Transcriptome (T) with separator

t_1 t_2 t_3 t_4 t_5 t_6

Maximum Mappable Prefix — **MMP$_i$**

Next Informative Position — **NIP(MMP$_i$)**

k-mer — **k$_i$**

ATTGAGTATA - - - - - - -

Read

Hash Table — h

k$_i$	[b, e)

Suffix Array(T)

b   b'   e'   e