

- **Virus Identification**
(Genomad, VS2, CT3)

- fasta and GFF files



Secondary Structure Detection

Stem-Loop

- input: fasta and GFF
- motif pattern
- + StemLoop-Finder

Iteron

- input: fasta and GFF
- + CRUISE



output/

- |— stemloop_finder.log # Log file capturing the analysis process
- |— output.gff # GFF file with stem-loop annotations
- |— output.csv # (Optional) CSV file with detailed candidate data

output/

- |— cruise.log
- |— finaloutput.gff