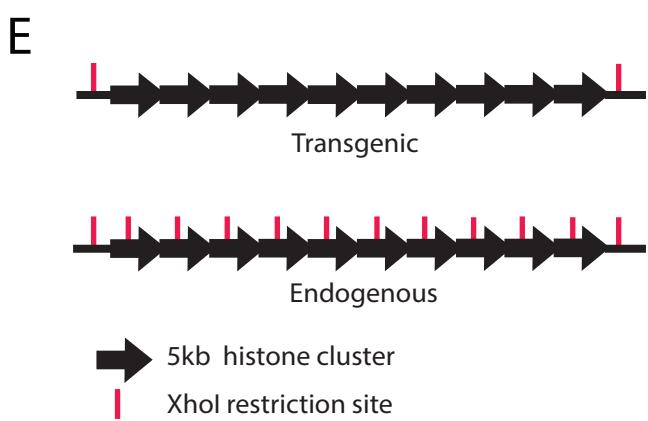
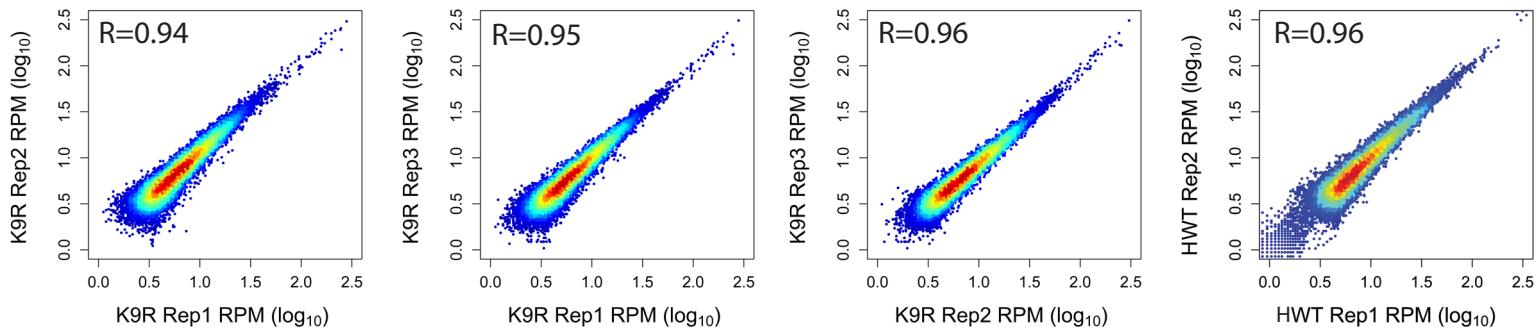


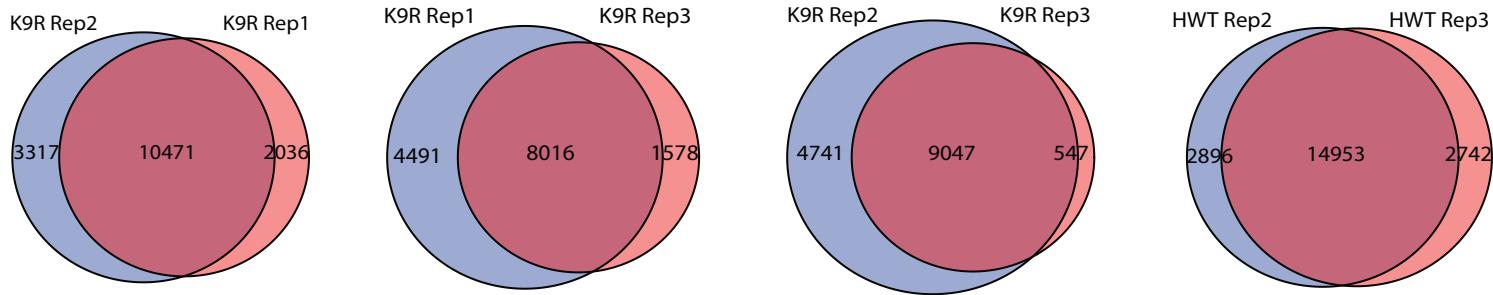
Transgenic array length	0	12	4	8	12	0	12	4	8	12
Endogenous histones	+	-	+	+	+	+	+	+	+	+
Genotype	HWT					K9R				



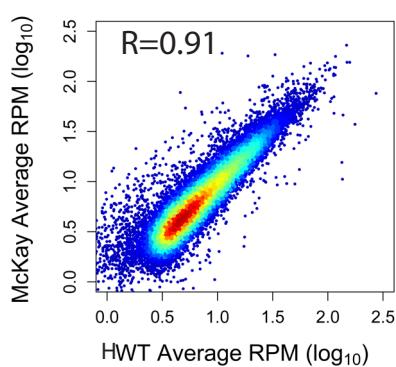
A



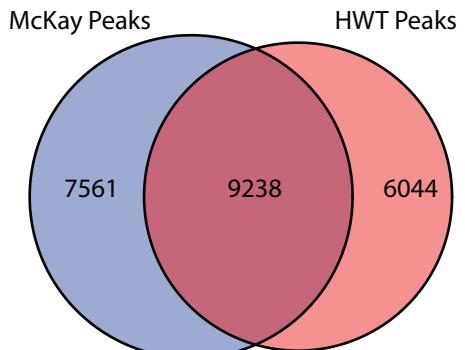
B



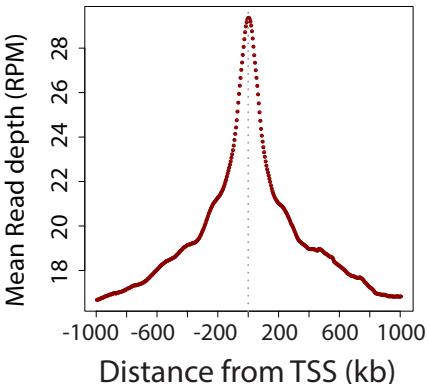
C



D

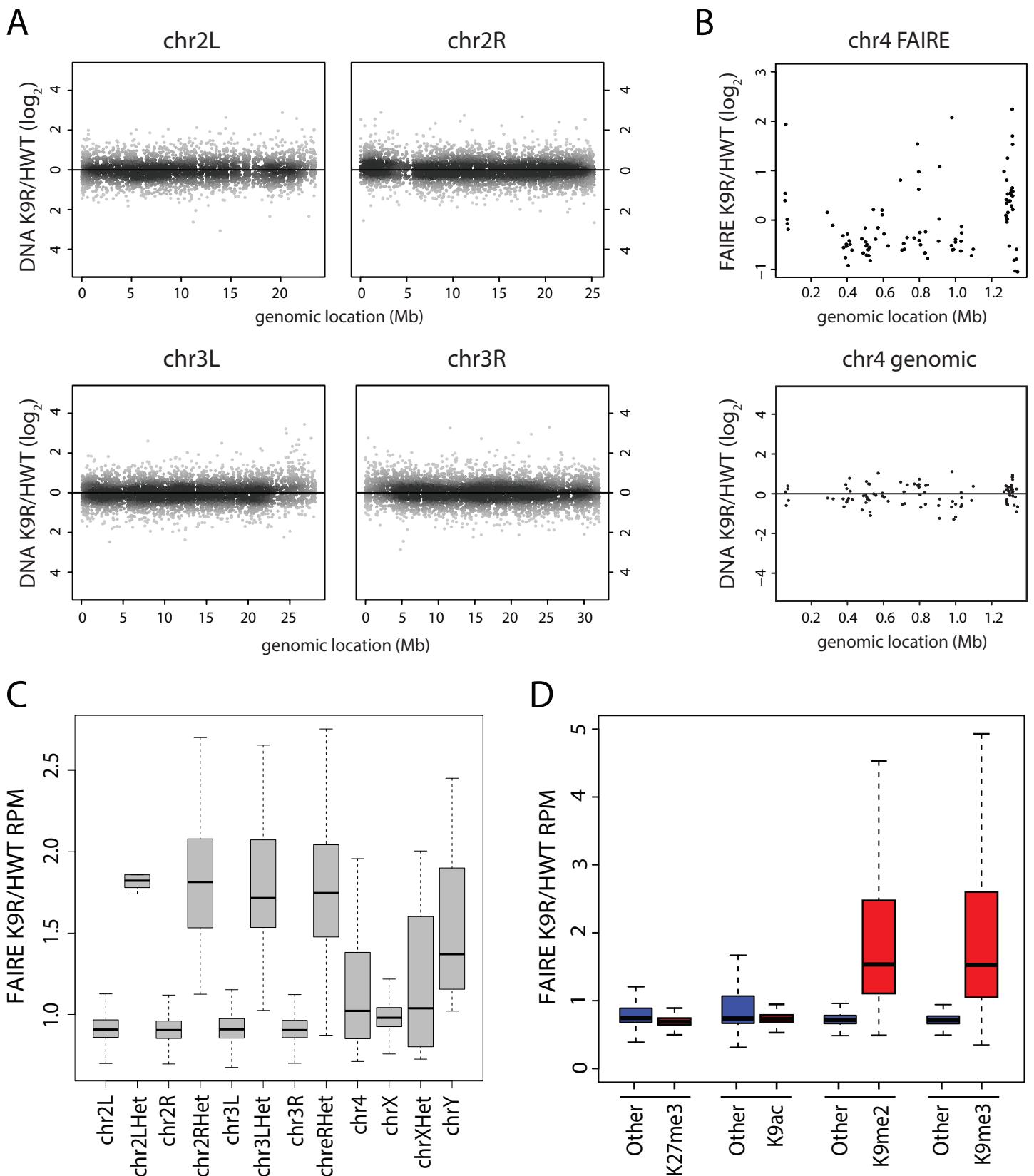


E

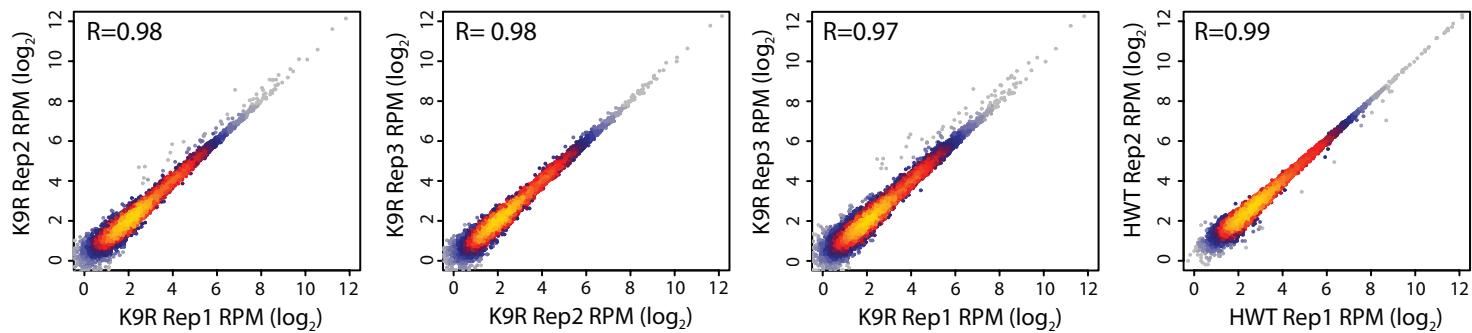


F

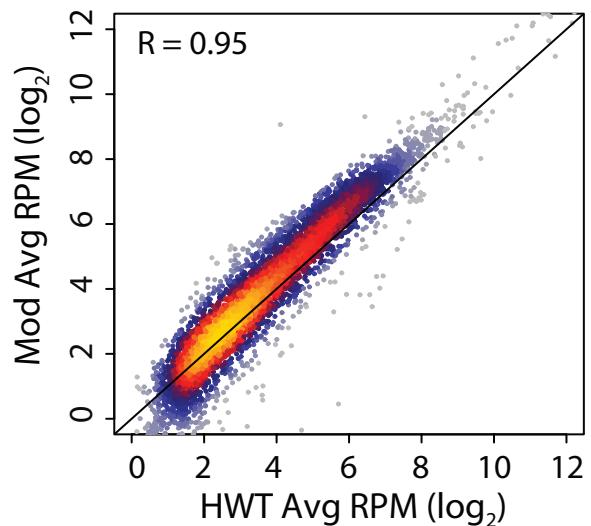
Replicate	Unique(%)	Multiple (%)
HWT 1	80.0	20.0
HWT 2	80.0	20.0
K9R 1	68.0	32.0
K9R 2	69.8	30.2
K9R 3	71.0	28.9



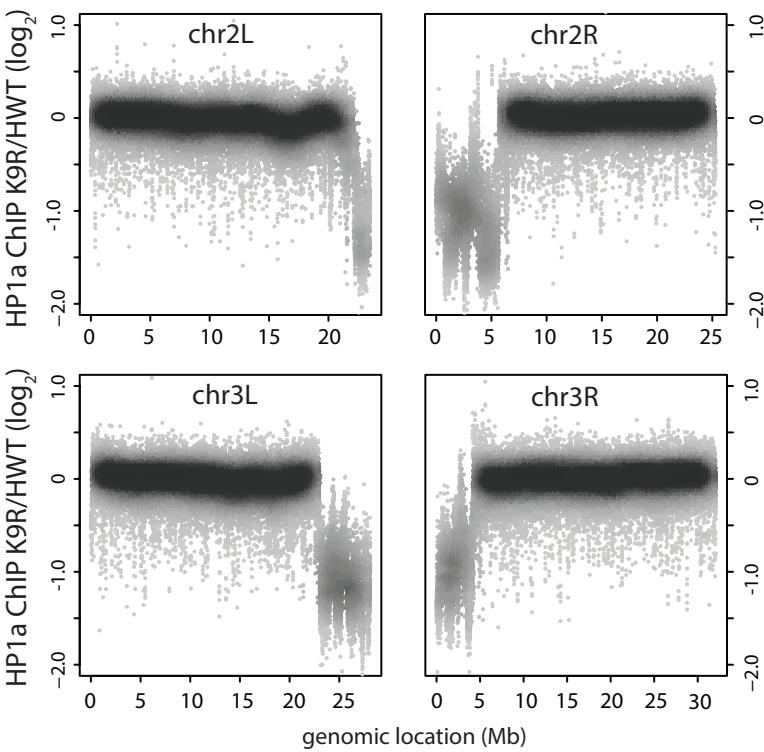
A



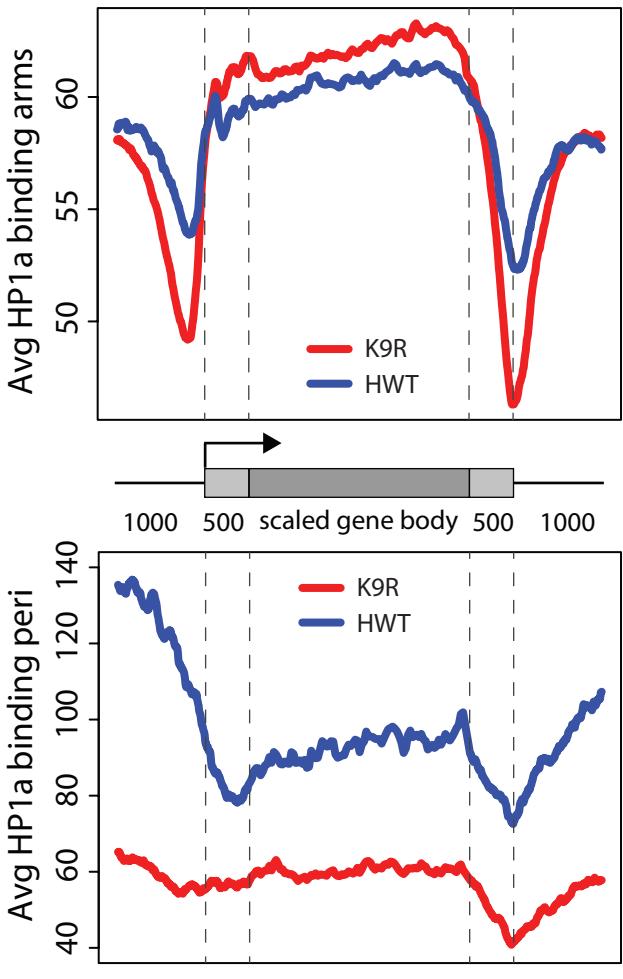
B



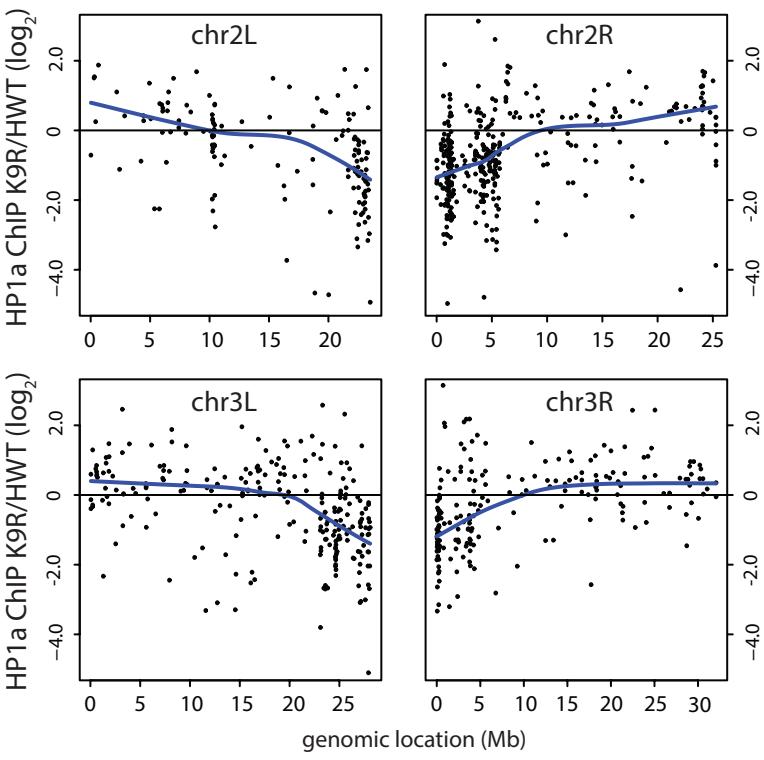
C

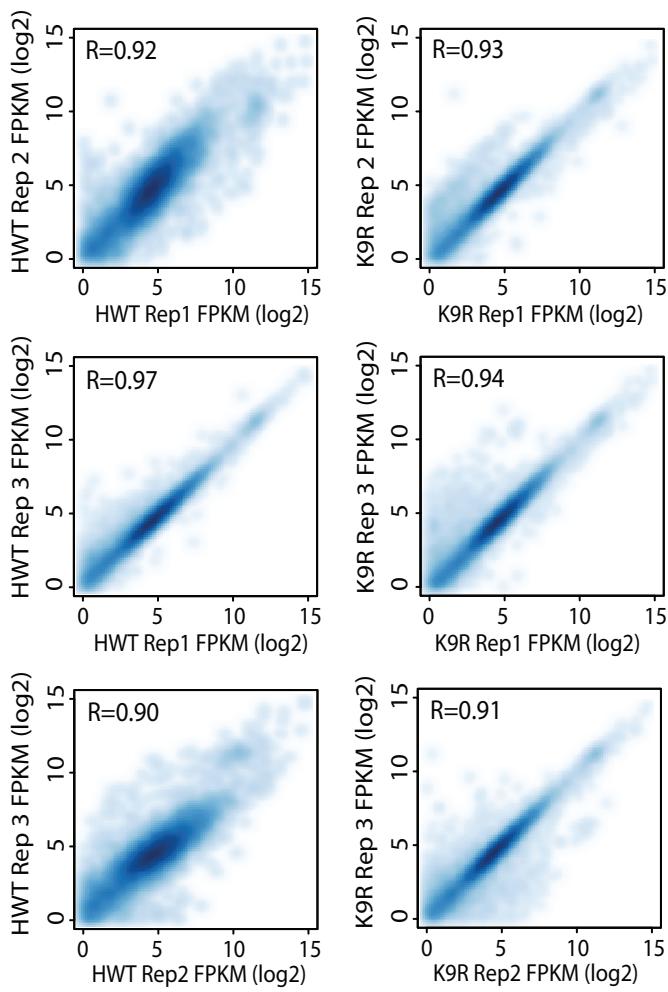
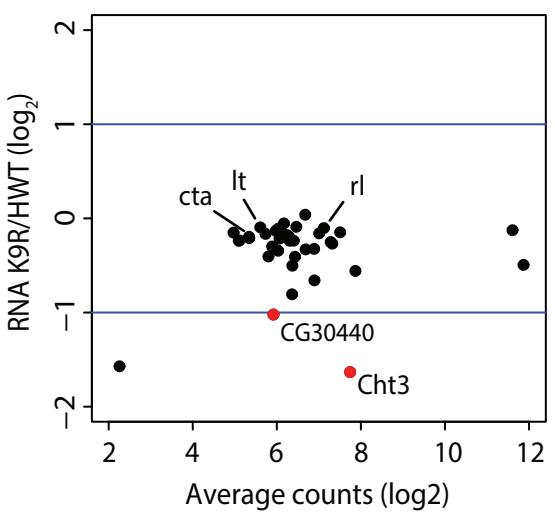
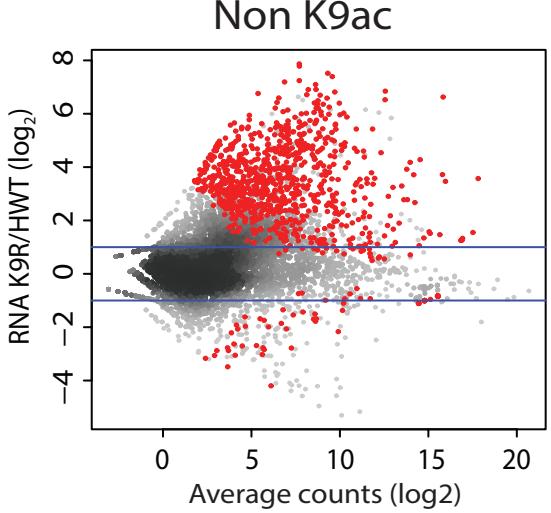
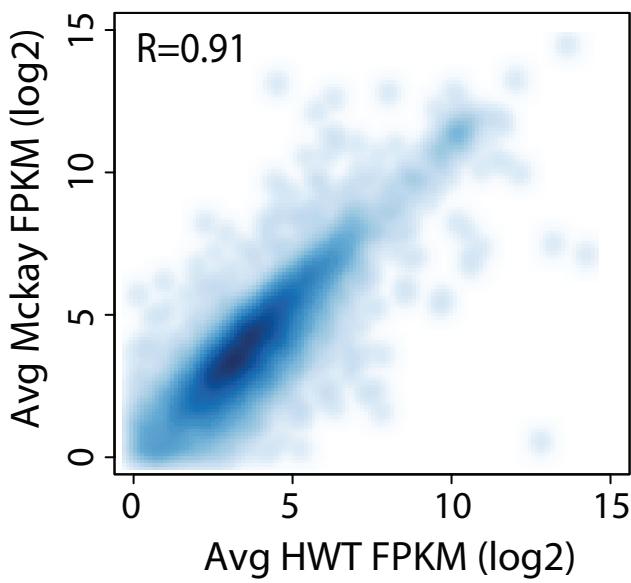
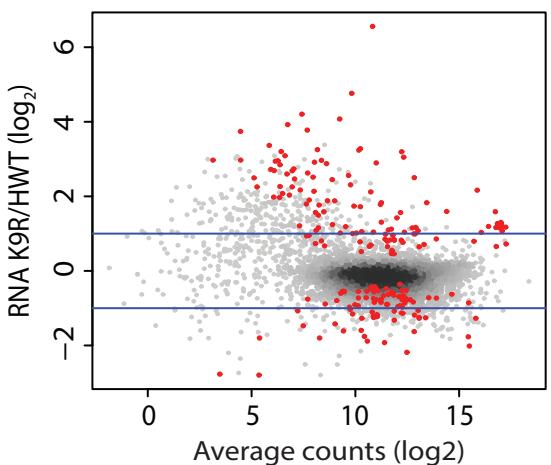


D

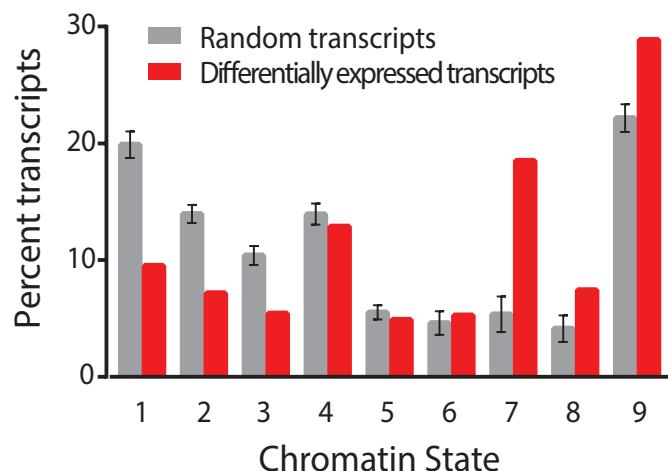


E



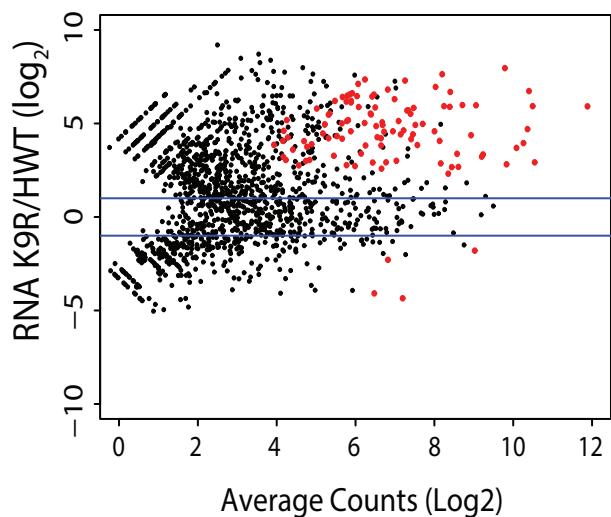
A**C****D****B****K9ac**

A

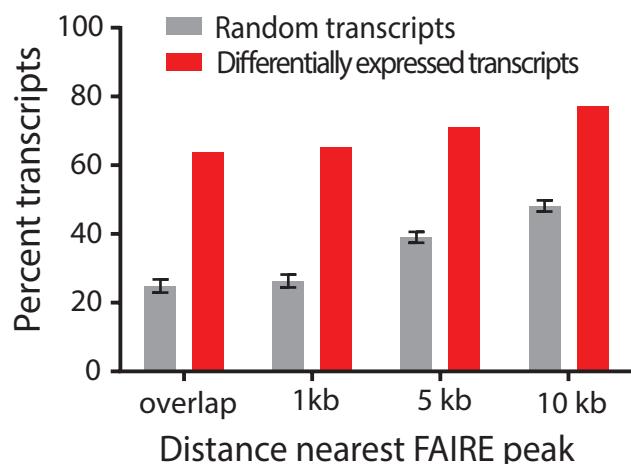


- 1) H3K4me2/3, H3K9ac 4) H3K36me 7) H3K9me2/3
 2) H3K36me3 5) H4K16ac, H3K36me3 8) Low H3K9me2/3
 3) H3K18ac,H3K27ac 6) H3K27me2/3 9) H1, SUUR

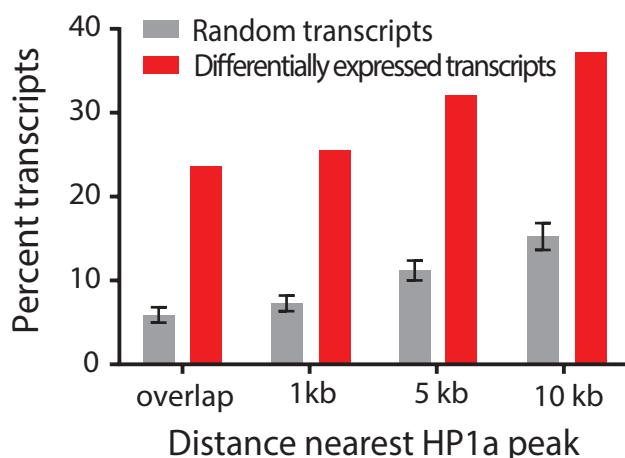
B



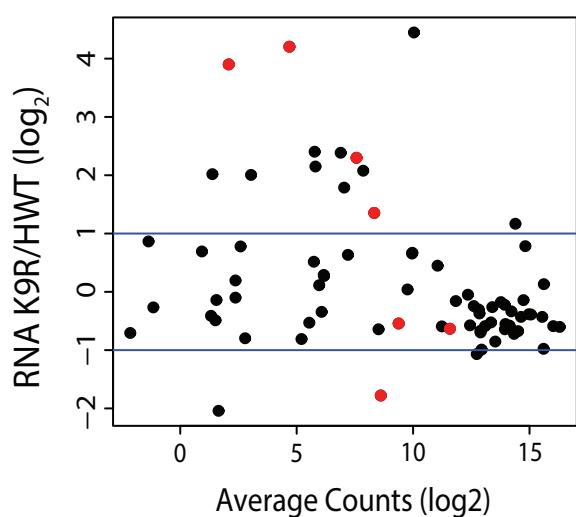
C



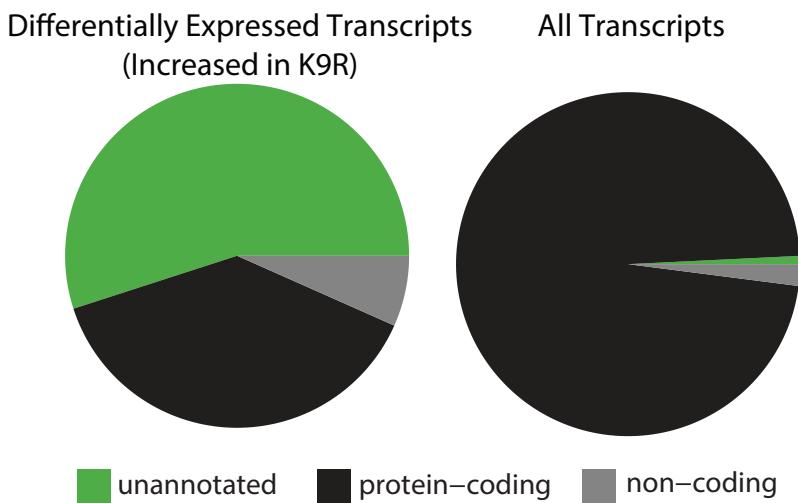
D

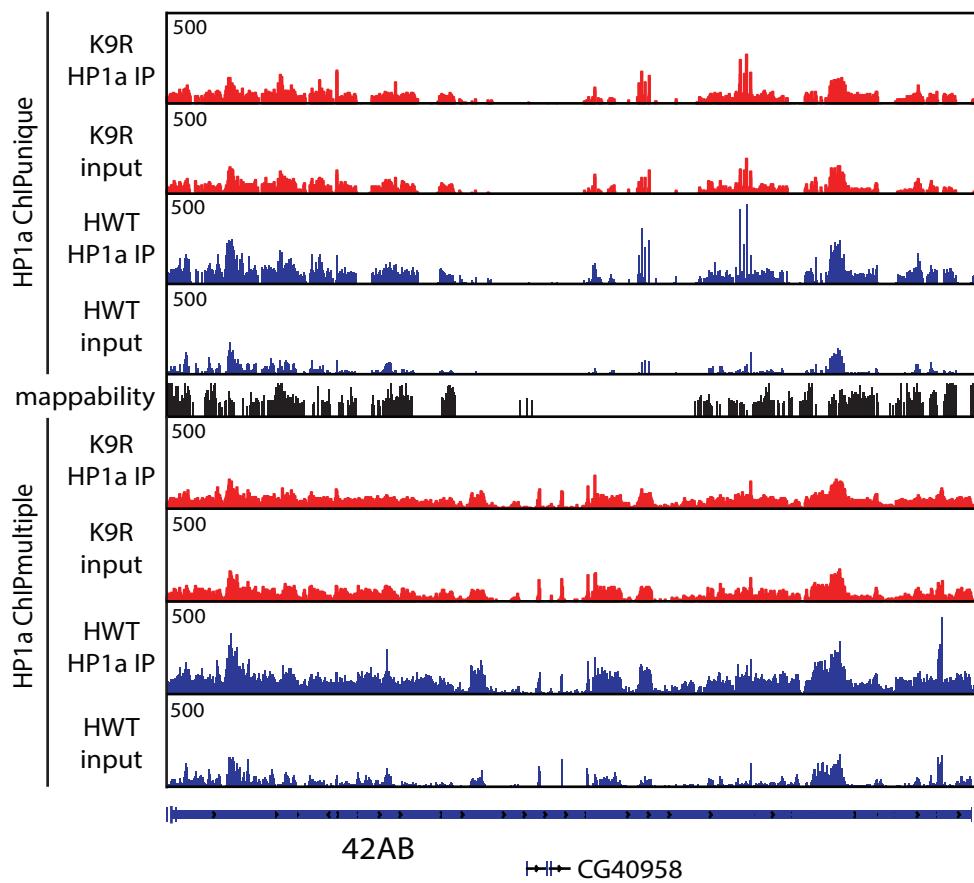
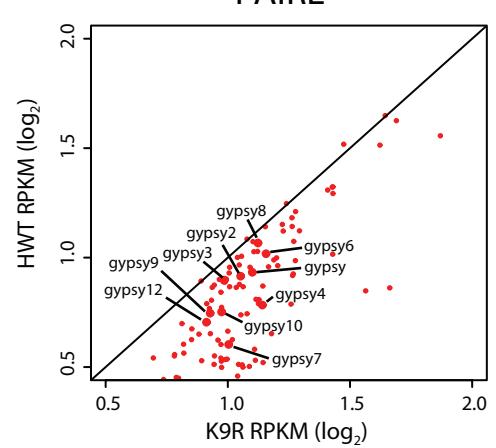
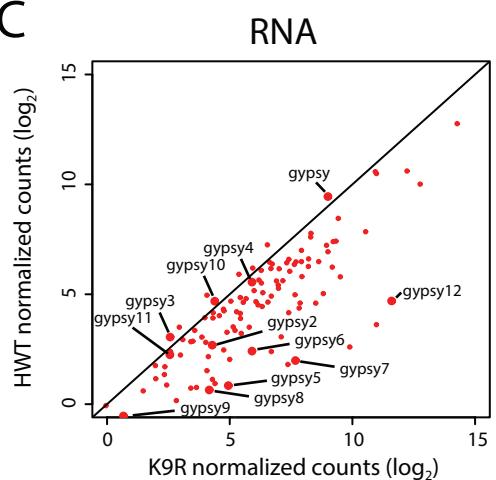
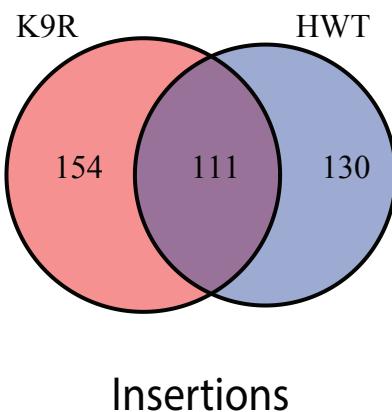
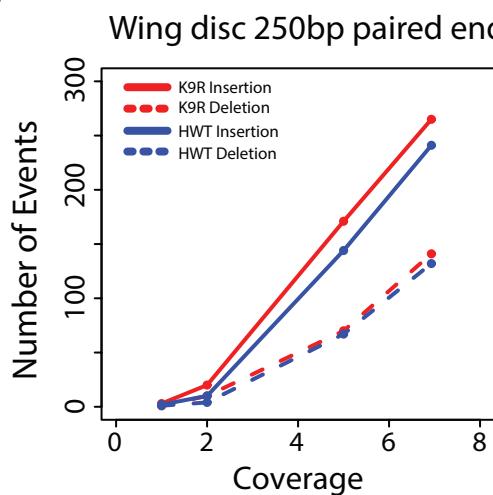


E

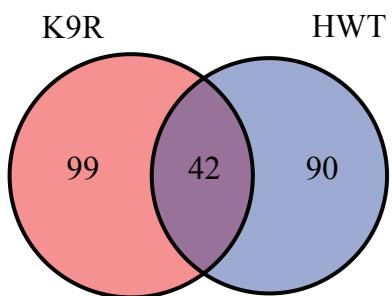


F



A**B****C****D**

Insertions



Deletions

E