



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 6961 Report

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

Pham number 6961 has 7 members, 0 are drafts.

Phages represented in each track:

- Track 1 : Zerg_80, Coco12_76, Sarma624_78, BigPhil_73, Velveteen_74, Cerasum_73, VRedHorse_84

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

Genes that call this "Most Annotated" start:

- BigPhil_73, Cerasum_73, Coco12_76, Sarma624_78, VRedHorse_84, Velveteen_74, Zerg_80,

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 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: BigPhil_73 (F1), Cerasum_73 (F1), Coco12_76 (F1), Sarma624_78 (F1), VRedHorse_84 (F1), Velveteen_74 (F1), Zerg_80 (F1),

Summary by clusters:

There is one cluster represented in this pham: F1

Info for manual annotations of cluster F1:

- Start number 1 was manually annotated 7 times for cluster F1.

Gene Information:

Gene: BigPhil_73 Start: 44623, Stop: 44790, Start Num: 1

Candidate Starts for BigPhil_73:

(Start: 1 @44623 has 7 MA's), (2, 44632), (3, 44644), (4, 44680),

Gene: Cerasum_73 Start: 44641, Stop: 44808, Start Num: 1

Candidate Starts for Cerasum_73:

(Start: 1 @44641 has 7 MA's), (2, 44650), (3, 44662), (4, 44698),

Gene: Coco12_76 Start: 48698, Stop: 48865, Start Num: 1

Candidate Starts for Coco12_76:

(Start: 1 @48698 has 7 MA's), (2, 48707), (3, 48719), (4, 48755),

Gene: Sarma624_78 Start: 45317, Stop: 45484, Start Num: 1

Candidate Starts for Sarma624_78:

(Start: 1 @45317 has 7 MA's), (2, 45326), (3, 45338), (4, 45374),

Gene: VRedHorse_84 Start: 47526, Stop: 47693, Start Num: 1

Candidate Starts for VRedHorse_84:

(Start: 1 @47526 has 7 MA's), (2, 47535), (3, 47547), (4, 47583),

Gene: Velveteen_74 Start: 45317, Stop: 45484, Start Num: 1

Candidate Starts for Velveteen_74:

(Start: 1 @45317 has 7 MA's), (2, 45326), (3, 45338), (4, 45374),

Gene: Zerg_80 Start: 48868, Stop: 49035, Start Num: 1

Candidate Starts for Zerg_80:

(Start: 1 @48868 has 7 MA's), (2, 48877), (3, 48889), (4, 48925),