



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 193238 Report

This analysis was run 11/02/24 on database version 579.

Pham number 193238 has 5 members, 0 are drafts.

Phages represented in each track:

- Track 1 : KatChan_43, Chepli_43, Luna18_43
- Track 2 : Morrigan_43
- Track 3 : Triscuit_17

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

The start number called the most often in the published annotations is 4, it was called in 3 of the 5 non-draft genes in the pham.

Genes that call this "Most Annotated" start:

- Chepli_43, KatChan_43, Luna18_43,

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

-

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

- Morrigan_43, Triscuit_17,

Summary by start number:

Start 3:

- Found in 1 of 5 (20.0%) of genes in pham
- Manual Annotations of this start: 1 of 5
- Called 100.0% of time when present
- Phage (with cluster) where this start called: Triscuit_17 (GG),

Start 4:

- Found in 3 of 5 (60.0%) of genes in pham
- Manual Annotations of this start: 3 of 5
- Called 100.0% of time when present
- Phage (with cluster) where this start called: Chepli_43 (EA6), KatChan_43 (EA6), Luna18_43 (EA6),

Start 5:

- Found in 1 of 5 (20.0%) of genes in pham
- Manual Annotations of this start: 1 of 5
- Called 100.0% of time when present
- Phage (with cluster) where this start called: Morrigan_43 (EA6),

Summary by clusters:

There are 2 clusters represented in this pham: GG, EA6,

Info for manual annotations of cluster EA6:

- Start number 4 was manually annotated 3 times for cluster EA6.
- Start number 5 was manually annotated 1 time for cluster EA6.

Info for manual annotations of cluster GG:

- Start number 3 was manually annotated 1 time for cluster GG.

Gene Information:

Gene: Chepli_43 Start: 30856, Stop: 30521, Start Num: 4

Candidate Starts for Chepli_43:

(Start: 4 @30856 has 3 MA's), (7, 30739), (11, 30664), (17, 30610),

Gene: KatChan_43 Start: 30868, Stop: 30533, Start Num: 4

Candidate Starts for KatChan_43:

(Start: 4 @30868 has 3 MA's), (7, 30751), (11, 30676), (17, 30622),

Gene: Luna18_43 Start: 30868, Stop: 30533, Start Num: 4

Candidate Starts for Luna18_43:

(Start: 4 @30868 has 3 MA's), (7, 30751), (11, 30676), (17, 30622),

Gene: Morrigan_43 Start: 30869, Stop: 30540, Start Num: 5

Candidate Starts for Morrigan_43:

(1, 30908), (2, 30905), (Start: 5 @30869 has 1 MA's), (7, 30752), (8, 30695), (9, 30692), (10, 30686), (11, 30680), (13, 30671), (14, 30665), (16, 30662),

Gene: Triscuit_17 Start: 6303, Stop: 6596, Start Num: 3

Candidate Starts for Triscuit_17:

(Start: 3 @6303 has 1 MA's), (6, 6432), (12, 6513), (15, 6528),