



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 88637 Report

This analysis was run 04/05/24 on database version 557.

Pham number 88637 has 5 members, 0 are drafts.

Phages represented in each track:

- Track 1 : EniyanLRS_71, Azrael100_74, MaryV_75
- Track 2 : Wildcat_75
- Track 3 : Cosmo_75

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

The start number called the most often in the published annotations is 2, it was called in 3 of the 5 non-draft genes in the pham.

Genes that call this "Most Annotated" start:

- Azrael100_74, EniyanLRS_71, MaryV_75,

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

- Cosmo_75, Wildcat_75,

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

-

Summary by start number:

Start 1:

- Found in 5 of 5 (100.0%) of genes in pham
- Manual Annotations of this start: 1 of 5
- Called 20.0% of time when present
- Phage (with cluster) where this start called: Cosmo_75 (V),

Start 2:

- Found in 5 of 5 (100.0%) of genes in pham
- Manual Annotations of this start: 3 of 5
- Called 60.0% of time when present
- Phage (with cluster) where this start called: Azrael100_74 (V), EniyanLRS_71 (V), MaryV_75 (V),

Start 7:

- Found in 5 of 5 (100.0%) of genes in pham
- Manual Annotations of this start: 1 of 5
- Called 20.0% of time when present
- Phage (with cluster) where this start called: Wildcat_75 (V),

Summary by clusters:

There is one cluster represented in this pham: V

Info for manual annotations of cluster V:

- Start number 1 was manually annotated 1 time for cluster V.
- Start number 2 was manually annotated 3 times for cluster V.
- Start number 7 was manually annotated 1 time for cluster V.

Gene Information:

Gene: Azrael100_74 Start: 48048, Stop: 48575, Start Num: 2

Candidate Starts for Azrael100_74:

(Start: 1 @48018 has 1 MA's), (Start: 2 @48048 has 3 MA's), (3, 48069), (4, 48111), (5, 48132), (6, 48162), (Start: 7 @48174 has 1 MA's), (8, 48249), (9, 48255), (10, 48282), (11, 48384), (12, 48507),

Gene: Cosmo_75 Start: 48019, Stop: 48576, Start Num: 1

Candidate Starts for Cosmo_75:

(Start: 1 @48019 has 1 MA's), (Start: 2 @48049 has 3 MA's), (3, 48070), (4, 48112), (5, 48133), (6, 48163), (Start: 7 @48175 has 1 MA's), (8, 48250), (9, 48256), (10, 48283), (11, 48385), (12, 48508),

Gene: EniyanLRS_71 Start: 47788, Stop: 48315, Start Num: 2

Candidate Starts for EniyanLRS_71:

(Start: 1 @47758 has 1 MA's), (Start: 2 @47788 has 3 MA's), (3, 47809), (4, 47851), (5, 47872), (6, 47902), (Start: 7 @47914 has 1 MA's), (8, 47989), (9, 47995), (10, 48022), (11, 48124), (12, 48247),

Gene: MaryV_75 Start: 47870, Stop: 48397, Start Num: 2

Candidate Starts for MaryV_75:

(Start: 1 @47840 has 1 MA's), (Start: 2 @47870 has 3 MA's), (3, 47891), (4, 47933), (5, 47954), (6, 47984), (Start: 7 @47996 has 1 MA's), (8, 48071), (9, 48077), (10, 48104), (11, 48206), (12, 48329),

Gene: Wildcat_75 Start: 48006, Stop: 48407, Start Num: 7

Candidate Starts for Wildcat_75:

(Start: 1 @47850 has 1 MA's), (Start: 2 @47880 has 3 MA's), (3, 47901), (4, 47943), (5, 47964), (6, 47994), (Start: 7 @48006 has 1 MA's), (8, 48081), (9, 48087), (10, 48114), (11, 48216), (12, 48339),