



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 28093 Report

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

Pham number 28093 has 6 members, 0 are drafts.

Phages represented in each track:

- Track 1 : Fryberger_115, Guey18_120, Ronaldo_117, Volt_119
- Track 2 : Keelan_116
- Track 3 : Ziko_118

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 5 of the 6 non-draft genes in the pham.

Genes that call this "Most Annotated" start:

- Fryberger_115, Guey18_120, Keelan_116, Ronaldo_117, Volt_119,

Genes that have the "Most Annotated" start but do not call it:

- Ziko_118,

Genes that do not have the "Most Annotated" start:

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Summary by start number:

Start 1:

- Found in 6 of 6 (100.0%) of genes in pham
- Manual Annotations of this start: 5 of 6
- Called 83.3% of time when present
- Phage (with cluster) where this start called: Fryberger_115 (DP), Guey18_120 (DP), Keelan_116 (DP), Ronaldo_117 (DP), Volt_119 (DP),

Start 2:

- Found in 5 of 6 (83.3%) of genes in pham
- Manual Annotations of this start: 1 of 6
- Called 20.0% of time when present
- Phage (with cluster) where this start called: Ziko_118 (DP),

Summary by clusters:

There is one cluster represented in this pham: DP

Info for manual annotations of cluster DP:

- Start number 1 was manually annotated 5 times for cluster DP.
- Start number 2 was manually annotated 1 time for cluster DP.

Gene Information:

Gene: Fryberger_115 Start: 55508, Stop: 55876, Start Num: 1

Candidate Starts for Fryberger_115:

(Start: 1 @55508 has 5 MA's), (Start: 2 @55532 has 1 MA's), (4, 55688), (5, 55769), (8, 55826), (9, 55868),

Gene: Guey18_120 Start: 56831, Stop: 57199, Start Num: 1

Candidate Starts for Guey18_120:

(Start: 1 @56831 has 5 MA's), (Start: 2 @56855 has 1 MA's), (4, 57011), (5, 57092), (8, 57149), (9, 57191),

Gene: Keelan_116 Start: 56367, Stop: 56732, Start Num: 1

Candidate Starts for Keelan_116:

(Start: 1 @56367 has 5 MA's), (3, 56448), (4, 56544), (6, 56655), (7, 56661), (8, 56682), (9, 56724),

Gene: Ronaldo_117 Start: 56413, Stop: 56781, Start Num: 1

Candidate Starts for Ronaldo_117:

(Start: 1 @56413 has 5 MA's), (Start: 2 @56437 has 1 MA's), (4, 56593), (5, 56674), (8, 56731), (9, 56773),

Gene: Volt_119 Start: 56577, Stop: 56945, Start Num: 1

Candidate Starts for Volt_119:

(Start: 1 @56577 has 5 MA's), (Start: 2 @56601 has 1 MA's), (4, 56757), (5, 56838), (8, 56895), (9, 56937),

Gene: Ziko_118 Start: 56443, Stop: 56787, Start Num: 2

Candidate Starts for Ziko_118:

(Start: 1 @56419 has 5 MA's), (Start: 2 @56443 has 1 MA's), (4, 56599), (5, 56680), (8, 56737), (9, 56779),