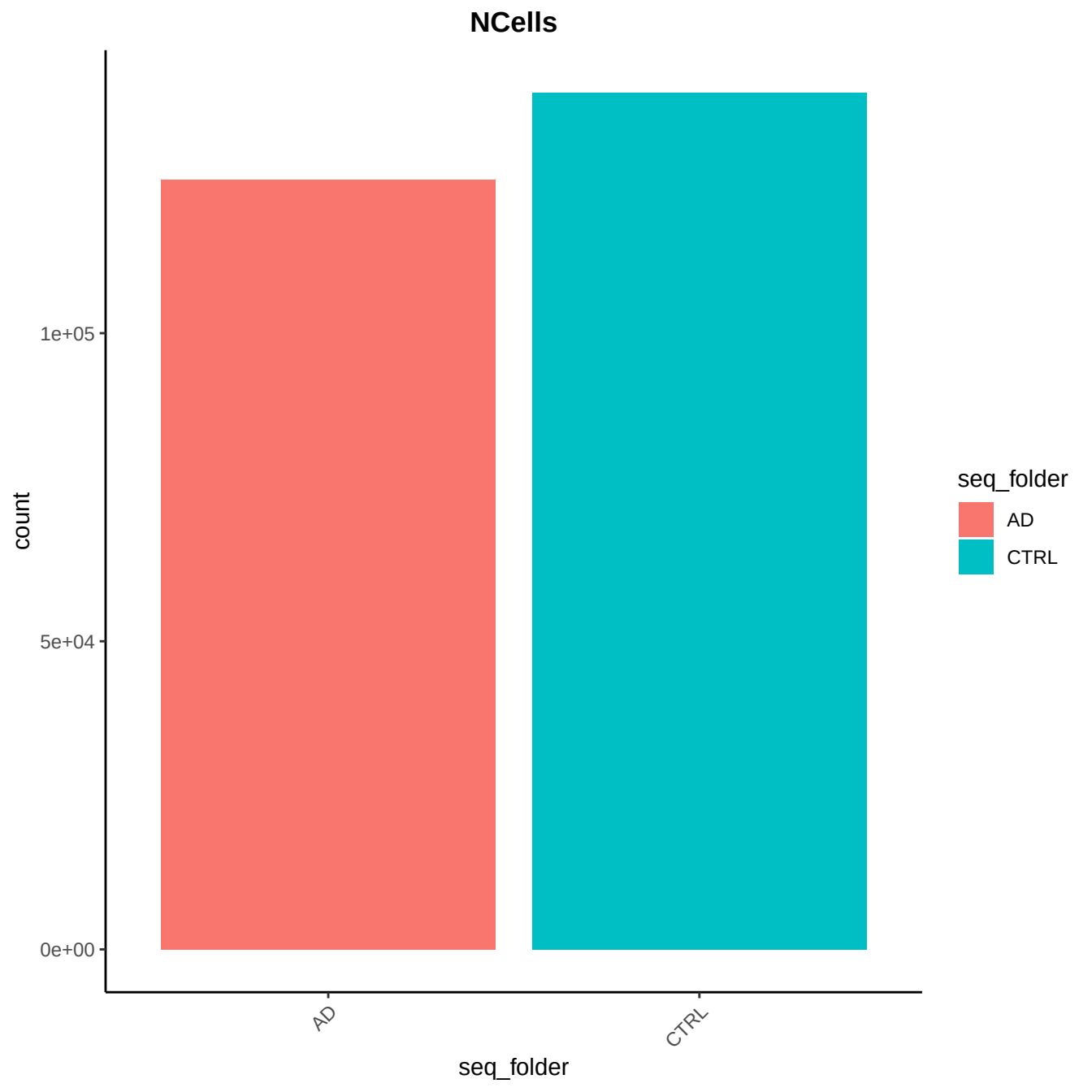
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count
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5e+04
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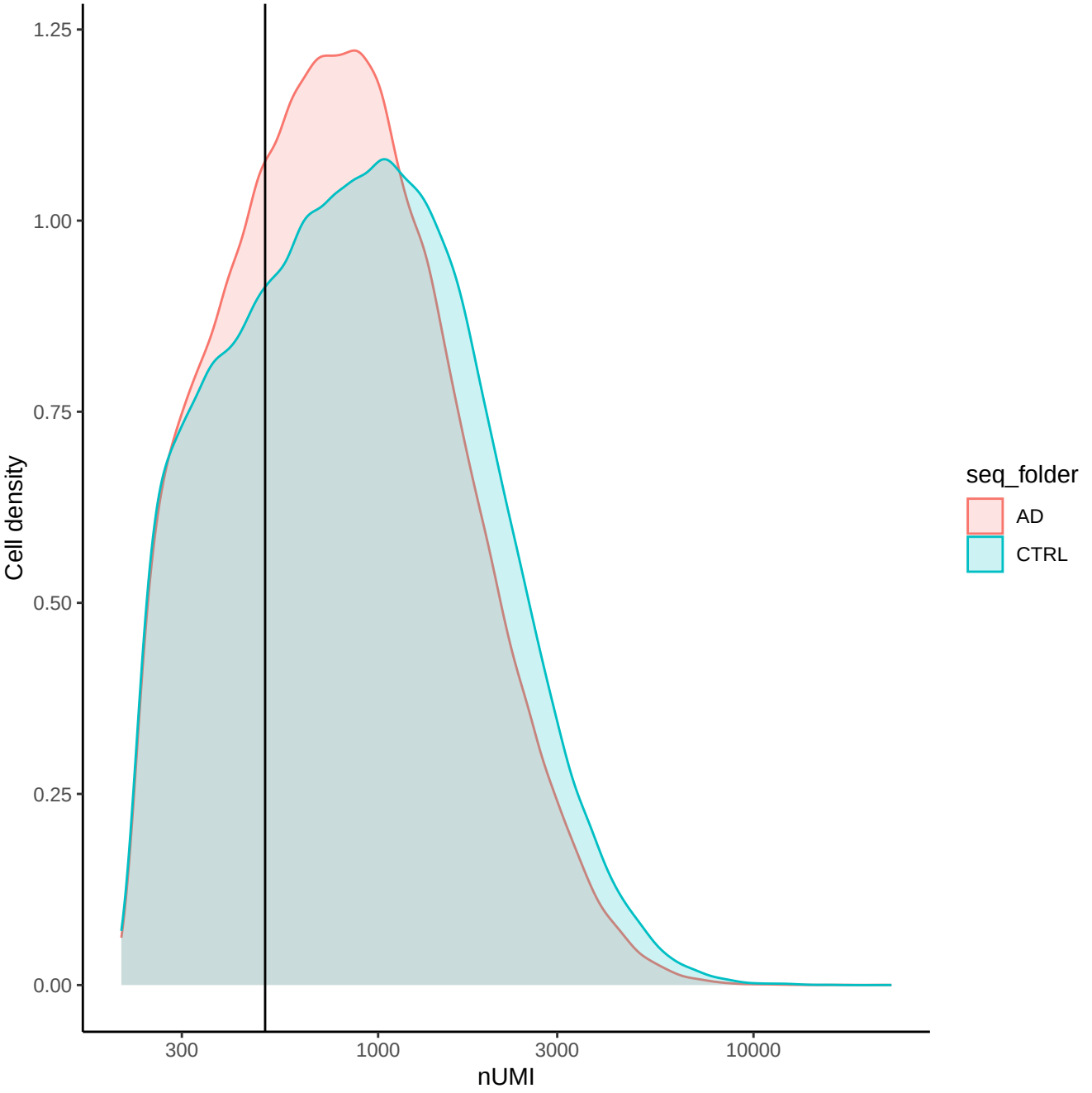
AD
CTRL
seq_folder
seq_folder
AD
CTRL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Cell density
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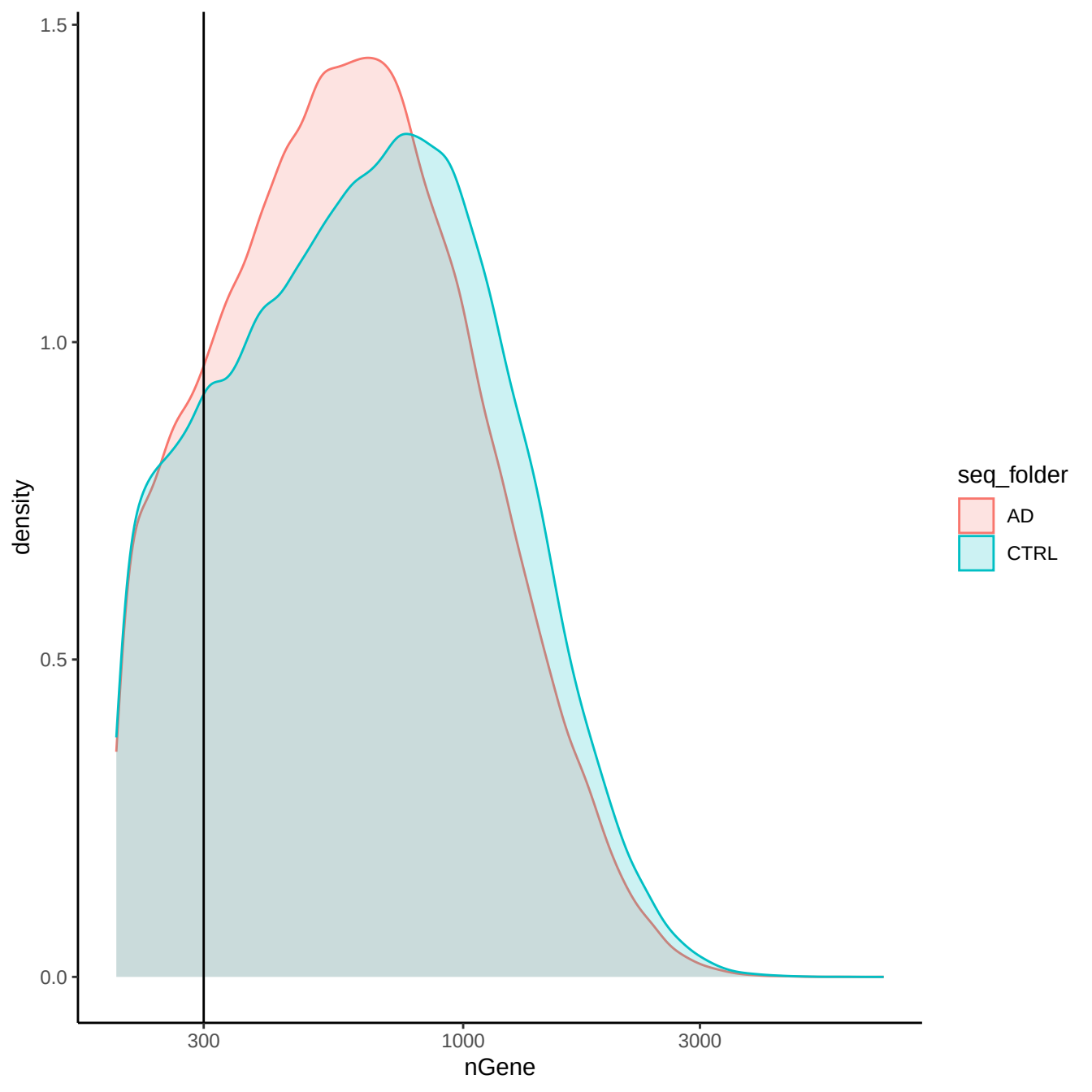
AD
CTRL


1.5
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0.5
0.0
density
300
1000
3000
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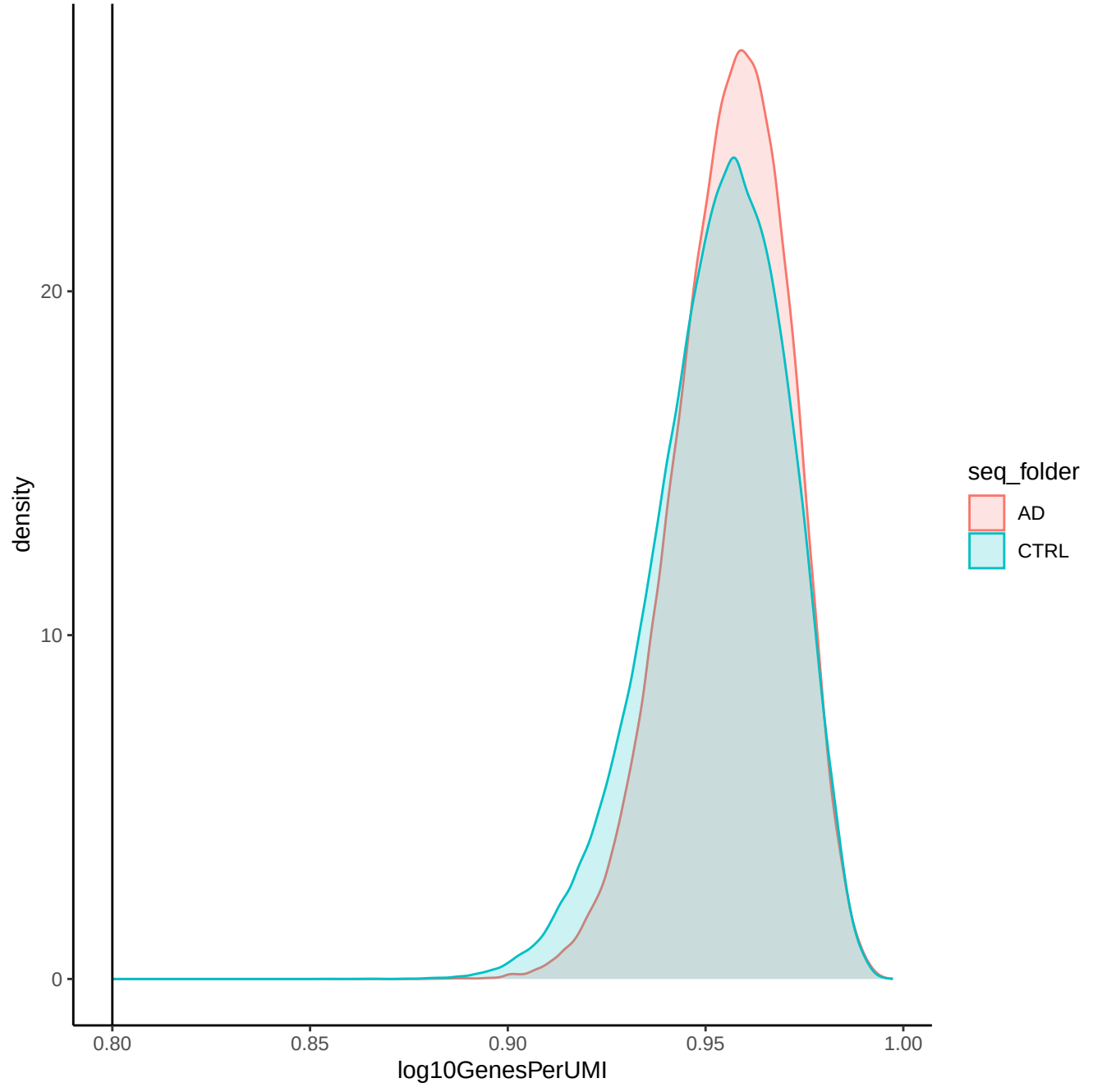
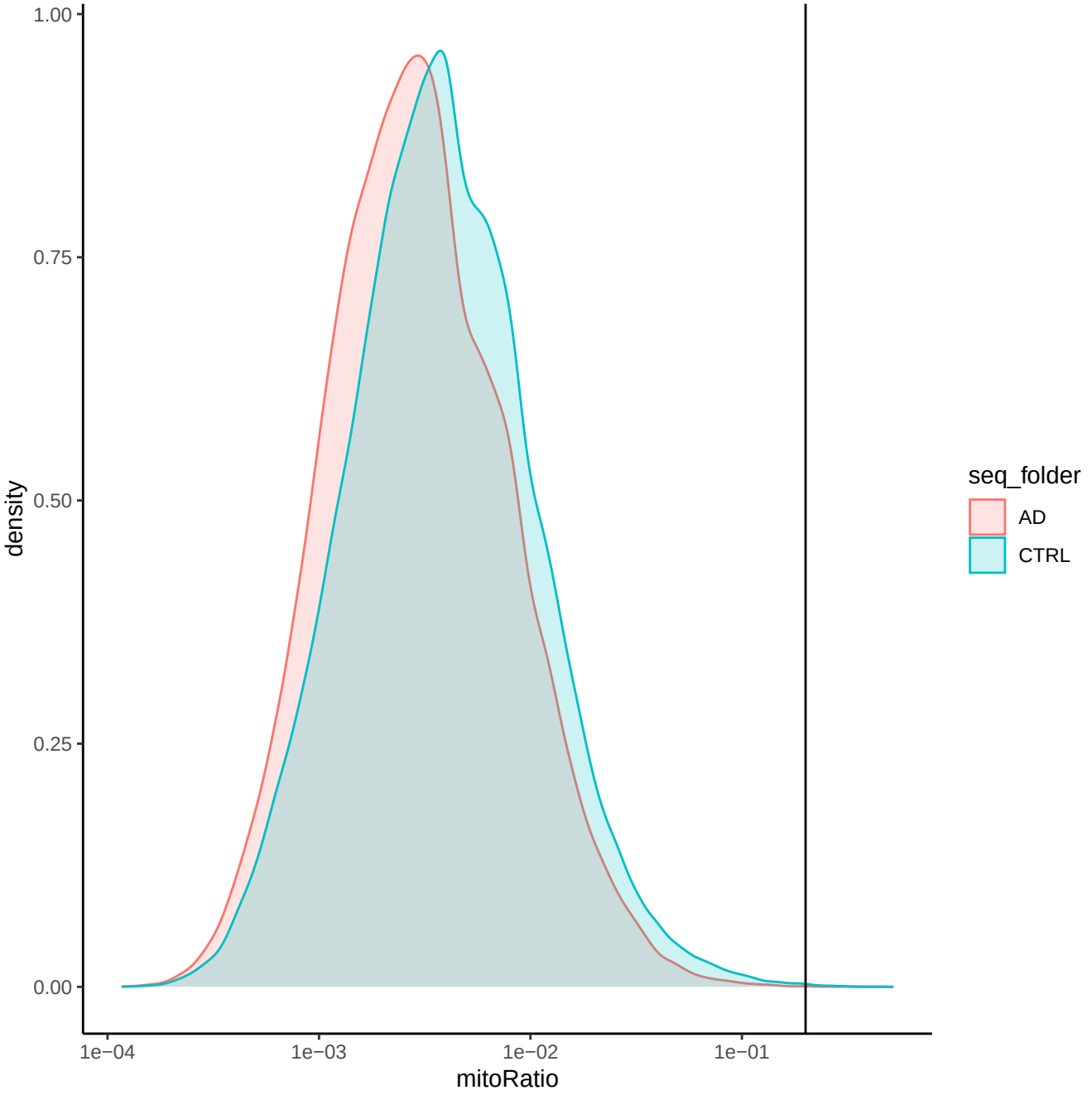
AD
CTRL


density
20
10
0
0.80
0.85
0.90
0.95
1.00
log10GenesPerUMI
seq_folder
AD
CTRL


1.00
0.75
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0.25
0.00
density
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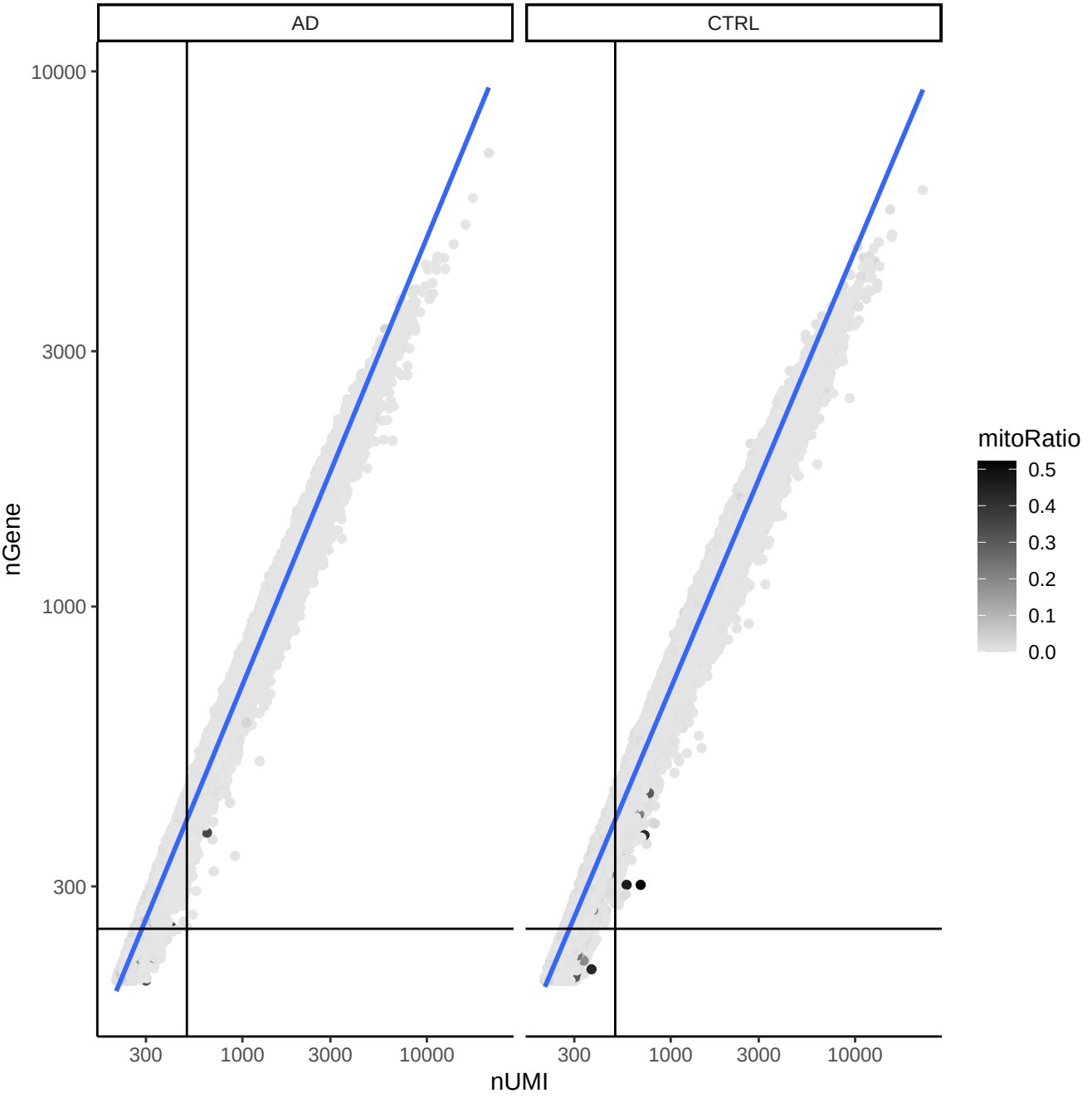
1e−03
1e−02
1e−01
mitoRatio
seq_folder
AD
CTRL


AD
CTRL
10000
3000
1000
300
nGene
300
1000
3000
10000
nUMI
mitoRatio
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0.4
0.3
0.2
0.1
0.0