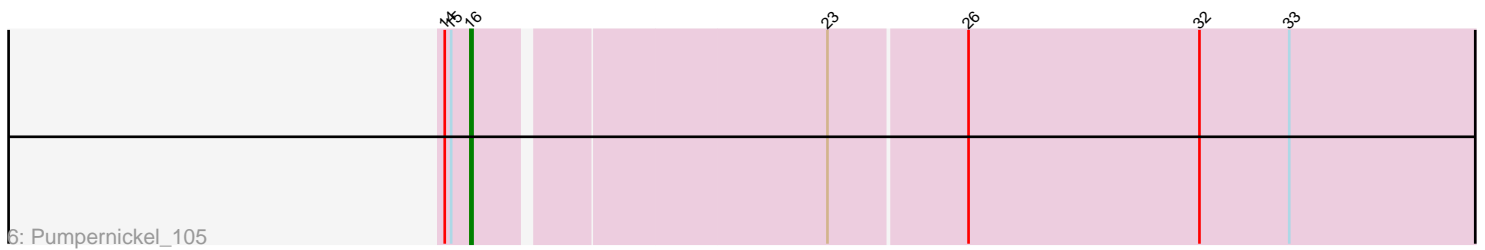
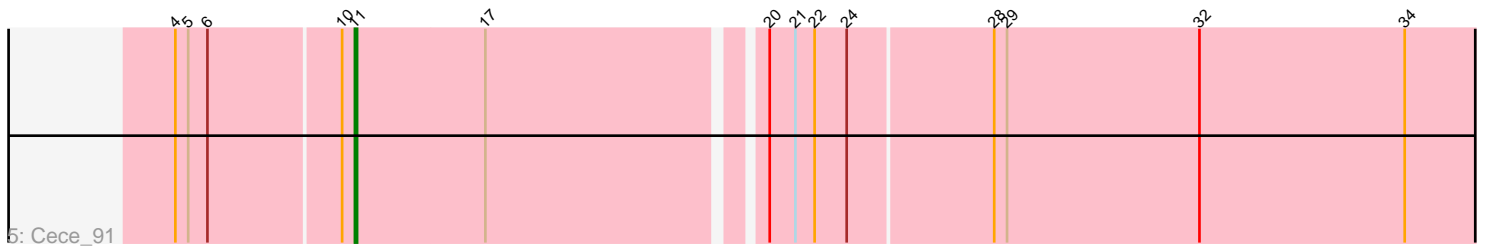
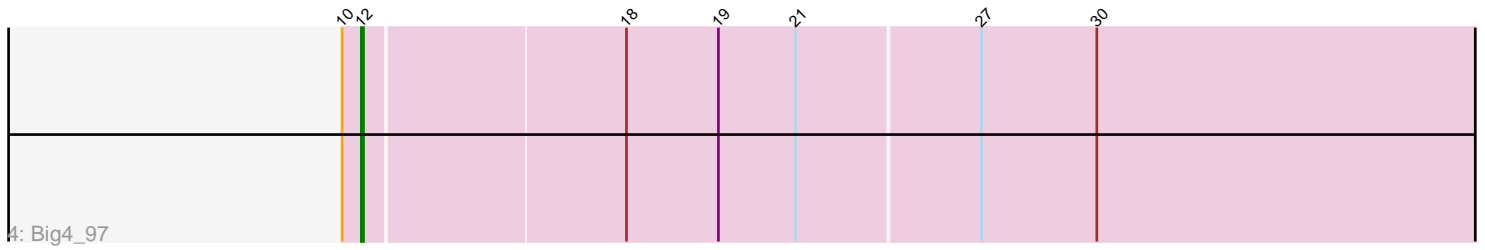
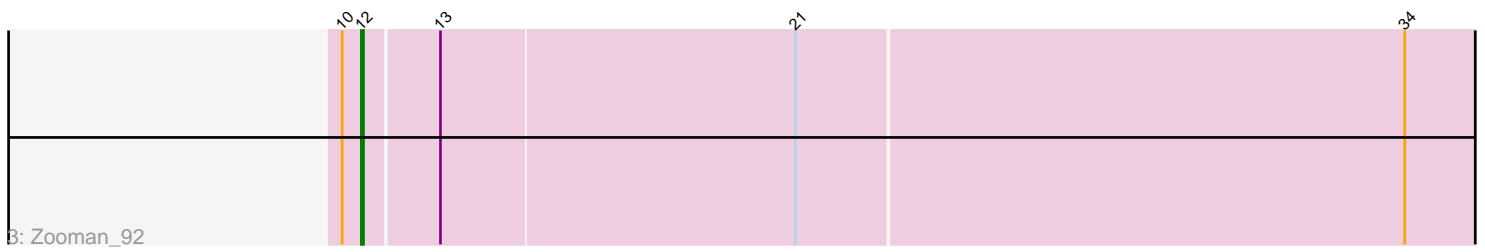
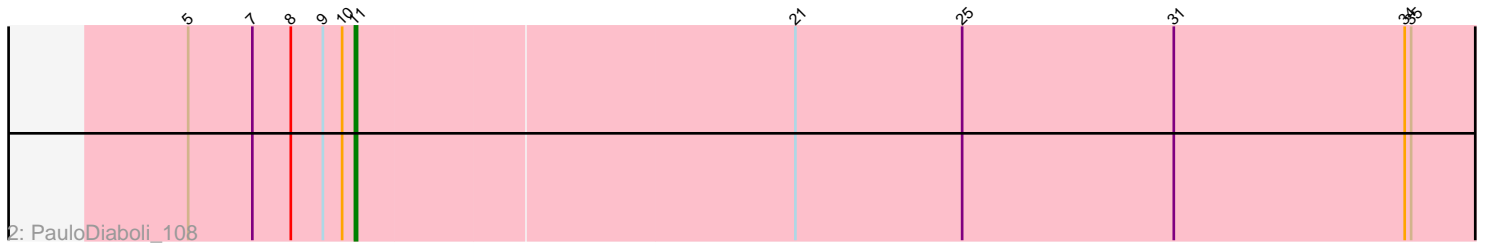
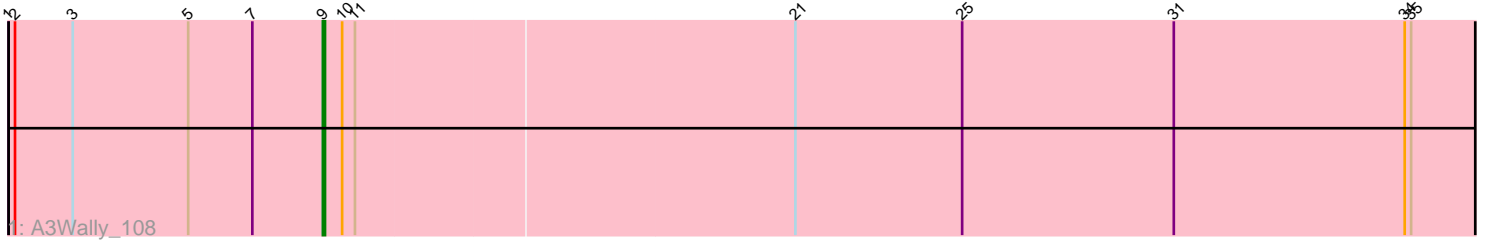


Pham 170670



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 170670 Report

This analysis was run 07/09/24 on database version 566.

Pham number 170670 has 6 members, 0 are drafts.

Phages represented in each track:

- Track 1 : A3Wally_108
- Track 2 : PauloDiaboli_108
- Track 3 : Zooman_92
- Track 4 : Big4_97
- Track 5 : Cece_91
- Track 6 : Pumpernickel_105

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

The start number called the most often in the published annotations is 11, it was called in 2 of the 6 non-draft genes in the pham.

Genes that call this "Most Annotated" start:

- Cece_91, PauloDiaboli_108,

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

- A3Wally_108,

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

- Big4_97, Pumpernickel_105, Zooman_92,

Summary by start number:

Start 9:

- Found in 2 of 6 (33.3%) of genes in pham
- Manual Annotations of this start: 1 of 6
- Called 50.0% of time when present
- Phage (with cluster) where this start called: A3Wally_108 (GD1),

Start 11:

- Found in 3 of 6 (50.0%) of genes in pham
- Manual Annotations of this start: 2 of 6
- Called 66.7% of time when present

- Phage (with cluster) where this start called: Cece_91 (GD3), PauloDiaboli_108 (GD1),

Start 12:

- Found in 2 of 6 (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: Big4_97 (GD2), Zooman_92 (GD2),

Start 16:

- Found in 1 of 6 (16.7%) 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: Pumpernickel_105 (GD4),

Summary by clusters:

There are 4 clusters represented in this pham: GD1, GD2, GD3, GD4,

Info for manual annotations of cluster GD1:

- Start number 9 was manually annotated 1 time for cluster GD1.
- Start number 11 was manually annotated 1 time for cluster GD1.

Info for manual annotations of cluster GD2:

- Start number 12 was manually annotated 2 times for cluster GD2.

Info for manual annotations of cluster GD3:

- Start number 11 was manually annotated 1 time for cluster GD3.

Info for manual annotations of cluster GD4:

- Start number 16 was manually annotated 1 time for cluster GD4.

Gene Information:

Gene: A3Wally_108 Start: 73992, Stop: 73408, Start Num: 9

Candidate Starts for A3Wally_108:

(1, 74139), (2, 74136), (3, 74109), (5, 74055), (7, 74025), (Start: 9 @73992 has 1 MA's), (10, 73983), (Start: 11 @73977 has 2 MA's), (21, 73776), (25, 73698), (31, 73599), (34, 73491), (35, 73488),

Gene: Big4_97 Start: 73090, Stop: 72530, Start Num: 12

Candidate Starts for Big4_97:

(10, 73099), (Start: 12 @73090 has 2 MA's), (18, 72973), (19, 72931), (21, 72895), (27, 72811), (30, 72757),

Gene: Cece_91 Start: 76156, Stop: 75599, Start Num: 11

Candidate Starts for Cece_91:

(4, 76237), (5, 76231), (6, 76222), (10, 76162), (Start: 11 @76156 has 2 MA's), (17, 76096), (20, 75976), (21, 75964), (22, 75955), (24, 75940), (28, 75874), (29, 75868), (32, 75778), (34, 75682),

Gene: PauloDiaboli_108 Start: 73334, Stop: 72765, Start Num: 11

Candidate Starts for PauloDiaboli_108:

(5, 73412), (7, 73382), (8, 73364), (Start: 9 @73349 has 1 MA's), (10, 73340), (Start: 11 @73334 has 2 MA's), (21, 73133), (25, 73055), (31, 72956), (34, 72848), (35, 72845),

Gene: Pumpernickel_105 Start: 74888, Stop: 74379, Start Num: 16

Candidate Starts for Pumpernickel_105:

(14, 74900), (15, 74897), (Start: 16 @74888 has 1 MA's), (23, 74732), (26, 74669), (32, 74561), (33, 74519),

Gene: Zooman_92 Start: 70938, Stop: 70378, Start Num: 12

Candidate Starts for Zooman_92:

(10, 70947), (Start: 12 @70938 has 2 MA's), (13, 70905), (21, 70743), (34, 70461),