



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 223709 Report

This analysis was run 03/28/25 on database version 593.

Pham number 223709 has 7 members, 2 are drafts.

Phages represented in each track:

- Track 1 : Razzleberry_4
- Track 2 : Ellison17_4, Prophecy_4, Momos_4, Mimi16_4, Grotle_4
- Track 3 : Ayka_4

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 5 of the 5 non-draft genes in the pham.

Genes that call this "Most Annotated" start:

- Ayka_4, Ellison17_4, Grotle_4, Mimi16_4, Momos_4, Prophecy_4, Razzleberry_4,

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

-

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: 5 of 5
- Called 100.0% of time when present
- Phage (with cluster) where this start called: Ayka_4 (UNK), Ellison17_4 (JB), Grotle_4 (JB), Mimi16_4 (JB), Momos_4 (JB), Prophecy_4 (JB), Razzleberry_4 (JB),

Summary by clusters:

There are 2 clusters represented in this pham: UNK, JB,

Info for manual annotations of cluster JB:

- Start number 1 was manually annotated 5 times for cluster JB.

Gene Information:

Gene: Ayka_4 Start: 3953, Stop: 4189, Start Num: 1

Candidate Starts for Ayka_4:

(Start: 1 @3953 has 5 MA's), (2, 4061), (3, 4103),

Gene: Ellison17_4 Start: 3732, Stop: 3974, Start Num: 1

Candidate Starts for Ellison17_4:

(Start: 1 @3732 has 5 MA's), (5, 3927),

Gene: Grotle_4 Start: 3691, Stop: 3933, Start Num: 1

Candidate Starts for Grotle_4:

(Start: 1 @3691 has 5 MA's), (5, 3886),

Gene: Mimi16_4 Start: 3742, Stop: 3984, Start Num: 1

Candidate Starts for Mimi16_4:

(Start: 1 @3742 has 5 MA's), (5, 3937),

Gene: Momos_4 Start: 3732, Stop: 3974, Start Num: 1

Candidate Starts for Momos_4:

(Start: 1 @3732 has 5 MA's), (5, 3927),

Gene: Prophecy_4 Start: 3742, Stop: 3984, Start Num: 1

Candidate Starts for Prophecy_4:

(Start: 1 @3742 has 5 MA's), (5, 3937),

Gene: Razzleberry_4 Start: 3717, Stop: 3950, Start Num: 1

Candidate Starts for Razzleberry_4:

(Start: 1 @3717 has 5 MA's), (4, 3888), (5, 3906),