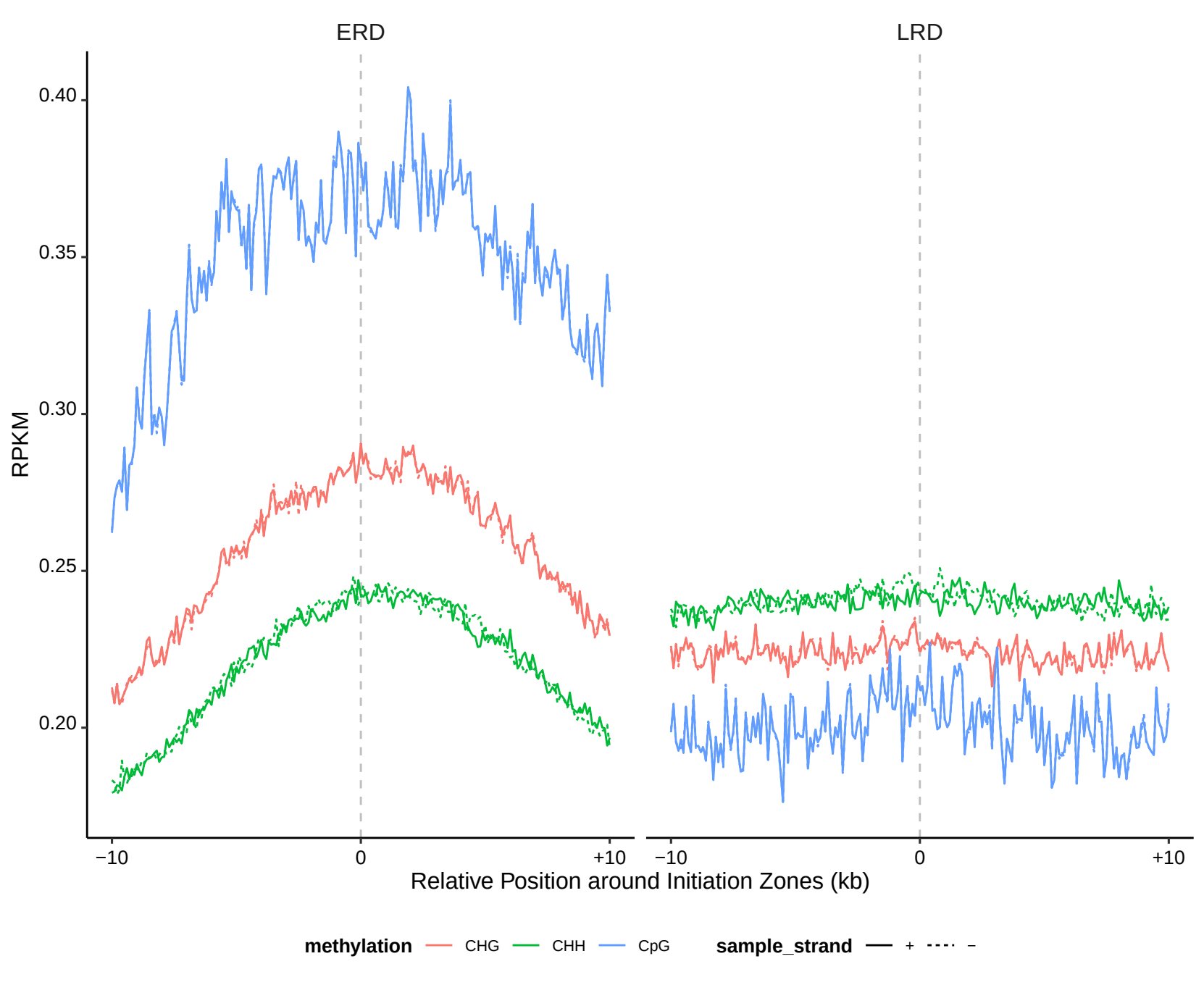
ERD
LRD
RPKM
0.40
0.35
0.30
0.25
0.20
−10
0
+10
Relative Position around Initiation Zones (kb)
methylation
CHG
CHH
CpG
sample_strand
+
−