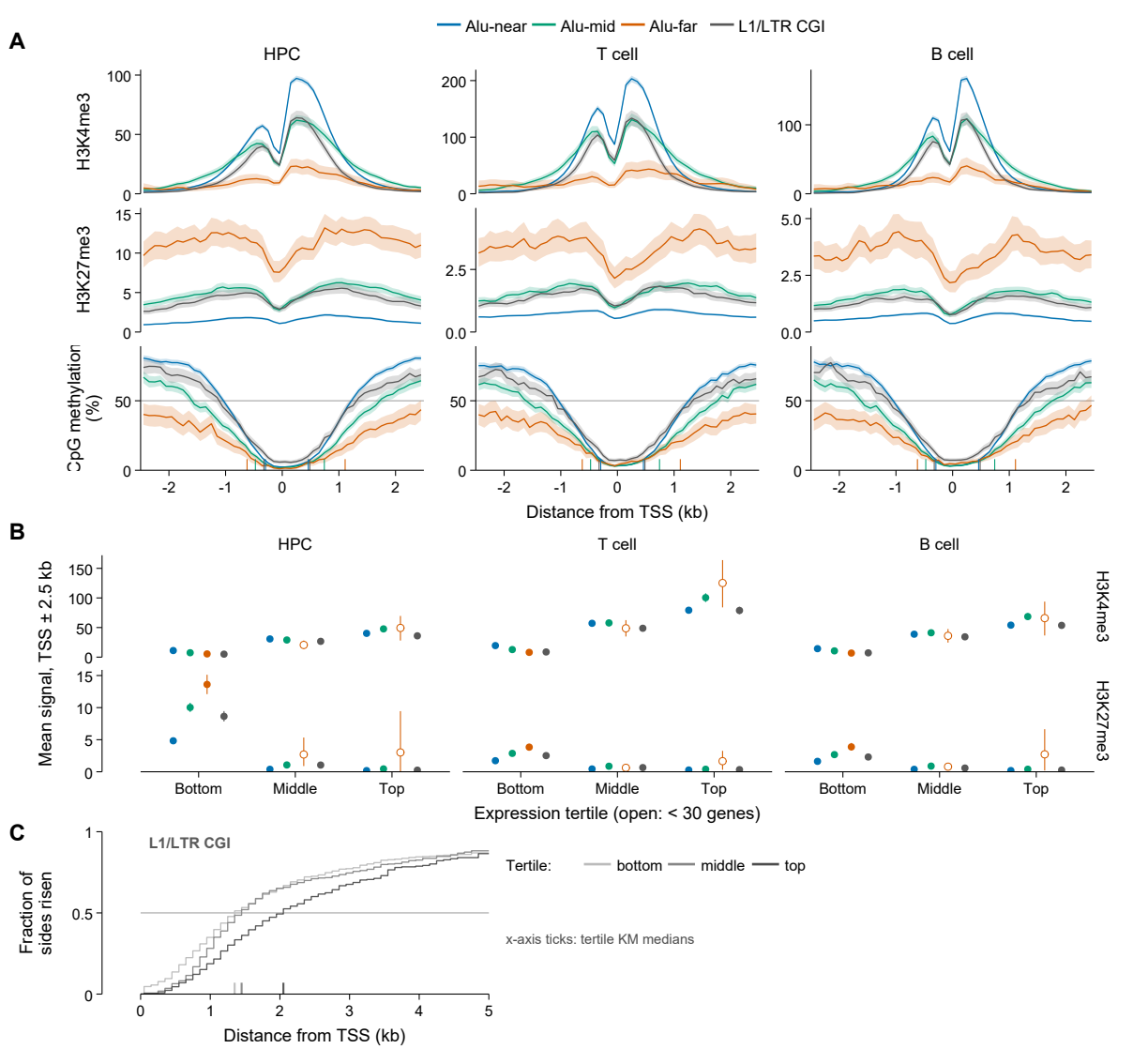

A
Alu-near
Alu-mid
Alu-far
L1/LTR CGI
HPC
T cell
B cell
H3K4me3
H3K27me3
CpG methylation (%)
Distance from TSS (kb)
B
Mean signal, TSS ± 2.5 kb
H3K4me3
H3K27me3
Bottom
Middle
Top
Expression tertile (open: < 30 genes)
C
L1/LTR CGI
Fraction of sides risen
Tertile:
bottom
middle
top
x-axis ticks: tertile KM medians
Distance from TSS (kb)