



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 107289 Report

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

Pham number 107289 has 6 members, 1 are drafts.

Phages represented in each track:

Track 1 : Blarby_50

Track 2: Estave1_39, Renaud18_42, Taj_41, DeadP_42, Wee_42

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 5 of the 5 non-draft genes in the pham.

Genes that call this "Most Annotated" start:

DeadP_42, Estave1_39, Renaud18_42, Taj_41, Wee_42,

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

Blarby_50,

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

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Summary by start number:

Start 1:

- Found in 6 of 6 (100.0%) of genes in pham
- Manual Annotations of this start: 5 of 5
- Called 83.3% of time when present
- Phage (with cluster) where this start called: DeadP_42 (F1), Estave1_39 (F1), Renaud18_42 (F4), Tai_41 (F1), Wee_42 (F1),

Start 3:

- Found in 6 of 6 (100.0%) of genes in pham
- No Manual Annotations of this start.
- Called 16.7% of time when present
- Phage (with cluster) where this start called: Blarby_50 (F),

Summary by clusters:

There are 3 clusters represented in this pham: F1, F4, F,

Info for manual annotations of cluster F1:

•Start number 1 was manually annotated 4 times for cluster F1.

Info for manual annotations of cluster F4:

•Start number 1 was manually annotated 1 time for cluster F4.

Gene Information:

Gene: Blarby_50 Start: 31964, Stop: 31815, Start Num: 3

Candidate Starts for Blarby_50:

(Start: 1 @31985 has 5 MA's), (2, 31970), (3, 31964), (4, 31940), (5, 31931), (6, 31928), (7, 31898),

Gene: DeadP_42 Start: 31682, Stop: 31512, Start Num: 1

Candidate Starts for DeadP_42:

(Start: 1 @31682 has 5 MA's), (2, 31667), (3, 31661), (4, 31637), (5, 31628), (6, 31625), (7, 31595),

Gene: Estave1 39 Start: 30985, Stop: 30815, Start Num: 1

Candidate Starts for Estave1_39:

(Start: 1 @ 30985 has 5 MA's), (2, 30970), (3, 30964), (4, 30940), (5, 30931), (6, 30928), (7, 30898),

Gene: Renaud18 42 Start: 31761, Stop: 31591, Start Num: 1

Candidate Starts for Renaud18 42:

(Start: 1 @31761 has 5 MA's), (2, 31746), (3, 31740), (4, 31716), (5, 31707), (6, 31704), (7, 31674),

Gene: Taj_41 Start: 31684, Stop: 31514, Start Num: 1

Candidate Starts for Taj_41:

(Start: 1 @31684 has 5 MA's), (2, 31669), (3, 31663), (4, 31639), (5, 31630), (6, 31627), (7, 31597),

Gene: Wee 42 Start: 32258, Stop: 32088, Start Num: 1

Candidate Starts for Wee 42:

(Start: 1 @32258 has 5 MA's), (2, 32243), (3, 32237), (4, 32213), (5, 32204), (6, 32201), (7, 32171),