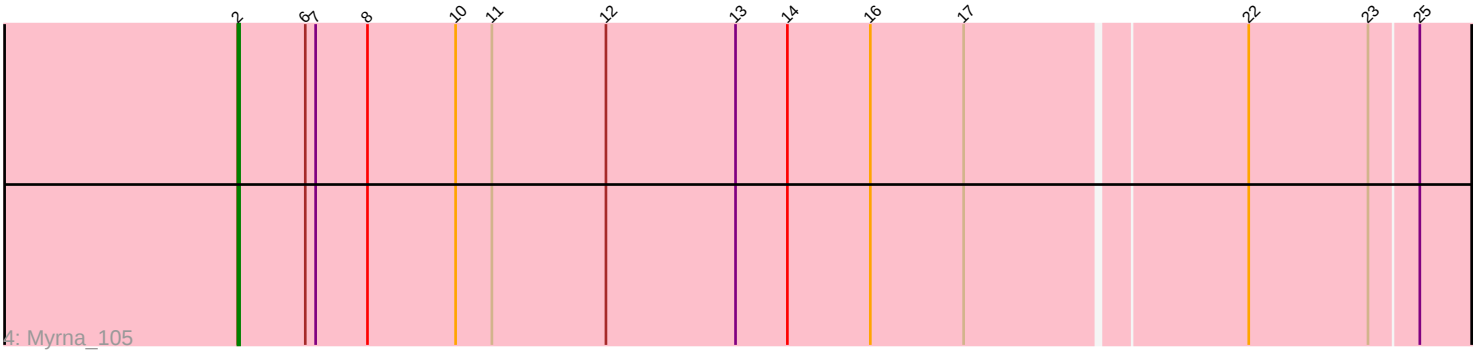
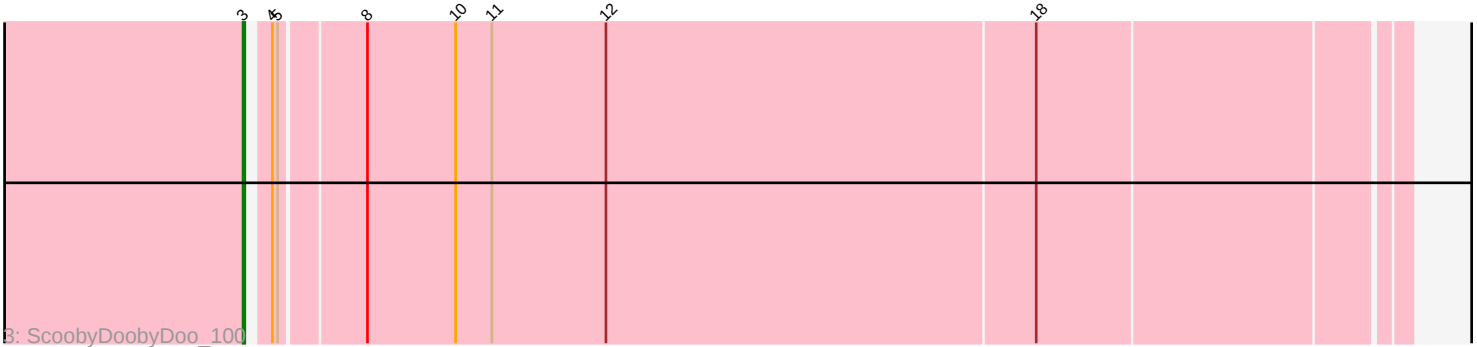
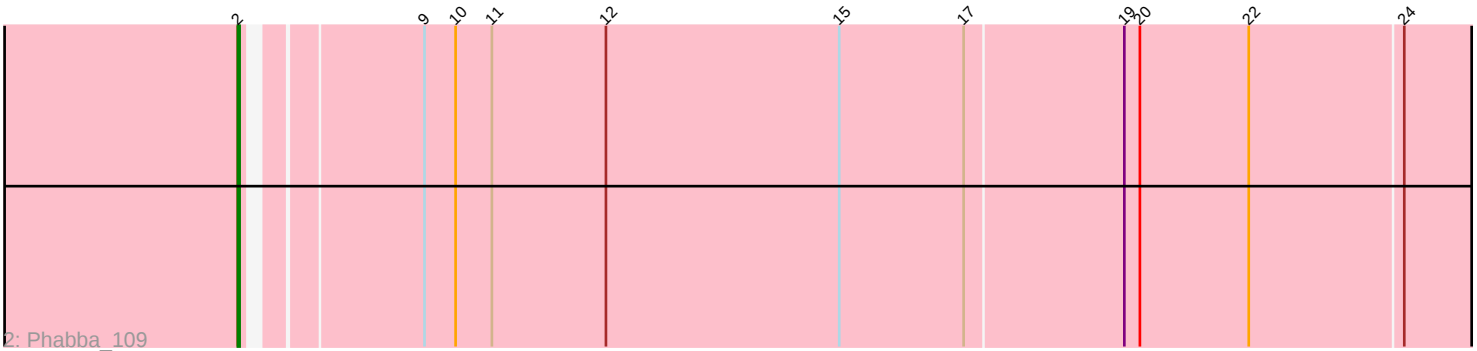
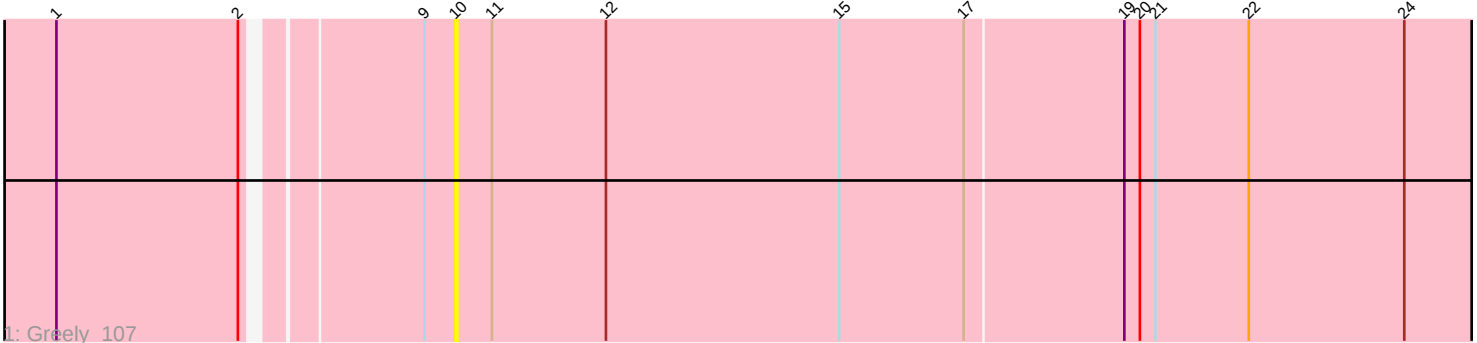


Pham 310397



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 310397 Report

This analysis was run 09/05/26 on database version 662.

Pham number 310397 has 4 members, 1 are drafts.

Phages represented in each track:

- Track 1 : Greely_107
- Track 2 : Phabba_109
- Track 3 : ScoobyDoobyDoo_100
- Track 4 : Myrna_105

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 2 of the 3 non-draft genes in the pham.

Genes that call this "Most Annotated" start:

- Myrna_105, Phabba_109,

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

- Greely_107,

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

- ScoobyDoobyDoo_100,

Summary by start number:

Start 2:

- Found in 3 of 4 (75.0%) of genes in pham
- Manual Annotations of this start: 2 of 3
- Called 66.7% of time when present
- Phage (with cluster) where this start called: Myrna_105 (C2), Phabba_109 (C2),

Start 3:

- Found in 1 of 4 (25.0%) of genes in pham
- Manual Annotations of this start: 1 of 3
- Called 100.0% of time when present
- Phage (with cluster) where this start called: ScoobyDoobyDoo_100 (C2),

Start 10:

- Found in 4 of 4 (100.0%) of genes in pham
- No Manual Annotations of this start.
- Called 25.0% of time when present
- Phage (with cluster) where this start called: Greely_107 (C2),

Summary by clusters:

There is one cluster represented in this pham: C2

Info for manual annotations of cluster C2:

- Start number 2 was manually annotated 2 times for cluster C2.
- Start number 3 was manually annotated 1 time for cluster C2.

Gene Information:

Gene: Greely_107 Start: 54382, Stop: 55014, Start Num: 10

Candidate Starts for Greely_107:

(1, 54169), (Start: 2 @54274 has 2 MA's), (9, 54364), (10, 54382), (11, 54403), (12, 54469), (15, 54604), (17, 54676), (19, 54766), (20, 54775), (21, 54784), (22, 54838), (24, 54928),

Gene: Myrna_105 Start: 55086, Stop: 55967, Start Num: 2

Candidate Starts for Myrna_105:

(Start: 2 @55086 has 2 MA's), (6, 55125), (7, 55131), (8, 55161), (10, 55212), (11, 55233), (12, 55299), (13, 55374), (14, 55404), (16, 55452), (17, 55506), (22, 55662), (23, 55731), (25, 55758),

Gene: Phabba_109 Start: 54030, Stop: 54767, Start Num: 2

Candidate Starts for Phabba_109:

(Start: 2 @54030 has 2 MA's), (9, 54120), (10, 54138), (11, 54159), (12, 54225), (15, 54360), (17, 54432), (19, 54522), (20, 54531), (22, 54594), (24, 54681),

Gene: ScoobyDoobyDoo_100 Start: 53713, Stop: 54357, Start Num: 3

Candidate Starts for ScoobyDoobyDoo_100:

(Start: 3 @53713 has 1 MA's), (4, 53722), (5, 53725), (8, 53770), (10, 53821), (11, 53842), (12, 53908), (18, 54154),