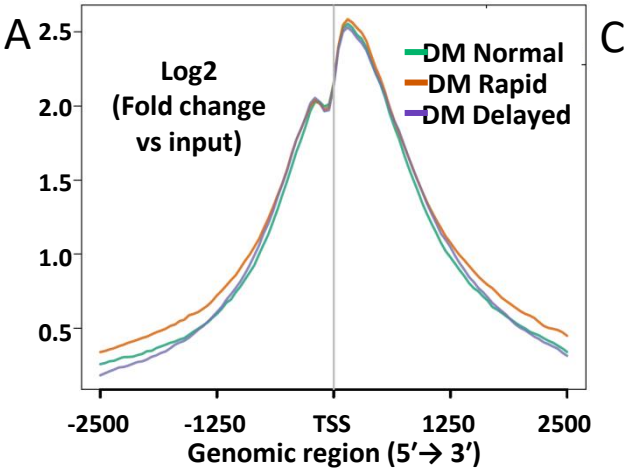
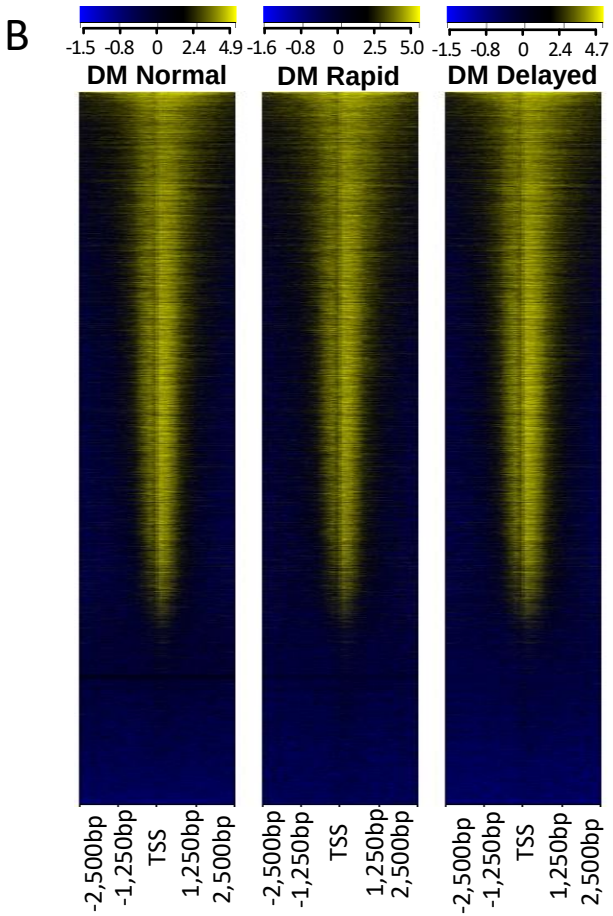
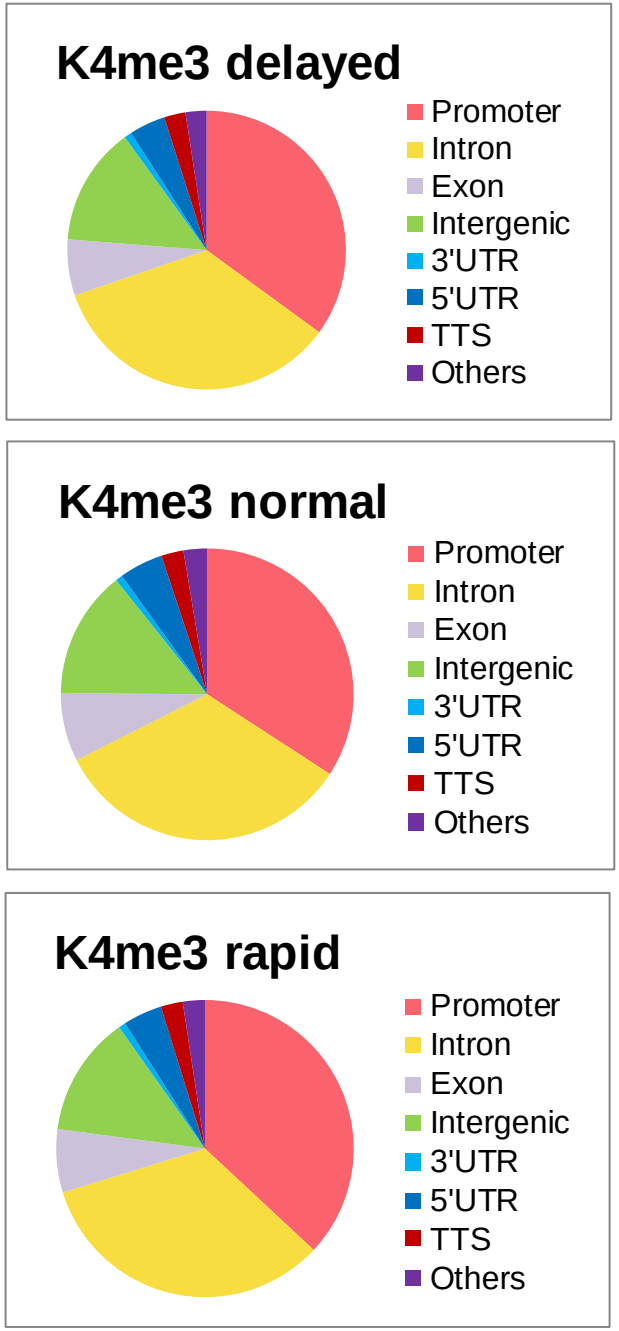

A
Log2
(Fold change
vs input)
DM Normal
DM Rapid
DM Delayed
2.5
2.0
1.5
1.0
0.5
-2500
-1250
TSS
1250
2500
Genomic region (5'→ 3')


B
-1.5 -0.8 0 2.4 4.9
DM Normal
-1.6 -0.8 0 2.5 5.0
DM Rapid
-1.5 -0.8 0 2.4 4.7
DM Delayed
-2,500bp
-1,250bp
TSS
1,250bp
2,500bp


C
K4me3 delayed
K4me3 normal
K4me3 rapid
Promoter
Intron
Exon
Intergenic
3'UTR
5'UTR
TTS
Others