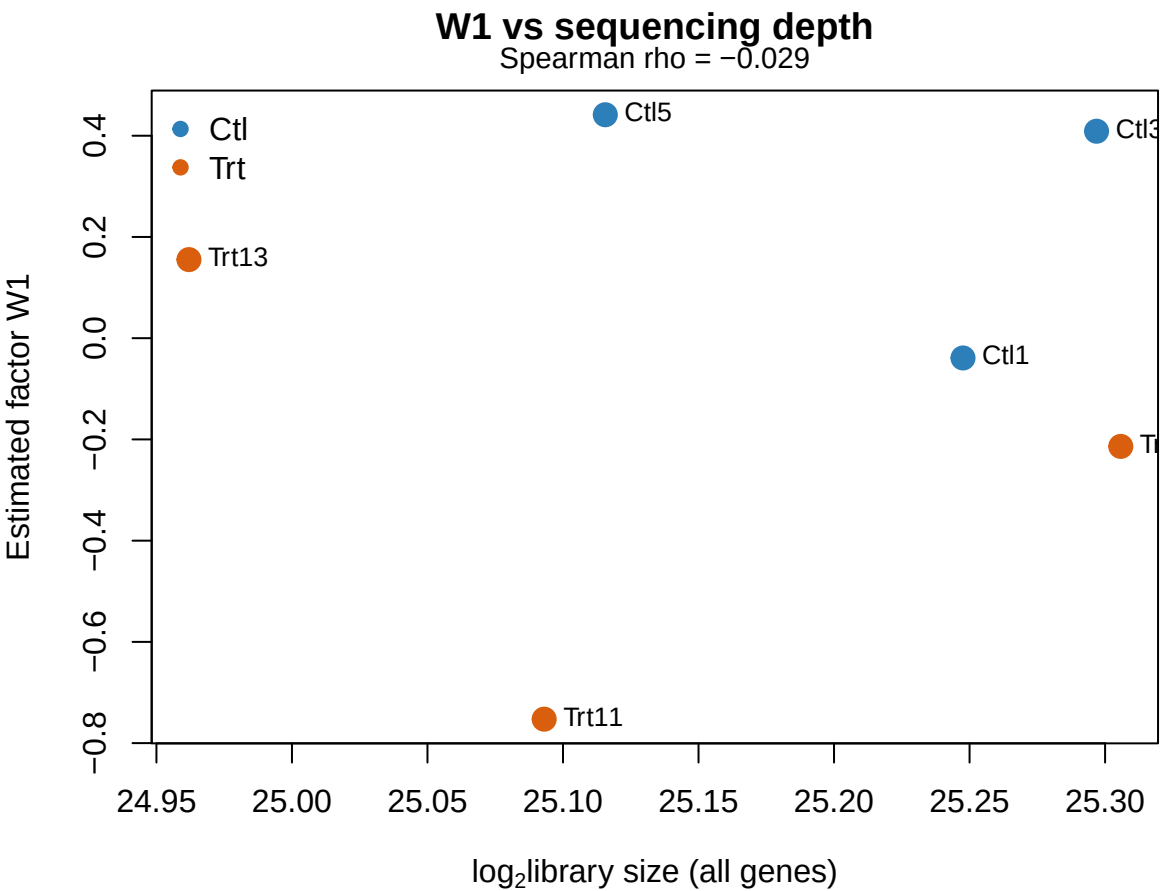

W1 vs sequencing depth
Spearman rho = −0.029
Ctl
Trt
Ctl5
Ctl3
Trt13
Ctl1
Tr
Trt11
Estimated factor W1
0.4
0.2
0.0
−0.2
−0.4
−0.6
−0.8
24.95
25.00
25.05
25.10
25.15
25.20
25.25
25.30
log2library size (all genes)