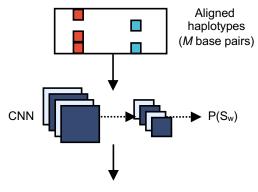
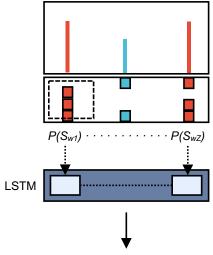
Simulated genomic windows (*N* base pairs)



Sliding window analysis on full simulated non-recombining genomes



Detecting and quantifying strength of positive selection, P(Selection) and (1 + s)