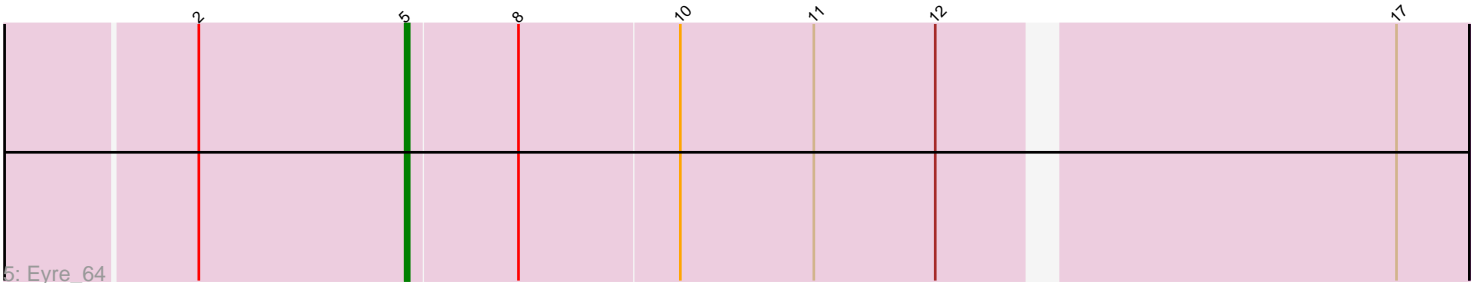
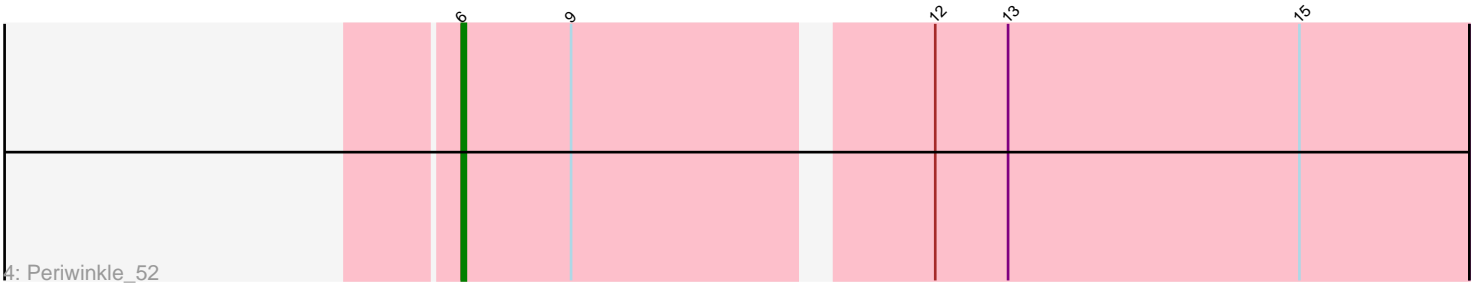
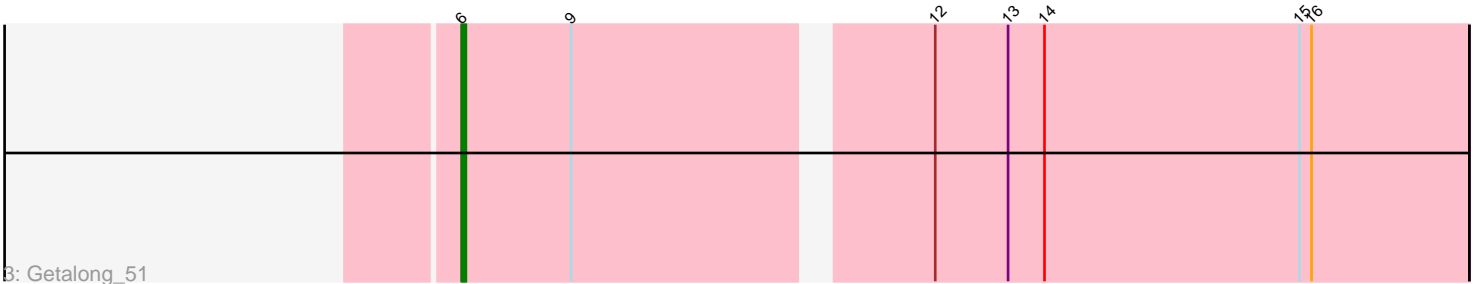
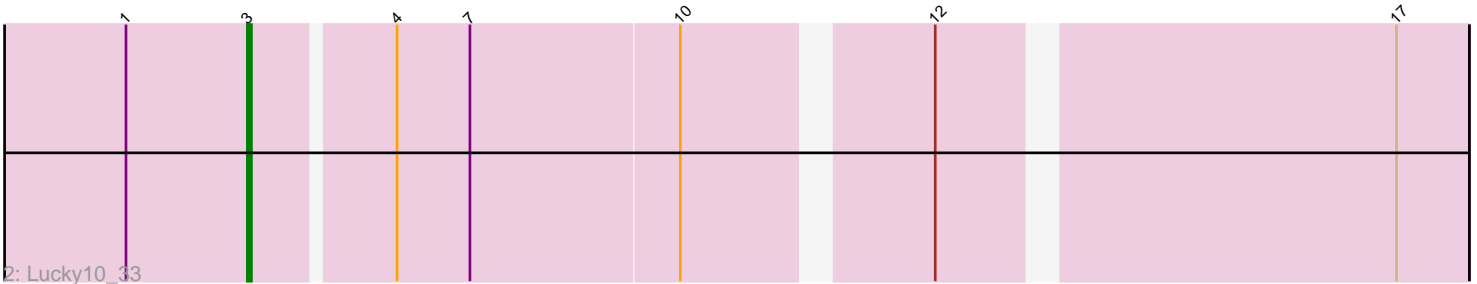
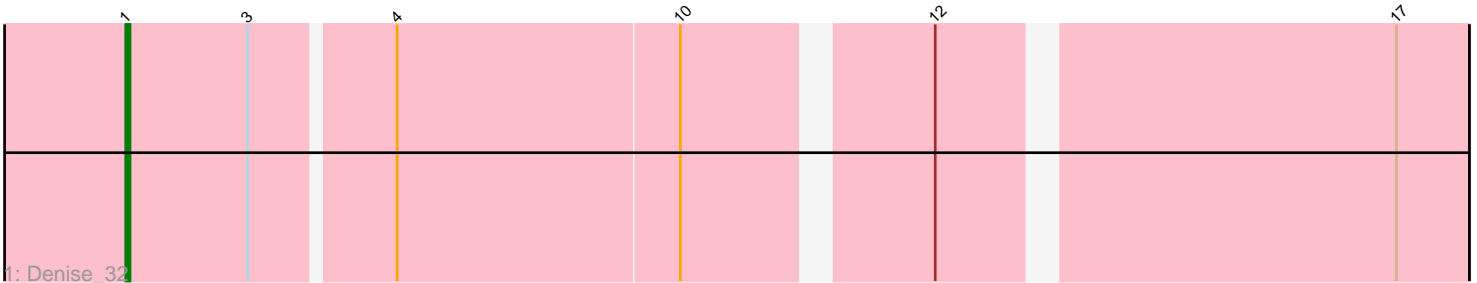


Pham 88683



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 88683 Report

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

Pham number 88683 has 5 members, 0 are drafts.

Phages represented in each track:

- Track 1 : Denise_32
- Track 2 : Lucky10_33
- Track 3 : Getalong_51
- Track 4 : Periwinkle_52
- Track 5 : Eyre_64

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

The start number called the most often in the published annotations is 6, it was called in 2 of the 5 non-draft genes in the pham.

Genes that call this "Most Annotated" start:

- Getalong_51, Periwinkle_52,

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

-

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

- Denise_32, Eyre_64, Lucky10_33,

Summary by start number:

Start 1:

- Found in 2 of 5 (40.0%) of genes in pham
- Manual Annotations of this start: 1 of 5
- Called 50.0% of time when present
- Phage (with cluster) where this start called: Denise_32 (CZ5),

Start 3:

- Found in 2 of 5 (40.0%) of genes in pham
- Manual Annotations of this start: 1 of 5
- Called 50.0% of time when present
- Phage (with cluster) where this start called: Lucky10_33 (DH),

Start 5:

- Found in 1 of 5 (20.0%) of genes in pham
- Manual Annotations of this start: 1 of 5
- Called 100.0% of time when present
- Phage (with cluster) where this start called: Eyre_64 (singleton),

Start 6:

- Found in 2 of 5 (40.0%) of genes in pham
- Manual Annotations of this start: 2 of 5
- Called 100.0% of time when present
- Phage (with cluster) where this start called: Getalong_51 (DN1), Periwinkle_52 (DN1),

Summary by clusters:

There are 4 clusters represented in this pham: singleton, DN1, DH, CZ5,

Info for manual annotations of cluster CZ5:

- Start number 1 was manually annotated 1 time for cluster CZ5.

Info for manual annotations of cluster DH:

- Start number 3 was manually annotated 1 time for cluster DH.

Info for manual annotations of cluster DN1:

- Start number 6 was manually annotated 2 times for cluster DN1.

Gene Information:

Gene: Denise_32 Start: 26430, Stop: 26738, Start Num: 1

Candidate Starts for Denise_32:

(Start: 1 @26430 has 1 MA's), (Start: 3 @26460 has 1 MA's), (4, 26493), (10, 26562), (12, 26616), (17, 26721),

Gene: Eyre_64 Start: 41393, Stop: 41644, Start Num: 5

Candidate Starts for Eyre_64:

(2, 41342), (Start: 5 @41393 has 1 MA's), (8, 41420), (10, 41459), (11, 41492), (12, 41522), (17, 41627),

Gene: Getalong_51 Start: 33873, Stop: 34112, Start Num: 6

Candidate Starts for Getalong_51:

(Start: 6 @33873 has 2 MA's), (9, 33900), (12, 33981), (13, 33999), (14, 34008), (15, 34071), (16, 34074),

Gene: Lucky10_33 Start: 27622, Stop: 27900, Start Num: 3

Candidate Starts for Lucky10_33:

(Start: 1 @27592 has 1 MA's), (Start: 3 @27622 has 1 MA's), (4, 27655), (7, 27673), (10, 27724), (12, 27778), (17, 27883),

Gene: Periwinkle_52 Start: 33407, Stop: 33646, Start Num: 6

Candidate Starts for Periwinkle_52:

(Start: 6 @33407 has 2 MA's), (9, 33434), (12, 33515), (13, 33533), (15, 33605),

