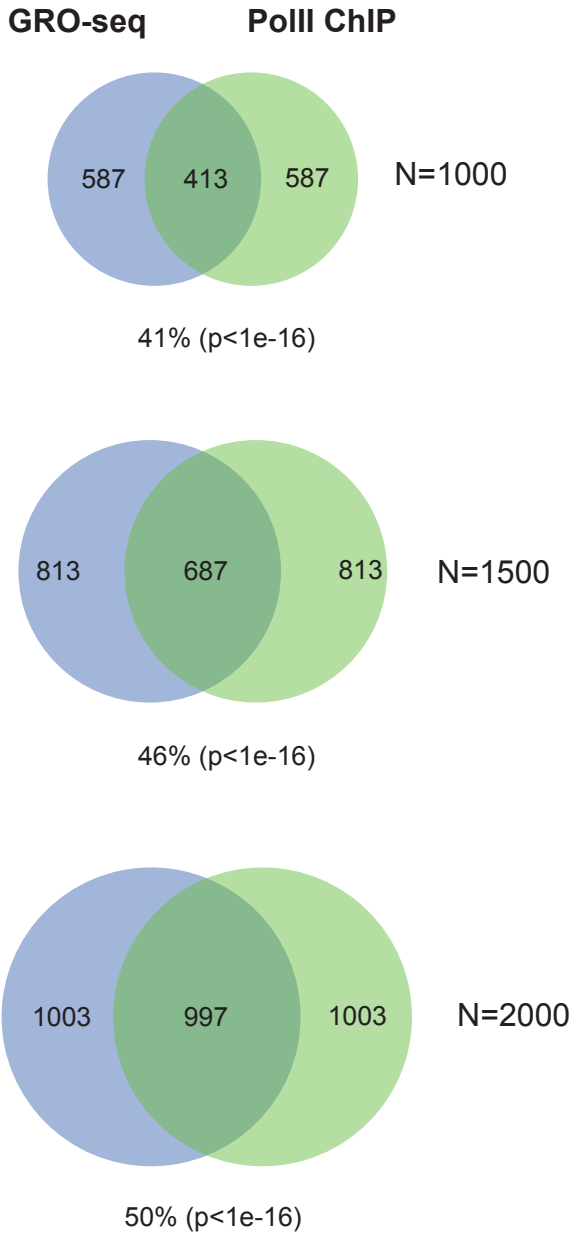

GRO-seq
PolII ChIP
587
413
587
N=1000
41% (p<1e-16)
813
687
813
N=1500
46% (p<1e-16)
1003
997
1003
N=2000
50% (p<1e-16)

Figure S8