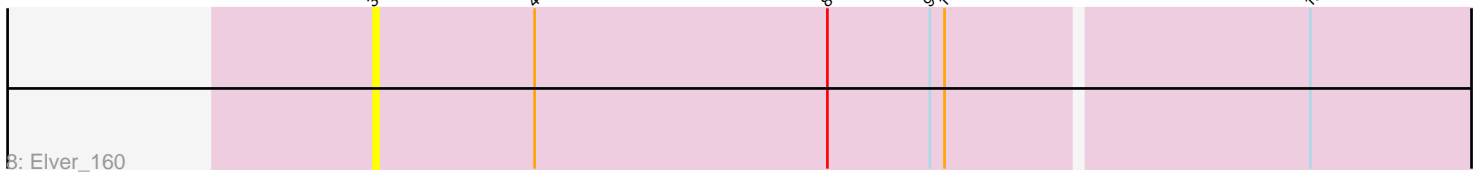
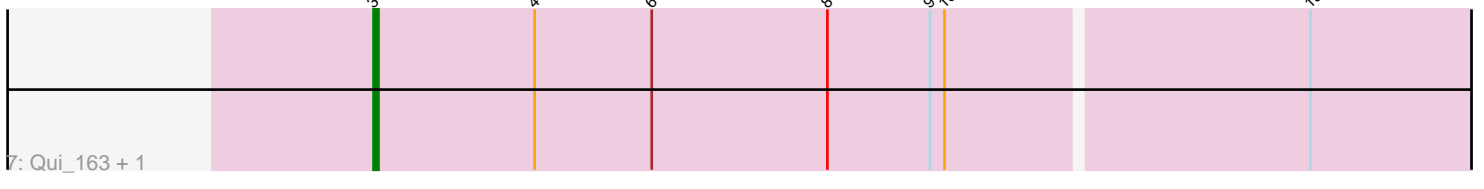
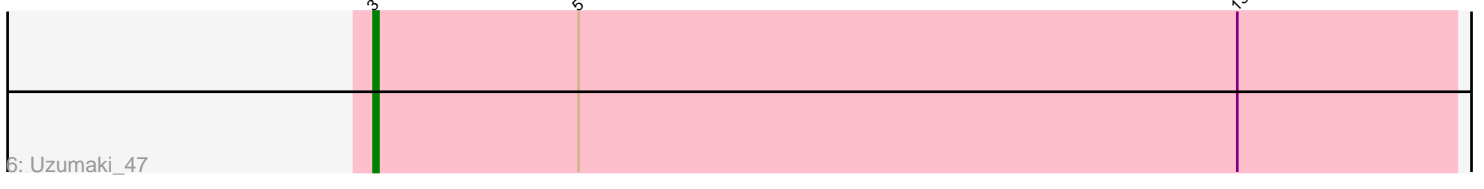
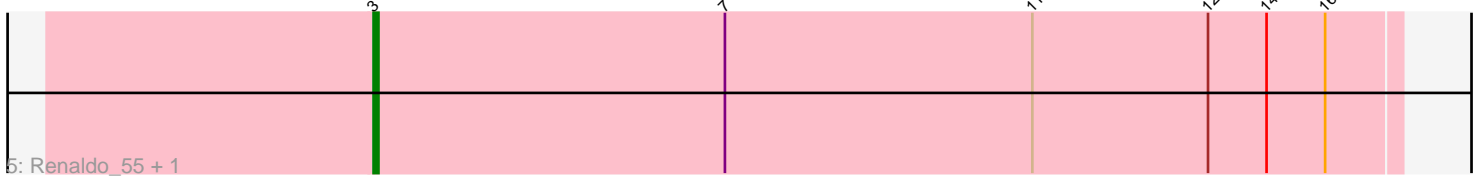
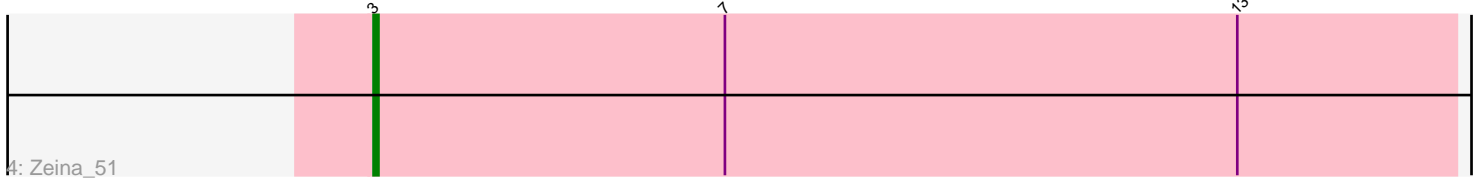
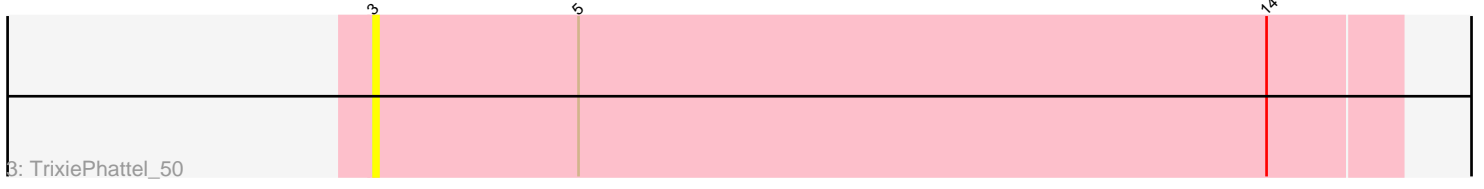
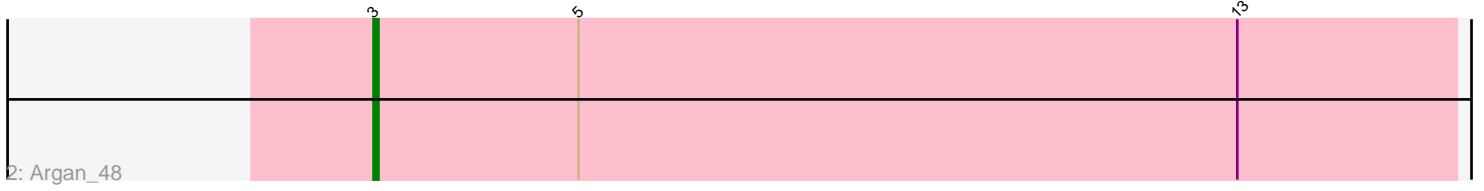
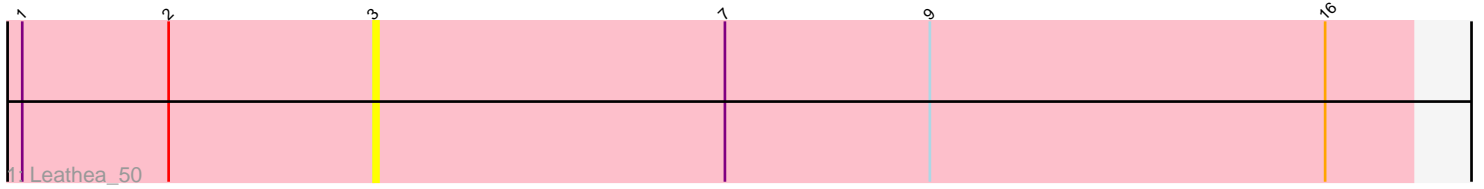


Pham 194560



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 194560 Report

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

Pham number 194560 has 10 members, 4 are drafts.

Phages represented in each track:

- Track 1 : Leathea_50
- Track 2 : Argan_48
- Track 3 : TrixiePhattel_50
- Track 4 : Zeina_51
- Track 5 : Renaldo_55, Lewando_52
- Track 6 : Uzumaki_47
- Track 7 : Qui_163, Paella_163
- Track 8 : Elver_160

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

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

Genes that call this "Most Annotated" start:

- Argan_48, Elver_160, Leathea_50, Lewando_52, Paella_163, Qui_163, Renaldo_55, TrixiePhattel_50, Uzumaki_47, Zeina_51,

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 3:

- Found in 10 of 10 (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: Argan_48 (AU6), Elver_160 (FK), Leathea_50 (AU6), Lewando_52 (AU6), Paella_163 (FK), Qui_163 (FK), Renaldo_55 (AU6), TrixiePhattel_50 (AU6), Uzumaki_47 (AU6), Zeina_51 (AU6),

Summary by clusters:

There are 2 clusters represented in this pham: FK, AU6,

Info for manual annotations of cluster AU6:

- Start number 3 was manually annotated 4 times for cluster AU6.

Info for manual annotations of cluster FK:

- Start number 3 was manually annotated 2 times for cluster FK.

Gene Information:

Gene: Argan_48 Start: 33040, Stop: 33261, Start Num: 3

Candidate Starts for Argan_48:

(Start: 3 @33040 has 6 MA's), (5, 33082), (13, 33217),

Gene: Elver_160 Start: 81364, Stop: 81597, Start Num: 3

Candidate Starts for Elver_160:

(Start: 3 @81364 has 6 MA's), (4, 81397), (8, 81457), (9, 81478), (10, 81481), (15, 81553),

Gene: Leathea_50 Start: 33107, Stop: 33319, Start Num: 3

Candidate Starts for Leathea_50:

(1, 33035), (2, 33065), (Start: 3 @33107 has 6 MA's), (7, 33179), (9, 33221), (16, 33302),

Gene: Lewando_52 Start: 34682, Stop: 34891, Start Num: 3

Candidate Starts for Lewando_52:

(Start: 3 @34682 has 6 MA's), (7, 34754), (11, 34817), (12, 34853), (14, 34865), (16, 34877),

Gene: Paella_163 Start: 82235, Stop: 82468, Start Num: 3

Candidate Starts for Paella_163:

(Start: 3 @82235 has 6 MA's), (4, 82268), (6, 82292), (8, 82328), (9, 82349), (10, 82352), (15, 82424),

Gene: Qui_163 Start: 82235, Stop: 82468, Start Num: 3

Candidate Starts for Qui_163:

(Start: 3 @82235 has 6 MA's), (4, 82268), (6, 82292), (8, 82328), (9, 82349), (10, 82352), (15, 82424),

Gene: Renaldo_55 Start: 34991, Stop: 35200, Start Num: 3

Candidate Starts for Renaldo_55:

(Start: 3 @34991 has 6 MA's), (7, 35063), (11, 35126), (12, 35162), (14, 35174), (16, 35186),

Gene: TrixiePhattel_50 Start: 33138, Stop: 33347, Start Num: 3

Candidate Starts for TrixiePhattel_50:

(Start: 3 @33138 has 6 MA's), (5, 33180), (14, 33321),

Gene: Uzumaki_47 Start: 33233, Stop: 33454, Start Num: 3

Candidate Starts for Uzumaki_47:

(Start: 3 @33233 has 6 MA's), (5, 33275), (13, 33410),

Gene: Zeina_51 Start: 33783, Stop: 34004, Start Num: 3

Candidate Starts for Zeina_51:

(Start: 3 @33783 has 6 MA's), (7, 33855), (13, 33960),

