



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 106805 Report

This analysis was run 04/05/24 on database version 557.

Pham number 106805 has 10 members, 0 are drafts.

Phages represented in each track:

- Track 1 : Cheesy_2, Tribby_2, Heisenberger_2, Circum_2, NapoleonB_2, Dynamite_2, JEGGS_2, Correa_2, Mudcat_2
- Track 2 : BenitoAntonio_2

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 10 of the 10 non-draft genes in the pham.

Genes that call this "Most Annotated" start:

- BenitoAntonio_2, Cheesy_2, Circum_2, Correa_2, Dynamite_2, Heisenberger_2, JEGGS_2, Mudcat_2, NapoleonB_2, Tribby_2,

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: 10 of 10
- Called 100.0% of time when present
- Phage (with cluster) where this start called: BenitoAntonio_2 (AM), Cheesy_2 (AM), Circum_2 (AM), Correa_2 (AM), Dynamite_2 (AM), Heisenberger_2 (AM), JEGGS_2 (AM), Mudcat_2 (AM), NapoleonB_2 (AM), Tribby_2 (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 10 times for cluster AM.

Gene Information:

Gene: BenitoAntonio_2 Start: 567, Stop: 334, Start Num: 2

Candidate Starts for BenitoAntonio_2:

(1, 657), (Start: 2 @567 has 10 MA's), (3, 555), (4, 549), (5, 498), (6, 363),

Gene: Cheesy_2 Start: 569, Stop: 372, Start Num: 2

Candidate Starts for Cheesy_2:

(1, 659), (Start: 2 @569 has 10 MA's), (3, 557), (4, 551), (5, 500),

Gene: Circum_2 Start: 569, Stop: 372, Start Num: 2

Candidate Starts for Circum_2:

(1, 659), (Start: 2 @569 has 10 MA's), (3, 557), (4, 551), (5, 500),

Gene: Correa_2 Start: 569, Stop: 372, Start Num: 2

Candidate Starts for Correa_2:

(1, 659), (Start: 2 @569 has 10 MA's), (3, 557), (4, 551), (5, 500),

Gene: Dynamite_2 Start: 569, Stop: 372, Start Num: 2

Candidate Starts for Dynamite_2:

(1, 659), (Start: 2 @569 has 10 MA's), (3, 557), (4, 551), (5, 500),

Gene: Heisenberger_2 Start: 569, Stop: 372, Start Num: 2

Candidate Starts for Heisenberger_2:

(1, 659), (Start: 2 @569 has 10 MA's), (3, 557), (4, 551), (5, 500),

Gene: JEGGS_2 Start: 571, Stop: 374, Start Num: 2

Candidate Starts for JEGGS_2:

(1, 661), (Start: 2 @571 has 10 MA's), (3, 559), (4, 553), (5, 502),

Gene: Mudcat_2 Start: 568, Stop: 371, Start Num: 2

Candidate Starts for Mudcat_2:

(1, 658), (Start: 2 @568 has 10 MA's), (3, 556), (4, 550), (5, 499),

Gene: NapoleonB_2 Start: 569, Stop: 372, Start Num: 2

Candidate Starts for NapoleonB_2:

(1, 659), (Start: 2 @569 has 10 MA's), (3, 557), (4, 551), (5, 500),

Gene: Tribby_2 Start: 569, Stop: 372, Start Num: 2

Candidate Starts for Tribby_2:

(1, 659), (Start: 2 @569 has 10 MA's), (3, 557), (4, 551), (5, 500),