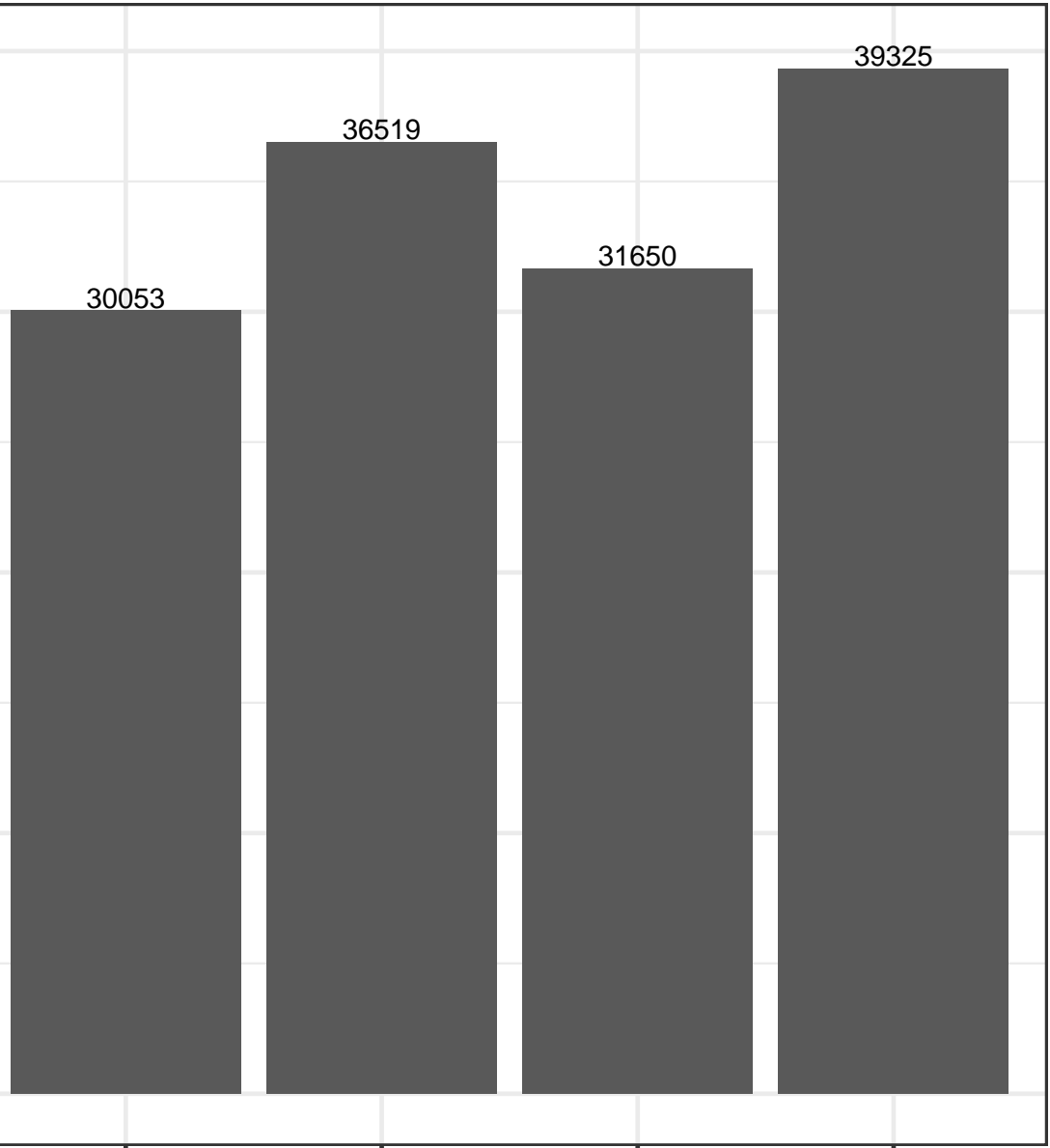


Total number of transcripts sequences

40000
30000
20000
10000
0



WT_1

WT_2

KD_1

KD_2

Sample