

Alignment Input



Condensed Repr. ($\times n$ threads)



Estimated RNA Mod. Location

Alignment Operation

PHRED Score Quality Filter

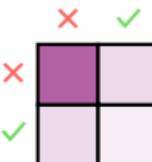
Filter

Process 3' to 5'

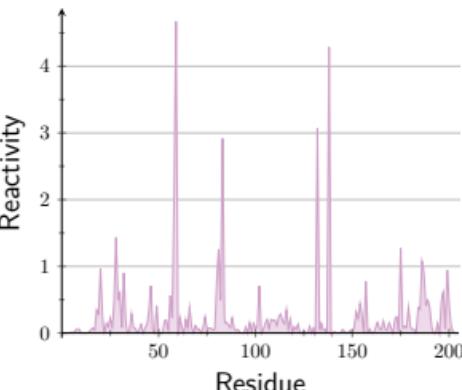
Per-Base Statistics

	A	C	G	T	del	ins	rm
A	dark purple						
C		dark purple					
G			dark purple				
T				dark purple			

Pairwise Statistics



Normalized Profiles



Mutual Information

