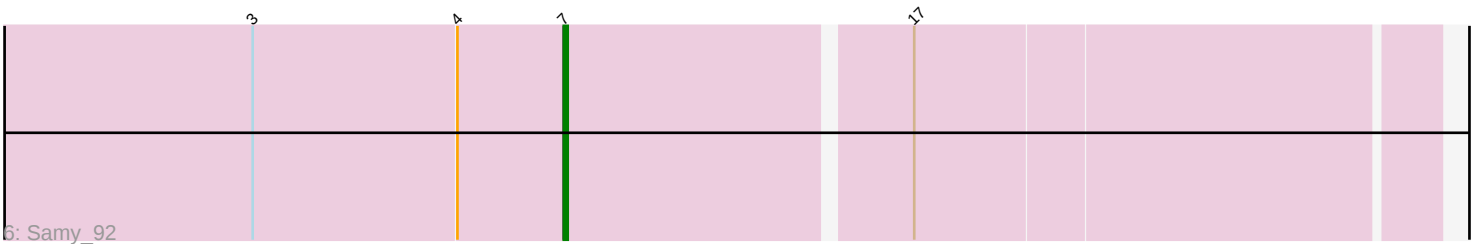
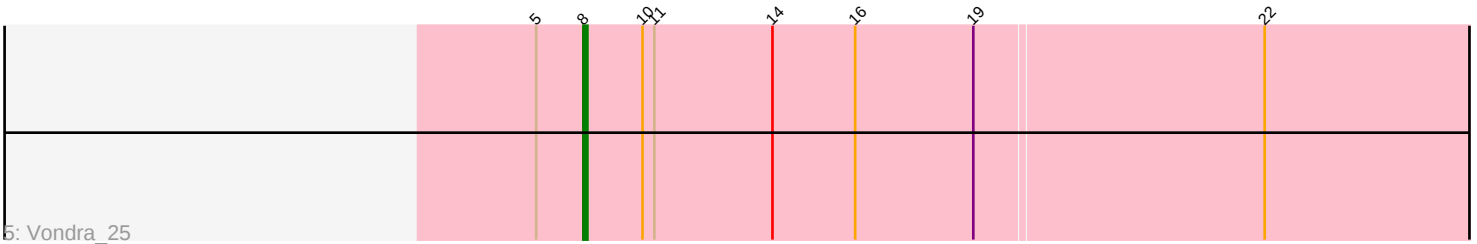
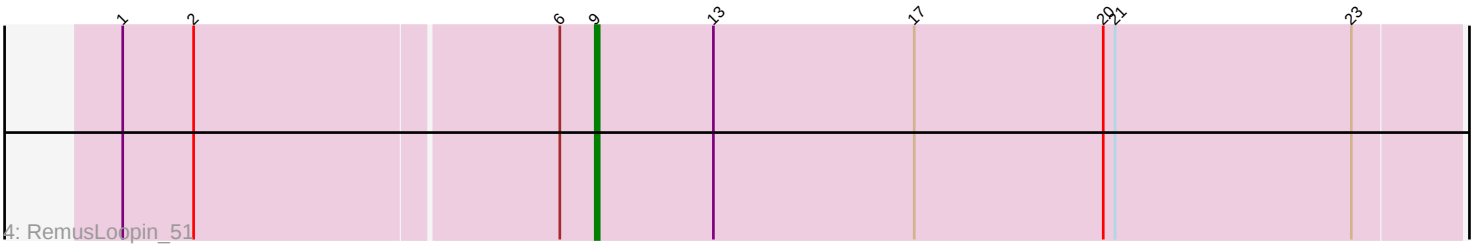
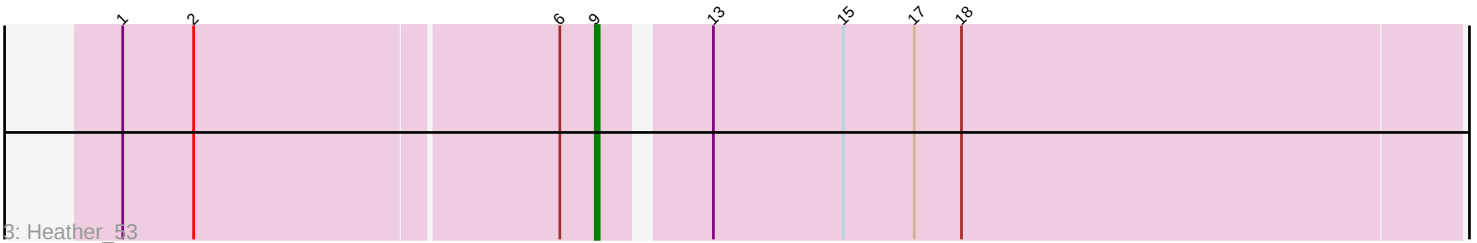
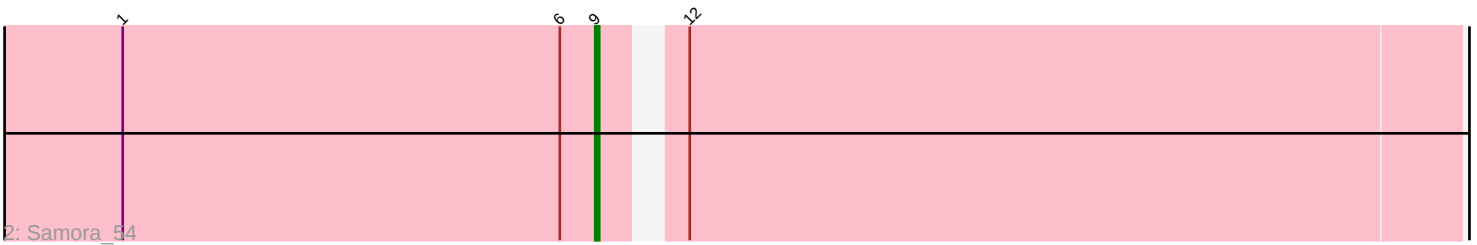
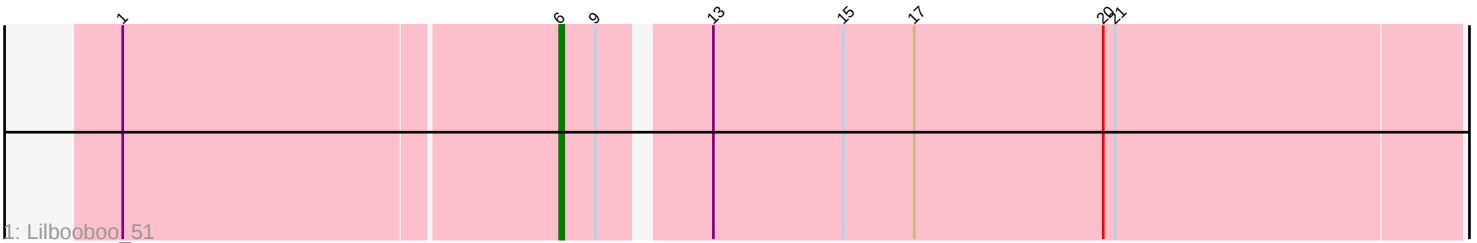


Pham 310148



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 310148 Report

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

Pham number 310148 has 6 members, 0 are drafts.

Phages represented in each track:

- Track 1 : Lilbooboo_51
- Track 2 : Samora_54
- Track 3 : Heather_53
- Track 4 : RemusLoopin_51
- Track 5 : Vondra_25
- Track 6 : Samy_92

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

The start number called the most often in the published annotations is 9, it was called in 3 of the 6 non-draft genes in the pham.

Genes that call this "Most Annotated" start:

- Heather_53, RemusLoopin_51, Samora_54,

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

- Lilbooboo_51,

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

- Samy_92, Vondra_25,

Summary by start number:

Start 6:

- Found in 4 of 6 (66.7%) of genes in pham
- Manual Annotations of this start: 1 of 6
- Called 25.0% of time when present
- Phage (with cluster) where this start called: Lilbooboo_51 (BB1),

Start 7:

- Found in 1 of 6 (16.7%) of genes in pham
- Manual Annotations of this start: 1 of 6
- Called 100.0% of time when present
- Phage (with cluster) where this start called: Samy_92 (singleton),

Start 8:

- Found in 1 of 6 (16.7%) of genes in pham
- Manual Annotations of this start: 1 of 6
- Called 100.0% of time when present
- Phage (with cluster) where this start called: Vondra_25 (BP),

Start 9:

- Found in 4 of 6 (66.7%) of genes in pham
- Manual Annotations of this start: 3 of 6
- Called 75.0% of time when present
- Phage (with cluster) where this start called: Heather_53 (BB2), RemusLoopin_51 (BB2), Samora_54 (BB1),

Summary by clusters:

There are 4 clusters represented in this pham: singleton, BP, BB2, BB1,

Info for manual annotations of cluster BB1:

- Start number 6 was manually annotated 1 time for cluster BB1.
- Start number 9 was manually annotated 1 time for cluster BB1.

Info for manual annotations of cluster BB2:

- Start number 9 was manually annotated 2 times for cluster BB2.

Info for manual annotations of cluster BP:

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

Gene Information:

Gene: Heather_53 Start: 37446, Stop: 37658, Start Num: 9

Candidate Starts for Heather_53:

(1, 37329), (2, 37347), (Start: 6 @37437 has 1 MA's), (Start: 9 @37446 has 3 MA's), (13, 37470), (15, 37503), (17, 37521), (18, 37533),

Gene: Lilbooboo_51 Start: 36992, Stop: 37213, Start Num: 6

Candidate Starts for Lilbooboo_51:

(1, 36884), (Start: 6 @36992 has 1 MA's), (Start: 9 @37001 has 3 MA's), (13, 37025), (15, 37058), (17, 37076), (20, 37124), (21, 37127),

Gene: RemusLoopin_51 Start: 37583, Stop: 37801, Start Num: 9

Candidate Starts for RemusLoopin_51:

(1, 37466), (2, 37484), (Start: 6 @37574 has 1 MA's), (Start: 9 @37583 has 3 MA's), (13, 37613), (17, 37664), (20, 37712), (21, 37715), (23, 37775),

Gene: Samora_54 Start: 37813, Stop: 38022, Start Num: 9

Candidate Starts for Samora_54:

(1, 37693), (Start: 6 @37804 has 1 MA's), (Start: 9 @37813 has 3 MA's), (12, 37828),

Gene: Samy_92 Start: 56116, Stop: 56328, Start Num: 7

Candidate Starts for Samy_92:

(3, 56038), (4, 56089), (Start: 7 @56116 has 1 MA's), (17, 56200),

Gene: Vondra_25 Start: 19991, Stop: 20215, Start Num: 8

Candidate Starts for Vondra_25:

(5, 19979), (Start: 8 @19991 has 1 MA's), (10, 20006), (11, 20009), (14, 20039), (16, 20060), (19, 20090), (22, 20162),