

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 134256 Report

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

Pham number 134256 has 9 members, 2 are drafts.

Phages represented in each track:

Track 1 : E1_48, Anatole_48, Doucette_54

Track 2: B3_50
Track 3: G4_65
Track 4: E6_54
Track 5: B22 57

• Track 6: PFR1 52, PFR2 54

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

The start number called the most often in the published annotations is 4, it was called in 7 of the 7 non-draft genes in the pham.

Genes that call this "Most Annotated" start:

• Anatole_48, B22_57, B3_50, Doucette_54, E1_48, E6_54, G4_65, PFR1_52, PFR2_54.

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:

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Summary by start number:

Start 4:

- Found in 9 of 9 (100.0%) of genes in pham
- Manual Annotations of this start: 7 of 7
- Called 100.0% of time when present
- Phage (with cluster) where this start called: Anatole_48 (BV), B22_57 (BW), B3_50 (BV), Doucette_54 (BW), E1_48 (BV), E6_54 (BW), G4_65 (BW), PFR1_52 (BX), PFR2_54 (BX),

Summary by clusters:

There are 3 clusters represented in this pham: BV, BW, BX,

Info for manual annotations of cluster BV:

Start number 4 was manually annotated 3 times for cluster BV.

Info for manual annotations of cluster BW:

Start number 4 was manually annotated 4 times for cluster BW.

Gene Information:

Gene: Anatole_48 Start: 31658, Stop: 31888, Start Num: 4

Candidate Starts for Anatole 48:

(Start: 4 @31658 has 7 MA's), (5, 31724), (6, 31730), (7, 31757), (8, 31772), (9, 31784), (10, 31796),

Gene: B22 57 Start: 35089, Stop: 35319, Start Num: 4

Candidate Starts for B22 57:

(1, 34864), (Start: 4 @ 35089 has 7 MA's), (5, 35155), (7, 35188), (8, 35203), (10, 35227), (13, 35311),

Gene: B3 50 Start: 32321, Stop: 32551, Start Num: 4

Candidate Starts for B3 50:

(Start: 4 @32321 has 7 MA's), (5, 32387), (6, 32393), (7, 32420), (8, 32435), (10, 32459),

Gene: Doucette 54 Start: 34767, Stop: 34997, Start Num: 4

Candidate Starts for Doucette 54:

(Start: 4 @ 34767 has 7 MA's), (5, 34833), (6, 34839), (7, 34866), (8, 34881), (9, 34893), (10, 34905),

Gene: E1_48 Start: 31658, Stop: 31888, Start Num: 4

Candidate Starts for E1_48:

(Start: 4 @31658 has 7 MA's), (5, 31724), (6, 31730), (7, 31757), (8, 31772), (9, 31784), (10, 31796),

Gene: E6 54 Start: 34589, Stop: 34819, Start Num: 4

Candidate Starts for E6 54:

(1, 34364), (2, 34466), (3, 34511), (Start: 4 @ 34589 has 7 MA's), (5, 34655), (6, 34661), (7, 34688), (8, 34703), (9, 34715), (10, 34727), (11, 34769), (12, 34787),

Gene: G4 65 Start: 36425, Stop: 36655, Start Num: 4

Candidate Starts for G4 65:

(2, 36302), (Start: 4 @ 36425 has 7 MA's), (5, 36491), (6, 36497), (7, 36524), (8, 36539), (10, 36563),

Gene: PFR1 52 Start: 35760, Stop: 35990, Start Num: 4

Candidate Starts for PFR1_52:

(Start: 4 @ 35760 has 7 MA's), (5, 35826), (7, 35859), (8, 35874), (9, 35886), (10, 35898), (11, 35940),

Gene: PFR2_54 Start: 37329, Stop: 37559, Start Num: 4

Candidate Starts for PFR2 54:

(Start: 4 @ 37329 has 7 MA's), (5, 37395), (7, 37428), (8, 37443), (9, 37455), (10, 37467), (11, 37509),