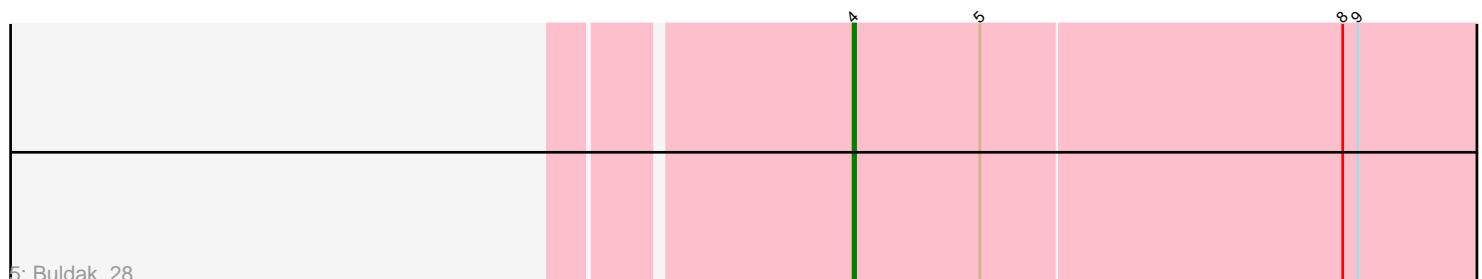
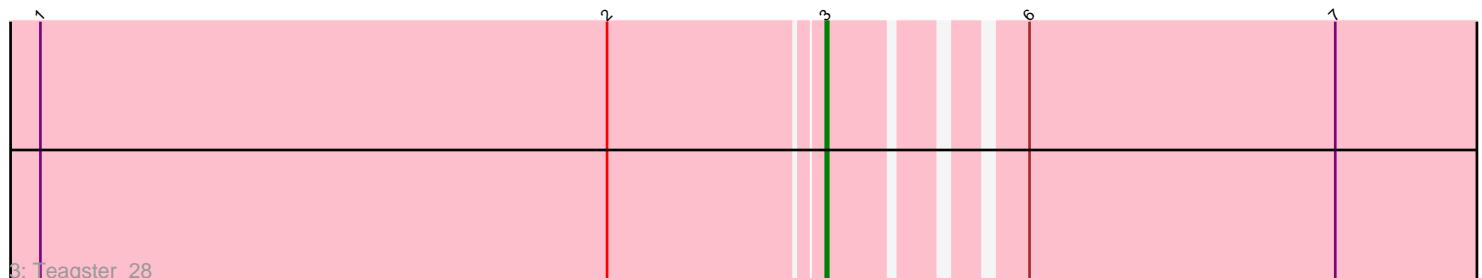
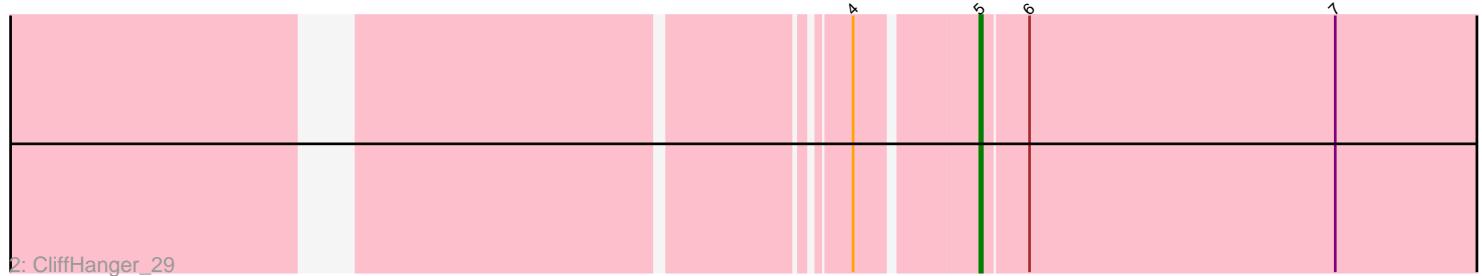
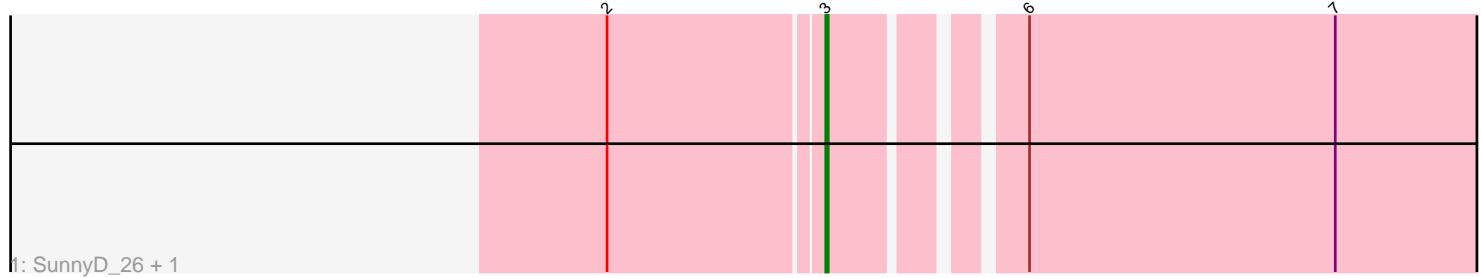


Pham 263851



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 263851 Report

This analysis was run 02/07/26 on database version 634.

Pham number 263851 has 7 members, 0 are drafts.

Phages represented in each track:

- Track 1 : SunnyD\_26, Shayna\_26
- Track 2 : CliffHanger\_29
- Track 3 : Teagster\_28
- Track 4 : Olliecat\_28, Squircle\_28
- Track 5 : Buldak\_28

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

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

Genes that call this "Most Annotated" start:

- Shayna\_26, SunnyD\_26, Teagster\_28,

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

- 

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

- Buldak\_28, CliffHanger\_29, Olliecat\_28, Squircle\_28,

### **Summary by start number:**

Start 3:

- 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\_26 (EB), SunnyD\_26 (EB), Teagster\_28 (EB),

Start 4:

- 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: Buldak\_28 (EB), Olliecat\_28 (EB), Squircle\_28 (EB),

Start 5:

- Found in 4 of 7 ( 57.1% ) of genes in pham
- Manual Annotations of this start: 1 of 7
- Called 25.0% of time when present
- Phage (with cluster) where this start called: CliffHanger\_29 (EB),

### **Summary by clusters:**

There is one cluster represented in this pham: EB

Info for manual annotations of cluster EB:

- Start number 3 was manually annotated 3 times for cluster EB.
- Start number 4 was manually annotated 3 times for cluster EB.
- Start number 5 was manually annotated 1 time for cluster EB.

### **Gene Information:**

Gene: Buldak\_28 Start: 20601, Stop: 20870, Start Num: 4

Candidate Starts for Buldak\_28:

(Start: 4 @20601 has 3 MA's), (Start: 5 @20652 has 1 MA's), (8, 20796), (9, 20802),

Gene: CliffHanger\_29 Start: 20337, Stop: 20555, Start Num: 5

Candidate Starts for CliffHanger\_29:

(Start: 4 @20292 has 3 MA's), (Start: 5 @20337 has 1 MA's), (6, 20355), (7, 20478),

Gene: Olliecat\_28 Start: 20581, Stop: 20850, Start Num: 4

Candidate Starts for Olliecat\_28:

(Start: 4 @20581 has 3 MA's), (Start: 5 @20632 has 1 MA's), (8, 20776), (9, 20782), (10, 20800),

Gene: Shayna\_26 Start: 20524, Stop: 20787, Start Num: 3

Candidate Starts for Shayna\_26:

(2, 20443), (Start: 3 @20524 has 3 MA's), (6, 20587), (7, 20710),

Gene: Squircle\_28 Start: 20580, Stop: 20849, Start Num: 4

Candidate Starts for Squircle\_28:

(Start: 4 @20580 has 3 MA's), (Start: 5 @20631 has 1 MA's), (8, 20775), (9, 20781), (10, 20799),

Gene: SunnyD\_26 Start: 20566, Stop: 20829, Start Num: 3

Candidate Starts for SunnyD\_26:

(2, 20485), (Start: 3 @20566 has 3 MA's), (6, 20629), (7, 20752),

Gene: Teagster\_28 Start: 21485, Stop: 21748, Start Num: 3

Candidate Starts for Teagster\_28:

(1, 21176), (2, 21404), (Start: 3 @21485 has 3 MA's), (6, 21548), (7, 21671),