



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 131305 Report

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

Pham number 131305 has 7 members, 1 are drafts.

Phages represented in each track:

- Track 1 : Frankenweenie_285
- Track 2 : Kradal_264, EhyElimayoE_267
- Track 3 : Kela_264, JustBecause_264
- Track 4 : Nirvana_301
- Track 5 : Satis_264

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 4 of the 6 non-draft genes in the pham.

Genes that call this "Most Annotated" start:

- EhyElimayoE_267, Frankenweenie_285, Kradal_264, Satis_264,

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

- JustBecause_264, Kela_264, Nirvana_301,

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

-

Summary by start number:

Start 2:

- Found in 2 of 7 (28.6%) of genes in pham
- Manual Annotations of this start: 2 of 6
- Called 100.0% of time when present
- Phage (with cluster) where this start called: JustBecause_264 (BM), Kela_264 (BM),

Start 3:

- Found in 7 of 7 (100.0%) of genes in pham
- Manual Annotations of this start: 4 of 6
- Called 57.1% of time when present
- Phage (with cluster) where this start called: EhyElimayoE_267 (BM), Frankenweenie_285 (BM), Kradal_264 (BM), Satis_264 (BM),

Start 4:

- Found in 7 of 7 (100.0%) of genes in pham
- No Manual Annotations of this start.
- Called 14.3% of time when present
- Phage (with cluster) where this start called: Nirvana_301 (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 2 times for cluster BM.
- Start number 3 was manually annotated 4 times for cluster BM.

Gene Information:

Gene: EhyElimayoE_267 Start: 152429, Stop: 151833, Start Num: 3

Candidate Starts for EhyElimayoE_267:

(Start: 3 @152429 has 4 MA's), (4, 152423), (8, 152327), (9, 152153), (10, 152096), (12, 151994), (15, 151862), (16, 151859),

Gene: Frankenweenie_285 Start: 163567, Stop: 162962, Start Num: 3

Candidate Starts for Frankenweenie_285:

(1, 163804), (Start: 3 @163567 has 4 MA's), (4, 163561), (5, 163525), (6, 163513), (7, 163483), (8, 163465), (9, 163291), (10, 163234), (11, 163138), (13, 163093), (14, 163072), (17, 162994),

Gene: JustBecause_264 Start: 149134, Stop: 148508, Start Num: 2

Candidate Starts for JustBecause_264:

(Start: 2 @149134 has 2 MA's), (Start: 3 @149113 has 4 MA's), (4, 149107), (8, 149011), (9, 148837), (14, 148618),

Gene: Kela_264 Start: 150738, Stop: 150112, Start Num: 2

Candidate Starts for Kela_264:

(Start: 2 @150738 has 2 MA's), (Start: 3 @150717 has 4 MA's), (4, 150711), (8, 150615), (9, 150441), (14, 150222),

Gene: Kradal_264 Start: 152426, Stop: 151830, Start Num: 3

Candidate Starts for Kradal_264:

(Start: 3 @152426 has 4 MA's), (4, 152420), (8, 152324), (9, 152150), (10, 152093), (12, 151991), (15, 151859), (16, 151856),

Gene: Nirvana_301 Start: 164015, Stop: 163416, Start Num: 4

Candidate Starts for Nirvana_301:

(Start: 3 @164021 has 4 MA's), (4, 164015), (5, 163979), (6, 163967), (7, 163937), (8, 163919), (9, 163745), (10, 163688), (14, 163526), (15, 163454), (17, 163448),

Gene: Satis_264 Start: 152763, Stop: 152167, Start Num: 3

Candidate Starts for Satis_264:

(Start: 3 @152763 has 4 MA's), (4, 152757), (5, 152721), (8, 152661), (9, 152487), (10, 152430), (12, 152328), (15, 152196), (16, 152193),

