

Pham 107536

Diagram illustrating a 2D array structure with 2 rows and 5 columns. The array is divided into three main color-coded regions: a light gray region (leftmost column), a light pink region (second column), and a light red region (remaining three columns). A green vertical line separates the first and second columns, and a brown vertical line separates the fourth and fifth columns. The bottom-left cell is labeled "1: Trooper_38 + 1". The top-left cell is labeled "1" and the bottom-right cell is labeled "5".

2: Acolyte_36

4: NothingSpecial_37

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 107536 Report

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

Pham number 107536 has 5 members, 1 are drafts.

Phages represented in each track:

- Track 1 : Trooper_38, TarsusIV_36
- Track 2 : Acolyte_36
- Track 3 : BabyBack_36
- Track 4 : NothingSpecial_37

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 4 of the 4 non-draft genes in the pham.

Genes that call this "Most Annotated" start:

- Acolyte_36, BabyBack_36, NothingSpecial_37, TarsusIV_36, Trooper_38,

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 5 of 5 (100.0%) of genes in pham
- Manual Annotations of this start: 4 of 4
- Called 100.0% of time when present
- Phage (with cluster) where this start called: Acolyte_36 (A2), BabyBack_36 (A2), NothingSpecial_37 (A2), TarsusIV_36 (A2), Trooper_38 (A2),

Summary by clusters:

There is one cluster represented in this pham: A2

Info for manual annotations of cluster A2:

- Start number 1 was manually annotated 4 times for cluster A2.

Gene Information:

Gene: Acolyte_36 Start: 28172, Stop: 28041, Start Num: 1

Candidate Starts for Acolyte_36:

(Start: 1 @28172 has 4 MA's), (3, 28097), (4, 28055),

Gene: BabyBack_36 Start: 28811, Stop: 28680, Start Num: 1

Candidate Starts for BabyBack_36:

(Start: 1 @28811 has 4 MA's), (2, 28748), (3, 28736), (5, 28685),

Gene: NothingSpecial_37 Start: 28979, Stop: 28848, Start Num: 1

Candidate Starts for NothingSpecial_37:

(Start: 1 @28979 has 4 MA's),

Gene: TarsusIV_36 Start: 27903, Stop: 27772, Start Num: 1

Candidate Starts for TarsusIV_36:

(Start: 1 @27903 has 4 MA's), (5, 27777),

Gene: Trooper_38 Start: 28137, Stop: 28006, Start Num: 1

Candidate Starts for Trooper_38:

(Start: 1 @28137 has 4 MA's), (5, 28011),