

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 216784 Report

This analysis was run 02/22/25 on database version 588.

Pham number 216784 has 10 members, 0 are drafts.

Phages represented in each track:

Track 1 : YDN12_45

Track 2: Piccadilly_48, Eklok_48, Cumberbatch_49, Eastland_48

Track 3 : AxeJC 47

Track 4: HFrancette_48, Ignacio_47, Vondra_47

Track 5 : Gilgamesh_21

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

The start number called the most often in the published annotations is 7, it was called in 8 of the 10 non-draft genes in the pham.

Genes that call this "Most Annotated" start:

AxeJC_47, Cumberbatch_49, Eastland_48, Eklok_48, HFrancette_48, Ignacio_47, Piccadilly_48, Vondra_47,

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

•

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

Gilgamesh_21, YDN12_45,

Summary by start number:

Start 6:

- Found in 2 of 10 (20.0%) of genes in pham
- Manual Annotations of this start: 2 of 10
- Called 100.0% of time when present
- Phage (with cluster) where this start called: Gilgamesh_21 (singleton), YDN12_45 (BG),

Start 7:

- Found in 8 of 10 (80.0%) of genes in pham
- Manual Annotations of this start: 8 of 10
- Called 100.0% of time when present

 Phage (with cluster) where this start called: AxeJC_47 (BP), Cumberbatch_49 (BP), Eastland_48 (BP), Eklok_48 (BP), HFrancette_48 (BP), Ignacio_47 (BP), Piccadilly_48 (BP), Vondra_47 (BP),

Summary by clusters:

There are 3 clusters represented in this pham: singleton, BG, BP,

Info for manual annotations of cluster BG:

Start number 6 was manually annotated 1 time for cluster BG.

Info for manual annotations of cluster BP:

Start number 7 was manually annotated 8 times for cluster BP.

Gene Information:

Gene: AxeJC_47 Start: 32027, Stop: 32236, Start Num: 7

Candidate Starts for AxeJC_47: (Start: 7 @32027 has 8 MA's),

Gene: Cumberbatch_49 Start: 32249, Stop: 32467, Start Num: 7

Candidate Starts for Cumberbatch 49:

(1, 31925), (2, 31949), (3, 32156), (Start: 7 @32249 has 8 MA's), (11, 32399),

Gene: Eastland 48 Start: 32209, Stop: 32427, Start Num: 7

Candidate Starts for Eastland_48:

(1, 31885), (2, 31909), (3, 32116), (Start: 7 @32209 has 8 MA's), (11, 32359),

Gene: Eklok_48 Start: 32008, Stop: 32226, Start Num: 7

Candidate Starts for Eklok 48:

(1, 31684), (2, 31708), (3, 31915), (Start: 7 @32008 has 8 MA's), (11, 32158),

Gene: Gilgamesh 21 Start: 13106, Stop: 12825, Start Num: 6

Candidate Starts for Gilgamesh_21:

(3, 13196), (5, 13109), (Start: 6 @13106 has 2 MA's), (8, 13088),

Gene: HFrancette 48 Start: 32550, Stop: 32768, Start Num: 7

Candidate Starts for HFrancette 48:

(Start: 7 @32550 has 8 MA's), (11, 32700),

Gene: Ignacio 47 Start: 32453, Stop: 32671, Start Num: 7

Candidate Starts for Ignacio_47:

(Start: 7 @ 32453 has 8 MA's), (11, 32603),

Gene: Piccadilly_48 Start: 32208, Stop: 32426, Start Num: 7

Candidate Starts for Piccadilly 48:

(1, 31884), (2, 31908), (3, 32115), (Start: 7 @32208 has 8 MA's), (11, 32358),

Gene: Vondra 47 Start: 31794, Stop: 32012, Start Num: 7

Candidate Starts for Vondra 47:

(Start: 7 @31794 has 8 MA's), (11, 31944),

Gene: YDN12_45 Start: 34681, Stop: 34436, Start Num: 6

Candidate Starts for YDN12_45:

(4, 34759), (Start: 6 @34681 has 2 MA's), (8, 34663), (9, 34645), (10, 34549), (12, 34498), (13, 34477),

(14, 34462), (15, 34459), (16, 34447),