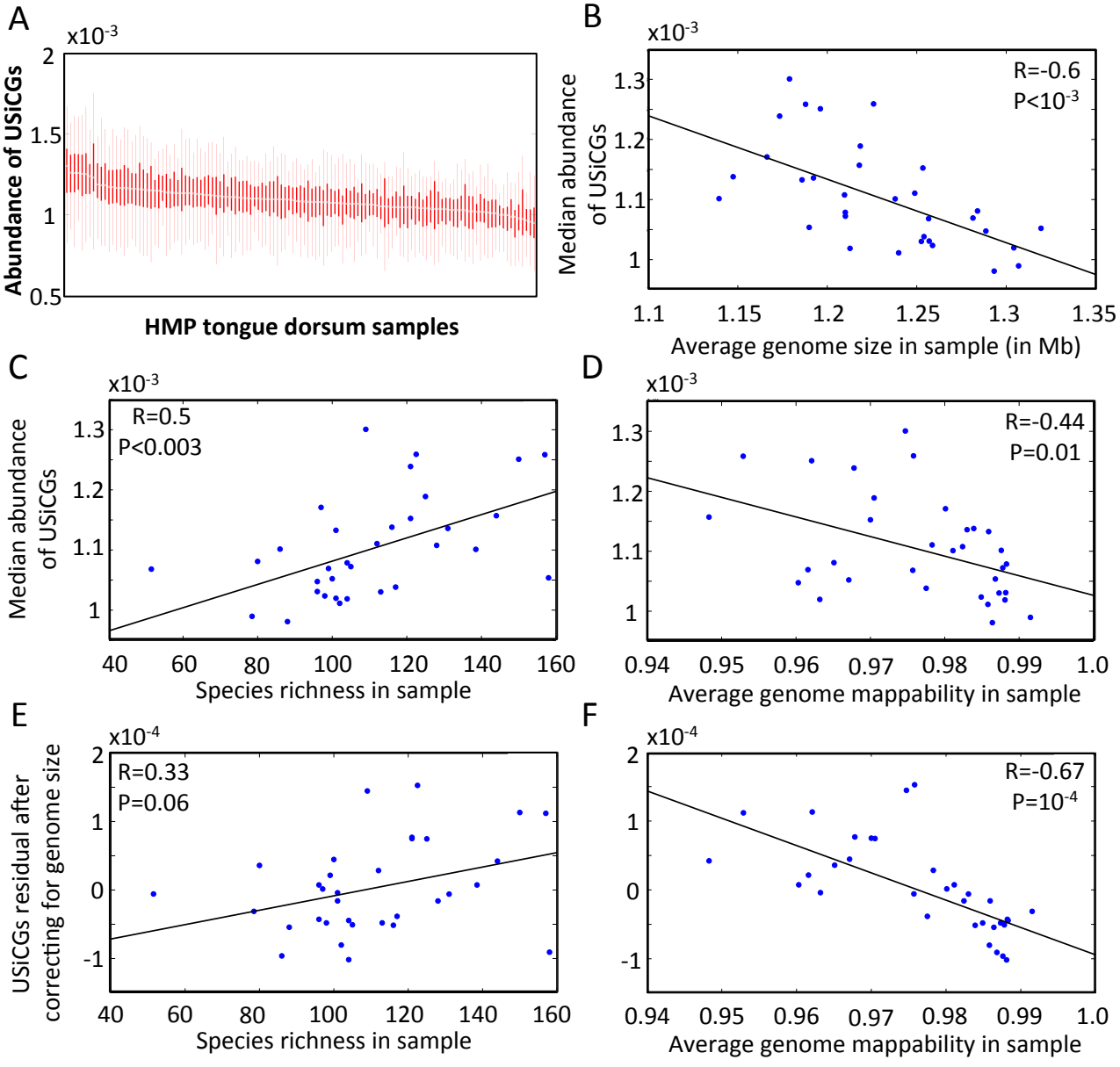
A
x10-3
Abundance of USiCGs
HMP tongue dorsum samples
B
x10-3
Median abundance of USiCGs
R=-0.6
P<10-3
Average genome size in sample (in Mb)
C
x10-3
Median abundance of USiCGs
R=0.5
P<0.003
Species richness in sample
D
x10-3
R=-0.44
P=0.01
Average genome mappability in sample
E
x10-4
USiCGs residual after correcting for genome size
R=0.33
P=0.06
Species richness in sample
F
x10-4
R=-0.67
P=10-4
Average genome mappability in sample