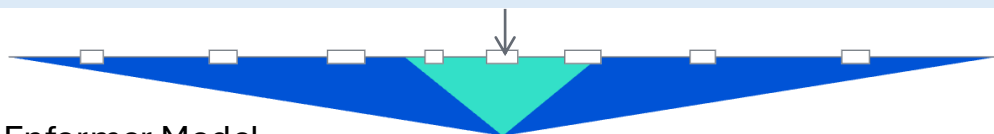


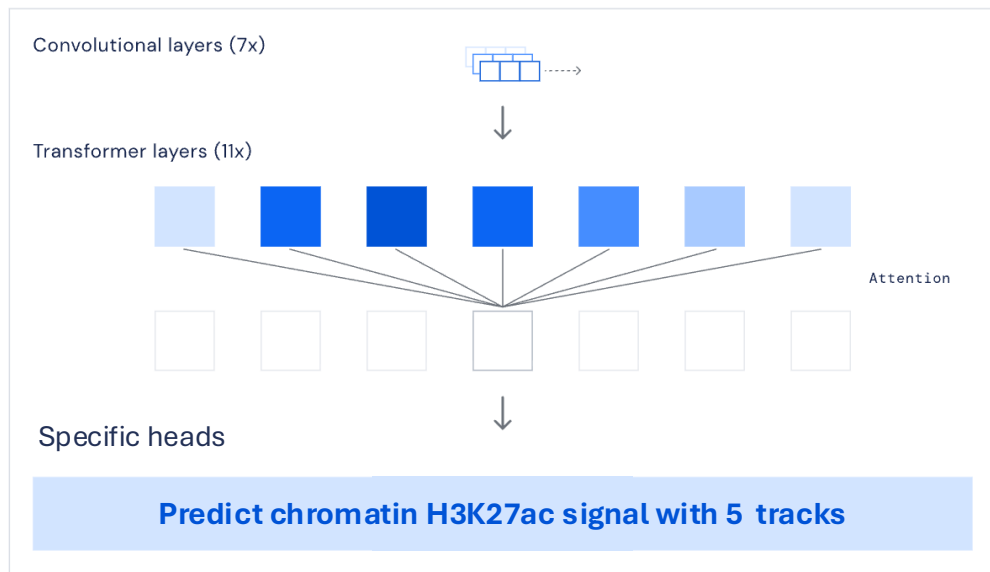
196,608 base pairs

Input: DNA sequence

CTAGTTCTATAGTTTTCTGAGGGTGT.....GCTTTTTTCCTTTTGAAGTGCCTAAAA



## Enformer Model



Output: Genomics tracks

Chromatin signal prediction for different stages

