



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 106904 Report

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

Pham number 106904 has 8 members, 0 are drafts.

Phages represented in each track:

- Track 1 : Espica_40, Nancinator_40, Shuman_40, Erik_40, UhSalsa_40, Takoda_40, Belenaria_40
- Track 2 : AngryOrchard_38

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 8 of the 8 non-draft genes in the pham.

Genes that call this "Most Annotated" start:

- AngryOrchard_38, Belenaria_40, Erik_40, Espica_40, Nancinator_40, Shuman_40, Takoda_40, UhSalsa_40,

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 8 of 8 (100.0%) of genes in pham
- Manual Annotations of this start: 8 of 8
- Called 100.0% of time when present
- Phage (with cluster) where this start called: AngryOrchard_38 (CA), Belenaria_40 (CA), Erik_40 (CA), Espica_40 (CA), Nancinator_40 (CA), Shuman_40 (CA), Takoda_40 (CA), UhSalsa_40 (CA),

Summary by clusters:

There is one cluster represented in this pham: CA

Info for manual annotations of cluster CA:

- Start number 2 was manually annotated 8 times for cluster CA.

Gene Information:

Gene: AngryOrchard_38 Start: 29897, Stop: 29739, Start Num: 2

Candidate Starts for AngryOrchard_38:

(1, 29909), (Start: 2 @29897 has 8 MA's), (4, 29792),

Gene: Belenaria_40 Start: 29965, Stop: 29807, Start Num: 2

Candidate Starts for Belenaria_40:

(1, 29977), (Start: 2 @29965 has 8 MA's), (3, 29884), (4, 29860),

Gene: Erik_40 Start: 29964, Stop: 29806, Start Num: 2

Candidate Starts for Erik_40:

(1, 29976), (Start: 2 @29964 has 8 MA's), (3, 29883), (4, 29859),

Gene: Espica_40 Start: 29965, Stop: 29807, Start Num: 2

Candidate Starts for Espica_40:

(1, 29977), (Start: 2 @29965 has 8 MA's), (3, 29884), (4, 29860),

Gene: Nancinator_40 Start: 29965, Stop: 29807, Start Num: 2

Candidate Starts for Nancinator_40:

(1, 29977), (Start: 2 @29965 has 8 MA's), (3, 29884), (4, 29860),

Gene: Shuman_40 Start: 29977, Stop: 29819, Start Num: 2

Candidate Starts for Shuman_40:

(1, 29989), (Start: 2 @29977 has 8 MA's), (3, 29896), (4, 29872),

Gene: Takoda_40 Start: 29939, Stop: 29781, Start Num: 2

Candidate Starts for Takoda_40:

(1, 29951), (Start: 2 @29939 has 8 MA's), (3, 29858), (4, 29834),

Gene: UhSalsa_40 Start: 29965, Stop: 29807, Start Num: 2

Candidate Starts for UhSalsa_40:

(1, 29977), (Start: 2 @29965 has 8 MA's), (3, 29884), (4, 29860),