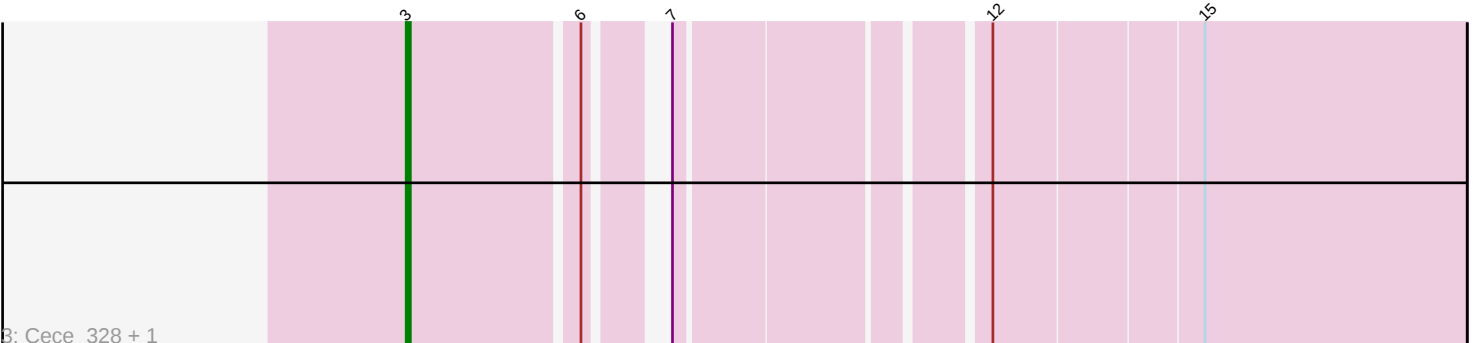
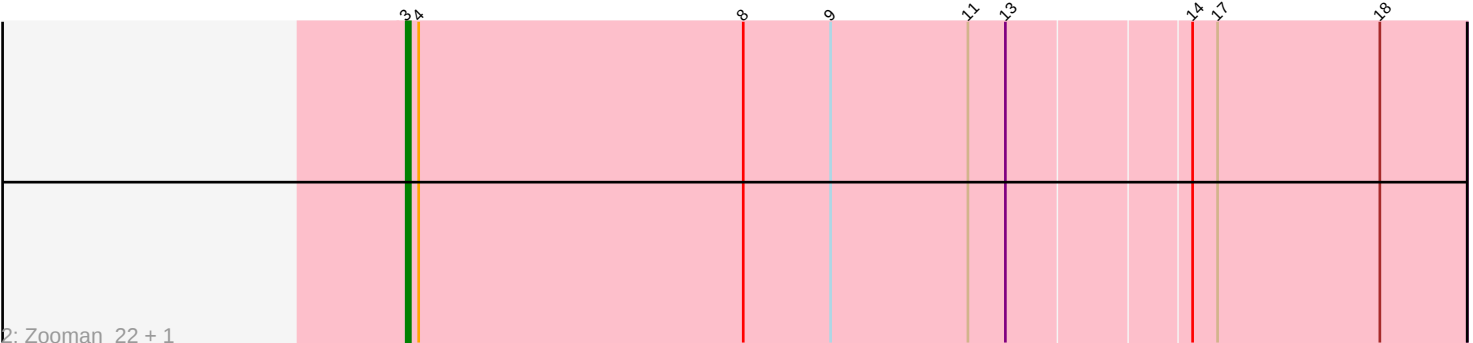
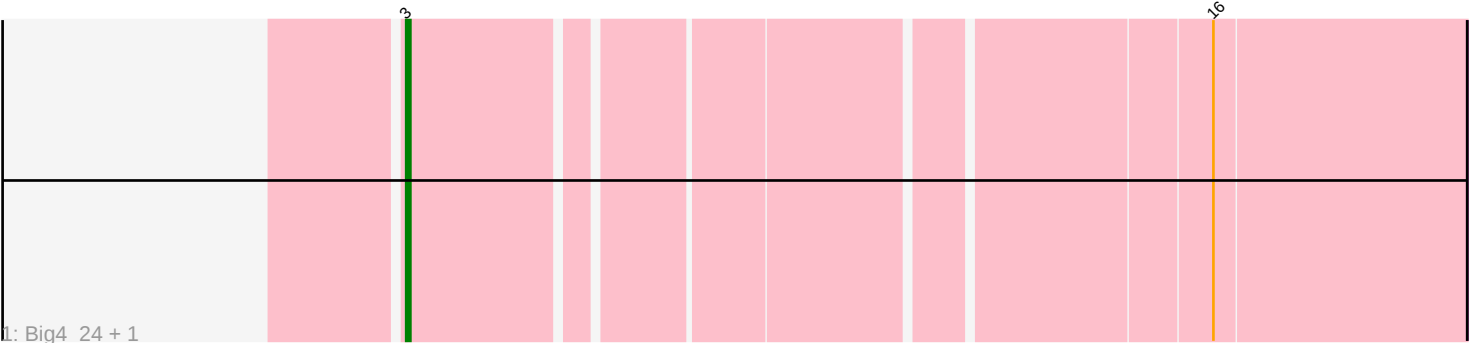




Note: Tracks are now grouped by subcluster and scaled. Switching in subcluster is indicated by changes in track color. Track scale is now set by default to display the region 30 bp upstream of start 1 to 30 bp downstream of the last possible start. If this default region is judged to be packed too tightly 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 309976 Report

This analysis was run 09/05/26 on database version 662.

Pham number 309976 has 8 members, 0 are drafts.

Phages represented in each track:

- Track 1 : Big4_24, Big4_350
- Track 2 : Zooman_22, Zooman_335
- Track 3 : Cece_328, Cece_26
- Track 4 : Pumpernickel_28, Pumpernickel_329

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 8 non-draft genes in the pham.

Genes that call this "Most Annotated" start:

- Big4_24, Big4_350, Cece_26, Cece_328, Zooman_22, Zooman_335,

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

-

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

- Pumpernickel_28, Pumpernickel_329,

Summary by start number:

Start 3:

- Found in 6 of 8 (75.0%) of genes in pham
- Manual Annotations of this start: 6 of 8
- Called 100.0% of time when present
- Phage (with cluster) where this start called: Big4_24 (GD2), Big4_350 (GD2), Cece_26 (GD3), Cece_328 (GD3), Zooman_22 (GD2), Zooman_335 (GD2),

Start 5:

- Found in 2 of 8 (25.0%) of genes in pham
- Manual Annotations of this start: 2 of 8
- Called 100.0% of time when present
- Phage (with cluster) where this start called: Pumpernickel_28 (GD4), Pumpernickel_329 (GD4),

Summary by clusters:

There are 3 clusters represented in this pham: GD2, GD3, GD4,

Info for manual annotations of cluster GD2:

- Start number 3 was manually annotated 4 times for cluster GD2.

Info for manual annotations of cluster GD3:

- Start number 3 was manually annotated 2 times for cluster GD3.

Info for manual annotations of cluster GD4:

- Start number 5 was manually annotated 2 times for cluster GD4.

Gene Information:

Gene: Big4_24 Start: 10420, Stop: 10656, Start Num: 3

Candidate Starts for Big4_24:

(Start: 3 @10420 has 6 MA's), (16, 10597),

Gene: Big4_350 Start: 185114, Stop: 185350, Start Num: 3

Candidate Starts for Big4_350:

(Start: 3 @185114 has 6 MA's), (16, 185291),

Gene: Cece_328 Start: 177963, Stop: 178190, Start Num: 3

Candidate Starts for Cece_328:

(Start: 3 @177963 has 6 MA's), (6, 178002), (7, 178014), (12, 178080), (15, 178128),

Gene: Cece_26 Start: 9529, Stop: 9756, Start Num: 3

Candidate Starts for Cece_26:

(Start: 3 @9529 has 6 MA's), (6, 9568), (7, 9580), (12, 9646), (15, 9694),

Gene: Pumpernickel_28 Start: 11370, Stop: 11594, Start Num: 5

Candidate Starts for Pumpernickel_28:

(1, 11277), (2, 11295), (Start: 5 @11370 has 2 MA's), (10, 11469), (19, 11583),

Gene: Pumpernickel_329 Start: 177502, Stop: 177726, Start Num: 5

Candidate Starts for Pumpernickel_329:

(1, 177409), (2, 177427), (Start: 5 @177502 has 2 MA's), (10, 177601), (19, 177715),

Gene: Zooman_22 Start: 9803, Stop: 10054, Start Num: 3

Candidate Starts for Zooman_22:

(Start: 3 @9803 has 6 MA's), (4, 9806), (8, 9884), (9, 9905), (11, 9938), (13, 9947), (14, 9989), (17, 9995), (18, 10034),

Gene: Zooman_335 Start: 185454, Stop: 185705, Start Num: 3

Candidate Starts for Zooman_335:

(Start: 3 @185454 has 6 MA's), (4, 185457), (8, 185535), (9, 185556), (11, 185589), (13, 185598), (14, 185640), (17, 185646), (18, 185685),