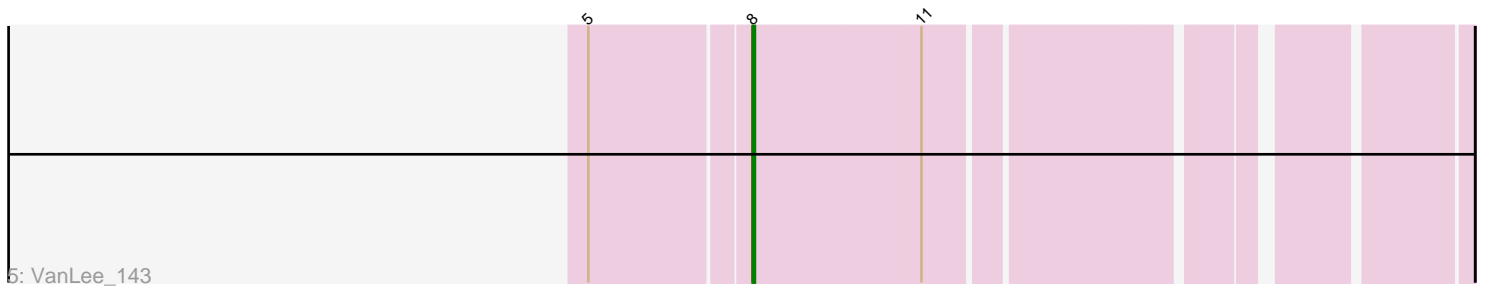
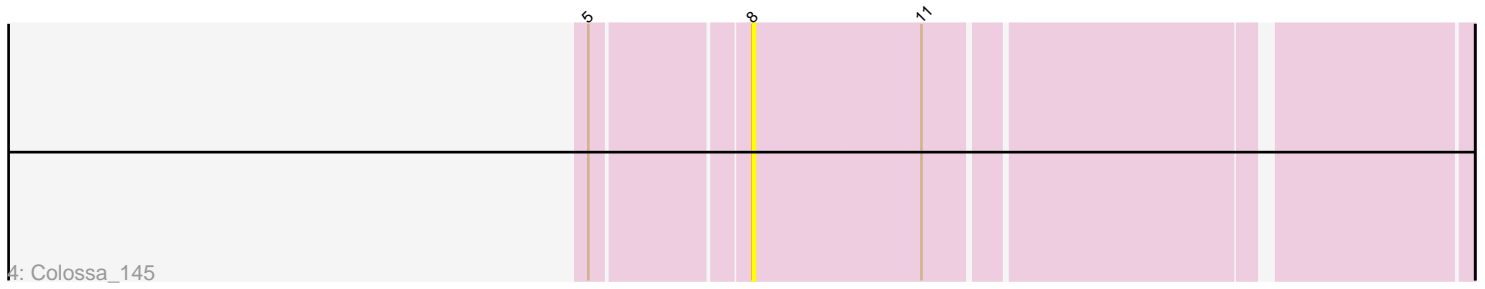
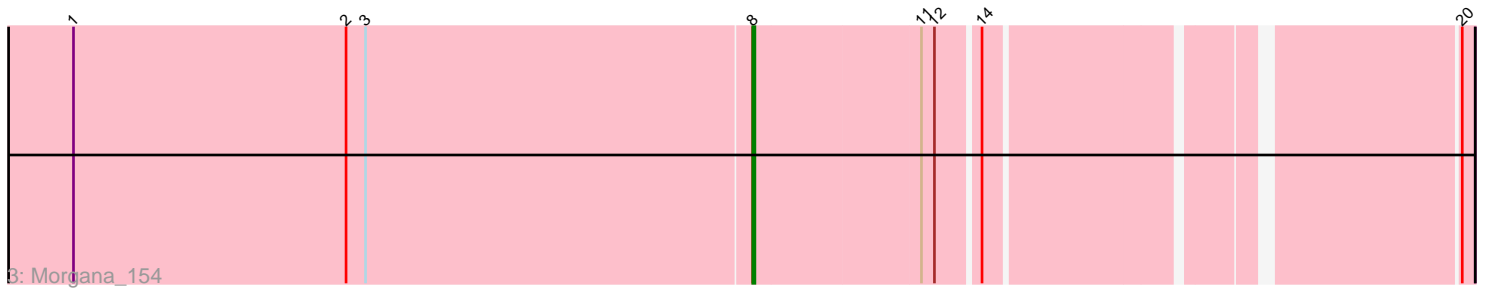
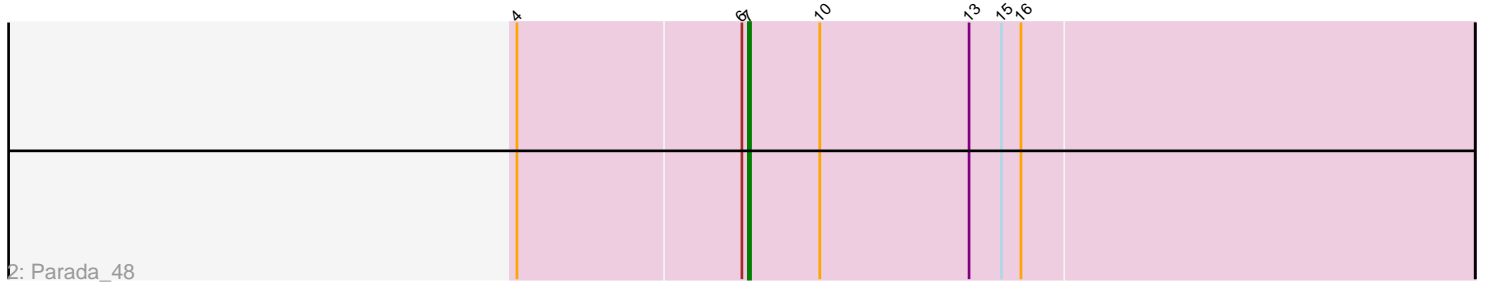
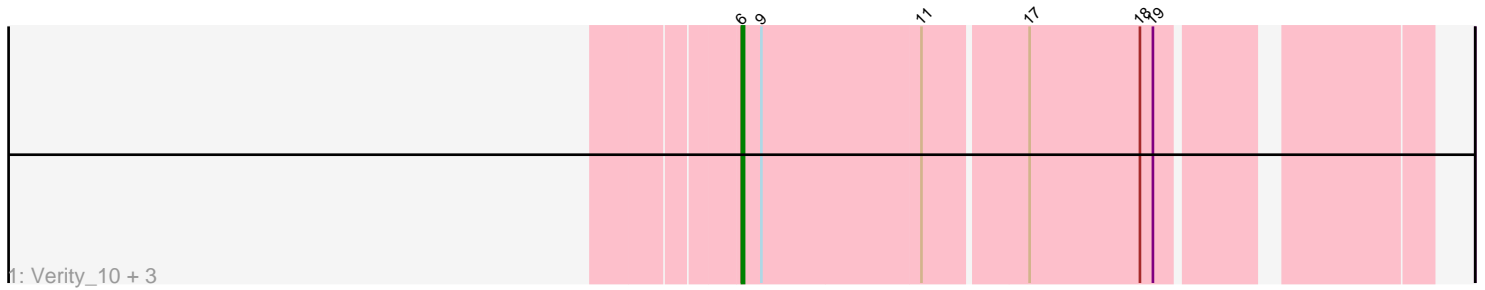


# Pham 297271



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 297271 Report

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

Pham number 297271 has 8 members, 1 are drafts.

Phages represented in each track:

- Track 1 : Verity\_10, Zipp\_10, Delrey21\_10, DoctorFroggo\_10
- Track 2 : Parada\_48
- Track 3 : Morgana\_154
- Track 4 : Colossa\_145
- Track 5 : VanLee\_143

### ***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 4 of the 7 non-draft genes in the pham.

Genes that call this "Most Annotated" start:

- Delrey21\_10, DoctorFroggo\_10, Verity\_10, Zipp\_10,

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

- Parada\_48,

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

- Colossa\_145, Morgana\_154, VanLee\_143,

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

Start 6:

- Found in 5 of 8 ( 62.5% ) of genes in pham
- Manual Annotations of this start: 4 of 7
- Called 80.0% of time when present
- Phage (with cluster) where this start called: Delrey21\_10 (DE4), DoctorFroggo\_10 (DE4), Verity\_10 (DE4), Zipp\_10 (DE4),

Start 7:

- Found in 1 of 8 ( 12.5% ) of genes in pham
- Manual Annotations of this start: 1 of 7
- Called 100.0% of time when present
- Phage (with cluster) where this start called: Parada\_48 (DI),

Start 8:

- Found in 3 of 8 ( 37.5% ) of genes in pham
- Manual Annotations of this start: 2 of 7
- Called 100.0% of time when present
- Phage (with cluster) where this start called: Colossa\_145 (KA), Morgana\_154 (DZ), VanLee\_143 (KA),

### **Summary by clusters:**

There are 4 clusters represented in this pham: DZ, KA, DE4, DI,

Info for manual annotations of cluster DE4:

- Start number 6 was manually annotated 4 times for cluster DE4.

Info for manual annotations of cluster DI:

- Start number 7 was manually annotated 1 time for cluster DI.

Info for manual annotations of cluster DZ:

- Start number 8 was manually annotated 1 time for cluster DZ.

Info for manual annotations of cluster KA:

- Start number 8 was manually annotated 1 time for cluster KA.

### **Gene Information:**

Gene: Colossa\_145 Start: 76064, Stop: 75759, Start Num: 8

Candidate Starts for Colossa\_145:

(5, 76130), (Start: 8 @76064 has 2 MA's), (11, 75989),

Gene: Delrey21\_10 Start: 5803, Stop: 6096, Start Num: 6

Candidate Starts for Delrey21\_10:

(Start: 6 @5803 has 4 MA's), (9, 5812), (11, 5884), (17, 5929), (18, 5980), (19, 5986),

Gene: DoctorFroggo\_10 Start: 5803, Stop: 6096, Start Num: 6

Candidate Starts for DoctorFroggo\_10:

(Start: 6 @5803 has 4 MA's), (9, 5812), (11, 5884), (17, 5929), (18, 5980), (19, 5986),

Gene: Morgana\_154 Start: 78675, Stop: 78376, Start Num: 8

Candidate Starts for Morgana\_154:

(1, 78987), (2, 78861), (3, 78852), (Start: 8 @78675 has 2 MA's), (11, 78600), (12, 78594), (14, 78576), (20, 78381),

Gene: Parada\_48 Start: 34923, Stop: 35255, Start Num: 7

Candidate Starts for Parada\_48:

(4, 34818), (Start: 6 @34920 has 4 MA's), (Start: 7 @34923 has 1 MA's), (10, 34956), (13, 35025), (15, 35040), (16, 35049),

Gene: VanLee\_143 Start: 75581, Stop: 75288, Start Num: 8

Candidate Starts for VanLee\_143:

(5, 75650), (Start: 8 @75581 has 2 MA's), (11, 75506),

Gene: Verity\_10 Start: 5803, Stop: 6096, Start Num: 6

Candidate Starts for Verity\_10:

(Start: 6 @5803 has 4 MA's), (9, 5812), (11, 5884), (17, 5929), (18, 5980), (19, 5986),

Gene: Zipp\_10 Start: 5948, Stop: 6241, Start Num: 6

Candidate Starts for Zipp\_10:

(Start: 6 @5948 has 4 MA's), (9, 5957), (11, 6029), (17, 6074), (18, 6125), (19, 6131),