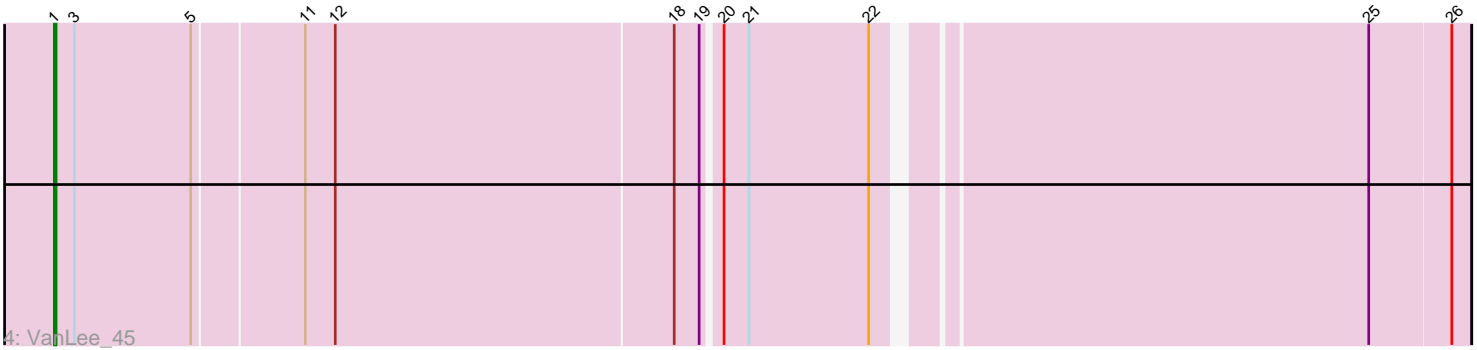
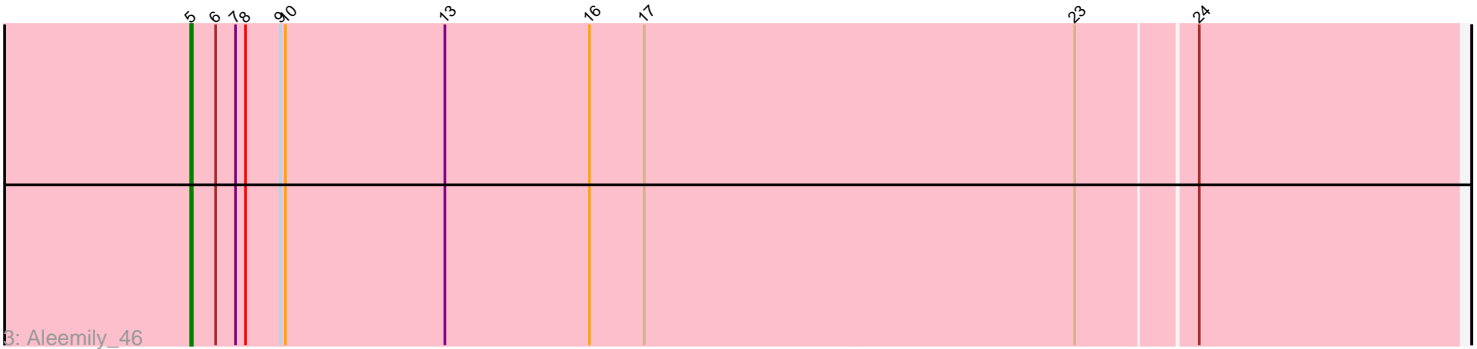
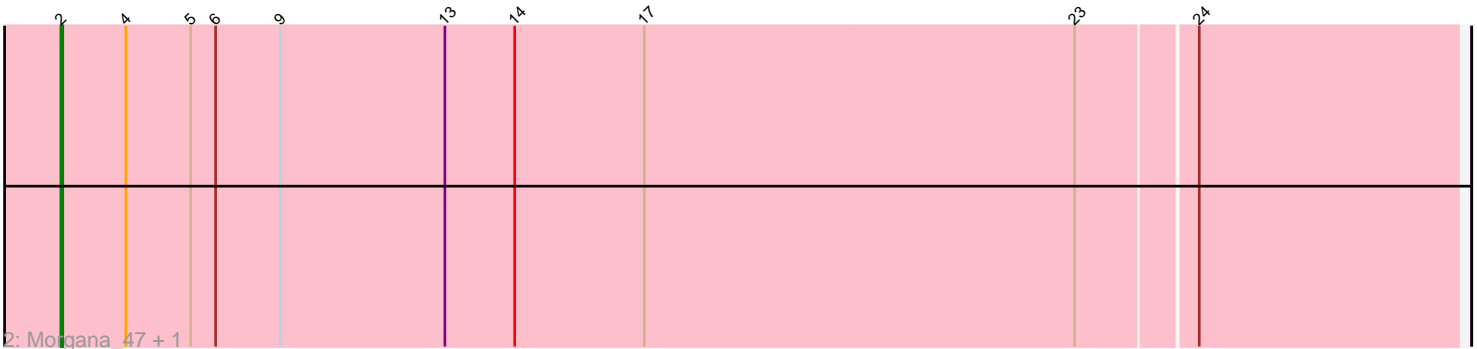
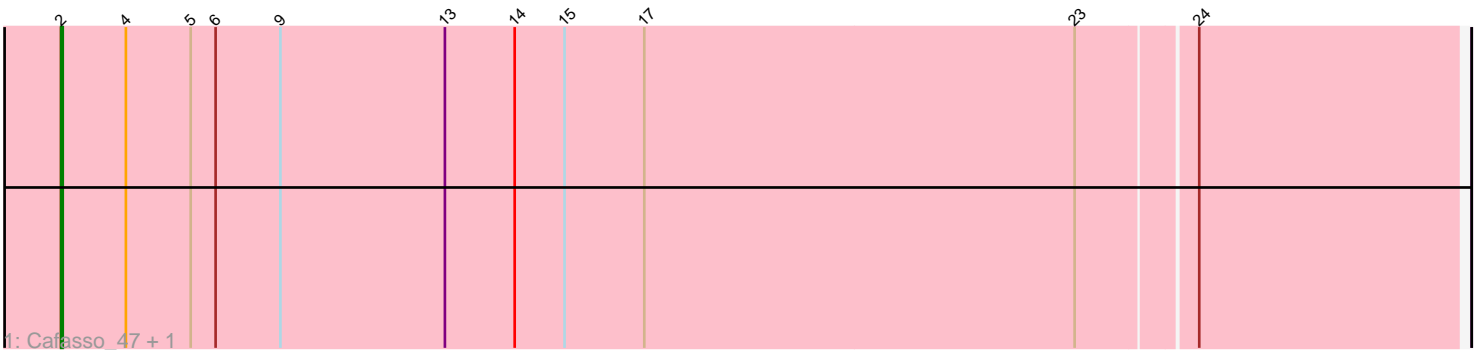


Pham 10305



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 10305 Report

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

Pham number 10305 has 6 members, 1 are drafts.

Phages represented in each track:

- Track 1 : Cafasso\_47, ModicumRichard\_47
- Track 2 : Morgana\_47, ObLaDi\_47
- Track 3 : Aleemily\_46
- Track 4 : VanLee\_45

### ***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 3 of the 5 non-draft genes in the pham.

Genes that call this "Most Annotated" start:

- Cafasso\_47, ModicumRichard\_47, Morgana\_47, ObLaDi\_47,

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

- 

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

- Aleemily\_46, VanLee\_45,

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

Start 1:

- Found in 1 of 6 ( 16.7% ) 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: VanLee\_45 (singleton),

Start 2:

- Found in 4 of 6 ( 66.7% ) of genes in pham
- Manual Annotations of this start: 3 of 5
- Called 100.0% of time when present
- Phage (with cluster) where this start called: Cafasso\_47 (DZ), ModicumRichard\_47 (DZ), Morgana\_47 (DZ), ObLaDi\_47 (DZ),

Start 5:

- Found in 6 of 6 ( 100.0% ) of genes in pham
- Manual Annotations of this start: 1 of 5
- Called 16.7% of time when present
- Phage (with cluster) where this start called: Aleemily\_46 (DZ),

### **Summary by clusters:**

There are 2 clusters represented in this pham: singleton, DZ,

Info for manual annotations of cluster DZ:

- Start number 2 was manually annotated 3 times for cluster DZ.
- Start number 5 was manually annotated 1 time for cluster DZ.

### **Gene Information:**

Gene: Aleemily\_46 Start: 35741, Stop: 36493, Start Num: 5

Candidate Starts for Aleemily\_46:

(Start: 5 @35741 has 1 MA's), (6, 35756), (7, 35768), (8, 35774), (9, 35795), (10, 35798), (13, 35894), (16, 35981), (17, 36014), (23, 36272), (24, 36338),

Gene: Cafasso\_47 Start: 36237, Stop: 37067, Start Num: 2

Candidate Starts for Cafasso\_47:

(Start: 2 @36237 has 3 MA's), (4, 36276), (Start: 5 @36315 has 1 MA's), (6, 36330), (9, 36369), (13, 36468), (14, 36510), (15, 36540), (17, 36588), (23, 36846), (24, 36912),

Gene: ModicumRichard\_47 Start: 36236, Stop: 37066, Start Num: 2

Candidate Starts for ModicumRichard\_47:

(Start: 2 @36236 has 3 MA's), (4, 36275), (Start: 5 @36314 has 1 MA's), (6, 36329), (9, 36368), (13, 36467), (14, 36509), (15, 36539), (17, 36587), (23, 36845), (24, 36911),

Gene: Morgana\_47 Start: 36190, Stop: 37020, Start Num: 2

Candidate Starts for Morgana\_47:

(Start: 2 @36190 has 3 MA's), (4, 36229), (Start: 5 @36268 has 1 MA's), (6, 36283), (9, 36322), (13, 36421), (14, 36463), (17, 36541), (23, 36799), (24, 36865),

Gene: ObLaDi\_47 Start: 36227, Stop: 37057, Start Num: 2

Candidate Starts for ObLaDi\_47:

(Start: 2 @36227 has 3 MA's), (4, 36266), (Start: 5 @36305 has 1 MA's), (6, 36320), (9, 36359), (13, 36458), (14, 36500), (17, 36578), (23, 36836), (24, 36902),

Gene: VanLee\_45 Start: 33944, Stop: 34756, Start Num: 1

Candidate Starts for VanLee\_45:

(Start: 1 @33944 has 1 MA's), (3, 33956), (Start: 5 @34025 has 1 MA's), (11, 34088), (12, 34106), (18, 34307), (19, 34322), (20, 34331), (21, 34346), (22, 34418), (25, 34697), (26, 34745),