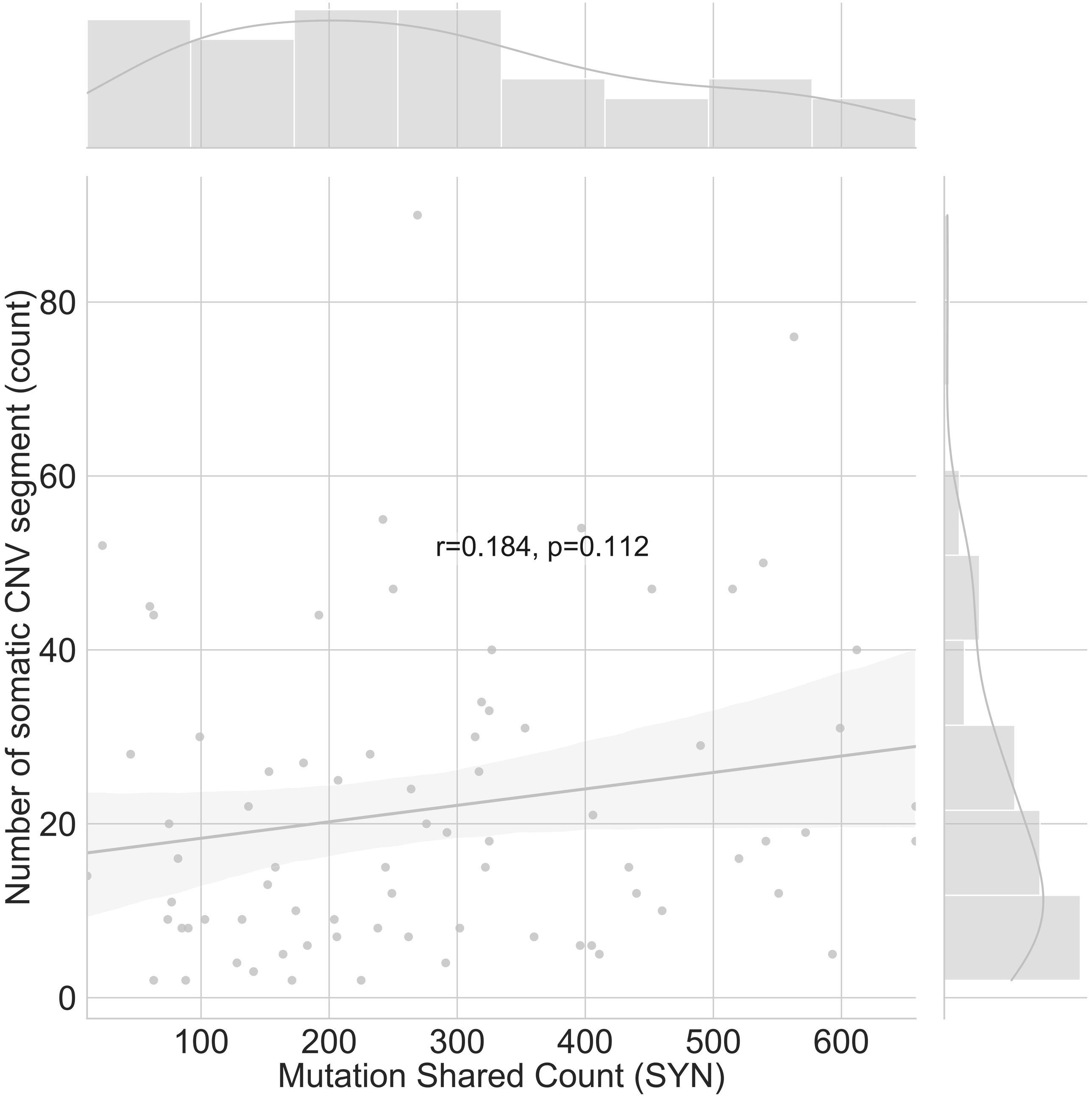
r=0.184, p=0.112
Number of somatic CNV segment (count)
0
20
40
60
80
100
200
300
400
500
600
Mutation Shared Count (SYN)