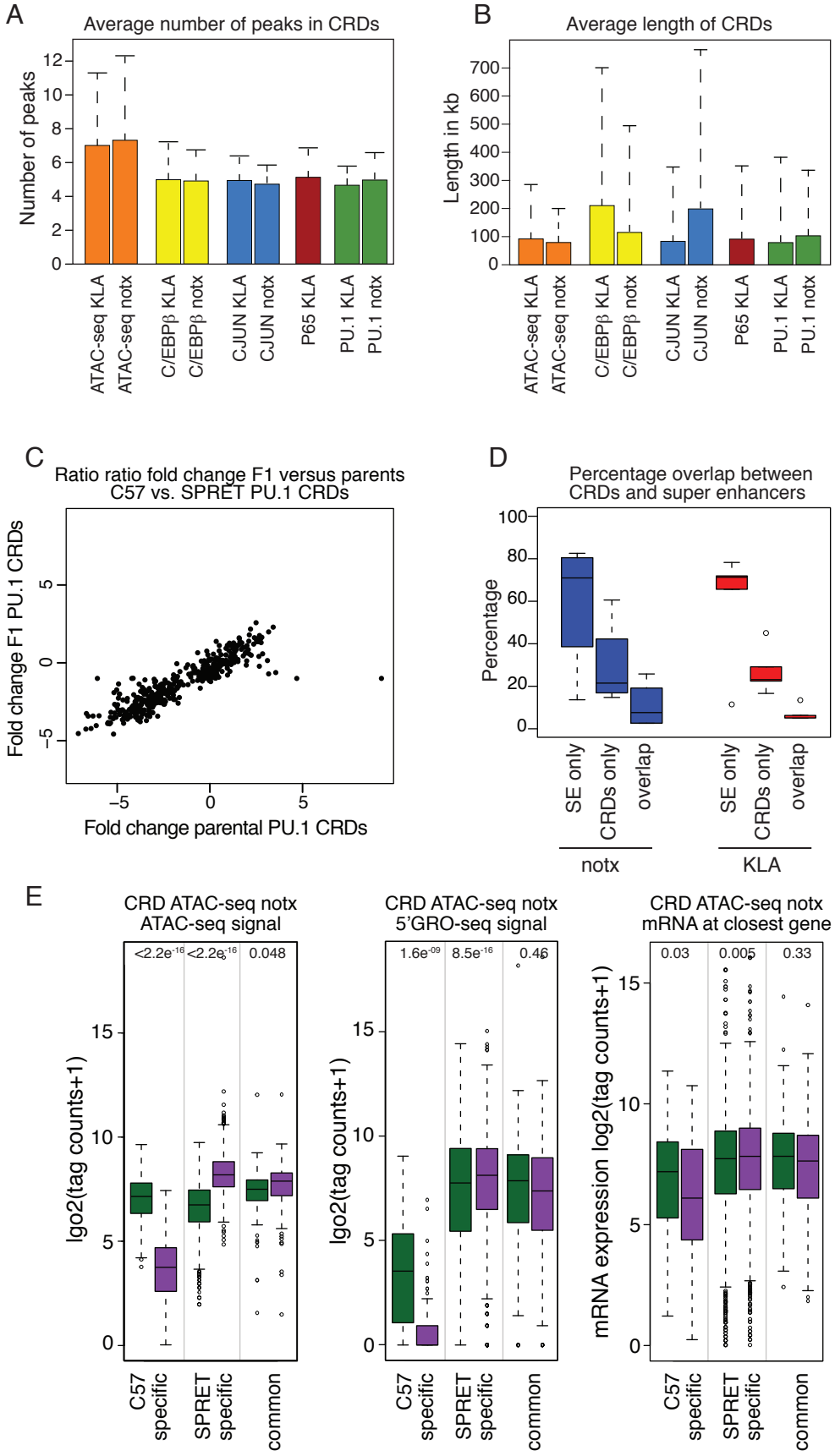
A
Average number of peaks in CRDs
Number of peaks
ATAC-seq KLA
ATAC-seq notx
C/EBPβ KLA
C/EBPβ notx
CJUN KLA
CJUN notx
P65 KLA
PU.1 KLA
PU.1 notx
B
Average length of CRDs
Length in kb
C
Ratio ratio fold change F1 versus parents
C57 vs. SPRET PU.1 CRDs
Fold change F1 PU.1 CRDs
Fold change parental PU.1 CRDs
D
Percentage overlap between
CRDs and super enhancers
Percentage
SE only
CRDs only
overlap
notx
KLA
E
CRD ATAC-seq notx
ATAC-seq signal
<2.2e-16
<2.2e-16
0.048
CRD ATAC-seq notx
5'GRO-seq signal
1.6e-09
8.5e-16
0.46
CRD ATAC-seq notx
mRNA at closest gene
0.03
0.005
0.33
lgo2(tag counts+1)
mRNA expression log2(tag counts+1)
C57 specific
SPRET specific
common

Figure S5