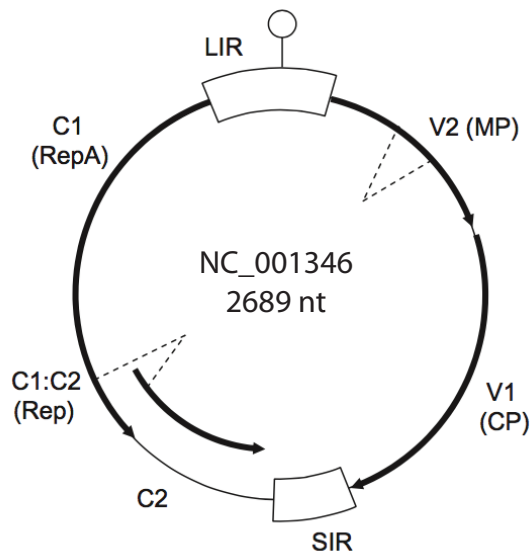


Overview of annotation pipeline for Maize Streak Virus

INPUT:

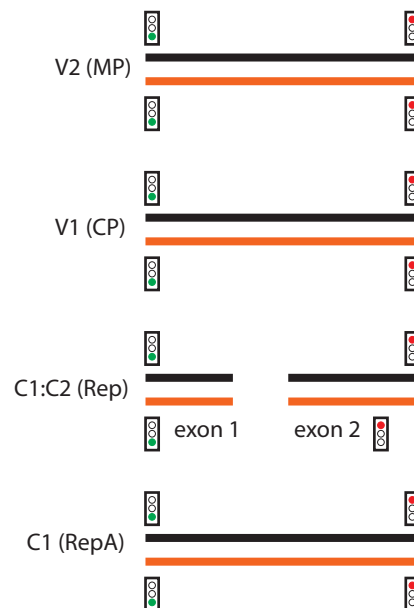
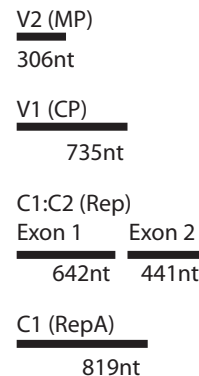
1. RefSeq annotation



2. Target sequence(s) to annotate:



Combine models into 'features', translate, find stop codons, and identify errors:



OUTPUT:

1. Tabular annotations of all features:

EU628610:MP:157-462
EU628610:CP:473-1207
EU628610:Rep:2530-1889,1796-1371
EU628610:RepA:2530-1712

2. List of all 'error codes':

EU628610:Rep:trc

3. Nucleotide and protein multiple alignments (optional)