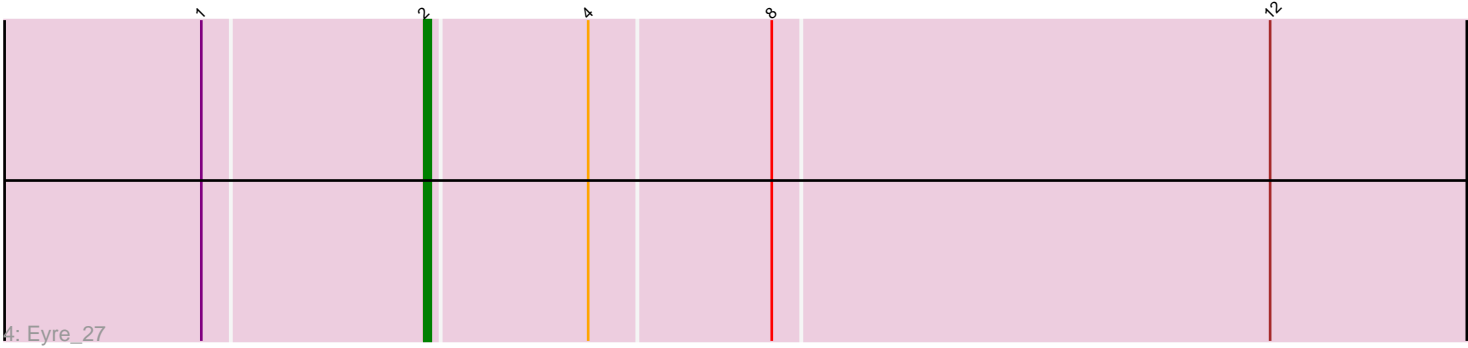
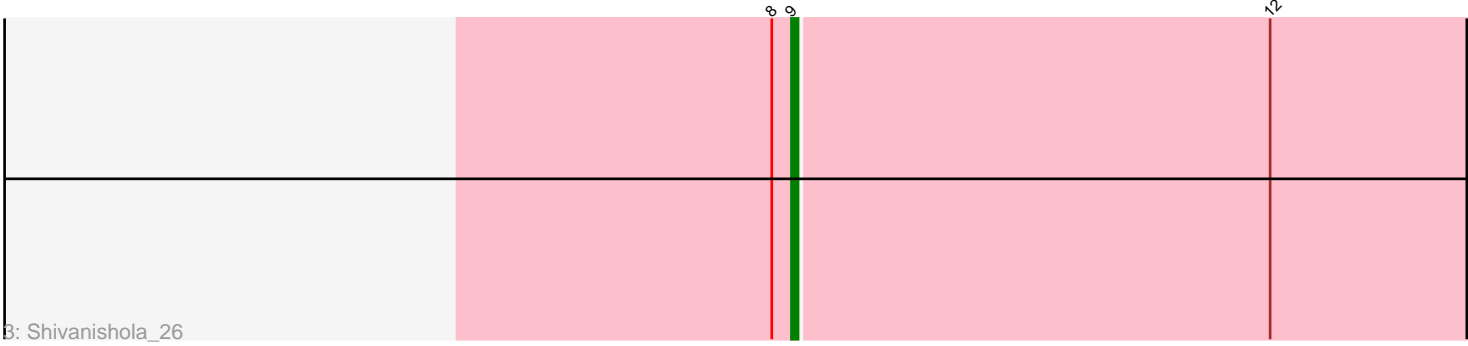
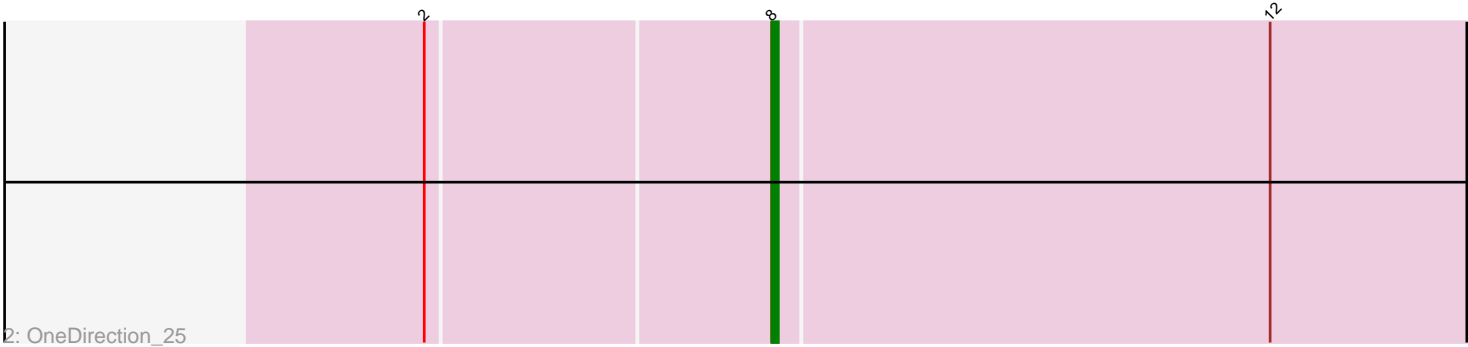
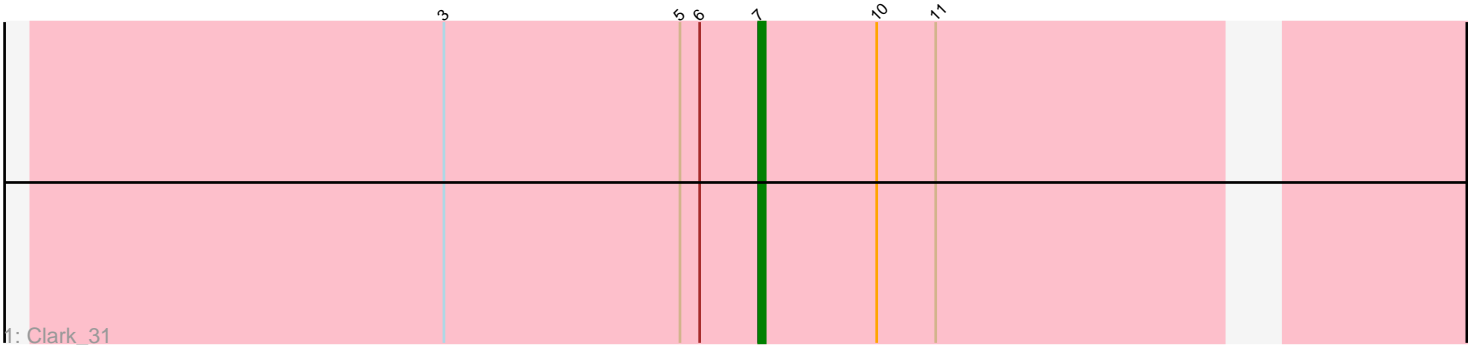


Pham 10334



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 10334 Report

This analysis was run 04/28/24 on database version 559.

Pham number 10334 has 4 members, 0 are drafts.

Phages represented in each track:

- Track 1 : Clark_31
- Track 2 : OneDirection_25
- Track 3 : Shivanishola_26
- Track 4 : Eyre_27

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

The start number called the most often in the published annotations is 7, it was called in 1 of the 4 non-draft genes in the pham.

Genes that call this "Most Annotated" start:

- Clark_31,

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

-

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

- Eyre_27, OneDirection_25, Shivanishola_26,

Summary by start number:

Start 2:

- Found in 2 of 4 (50.0%) of genes in pham
- Manual Annotations of this start: 1 of 4
- Called 50.0% of time when present
- Phage (with cluster) where this start called: Eyre_27 (singleton),

Start 7:

- Found in 1 of 4 (25.0%) of genes in pham
- Manual Annotations of this start: 1 of 4
- Called 100.0% of time when present
- Phage (with cluster) where this start called: Clark_31 (CZ4),

Start 8:

- Found in 3 of 4 (75.0%) of genes in pham
- Manual Annotations of this start: 1 of 4
- Called 33.3% of time when present
- Phage (with cluster) where this start called: OneDirection_25 (CZ6),

Start 9:

- Found in 1 of 4 (25.0%) of genes in pham
- Manual Annotations of this start: 1 of 4
- Called 100.0% of time when present
- Phage (with cluster) where this start called: Shivanishola_26 (DE1),

Summary by clusters:

There are 4 clusters represented in this pham: DE1, singleton, CZ6, CZ4,

Info for manual annotations of cluster CZ4:

- Start number 7 was manually annotated 1 time for cluster CZ4.

Info for manual annotations of cluster CZ6:

- Start number 8 was manually annotated 1 time for cluster CZ6.

Info for manual annotations of cluster DE1:

- Start number 9 was manually annotated 1 time for cluster DE1.

Gene Information:

Gene: Clark_31 Start: 25818, Stop: 25654, Start Num: 7

Candidate Starts for Clark_31:

(3, 25866), (5, 25830), (6, 25827), (Start: 7 @25818 has 1 MA's), (10, 25800), (11, 25791),

Gene: Eyre_27 Start: 24032, Stop: 23814, Start Num: 2

Candidate Starts for Eyre_27:

(1, 24065), (Start: 2 @24032 has 1 MA's), (4, 24008), (Start: 8 @23981 has 1 MA's), (12, 23906),

Gene: OneDirection_25 Start: 22884, Stop: 22717, Start Num: 8

Candidate Starts for OneDirection_25:

(Start: 2 @22935 has 1 MA's), (Start: 8 @22884 has 1 MA's), (12, 22809),

Gene: Shivanishola_26 Start: 21043, Stop: 20879, Start Num: 9

Candidate Starts for Shivanishola_26:

(Start: 8 @21046 has 1 MA's), (Start: 9 @21043 has 1 MA's), (12, 20971),