



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 88782 Report

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

Pham number 88782 has 6 members, 3 are drafts.

Phages represented in each track:

- Track 1 : Claus_67, Netyap_68, ZhongYanYuan_65, Hafay_68
- Track 2 : DrSeegs_67, Faith1_67

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 2 of the 3 non-draft genes in the pham.

Genes that call this "Most Annotated" start:

- DrSeegs_67, Faith1_67,

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

- Claus_67, Hafay_68, Netyap_68, ZhongYanYuan_65,

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

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Summary by start number:

Start 1:

- Found in 6 of 6 (100.0%) of genes in pham
- Manual Annotations of this start: 2 of 3
- Called 33.3% of time when present
- Phage (with cluster) where this start called: DrSeegs_67 (L2), Faith1_67 (L2),

Start 2:

- Found in 6 of 6 (100.0%) of genes in pham
- Manual Annotations of this start: 1 of 3
- Called 66.7% of time when present
- Phage (with cluster) where this start called: Claus_67 (L2), Hafay_68 (L2), Netyap_68 (L2), ZhongYanYuan_65 (L2),

Summary by clusters:

There is one cluster represented in this pham: L2

Info for manual annotations of cluster L2:

- Start number 1 was manually annotated 2 times for cluster L2.
- Start number 2 was manually annotated 1 time for cluster L2.

Gene Information:

Gene: Claus_67 Start: 46665, Stop: 46405, Start Num: 2

Candidate Starts for Claus_67:

(Start: 1 @46680 has 2 MA's), (Start: 2 @46665 has 1 MA's), (3, 46644), (4, 46635), (5, 46632), (6, 46617), (7, 46599), (8, 46536), (9, 46533), (10, 46467),

Gene: DrSeegs_67 Start: 46599, Stop: 46324, Start Num: 1

Candidate Starts for DrSeegs_67:

(Start: 1 @46599 has 2 MA's), (Start: 2 @46584 has 1 MA's), (3, 46563), (4, 46554), (5, 46551), (6, 46536), (7, 46518), (8, 46455), (9, 46452), (10, 46386),

Gene: Faith1_67 Start: 46599, Stop: 46324, Start Num: 1

Candidate Starts for Faith1_67:

(Start: 1 @46599 has 2 MA's), (Start: 2 @46584 has 1 MA's), (3, 46563), (4, 46554), (5, 46551), (6, 46536), (7, 46518), (8, 46455), (9, 46452), (10, 46386),

Gene: Hafay_68 Start: 46688, Stop: 46428, Start Num: 2

Candidate Starts for Hafay_68:

(Start: 1 @46703 has 2 MA's), (Start: 2 @46688 has 1 MA's), (3, 46667), (4, 46658), (5, 46655), (6, 46640), (7, 46622), (8, 46559), (9, 46556), (10, 46490),

Gene: Netyap_68 Start: 46585, Stop: 46325, Start Num: 2

Candidate Starts for Netyap_68:

(Start: 1 @46600 has 2 MA's), (Start: 2 @46585 has 1 MA's), (3, 46564), (4, 46555), (5, 46552), (6, 46537), (7, 46519), (8, 46456), (9, 46453), (10, 46387),

Gene: ZhongYanYuan_65 Start: 46479, Stop: 46219, Start Num: 2

Candidate Starts for ZhongYanYuan_65:

(Start: 1 @46494 has 2 MA's), (Start: 2 @46479 has 1 MA's), (3, 46458), (4, 46449), (5, 46446), (6, 46431), (7, 46413), (8, 46350), (9, 46347), (10, 46281),