

Overlap of intronic deletions with regulatory features.

log₂ Fold Change (p-value)	Sudmant (Nature)	Zarrei	Abyzov
CTCF binding site	-0.18 (0.0072)	-0.14 (0.11018)	-0.81 (0.0001)
Enhancer	-0.14 (0.0304)	-0.45 (0.0006)	-0.47 (0.0045)