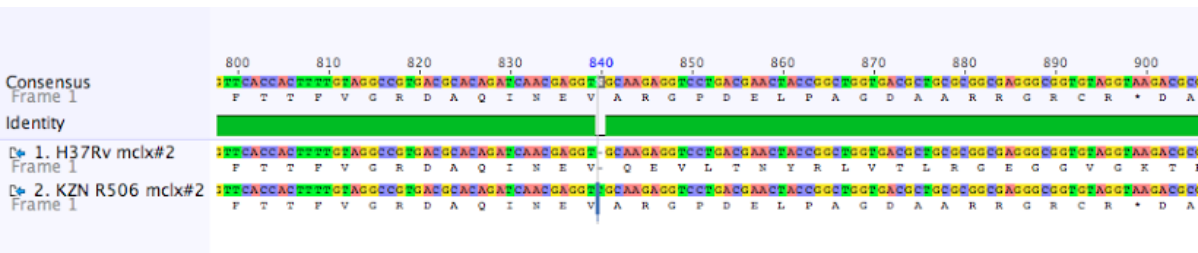
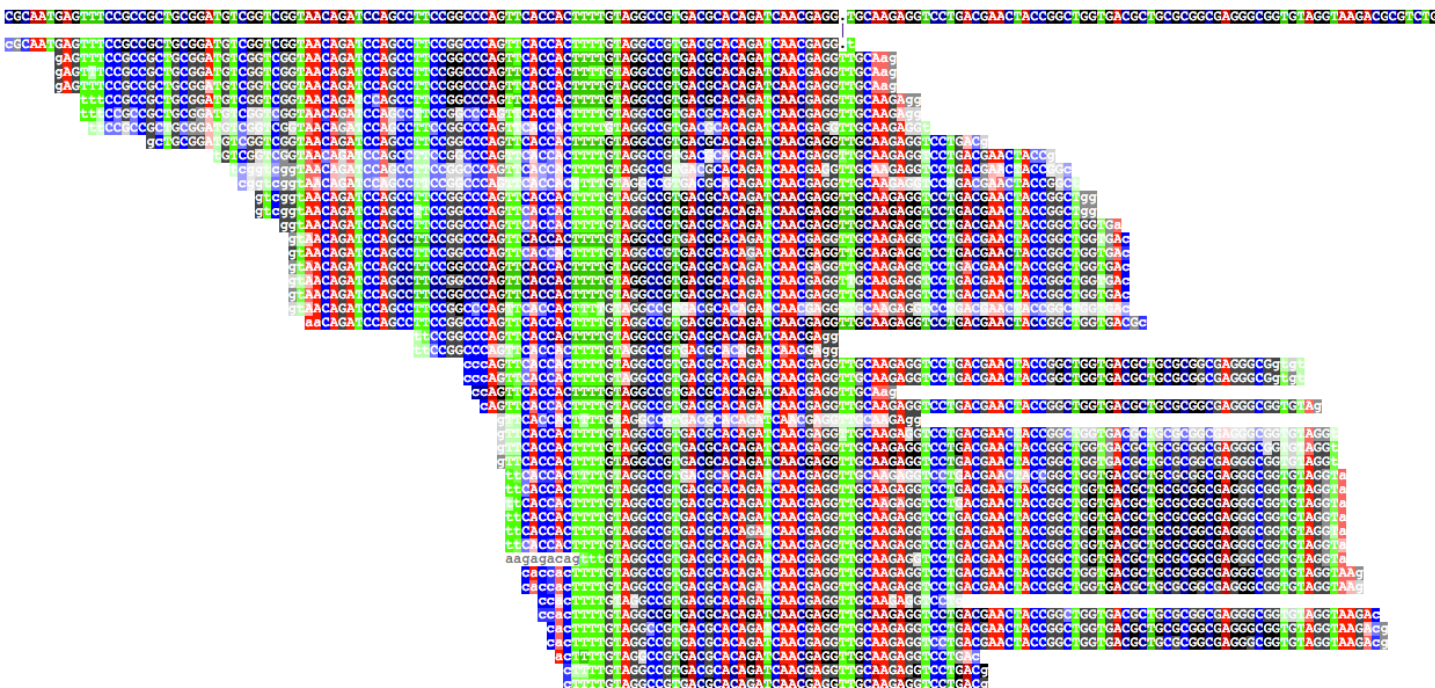


A**B**

Read alignment evidence...						
seq id	position	change	freq	score	reads	annotation genes product
NC_000962	1,527,449	1 → T	100.0%	213.6	61	coding (838/3480 nt) <i>Rv1358</i> transcriptional regulatory protein
Reads supporting (aligned to +/- strand): new base (27/34): ref base (0/0): total (27/34)						

**C**