



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 106939 Report

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

Pham number 106939 has 10 members, 2 are drafts.

Phages represented in each track:

- Track 1 : Mudcat_85, Hankly_90, Cheesy_91, Tribby_92, Kardesai_89, Heisenberger_89, Mooshroom_89, JEGGS_89, Benllo_89, BenitoAntonio_88

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

The start number called the most often in the published annotations is 2, it was called in 8 of the 8 non-draft genes in the pham.

Genes that call this "Most Annotated" start:

- BenitoAntonio_88, Benllo_89, Cheesy_91, Hankly_90, Heisenberger_89, JEGGS_89, Kardesai_89, Mooshroom_89, Mudcat_85, Tribby_92,

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 2:

- Found in 10 of 10 (100.0%) of genes in pham
- Manual Annotations of this start: 8 of 8
- Called 100.0% of time when present
- Phage (with cluster) where this start called: BenitoAntonio_88 (AM), Benllo_89 (AM), Cheesy_91 (AM), Hankly_90 (AM), Heisenberger_89 (AM), JEGGS_89 (AM), Kardesai_89 (AM), Mooshroom_89 (AM), Mudcat_85 (AM), Tribby_92 (AM),

Summary by clusters:

There is one cluster represented in this pham: AM

Info for manual annotations of cluster AM:

- Start number 2 was manually annotated 8 times for cluster AM.

Gene Information:

Gene: BenitoAntonio_88 Start: 51737, Stop: 52045, Start Num: 2
Candidate Starts for BenitoAntonio_88:
(1, 51734), (Start: 2 @51737 has 8 MA's), (3, 51884), (4, 51911), (5, 51926),

Gene: Benllo_89 Start: 52391, Stop: 52699, Start Num: 2
Candidate Starts for Benllo_89:
(1, 52388), (Start: 2 @52391 has 8 MA's), (3, 52538), (4, 52565), (5, 52580),

Gene: Cheesy_91 Start: 52482, Stop: 52790, Start Num: 2
Candidate Starts for Cheesy_91:
(1, 52479), (Start: 2 @52482 has 8 MA's), (3, 52629), (4, 52656), (5, 52671),

Gene: Hankly_90 Start: 51826, Stop: 52134, Start Num: 2
Candidate Starts for Hankly_90:
(1, 51823), (Start: 2 @51826 has 8 MA's), (3, 51973), (4, 52000), (5, 52015),

Gene: Heisenberger_89 Start: 51947, Stop: 52255, Start Num: 2
Candidate Starts for Heisenberger_89:
(1, 51944), (Start: 2 @51947 has 8 MA's), (3, 52094), (4, 52121), (5, 52136),

Gene: JEGGS_89 Start: 52026, Stop: 52334, Start Num: 2
Candidate Starts for JEGGS_89:
(1, 52023), (Start: 2 @52026 has 8 MA's), (3, 52173), (4, 52200), (5, 52215),

Gene: Kardesai_89 Start: 51430, Stop: 51738, Start Num: 2
Candidate Starts for Kardesai_89:
(1, 51427), (Start: 2 @51430 has 8 MA's), (3, 51577), (4, 51604), (5, 51619),

Gene: Mooshroom_89 Start: 51430, Stop: 51738, Start Num: 2
Candidate Starts for Mooshroom_89:
(1, 51427), (Start: 2 @51430 has 8 MA's), (3, 51577), (4, 51604), (5, 51619),

Gene: Mudcat_85 Start: 53181, Stop: 53489, Start Num: 2
Candidate Starts for Mudcat_85:
(1, 53178), (Start: 2 @53181 has 8 MA's), (3, 53328), (4, 53355), (5, 53370),

Gene: Tribby_92 Start: 52817, Stop: 53125, Start Num: 2
Candidate Starts for Tribby_92:
(1, 52814), (Start: 2 @52817 has 8 MA's), (3, 52964), (4, 52991), (5, 53006),