



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 284537 Report

This analysis was run 02/23/26 on database version 636.

Pham number 284537 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:

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### **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),