



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 106942 Report

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

Pham number 106942 has 8 members, 0 are drafts.

Phages represented in each track:

- Track 1 : Pinto_58, PhrostyMug_57, DrFeelGood_55, Kykar_57
- Track 2 : KyMonks1A_60
- Track 3 : Scowl_58
- Track 4 : Jorgensen_58
- Track 5 : GageAP_58

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

The start number called the most often in the published annotations is 2, it was called in 8 of the 8 non-draft genes in the pham.

Genes that call this "Most Annotated" start:

- DrFeelGood_55, GageAP_58, Jorgensen_58, KyMonks1A_60, Kykar_57, PhrostyMug_57, Pinto_58, Scowl_58,

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 2:

- 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: DrFeelGood_55 (A1), GageAP_58 (A1), Jorgensen_58 (A1), KyMonks1A_60 (A1), Kykar_57 (A1), PhrostyMug_57 (A1), Pinto_58 (A1), Scowl_58 (A1),

Summary by clusters:

There is one cluster represented in this pham: A1

Info for manual annotations of cluster A1:

- Start number 2 was manually annotated 8 times for cluster A1.

Gene Information:

Gene: DrFeelGood_55 Start: 39103, Stop: 39038, Start Num: 2

Candidate Starts for DrFeelGood_55:

(1, 39112), (Start: 2 @39103 has 8 MA's), (4, 39049), (5, 39046),

Gene: GageAP_58 Start: 40190, Stop: 40125, Start Num: 2

Candidate Starts for GageAP_58:

(1, 40199), (Start: 2 @40190 has 8 MA's), (5, 40133),

Gene: Jorgensen_58 Start: 39649, Stop: 39584, Start Num: 2

Candidate Starts for Jorgensen_58:

(1, 39658), (Start: 2 @39649 has 8 MA's), (3, 39640), (5, 39592),

Gene: KyMonks1A_60 Start: 39860, Stop: 39795, Start Num: 2

Candidate Starts for KyMonks1A_60:

(1, 39869), (Start: 2 @39860 has 8 MA's), (4, 39806), (5, 39803),

Gene: Kykar_57 Start: 39306, Stop: 39241, Start Num: 2

Candidate Starts for Kykar_57:

(1, 39315), (Start: 2 @39306 has 8 MA's), (4, 39252), (5, 39249),

Gene: PhrostyMug_57 Start: 40431, Stop: 40366, Start Num: 2

Candidate Starts for PhrostyMug_57:

(1, 40440), (Start: 2 @40431 has 8 MA's), (4, 40377), (5, 40374),

Gene: Pinto_58 Start: 40016, Stop: 39951, Start Num: 2

Candidate Starts for Pinto_58:

(1, 40025), (Start: 2 @40016 has 8 MA's), (4, 39962), (5, 39959),

Gene: Scowl_58 Start: 39760, Stop: 39695, Start Num: 2

Candidate Starts for Scowl_58:

(1, 39769), (Start: 2 @39760 has 8 MA's), (3, 39751), (5, 39703),