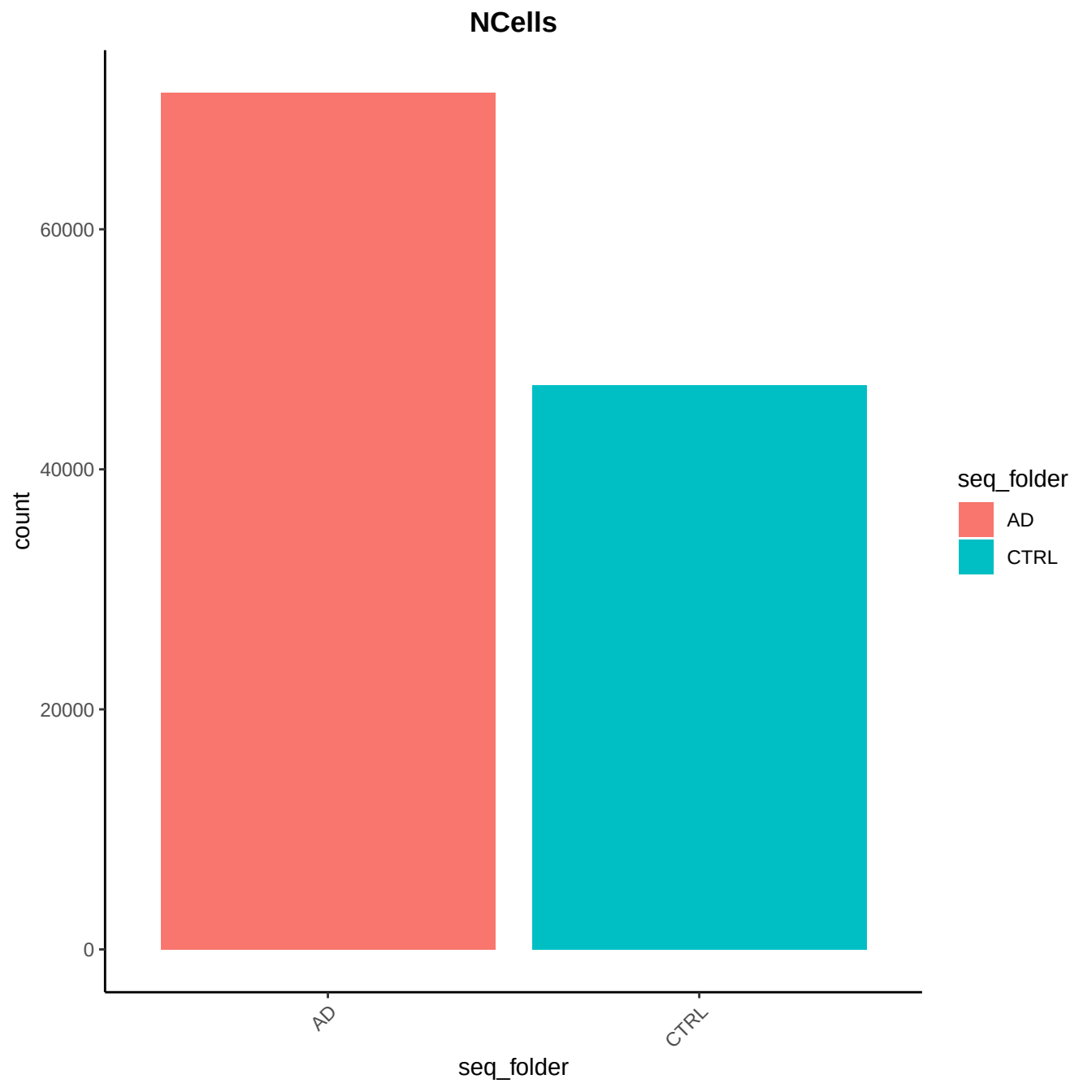
NCells
count
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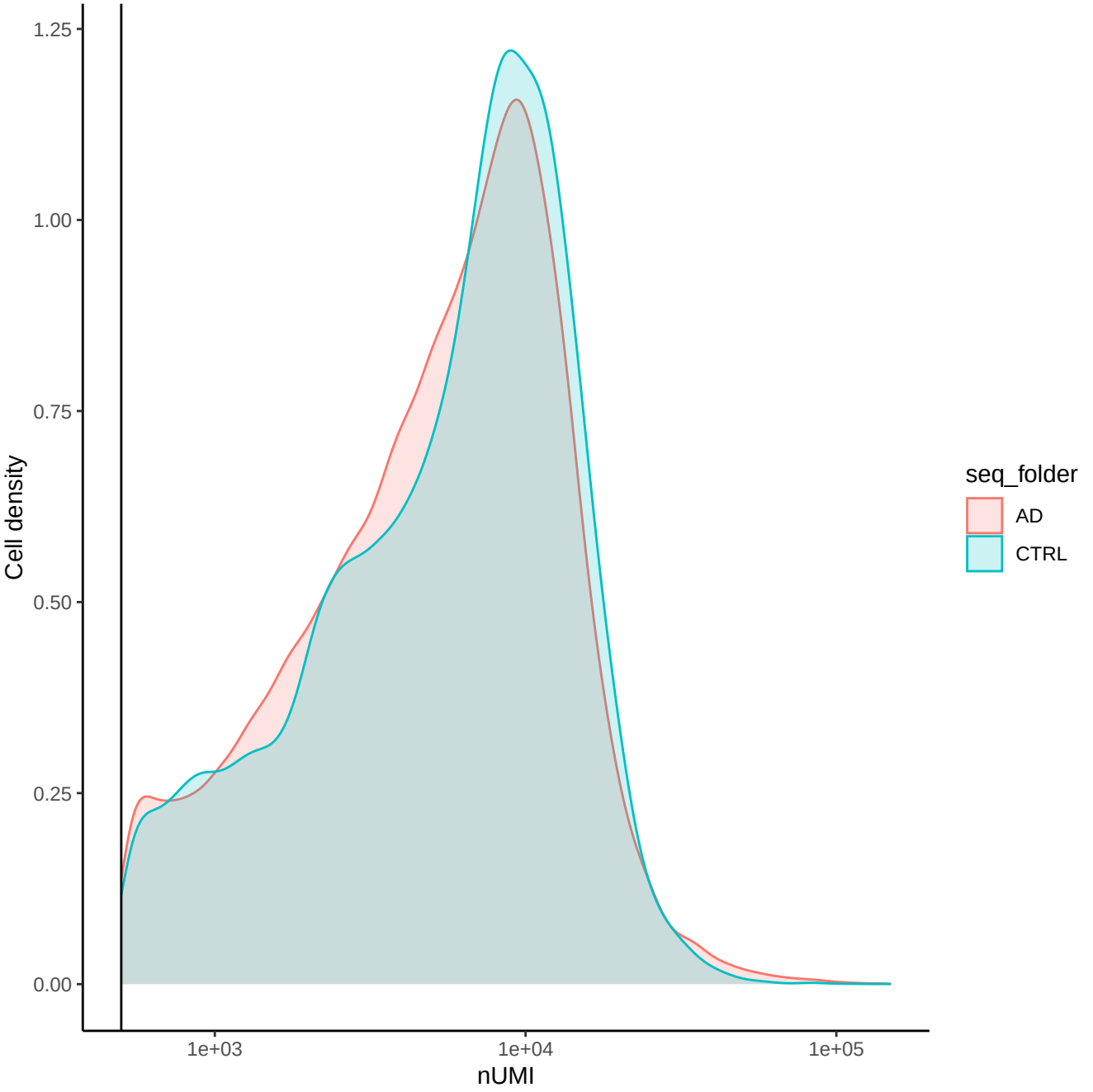
40000
20000
0
AD
CTRL
seq_folder
seq_folder
AD
CTRL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Cell density
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1e+04
1e+05
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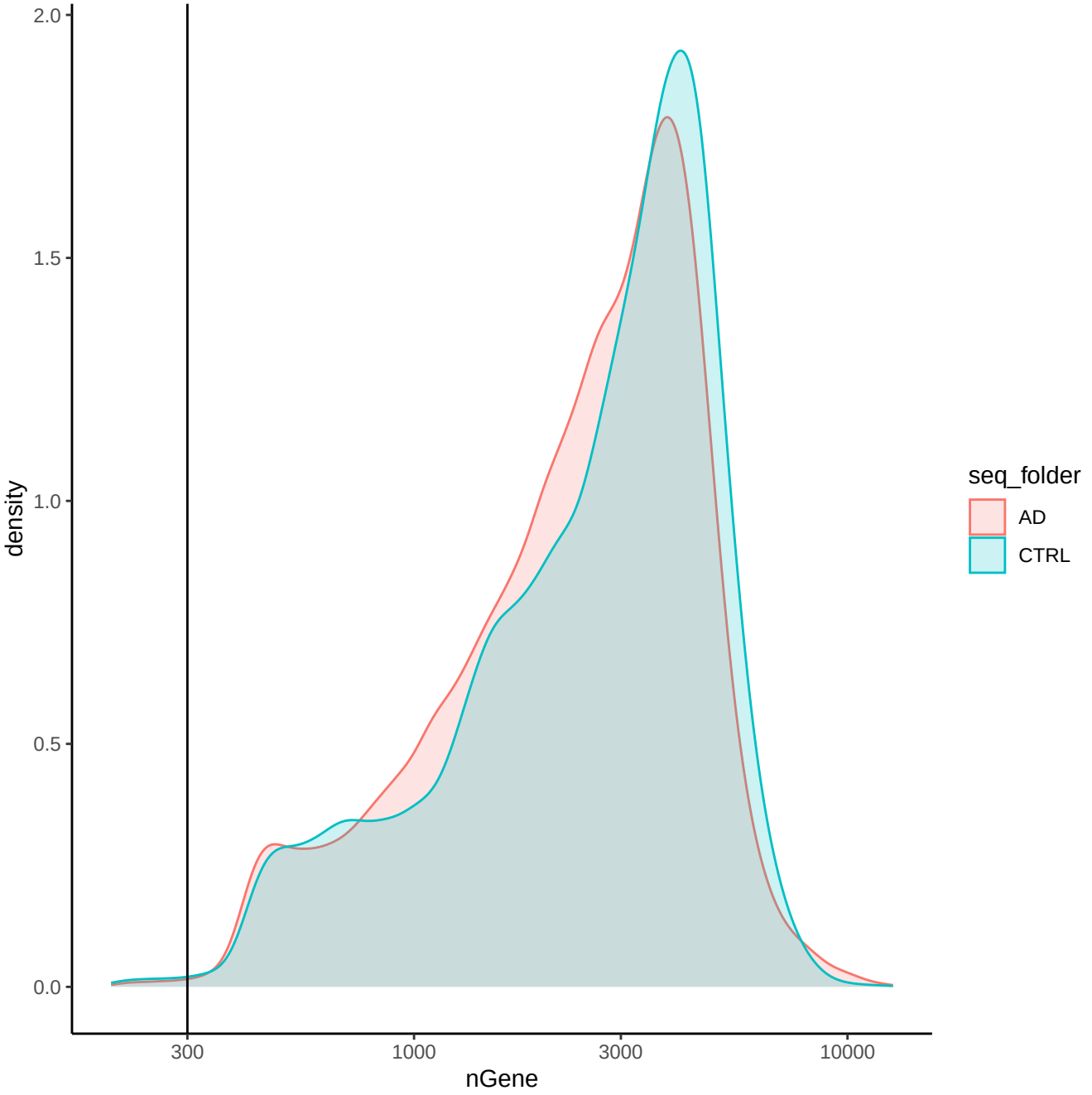
AD
CTRL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0.0
density
300
1000
3000
10000
nGene
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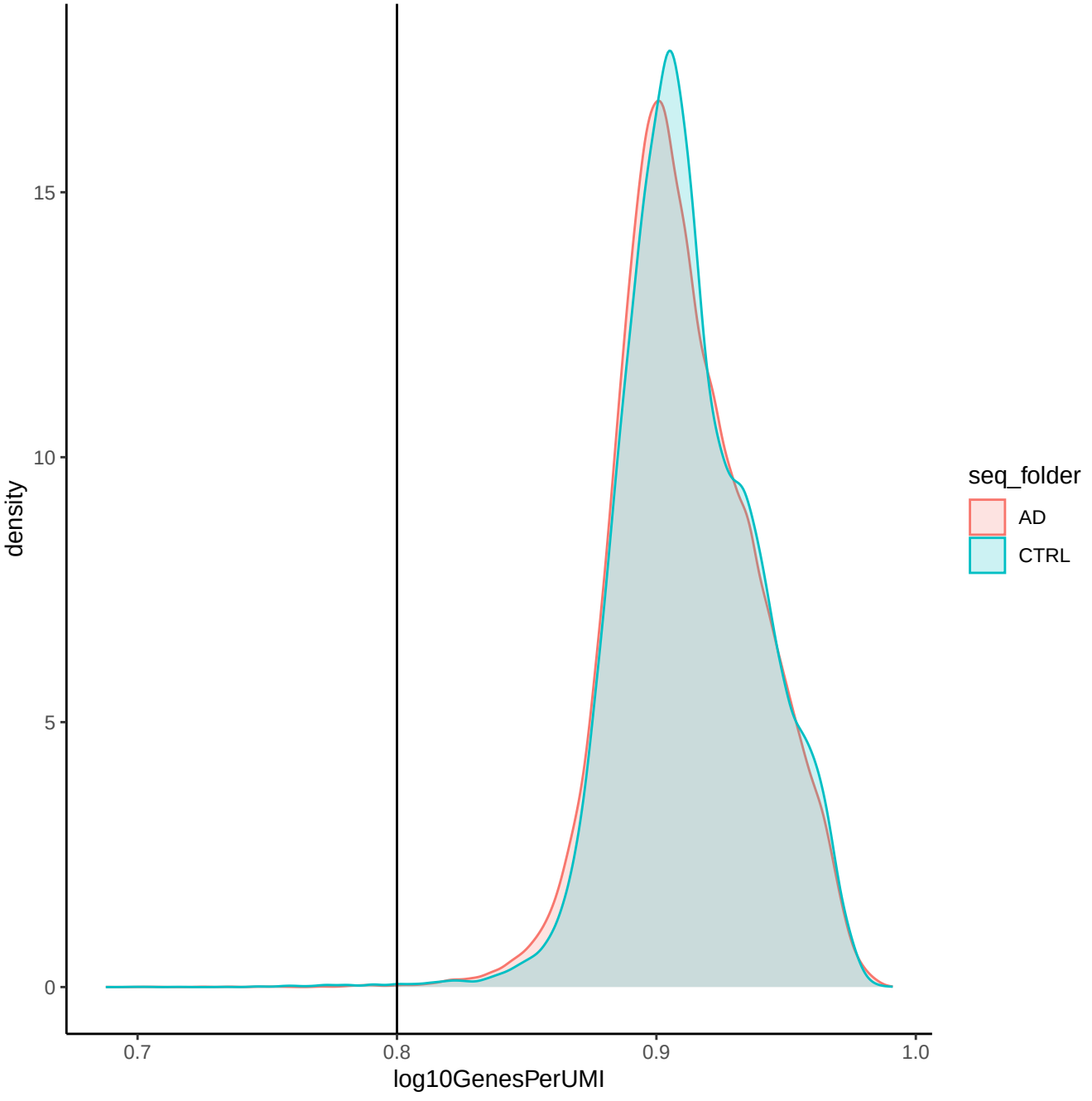
AD
CTRL


density
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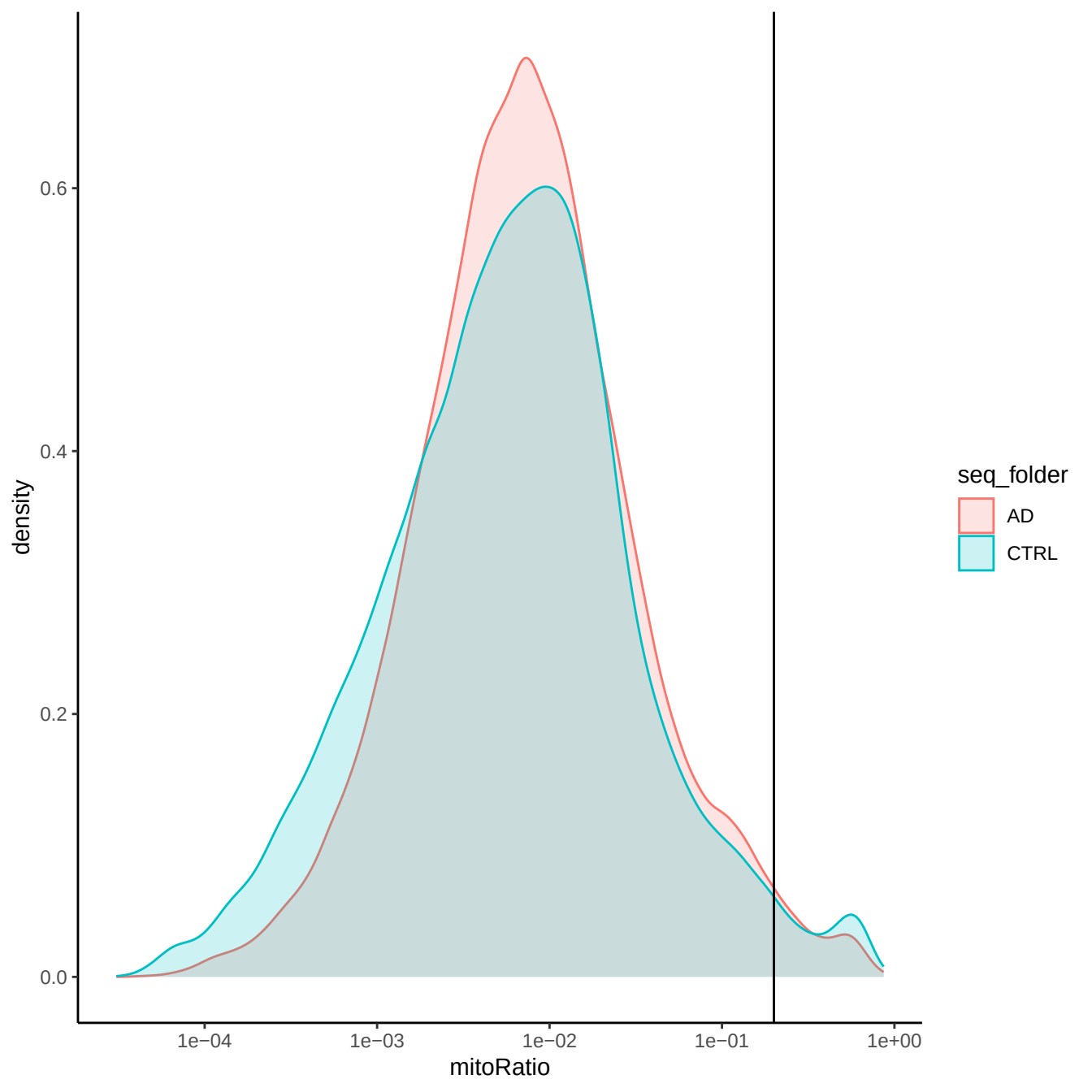
10
5
0
0.7
0.8
0.9
1.0
log10GenesPerUMI
seq_folder
AD
CTRL


density
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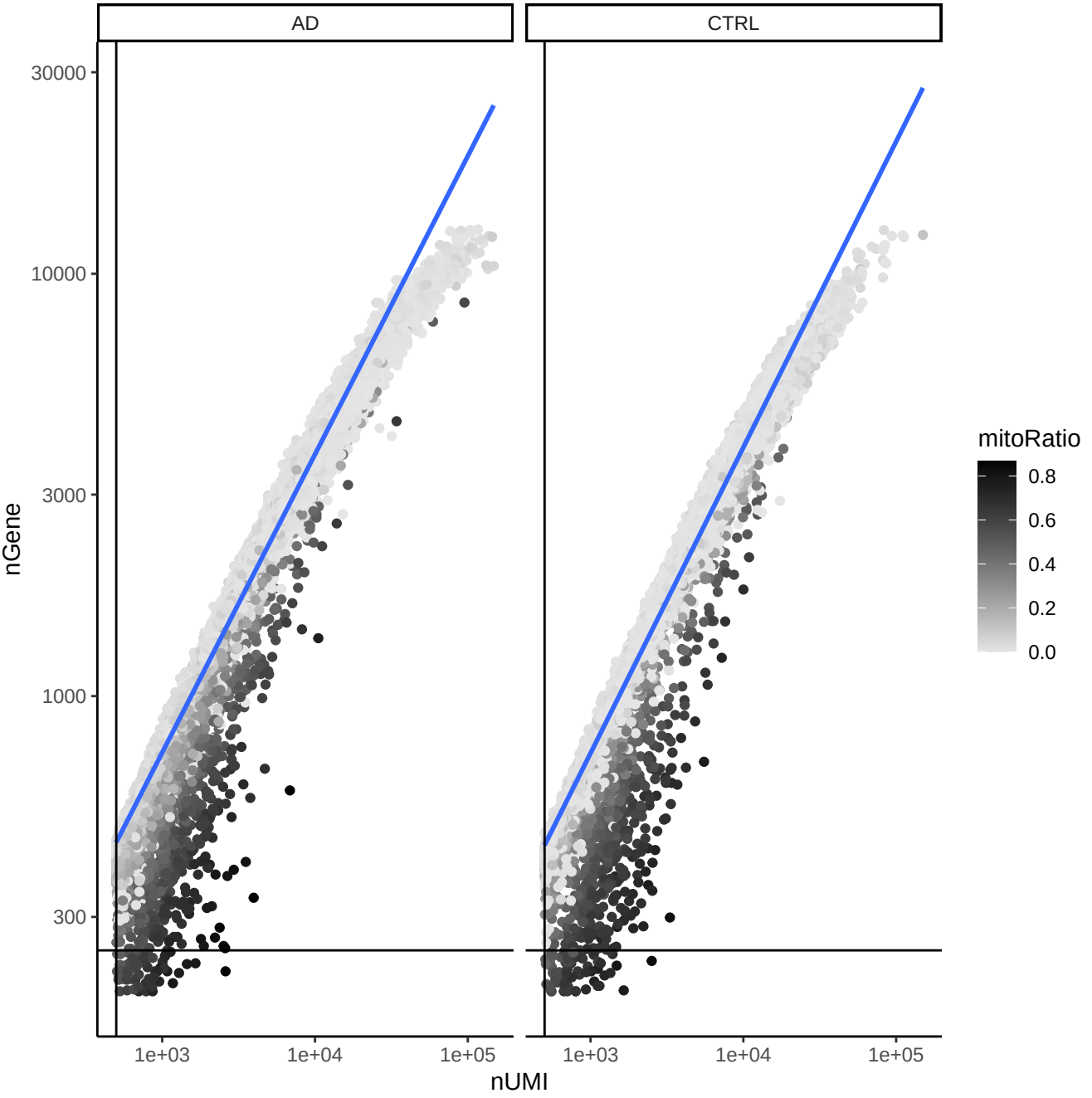
0.4
0.2
0.0
1e−04
1e−03
1e−02
1e−01
1e+00
mitoRatio
seq_folder
AD
CTRL


AD
CTRL
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10000
3000
1000
300
nGene
1e+03
1e+04
1e+05
nUMI
mitoRatio
0.8
0.6
0.4
0.2
0.0