

Bin\_1

coverage per genome

0.10

0.05

0.00

sample1

sample2

sample3

sample4

sample5

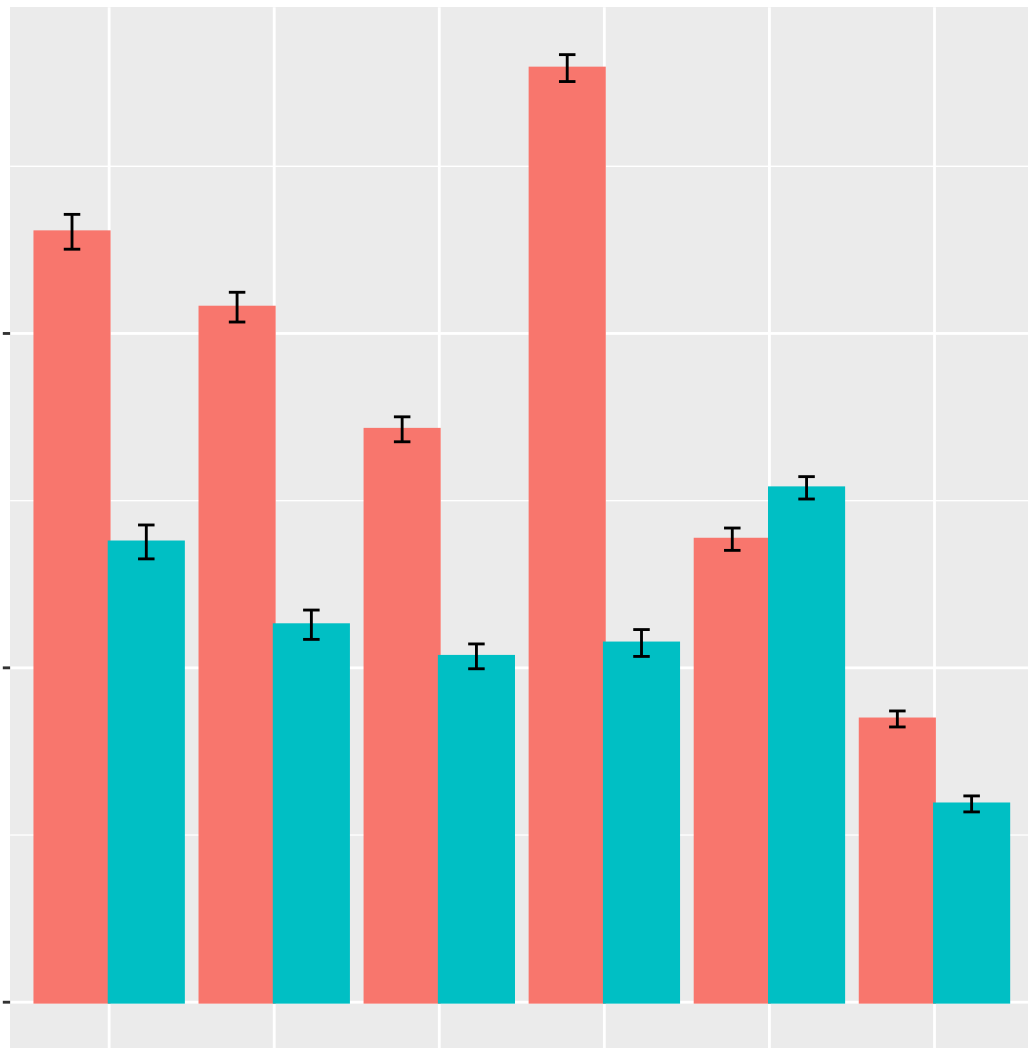
sample6

Samples

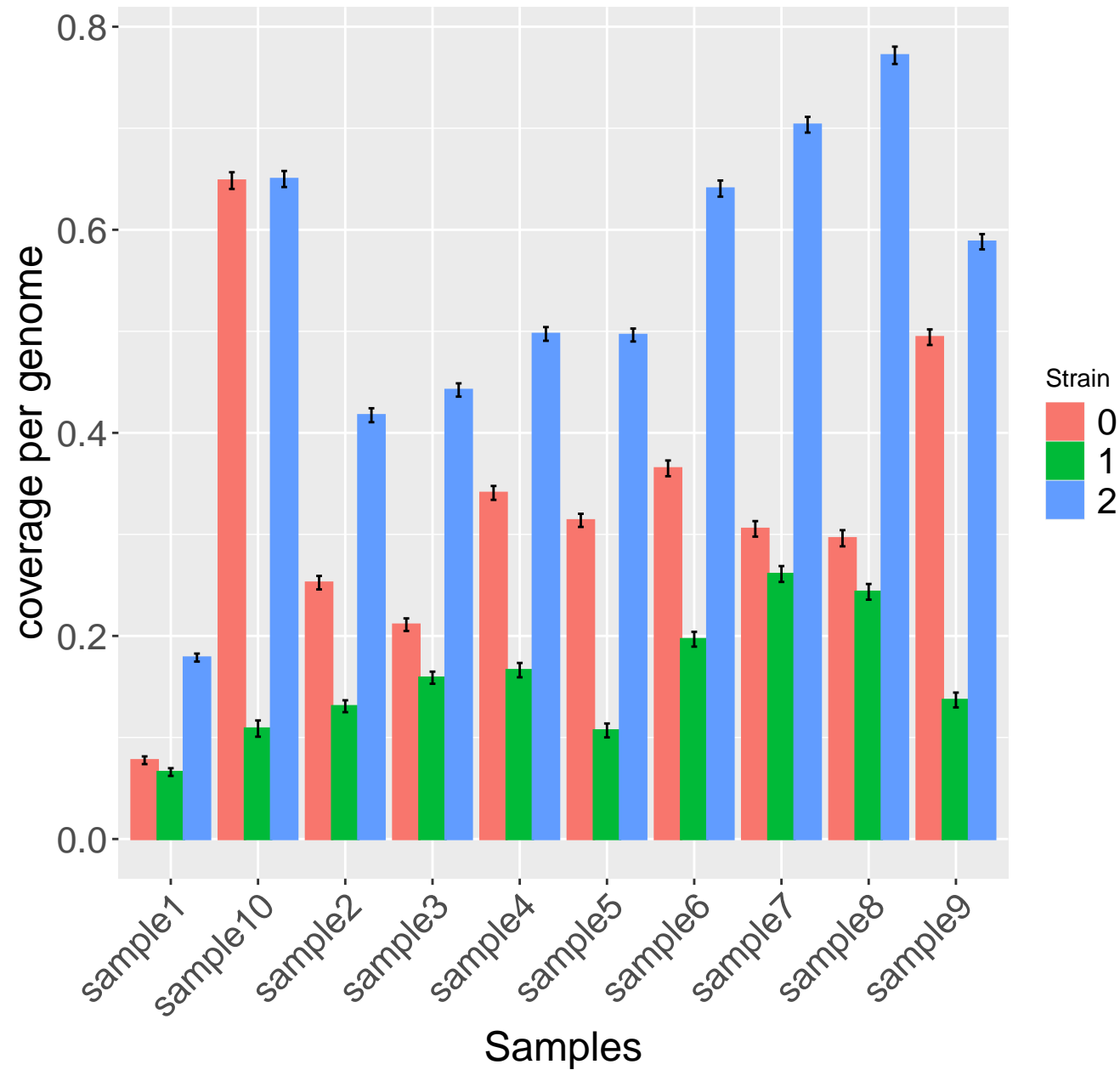
Strain

0

1



Bin\_2



Bin\_8

