

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 222213 Report

This analysis was run 03/28/25 on database version 593.

Pham number 222213 has 6 members, 0 are drafts.

Phages represented in each track:

Track 1 : Indlulamithi_105Track 2 : MrMiyagi_115

Track 3 : Fowlmouth_116

Track 4 : Predator_6Track 5 : DrLupo_5

Track 6 : Barnyard_5

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 3 of the 6 non-draft genes in the pham.

Genes that call this "Most Annotated" start:

• Fowlmouth_116, Indlulamithi_105, MrMiyagi_115,

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

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

Barnyard_5, DrLupo_5, Predator_6,

Summary by start number:

Start 2:

- Found in 3 of 6 (50.0%) of genes in pham
- Manual Annotations of this start: 3 of 6
- Called 100.0% of time when present
- Phage (with cluster) where this start called: Fowlmouth_116 (AC), Indlulamithi_105 (AC), MrMiyagi_115 (AC),

Start 3:

- Found in 2 of 6 (33.3%) of genes in pham
- Manual Annotations of this start: 2 of 6
- Called 100.0% of time when present

Phage (with cluster) where this start called: Barnyard_5 (H2), Predator_6 (H1),

Start 4:

- Found in 1 of 6 (16.7%) of genes in pham
- Manual Annotations of this start: 1 of 6
- Called 100.0% of time when present
- Phage (with cluster) where this start called: DrLupo_5 (H2),

Summary by clusters:

There are 3 clusters represented in this pham: H2, AC, H1,

Info for manual annotations of cluster AC:

•Start number 2 was manually annotated 3 times for cluster AC.

Info for manual annotations of cluster H1:

•Start number 3 was manually annotated 1 time for cluster H1.

Info for manual annotations of cluster H2:

- •Start number 3 was manually annotated 1 time for cluster H2.
- •Start number 4 was manually annotated 1 time for cluster H2.

Gene Information:

Gene: Barnyard_5 Start: 2336, Stop: 2662, Start Num: 3

Candidate Starts for Barnyard_5:

(Start: 3 @2336 has 2 MA's), (6, 2396), (14, 2513), (17, 2534), (20, 2552), (22, 2561), (25, 2597), (30, 2651),

Gene: DrLupo 5 Start: 2304, Stop: 2633, Start Num: 4

Candidate Starts for DrLupo 5:

(Start: 4 @ 2304 has 1 MA's), (5, 2361), (8, 2394), (12, 2457), (14, 2481), (17, 2502), (26, 2571), (28, 2604), (30, 2622),

Gene: Fowlmouth_116 Start: 66878, Stop: 67264, Start Num: 2

Candidate Starts for Fowlmouth_116:

(1, 66869), (Start: 2 @66878 has 3 MA's), (6, 66941), (11, 67031), (15, 67061), (16, 67067), (18, 67088), (19, 67115), (21, 67127), (24, 67142), (27, 67202), (29, 67214),

Gene: Indlulamithi_105 Start: 68264, Stop: 68617, Start Num: 2

Candidate Starts for Indlulamithi 105:

(Start: 2 @68264 has 3 MA's), (5, 68324), (7, 68351), (10, 68402), (11, 68417), (13, 68432), (21, 68489), (31, 68600), (32, 68603),

Gene: MrMiyagi_115 Start: 67159, Stop: 67545, Start Num: 2

Candidate Starts for MrMiyagi 115:

(1, 67150), (Start: 2 @67159 has 3 MA's), (6, 67222), (11, 67312), (15, 67342), (16, 67348), (18, 67369), (19, 67396), (21, 67408), (24, 67423), (29, 67495),

Gene: Predator 6 Start: 2730, Stop: 3125, Start Num: 3

Candidate Starts for Predator_6:

(Start: 3 @2730 has 2 MA's), (9, 2826), (12, 2883), (13, 2895), (19, 2961), (21, 2973), (23, 2982),