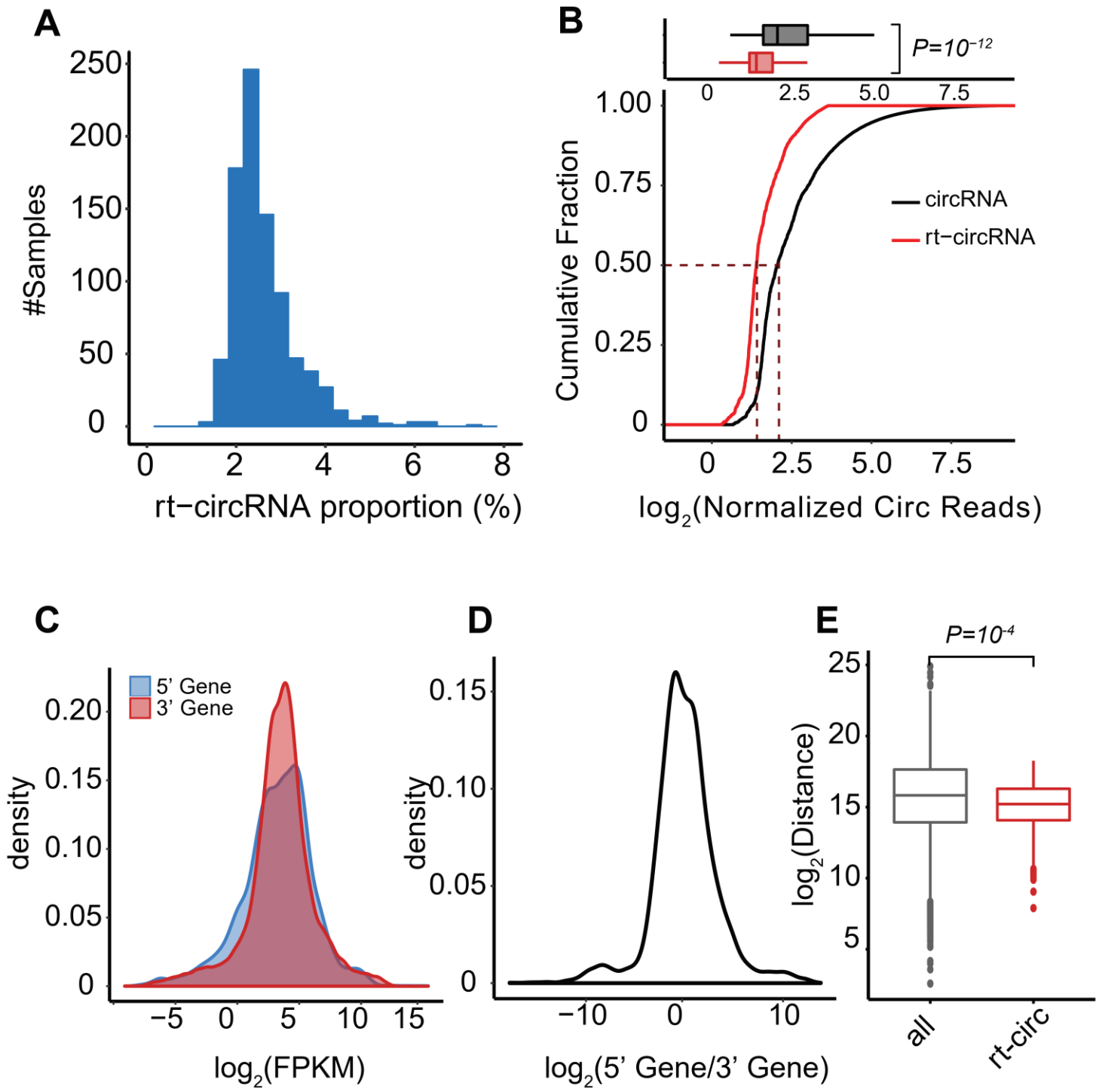
# Figure S3


A
#Samples
250
200
150
100
50
0
0
2
4
6
8
rt−circRNA proportion (%)
B
P=10−12
0
2.5
5.0
7.5
Cumulative Fraction
1.00
0.75
0.50
0.25
0
circRNA
rt−circRNA
log2(Normalized Circ Reads)
C
5' Gene
3' Gene
density
0.20
0.15
0.10
0.05
0
−5
0
5
10
15
log2(FPKM)
D
density
0.15
0.10
0.05
0
−10
0
10
log2(5' Gene/3' Gene)
E
P=10-4
log2(Distance)
25
20
15
10
5
all
rt-circ