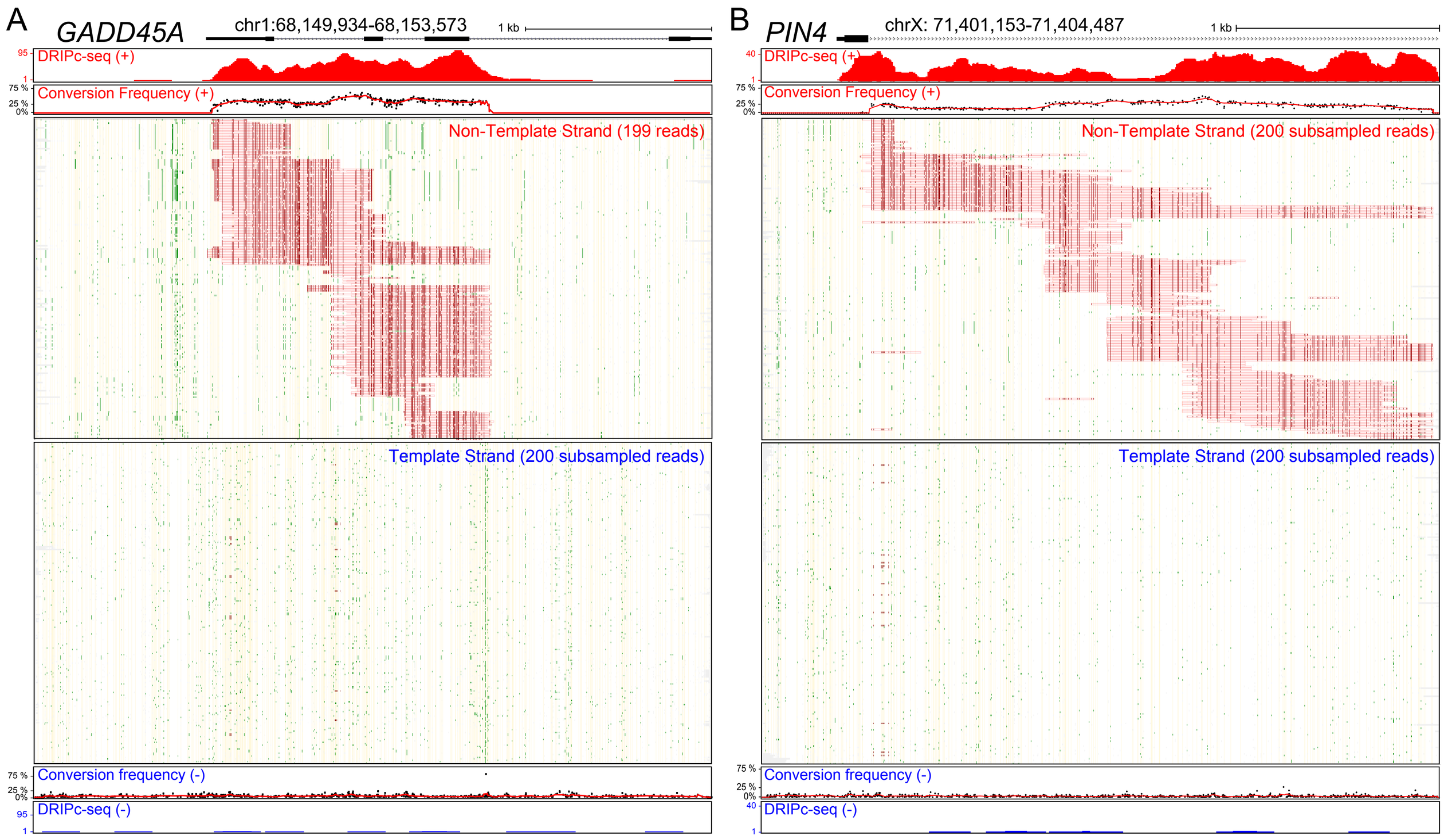
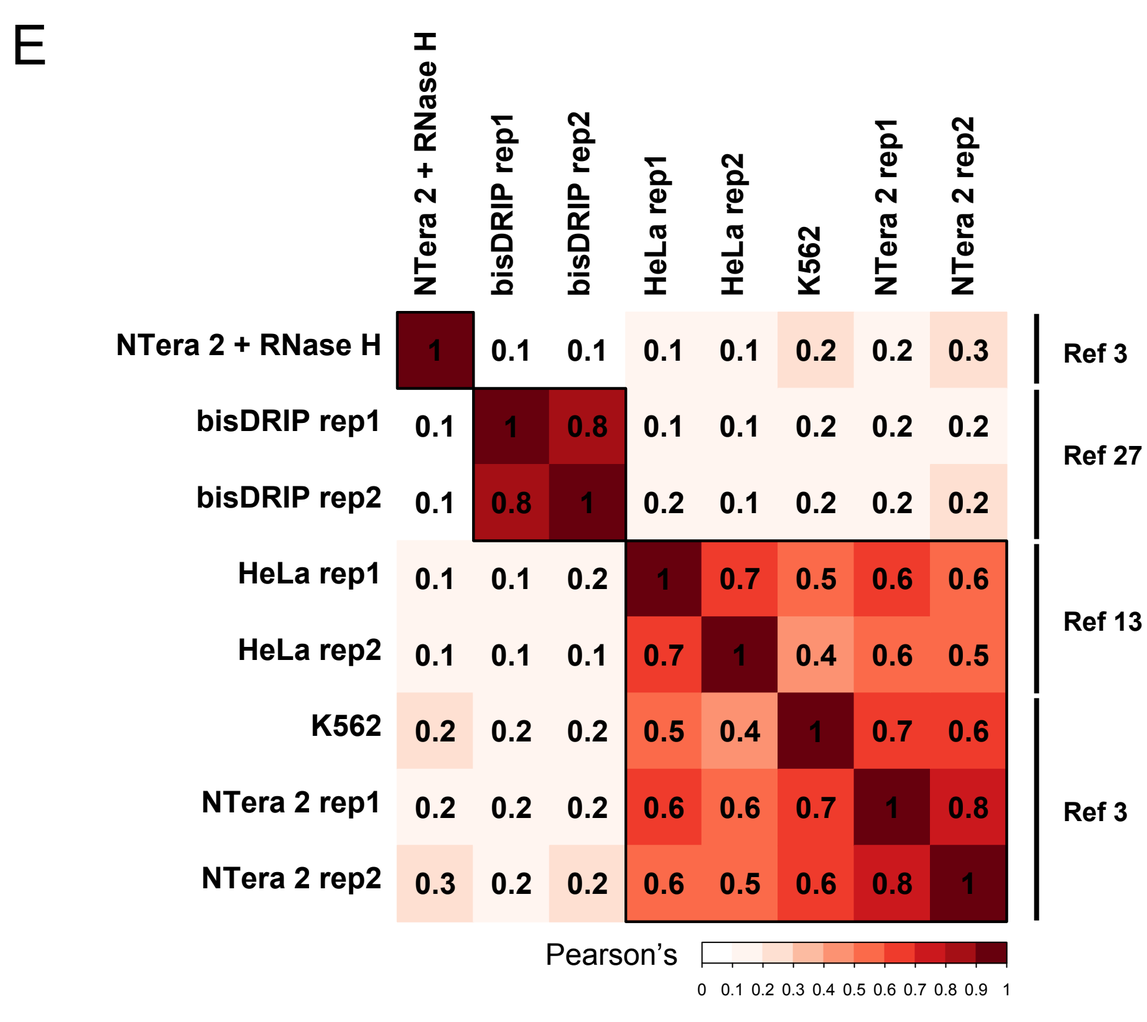
Figure S6


A
GADD45A
chr1:68,149,934-68,153,573
1 kb
DRIPc-seq (+)
Conversion Frequency (+)
Non-Template Strand (199 reads)
Template Strand (200 subsampled reads)
Conversion frequency (-)
DRIPc-seq (-)
B
PIN4
chrX: 71,401,153-71,404,487
1 kb
DRIPc-seq (+)
Conversion Frequency (+)
Non-Template Strand (200 subsampled reads)
Template Strand (200 subsampled reads)
Conversion frequency (-)
DRIPc-seq (-)

C
BTF3
chr5:72,794,135-72,797,124
1 kb
DRIPc-seq (+)
Non-Template Strand
(1,350 reads)

D
MRFAP1L1
chr4:6,708,655-6,711,808
1 kb
DRIPc-seq (-)
Non-Template Strand
(804 reads)


E
NTera 2 + RNase H
bisDRIP rep1
bisDRIP rep2
HeLa rep1
HeLa rep2
K562
NTera 2 rep1
NTera 2 rep2
Ref 3
Ref 27
Ref 13
Ref 3
Pearson's
0 0.1 0.2 0.3 0.4 0.5 0.6 0.7 0.8 0.9 1