



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 214975 Report

This analysis was run 02/22/25 on database version 588.

Pham number 214975 has 7 members, 0 are drafts.

Phages represented in each track:

- Track 1 : CarolAnn_23, KappaFarmDelta_24, Blino_23, Obliviate_22, ZiggyZoo_22
- Track 2 : JuJu_23
- Track 3 : Clown_47

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 7 of the 7 non-draft genes in the pham.

Genes that call this "Most Annotated" start:

- Blino_23, CarolAnn_23, Clown_47, JuJu_23, KappaFarmDelta_24, Obliviate_22, ZiggyZoo_22,

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: 7 of 7
- Called 100.0% of time when present
- Phage (with cluster) where this start called: Blino_23 (CV), CarolAnn_23 (CV), Clown_47 (DC2), JuJu_23 (CV), KappaFarmDelta_24 (CV), Obliviate_22 (CV), ZiggyZoo_22 (CV),

Summary by clusters:

There are 2 clusters represented in this pham: DC2, CV,

Info for manual annotations of cluster CV:

- Start number 1 was manually annotated 6 times for cluster CV.

Info for manual annotations of cluster DC2:

- Start number 1 was manually annotated 1 time for cluster DC2.

Gene Information:

Gene: Blino_23 Start: 20354, Stop: 20677, Start Num: 1

Candidate Starts for Blino_23:

(Start: 1 @20354 has 7 MA's), (3, 20408), (4, 20459), (6, 20492), (10, 20630),

Gene: CarolAnn_23 Start: 20353, Stop: 20676, Start Num: 1

Candidate Starts for CarolAnn_23:

(Start: 1 @20353 has 7 MA's), (3, 20407), (4, 20458), (6, 20491), (10, 20629),

Gene: Clown_47 Start: 37663, Stop: 37989, Start Num: 1

Candidate Starts for Clown_47:

(Start: 1 @37663 has 7 MA's), (2, 37675), (4, 37768), (5, 37771), (6, 37801), (9, 37933), (10, 37939),

Gene: JuJu_23 Start: 20326, Stop: 20649, Start Num: 1

Candidate Starts for JuJu_23:

(Start: 1 @20326 has 7 MA's), (3, 20380), (4, 20431), (6, 20464), (7, 20497), (8, 20548), (9, 20596), (10, 20602),

Gene: KappaFarmDelta_24 Start: 20754, Stop: 21077, Start Num: 1

Candidate Starts for KappaFarmDelta_24:

(Start: 1 @20754 has 7 MA's), (3, 20808), (4, 20859), (6, 20892), (10, 21030),

Gene: Oblivate_22 Start: 20176, Stop: 20499, Start Num: 1

Candidate Starts for Oblivate_22:

(Start: 1 @20176 has 7 MA's), (3, 20230), (4, 20281), (6, 20314), (10, 20452),

Gene: ZiggyZoo_22 Start: 20176, Stop: 20499, Start Num: 1

Candidate Starts for ZiggyZoo_22:

(Start: 1 @20176 has 7 MA's), (3, 20230), (4, 20281), (6, 20314), (10, 20452),