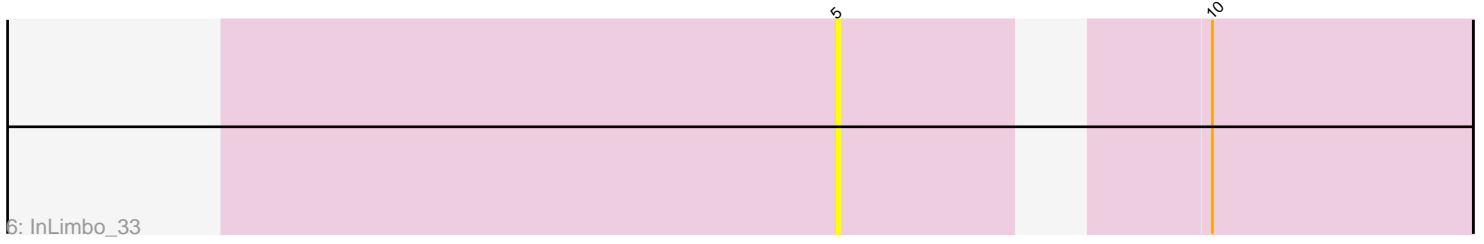
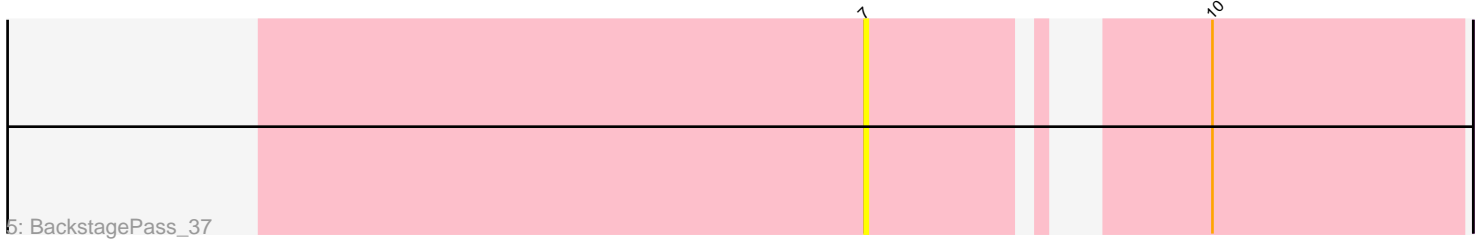
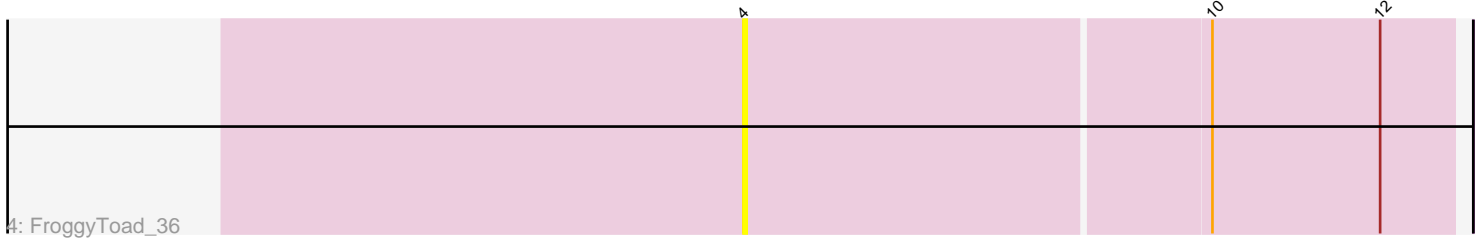
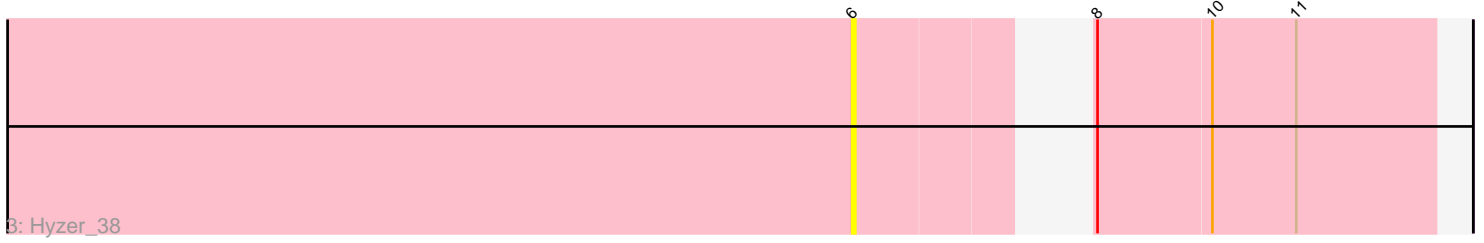
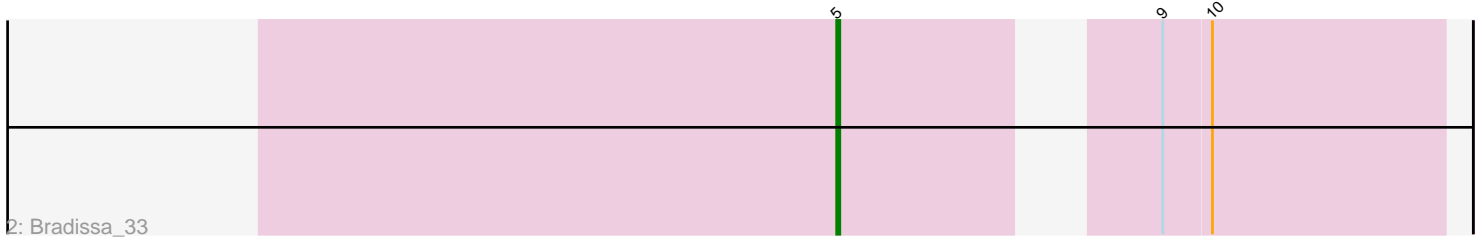
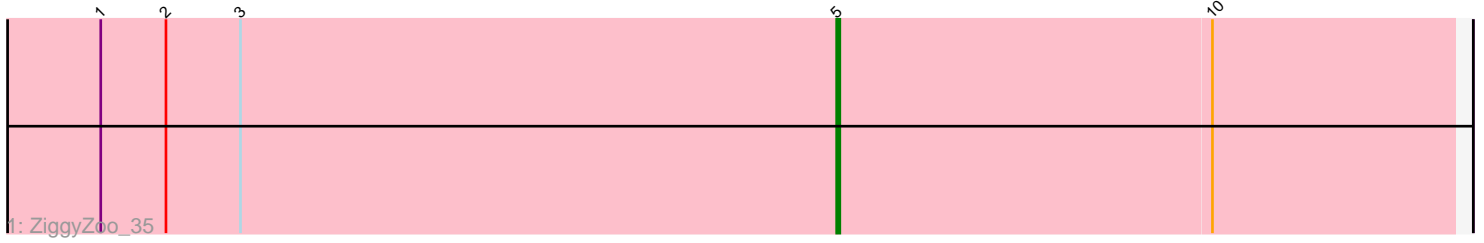


Pham 297398



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 297398 Report

This analysis was run 04/25/26 on database version 644.

Pham number 297398 has 6 members, 4 are drafts.

Phages represented in each track:

- Track 1 : ZiggyZoo_35
- Track 2 : Bradissa_33
- Track 3 : Hyzer_38
- Track 4 : FroggyToad_36
- Track 5 : BackstagePass_37
- Track 6 : InLimbo_33

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

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

Genes that call this "Most Annotated" start:

- Bradissa_33, InLimbo_33, ZiggyZoo_35,

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

-

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

- BackstagePass_37, FroggyToad_36, Hyzer_38,

Summary by start number:

Start 4:

- Found in 1 of 6 (16.7%) of genes in pham
- No Manual Annotations of this start.
- Called 100.0% of time when present
- Phage (with cluster) where this start called: FroggyToad_36 (CZ2),

Start 5:

- Found in 3 of 6 (50.0%) of genes in pham
- Manual Annotations of this start: 2 of 2
- Called 100.0% of time when present

- Phage (with cluster) where this start called: Bradissa_33 (CY1), InLimbo_33 (DH), ZiggyZoo_35 (CV),

Start 6:

- Found in 1 of 6 (16.7%) of genes in pham
- No Manual Annotations of this start.
- Called 100.0% of time when present
- Phage (with cluster) where this start called: Hyzer_38 (CZ1),

Start 7:

- Found in 1 of 6 (16.7%) of genes in pham
- No Manual Annotations of this start.
- Called 100.0% of time when present
- Phage (with cluster) where this start called: BackstagePass_37 (DB),

Summary by clusters:

There are 6 clusters represented in this pham: CY1, DH, CZ2, CZ1, DB, CV,

Info for manual annotations of cluster CV:

- Start number 5 was manually annotated 1 time for cluster CV.

Info for manual annotations of cluster CY1:

- Start number 5 was manually annotated 1 time for cluster CY1.

Gene Information:

Gene: BackstagePass_37 Start: 29116, Stop: 29283, Start Num: 7

Candidate Starts for BackstagePass_37:

(7, 29116), (10, 29203),

Gene: Bradissa_33 Start: 28417, Stop: 28587, Start Num: 5

Candidate Starts for Bradissa_33:

(Start: 5 @28417 has 2 MA's), (9, 28498), (10, 28513),

Gene: FroggyToad_36 Start: 27661, Stop: 27885, Start Num: 4

Candidate Starts for FroggyToad_36:

(4, 27661), (10, 27808), (12, 27862),

Gene: Hyzer_38 Start: 31750, Stop: 31908, Start Num: 6

Candidate Starts for Hyzer_38:

(6, 31750), (8, 31801), (10, 31837), (11, 31864),

Gene: InLimbo_33 Start: 25611, Stop: 25790, Start Num: 5

Candidate Starts for InLimbo_33:

(Start: 5 @25611 has 2 MA's), (10, 25707),

Gene: ZiggyZoo_35 Start: 28894, Stop: 29091, Start Num: 5

Candidate Starts for ZiggyZoo_35:

(1, 28657), (2, 28678), (3, 28702), (Start: 5 @28894 has 2 MA's), (10, 29014),