



Supplemental Figure S22 Proportions of TSSs harboring regulatory variants within TSS \pm 1kb in different TSS groups, based on variants with higher thresholds of derived allele frequency (DAF). (A) Results based on variants with DAF ≥ 0.1 . (B) Results based on variants with DAF ≥ 0.5 . Above the bars are the numbers of TSSs with regulatory variants. Note that for the results based on variants with DAF ≥ 0.5 , the numbers of regulatory variants found in the TSS groups/subgroups are very small, so the changing trends shown in the some panels for different transcript types may not accurately reflect actual trends. LCLs, lymphoblastoid cell lines.