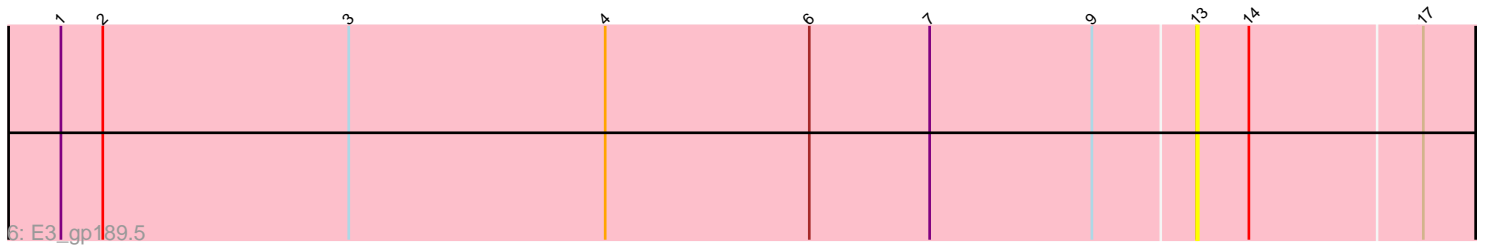
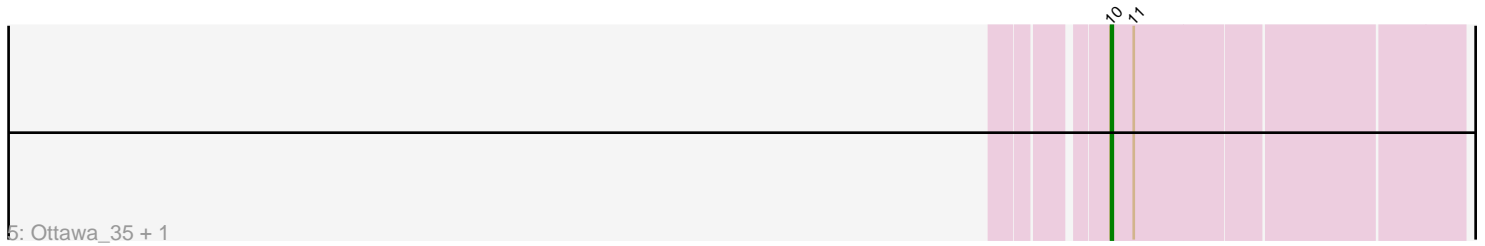
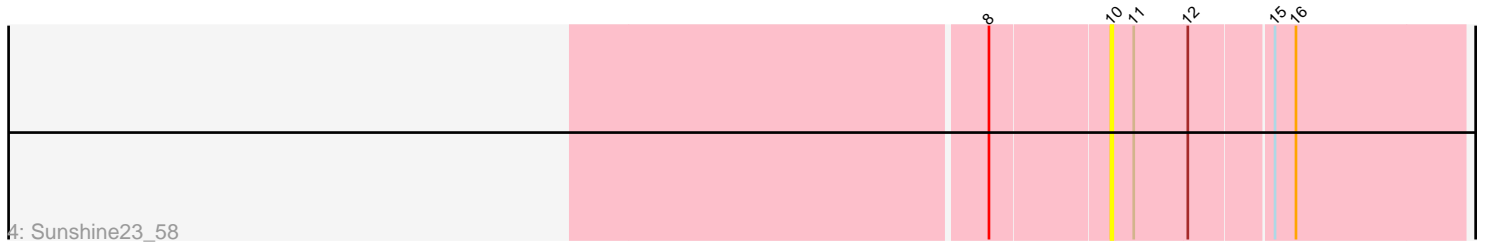
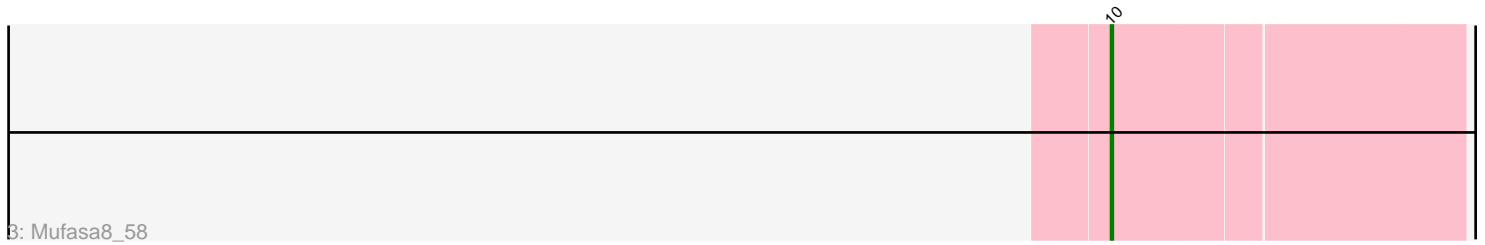
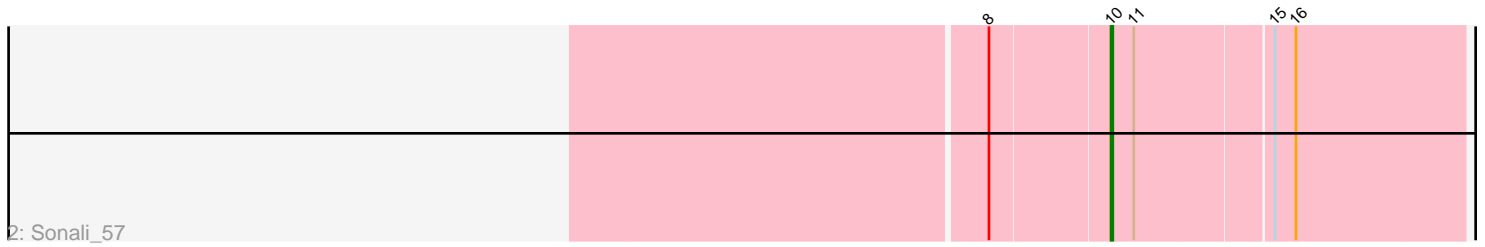
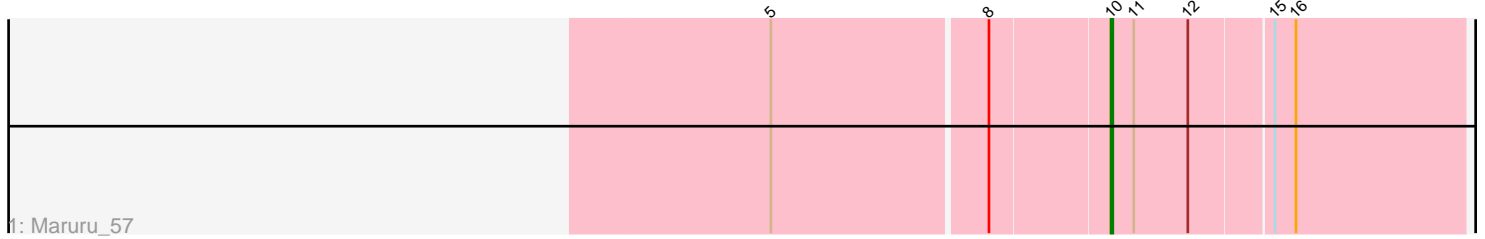


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Note: Tracks are now grouped by subcluster and scaled. Switching in subcluster is indicated by changes in track color. Track scale is now set by default to display the region 30 bp upstream of start 1 to 30 bp downstream of the last possible start. 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 297352 Report

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

Pham number 297352 has 7 members, 2 are drafts.

Phages represented in each track:

- Track 1 : Maruru\_57
- Track 2 : Sonali\_57
- Track 3 : Mufasa8\_58
- Track 4 : Sunshine23\_58
- Track 5 : Ottawa\_35, Kharcho\_35
- Track 6 : E3\_gp189.5

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

Genes that call this "Most Annotated" start:

- Kharcho\_35, Maruru\_57, Mufasa8\_58, Ottawa\_35, Sonali\_57, Sunshine23\_58,

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

- 

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

- E3\_gp189.5,

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

Start 10:

- Found in 6 of 7 ( 85.7% ) of genes in pham
- Manual Annotations of this start: 5 of 5
- Called 100.0% of time when present
- Phage (with cluster) where this start called: Kharcho\_35 (FM), Maruru\_57 (FG), Mufasa8\_58 (FG), Ottawa\_35 (FM), Sonali\_57 (FG), Sunshine23\_58 (FG),

Start 13:

- Found in 1 of 7 ( 14.3% ) of genes in pham
- No Manual Annotations of this start.
- Called 100.0% of time when present

- Phage (with cluster) where this start called: E3\_gp189.5 (singleton),

### **Summary by clusters:**

There are 3 clusters represented in this pham: singleton, FM, FG,

Info for manual annotations of cluster FG:

- Start number 10 was manually annotated 3 times for cluster FG.

Info for manual annotations of cluster FM:

- Start number 10 was manually annotated 2 times for cluster FM.

### **Gene Information:**

Gene: E3\_gp189.5 Start: 125789, Stop: 125962, Start Num: 13

Candidate Starts for E3\_gp189.5:

(1, 125141), (2, 125165), (3, 125306), (4, 125453), (6, 125570), (7, 125639), (9, 125732), (13, 125789), (14, 125819), (17, 125915),

Gene: Kharcho\_35 Start: 14505, Stop: 14696, Start Num: 10

Candidate Starts for Kharcho\_35:

(Start: 10 @14505 has 5 MA's), (11, 14517),

Gene: Maruru\_57 Start: 42255, Stop: 42449, Start Num: 10

Candidate Starts for Maruru\_57:

(5, 42069), (8, 42189), (Start: 10 @42255 has 5 MA's), (11, 42267), (12, 42297), (15, 42342), (16, 42354),

Gene: Mufasa8\_58 Start: 41034, Stop: 41228, Start Num: 10

Candidate Starts for Mufasa8\_58:

(Start: 10 @41034 has 5 MA's),

Gene: Ottawa\_35 Start: 14503, Stop: 14694, Start Num: 10

Candidate Starts for Ottawa\_35:

(Start: 10 @14503 has 5 MA's), (11, 14515),

Gene: Sonali\_57 Start: 42711, Stop: 42905, Start Num: 10

Candidate Starts for Sonali\_57:

(8, 42645), (Start: 10 @42711 has 5 MA's), (11, 42723), (15, 42798), (16, 42810),

Gene: Sunshine23\_58 Start: 42392, Stop: 42586, Start Num: 10

Candidate Starts for Sunshine23\_58:

(8, 42326), (Start: 10 @42392 has 5 MA's), (11, 42404), (12, 42434), (15, 42479), (16, 42491),