



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 282555 Report

This analysis was run 02/23/26 on database version 636.

Pham number 282555 has 15 members, 0 are drafts.

Phages represented in each track:

- Track 1 : BetterKatz_56, Mulch_56, Nadeem_56, Ayotoya_57, Bock_57, Pimento_58, WheatThin_55, GrandSlam_57, DelRio_57, Hamood_57, Parada_56, Chop_57, Brylie_56, NancyRae_55, Francois_56

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 15 of the 15 non-draft genes in the pham.

Genes that call this "Most Annotated" start:

- Ayotoya_57, BetterKatz_56, Bock_57, Brylie_56, Chop_57, DelRio_57, Francois_56, GrandSlam_57, Hamood_57, Mulch_56, Nadeem_56, NancyRae_55, Parada_56, Pimento_58, WheatThin_55,

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 15 of 15 (100.0%) of genes in pham
- Manual Annotations of this start: 15 of 15
- Called 100.0% of time when present
- Phage (with cluster) where this start called: Ayotoya_57 (DI), BetterKatz_56 (DI), Bock_57 (DI), Brylie_56 (DI), Chop_57 (DI), DelRio_57 (DI), Francois_56 (DI), GrandSlam_57 (DI), Hamood_57 (DI), Mulch_56 (DI), Nadeem_56 (DI), NancyRae_55 (DI), Parada_56 (DI), Pimento_58 (DI), WheatThin_55 (DI),

Summary by clusters:

There is one cluster represented in this pham: DI

Info for manual annotations of cluster DI:

•Start number 1 was manually annotated 15 times for cluster DI.

Gene Information:

Gene: Ayotoya_57 Start: 39666, Stop: 39851, Start Num: 1

Candidate Starts for Ayotoya_57:

(Start: 1 @39666 has 15 MA's), (2, 39705), (3, 39756), (4, 39774),

Gene: BetterKatz_56 Start: 38865, Stop: 39053, Start Num: 1

Candidate Starts for BetterKatz_56:

(Start: 1 @38865 has 15 MA's), (2, 38904), (3, 38955), (4, 38973),

Gene: Bock_57 Start: 38898, Stop: 39083, Start Num: 1

Candidate Starts for Bock_57:

(Start: 1 @38898 has 15 MA's), (2, 38937), (3, 38988), (4, 39006),

Gene: Brylie_56 Start: 38646, Stop: 38834, Start Num: 1

Candidate Starts for Brylie_56:

(Start: 1 @38646 has 15 MA's), (2, 38685), (3, 38736), (4, 38754),

Gene: Chop_57 Start: 39417, Stop: 39602, Start Num: 1

Candidate Starts for Chop_57:

(Start: 1 @39417 has 15 MA's), (2, 39456), (3, 39507), (4, 39525),

Gene: DelRio_57 Start: 39828, Stop: 40013, Start Num: 1

Candidate Starts for DelRio_57:

(Start: 1 @39828 has 15 MA's), (2, 39867), (3, 39918), (4, 39936),

Gene: Francois_56 Start: 38904, Stop: 39092, Start Num: 1

Candidate Starts for Francois_56:

(Start: 1 @38904 has 15 MA's), (2, 38943), (3, 38994), (4, 39012),

Gene: GrandSlam_57 Start: 39417, Stop: 39602, Start Num: 1

Candidate Starts for GrandSlam_57:

(Start: 1 @39417 has 15 MA's), (2, 39456), (3, 39507), (4, 39525),

Gene: Hamood_57 Start: 39417, Stop: 39602, Start Num: 1

Candidate Starts for Hamood_57:

(Start: 1 @39417 has 15 MA's), (2, 39456), (3, 39507), (4, 39525),

Gene: Mulch_56 Start: 38646, Stop: 38834, Start Num: 1

Candidate Starts for Mulch_56:

(Start: 1 @38646 has 15 MA's), (2, 38685), (3, 38736), (4, 38754),

Gene: Nadeem_56 Start: 38634, Stop: 38822, Start Num: 1

Candidate Starts for Nadeem_56:

(Start: 1 @38634 has 15 MA's), (2, 38673), (3, 38724), (4, 38742),

Gene: NancyRae_55 Start: 38632, Stop: 38817, Start Num: 1

Candidate Starts for NancyRae_55:

(Start: 1 @38632 has 15 MA's), (2, 38671), (3, 38722), (4, 38740),

Gene: Parada_56 Start: 38646, Stop: 38834, Start Num: 1

Candidate Starts for Parada_56:

(Start: 1 @38646 has 15 MA's), (2, 38685), (3, 38736), (4, 38754),

Gene: Pimento_58 Start: 38545, Stop: 38730, Start Num: 1

Candidate Starts for Pimento_58:

(Start: 1 @38545 has 15 MA's), (2, 38584), (3, 38635), (4, 38653),

Gene: WheatThin_55 Start: 38634, Stop: 38822, Start Num: 1

Candidate Starts for WheatThin_55:

(Start: 1 @38634 has 15 MA's), (2, 38673), (3, 38724), (4, 38742),