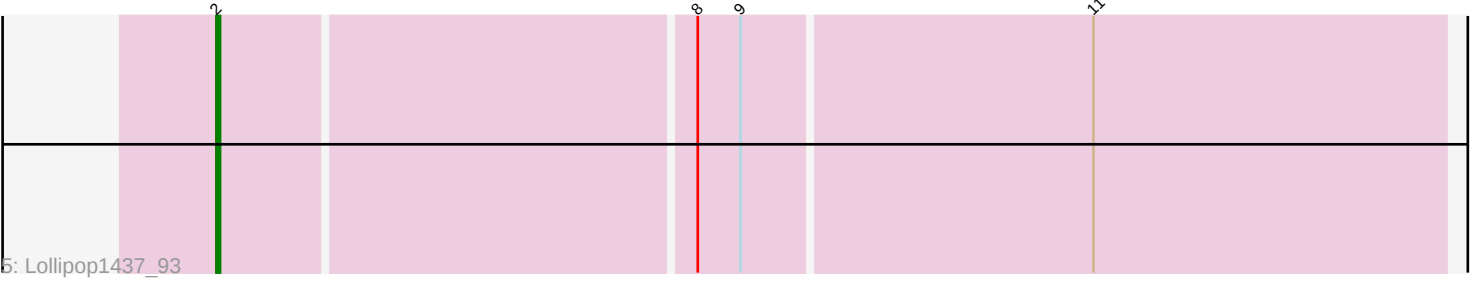
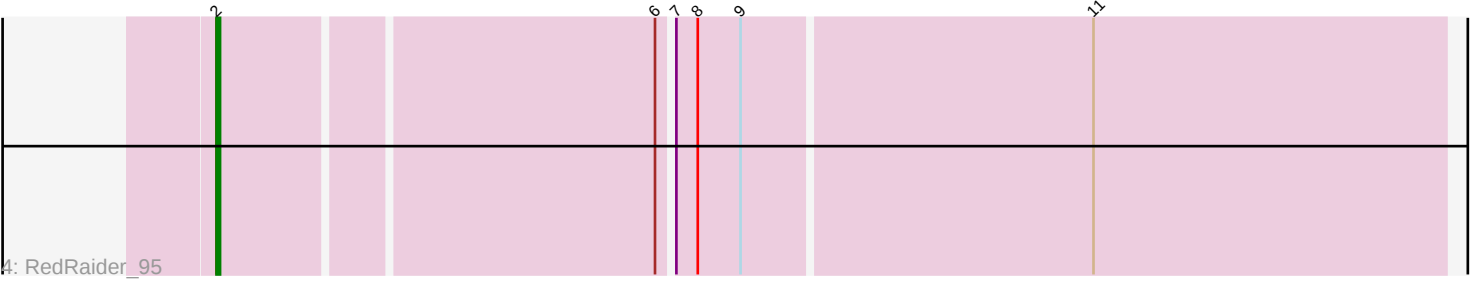
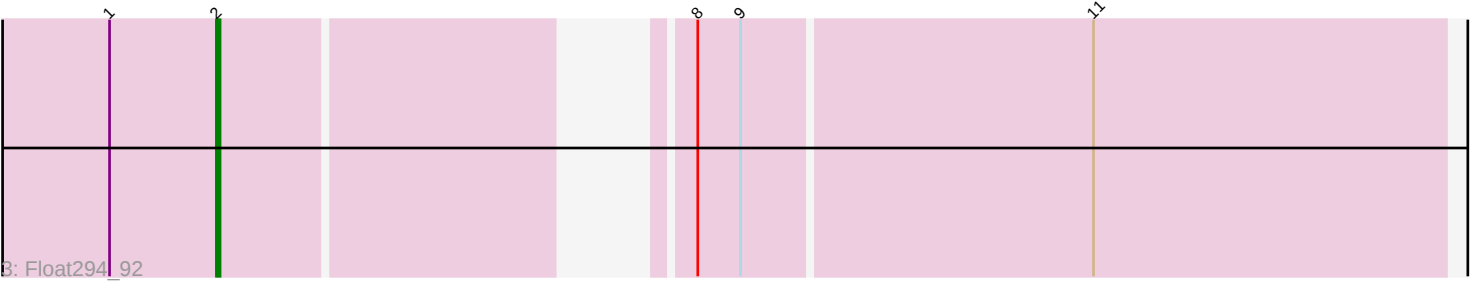
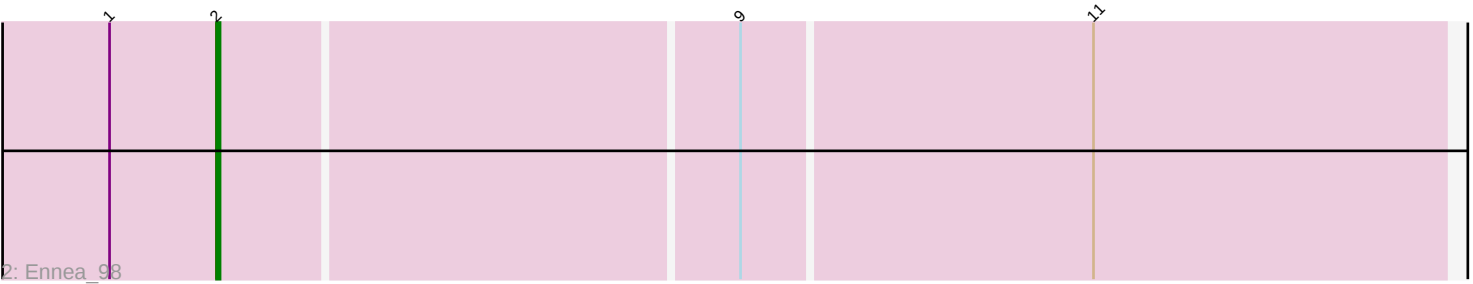


Pham 331746



Note: Tracks are now grouped by subcluster and scaled. Switching in subcluster is indicated by changes in track color. Track scale is now set by default to display the region 30 bp upstream of start 1 to 30 bp downstream of the last possible start. If this default region is judged to be packed too tightly with annotated starts, the track will be further scaled to only show that region of the ORF with annotated starts. This action will be indicated by adding "Zoomed" to the title. For starts, yellow indicates the location of called starts comprised solely of Glimmer/GeneMark auto-annotations, green indicates the location of called starts with at least 1 manual gene annotation.

Pham 331746 Report

This analysis was run 09/05/26 on database version 662.

Pham number 331746 has 5 members, 0 are drafts.

Phages represented in each track:

- Track 1 : Commandaria_92
- Track 2 : Ennea_98
- Track 3 : Float294_92
- Track 4 : RedRaider_95
- Track 5 : Lollipop1437_93

Summary of Final Annotations (See graph section above for start numbers):

The start number called the most often in the published annotations is 2, it was called in 4 of the 5 non-draft genes in the pham.

Genes that call this "Most Annotated" start:

- Ennea_98, Float294_92, Lollipop1437_93, RedRaider_95,

Genes that have the "Most Annotated" start but do not call it:

-

Genes that do not have the "Most Annotated" start:

- Commandaria_92,

Summary by start number:

Start 2:

- Found in 4 of 5 (80.0%) of genes in pham
- Manual Annotations of this start: 4 of 5
- Called 100.0% of time when present
- Phage (with cluster) where this start called: Ennea_98 (CR3), Float294_92 (CR3), Lollipop1437_93 (CR3), RedRaider_95 (CR3),

Start 3:

- Found in 1 of 5 (20.0%) of genes in pham
- Manual Annotations of this start: 1 of 5
- Called 100.0% of time when present
- Phage (with cluster) where this start called: Commandaria_92 (CR2),

Summary by clusters:

There are 2 clusters represented in this pham: CR2, CR3,

Info for manual annotations of cluster CR2:

- Start number 3 was manually annotated 1 time for cluster CR2.

Info for manual annotations of cluster CR3:

- Start number 2 was manually annotated 4 times for cluster CR3.

Gene Information:

Gene: Commandaria_92 Start: 66542, Stop: 66904, Start Num: 3

Candidate Starts for Commandaria_92:

(Start: 3 @66542 has 1 MA's), (4, 66599), (5, 66626), (10, 66737), (12, 66845),

Gene: Ennea_98 Start: 66466, Stop: 66801, Start Num: 2

Candidate Starts for Ennea_98:

(1, 66436), (Start: 2 @66466 has 4 MA's), (9, 66607), (11, 66703),

Gene: Float294_92 Start: 66408, Stop: 66716, Start Num: 2

Candidate Starts for Float294_92:

(1, 66378), (Start: 2 @66408 has 4 MA's), (8, 66510), (9, 66522), (11, 66618),

Gene: Lollipop1437_93 Start: 66146, Stop: 66481, Start Num: 2

Candidate Starts for Lollipop1437_93:

(Start: 2 @66146 has 4 MA's), (8, 66275), (9, 66287), (11, 66383),

Gene: RedRaider_95 Start: 67268, Stop: 67600, Start Num: 2

Candidate Starts for RedRaider_95:

(Start: 2 @67268 has 4 MA's), (6, 67385), (7, 67388), (8, 67394), (9, 67406), (11, 67502),