



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 107026 Report

This analysis was run 03/30/24 on database version 556.

Pham number 107026 has 7 members, 0 are drafts.

Phages represented in each track:

- Track 1 : PherrisBueller_3, Dreamboat_3, Jasper_3
- Track 2 : STLscum_3, Pepe_3
- Track 3 : Alsfro_3
- Track 4 : SkiPole_3

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

The start number called the most often in the published annotations is 3, it was called in 7 of the 7 non-draft genes in the pham.

Genes that call this "Most Annotated" start:

- Alsfro_3, Dreamboat_3, Jasper_3, Pepe_3, PherrisBueller_3, STLscum_3, SkiPole_3,

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 3:

- 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: Alsfro_3 (A1), Dreamboat_3 (A1), Jasper_3 (A1), Pepe_3 (A1), PherrisBueller_3 (A1), STLscum_3 (A1), SkiPole_3 (A1),

Summary by clusters:

There is one cluster represented in this pham: A1

Info for manual annotations of cluster A1:

- Start number 3 was manually annotated 7 times for cluster A1.

Gene Information:

Gene: Alsfro_3 Start: 1080, Stop: 1217, Start Num: 3

Candidate Starts for Alsfro_3:

(1, 960), (2, 969), (Start: 3 @1080 has 7 MA's), (4, 1137), (5, 1140), (6, 1188), (7, 1209),

Gene: Dreamboat_3 Start: 1168, Stop: 1305, Start Num: 3

Candidate Starts for Dreamboat_3:

(2, 1054), (Start: 3 @1168 has 7 MA's), (4, 1225), (5, 1228), (6, 1276), (7, 1297),

Gene: Jasper_3 Start: 1168, Stop: 1305, Start Num: 3

Candidate Starts for Jasper_3:

(2, 1054), (Start: 3 @1168 has 7 MA's), (4, 1225), (5, 1228), (6, 1276), (7, 1297),

Gene: Pepe_3 Start: 1162, Stop: 1299, Start Num: 3

Candidate Starts for Pepe_3:

(1, 1042), (2, 1051), (Start: 3 @1162 has 7 MA's), (4, 1219), (5, 1222), (6, 1270), (7, 1291),

Gene: PherrisBueller_3 Start: 1168, Stop: 1305, Start Num: 3

Candidate Starts for PherrisBueller_3:

(2, 1054), (Start: 3 @1168 has 7 MA's), (4, 1225), (5, 1228), (6, 1276), (7, 1297),

Gene: STLscum_3 Start: 1163, Stop: 1300, Start Num: 3

Candidate Starts for STLscum_3:

(1, 1043), (2, 1052), (Start: 3 @1163 has 7 MA's), (4, 1220), (5, 1223), (6, 1271), (7, 1292),

Gene: SkiPole_3 Start: 1163, Stop: 1300, Start Num: 3

Candidate Starts for SkiPole_3:

(2, 1052), (Start: 3 @1163 has 7 MA's), (4, 1220), (5, 1223), (6, 1271), (7, 1292),