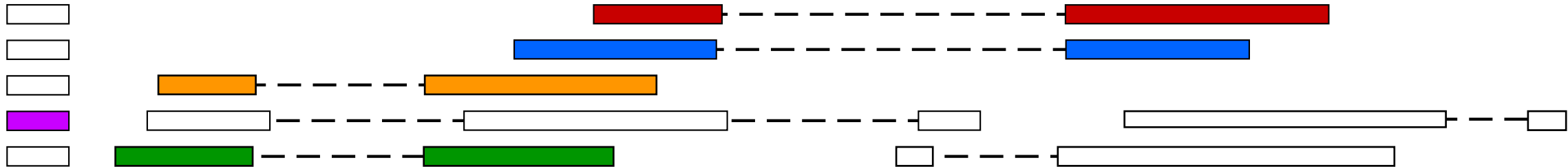


(Not
Aligned)

Alignment Qualities:
 e errors in n bases
 m reads are not aligned



Reference Genome