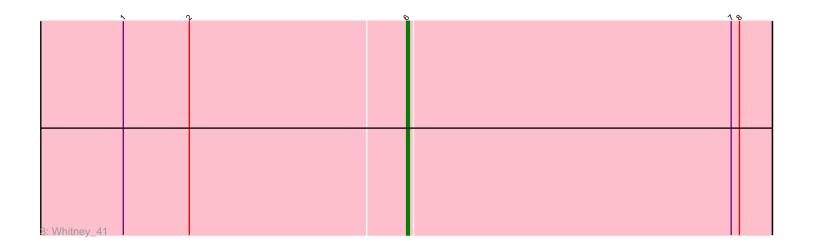
Pham 89085

	ი	6	18
1: Gambino_37 + 1			

	×	ଚ	
2: Lton_28			



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 89085 Report

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

Pham number 89085 has 4 members, 0 are drafts.

Phages represented in each track:

- Track 1 : Gambino_37, Azula_36
- Track 2 : Lton_28
- Track 3 : Whitney_41

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

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

Genes that call this "Most Annotated" start: • Azula 36, Gambino 37,

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

Genes that do not have the "Most Annotated" start: • Lton_28, Whitney_41,

Summary by start number:

Start 4:

- 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: Lton_28 (CZ),

Start 5:

- Found in 2 of 4 (50.0%) of genes in pham
- Manual Annotations of this start: 2 of 4
- Called 100.0% of time when present
- Phage (with cluster) where this start called: Azula_36 (CV), Gambino_37 (CV),

Start 6:

• 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: Whitney_41 (DN1),

Summary by clusters:

There are 3 clusters represented in this pham: CZ, DN1, CV,

Info for manual annotations of cluster CV: •Start number 5 was manually annotated 2 times for cluster CV.

Info for manual annotations of cluster CZ: •Start number 4 was manually annotated 1 time for cluster CZ.

Info for manual annotations of cluster DN1: •Start number 6 was manually annotated 1 time for cluster DN1.

Gene Information:

Gene: Azula_36 Start: 29166, Stop: 29023, Start Num: 5 Candidate Starts for Azula_36: (3, 29193), (Start: 5 @29166 has 2 MA's), (7, 29037), (8, 29034),

Gene: Gambino_37 Start: 29166, Stop: 29023, Start Num: 5 Candidate Starts for Gambino_37: (3, 29193), (Start: 5 @29166 has 2 MA's), (7, 29037), (8, 29034),

Gene: Lton_28 Start: 23301, Stop: 23146, Start Num: 4 Candidate Starts for Lton_28: (Start: 4 @23301 has 1 MA's), (8, 23157),

Gene: Whitney_41 Start: 32279, Stop: 32148, Start Num: 6 Candidate Starts for Whitney_41: (1, 32381), (2, 32357), (Start: 6 @32279 has 1 MA's), (7, 32162), (8, 32159),