



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 107127 Report

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

Pham number 107127 has 6 members, 0 are drafts.

Phages represented in each track:

- Track 1 : Fireman_92, RobsFeet_95
- Track 2 : Gina_95, Teamocil_95
- Track 3 : Smarties_98, Ariadne_98

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

The start number called the most often in the published annotations is 4, it was called in 6 of the 6 non-draft genes in the pham.

Genes that call this "Most Annotated" start:

- Ariadne_98, Fireman_92, Gina_95, RobsFeet_95, Smarties_98, Teamocil_95,

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 4:

- Found in 6 of 6 (100.0%) of genes in pham
- Manual Annotations of this start: 6 of 6
- Called 100.0% of time when present
- Phage (with cluster) where this start called: Ariadne_98 (EC), Fireman_92 (EC), Gina_95 (EC), RobsFeet_95 (EC), Smarties_98 (EC), Teamocil_95 (EC),

Summary by clusters:

There is one cluster represented in this pham: EC

Info for manual annotations of cluster EC:

- Start number 4 was manually annotated 6 times for cluster EC.

Gene Information:

Gene: Ariadne_98 Start: 53358, Stop: 53744, Start Num: 4

Candidate Starts for Ariadne_98:

(Start: 4 @53358 has 6 MA's), (5, 53394), (6, 53430), (8, 53523), (9, 53595), (10, 53607), (12, 53667), (14, 53685),

Gene: Fireman_92 Start: 53474, Stop: 53860, Start Num: 4

Candidate Starts for Fireman_92:

(3, 53468), (Start: 4 @53474 has 6 MA's), (5, 53510), (7, 53558), (8, 53639), (9, 53711), (10, 53723), (11, 53774), (12, 53783), (14, 53801),

Gene: Gina_95 Start: 52671, Stop: 53051, Start Num: 4

Candidate Starts for Gina_95:

(1, 52251), (2, 52305), (Start: 4 @52671 has 6 MA's), (6, 52740), (9, 52905), (11, 52968), (12, 52977), (13, 52992),

Gene: RobsFeet_95 Start: 53084, Stop: 53470, Start Num: 4

Candidate Starts for RobsFeet_95:

(3, 53078), (Start: 4 @53084 has 6 MA's), (5, 53120), (7, 53168), (8, 53249), (9, 53321), (10, 53333), (11, 53384), (12, 53393), (14, 53411),

Gene: Smarties_98 Start: 53355, Stop: 53741, Start Num: 4

Candidate Starts for Smarties_98:

(Start: 4 @53355 has 6 MA's), (5, 53391), (6, 53427), (8, 53520), (9, 53592), (10, 53604), (12, 53664), (14, 53682),

Gene: Teamocil_95 Start: 52764, Stop: 53144, Start Num: 4

Candidate Starts for Teamocil_95:

(1, 52344), (2, 52398), (Start: 4 @52764 has 6 MA's), (6, 52833), (9, 52998), (11, 53061), (12, 53070), (13, 53085),