



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 106905 Report

This analysis was run 04/28/24 on database version 559.

Pham number 106905 has 9 members, 1 are drafts.

Phages represented in each track:

- Track 1 : TunaTartare\_108, Sham\_105, Annadreamy\_98, Blueeyedbeauty\_108
- Track 2 : Limpid\_107
- Track 3 : Beuffert\_105
- Track 4 : Faust\_108
- Track 5 : SeresaTree\_108
- Track 6 : Stigma\_107

### ***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 6 of the 8 non-draft genes in the pham.

Genes that call this "Most Annotated" start:

- Annadreamy\_98, Beuffert\_105, Blueeyedbeauty\_108, Faust\_108, Sham\_105, TunaTartare\_108,

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

- Limpid\_107, SeresaTree\_108, Stigma\_107,

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

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### **Summary by start number:**

Start 2:

- Found in 9 of 9 ( 100.0% ) of genes in pham
- Manual Annotations of this start: 6 of 8
- Called 66.7% of time when present
- Phage (with cluster) where this start called: Annadreamy\_98 (BK1), Beuffert\_105 (BK1), Blueeyedbeauty\_108 (BK1), Faust\_108 (BK1), Sham\_105 (BK1), TunaTartare\_108 (BK1),

Start 3:

- Found in 9 of 9 ( 100.0% ) of genes in pham

- Manual Annotations of this start: 2 of 8
- Called 33.3% of time when present
- Phage (with cluster) where this start called: Limpid\_107 (BK1), SeresaTree\_108 (BK1), Stigma\_107 (BK1),

### **Summary by clusters:**

There is one cluster represented in this pham: BK1

Info for manual annotations of cluster BK1:

- Start number 2 was manually annotated 6 times for cluster BK1.
- Start number 3 was manually annotated 2 times for cluster BK1.

### **Gene Information:**

Gene: Annadreamy\_98 Start: 67306, Stop: 67437, Start Num: 2

Candidate Starts for Annadreamy\_98:

(Start: 2 @67306 has 6 MA's), (Start: 3 @67318 has 2 MA's),

Gene: Beuffert\_105 Start: 72126, Stop: 72257, Start Num: 2

Candidate Starts for Beuffert\_105:

(1, 72096), (Start: 2 @72126 has 6 MA's), (Start: 3 @72138 has 2 MA's),

Gene: Blueeyedbeauty\_108 Start: 71223, Stop: 71354, Start Num: 2

Candidate Starts for Blueeyedbeauty\_108:

(Start: 2 @71223 has 6 MA's), (Start: 3 @71235 has 2 MA's),

Gene: Faust\_108 Start: 72794, Stop: 72925, Start Num: 2

Candidate Starts for Faust\_108:

(1, 72764), (Start: 2 @72794 has 6 MA's), (Start: 3 @72806 has 2 MA's),

Gene: Limpid\_107 Start: 72561, Stop: 72680, Start Num: 3

Candidate Starts for Limpid\_107:

(Start: 2 @72549 has 6 MA's), (Start: 3 @72561 has 2 MA's),

Gene: SeresaTree\_108 Start: 72188, Stop: 72307, Start Num: 3

Candidate Starts for SeresaTree\_108:

(1, 72146), (Start: 2 @72176 has 6 MA's), (Start: 3 @72188 has 2 MA's),

Gene: Sham\_105 Start: 72304, Stop: 72435, Start Num: 2

Candidate Starts for Sham\_105:

(Start: 2 @72304 has 6 MA's), (Start: 3 @72316 has 2 MA's),

Gene: Stigma\_107 Start: 72792, Stop: 72911, Start Num: 3

Candidate Starts for Stigma\_107:

(Start: 2 @72786 has 6 MA's), (Start: 3 @72792 has 2 MA's), (4, 72885),

Gene: TunaTartare\_108 Start: 74045, Stop: 74176, Start Num: 2

Candidate Starts for TunaTartare\_108:

(Start: 2 @74045 has 6 MA's), (Start: 3 @74057 has 2 MA's),