



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 87982 Report

This analysis was run 04/05/24 on database version 557.

Pham number 87982 has 7 members, 0 are drafts.

Phages represented in each track:

- Track 1 : Ariel_52, Lucky2013_52, Superphikiman_52, MiaZeal_52, Squint_51, Porcelain_51, Courthouse_52

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 7 of the 7 non-draft genes in the pham.

Genes that call this "Most Annotated" start:

- Ariel_52, Courthouse_52, Lucky2013_52, MiaZeal_52, Porcelain_51, Squint_51, Superphikiman_52,

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 7 of 7 (100.0%) of genes in pham
- Manual Annotations of this start: 7 of 7
- Called 100.0% of time when present
- Phage (with cluster) where this start called: Ariel_52 (J), Courthouse_52 (J), Lucky2013_52 (J), MiaZeal_52 (J), Porcelain_51 (J), Squint_51 (J), Superphikiman_52 (J),

Summary by clusters:

There is one cluster represented in this pham: J

Info for manual annotations of cluster J:

- Start number 1 was manually annotated 7 times for cluster J.

Gene Information:

Gene: Ariel_52 Start: 40978, Stop: 40742, Start Num: 1

Candidate Starts for Ariel_52:

(Start: 1 @40978 has 7 MA's), (2, 40873),

Gene: Courthouse_52 Start: 41382, Stop: 41146, Start Num: 1

Candidate Starts for Courthouse_52:

(Start: 1 @41382 has 7 MA's), (2, 41277),

Gene: Lucky2013_52 Start: 41518, Stop: 41282, Start Num: 1

Candidate Starts for Lucky2013_52:

(Start: 1 @41518 has 7 MA's), (2, 41413),

Gene: MiaZeal_52 Start: 41193, Stop: 40957, Start Num: 1

Candidate Starts for MiaZeal_52:

(Start: 1 @41193 has 7 MA's), (2, 41088),

Gene: Porcelain_51 Start: 41193, Stop: 40957, Start Num: 1

Candidate Starts for Porcelain_51:

(Start: 1 @41193 has 7 MA's), (2, 41088),

Gene: Squint_51 Start: 41312, Stop: 41076, Start Num: 1

Candidate Starts for Squint_51:

(Start: 1 @41312 has 7 MA's), (2, 41207),

Gene: Superphikiman_52 Start: 41384, Stop: 41148, Start Num: 1

Candidate Starts for Superphikiman_52:

(Start: 1 @41384 has 7 MA's), (2, 41279),