

Pham 7440

f: GenevaB15_9 + 1

2: MrMagoo_9 + 1

The diagram consists of a horizontal bar divided into two main sections. The left section is light gray and contains the text 'B: Estes_9' at the bottom left. The right section is light pink and contains two vertical lines, one green and one purple, with labels '6' and '1' above them respectively. A horizontal line runs across the middle of the bar, separating the top and bottom halves.

4: Rey_8

4

6

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 7440 Report

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

Pham number 7440 has 6 members, 0 are drafts.

Phages represented in each track:

- Track 1 : GenevaB15_9, Aziz_9
- Track 2 : MrMagoo_9, GardenSalsa_9
- Track 3 : Estes_9
- Track 4 : Rey_8

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

The start number called the most often in the published annotations is 6, it was called in 5 of the 6 non-draft genes in the pham.

Genes that call this "Most Annotated" start:

- Aziz_9, Estes_9, GardenSalsa_9, GenevaB15_9, MrMagoo_9,

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

- Rey_8,

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

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Summary by start number:

Start 4:

- Found in 1 of 6 (16.7%) of genes in pham
- Manual Annotations of this start: 1 of 6
- Called 100.0% of time when present
- Phage (with cluster) where this start called: Rey_8 (M2),

Start 6:

- Found in 6 of 6 (100.0%) of genes in pham
- Manual Annotations of this start: 5 of 6
- Called 83.3% of time when present
- Phage (with cluster) where this start called: Aziz_9 (M2), Estes_9 (M2), GardenSalsa_9 (M2), GenevaB15_9 (M2), MrMagoo_9 (M2),

Summary by clusters:

There is one cluster represented in this pham: M2

Info for manual annotations of cluster M2:

- Start number 4 was manually annotated 1 time for cluster M2.
- Start number 6 was manually annotated 5 times for cluster M2.

Gene Information:

Gene: Aziz_9 Start: 3005, Stop: 2856, Start Num: 6

Candidate Starts for Aziz_9:

(5, 3029), (Start: 6 @3005 has 5 MA's),

Gene: Estes_9 Start: 3010, Stop: 2861, Start Num: 6

Candidate Starts for Estes_9:

(Start: 6 @3010 has 5 MA's), (7, 2974),

Gene: GardenSalsa_9 Start: 3029, Stop: 2880, Start Num: 6

Candidate Starts for GardenSalsa_9:

(1, 3437), (2, 3269), (3, 3146), (5, 3053), (Start: 6 @3029 has 5 MA's),

Gene: GenevaB15_9 Start: 3005, Stop: 2856, Start Num: 6

Candidate Starts for GenevaB15_9:

(5, 3029), (Start: 6 @3005 has 5 MA's),

Gene: MrMagoo_9 Start: 3029, Stop: 2880, Start Num: 6

Candidate Starts for MrMagoo_9:

(1, 3437), (2, 3269), (3, 3146), (5, 3053), (Start: 6 @3029 has 5 MA's),

Gene: Rey_8 Start: 3067, Stop: 2867, Start Num: 4

Candidate Starts for Rey_8:

(Start: 4 @3067 has 1 MA's), (Start: 6 @3016 has 5 MA's),