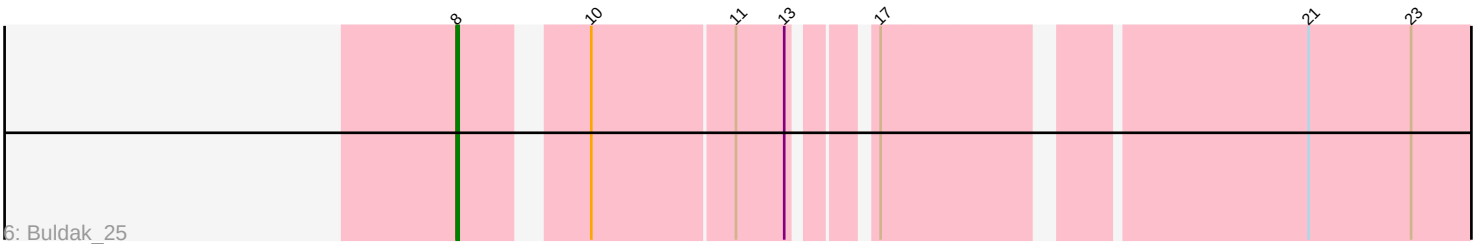
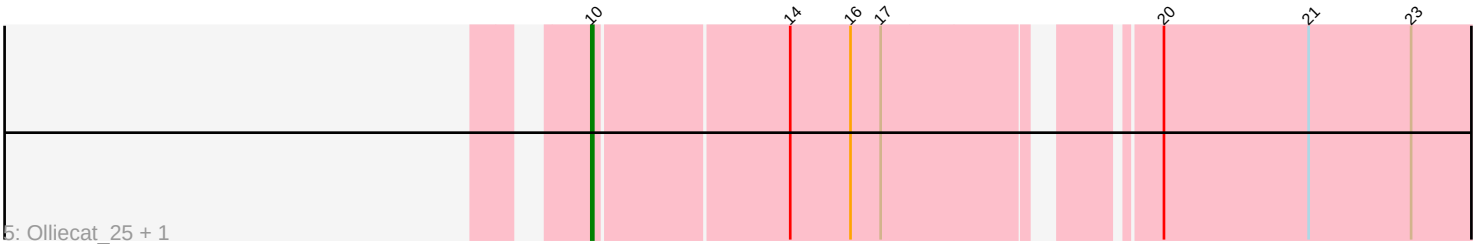
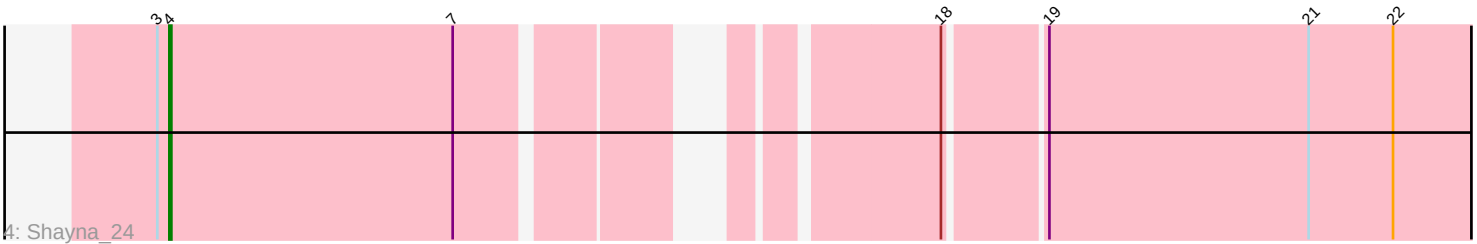
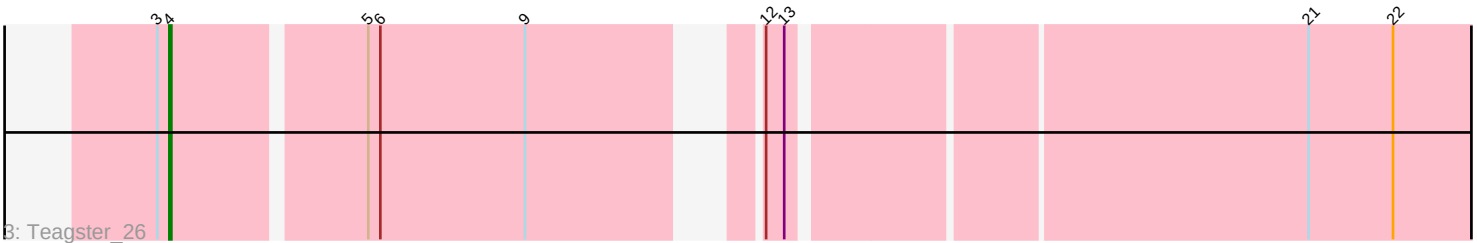
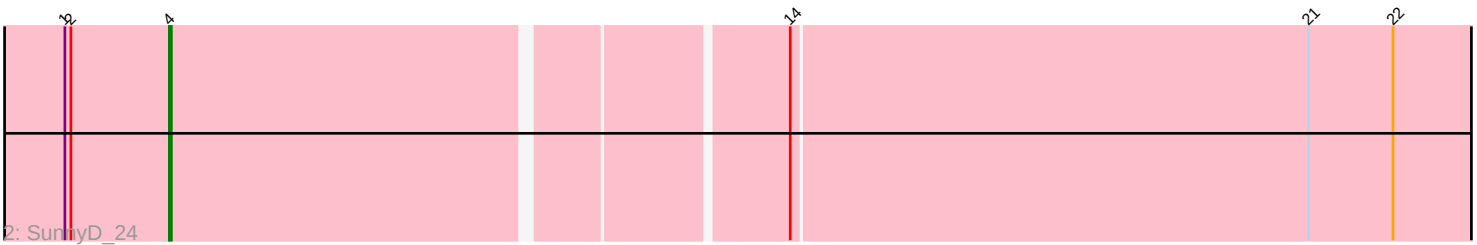
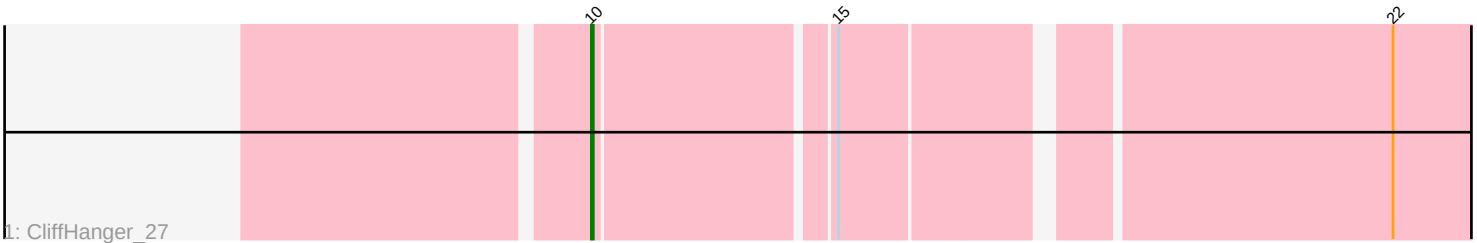


Pham 331588



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 331588 Report

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

Pham number 331588 has 7 members, 0 are drafts.

Phages represented in each track:

- Track 1 : CliffHanger_27
- Track 2 : SunnyD_24
- Track 3 : Teagster_26
- Track 4 : Shayna_24
- Track 5 : Olliecat_25, Squiracle_25
- Track 6 : Buldak_25

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

The start number called the most often in the published annotations is 4, it was called in 3 of the 7 non-draft genes in the pham.

Genes that call this "Most Annotated" start:

- Shayna_24, SunnyD_24, Teagster_26,

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

-

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

- Buldak_25, CliffHanger_27, Olliecat_25, Squiracle_25,

Summary by start number:

Start 4:

- Found in 3 of 7 (42.9%) of genes in pham
- Manual Annotations of this start: 3 of 7
- Called 100.0% of time when present
- Phage (with cluster) where this start called: Shayna_24 (EB), SunnyD_24 (EB), Teagster_26 (EB),

Start 8:

- Found in 1 of 7 (14.3%) of genes in pham
- Manual Annotations of this start: 1 of 7
- Called 100.0% of time when present

- Phage (with cluster) where this start called: Buldak_25 (EB),

Start 10:

- Found in 4 of 7 (57.1%) of genes in pham
- Manual Annotations of this start: 3 of 7
- Called 75.0% of time when present
- Phage (with cluster) where this start called: CliffHanger_27 (EB), Olliecat_25 (EB), Squiracle_25 (EB),

Summary by clusters:

There is one cluster represented in this pham: EB

Info for manual annotations of cluster EB:

- Start number 4 was manually annotated 3 times for cluster EB.
- Start number 8 was manually annotated 1 time for cluster EB.
- Start number 10 was manually annotated 3 times for cluster EB.

Gene Information:

Gene: Buldak_25 Start: 19698, Stop: 20216, Start Num: 8

Candidate Starts for Buldak_25:

(Start: 8 @19698 has 1 MA's), (Start: 10 @19746 has 3 MA's), (11, 19815), (13, 19839), (17, 19869), (21, 20061), (23, 20112),

Gene: CliffHanger_27 Start: 19625, Stop: 20101, Start Num: 10

Candidate Starts for CliffHanger_27:

(Start: 10 @19625 has 3 MA's), (15, 19736), (22, 19988),

Gene: Olliecat_25 Start: 19714, Stop: 20199, Start Num: 10

Candidate Starts for Olliecat_25:

(Start: 10 @19714 has 3 MA's), (14, 19807), (16, 19837), (17, 19852), (20, 19966), (21, 20038), (23, 20089),

Gene: Shayna_24 Start: 19653, Stop: 20309, Start Num: 4

Candidate Starts for Shayna_24:

(3, 19647), (Start: 4 @19653 has 3 MA's), (7, 19794), (18, 19983), (19, 20028), (21, 20157), (22, 20199),

Gene: Squiracle_25 Start: 19713, Stop: 20198, Start Num: 10

Candidate Starts for Squiracle_25:

(Start: 10 @19713 has 3 MA's), (14, 19806), (16, 19836), (17, 19851), (20, 19965), (21, 20037), (23, 20088),

Gene: SunnyD_24 Start: 19653, Stop: 20351, Start Num: 4

Candidate Starts for SunnyD_24:

(1, 19602), (2, 19605), (Start: 4 @19653 has 3 MA's), (14, 19944), (21, 20199), (22, 20241),

Gene: Teagster_26 Start: 20611, Stop: 21270, Start Num: 4

Candidate Starts for Teagster_26:

(3, 20605), (Start: 4 @20611 has 3 MA's), (5, 20701), (6, 20707), (9, 20779), (12, 20866), (13, 20875),
(21, 21118), (22, 21160),