



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 168862 Report

This analysis was run 07/09/24 on database version 566.

Pham number 168862 has 7 members, 0 are drafts.

Phages represented in each track:

- Track 1 : Goose_73, OKCentral2016_71
- Track 2 : Eaglepride_75
- Track 3 : Poompha_69, Severus_70
- Track 4 : WalterMcMickey_71, Twister_72

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

Genes that call this "Most Annotated" start:

- Goose_73, OKCentral2016_71, Poompha_69, Severus_70, Twister_72, WalterMcMickey_71,

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

- Eaglepride_75,

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

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Summary by start number:

Start 3:

- Found in 7 of 7 (100.0%) of genes in pham
- Manual Annotations of this start: 6 of 7
- Called 85.7% of time when present
- Phage (with cluster) where this start called: Goose_73 (A10), OKCentral2016_71 (A10), Poompha_69 (A10), Severus_70 (A10), Twister_72 (A10), WalterMcMickey_71 (A10),

Start 4:

- Found in 3 of 7 (42.9%) of genes in pham
- Manual Annotations of this start: 1 of 7
- Called 33.3% of time when present

- Phage (with cluster) where this start called: Eaglepride_75 (A10),

Summary by clusters:

There is one cluster represented in this pham: A10

Info for manual annotations of cluster A10:

- Start number 3 was manually annotated 6 times for cluster A10.
- Start number 4 was manually annotated 1 time for cluster A10.

Gene Information:

Gene: Eaglepride_75 Start: 44715, Stop: 44551, Start Num: 4

Candidate Starts for Eaglepride_75:

(1, 44940), (Start: 3 @44733 has 6 MA's), (Start: 4 @44715 has 1 MA's), (5, 44568),

Gene: Goose_73 Start: 44452, Stop: 44270, Start Num: 3

Candidate Starts for Goose_73:

(1, 44659), (Start: 3 @44452 has 6 MA's), (Start: 4 @44434 has 1 MA's), (5, 44287), (6, 44281),

Gene: OKCentral2016_71 Start: 43879, Stop: 43697, Start Num: 3

Candidate Starts for OKCentral2016_71:

(1, 44086), (Start: 3 @43879 has 6 MA's), (Start: 4 @43861 has 1 MA's), (5, 43714), (6, 43708),

Gene: Poompha_69 Start: 43781, Stop: 43587, Start Num: 3

Candidate Starts for Poompha_69:

(Start: 3 @43781 has 6 MA's), (7, 43592),

Gene: Severus_70 Start: 43781, Stop: 43587, Start Num: 3

Candidate Starts for Severus_70:

(Start: 3 @43781 has 6 MA's), (7, 43592),

Gene: Twister_72 Start: 44636, Stop: 44454, Start Num: 3

Candidate Starts for Twister_72:

(1, 44843), (2, 44693), (Start: 3 @44636 has 6 MA's), (6, 44465),

Gene: WalterMcMickey_71 Start: 44636, Stop: 44454, Start Num: 3

Candidate Starts for WalterMcMickey_71:

(1, 44843), (2, 44693), (Start: 3 @44636 has 6 MA's), (6, 44465),