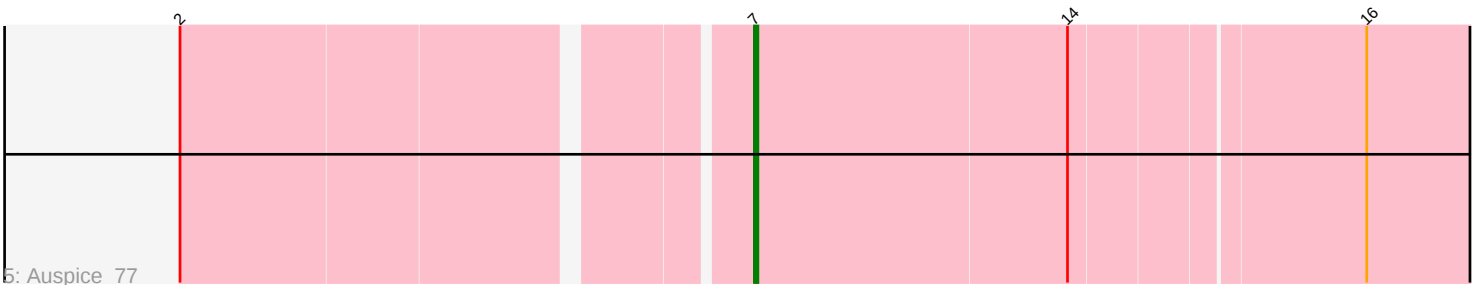
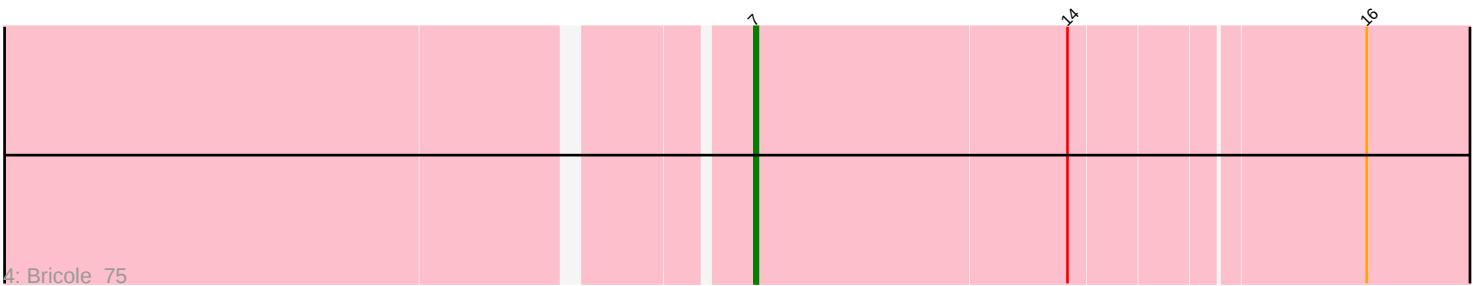
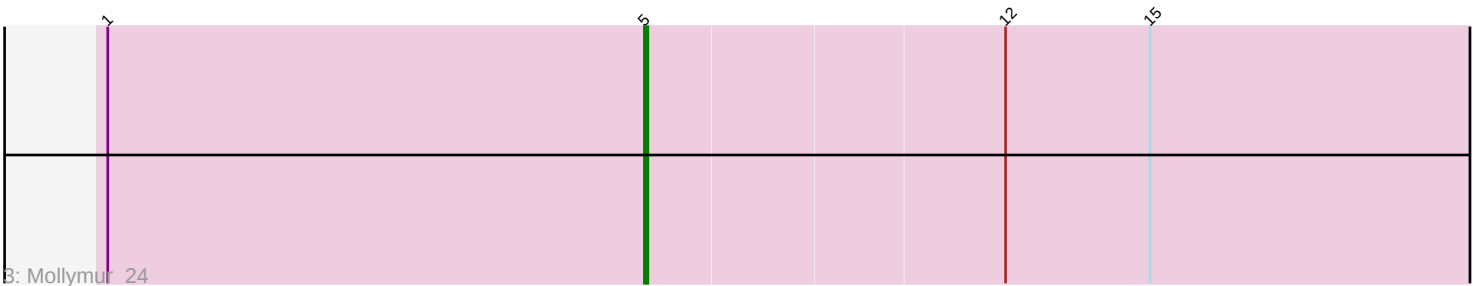
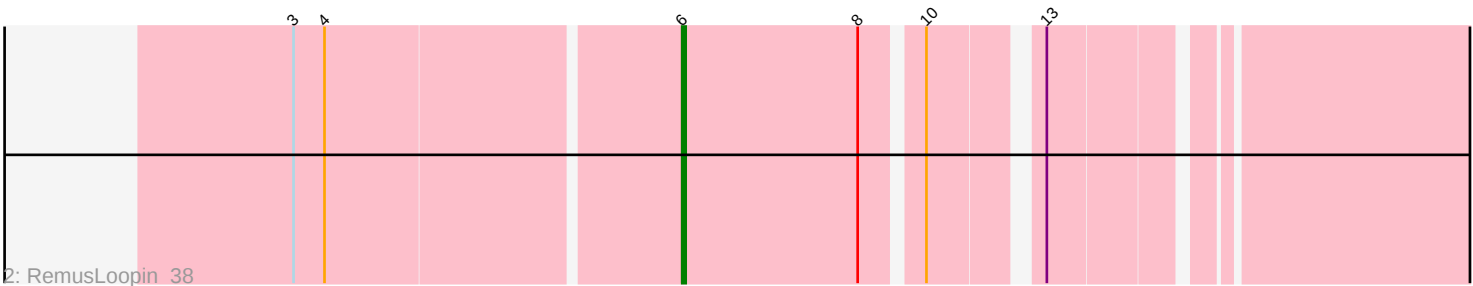
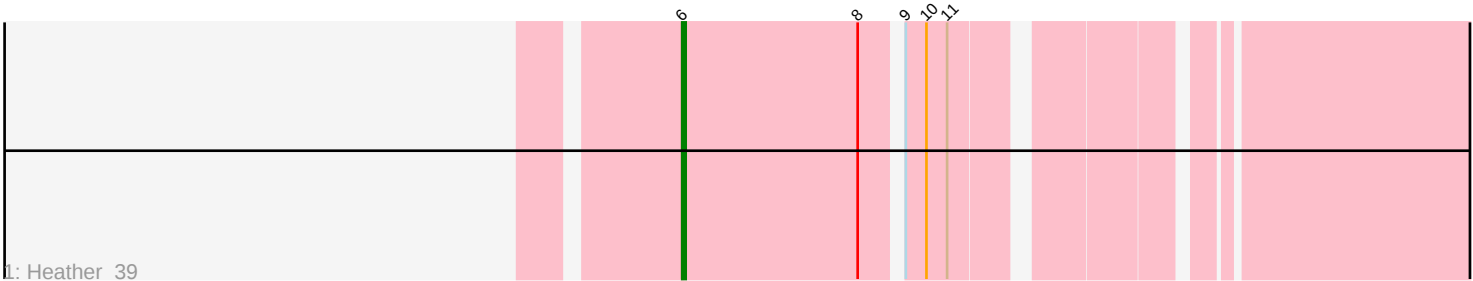


Pham 331756



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 331756 Report

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

Pham number 331756 has 5 members, 0 are drafts.

Phages represented in each track:

- Track 1 : Heather_39
- Track 2 : RemusLoopin_38
- Track 3 : Mollymur_24
- Track 4 : Bricole_75
- Track 5 : Auspice_77

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:

- Auspice_77, Bricole_75,

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

-

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

- Heather_39, Mollymur_24, RemusLoopin_38,

Summary by start number:

Start 5:

- Found in 1 of 5 (20.0%) 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: Mollymur_24 (DL),

Start 6:

- Found in 2 of 5 (40.0%) 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: Heather_39 (BB2), RemusLoopin_38 (BB2),

Start 7:

- Found in 2 of 5 (40.0%) 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: Auspice_77 (M1), Bricole_75 (M1),

Summary by clusters:

There are 3 clusters represented in this pham: DL, M1, BB2,

Info for manual annotations of cluster BB2:

- Start number 6 was manually annotated 2 times for cluster BB2.

Info for manual annotations of cluster DL:

- Start number 5 was manually annotated 1 time for cluster DL.

Info for manual annotations of cluster M1:

- Start number 7 was manually annotated 2 times for cluster M1.

Gene Information:

Gene: Auspice_77 Start: 49327, Stop: 49590, Start Num: 7

Candidate Starts for Auspice_77:

(2, 49174), (Start: 7 @49327 has 2 MA's), (14, 49417), (16, 49498),

Gene: Bricole_75 Start: 48816, Stop: 49079, Start Num: 7

Candidate Starts for Bricole_75:

(Start: 7 @48816 has 2 MA's), (14, 48906), (16, 48987),

Gene: Heather_39 Start: 31344, Stop: 31631, Start Num: 6

Candidate Starts for Heather_39:

(Start: 6 @31344 has 2 MA's), (8, 31395), (9, 31404), (10, 31410), (11, 31416),

Gene: Mollymur_24 Start: 15771, Stop: 16076, Start Num: 5

Candidate Starts for Mollymur_24:

(1, 15615), (Start: 5 @15771 has 1 MA's), (12, 15873), (15, 15915),

Gene: RemusLoopin_38 Start: 31565, Stop: 31852, Start Num: 6

Candidate Starts for RemusLoopin_38:

(3, 31457), (4, 31466), (Start: 6 @31565 has 2 MA's), (8, 31616), (10, 31631), (13, 31658),