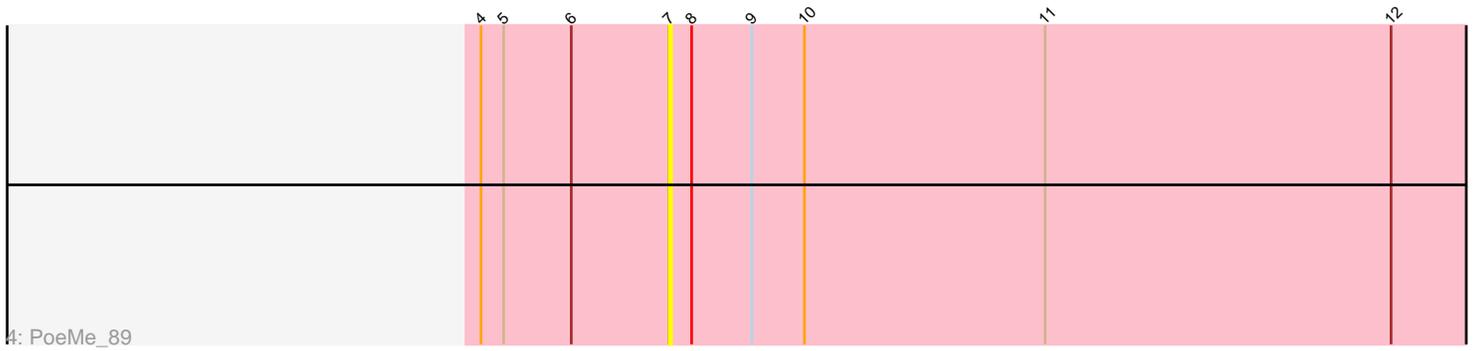
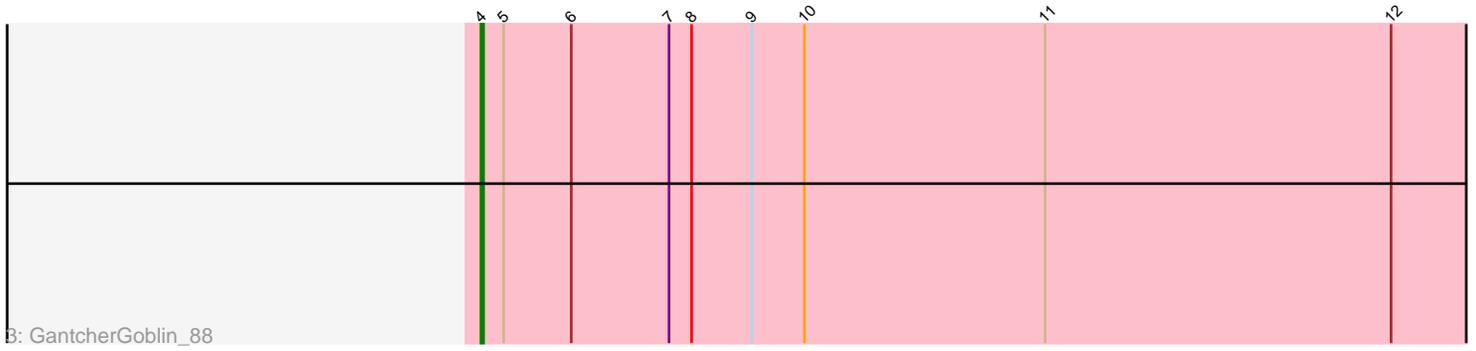
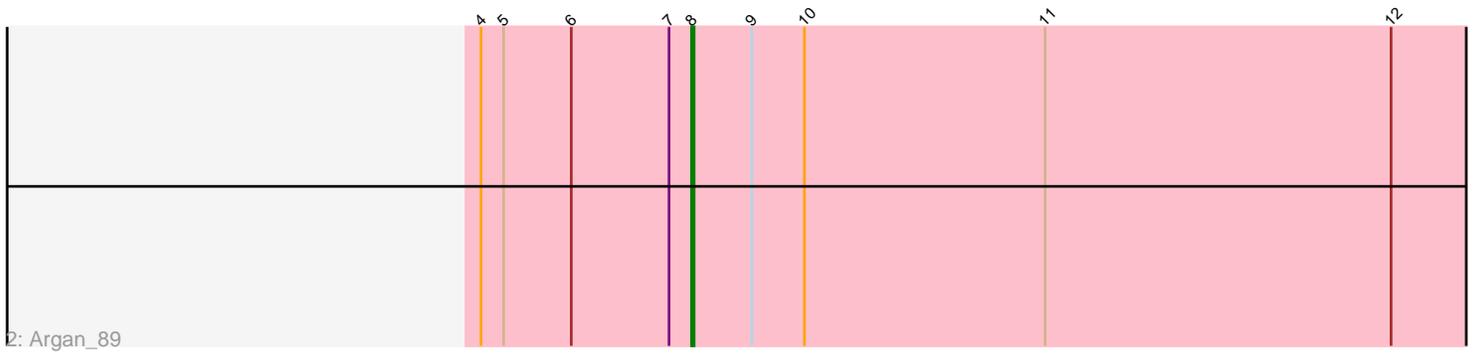
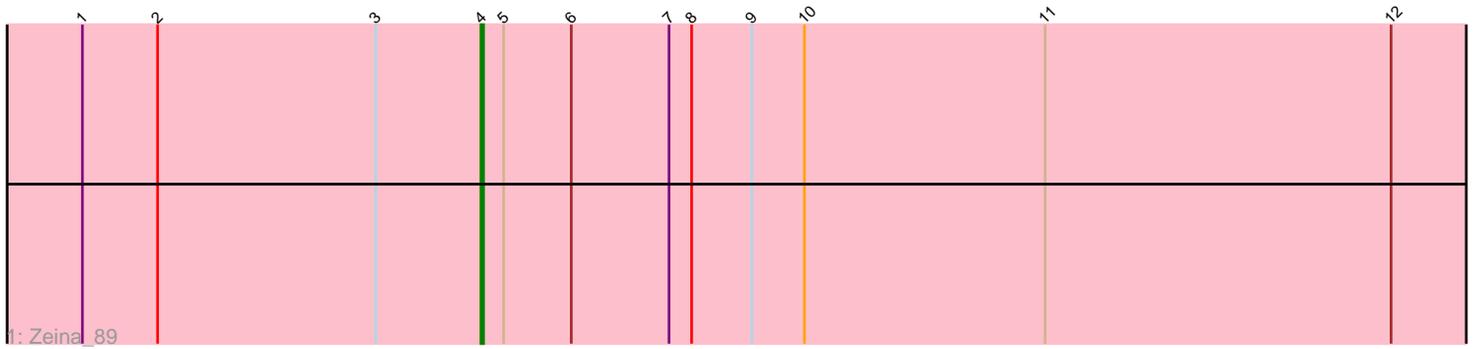


Pham 291982



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 291982 Report

This analysis was run 03/28/26 on database version 641.

Pham number 291982 has 4 members, 1 are drafts.

Phages represented in each track:

- Track 1 : Zeina_89
- Track 2 : Argan_89
- Track 3 : GantcherGoblin_88
- Track 4 : PoeMe_89

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 2 of the 3 non-draft genes in the pham.

Genes that call this "Most Annotated" start:

- GantcherGoblin_88, Zeina_89,

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

- Argan_89, PoeMe_89,

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

-

Summary by start number:

Start 4:

- Found in 4 of 4 (100.0%) of genes in pham
- Manual Annotations of this start: 2 of 3
- Called 50.0% of time when present
- Phage (with cluster) where this start called: GantcherGoblin_88 (AU6), Zeina_89 (AU6),

Start 7:

- Found in 4 of 4 (100.0%) of genes in pham
- No Manual Annotations of this start.
- Called 25.0% of time when present
- Phage (with cluster) where this start called: PoeMe_89 (AU6),

Start 8:

- Found in 4 of 4 (100.0%) of genes in pham
- Manual Annotations of this start: 1 of 3
- Called 25.0% of time when present
- Phage (with cluster) where this start called: Argan_89 (AU6),

Summary by clusters:

There is one cluster represented in this pham: AU6

Info for manual annotations of cluster AU6:

- Start number 4 was manually annotated 2 times for cluster AU6.
- Start number 8 was manually annotated 1 time for cluster AU6.

Gene Information:

Gene: Argan_89 Start: 51286, Stop: 51648, Start Num: 8

Candidate Starts for Argan_89:

(Start: 4 @51202 has 2 MA's), (5, 51211), (6, 51238), (7, 51277), (Start: 8 @51286 has 1 MA's), (9, 51310), (10, 51331), (11, 51427), (12, 51565),

Gene: GantcherGoblin_88 Start: 51349, Stop: 51795, Start Num: 4

Candidate Starts for GantcherGoblin_88:

(Start: 4 @51349 has 2 MA's), (5, 51358), (6, 51385), (7, 51424), (Start: 8 @51433 has 1 MA's), (9, 51457), (10, 51478), (11, 51574), (12, 51712),

Gene: PoeMe_89 Start: 51717, Stop: 52088, Start Num: 7

Candidate Starts for PoeMe_89:

(Start: 4 @51642 has 2 MA's), (5, 51651), (6, 51678), (7, 51717), (Start: 8 @51726 has 1 MA's), (9, 51750), (10, 51771), (11, 51867), (12, 52005),

Gene: Zeina_89 Start: 50998, Stop: 51444, Start Num: 4

Candidate Starts for Zeina_89:

(1, 50839), (2, 50869), (3, 50956), (Start: 4 @50998 has 2 MA's), (5, 51007), (6, 51034), (7, 51073), (Start: 8 @51082 has 1 MA's), (9, 51106), (10, 51127), (11, 51223), (12, 51361),