



1: Phrann_62 + B

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 106854 Report

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

Pham number 106854 has 9 members, 0 are drafts.

Phages represented in each track:

- Track 1 : Phrann_62, Raymond7_56, Nenae_63, Purgamenstris_63, ShrimpFriedEgg_63, Snekmaggedon_58, SpongeBob_58, Jamie19_58, BabeRuth_63

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 9 of the 9 non-draft genes in the pham.

Genes that call this "Most Annotated" start:

- BabeRuth_63, Jamie19_58, Nenae_63, Phrann_62, Purgamenstris_63, Raymond7_56, ShrimpFriedEgg_63, Snekmaggedon_58, SpongeBob_58,

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 2:

- Found in 9 of 9 (100.0%) of genes in pham
- Manual Annotations of this start: 9 of 9
- Called 100.0% of time when present
- Phage (with cluster) where this start called: BabeRuth_63 (N), Jamie19_58 (N), Nenae_63 (N), Phrann_62 (N), Purgamenstris_63 (N), Raymond7_56 (N), ShrimpFriedEgg_63 (N), Snekmaggedon_58 (N), SpongeBob_58 (N),

Summary by clusters:

There is one cluster represented in this pham: N

Info for manual annotations of cluster N:

- Start number 2 was manually annotated 9 times for cluster N.

Gene Information:

Gene: BabeRuth_63 Start: 39742, Stop: 39858, Start Num: 2

Candidate Starts for BabeRuth_63:

(1, 39571), (Start: 2 @39742 has 9 MA's), (3, 39772), (4, 39799),

Gene: Jamie19_58 Start: 38422, Stop: 38538, Start Num: 2

Candidate Starts for Jamie19_58:

(1, 38251), (Start: 2 @38422 has 9 MA's), (3, 38452), (4, 38479),

Gene: Nenae_63 Start: 39744, Stop: 39860, Start Num: 2

Candidate Starts for Nenae_63:

(1, 39573), (Start: 2 @39744 has 9 MA's), (3, 39774), (4, 39801),

Gene: Phrann_62 Start: 42287, Stop: 42403, Start Num: 2

Candidate Starts for Phrann_62:

(1, 42116), (Start: 2 @42287 has 9 MA's), (3, 42317), (4, 42344),

Gene: Purgamenstris_63 Start: 39742, Stop: 39858, Start Num: 2

Candidate Starts for Purgamenstris_63:

(1, 39571), (Start: 2 @39742 has 9 MA's), (3, 39772), (4, 39799),

Gene: Raymond7_56 Start: 39530, Stop: 39646, Start Num: 2

Candidate Starts for Raymond7_56:

(1, 39359), (Start: 2 @39530 has 9 MA's), (3, 39560), (4, 39587),

Gene: ShrimpFriedEgg_63 Start: 39741, Stop: 39857, Start Num: 2

Candidate Starts for ShrimpFriedEgg_63:

(1, 39570), (Start: 2 @39741 has 9 MA's), (3, 39771), (4, 39798),

Gene: Snekmaggedon_58 Start: 38422, Stop: 38538, Start Num: 2

Candidate Starts for Snekmaggedon_58:

(1, 38251), (Start: 2 @38422 has 9 MA's), (3, 38452), (4, 38479),

Gene: SpongeBob_58 Start: 38422, Stop: 38538, Start Num: 2

Candidate Starts for SpongeBob_58:

(1, 38251), (Start: 2 @38422 has 9 MA's), (3, 38452), (4, 38479),