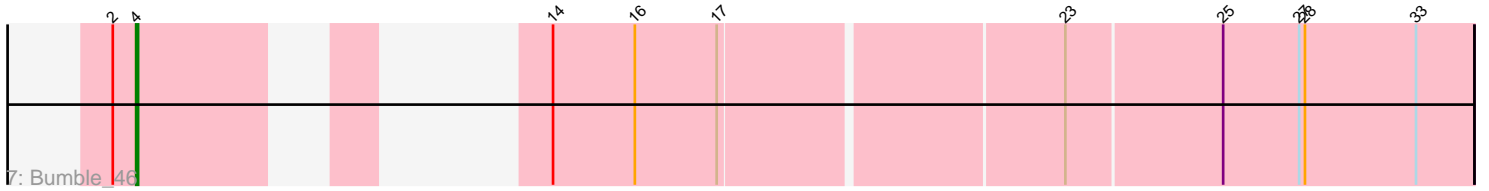
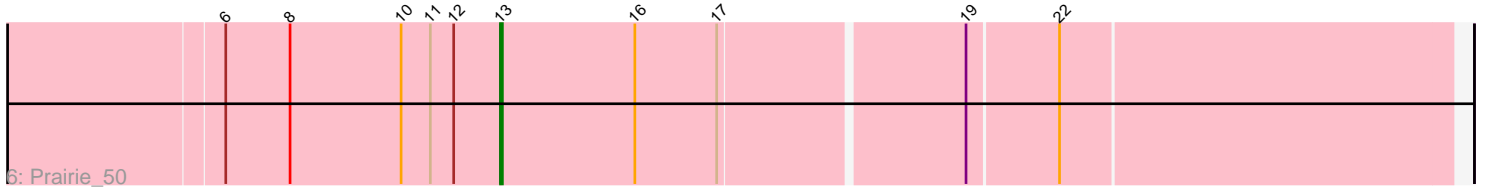
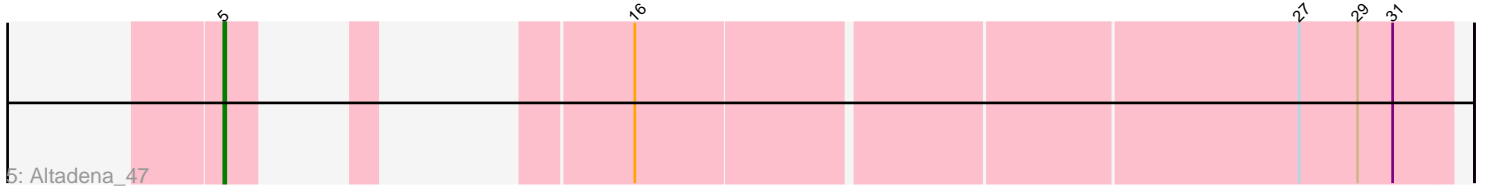
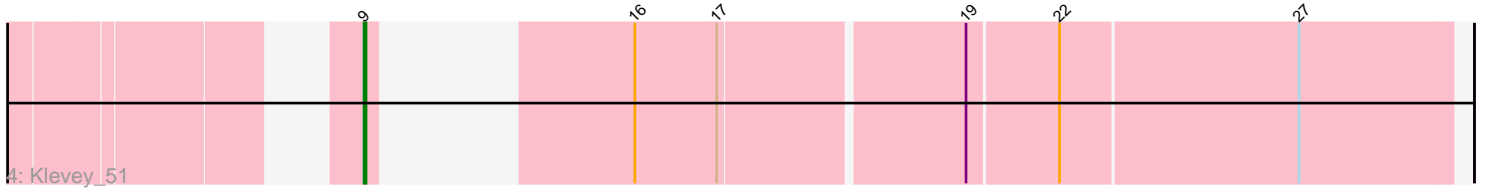
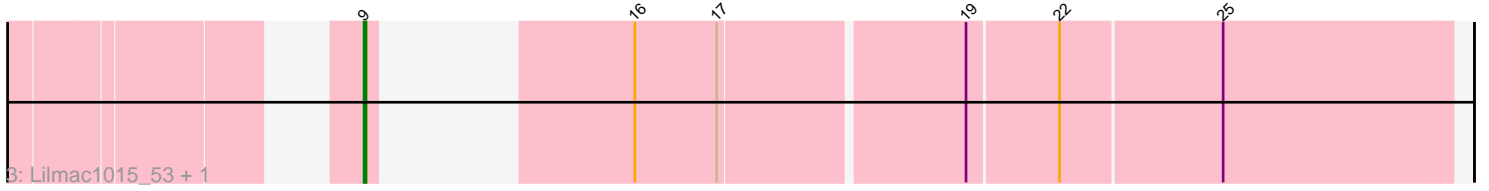
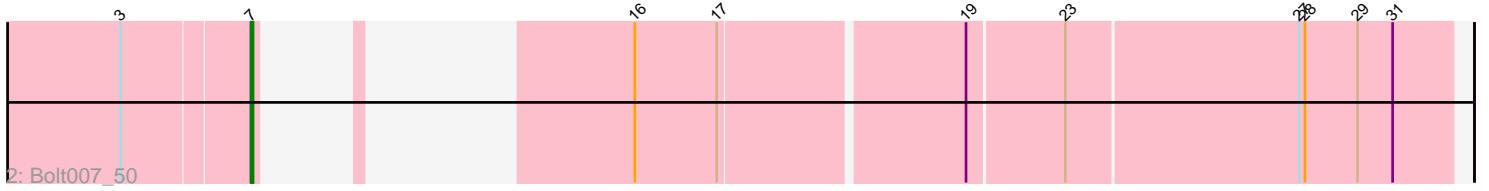
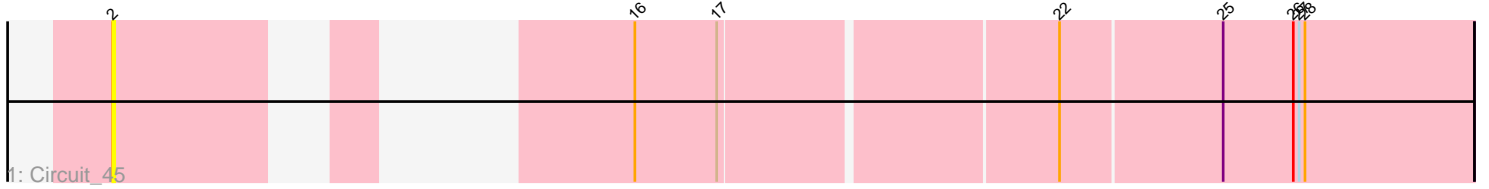


Pham 222044



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 222044 Report

This analysis was run 03/28/25 on database version 593.

Pham number 222044 has 9 members, 3 are drafts.

Phages represented in each track:

- Track 1 : Circuit_45
- Track 2 : Bolt007_50
- Track 3 : Lilmac1015_53, CalWood4100_53
- Track 4 : Klevey_51
- Track 5 : Altadena_47
- Track 6 : Prairie_50
- Track 7 : Bumble_46
- Track 8 : Lukepolites_43

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 2 of the 6 non-draft genes in the pham.

Genes that call this "Most Annotated" start:

- CalWood4100_53, Klevey_51, Lilmac1015_53,

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

-

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

- Altadena_47, Bolt007_50, Bumble_46, Circuit_45, Lukepolites_43, Prairie_50,

Summary by start number:

Start 2:

- Found in 2 of 9 (22.2%) of genes in pham
- No Manual Annotations of this start.
- Called 50.0% of time when present
- Phage (with cluster) where this start called: Circuit_45 (FH),

Start 4:

- Found in 1 of 9 (11.1%) 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: Bumble_46 (FH),

Start 5:

- Found in 1 of 9 (11.1%) 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: Altadena_47 (FH),

Start 7:

- Found in 1 of 9 (11.1%) 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: Bolt007_50 (FH),

Start 9:

- Found in 3 of 9 (33.3%) of genes in pham
- Manual Annotations of this start: 2 of 6
- Called 100.0% of time when present
- Phage (with cluster) where this start called: CalWood4100_53 (FH), Klevey_51 (FH), Lilmac1015_53 (FH),

Start 13:

- Found in 1 of 9 (11.1%) 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: Prairie_50 (FH),

Start 15:

- Found in 1 of 9 (11.1%) of genes in pham
- No Manual Annotations of this start.
- Called 100.0% of time when present
- Phage (with cluster) where this start called: Lukepolites_43 (singleton),

Summary by clusters:

There are 2 clusters represented in this pham: FH, singleton,

Info for manual annotations of cluster FH:

- Start number 4 was manually annotated 1 time for cluster FH.
- Start number 5 was manually annotated 1 time for cluster FH.
- Start number 7 was manually annotated 1 time for cluster FH.
- Start number 9 was manually annotated 2 times for cluster FH.
- Start number 13 was manually annotated 1 time for cluster FH.

Gene Information:

Gene: Altadena_47 Start: 32540, Stop: 33031, Start Num: 5

Candidate Starts for Altadena_47:

(Start: 5 @32540 has 1 MA's), (16, 32627), (27, 32954), (29, 32984), (31, 33002),

Gene: Bolt007_50 Start: 35519, Stop: 35992, Start Num: 7

Candidate Starts for Bolt007_50:

(3, 35456), (Start: 7 @35519 has 1 MA's), (16, 35588), (17, 35630), (19, 35750), (23, 35798), (27, 35915), (28, 35918), (29, 35945), (31, 35963),

Gene: Bumble_46 Start: 32813, Stop: 33379, Start Num: 4

Candidate Starts for Bumble_46:

(2, 32801), (Start: 4 @32813 has 1 MA's), (14, 32921), (16, 32963), (17, 33005), (23, 33173), (25, 33251), (27, 33290), (28, 33293), (33, 33350),

Gene: CalWood4100_53 Start: 35216, Stop: 35686, Start Num: 9

Candidate Starts for CalWood4100_53:

(Start: 9 @35216 has 2 MA's), (16, 35282), (17, 35324), (19, 35444), (22, 35489), (25, 35570),

Gene: Circuit_45 Start: 33675, Stop: 34253, Start Num: 2

Candidate Starts for Circuit_45:

(2, 33675), (16, 33837), (17, 33879), (22, 34044), (25, 34125), (26, 34161), (27, 34164), (28, 34167),

Gene: Klevey_51 Start: 34652, Stop: 35122, Start Num: 9

Candidate Starts for Klevey_51:

(Start: 9 @34652 has 2 MA's), (16, 34718), (17, 34760), (19, 34880), (22, 34925), (27, 35045),

Gene: Lilmac1015_53 Start: 35216, Stop: 35686, Start Num: 9

Candidate Starts for Lilmac1015_53:

(Start: 9 @35216 has 2 MA's), (16, 35282), (17, 35324), (19, 35444), (22, 35489), (25, 35570),

Gene: Lukepolites_43 Start: 33328, Stop: 33786, Start Num: 15

Candidate Starts for Lukepolites_43:

(1, 33172), (15, 33328), (18, 33499), (20, 33556), (21, 33574), (24, 33604), (30, 33742), (32, 33754), (34, 33775),

Gene: Prairie_50 Start: 34461, Stop: 34934, Start Num: 13

Candidate Starts for Prairie_50:

(6, 34320), (8, 34353), (10, 34410), (11, 34425), (12, 34437), (Start: 13 @34461 has 1 MA's), (16, 34530), (17, 34572), (19, 34692), (22, 34737),