



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 162494 Report

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

Pham number 162494 has 7 members, 0 are drafts.

Phages represented in each track:

- Track 1 : Coco12_80, Kingsley_87, Velveteen_77, Sarma624_82, BigPhil_76, Cerasum_77
- Track 2 : Latretium_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_76, Cerasum_77, Coco12_80, Kingsley_87, Latretium_84, Sarma624_82, Velveteen_77,

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_76 (F1), Cerasum_77 (F1), Coco12_80 (F1), Kingsley_87 (F1), Latretium_84 (F1), Sarma624_82 (F1), Velveteen_77 (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_76 Start: 45545, Stop: 45787, Start Num: 1

Candidate Starts for BigPhil_76:

(Start: 1 @45545 has 7 MA's), (2, 45569), (3, 45578), (4, 45611), (5, 45623), (6, 45626), (7, 45674), (8, 45701),

Gene: Cerasum_77 Start: 45563, Stop: 45805, Start Num: 1

Candidate Starts for Cerasum_77:

(Start: 1 @45563 has 7 MA's), (2, 45587), (3, 45596), (4, 45629), (5, 45641), (6, 45644), (7, 45692), (8, 45719),

Gene: Coco12_80 Start: 49620, Stop: 49862, Start Num: 1

Candidate Starts for Coco12_80:

(Start: 1 @49620 has 7 MA's), (2, 49644), (3, 49653), (4, 49686), (5, 49698), (6, 49701), (7, 49749), (8, 49776),

Gene: Kingsley_87 Start: 52227, Stop: 52469, Start Num: 1

Candidate Starts for Kingsley_87:

(Start: 1 @52227 has 7 MA's), (2, 52251), (3, 52260), (4, 52293), (5, 52305), (6, 52308), (7, 52356), (8, 52383),

Gene: Latretium_84 Start: 48945, Stop: 49187, Start Num: 1

Candidate Starts for Latretium_84:

(Start: 1 @48945 has 7 MA's), (2, 48969), (3, 48978), (4, 49011), (5, 49023), (6, 49026), (7, 49074), (8, 49101),

Gene: Sarma624_82 Start: 46239, Stop: 46481, Start Num: 1

Candidate Starts for Sarma624_82:

(Start: 1 @46239 has 7 MA's), (2, 46263), (3, 46272), (4, 46305), (5, 46317), (6, 46320), (7, 46368), (8, 46395),

Gene: Velveteen_77 Start: 46239, Stop: 46481, Start Num: 1

Candidate Starts for Velveteen_77:

(Start: 1 @46239 has 7 MA's), (2, 46263), (3, 46272), (4, 46305), (5, 46317), (6, 46320), (7, 46368), (8, 46395),