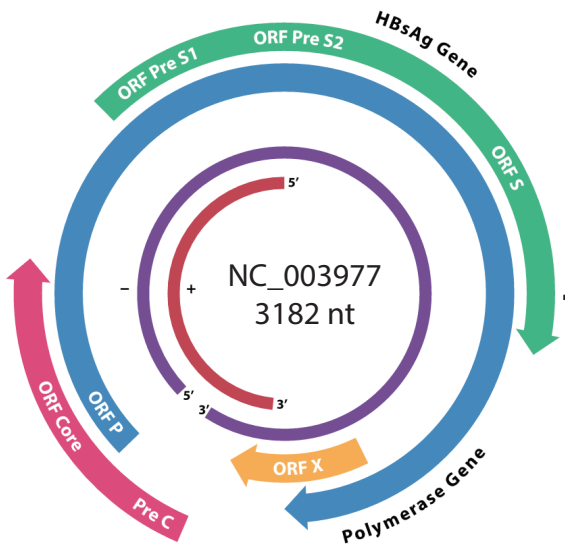


Overview of annotation pipeline for Hepatitis B Virus

INPUT:

1. RefSeq annotation



2. Target sequence(s) to annotate:



Build homology models and search duplicated genome targets:

