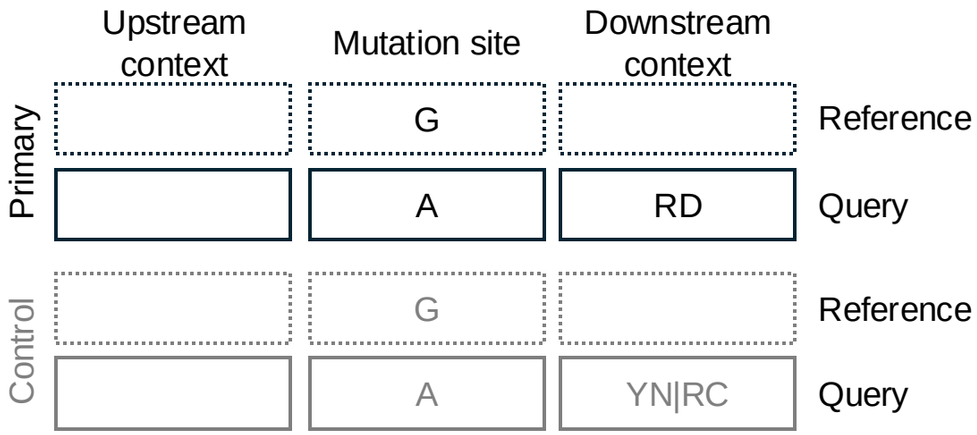
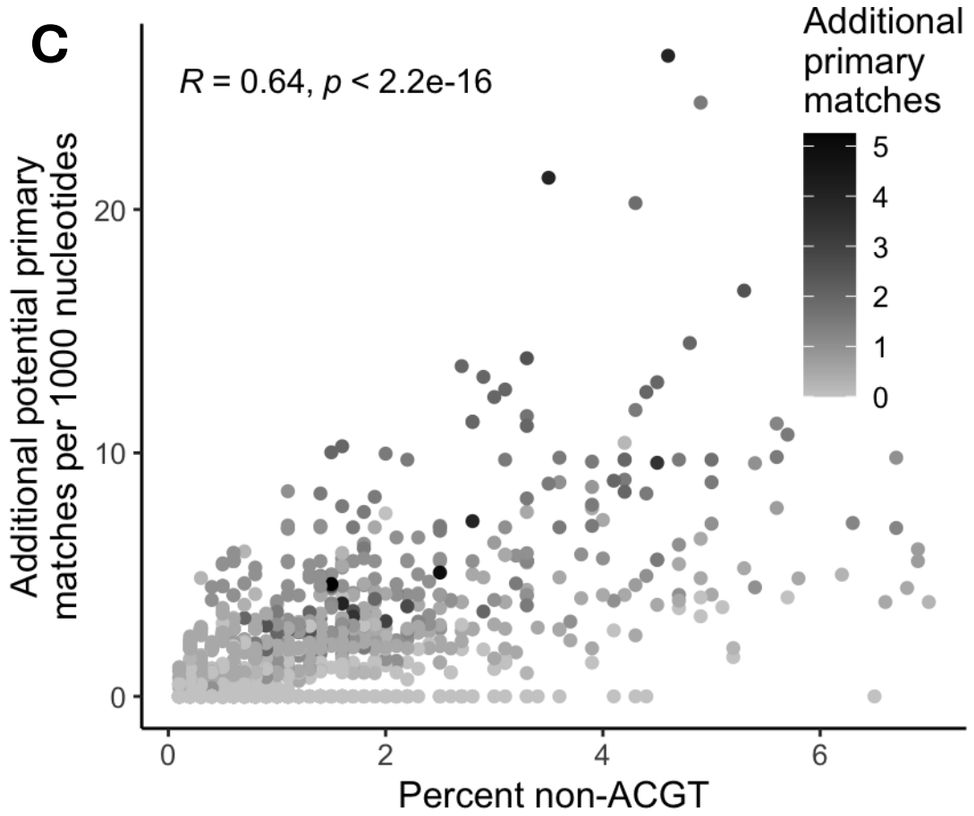
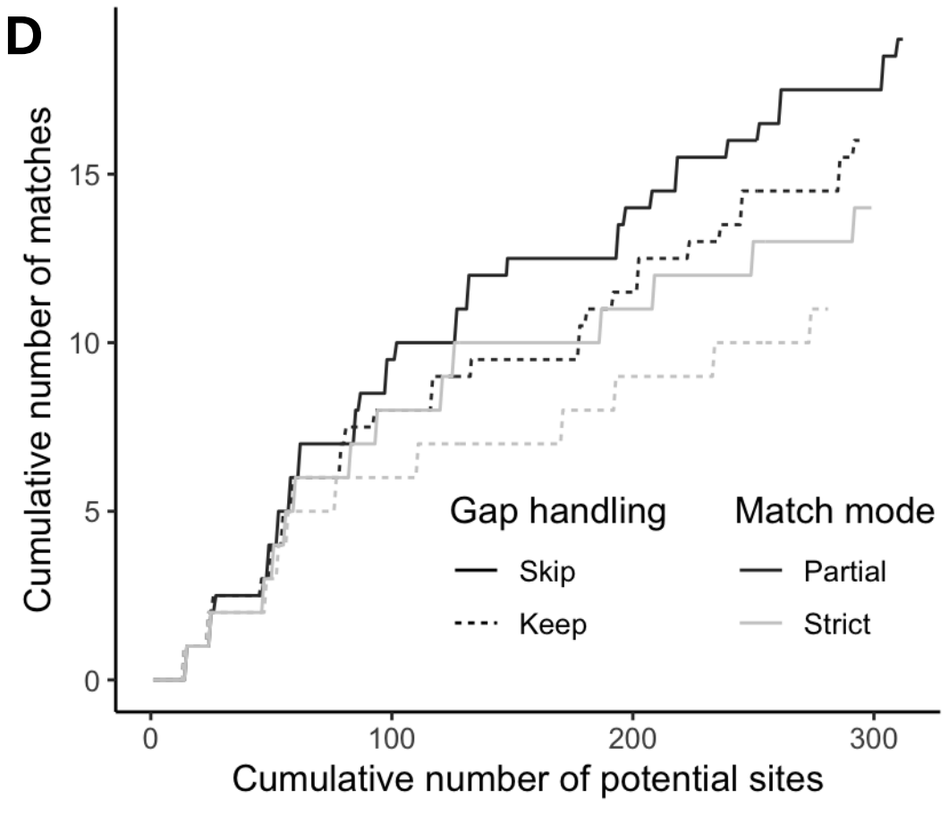
A



B

		Potential primary match			Primary match score		
ID	Seq	2.0	strict	partial	2.0	strict	partial
ref	GATC						
q1	**AGT**C	1	1	1	1	1	1
q2	T**GT**C	1	1	1	0	0	0
q3	**A**GCC	0	0	0	0	0	0
q4	-**GT**C	1	0	0	0	0	0
q5	**A**-**GT**	0	1	1	0	1	1
q6	**RGT**C	1	0	1	0	0	0.5
q7	**AGR**C	0	1	1	0	1	1
q8	**ANT**C	0	0	0.5	0	0	0.5
q9	**RNT**C	0	0	0.5	0	0	0.25

C



D