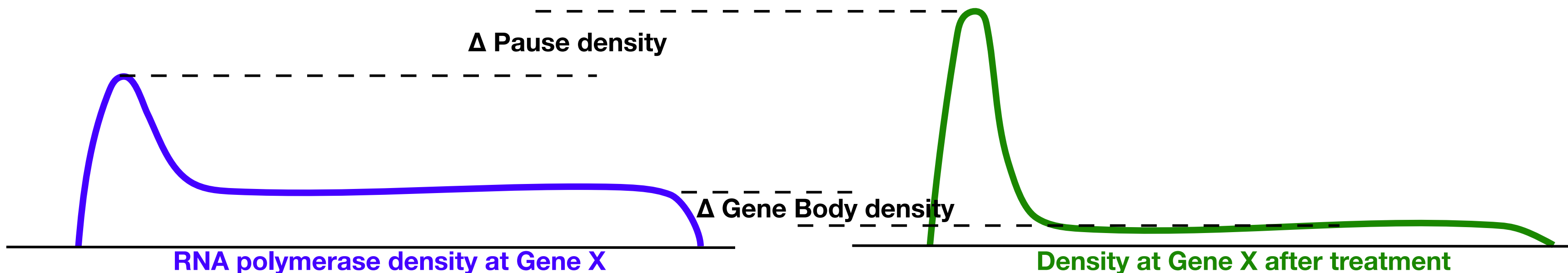
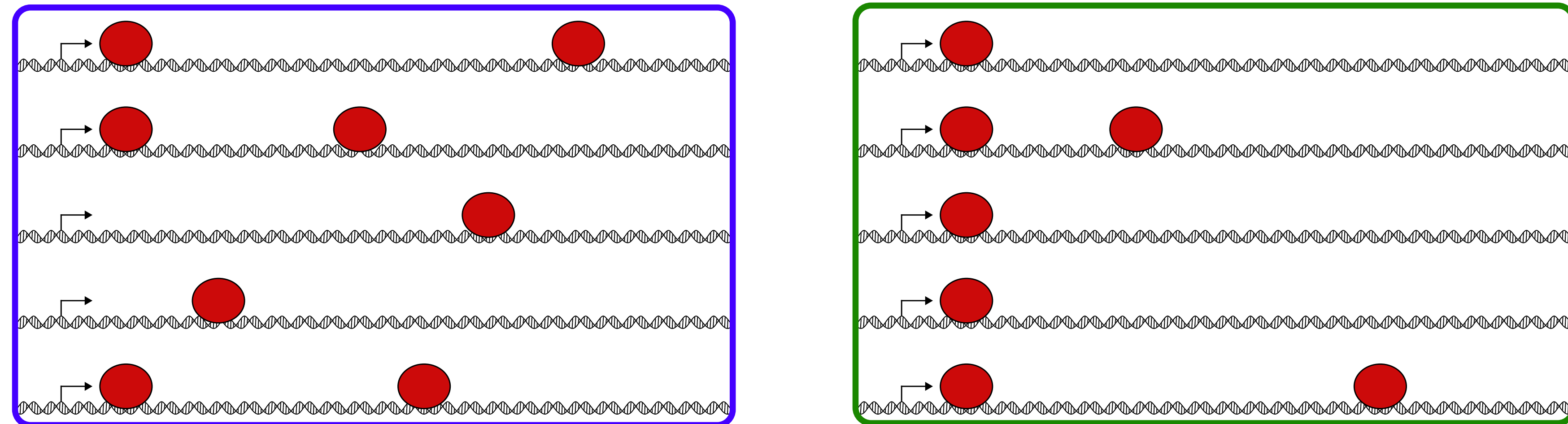
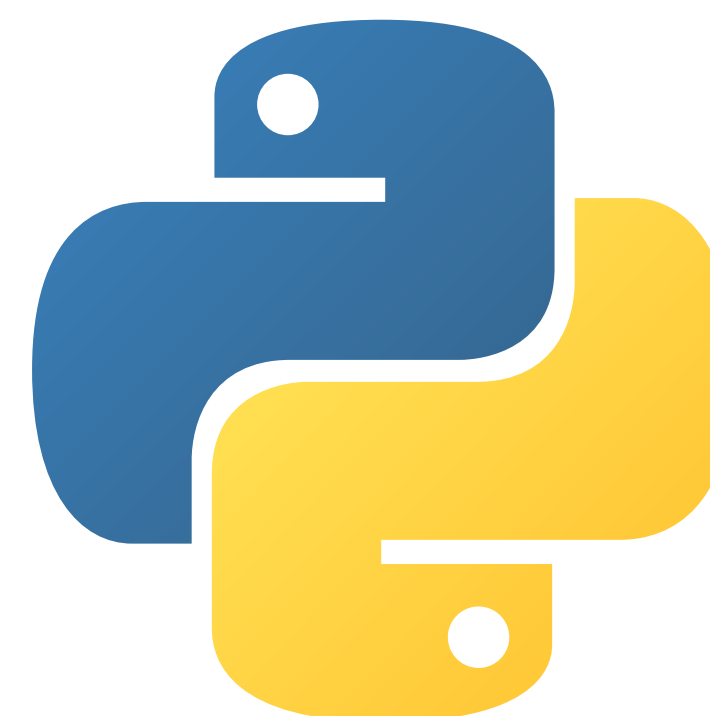


PRO-seq measures engaged **RNA Polymerase** density for all genes in **control** and **treatment** conditions



Input:
PRO-seq densities
in pause and gene
body regions



python3 compartmentModel

Output:
Rates of initiation,
pause release, and
premature termination
for all genes

