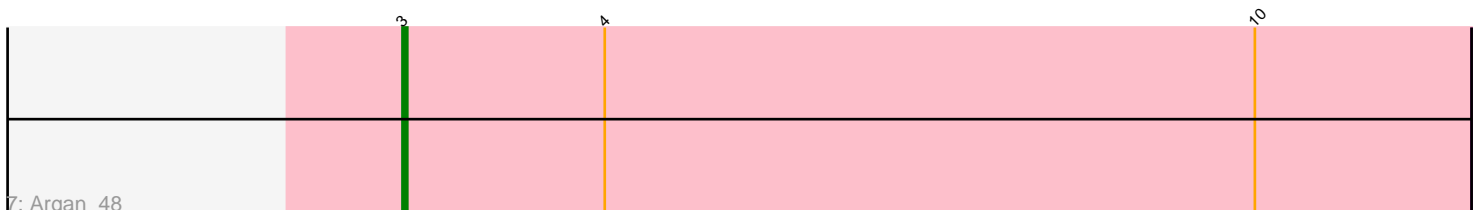
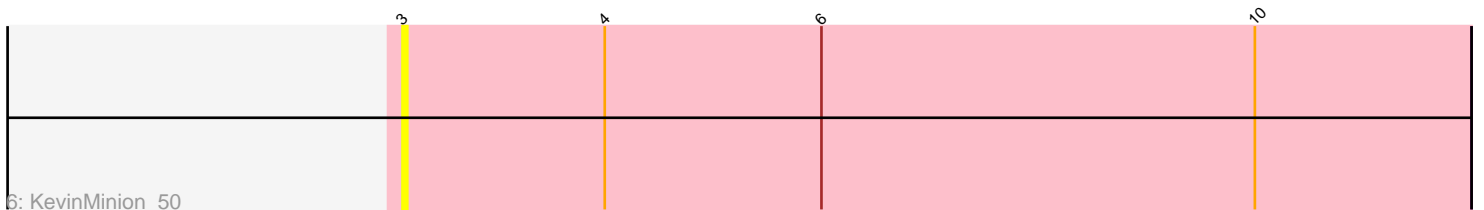
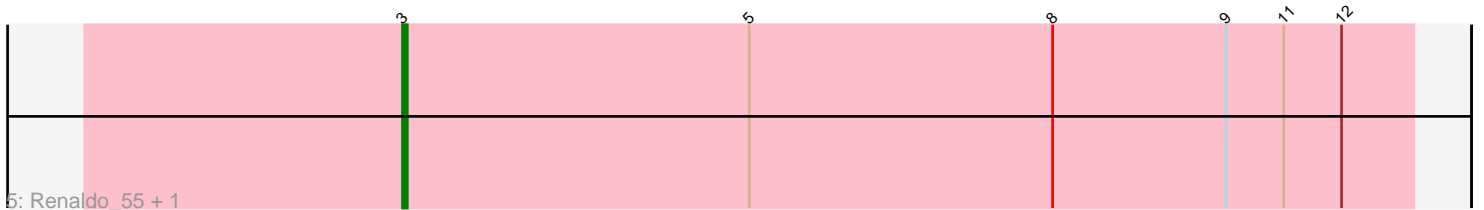
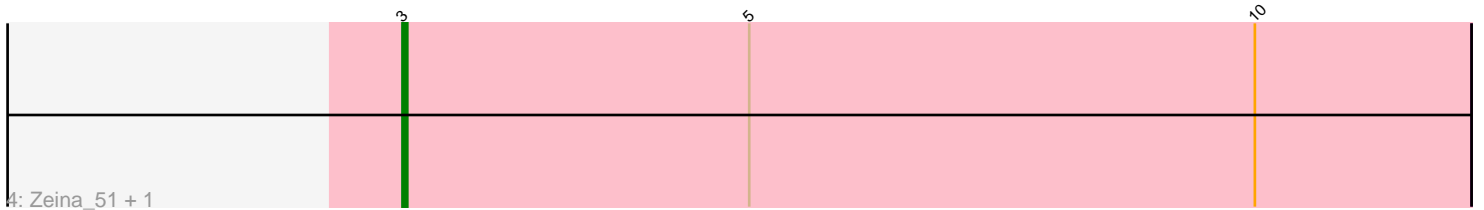
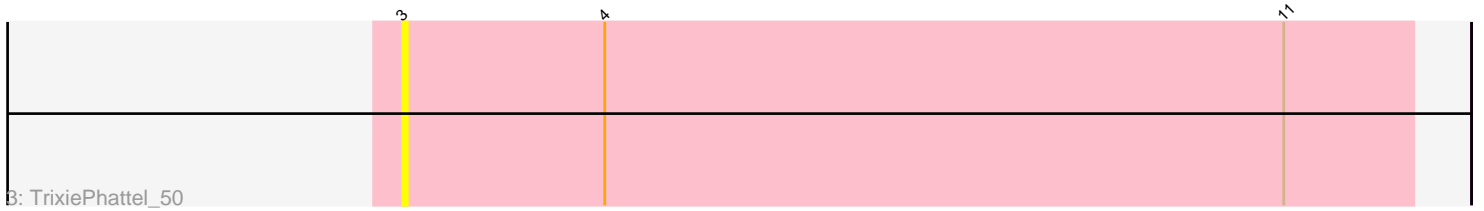
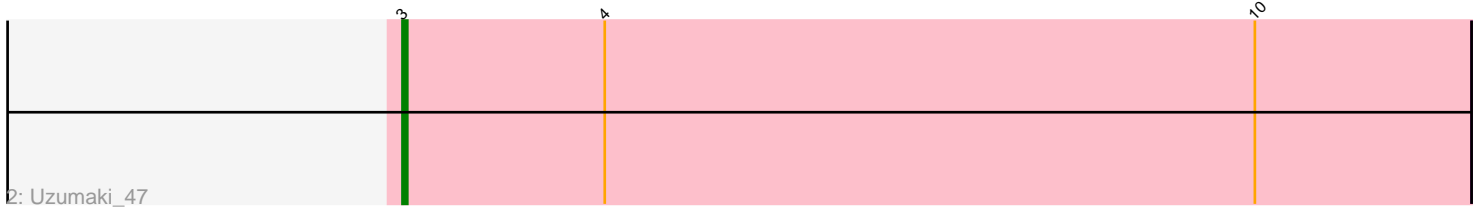
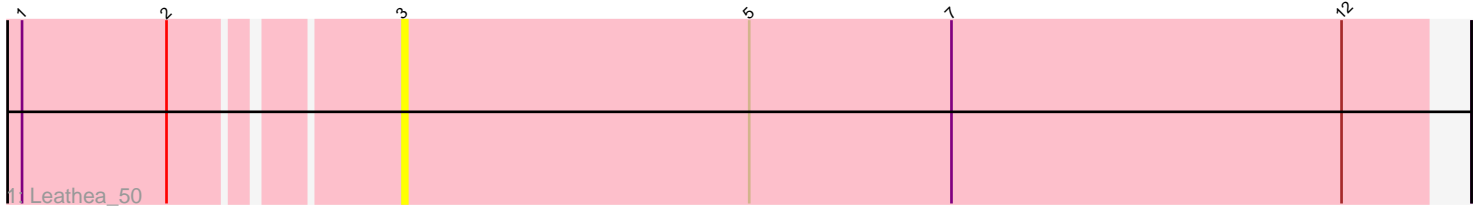


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Note: Tracks are now grouped by subcluster and scaled. Switching in subcluster is indicated by changes in track color. Track scale is now set by default to display the region 30 bp upstream of start 1 to 30 bp downstream of the last possible start. 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 225108 Report

This analysis was run 03/28/25 on database version 593.

Pham number 225108 has 9 members, 5 are drafts.

Phages represented in each track:

- Track 1 : Leathea\_50
- Track 2 : Uzumaki\_47
- Track 3 : TrixiePhattel\_50
- Track 4 : Zeina\_51, Tenney120\_52
- Track 5 : Renaldo\_55, Lewando\_52
- Track 6 : KevinMinion\_50
- Track 7 : Argan\_48

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

The start number called the most often in the published annotations is 3, it was called in 4 of the 4 non-draft genes in the pham.

Genes that call this "Most Annotated" start:

- Argan\_48, KevinMinion\_50, Leathea\_50, Lewando\_52, Renaldo\_55, Tenney120\_52, TrixiePhattel\_50, Uzumaki\_47, Zeina\_51,

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

- 

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

- 

### **Summary by start number:**

Start 3:

- Found in 9 of 9 ( 100.0% ) of genes in pham
- Manual Annotations of this start: 4 of 4
- Called 100.0% of time when present
- Phage (with cluster) where this start called: Argan\_48 (AU6), KevinMinion\_50 (AU6), Leathea\_50 (AU6), Lewando\_52 (AU6), Renaldo\_55 (AU6), Tenney120\_52 (AU6), TrixiePhattel\_50 (AU6), Uzumaki\_47 (AU6), Zeina\_51 (AU6),

### **Summary by clusters:**

There is one cluster represented in this pham: AU6

Info for manual annotations of cluster AU6:

•Start number 3 was manually annotated 4 times for cluster AU6.

**Gene Information:**

Gene: Argan\_48 Start: 33040, Stop: 33261, Start Num: 3

Candidate Starts for Argan\_48:

(Start: 3 @33040 has 4 MA's), (4, 33082), (10, 33217),

Gene: KevinMinion\_50 Start: 33991, Stop: 34212, Start Num: 3

Candidate Starts for KevinMinion\_50:

(Start: 3 @33991 has 4 MA's), (4, 34033), (6, 34078), (10, 34168),

Gene: Leathea\_50 Start: 33107, Stop: 33319, Start Num: 3

Candidate Starts for Leathea\_50:

(1, 33035), (2, 33065), (Start: 3 @33107 has 4 MA's), (5, 33179), (7, 33221), (12, 33302),

Gene: Lewando\_52 Start: 34682, Stop: 34891, Start Num: 3

Candidate Starts for Lewando\_52:

(Start: 3 @34682 has 4 MA's), (5, 34754), (8, 34817), (9, 34853), (11, 34865), (12, 34877),

Gene: Renaldo\_55 Start: 34991, Stop: 35200, Start Num: 3

Candidate Starts for Renaldo\_55:

(Start: 3 @34991 has 4 MA's), (5, 35063), (8, 35126), (9, 35162), (11, 35174), (12, 35186),

Gene: Tenney120\_52 Start: 33775, Stop: 33996, Start Num: 3

Candidate Starts for Tenney120\_52:

(Start: 3 @33775 has 4 MA's), (5, 33847), (10, 33952),

Gene: TrixiePhattel\_50 Start: 33138, Stop: 33347, Start Num: 3

Candidate Starts for TrixiePhattel\_50:

(Start: 3 @33138 has 4 MA's), (4, 33180), (11, 33321),

Gene: Uzumaki\_47 Start: 33233, Stop: 33454, Start Num: 3

Candidate Starts for Uzumaki\_47:

(Start: 3 @33233 has 4 MA's), (4, 33275), (10, 33410),

Gene: Zeina\_51 Start: 33783, Stop: 34004, Start Num: 3

Candidate Starts for Zeina\_51:

(Start: 3 @33783 has 4 MA's), (5, 33855), (10, 33960),