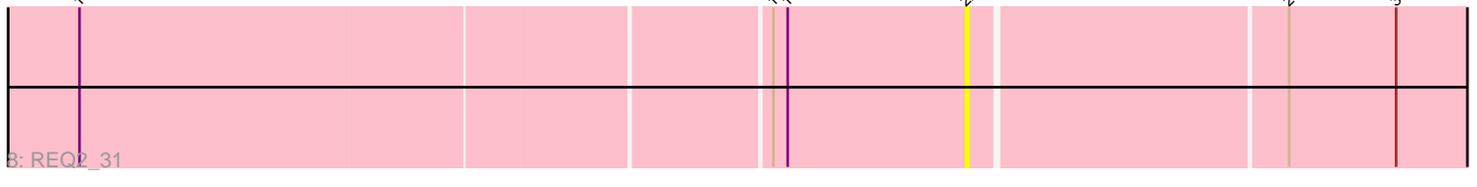
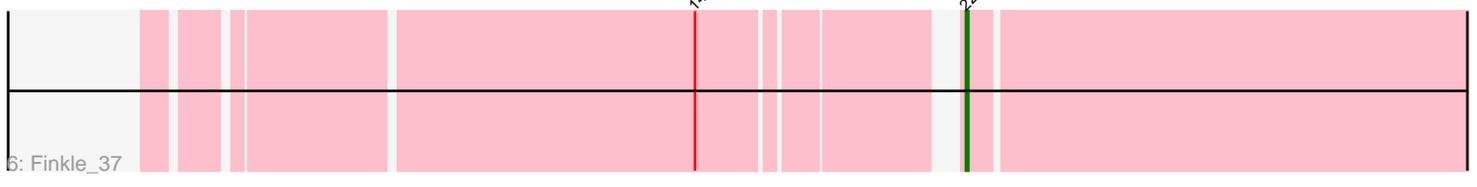
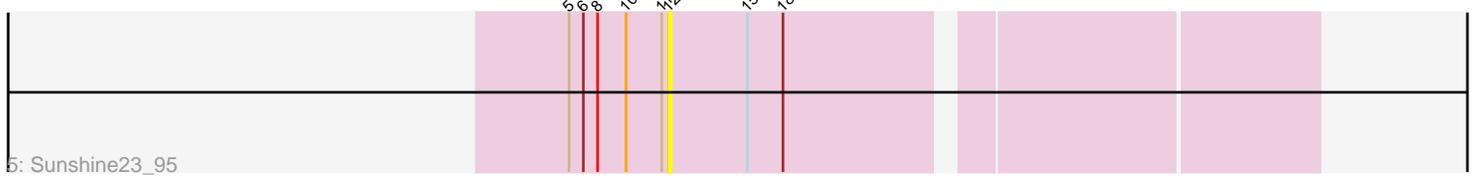
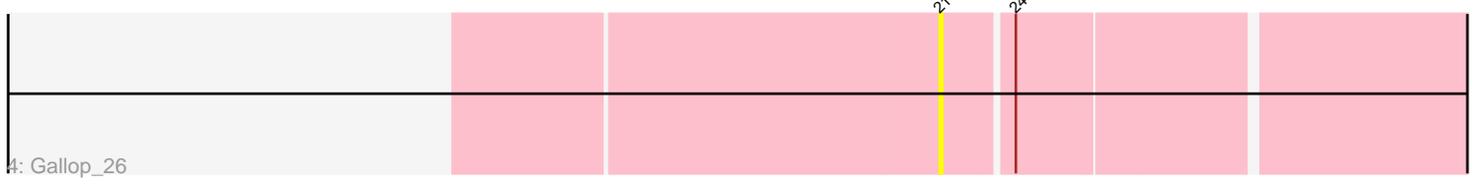
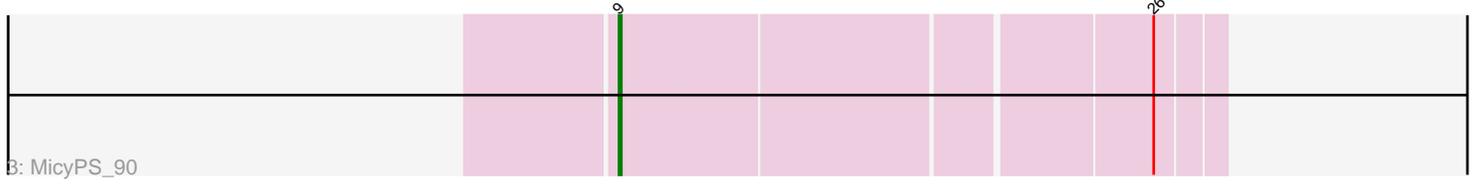
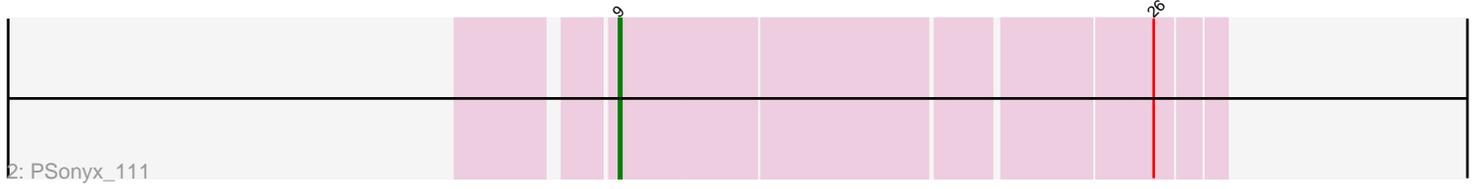
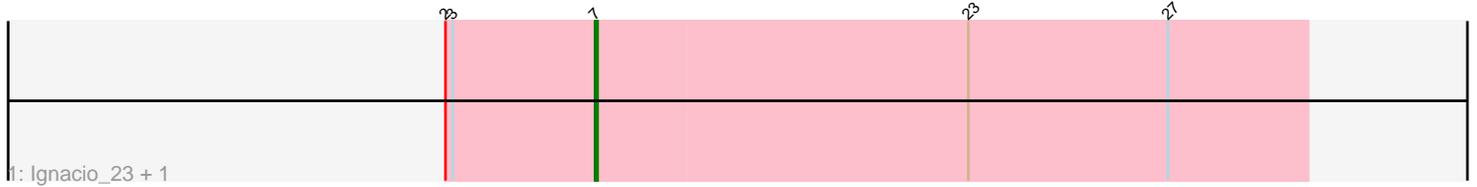


Pham 284463



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 284463 Report

This analysis was run 02/23/26 on database version 636.

Pham number 284463 has 9 members, 4 are drafts.

Phages represented in each track:

- Track 1 : Ignacio_23, HFrancette_23
- Track 2 : PSonyx_111
- Track 3 : MicyPS_90
- Track 4 : Gallop_26
- Track 5 : Sunshine23_95
- Track 6 : Finkle_37
- Track 7 : phiAsp2_39
- Track 8 : REQ2_31

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

The start number called the most often in the published annotations is 7, it was called in 2 of the 5 non-draft genes in the pham.

Genes that call this "Most Annotated" start:

- HFrancette_23, Ignacio_23,

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

•

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

- Finkle_37, Gallop_26, MicyPS_90, PSonyx_111, REQ2_31, Sunshine23_95, phiAsp2_39,

Summary by start number:

Start 7:

- Found in 2 of 9 (22.2%) of genes in pham
- Manual Annotations of this start: 2 of 5
- Called 100.0% of time when present
- Phage (with cluster) where this start called: HFrancette_23 (BP), Ignacio_23 (BP),

Start 9:

- Found in 3 of 9 (33.3%) of genes in pham

- Manual Annotations of this start: 2 of 5
- Called 100.0% of time when present
- Phage (with cluster) where this start called: MicyPS_90 (EQ), PSonyx_111 (EQ), phiAsp2_39 (singleton),

Start 12:

- Found in 1 of 9 (11.1%) of genes in pham
- No Manual Annotations of this start.
- Called 100.0% of time when present
- Phage (with cluster) where this start called: Sunshine23_95 (FG),

Start 21:

- Found in 1 of 9 (11.1%) of genes in pham
- No Manual Annotations of this start.
- Called 100.0% of time when present
- Phage (with cluster) where this start called: Gallop_26 (ES),

Start 22:

- Found in 2 of 9 (22.2%) 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: Finkle_37 (singleton), REQ2_31 (singleton),

Summary by clusters:

There are 5 clusters represented in this pham: EQ, singleton, BP, ES, FG,

Info for manual annotations of cluster BP:

- Start number 7 was manually annotated 2 times for cluster BP.

Info for manual annotations of cluster EQ:

- Start number 9 was manually annotated 2 times for cluster EQ.

Gene Information:

Gene: Finkle_37 Start: 29834, Stop: 29601, Start Num: 22

Candidate Starts for Finkle_37:

(14, 29927), (Start: 22 @29834 has 1 MA's),

Gene: Gallop_26 Start: 24284, Stop: 24048, Start Num: 21

Candidate Starts for Gallop_26:

(21, 24284), (24, 24257),

Gene: HFrancette_23 Start: 18663, Stop: 18959, Start Num: 7

Candidate Starts for HFrancette_23:

(2, 18600), (3, 18603), (Start: 7 @18663 has 2 MA's), (23, 18819), (27, 18903),

Gene: Ignacio_23 Start: 18663, Stop: 18959, Start Num: 7

Candidate Starts for Ignacio_23:

(2, 18600), (3, 18603), (Start: 7 @18663 has 2 MA's), (23, 18819), (27, 18903),

Gene: MicyPS_90 Start: 54573, Stop: 54337, Start Num: 9

Candidate Starts for MicyPS_90:

(Start: 9 @54573 has 2 MA's), (26, 54363),

Gene: PSonyx_111 Start: 59498, Stop: 59262, Start Num: 9

Candidate Starts for PSonyx_111:

(Start: 9 @59498 has 2 MA's), (26, 59288),

Gene: REQ2_31 Start: 26798, Stop: 26559, Start Num: 22

Candidate Starts for REQ2_31:

(1, 27161), (17, 26879), (19, 26873), (Start: 22 @26798 has 1 MA's), (29, 26669), (30, 26624),

Gene: Sunshine23_95 Start: 57525, Stop: 57268, Start Num: 12

Candidate Starts for Sunshine23_95:

(5, 57567), (6, 57561), (8, 57555), (10, 57543), (11, 57528), (12, 57525), (15, 57492), (18, 57477),

Gene: phiAsp2_39 Start: 33131, Stop: 33403, Start Num: 9

Candidate Starts for phiAsp2_39:

(4, 33095), (Start: 9 @33131 has 2 MA's), (13, 33155), (16, 33191), (20, 33251), (25, 33317), (28, 33392),