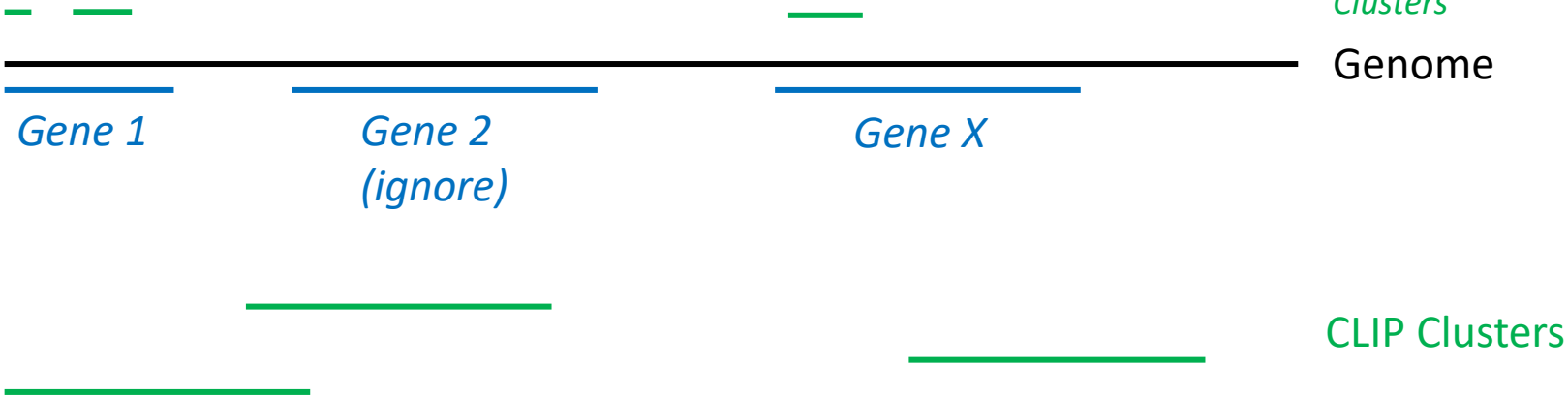


Find each gene overlapped by a CLIP cluster

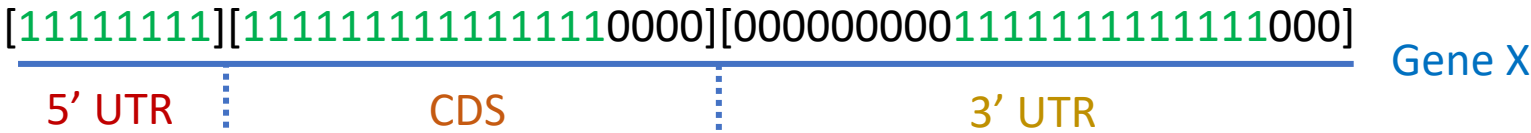


Optional

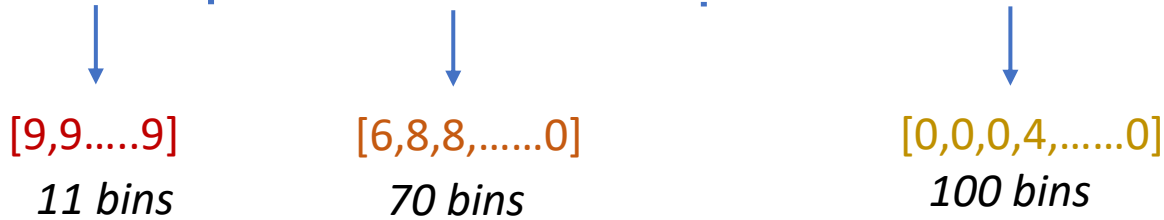
Select certain genes.

Filter clusters.

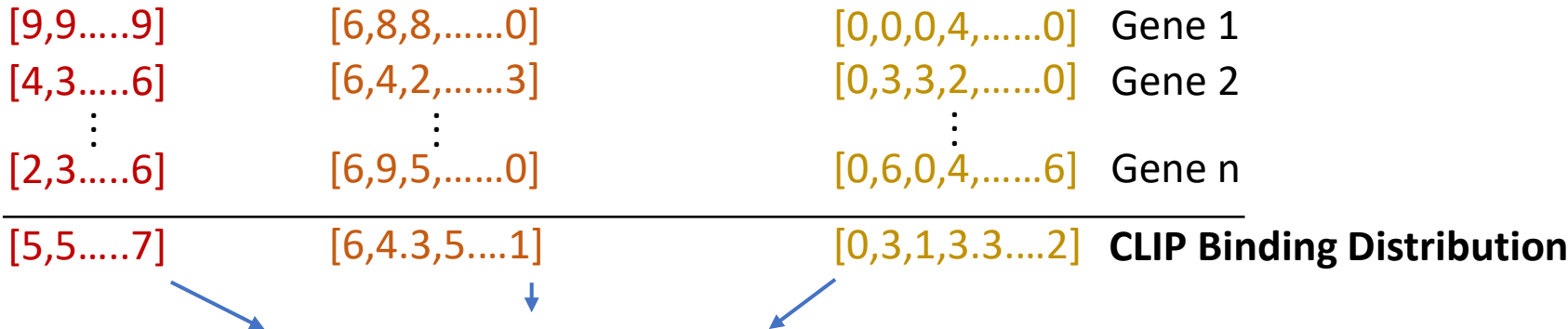
Nucleotide-Level Overlap (By region)



Binning



Averaging Gene Lists (all overlapped genes)



Scale the contribution of each gene by the number of reads aligned to that gene.