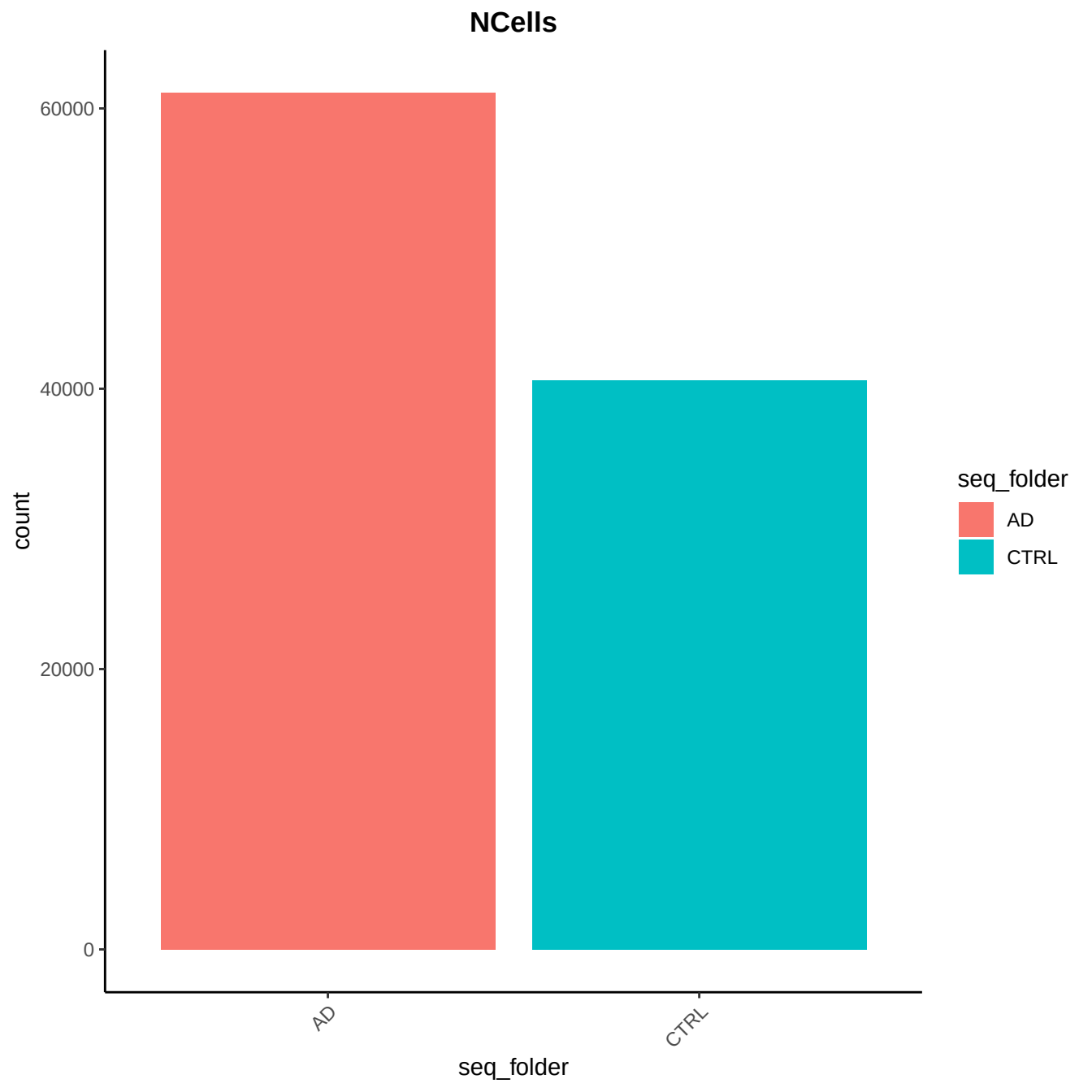
NCells
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40000
20000
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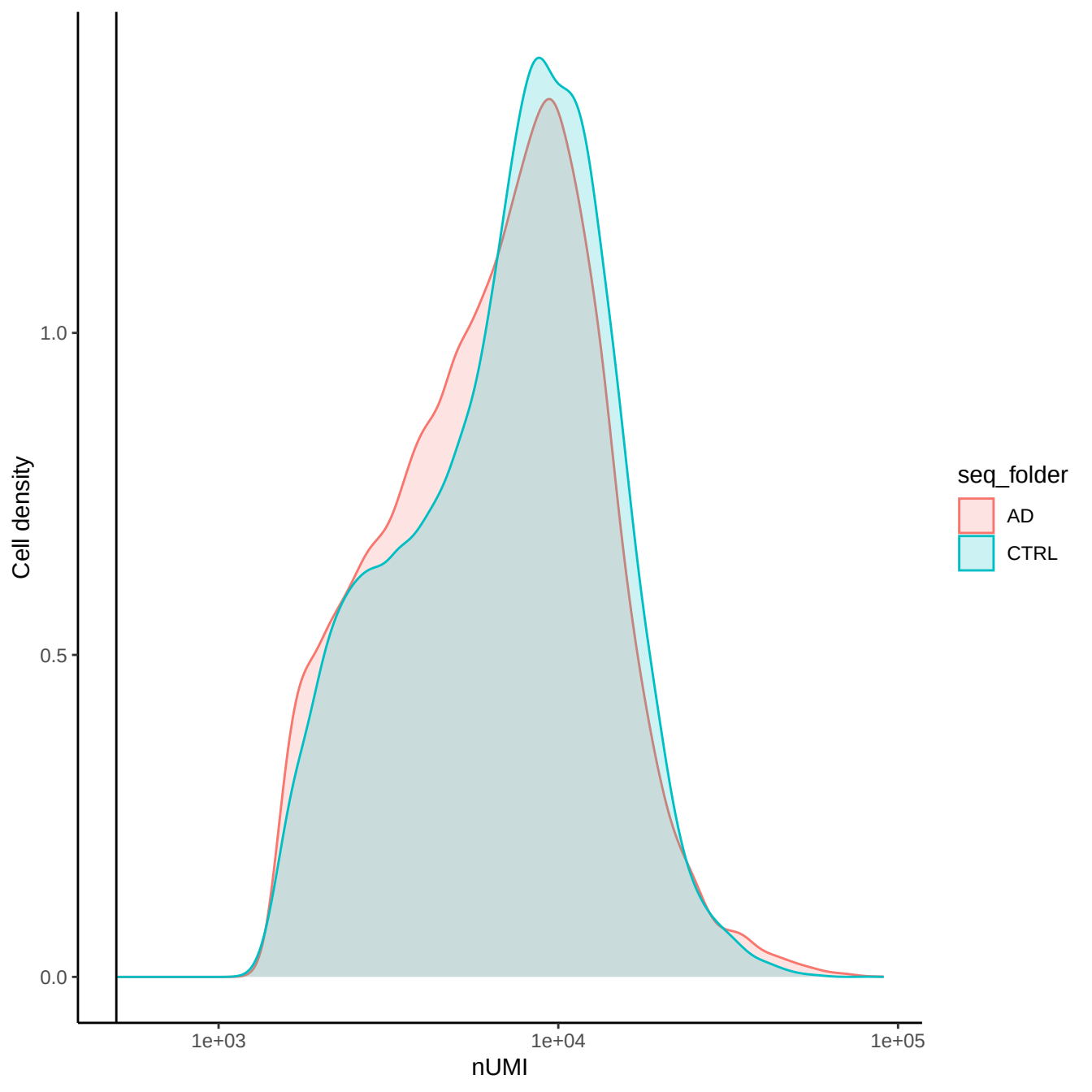
count
AD
CTRL
seq_folder
seq_folder
AD
CTRL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0.0
1e+03
1e+04
1e+05
nUMI
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AD
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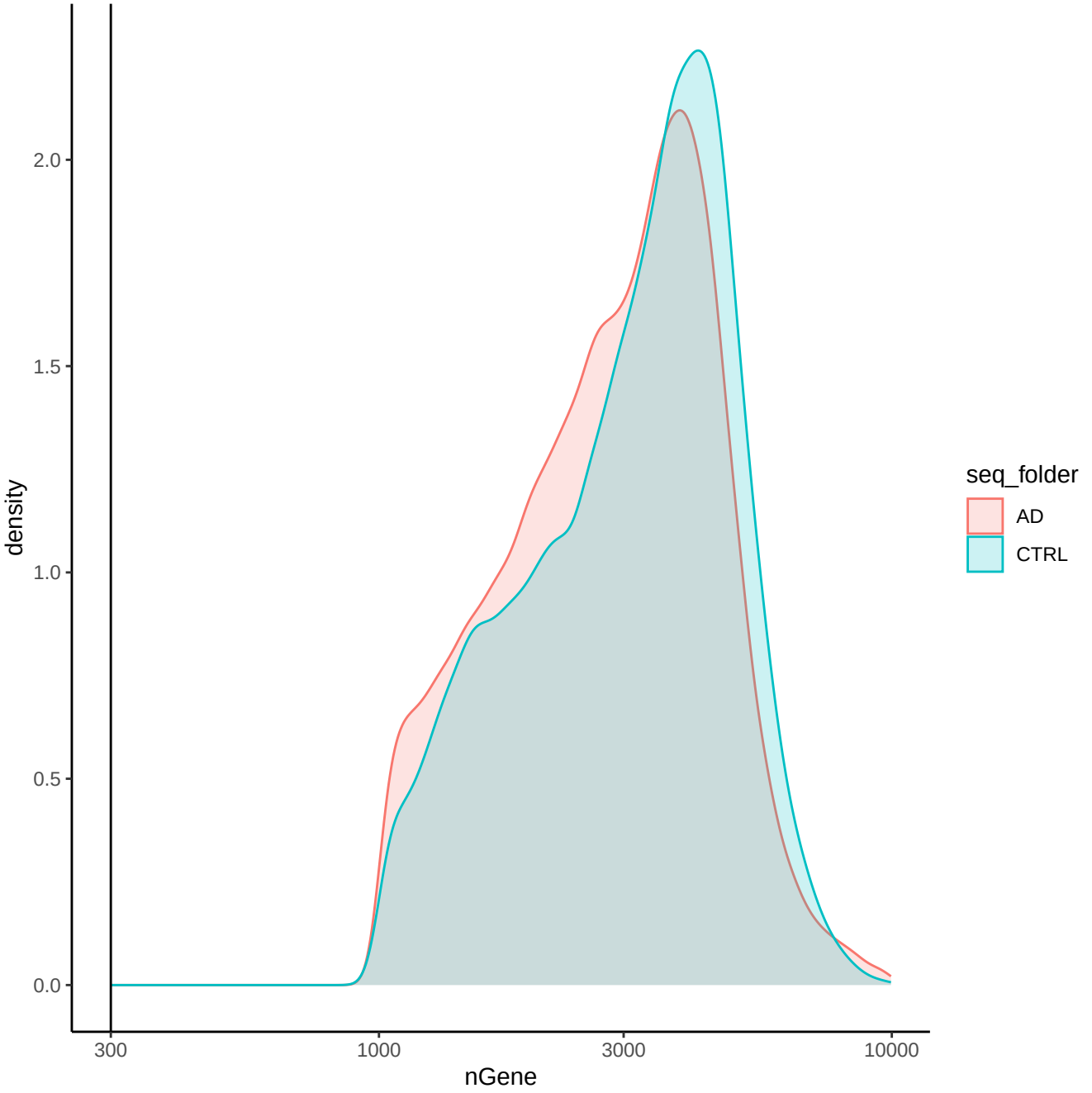
density
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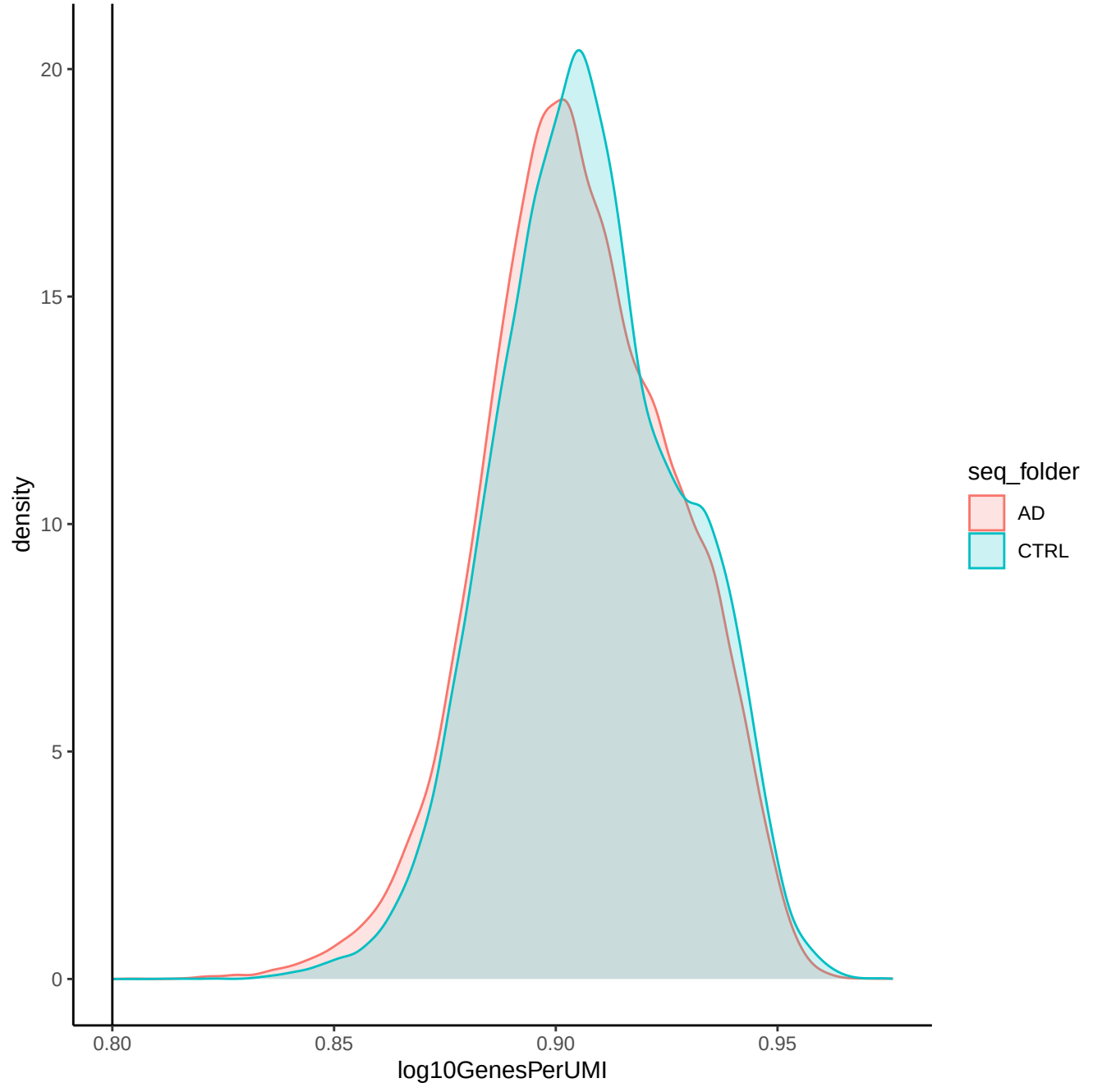
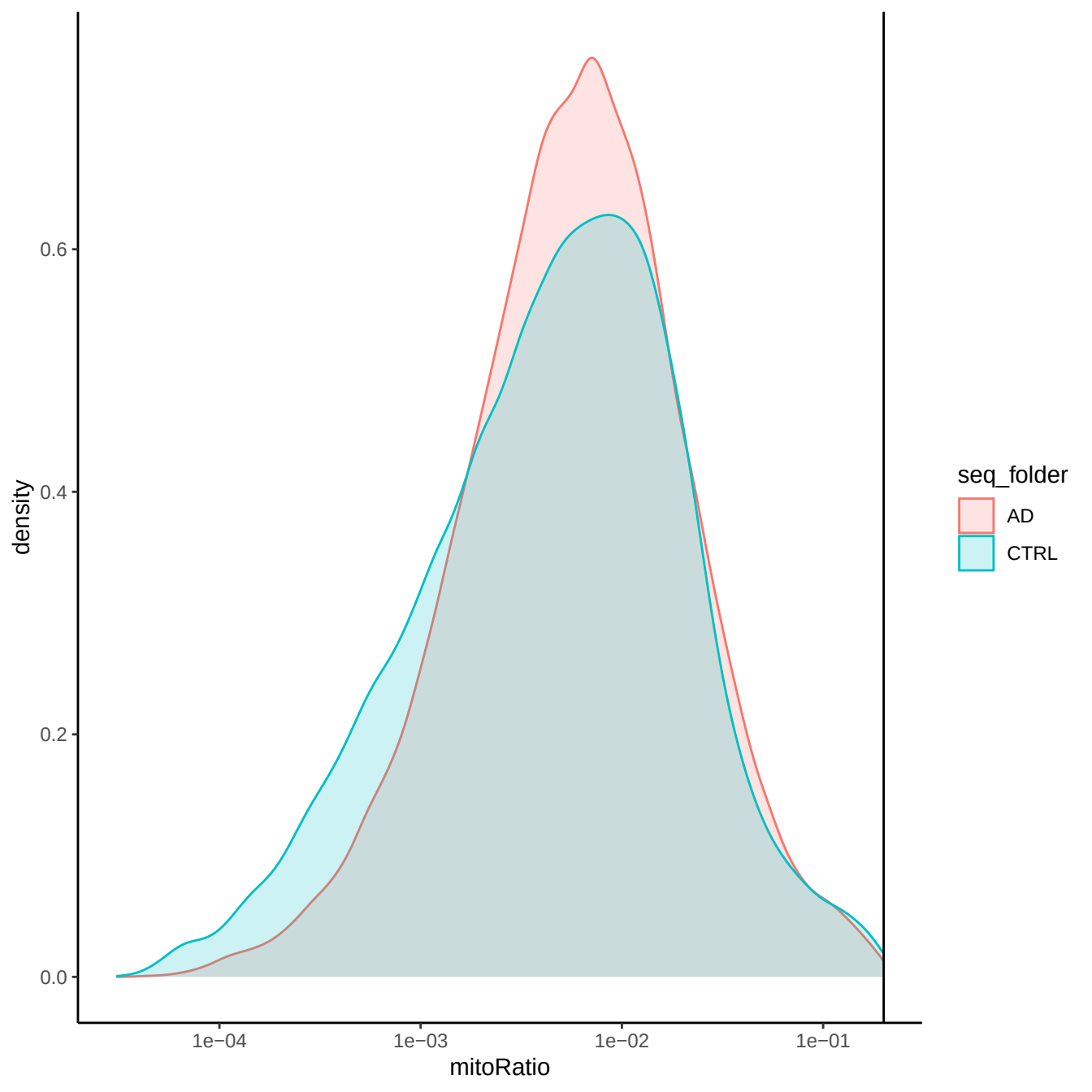
1.5
1.0
0.5
0.0
300
1000
3000
10000
nGene
seq_folder
AD
CTRL


density
20
15
10
5
0
0.80
0.85
0.90
0.95
log10GenesPerUMI
seq_folder
AD
CTRL


density
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0.4
0.2
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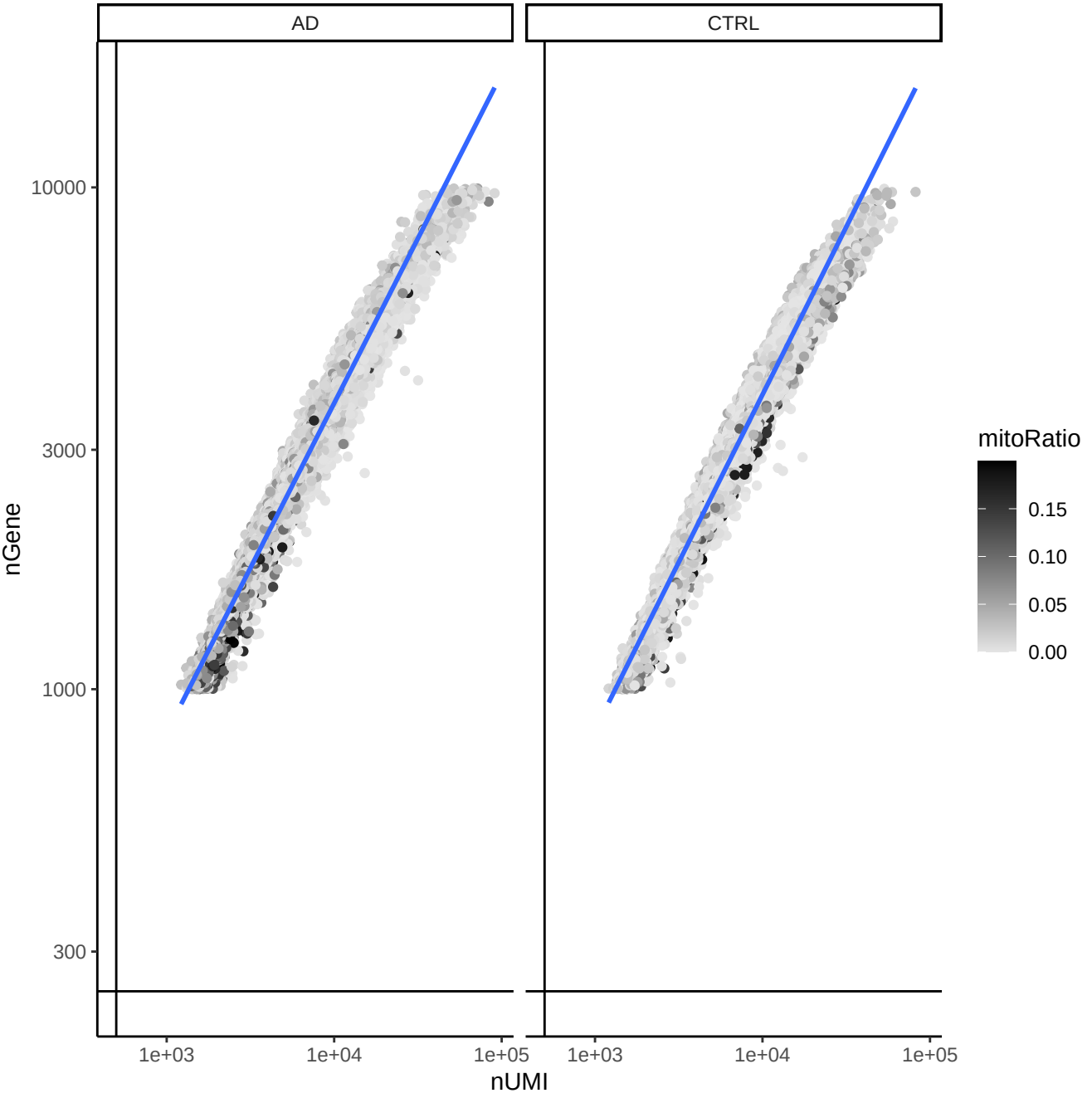
1e−04
1e−03
1e−02
1e−01
mitoRatio
seq_folder
AD
CTRL


AD
CTRL
10000
3000
1000
300
nGene
1e+03
1e+04
1e+05
nUMI
mitoRatio
0.15
0.10
0.05
0.00