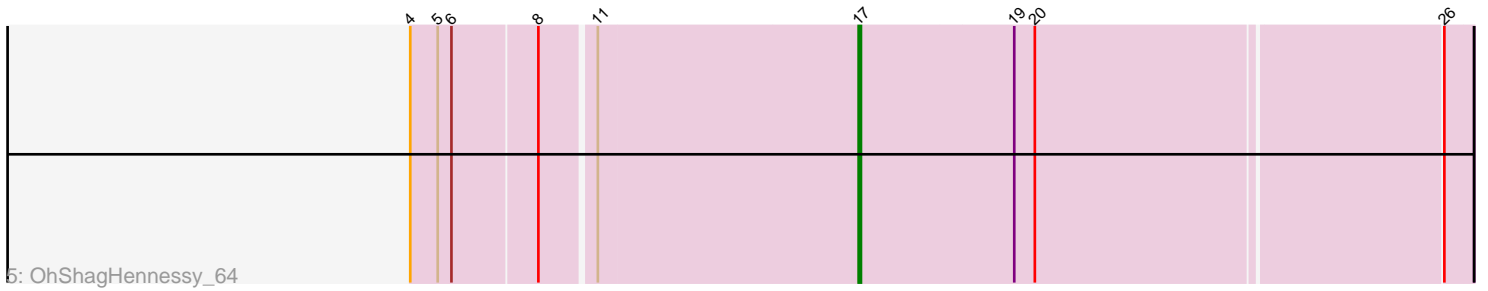
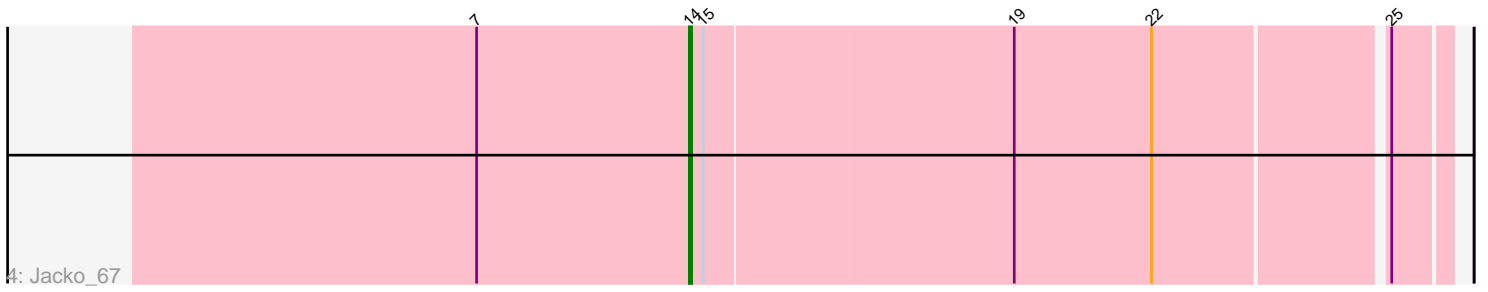
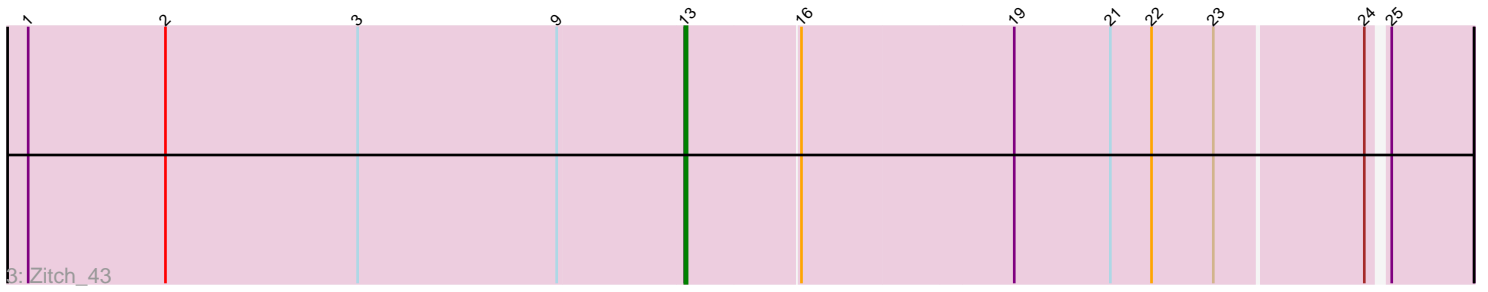
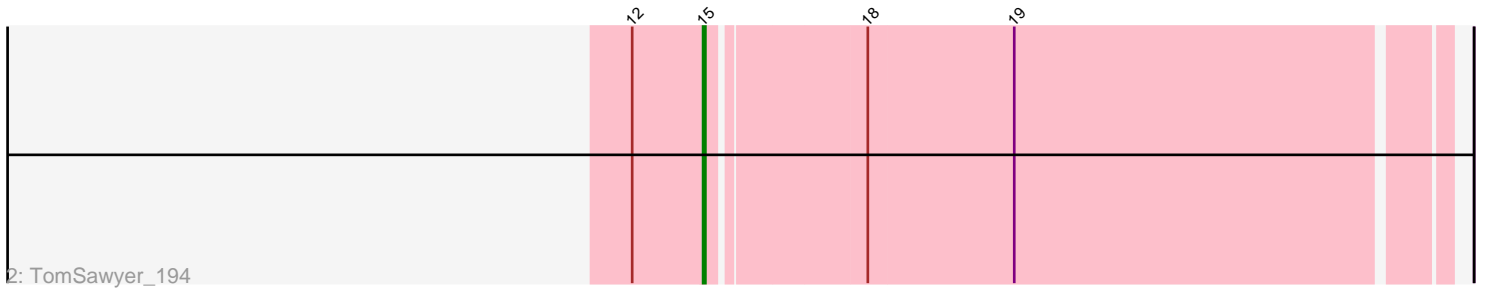
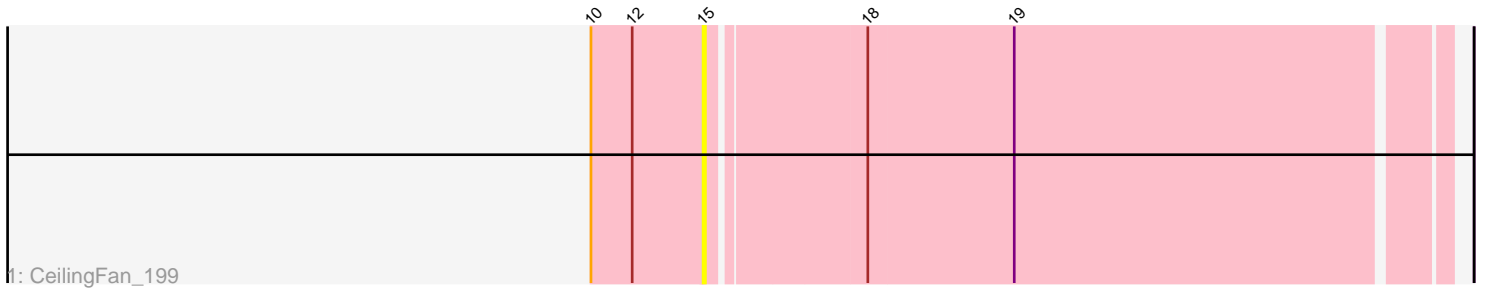


Pham 173643



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 173643 Report

This analysis was run 11/02/24 on database version 579.

Pham number 173643 has 5 members, 1 are drafts.

Phages represented in each track:

- Track 1 : CeilingFan_199
- Track 2 : TomSawyer_194
- Track 3 : Zitch_43
- Track 4 : Jacko_67
- Track 5 : OhShagHennessy_64

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

The start number called the most often in the published annotations is 15, it was called in 1 of the 4 non-draft genes in the pham.

Genes that call this "Most Annotated" start:

- CeilingFan_199, TomSawyer_194,

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

- Jacko_67,

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

- OhShagHennessy_64, Zitch_43,

Summary by start number:

Start 13:

- Found in 1 of 5 (20.0%) of genes in pham
- Manual Annotations of this start: 1 of 4
- Called 100.0% of time when present
- Phage (with cluster) where this start called: Zitch_43 (DE4),

Start 14:

- Found in 1 of 5 (20.0%) of genes in pham
- Manual Annotations of this start: 1 of 4
- Called 100.0% of time when present
- Phage (with cluster) where this start called: Jacko_67 (ED1),

Start 15:

- Found in 3 of 5 (60.0%) of genes in pham
- Manual Annotations of this start: 1 of 4
- Called 66.7% of time when present
- Phage (with cluster) where this start called: CeilingFan_199 (BE2), TomSawyer_194 (BE2),

Start 17:

- Found in 1 of 5 (20.0%) of genes in pham
- Manual Annotations of this start: 1 of 4
- Called 100.0% of time when present
- Phage (with cluster) where this start called: OhShagHennessy_64 (L1),

Summary by clusters:

There are 4 clusters represented in this pham: BE2, ED1, DE4, L1,

Info for manual annotations of cluster BE2:

- Start number 15 was manually annotated 1 time for cluster BE2.

Info for manual annotations of cluster DE4:

- Start number 13 was manually annotated 1 time for cluster DE4.

Info for manual annotations of cluster ED1:

- Start number 14 was manually annotated 1 time for cluster ED1.

Info for manual annotations of cluster L1:

- Start number 17 was manually annotated 1 time for cluster L1.

Gene Information:

Gene: CeilingFan_199 Start: 96969, Stop: 97277, Start Num: 15

Candidate Starts for CeilingFan_199:

(10, 96921), (12, 96939), (Start: 15 @96969 has 1 MA's), (18, 97032), (19, 97095),

Gene: Jacko_67 Start: 40024, Stop: 39710, Start Num: 14

Candidate Starts for Jacko_67:

(7, 40117), (Start: 14 @40024 has 1 MA's), (Start: 15 @40018 has 1 MA's), (19, 39889), (22, 39829), (25, 39733),

Gene: OhShagHennessy_64 Start: 43933, Stop: 44190, Start Num: 17

Candidate Starts for OhShagHennessy_64:

(4, 43747), (5, 43759), (6, 43765), (8, 43801), (11, 43822), (Start: 17 @43933 has 1 MA's), (19, 43999), (20, 44008), (26, 44179),

Gene: TomSawyer_194 Start: 98310, Stop: 98618, Start Num: 15

Candidate Starts for TomSawyer_194:

(12, 98280), (Start: 15 @98310 has 1 MA's), (18, 98373), (19, 98436),

Gene: Zitch_43 Start: 35193, Stop: 35522, Start Num: 13

Candidate Starts for Zitch_43:

(1, 34908), (2, 34968), (3, 35052), (9, 35139), (Start: 13 @35193 has 1 MA's), (16, 35241), (19, 35331), (21, 35373), (22, 35391), (23, 35418), (24, 35481), (25, 35487),