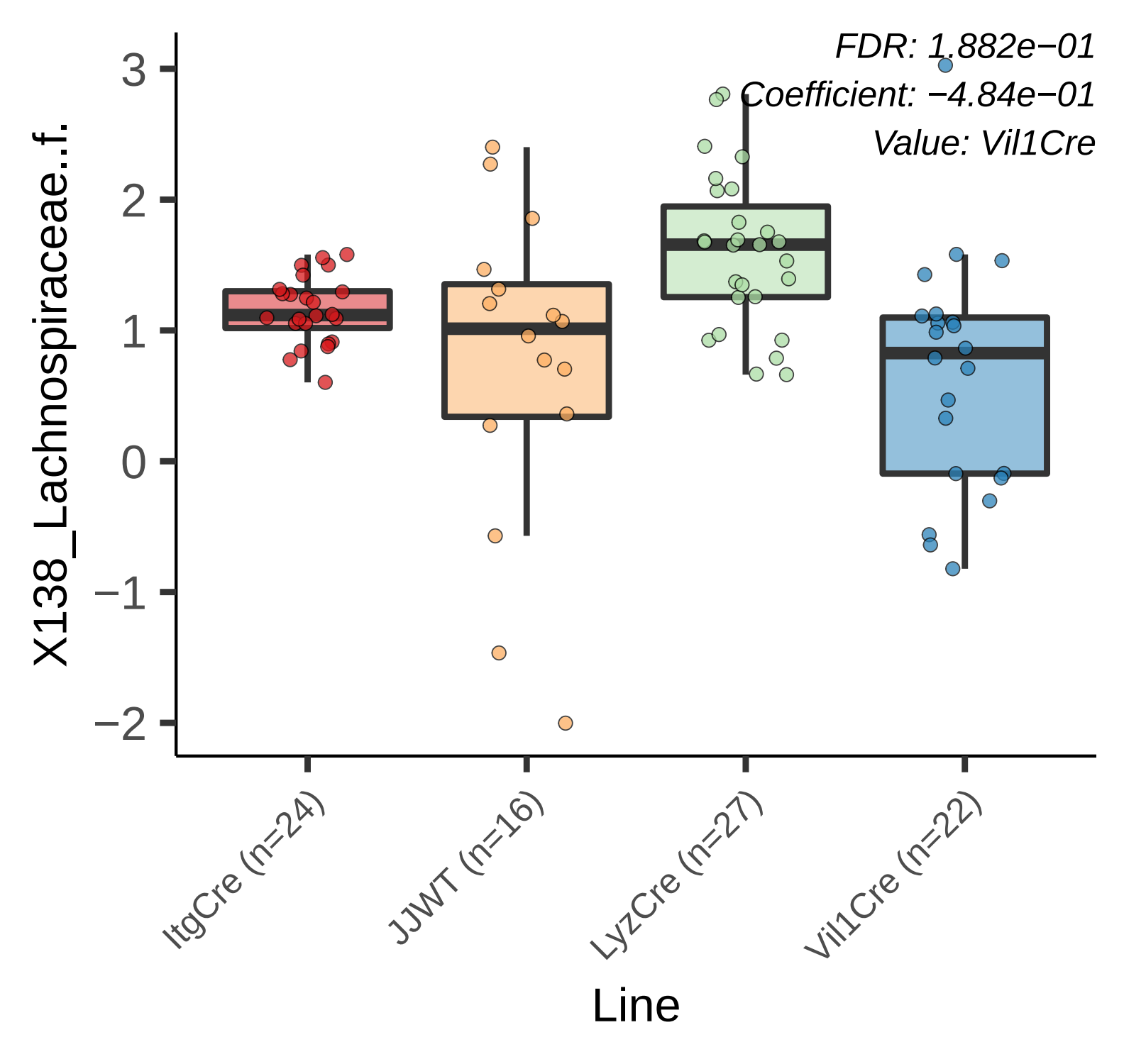

FDR: 1.771e−01
Coefficient: 1.50e+00
Value: JJWT
X128_GCA.900066575.
2
1
0
−1
−2
−3
ItgCre (n=24)
JJWT (n=16)
LyzCre (n=27)
Vil1Cre (n=22)
Line

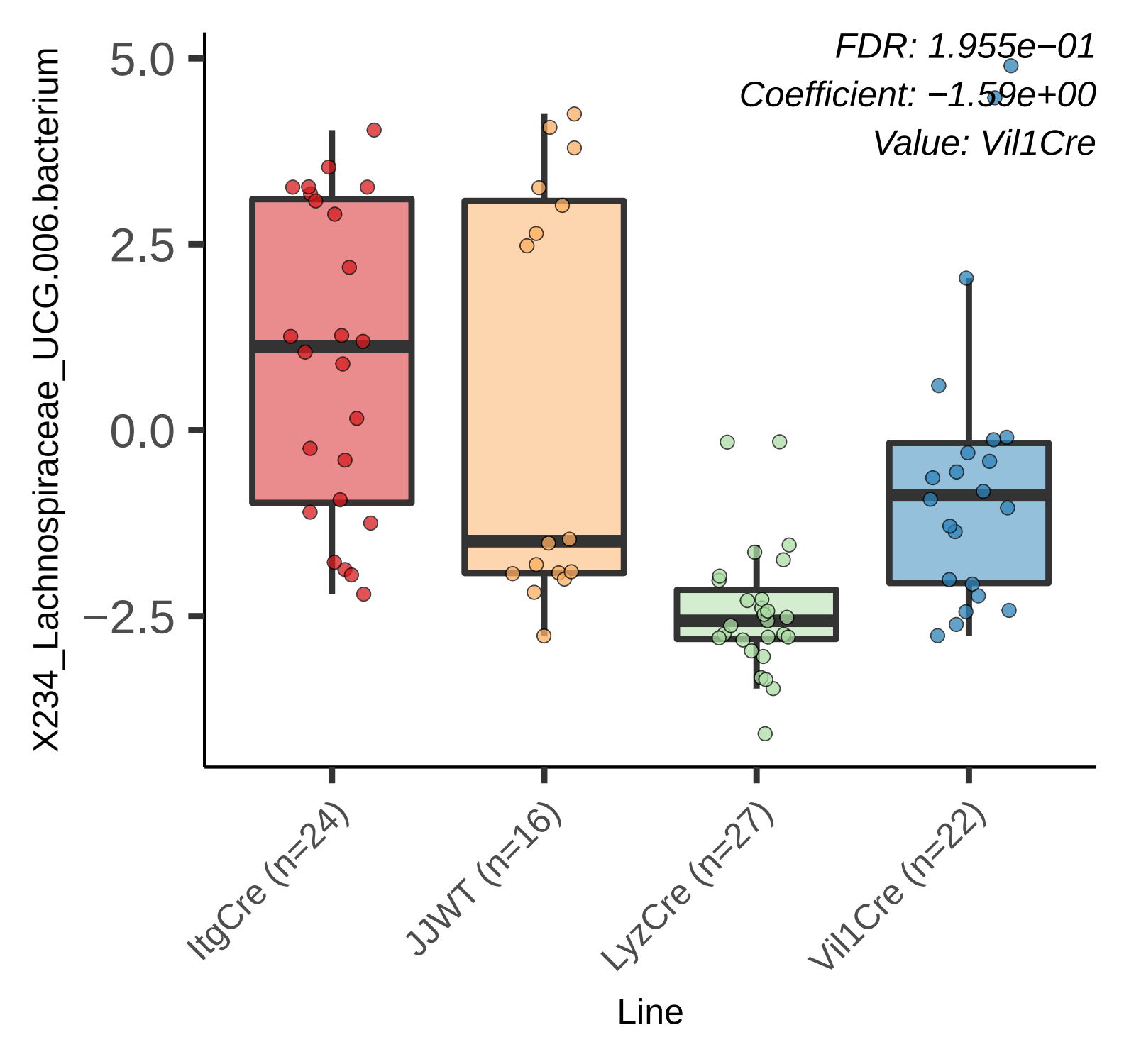

FDR: 1.772e−01
Coefficient: 1.06e+00
Value: JJWT
X220_Gastranaerophilales..o.
2
1
0
−1
−2
ItgCre (n=24)
JJWT (n=16)
LyzCre (n=27)
Vil1Cre (n=22)
Line

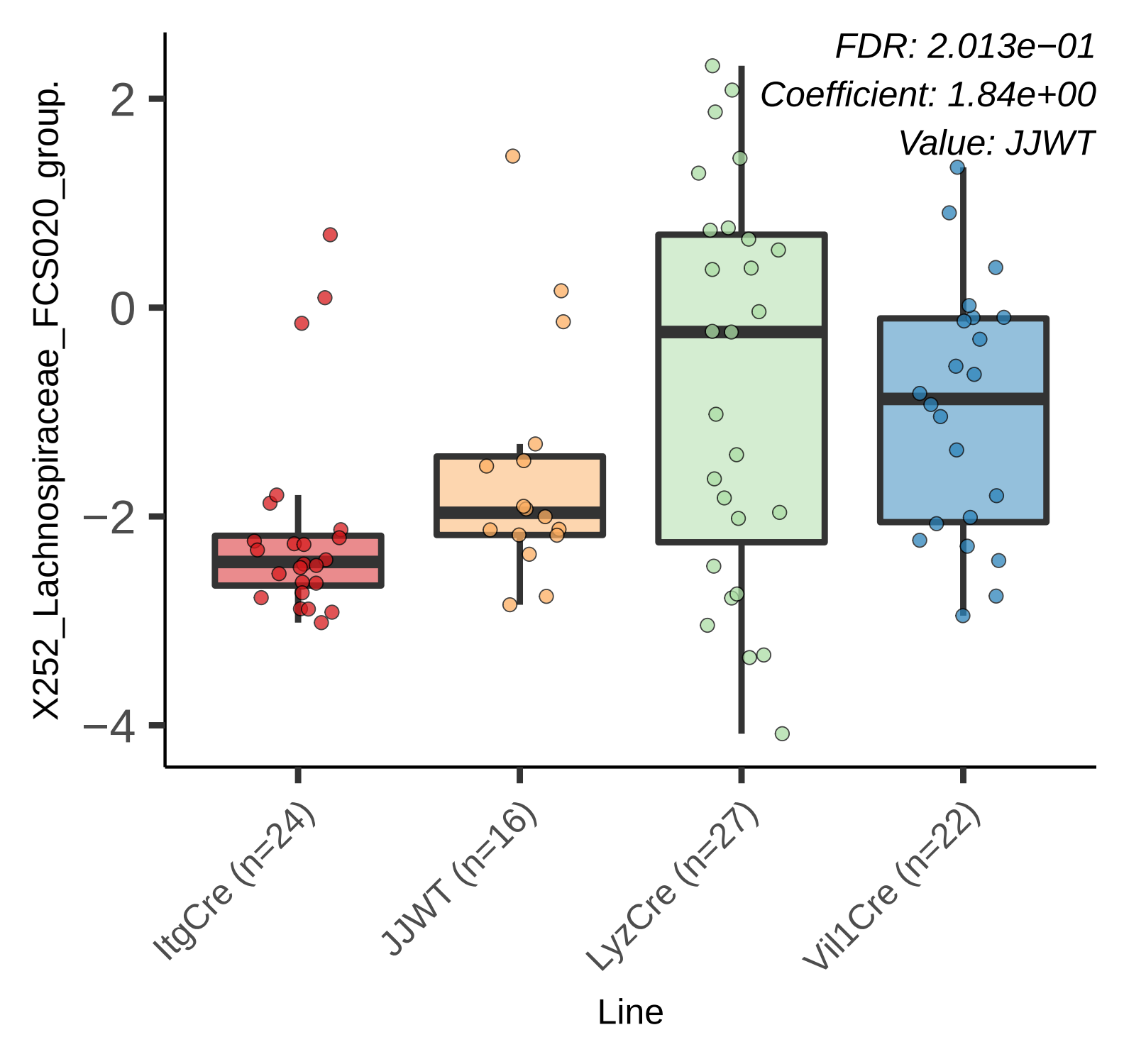

FDR: 1.791e−01
Coefficient: −2.04e+00
Value: JJWT
X1_Faecalibaculum.rodentium
10.0
7.5
5.0
2.5
ItgCre (n=24)
JJWT (n=16)
LyzCre (n=27)
Vil1Cre (n=22)
Line

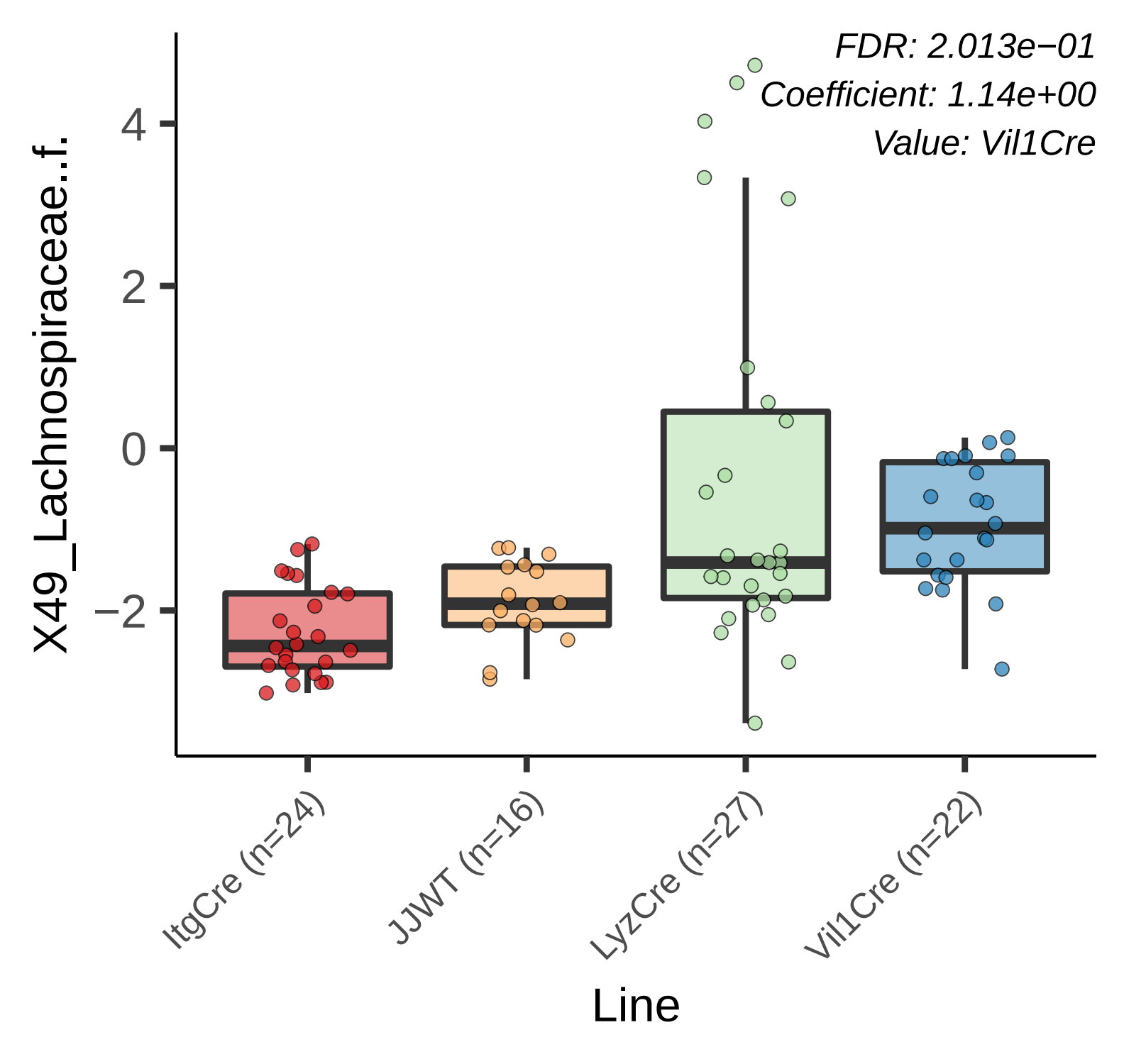

FDR: 1.791e−01
Coefficient: 8.32e−01
Value: LyzCre
X108_Lachnospiraceae.f.
2
0
−2
−4
ItgCre (n=24)
JJWT (n=16)
LyzCre (n=27)
Vil1Cre (n=22)
Line

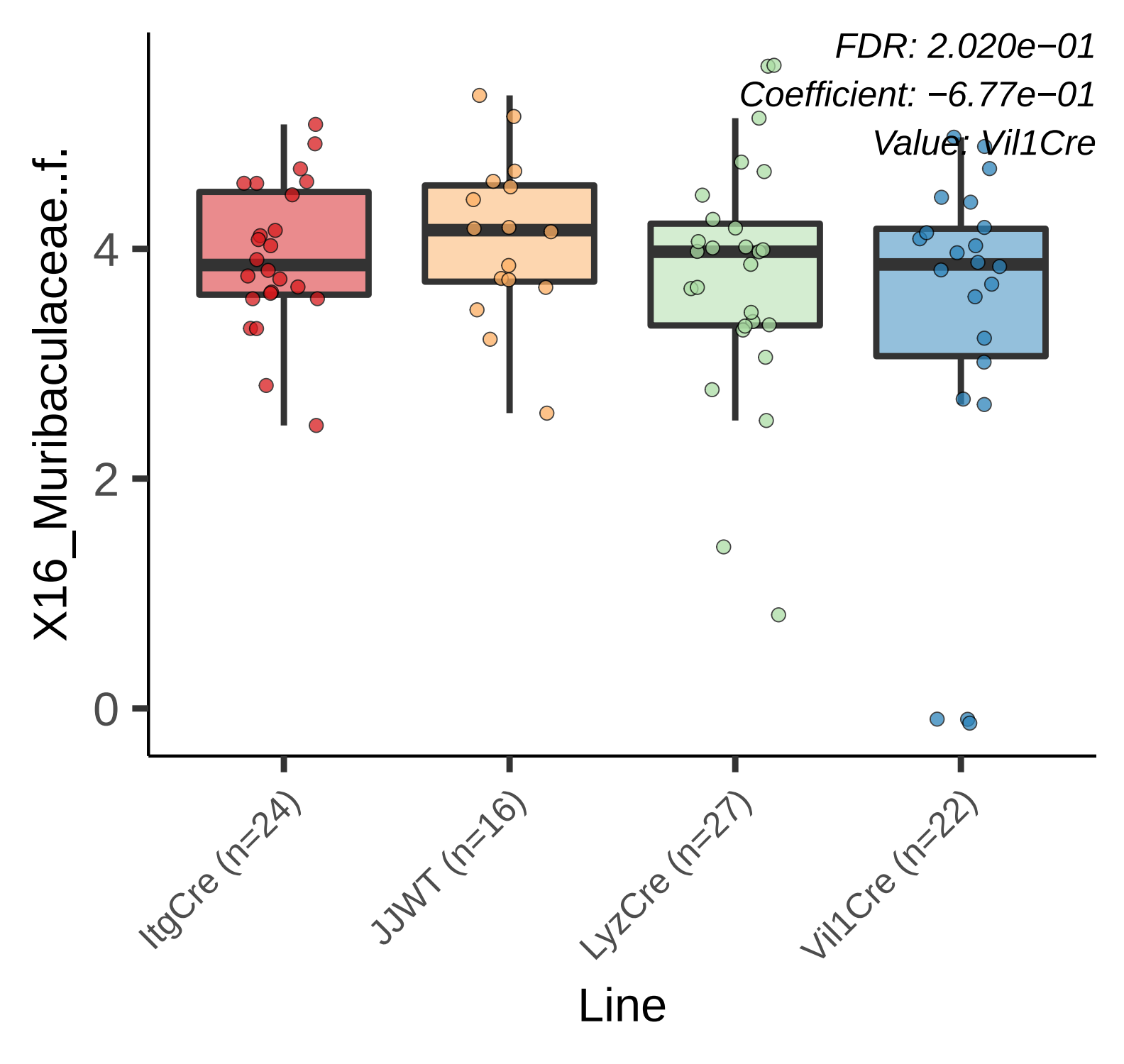

FDR: 1.856e−01
Coefficient: −8.43e−01
Value: LyzCre
X223_Muribaculaceae..f.
2
1
0
−1
−2
−3
ItgCre (n=24)
JJWT (n=16)
LyzCre (n=27)
Vil1Cre (n=22)
Line

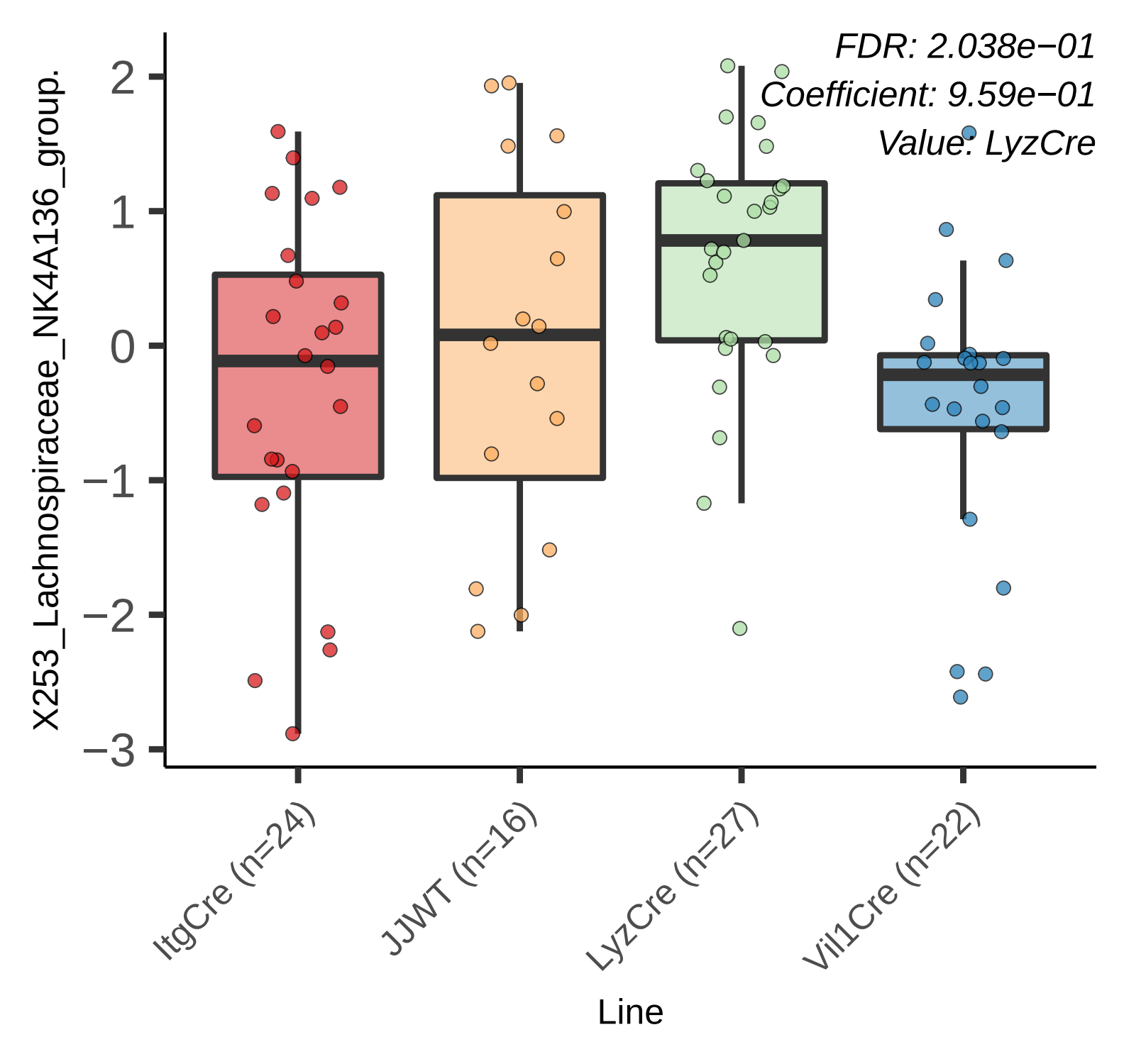

FDR: 1.882e−01
Coefficient: −4.84e−01
Value: Vil1Cre
X138_Lachnospiraceae.f.
3
2
1
0
−1
−2
ItgCre (n=24)
JJWT (n=16)
LyzCre (n=27)
Vil1Cre (n=22)
Line

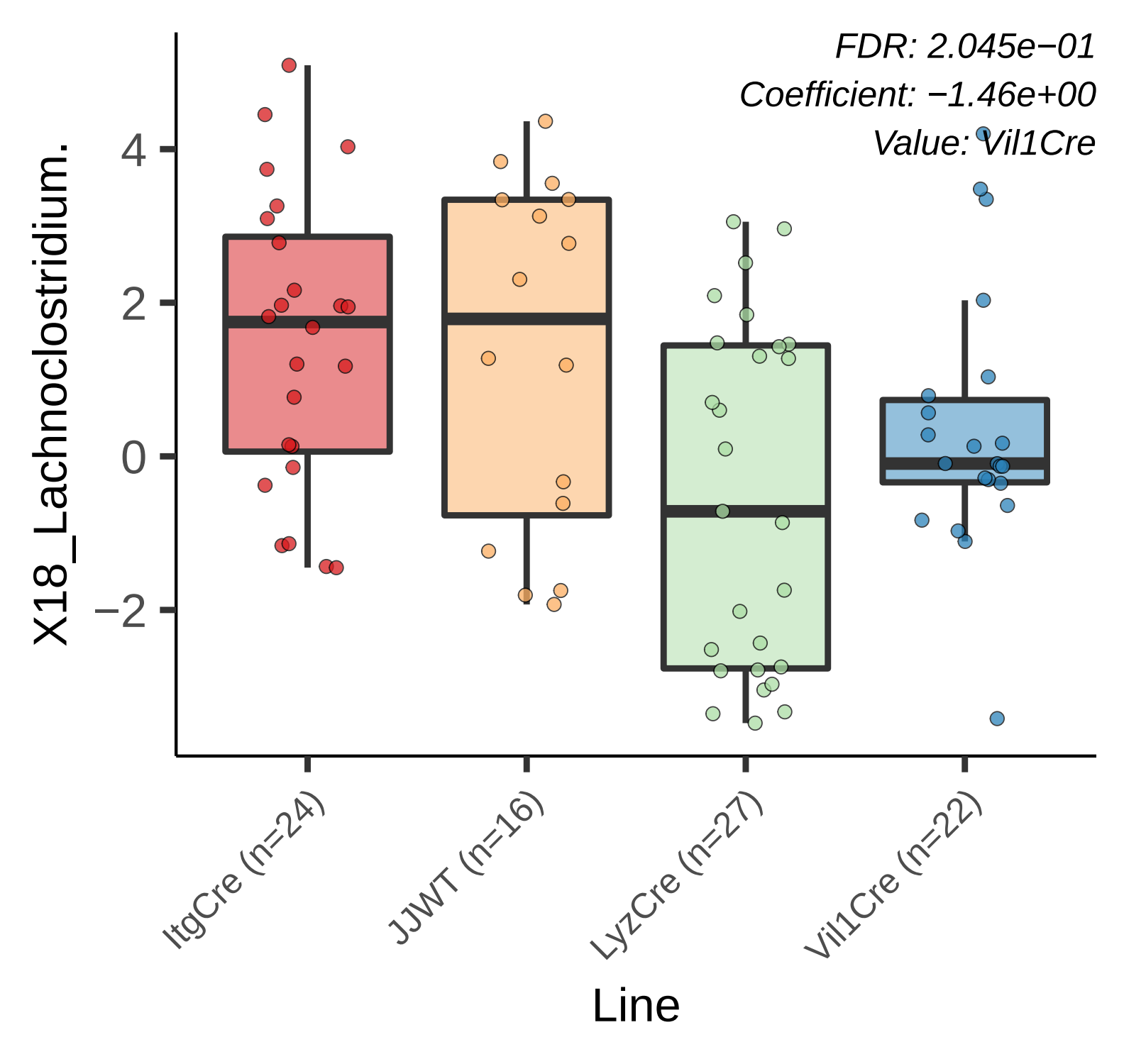

FDR: 1.955e−01
Coefficient: −1.59e+00
Value: Vil1Cre
X234_Lachnospiraceae_UCG.006.bacterium
5.0
2.5
0.0
−2.5
ItgCre (n=24)
JJWT (n=16)
LyzCre (n=27)
Vil1Cre (n=22)
Line

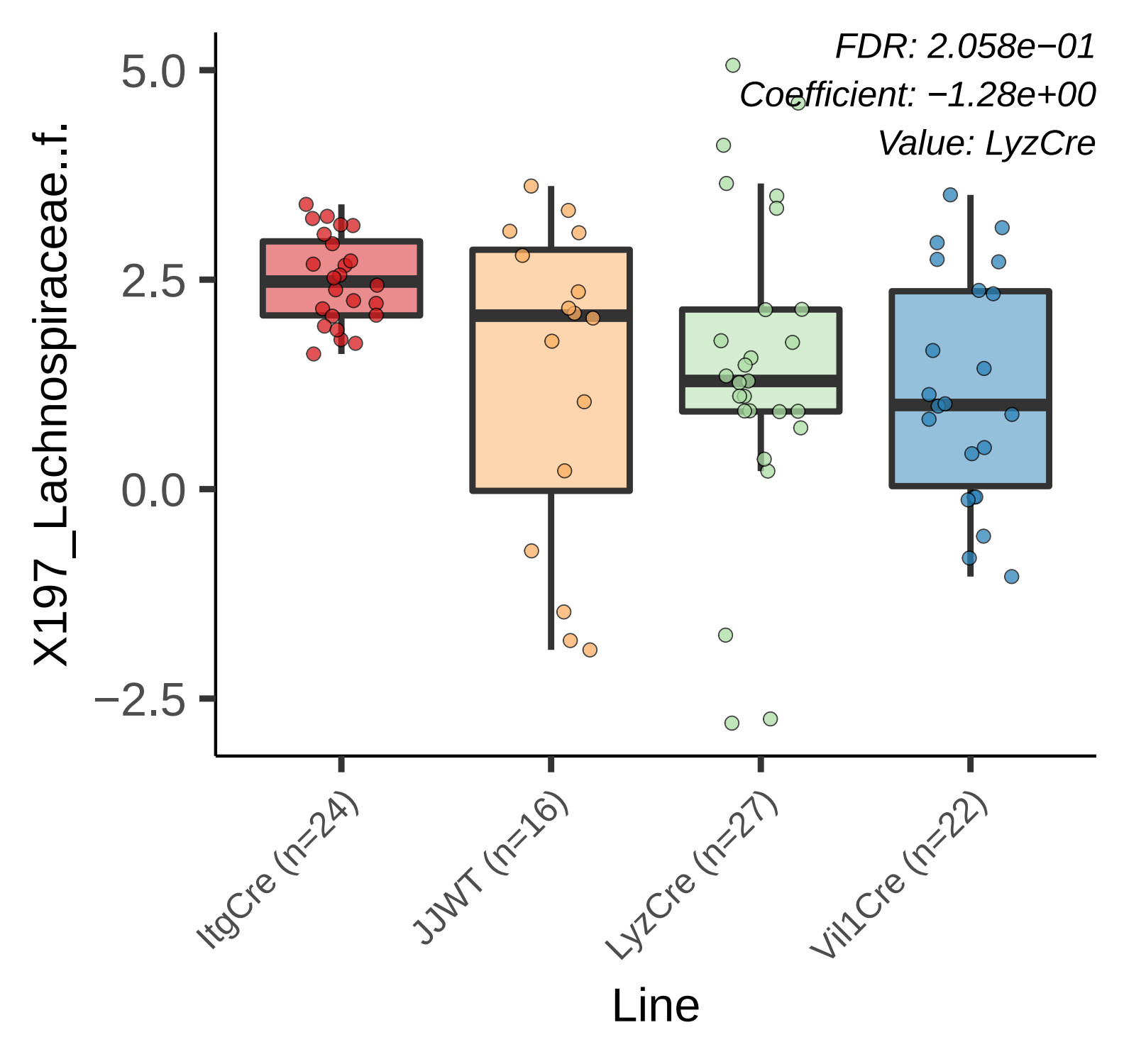

FDR: 2.013e−01
Coefficient: 1.84e+00
Value: JJWT
X252_Lachnospiraceae_FCS020_group.
2
0
−2
−4
ItgCre (n=24)
JJWT (n=16)
LyzCre (n=27)
Vil1Cre (n=22)
Line

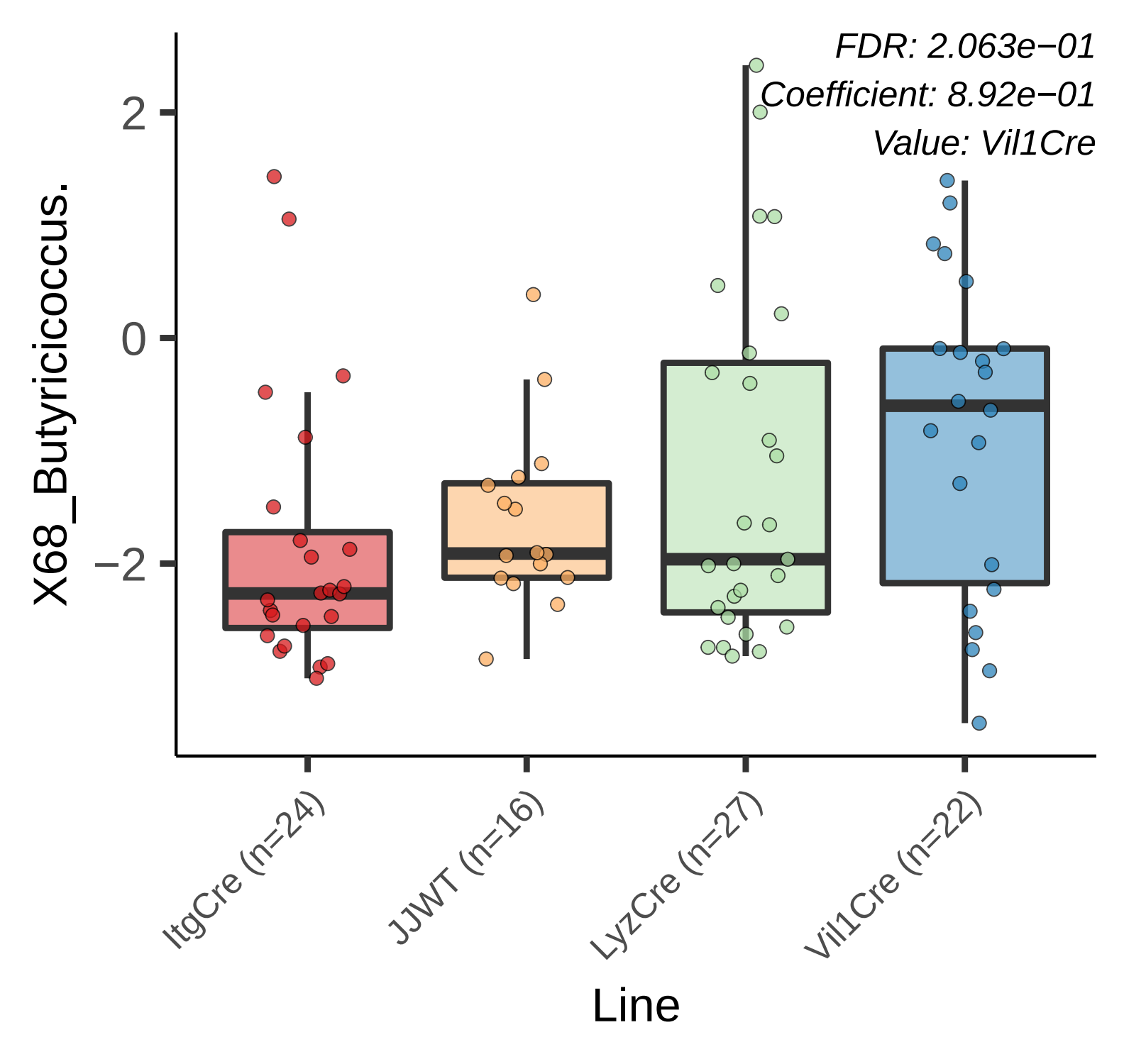

FDR: 2.013e−01
Coefficient: 1.14e+00
Value: Vil1Cre
X49_Lachnospiraceae..f.
4
2
0
−2
ItgCre (n=24)
JJWT (n=16)
LyzCre (n=27)
Vil1Cre (n=22)
Line

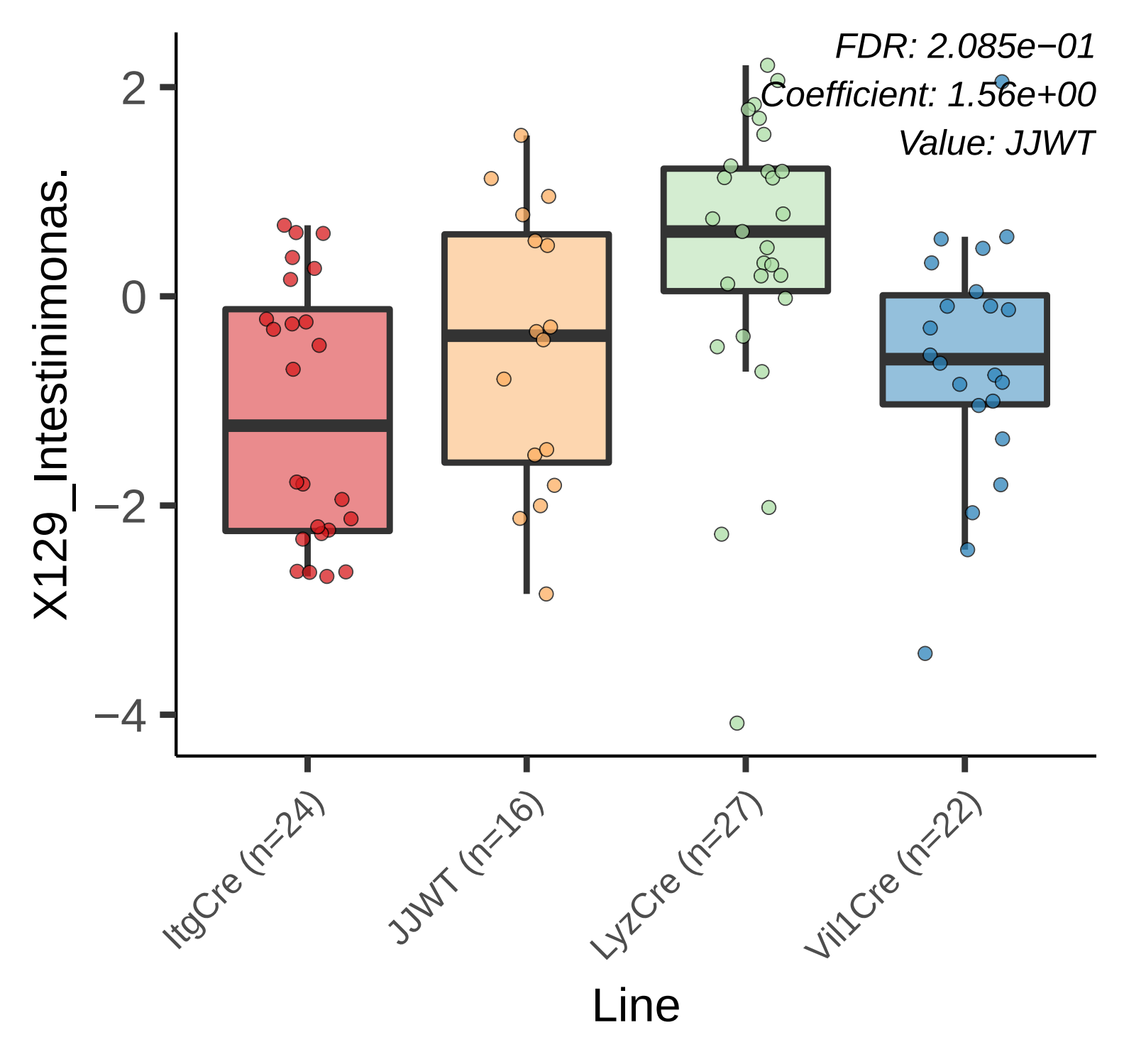

FDR: 2.020e−01
Coefficient: −6.77e−01
Value: Vil1Cre
X16_Muribaculaceae..f.
0
2
4
ItgCre (n=24)
JJWT (n=16)
LyzCre (n=27)
Vil1Cre (n=22)
Line

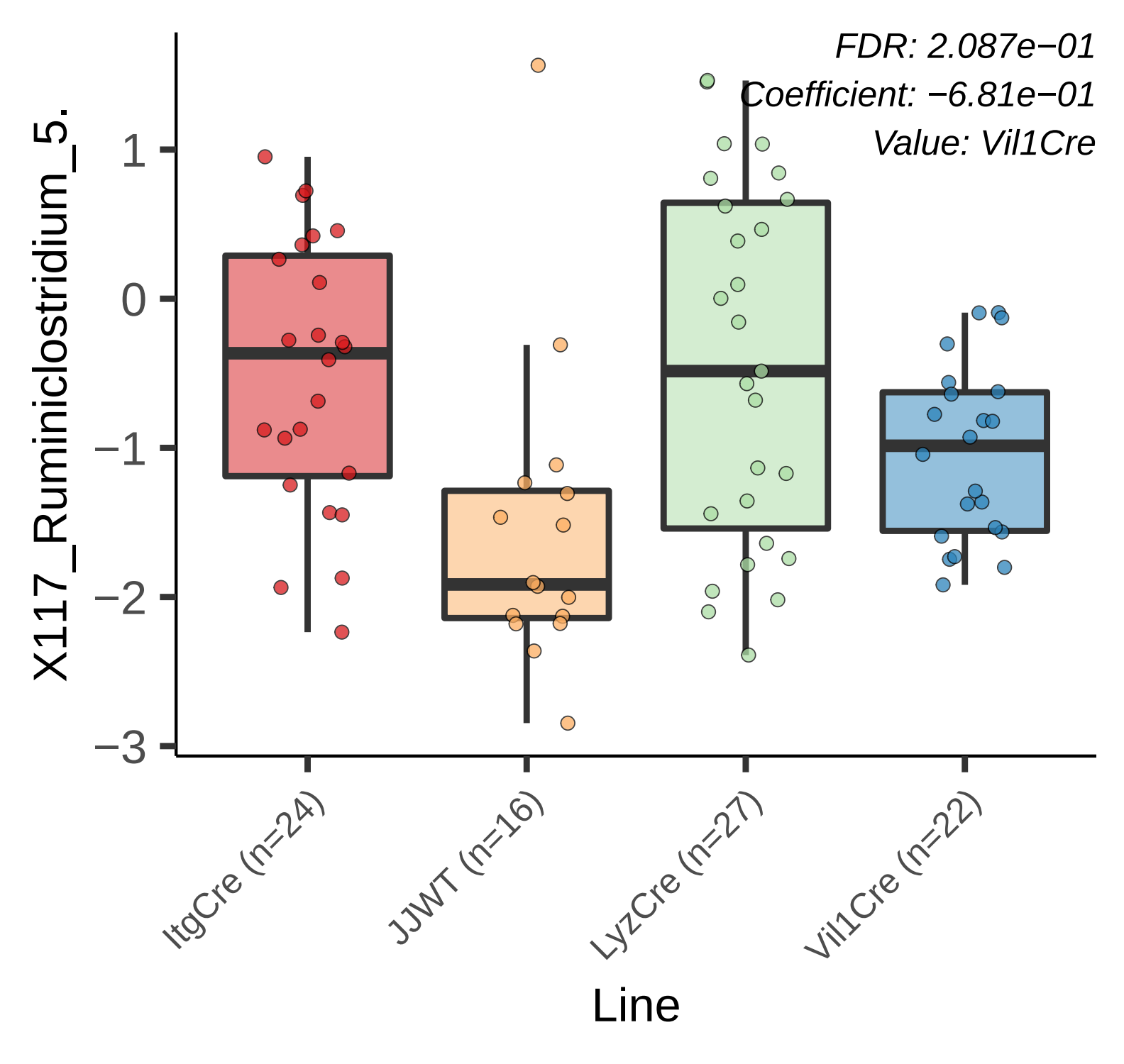

FDR: 2.038e−01
Coefficient: 9.59e−01
Value: LyzCre
X253_Lachnospiraceae_NK4A136_group.
2
1
0
−1
−2
−3
ItgCre (n=24)
JJWT (n=16)
LyzCre (n=27)
Vil1Cre (n=22)
Line

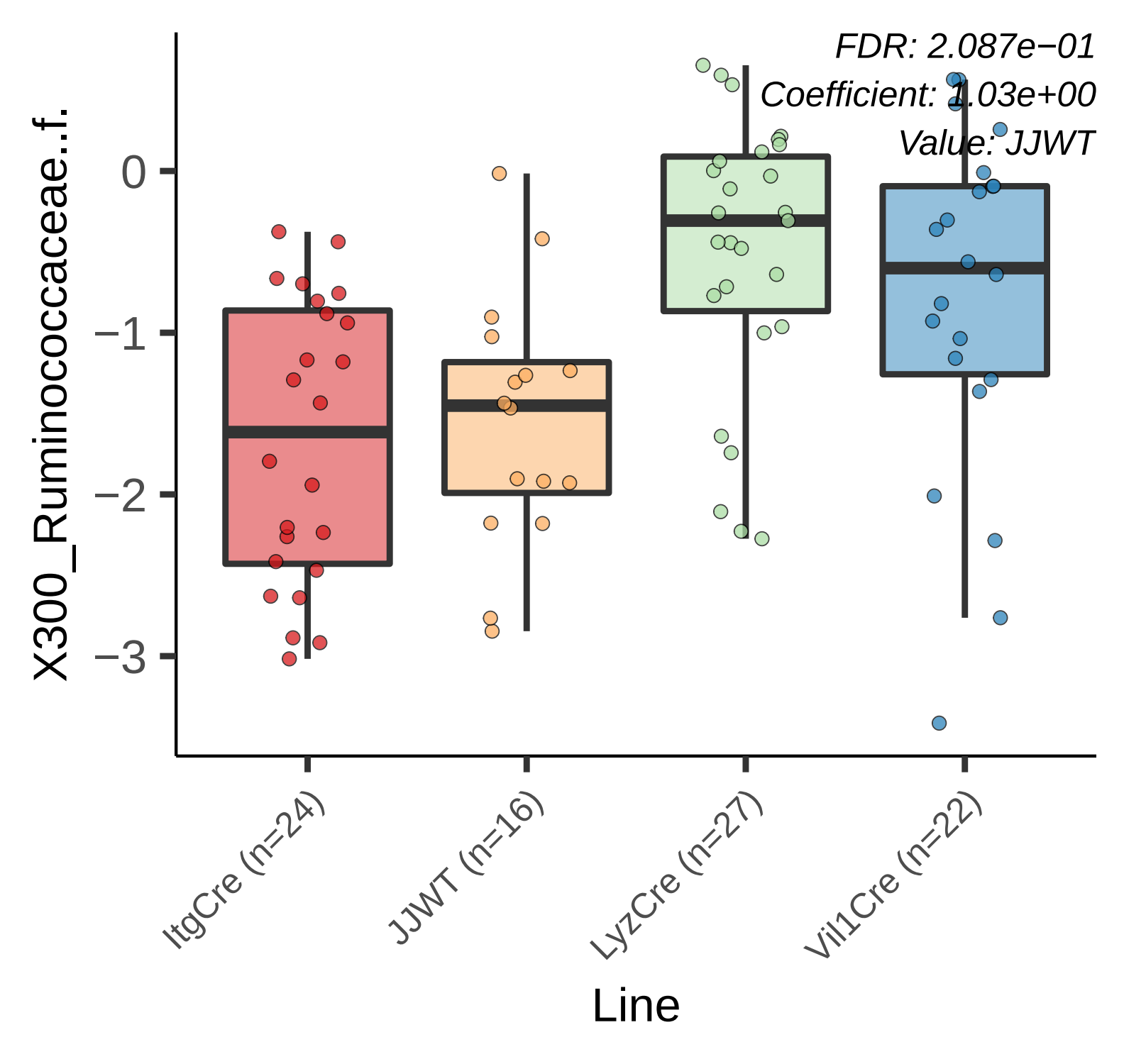

FDR: 2.045e−01
Coefficient: −1.46e+00
Value: Vil1Cre
X18_Lachnoclostridium.
4
2
0
−2
ItgCre (n=24)
JJWT (n=16)
LyzCre (n=27)
Vil1Cre (n=22)
Line

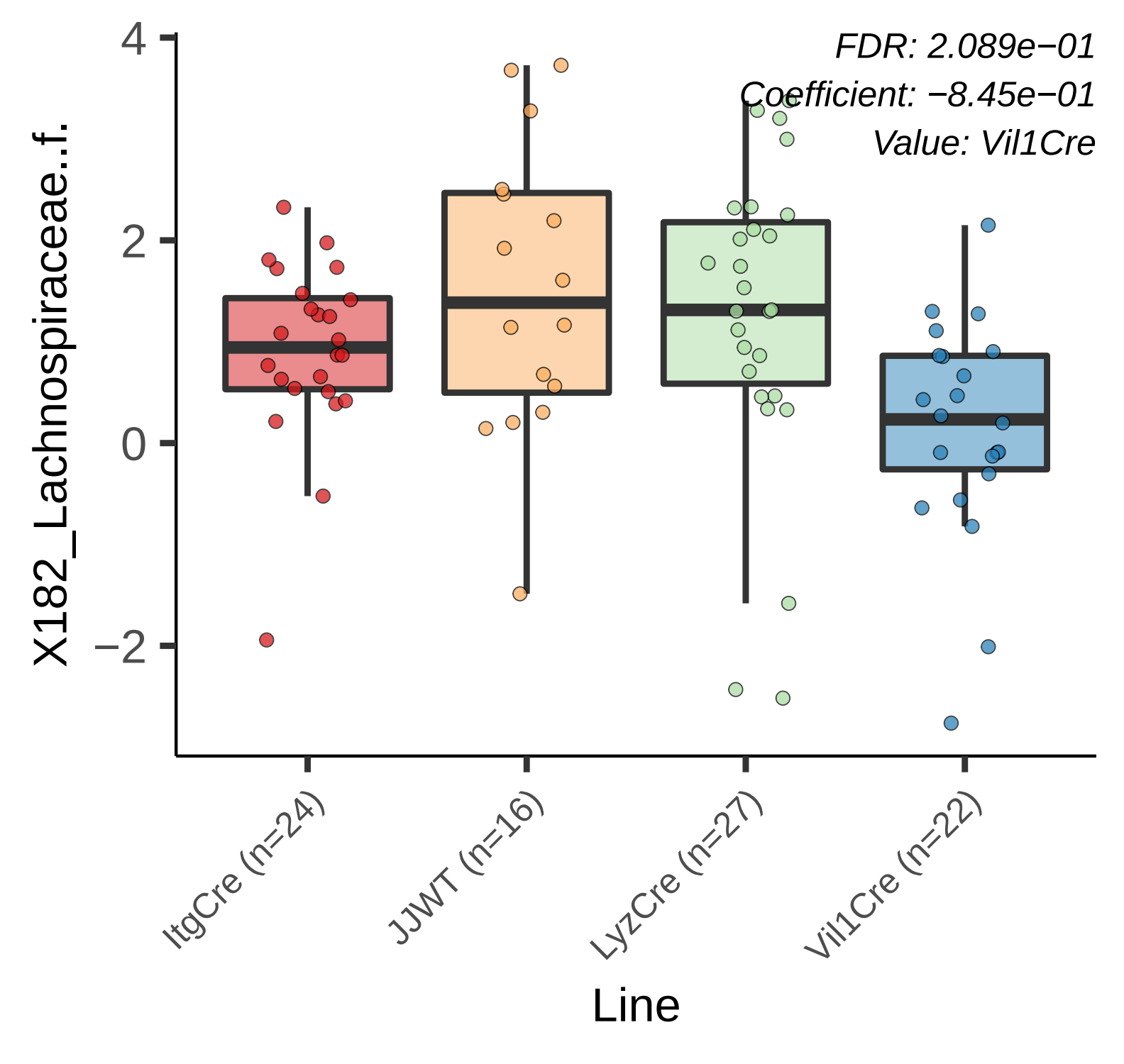

FDR: 2.058e−01
Coefficient: −1.28e+00
Value: LyzCre
X197_Lachnospiraceae.f.
5.0
2.5
0.0
−2.5
ItgCre (n=24)
JJWT (n=16)
LyzCre (n=27)
Vil1Cre (n=22)
Line

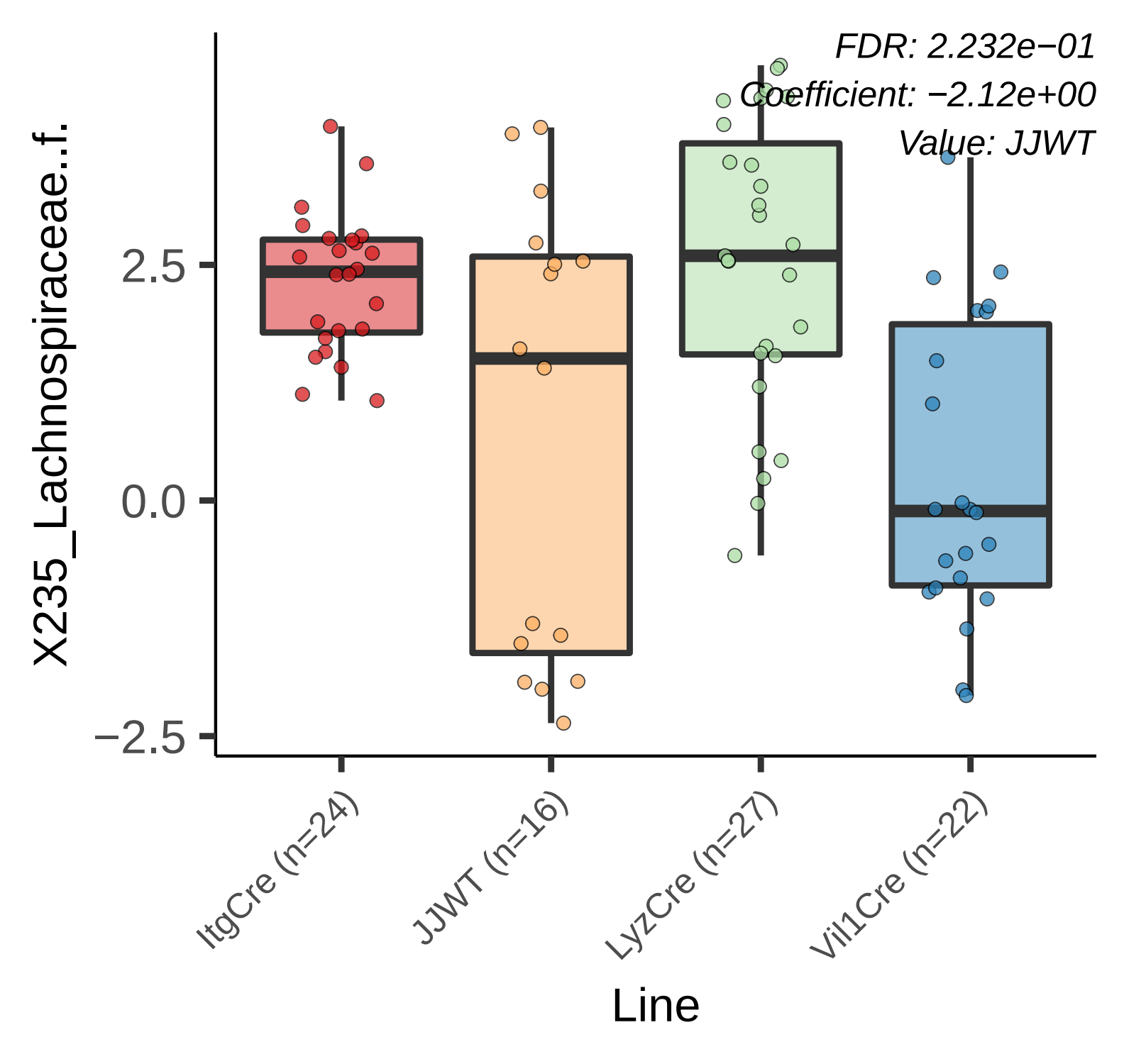

FDR: 2.063e−01
Coefficient: 8.92e−01
Value: Vil1Cre
X68_Butyricicoccus.
2
0
−2
ItgCre (n=24)
JJWT (n=16)
LyzCre (n=27)
Vil1Cre (n=22)
Line

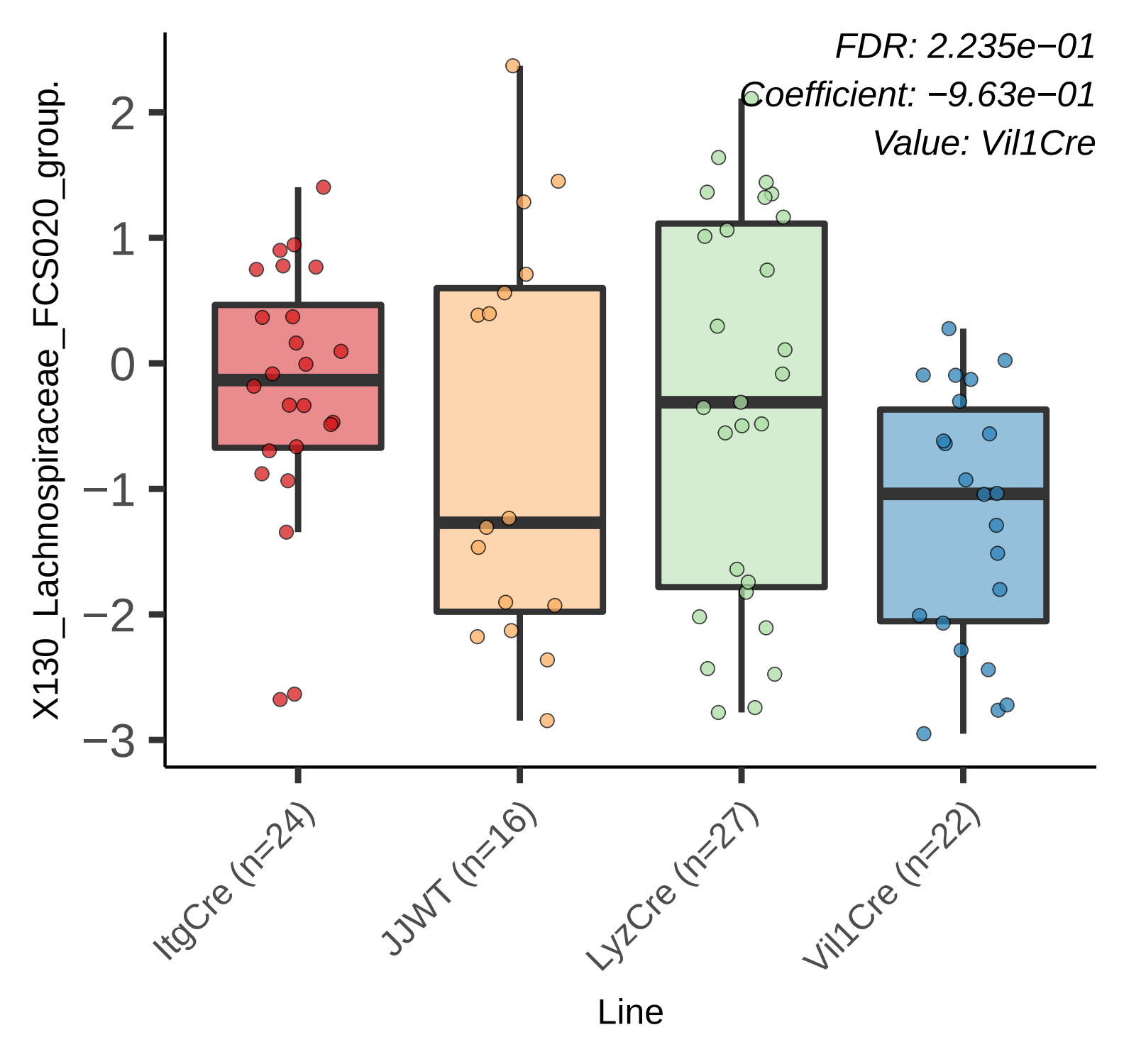

FDR: 2.085e−01
Coefficient: 1.56e+00
Value: JJWT
X129_Intestinimonas.
2
0
−2
−4
ItgCre (n=24)
JJWT (n=16)
LyzCre (n=27)
Vil1Cre (n=22)
Line

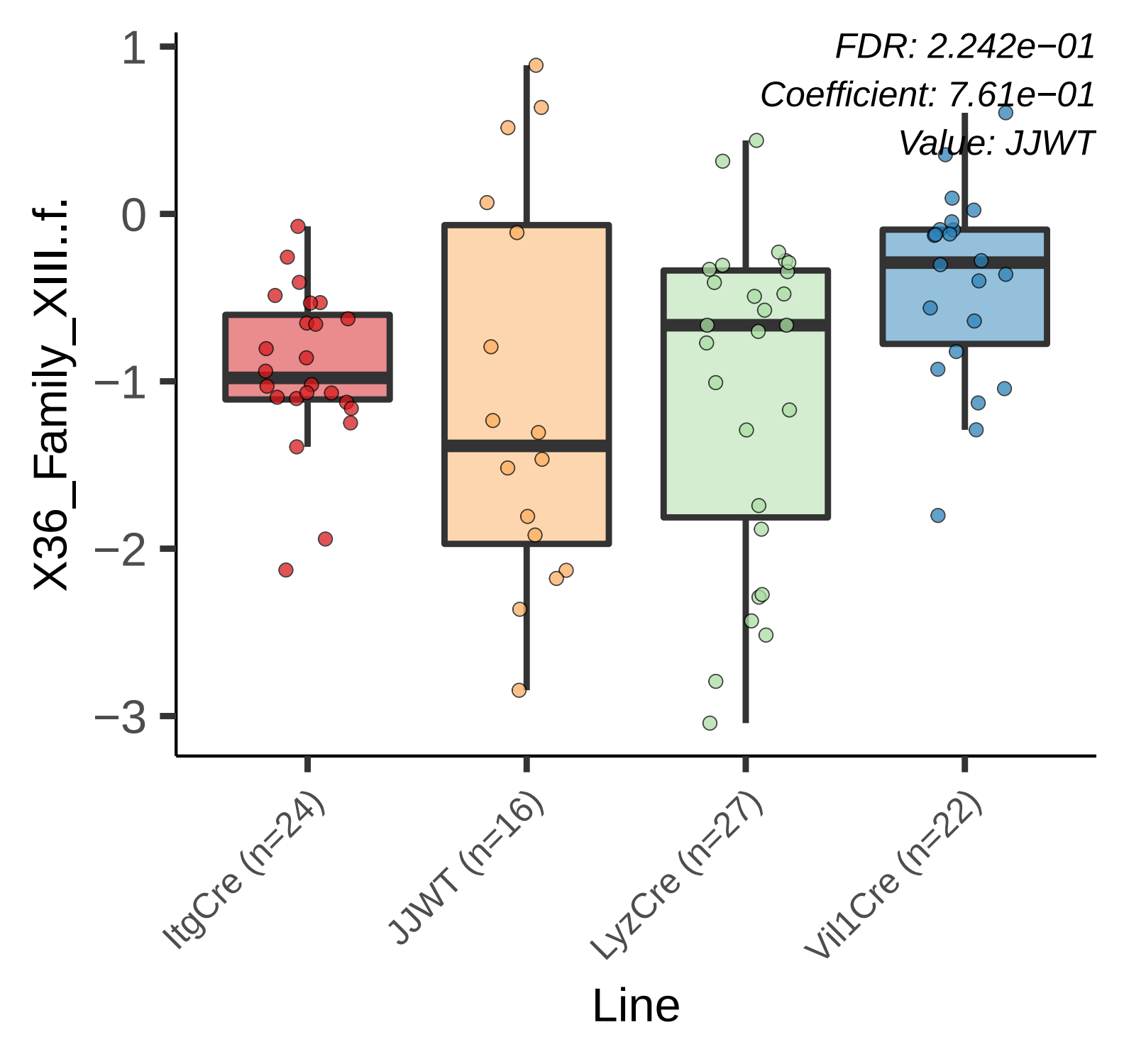

FDR: 2.087e−01
Coefficient: −6.81e−01
Value: Vil1Cre
X117_Ruminiclostridium_5.
1
0
−1
−2
−3
ItgCre (n=24)
JJWT (n=16)
LyzCre (n=27)
Vil1Cre (n=22)
Line

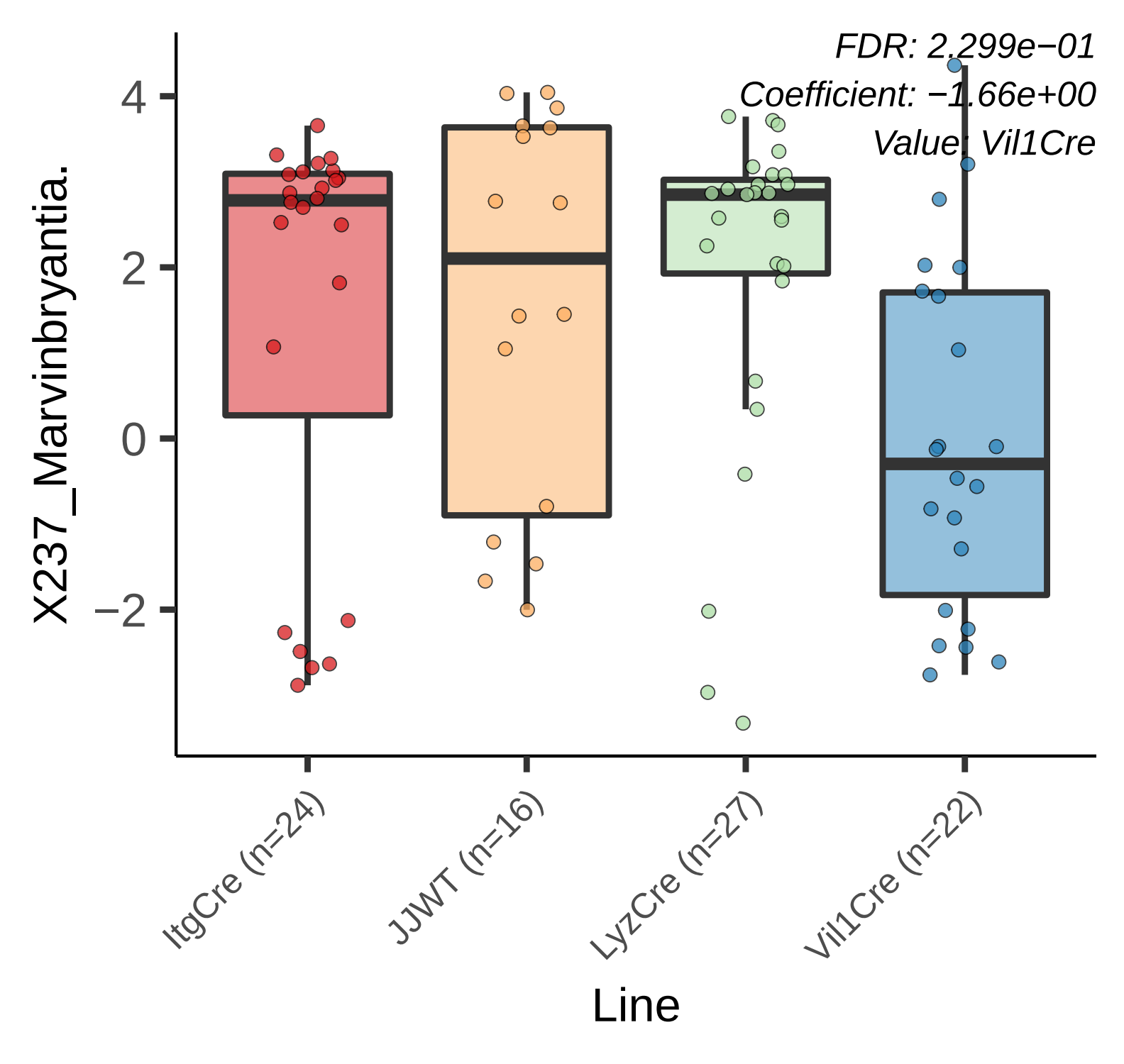

FDR: 2.087e−01
Coefficient: 1.03e+00
Value: JJWT
X300_Ruminococcaceae..f.
0
−1
−2
−3
ItgCre (n=24)
JJWT (n=16)
LyzCre (n=27)
Vil1Cre (n=22)
Line

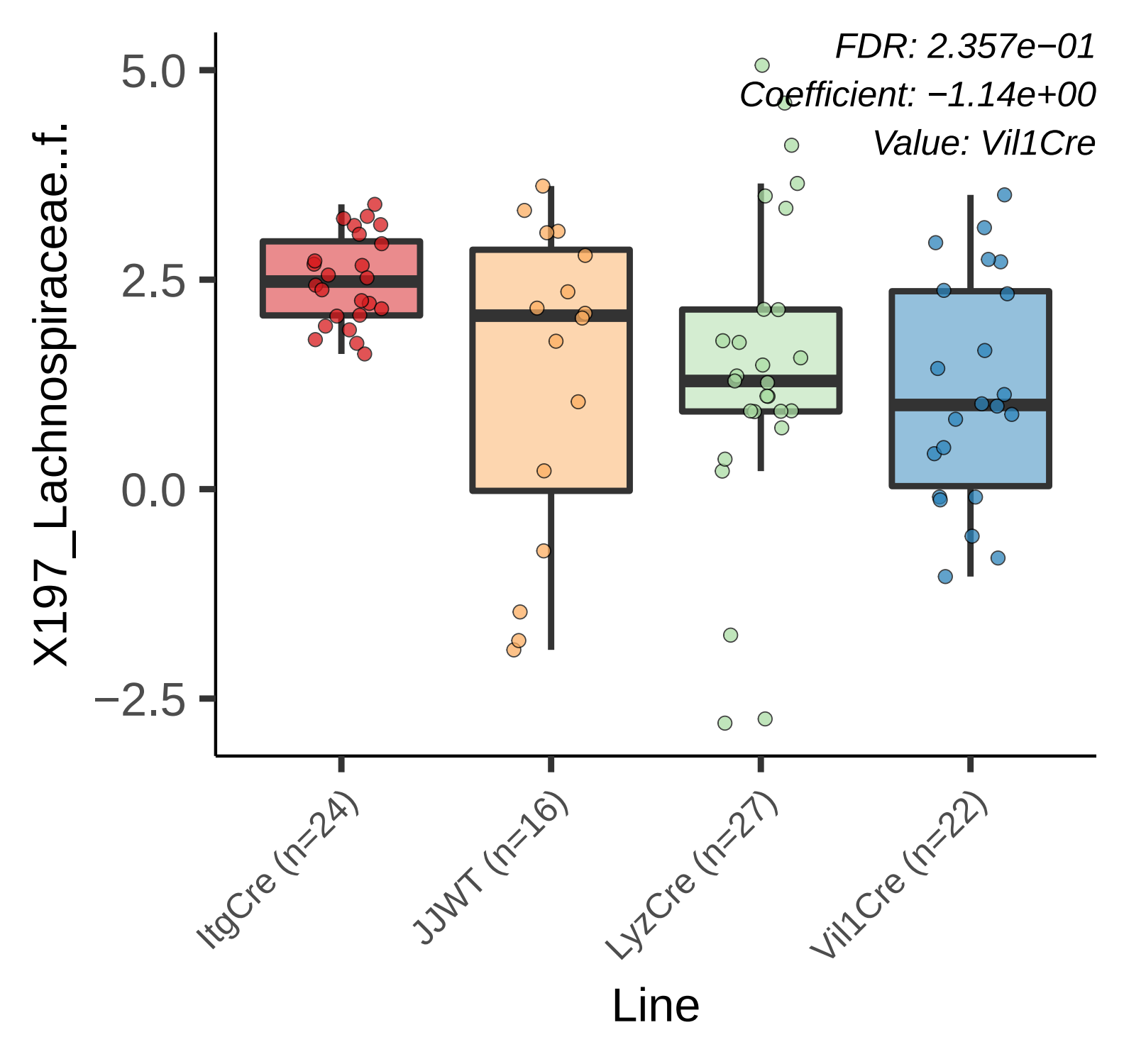

FDR: 2.089e−01
Coefficient: −8.45e−01
Value: Vil1Cre
X182_Lachnospiraceae.f.
4
2
0
−2
ItgCre (n=24)
JJWT (n=16)
LyzCre (n=27)
Vil1Cre (n=22)
Line

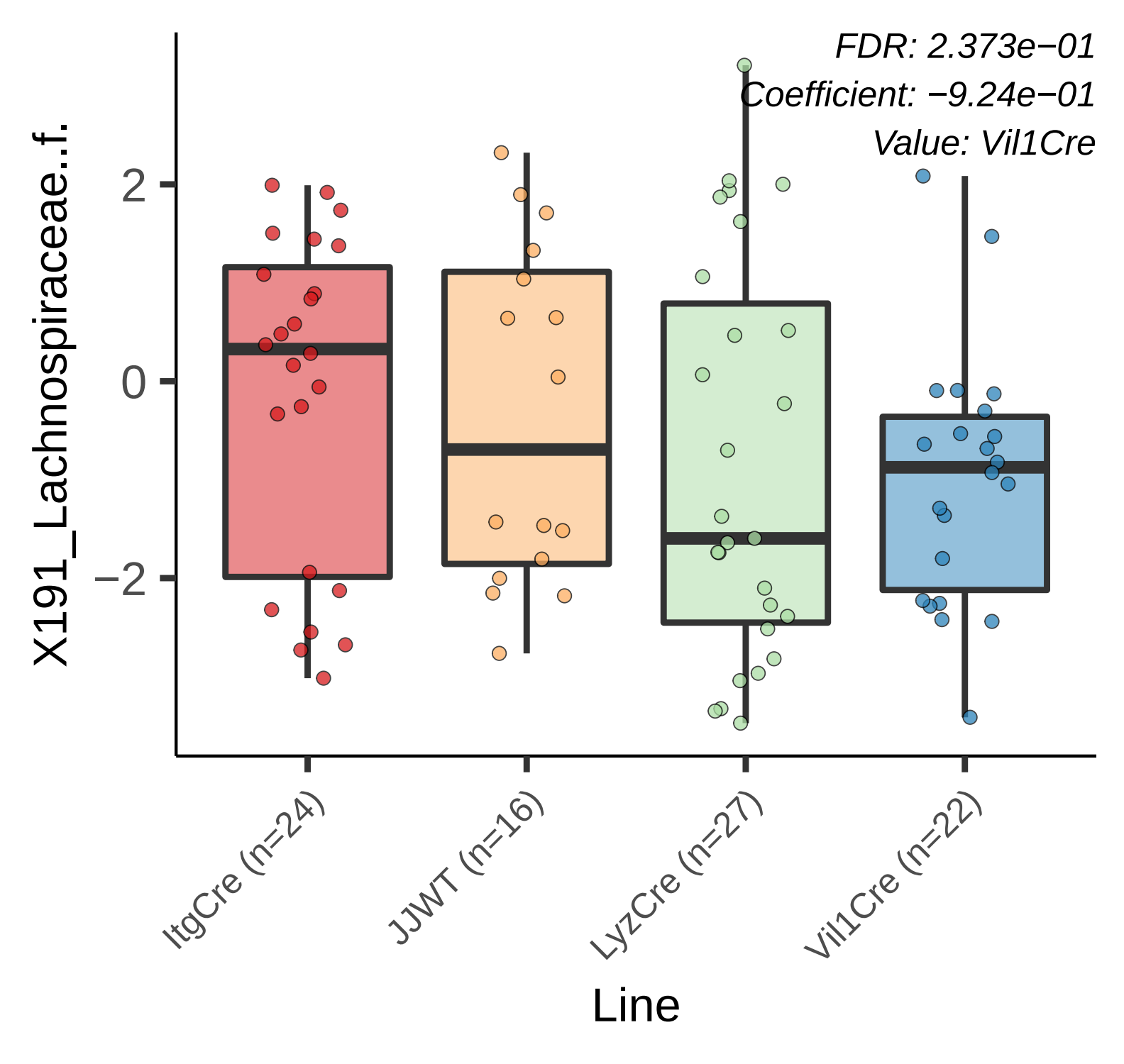

FDR: 2.232e−01
Coefficient: −2.12e+00
Value: JJWT
X235_Lachnospiraceae.f.
2.5
0.0
−2.5
ItgCre (n=24)
JJWT (n=16)
LyzCre (n=27)
Vil1Cre (n=22)
Line

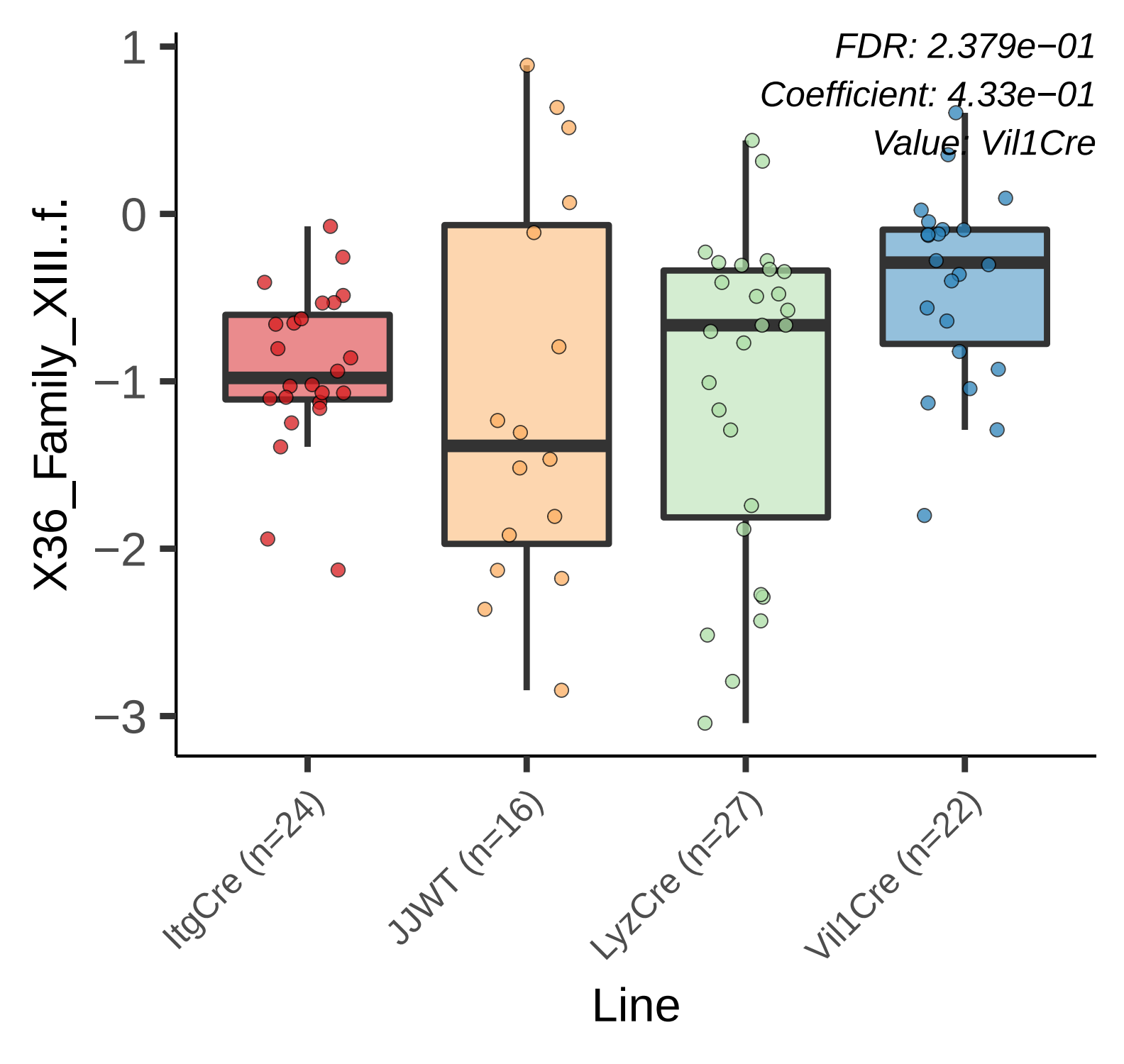

FDR: 2.235e−01
Coefficient: −9.63e−01
Value: Vil1Cre
X130_Lachnospiraceae_FCS020_group.
2
1
0
−1
−2
−3
ItgCre (n=24)
JJWT (n=16)
LyzCre (n=27)
Vil1Cre (n=22)
Line

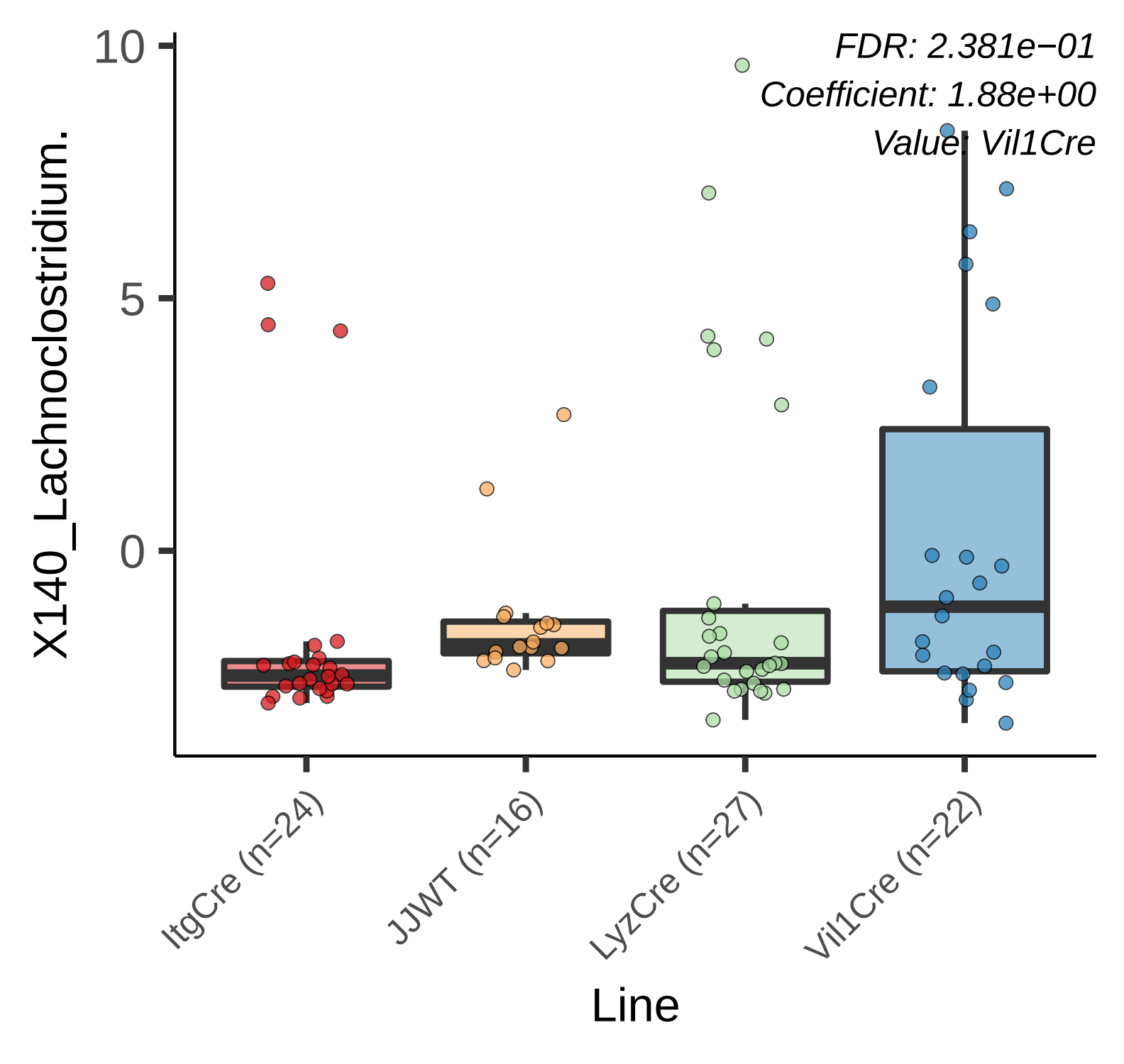

FDR: 2.242e−01
Coefficient: 7.61e−01
Value: JJWT
X36_Family_XIII..f.
1
0
−1
−2
−3
ItgCre (n=24)
JJWT (n=16)
LyzCre (n=27)
Vil1Cre (n=22)
Line

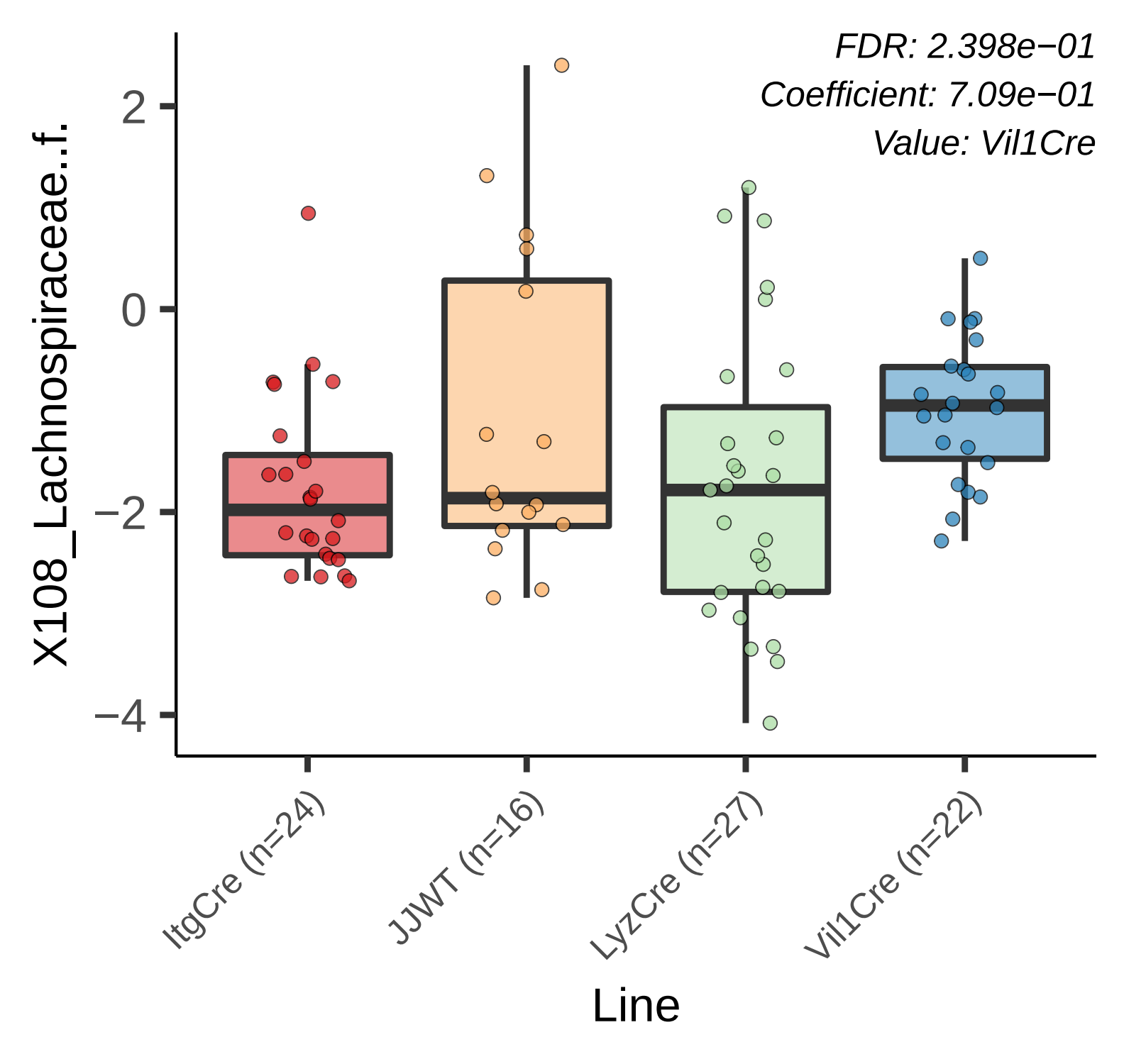

FDR: 2.299e−01
Coefficient: −1.66e+00
Value: Vil1Cre
X237_Marvinbryantia.
4
2
0
−2
ItgCre (n=24)
JJWT (n=16)
LyzCre (n=27)
Vil1Cre (n=22)
Line

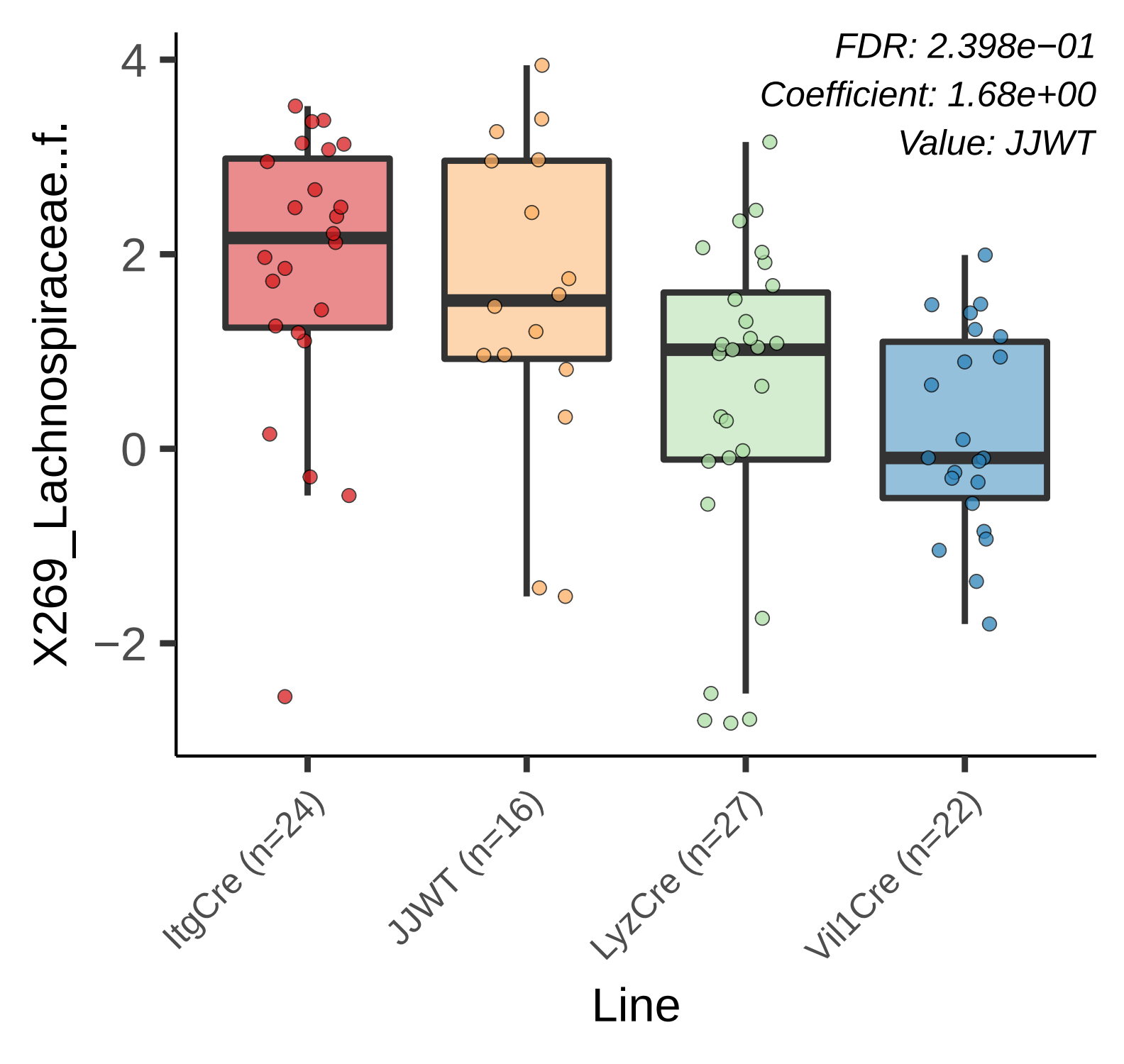

FDR: 2.357e−01
Coefficient: −1.14e+00
Value: Vil1Cre
X197_Lachnospiraceae.f.
5.0
2.5
0.0
−2.5
ItgCre (n=24)
JJWT (n=16)
LyzCre (n=27)
Vil1Cre (n=22)
Line

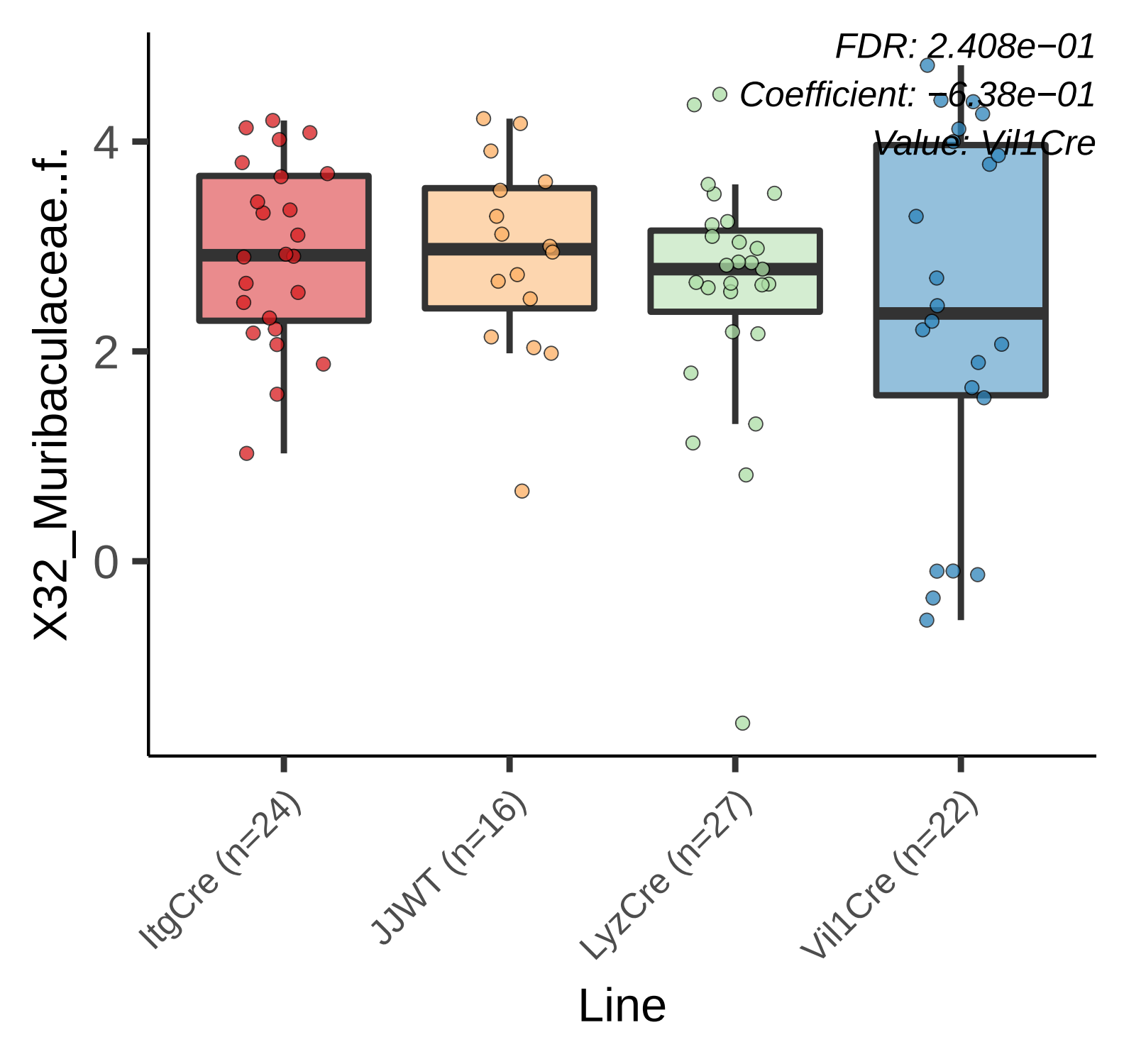

FDR: 2.373e−01
Coefficient: −9.24e−01
Value: Vil1Cre
X191_Lachnospiraceae.f.
2
0
−2
ItgCre (n=24)
JJWT (n=16)
LyzCre (n=27)
Vil1Cre (n=22)
Line

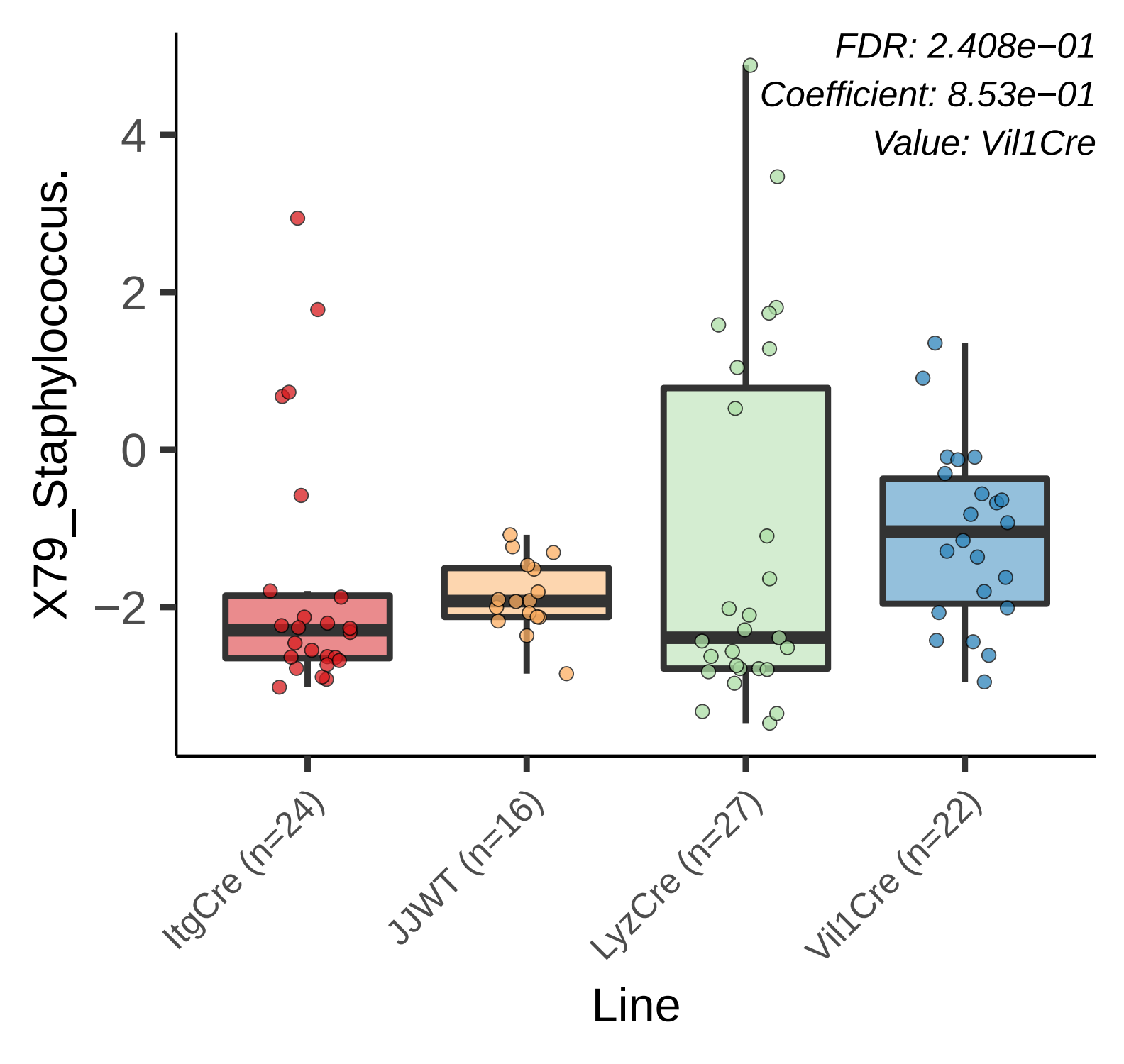

FDR: 2.379e−01
Coefficient: 4.33e−01
Value: Vil1Cre
X36_Family_XIII..f.
1
0
−1
−2
−3
ItgCre (n=24)
JJWT (n=16)
LyzCre (n=27)
Vil1Cre (n=22)
Line

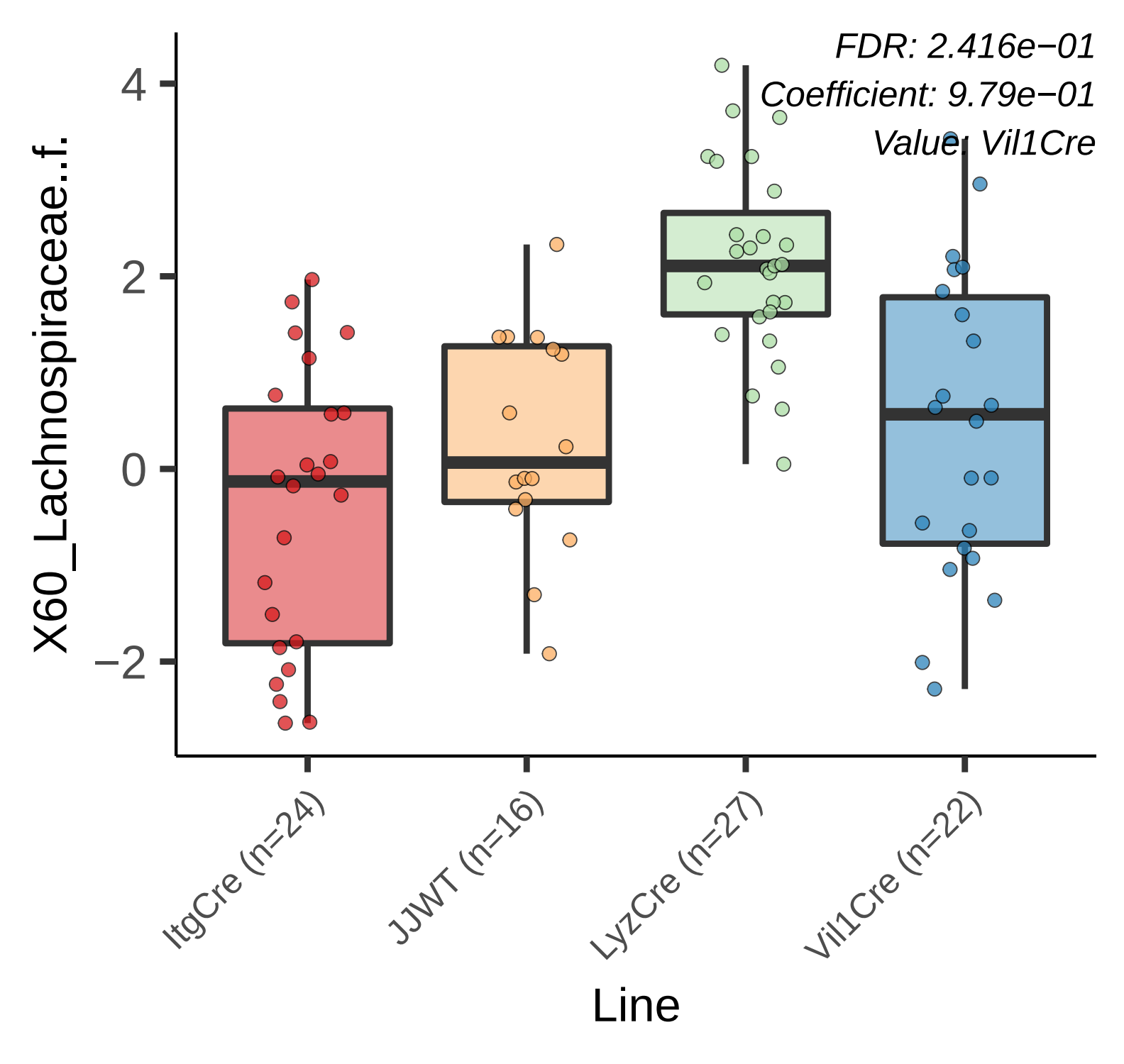

FDR: 2.381e−01
Coefficient: 1.88e+00
Value: Vil1Cre
X140_Lachnoclostridium.
10
5
0
ItgCre (n=24)
JJWT (n=16)
LyzCre (n=27)
Vil1Cre (n=22)
Line

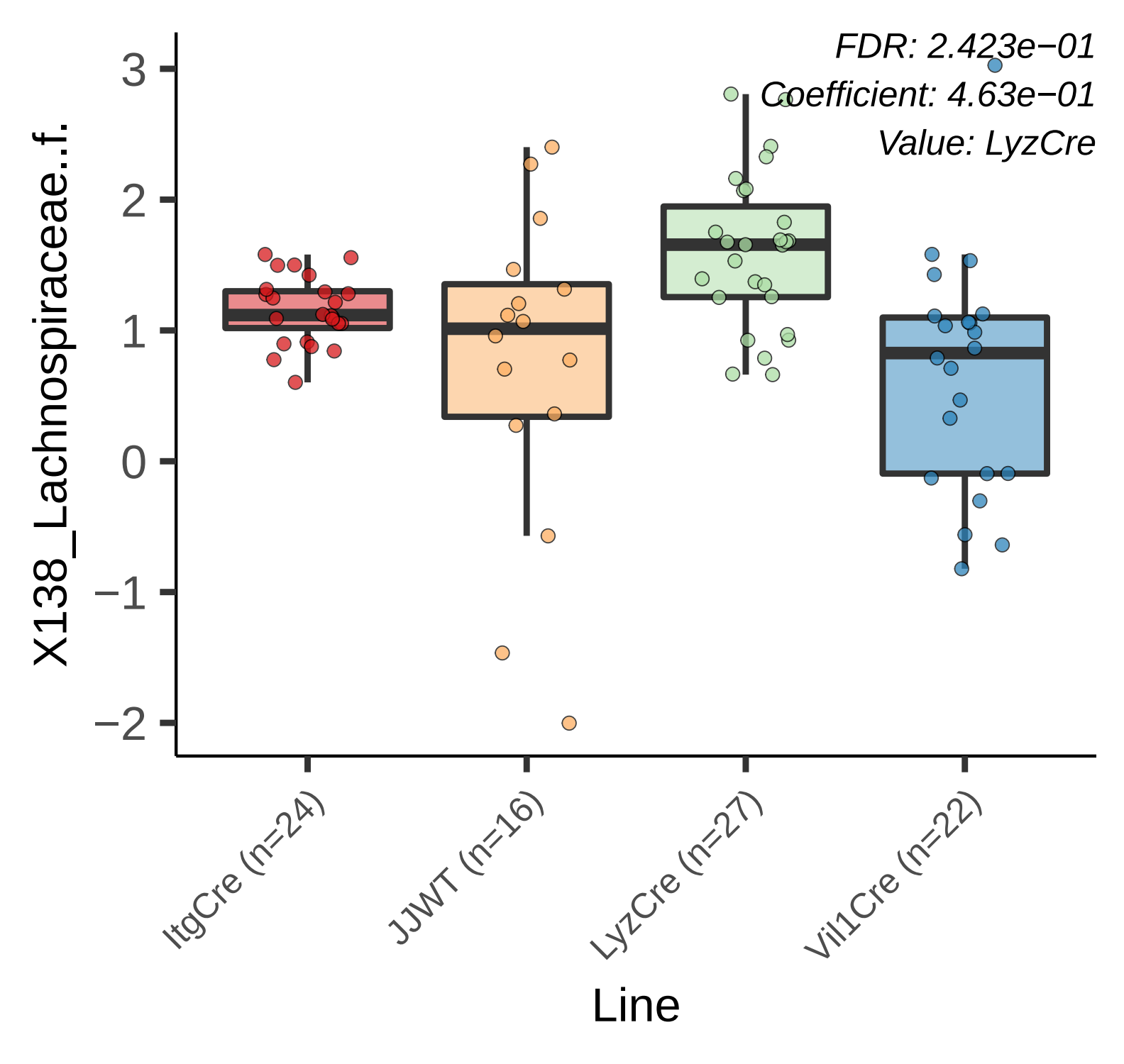

FDR: 2.398e−01
Coefficient: 7.09e−01
Value: Vil1Cre
X108_Lachnospiraceae.f.
2
0
−2
−4
ItgCre (n=24)
JJWT (n=16)
LyzCre (n=27)
Vil1Cre (n=22)
Line

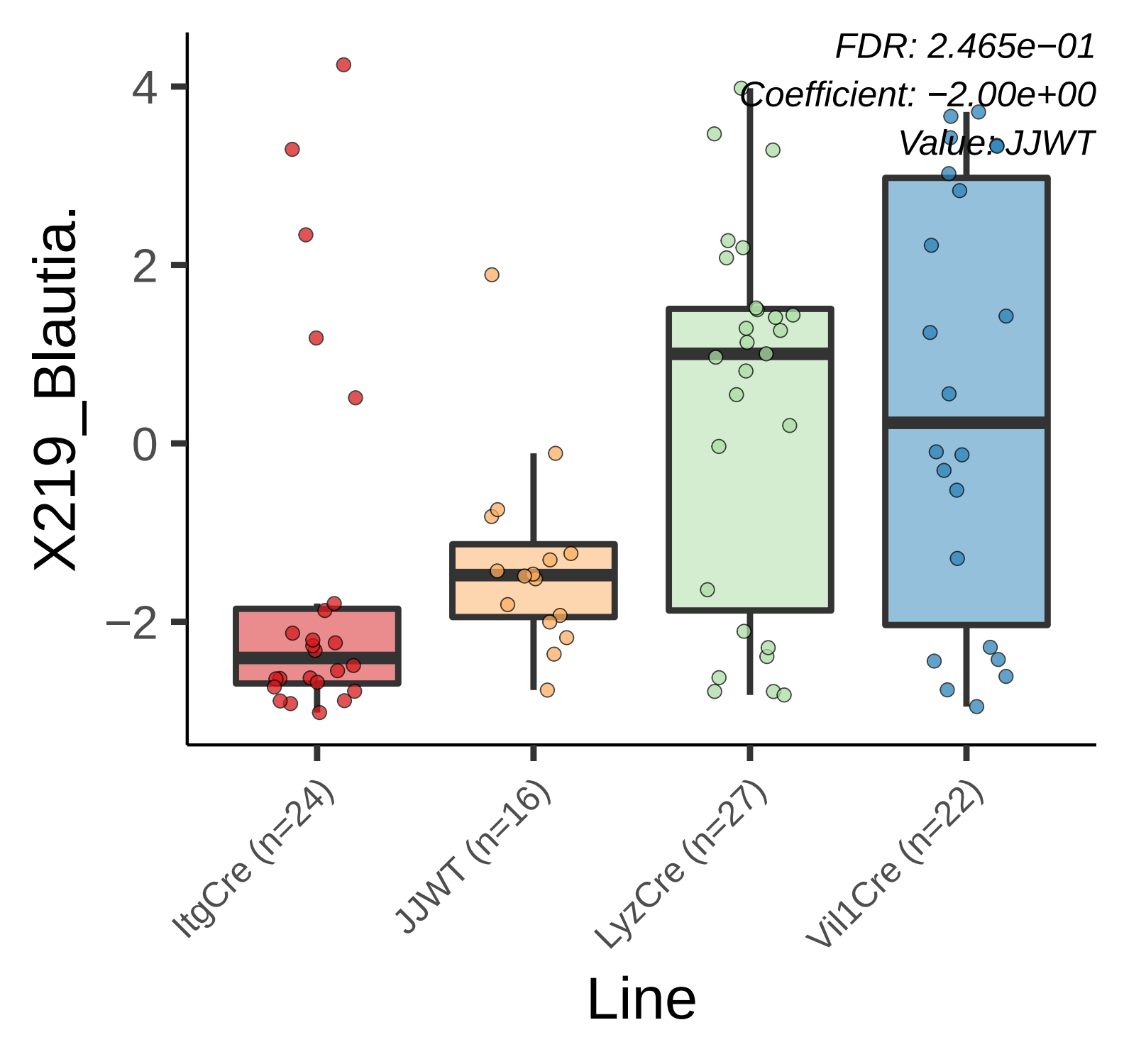

FDR: 2.398e−01
Coefficient: 1.68e+00
Value: JJWT
X269_Lachnospiraceae.f.
4
2
0
−2
ItgCre (n=24)
JJWT (n=16)
LyzCre (n=27)
Vil1Cre (n=22)
Line

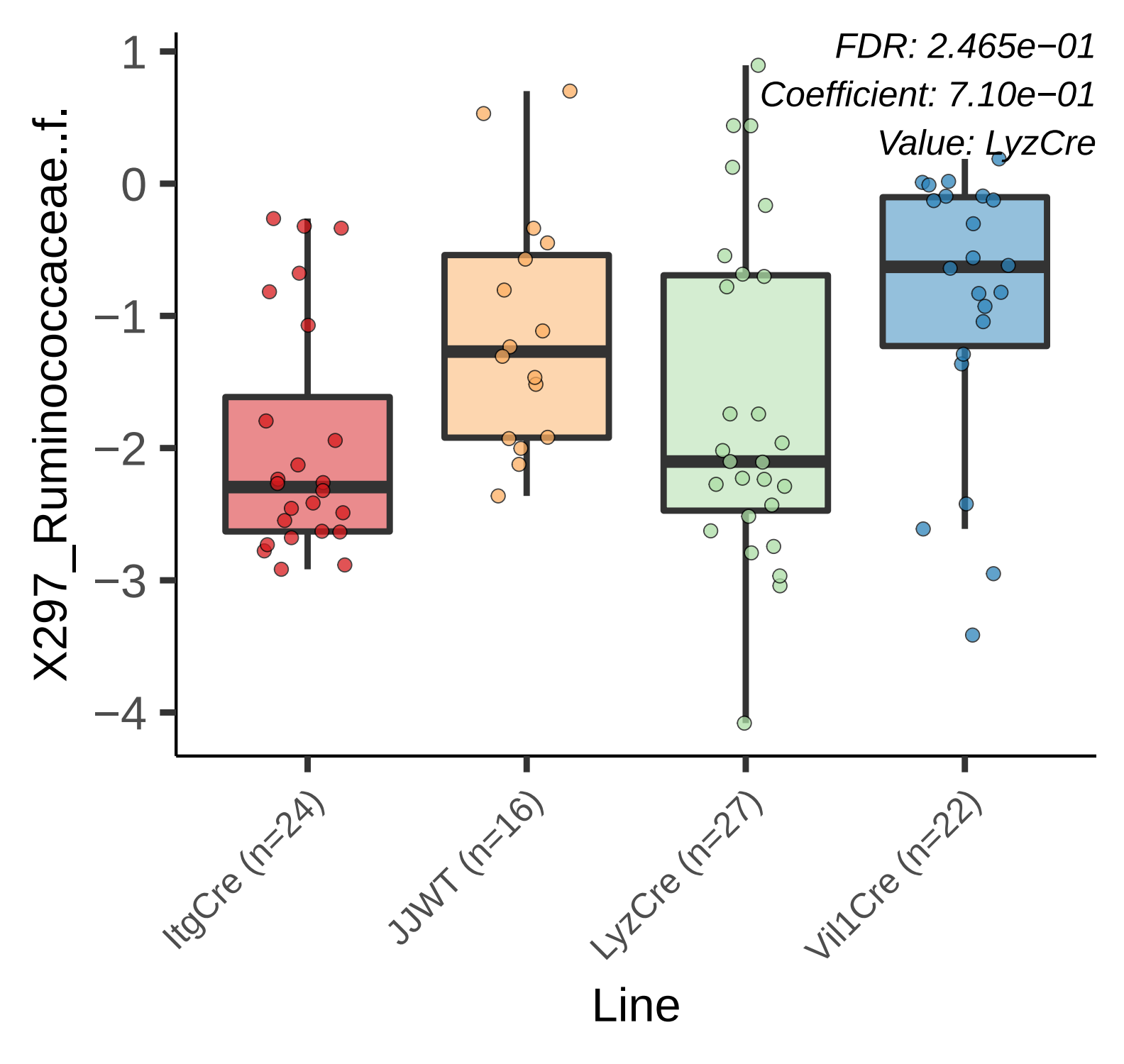

FDR: 2.408e−01
Coefficient: −6.38e−01
Value: Vil1Cre
X32_Muribaculaceae..f.
4
2
0
ItgCre (n=24)
JJWT (n=16)
LyzCre (n=27)
Vil1Cre (n=22)
Line

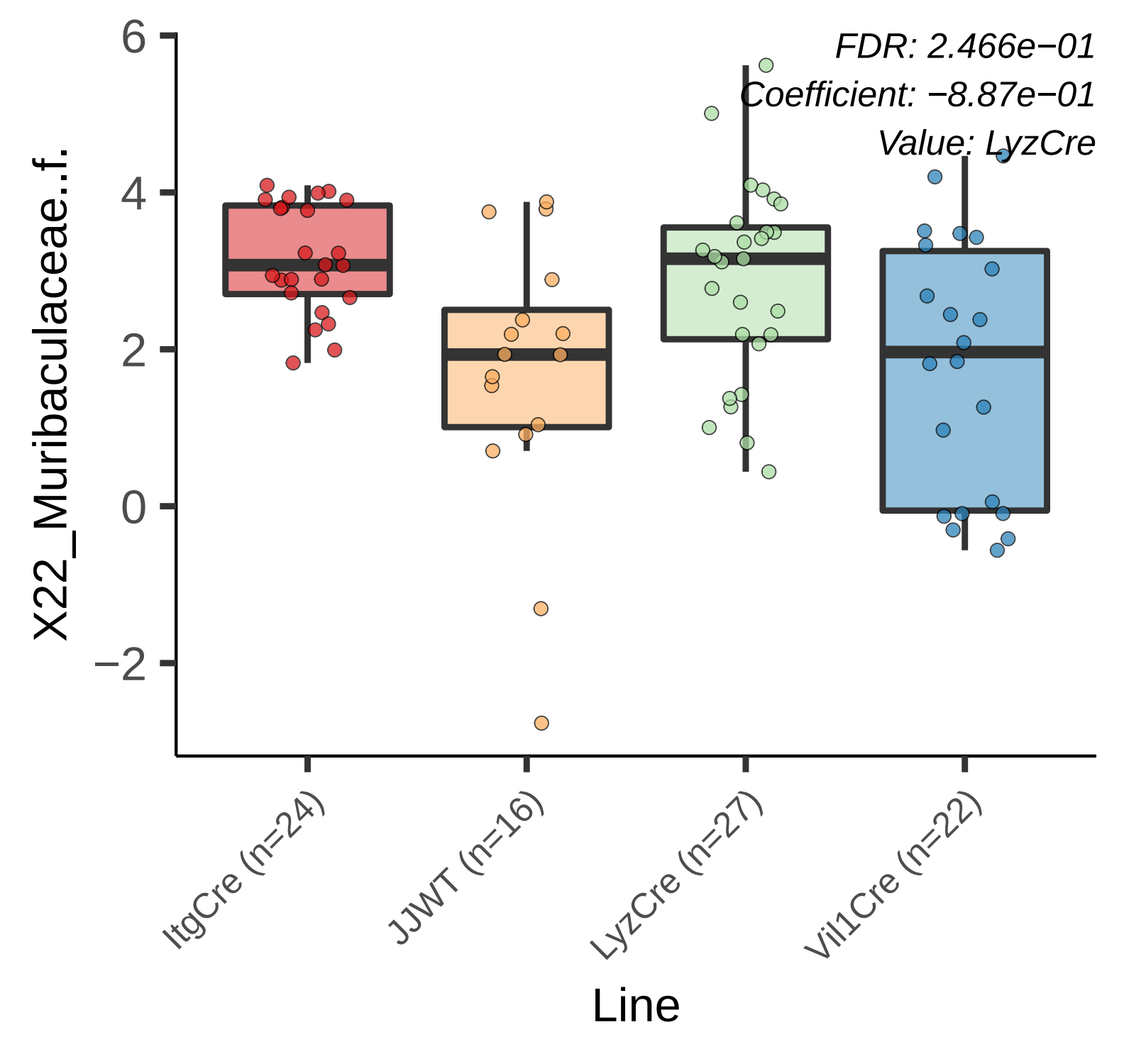

FDR: 2.408e−01
Coefficient: 8.53e−01
Value: Vil1Cre
X79_Staphylococcus.
4
2
0
−2
ItgCre (n=24)
JJWT (n=16)
LyzCre (n=27)
Vil1Cre (n=22)
Line

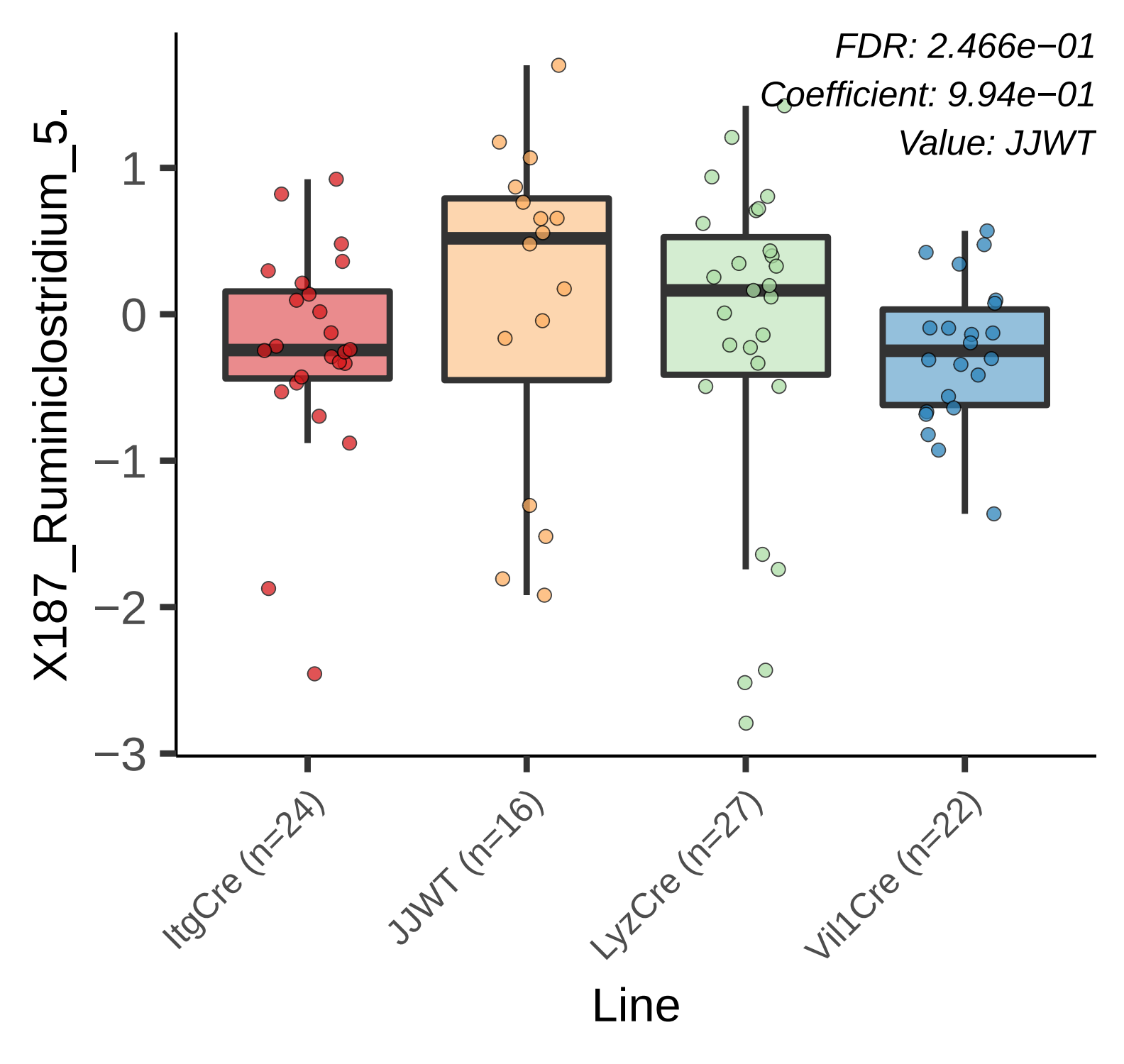

FDR: 2.416e−01
Coefficient: 9.79e−01
Value: Vil1Cre
X60_Lachnospiraceae..f.
4
2
0
−2
ItgCre (n=24)
JJWT (n=16)
LyzCre (n=27)
Vil1Cre (n=22)
Line

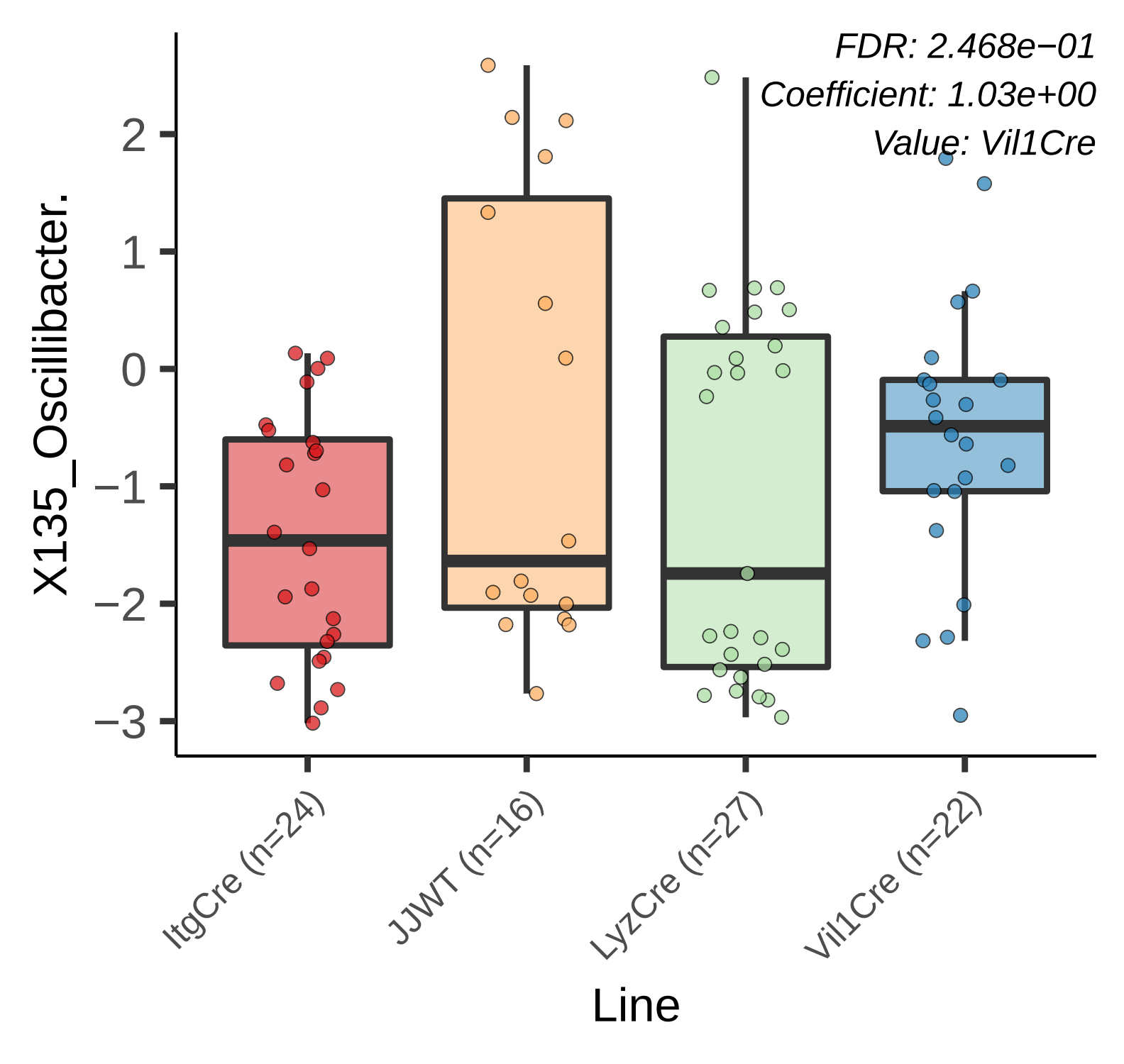

FDR: 2.423e−01
Coefficient: 4.63e−01
Value: LyzCre
X138_Lachnospiraceae.f.
3
2
1
0
−1
−2
ItgCre (n=24)
JJWT (n=16)
LyzCre (n=27)
Vil1Cre (n=22)
Line

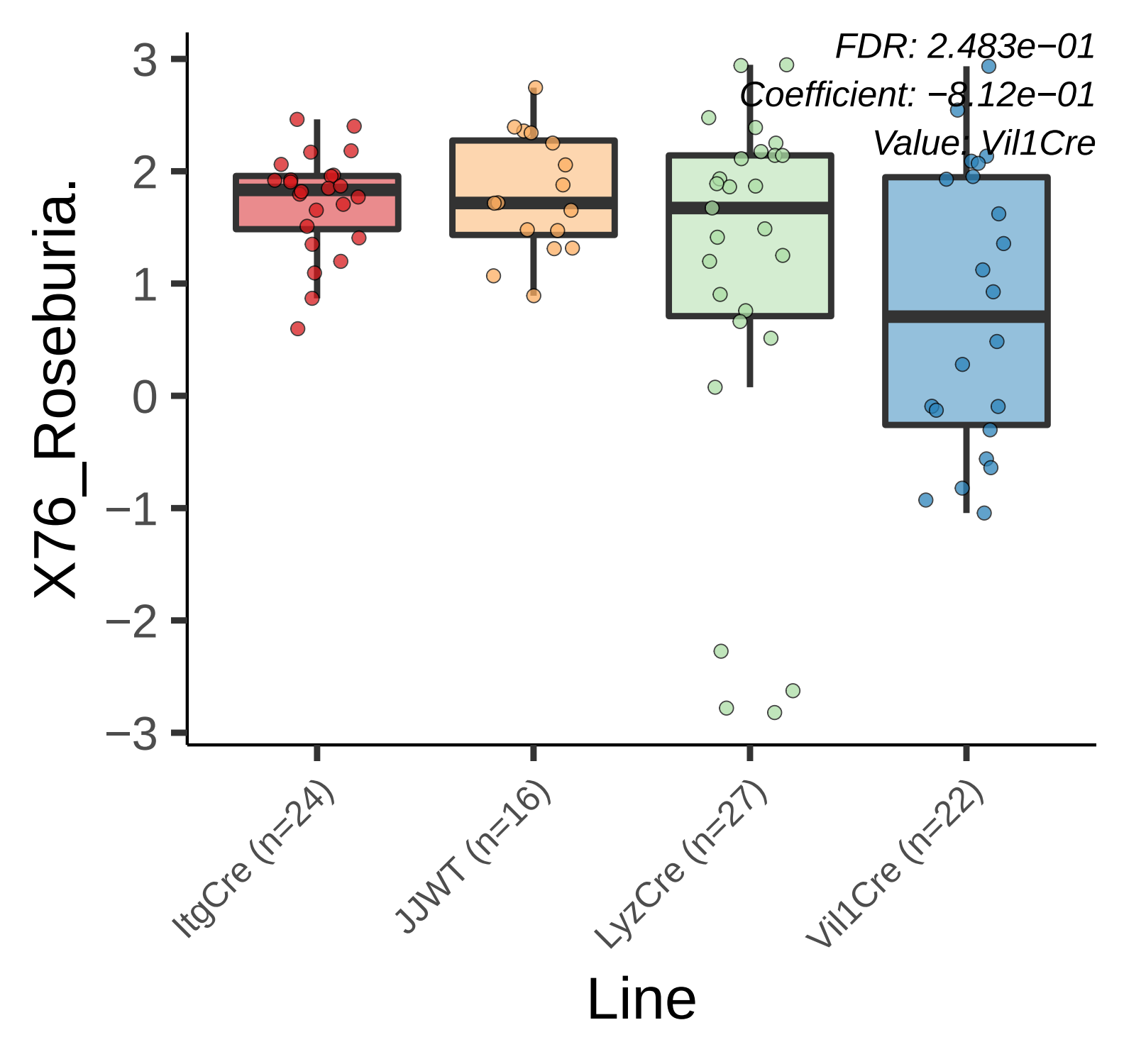

FDR: 2.465e−01
Coefficient: −2.00e+00
Value: JJWT
X219_Blautia.
4
2
0
−2
ItgCre (n=24)
JJWT (n=16)
LyzCre (n=27)
Vil1Cre (n=22)
Line


FDR: 2.465e−01
Coefficient: 7.10e−01
Value: LyzCre
X297_Ruminococcaceae..f.
1
0
−1
−2
−3
−4
ItgCre (n=24)
JJWT (n=16)
LyzCre (n=27)
Vil1Cre (n=22)
Line


FDR: 2.466e−01
Coefficient: −8.87e−01
Value: LyzCre
X22_Muribaculaceae..f.
6
4
2
0
−2
ItgCre (n=24)
JJWT (n=16)
LyzCre (n=27)
Vil1Cre (n=22)
Line


FDR: 2.466e−01
Coefficient: 9.94e−01
Value: JJWT
X187_Ruminiclostridium_5.
1
0
−1
−2
−3
ItgCre (n=24)
JJWT (n=16)
LyzCre (n=27)
Vil1Cre (n=22)
Line


FDR: 2.468e−01
Coefficient: 1.03e+00
Value: Vil1Cre
X135_Oscillibacter.
2
1
0
−1
−2
−3
ItgCre (n=24)
JJWT (n=16)
LyzCre (n=27)
Vil1Cre (n=22)
Line


FDR: 2.483e−01
Coefficient: −8.12e−01
Value: Vil1Cre
X76_Roseburia.
3
2
1
0
−1
−2
−3
ItgCre (n=24)
JJWT (n=16)
LyzCre (n=27)
Vil1Cre (n=22)
Line