

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 8012 Report

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

Pham number 8012 has 7 members, 1 are drafts.

Phages represented in each track:

• Track 1 : Kradal 114, Satis 115

Track 2: Nirvana 125

Track 3: Frankenweenie\_124

Track 4 : EhyElimayoE\_115

Track 5 : Kela\_114, JustBecause\_113

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

The start number called the most often in the published annotations is 3, it was called in 5 of the 6 non-draft genes in the pham.

Genes that call this "Most Annotated" start:

• Frankenweenie\_124, JustBecause\_113, Kela\_114, Kradal\_114, Nirvana\_125, Satis\_115,

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

EhyElimayoE\_115,

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

### Summary by start number:

#### Start 2:

- Found in 3 of 7 (42.9%) of genes in pham
- Manual Annotations of this start: 1 of 6
- Called 33.3% of time when present
- Phage (with cluster) where this start called: EhyElimayoE\_115 (BM),

#### Start 3:

- Found in 7 of 7 (100.0%) of genes in pham
- Manual Annotations of this start: 5 of 6
- Called 85.7% of time when present

Phage (with cluster) where this start called: Frankenweenie\_124 (BM),
JustBecause\_113 (BM), Kela\_114 (BM), Kradal\_114 (BM), Nirvana\_125 (BM),
Satis\_115 (BM),

### **Summary by clusters:**

There is one cluster represented in this pham: BM

Info for manual annotations of cluster BM:

- •Start number 2 was manually annotated 1 time for cluster BM.
- •Start number 3 was manually annotated 5 times for cluster BM.

### Gene Information:

Gene: EhyElimayoE 115 Start: 76255, Stop: 75923, Start Num: 2

Candidate Starts for EhyElimayoE\_115:

(Start: 2 @76255 has 1 MA's), (Start: 3 @76213 has 5 MA's), (8, 76126), (12, 76054), (13, 76036), (15, 75982), (16, 75979),

Gene: Frankenweenie\_124 Start: 81091, Stop: 80807, Start Num: 3

Candidate Starts for Frankenweenie\_124:

(1, 81244), (Start: 3 @81091 has 5 MA's), (8, 81001), (9, 80983), (11, 80968), (13, 80911), (15, 80866), (16, 80863), (17, 80836),

Gene: JustBecause\_113 Start: 75304, Stop: 75020, Start Num: 3

Candidate Starts for JustBecause\_113:

(Start: 3 @75304 has 5 MA's), (5, 75289), (7, 75253), (8, 75214), (9, 75196), (13, 75124), (14, 75103), (15, 75079), (17, 75049),

Gene: Kela 114 Start: 75184, Stop: 74900, Start Num: 3

Candidate Starts for Kela 114:

(Start: 3 @75184 has 5 MA's), (5, 75169), (7, 75133), (8, 75094), (9, 75076), (13, 75004), (14, 74983), (15, 74959), (17, 74929),

Gene: Kradal\_114 Start: 76213, Stop: 75923, Start Num: 3

Candidate Starts for Kradal\_114:

(Start: 2 @76255 has 1 MA's), (Start: 3 @76213 has 5 MA's), (8, 76126), (12, 76054), (13, 76036), (15, 75982), (16, 75979),

Gene: Nirvana\_125 Start: 79482, Stop: 79198, Start Num: 3

Candidate Starts for Nirvana 125:

(Start: 3 @79482 has 5 MA's), (4, 79470), (6, 79446), (8, 79392), (9, 79374), (10, 79365), (13, 79302), (15, 79257), (17, 79227),

Gene: Satis\_115 Start: 76209, Stop: 75919, Start Num: 3

Candidate Starts for Satis 115:

(Start: 2 @76251 has 1 MA's), (Start: 3 @76209 has 5 MA's), (8, 76122), (12, 76050), (13, 76032), (15, 75978), (16, 75975),