

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 158501 Report

This analysis was run 04/13/24 on database version 558.

Pham number 158501 has 6 members, 0 are drafts.

Phages represented in each track:

Track 1: Mahdia_38
Track 2: Gustav_40
Track 3: Upyo_40
Track 4: Morrissey_39
Track 5: Trine_38

• Track 6 : Widow 38

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

The start number called the most often in the published annotations is 15, it was called in 5 of the 6 non-draft genes in the pham.

Genes that call this "Most Annotated" start:

• Gustav_40, Mahdia_38, Morrissey_39, Trine_38, Widow_38,

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:

Upyo_40,

Summary by start number:

Start 15:

- Found in 5 of 6 (83.3%) of genes in pham
- Manual Annotations of this start: 5 of 6
- Called 100.0% of time when present
- Phage (with cluster) where this start called: Gustav_40 (CD), Mahdia_38 (CD), Morrissey_39 (CD), Trine_38 (CD), Widow_38 (CD),

Start 16:

- 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: Upyo_40 (CD),

Summary by clusters:

There is one cluster represented in this pham: CD

Info for manual annotations of cluster CD:

- •Start number 15 was manually annotated 5 times for cluster CD.
- •Start number 16 was manually annotated 1 time for cluster CD.

Gene Information:

Gene: Gustav_40 Start: 31883, Stop: 31605, Start Num: 15

Candidate Starts for Gustav_40:

(9, 32087), (10, 32084), (Start: 15 @ 31883 has 5 MA's), (21, 31640),

Gene: Mahdia_38 Start: 31109, Stop: 30855, Start Num: 15

Candidate Starts for Mahdia_38:

(1, 31604), (2, 31586), (3, 31484), (6, 31367), (8, 31355), (9, 31325), (10, 31322), (11, 31289), (12, 31586), (13, 31484), (13, 31484), (14, 31367), (15, 31385), (15, 31386), (15, 31886), (15, 3188

31250), (Start: 15 @31109 has 5 MA's), (19, 30938), (20, 30866),

Gene: Morrissey_39 Start: 31553, Stop: 31275, Start Num: 15

Candidate Starts for Morrissey_39:

(9, 31757), (10, 31754), (Start: 15 @31553 has 5 MA's), (18, 31430), (20, 31313),

Gene: Trine_38 Start: 31043, Stop: 30771, Start Num: 15

Candidate Starts for Trine_38:

(Start: 15 @31043 has 5 MA's), (18, 30914),

Gene: Upyo 40 Start: 32017, Stop: 31766, Start Num: 16

Candidate Starts for Upyo 40:

(4, 32392), (5, 32371), (13, 32068), (14, 32056), (Start: 16 @32017 has 1 MA's), (17, 31960), (20, 31804),

Gene: Widow_38 Start: 31430, Stop: 31161, Start Num: 15

Candidate Starts for Widow 38:

(6, 31676), (7, 31667), (9, 31634), (Start: 15 @31430 has 5 MA's),