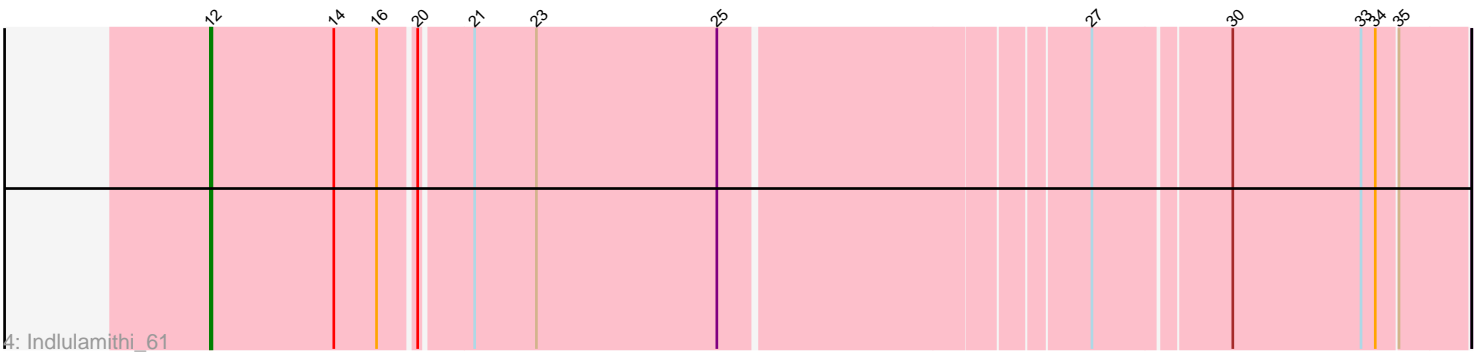
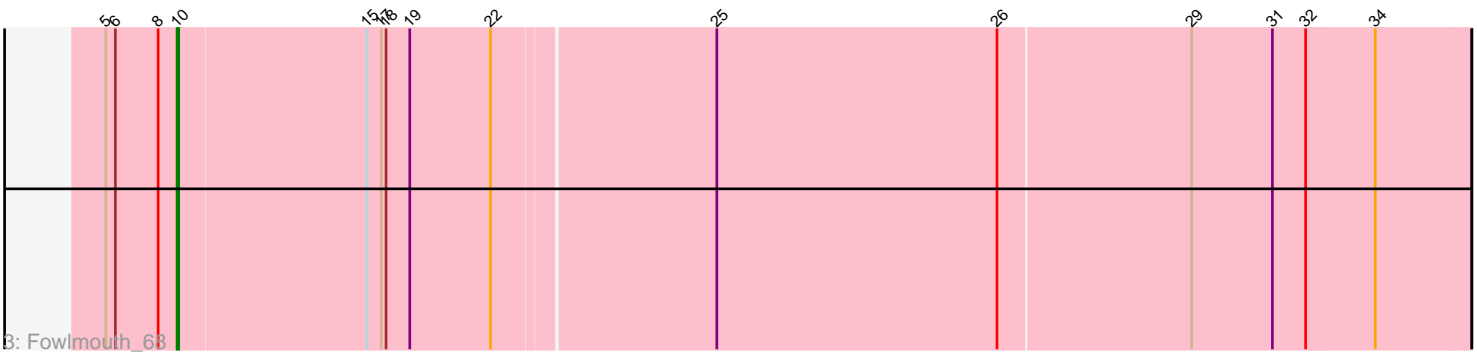
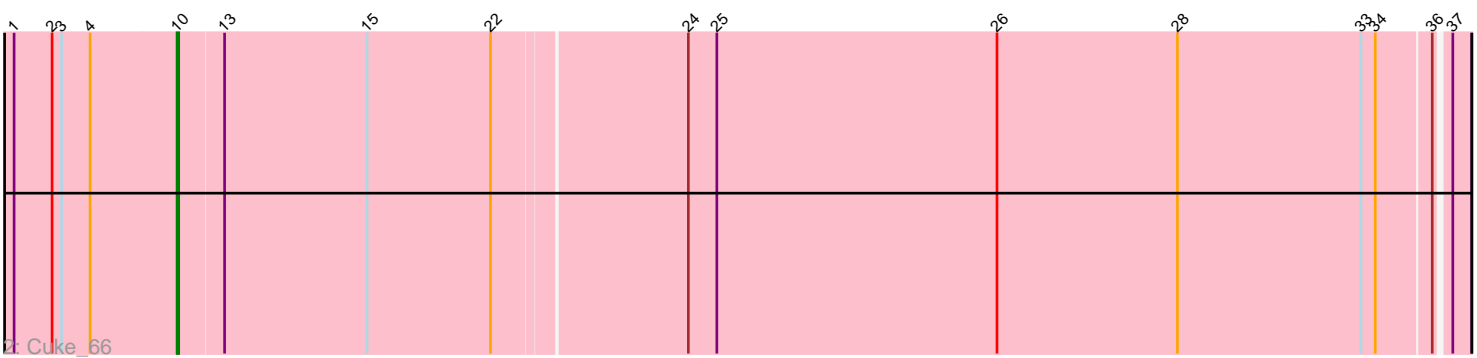
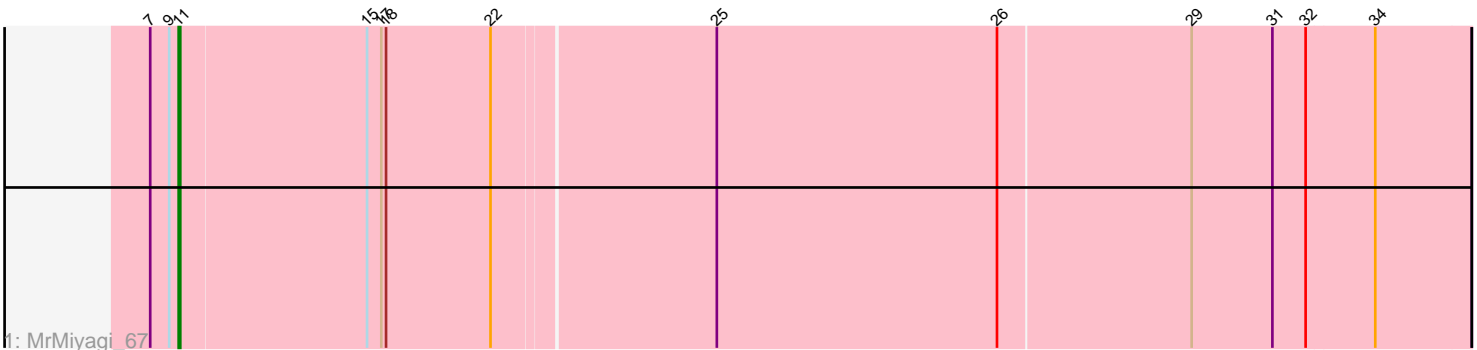


Pham 28413



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 28413 Report

This analysis was run 04/05/24 on database version 557.

Pham number 28413 has 4 members, 0 are drafts.

Phages represented in each track:

- Track 1 : MrMiyagi\_67
- Track 2 : Cuke\_66
- Track 3 : Fowlmouth\_68
- Track 4 : Indlulamithi\_61

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

The start number called the most often in the published annotations is 10, it was called in 2 of the 4 non-draft genes in the pham.

Genes that call this "Most Annotated" start:

- Cuke\_66, Fowlmouth\_68,

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

- 

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

- Indlulamithi\_61, MrMiyagi\_67,

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

Start 10:

- Found in 2 of 4 ( 50.0% ) of genes in pham
- Manual Annotations of this start: 2 of 4
- Called 100.0% of time when present
- Phage (with cluster) where this start called: Cuke\_66 (AC), Fowlmouth\_68 (AC),

Start 11:

- Found in 1 of 4 ( 25.0% ) of genes in pham
- Manual Annotations of this start: 1 of 4
- Called 100.0% of time when present
- Phage (with cluster) where this start called: MrMiyagi\_67 (AC),

Start 12:

- Found in 1 of 4 ( 25.0% ) of genes in pham
- Manual Annotations of this start: 1 of 4
- Called 100.0% of time when present
- Phage (with cluster) where this start called: Indlulamithi\_61 (AC),

### **Summary by clusters:**

There is one cluster represented in this pham: AC

Info for manual annotations of cluster AC:

- Start number 10 was manually annotated 2 times for cluster AC.
- Start number 11 was manually annotated 1 time for cluster AC.
- Start number 12 was manually annotated 1 time for cluster AC.

### **Gene Information:**

Gene: Cuke\_66 Start: 40166, Stop: 40960, Start Num: 10

Candidate Starts for Cuke\_66:

(1, 40064), (2, 40088), (3, 40094), (4, 40112), (Start: 10 @40166 has 2 MA's), (13, 40193), (15, 40283), (22, 40361), (24, 40478), (25, 40496), (26, 40673), (28, 40787), (33, 40901), (34, 40910), (36, 40943), (37, 40949),

Gene: Fowlmouth\_68 Start: 42995, Stop: 43795, Start Num: 10

Candidate Starts for Fowlmouth\_68:

(5, 42950), (6, 42956), (8, 42983), (Start: 10 @42995 has 2 MA's), (15, 43112), (17, 43121), (18, 43124), (19, 43139), (22, 43190), (25, 43325), (26, 43502), (29, 43622), (31, 43673), (32, 43694), (34, 43736),

Gene: Indlulamithi\_61 Start: 43580, Stop: 44332, Start Num: 12

Candidate Starts for Indlulamithi\_61:

(Start: 12 @43580 has 1 MA's), (14, 43658), (16, 43685), (20, 43706), (21, 43736), (23, 43775), (25, 43889), (27, 44108), (30, 44189), (33, 44270), (34, 44279), (35, 44291),

Gene: MrMiyagi\_67 Start: 43380, Stop: 44180, Start Num: 11

Candidate Starts for MrMiyagi\_67:

(7, 43362), (9, 43374), (Start: 11 @43380 has 1 MA's), (15, 43497), (17, 43506), (18, 43509), (22, 43575), (25, 43710), (26, 43887), (29, 44007), (31, 44058), (32, 44079), (34, 44121),