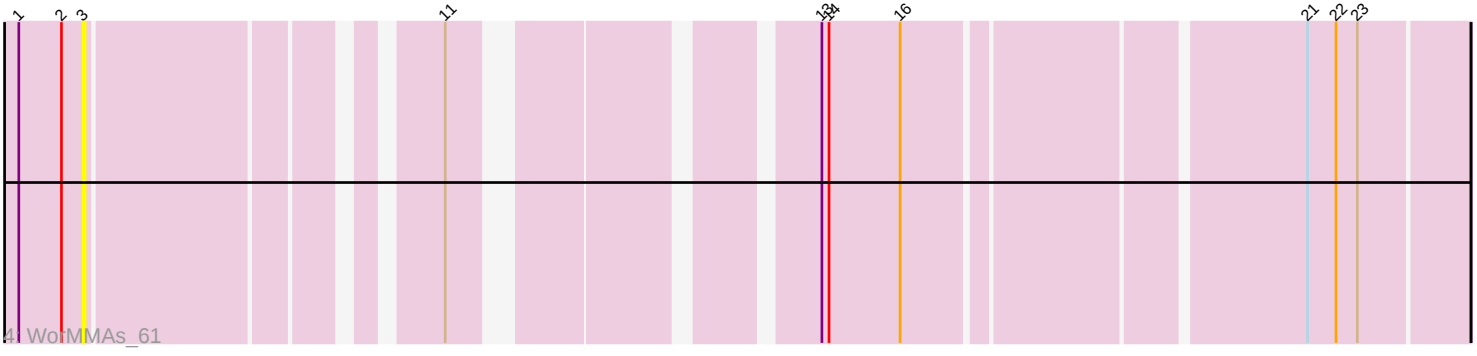
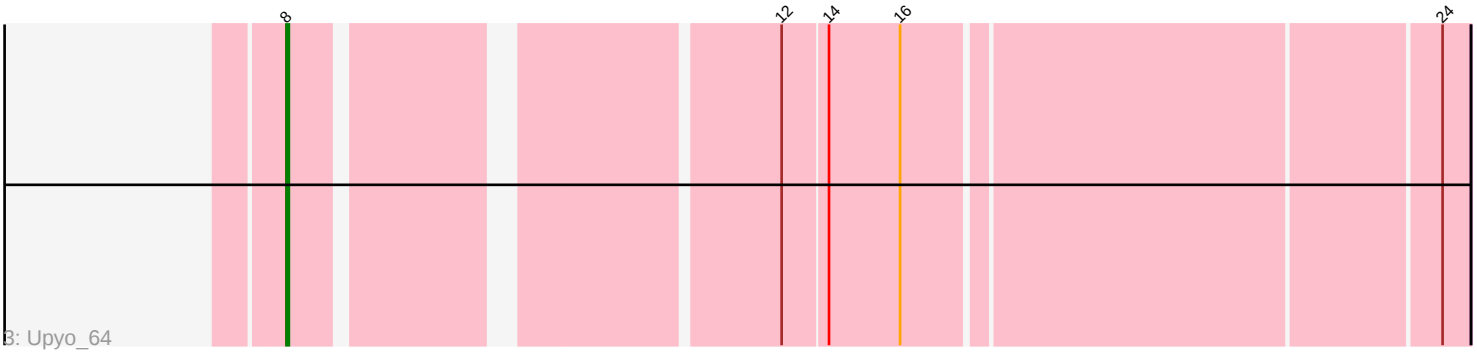
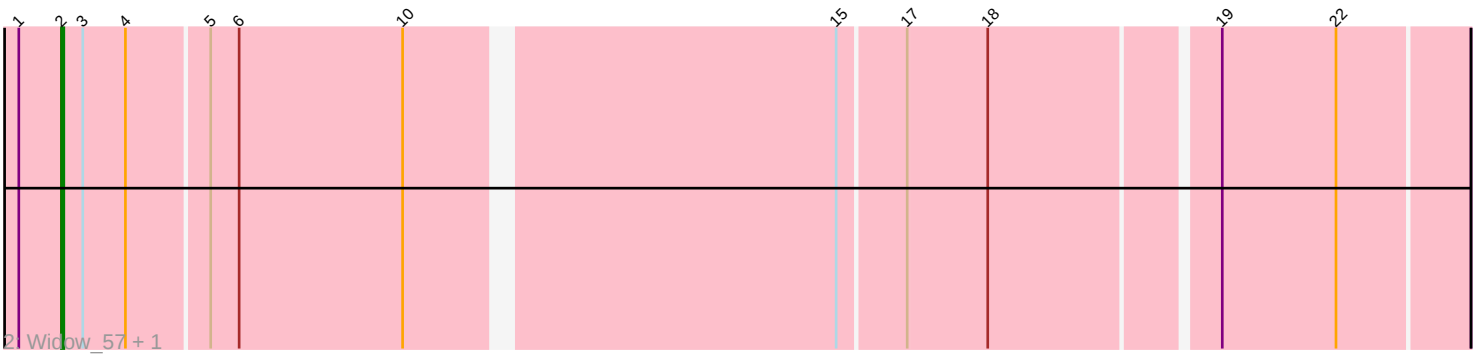
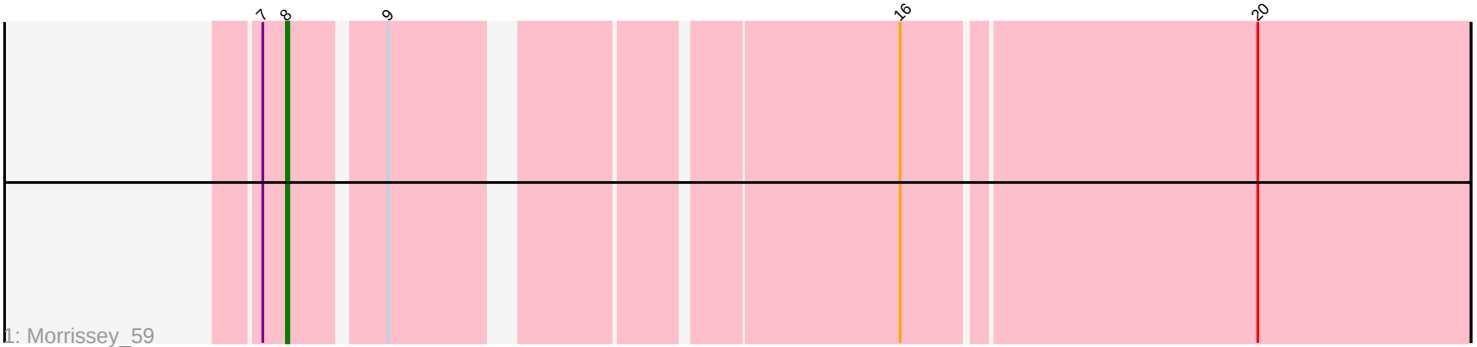


Pham 309944



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 309944 Report

This analysis was run 09/05/26 on database version 662.

Pham number 309944 has 5 members, 1 are drafts.

Phages represented in each track:

- Track 1 : Morrissey_59
- Track 2 : Widow_57, Puppies_57
- Track 3 : Upyo_64
- Track 4 : WorMMAs_61

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 2 of the 4 non-draft genes in the pham.

Genes that call this "Most Annotated" start:

- Puppies_57, Widow_57,

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

- WorMMAs_61,

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

- Morrissey_59, Upyo_64,

Summary by start number:

Start 2:

- Found in 3 of 5 (60.0%) of genes in pham
- Manual Annotations of this start: 2 of 4
- Called 66.7% of time when present
- Phage (with cluster) where this start called: Puppies_57 (CD), Widow_57 (CD),

Start 3:

- Found in 3 of 5 (60.0%) of genes in pham
- No Manual Annotations of this start.
- Called 33.3% of time when present
- Phage (with cluster) where this start called: WorMMAs_61 (UNK),

Start 8:

- Found in 2 of 5 (40.0%) of genes in pham
- Manual Annotations of this start: 2 of 4
- Called 100.0% of time when present
- Phage (with cluster) where this start called: Morrissey_59 (CD), Upyo_64 (CD),

Summary by clusters:

There are 2 clusters represented in this pham: UNK, CD,

Info for manual annotations of cluster CD:

- Start number 2 was manually annotated 2 times for cluster CD.
- Start number 8 was manually annotated 2 times for cluster CD.

Gene Information:

Gene: Morrissey_59 Start: 42085, Stop: 42543, Start Num: 8

Candidate Starts for Morrissey_59:

(7, 42076), (Start: 8 @42085 has 2 MA's), (9, 42121), (16, 42310), (20, 42454),

Gene: Puppers_57 Start: 39976, Stop: 40542, Start Num: 2

Candidate Starts for Puppers_57:

(1, 39958), (Start: 2 @39976 has 2 MA's), (3, 39985), (4, 40003), (5, 40036), (6, 40048), (10, 40114), (15, 40291), (17, 40318), (18, 40351), (19, 40441), (22, 40489),

Gene: Upyo_64 Start: 43304, Stop: 43759, Start Num: 8

Candidate Starts for Upyo_64:

(Start: 8 @43304 has 2 MA's), (12, 43484), (14, 43502), (16, 43532), (24, 43748),

Gene: Widow_57 Start: 40525, Stop: 41088, Start Num: 2

Candidate Starts for Widow_57:

(1, 40507), (Start: 2 @40525 has 2 MA's), (3, 40534), (4, 40552), (5, 40585), (6, 40597), (10, 40666), (15, 40837), (17, 40864), (18, 40897), (19, 40987), (22, 41035),

Gene: WorMMAs_61 Start: 40675, Stop: 41178, Start Num: 3

Candidate Starts for WorMMAs_61:

(1, 40648), (Start: 2 @40666 has 2 MA's), (3, 40675), (11, 40801), (13, 40924), (14, 40927), (16, 40957), (21, 41113), (22, 41125), (23, 41134),