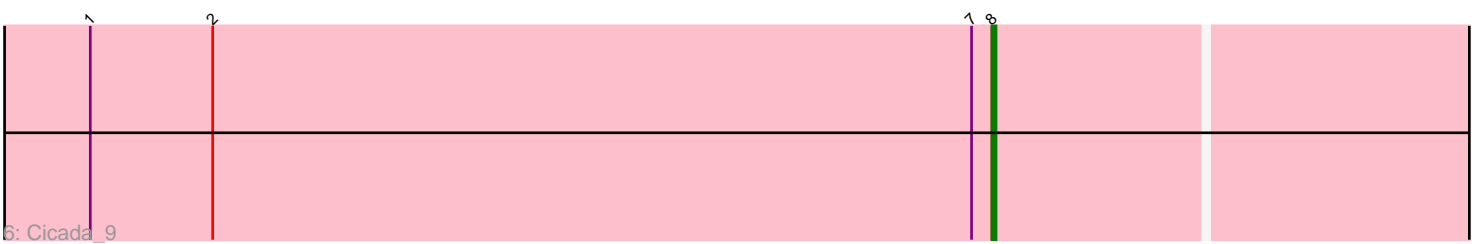
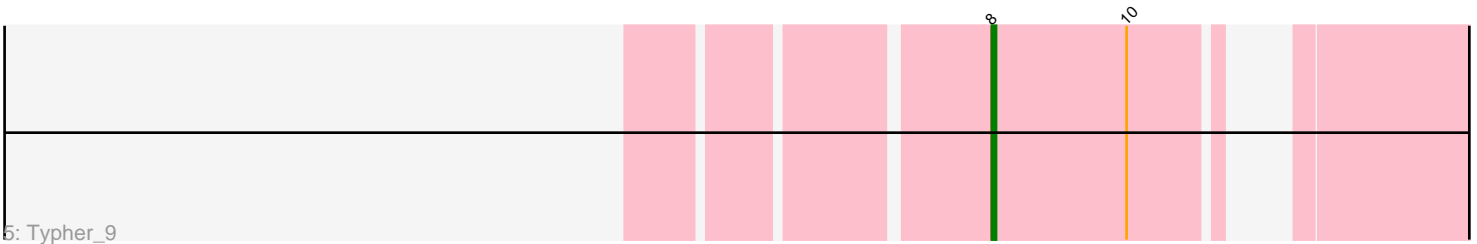
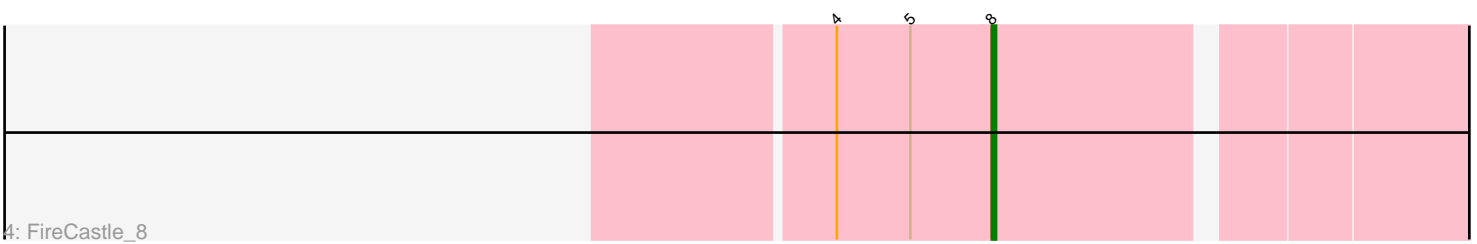
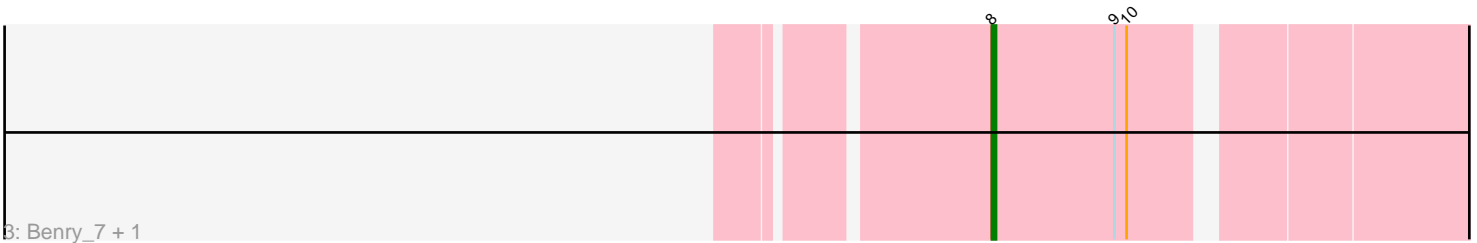
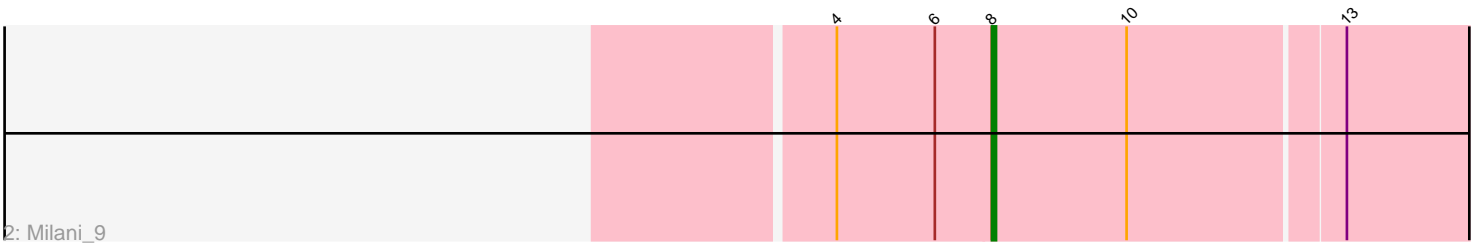
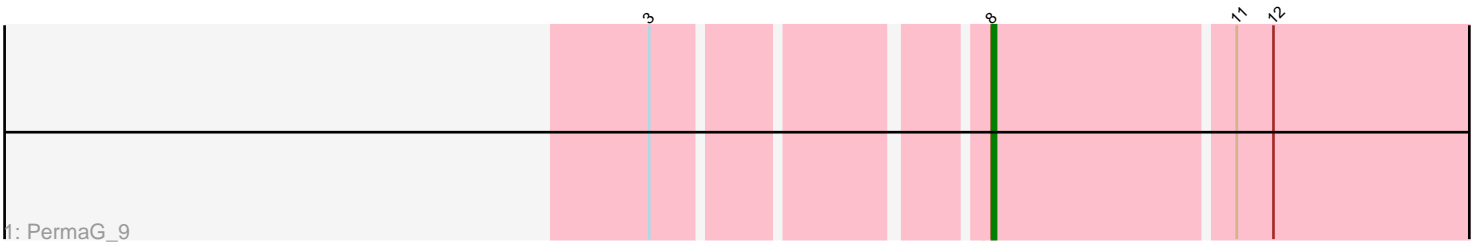


Pham 8872



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 8872 Report

This analysis was run 04/28/24 on database version 559.

Pham number 8872 has 7 members, 1 are drafts.

Phages represented in each track:

- Track 1 : PermaG_9
- Track 2 : Milani_9
- Track 3 : Benry_7, Sucha_7
- Track 4 : FireCastle_8
- Track 5 : Typher_9
- Track 6 : Cicada_9

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

The start number called the most often in the published annotations is 8, it was called in 6 of the 6 non-draft genes in the pham.

Genes that call this "Most Annotated" start:

- Benry_7, Cicada_9, FireCastle_8, Milani_9, PermaG_9, Sucha_7, Typher_9,

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

-

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

-

Summary by start number:

Start 8:

- Found in 7 of 7 (100.0%) of genes in pham
- Manual Annotations of this start: 6 of 6
- Called 100.0% of time when present
- Phage (with cluster) where this start called: Benry_7 (EJ), Cicada_9 (EJ), FireCastle_8 (EJ), Milani_9 (EJ), PermaG_9 (EJ), Sucha_7 (EJ), Typher_9 (EJ),

Summary by clusters:

There is one cluster represented in this pham: EJ

Info for manual annotations of cluster EJ:

- Start number 8 was manually annotated 6 times for cluster EJ.

Gene Information:

Gene: Benry_7 Start: 5842, Stop: 5696, Start Num: 8

Candidate Starts for Benry_7:

(Start: 8 @5842 has 6 MA's), (9, 5812), (10, 5809),

Gene: Cicada_9 Start: 8141, Stop: 7989, Start Num: 8

Candidate Starts for Cicada_9:

(1, 8360), (2, 8330), (7, 8144), (Start: 8 @8141 has 6 MA's),

Gene: FireCastle_8 Start: 7976, Stop: 7830, Start Num: 8

Candidate Starts for FireCastle_8:

(4, 8012), (5, 7994), (Start: 8 @7976 has 6 MA's),

Gene: Milani_9 Start: 6875, Stop: 6723, Start Num: 8

Candidate Starts for Milani_9:

(4, 6911), (6, 6887), (Start: 8 @6875 has 6 MA's), (10, 6842), (13, 6791),

Gene: PermaG_9 Start: 8156, Stop: 8001, Start Num: 8

Candidate Starts for PermaG_9:

(3, 8225), (Start: 8 @8156 has 6 MA's), (11, 8099), (12, 8090),

Gene: Sucha_7 Start: 5839, Stop: 5696, Start Num: 8

Candidate Starts for Sucha_7:

(Start: 8 @5839 has 6 MA's), (9, 5809), (10, 5806),

Gene: Typher_9 Start: 8100, Stop: 7966, Start Num: 8

Candidate Starts for Typher_9:

(Start: 8 @8100 has 6 MA's), (10, 8067),