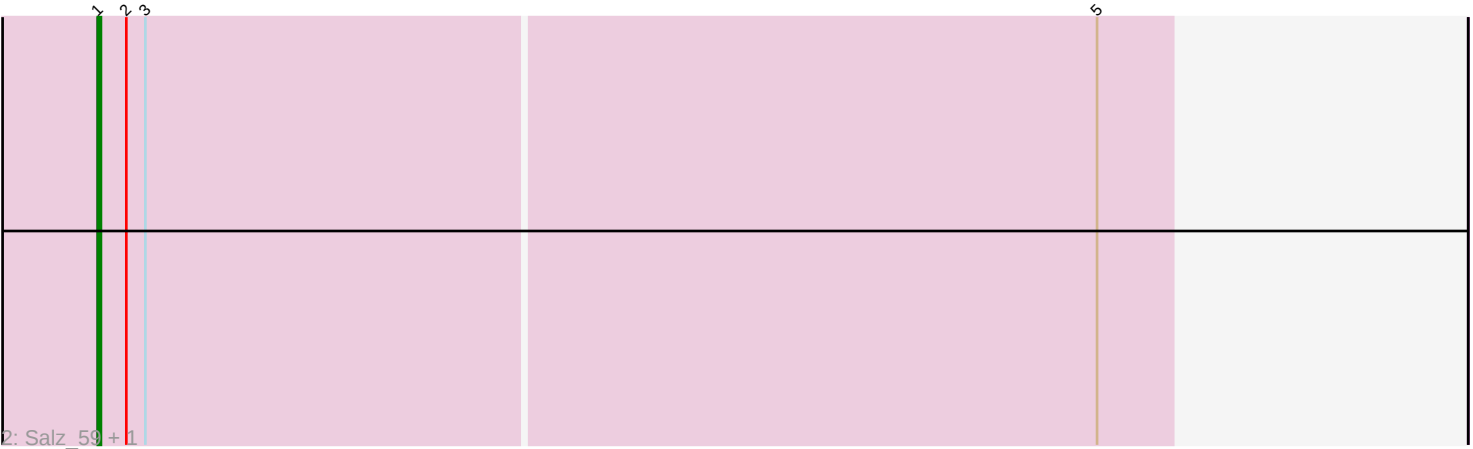
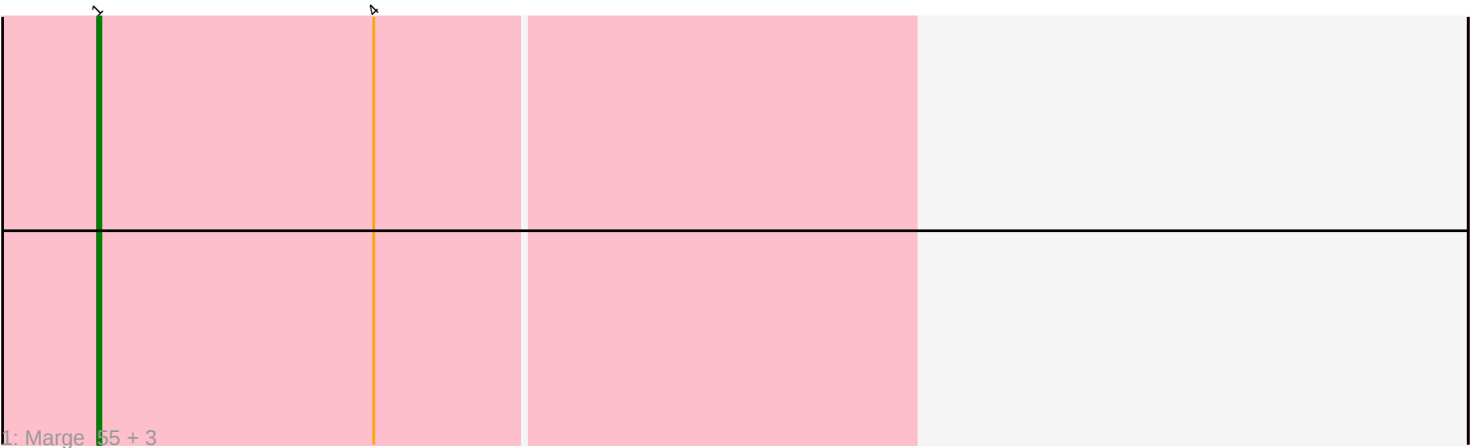




Note: Tracks are now grouped by subcluster and scaled. Switching in subcluster is indicated by changes in track color. Track scale is now set by default to display the region 30 bp upstream of start 1 to 30 bp downstream of the last possible start. If this default region is judged to be packed too tightly 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 310021 Report

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

Pham number 310021 has 7 members, 0 are drafts.

Phages represented in each track:

- Track 1 : Marge_55, Peterson_60, BaconJack_58, FrogHopper_53
- Track 2 : Salz_59, Et2Brutus_59
- Track 3 : Steamy_56

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

The start number called the most often in the published annotations is 1, it was called in 6 of the 7 non-draft genes in the pham.

Genes that call this "Most Annotated" start:

- BaconJack_58, Et2Brutus_59, FrogHopper_53, Marge_55, Peterson_60, Salz_59,

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

- Steamy_56,

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

-

Summary by start number:

Start 1:

- Found in 7 of 7 (100.0%) of genes in pham
- Manual Annotations of this start: 6 of 7
- Called 85.7% of time when present
- Phage (with cluster) where this start called: BaconJack_58 (A1), Et2Brutus_59 (A11), FrogHopper_53 (A1), Marge_55 (A1), Peterson_60 (A1), Salz_59 (A11),

Start 2:

- Found in 3 of 7 (42.9%) of genes in pham
- Manual Annotations of this start: 1 of 7
- Called 33.3% of time when present
- Phage (with cluster) where this start called: Steamy_56 (A12),

Summary by clusters:

There are 3 clusters represented in this pham: A1, A11, A12,

Info for manual annotations of cluster A1:

- Start number 1 was manually annotated 4 times for cluster A1.

Info for manual annotations of cluster A11:

- Start number 1 was manually annotated 2 times for cluster A11.

Info for manual annotations of cluster A12:

- Start number 2 was manually annotated 1 time for cluster A12.

Gene Information:

Gene: BaconJack_58 Start: 40981, Stop: 40727, Start Num: 1

Candidate Starts for BaconJack_58:

(Start: 1 @40981 has 6 MA's), (4, 40894),

Gene: Et2Brutus_59 Start: 38337, Stop: 38002, Start Num: 1

Candidate Starts for Et2Brutus_59:

(Start: 1 @38337 has 6 MA's), (Start: 2 @38328 has 1 MA's), (3, 38322), (5, 38025),

Gene: Froghopper_53 Start: 38148, Stop: 37894, Start Num: 1

Candidate Starts for Froghopper_53:

(Start: 1 @38148 has 6 MA's), (4, 38061),

Gene: Marge_55 Start: 39504, Stop: 39250, Start Num: 1

Candidate Starts for Marge_55:

(Start: 1 @39504 has 6 MA's), (4, 39417),

Gene: Peterson_60 Start: 41680, Stop: 41426, Start Num: 1

Candidate Starts for Peterson_60:

(Start: 1 @41680 has 6 MA's), (4, 41593),

Gene: Salz_59 Start: 38322, Stop: 37987, Start Num: 1

Candidate Starts for Salz_59:

(Start: 1 @38322 has 6 MA's), (Start: 2 @38313 has 1 MA's), (3, 38307), (5, 38010),

Gene: Steamy_56 Start: 38769, Stop: 38350, Start Num: 2

Candidate Starts for Steamy_56:

(Start: 1 @38778 has 6 MA's), (Start: 2 @38769 has 1 MA's), (5, 38466), (6, 38400), (7, 38373),