

All the genes in the
annotation file

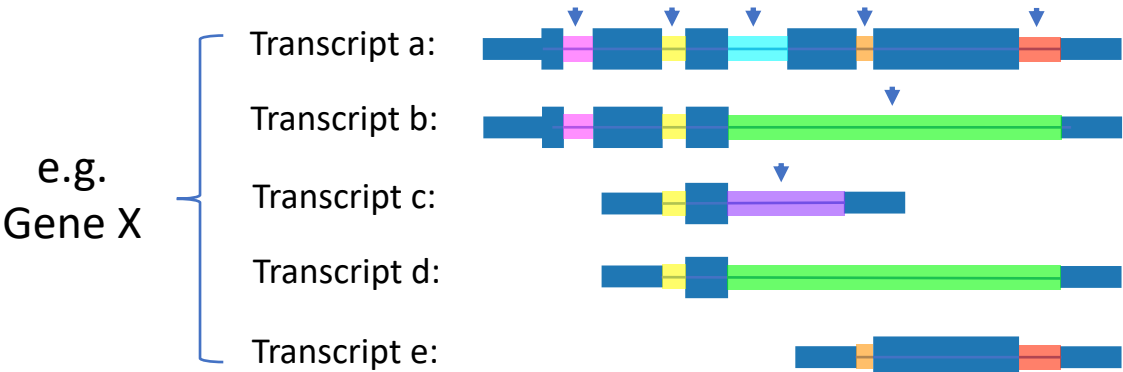


Select the genes that have
expression in ≥ 1 sample(s)

1	0	0	3	0	1	0
0	0	0	0	0	0	0
5	0	10	0	0	0	2
0	0	0	0	0	0	0



Get the set of unique introns from
all the transcripts of each gene



Keep the introns with length > 30



Total number: 335,436