



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 106951 Report

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

Pham number 106951 has 8 members, 0 are drafts.

Phages represented in each track:

- Track 1 : C3PO_51, Kimchi1738_52, Stickynote_53, PotatoChip_53, Zion_53, PeteyPab_52, Cruella_51
- Track 2 : Darwin_55

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

Genes that call this "Most Annotated" start:

- C3PO_51, Cruella_51, Darwin_55, Kimchi1738_52, PeteyPab_52, PotatoChip_53, Stickynote_53, Zion_53,

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 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: C3PO_51 (EN), Cruella_51 (EN), Darwin_55 (EN), Kimchi1738_52 (EN), PeteyPab_52 (EN), PotatoChip_53 (EN), Stickynote_53 (EN), Zion_53 (EN),

Summary by clusters:

There is one cluster represented in this pham: EN

Info for manual annotations of cluster EN:

- Start number 1 was manually annotated 8 times for cluster EN.

Gene Information:

Gene: C3PO_51 Start: 44310, Stop: 43690, Start Num: 1

Candidate Starts for C3PO_51:

(Start: 1 @44310 has 8 MA's), (2, 44202), (3, 44073), (4, 44067), (5, 43884), (6, 43806), (7, 43746), (8, 43707),

Gene: Cruella_51 Start: 44310, Stop: 43690, Start Num: 1

Candidate Starts for Cruella_51:

(Start: 1 @44310 has 8 MA's), (2, 44202), (3, 44073), (4, 44067), (5, 43884), (6, 43806), (7, 43746), (8, 43707),

Gene: Darwin_55 Start: 43791, Stop: 43171, Start Num: 1

Candidate Starts for Darwin_55:

(Start: 1 @43791 has 8 MA's), (2, 43683), (3, 43554), (4, 43548), (5, 43365), (7, 43227), (8, 43188),

Gene: Kimchi1738_52 Start: 44287, Stop: 43667, Start Num: 1

Candidate Starts for Kimchi1738_52:

(Start: 1 @44287 has 8 MA's), (2, 44179), (3, 44050), (4, 44044), (5, 43861), (6, 43783), (7, 43723), (8, 43684),

Gene: PeteyPab_52 Start: 44571, Stop: 43951, Start Num: 1

Candidate Starts for PeteyPab_52:

(Start: 1 @44571 has 8 MA's), (2, 44463), (3, 44334), (4, 44328), (5, 44145), (6, 44067), (7, 44007), (8, 43968),

Gene: PotatoChip_53 Start: 44573, Stop: 43953, Start Num: 1

Candidate Starts for PotatoChip_53:

(Start: 1 @44573 has 8 MA's), (2, 44465), (3, 44336), (4, 44330), (5, 44147), (6, 44069), (7, 44009), (8, 43970),

Gene: Stickynote_53 Start: 44577, Stop: 43957, Start Num: 1

Candidate Starts for Stickynote_53:

(Start: 1 @44577 has 8 MA's), (2, 44469), (3, 44340), (4, 44334), (5, 44151), (6, 44073), (7, 44013), (8, 43974),

Gene: Zion_53 Start: 44571, Stop: 43951, Start Num: 1

Candidate Starts for Zion_53:

(Start: 1 @44571 has 8 MA's), (2, 44463), (3, 44334), (4, 44328), (5, 44145), (6, 44067), (7, 44007), (8, 43968),