



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 107505 Report

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

Pham number 107505 has 4 members, 0 are drafts.

Phages represented in each track:

- Track 1 : Frogghopper_1, Gyzlar_1, Petruchio_1
- Track 2 : Marcell_1

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 3 of the 4 non-draft genes in the pham.

Genes that call this "Most Annotated" start:

- Frogghopper_1, Gyzlar_1, Petruchio_1,

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

- Marcell_1,

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

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Summary by start number:

Start 1:

- Found in 4 of 4 (100.0%) of genes in pham
- Manual Annotations of this start: 3 of 4
- Called 75.0% of time when present
- Phage (with cluster) where this start called: Frogghopper_1 (A1), Gyzlar_1 (A1), Petruchio_1 (A1),

Start 2:

- Found in 4 of 4 (100.0%) of genes in pham
- Manual Annotations of this start: 1 of 4
- Called 25.0% of time when present
- Phage (with cluster) where this start called: Marcell_1 (A1),

Summary by clusters:

There is one cluster represented in this pham: A1

Info for manual annotations of cluster A1:

- Start number 1 was manually annotated 3 times for cluster A1.
- Start number 2 was manually annotated 1 time for cluster A1.

Gene Information:

Gene: Froghopper_1 Start: 609, Stop: 749, Start Num: 1

Candidate Starts for Froghopper_1:

(Start: 1 @609 has 3 MA's), (Start: 2 @621 has 1 MA's), (3, 630), (4, 711), (5, 726),

Gene: Gyzlar_1 Start: 602, Stop: 742, Start Num: 1

Candidate Starts for Gyzlar_1:

(Start: 1 @602 has 3 MA's), (Start: 2 @614 has 1 MA's), (3, 623), (4, 704), (5, 719),

Gene: Marcell_1 Start: 615, Stop: 743, Start Num: 2

Candidate Starts for Marcell_1:

(Start: 1 @603 has 3 MA's), (Start: 2 @615 has 1 MA's), (3, 624), (4, 705), (5, 720),

Gene: Petruchio_1 Start: 609, Stop: 749, Start Num: 1

Candidate Starts for Petruchio_1:

(Start: 1 @609 has 3 MA's), (Start: 2 @621 has 1 MA's), (3, 630), (4, 711), (5, 726),