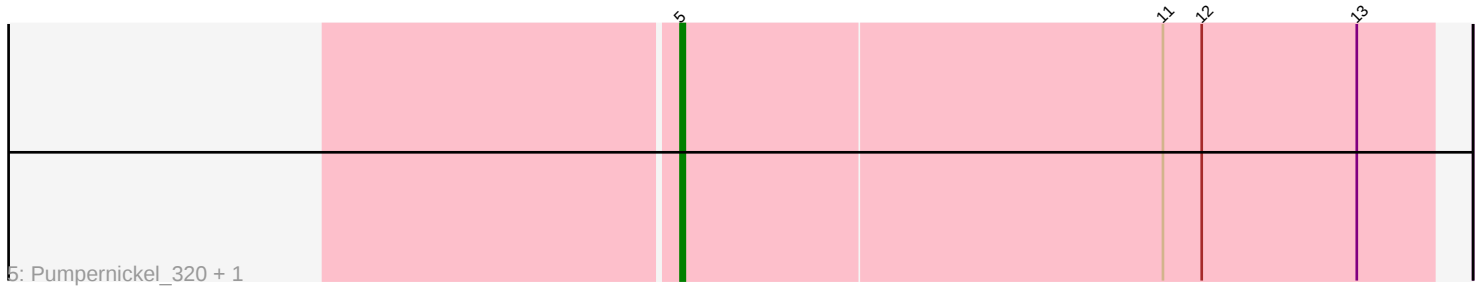
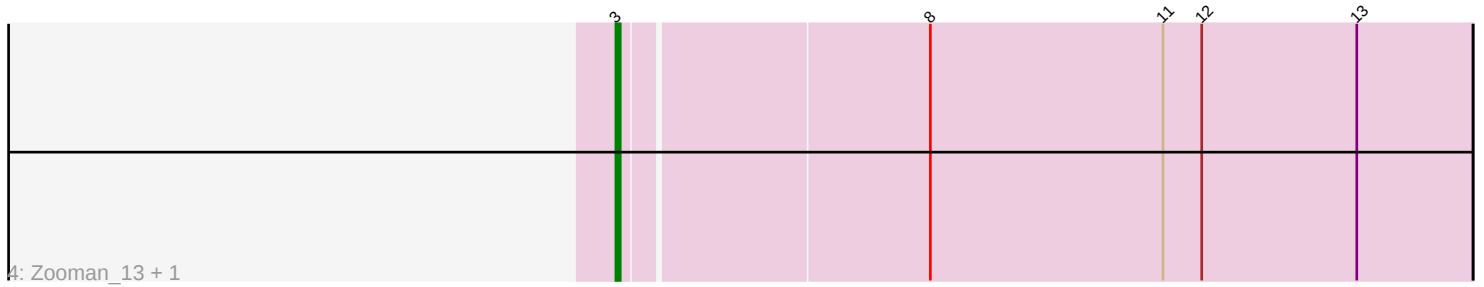
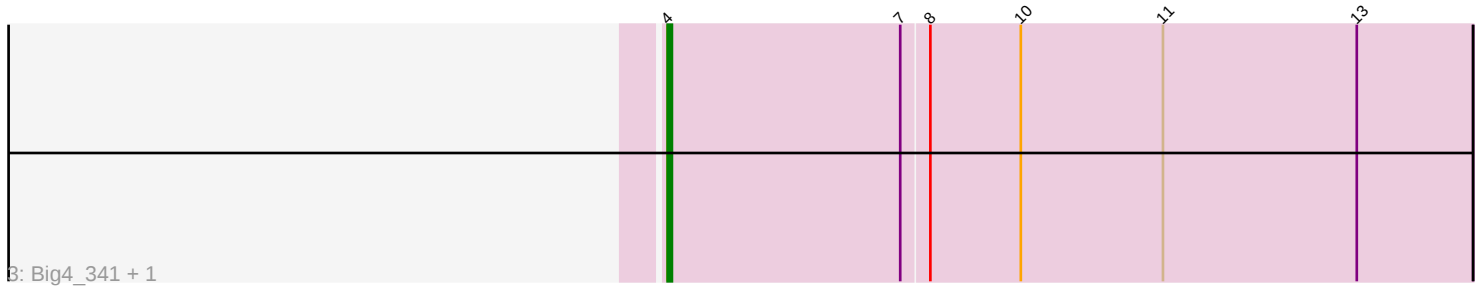
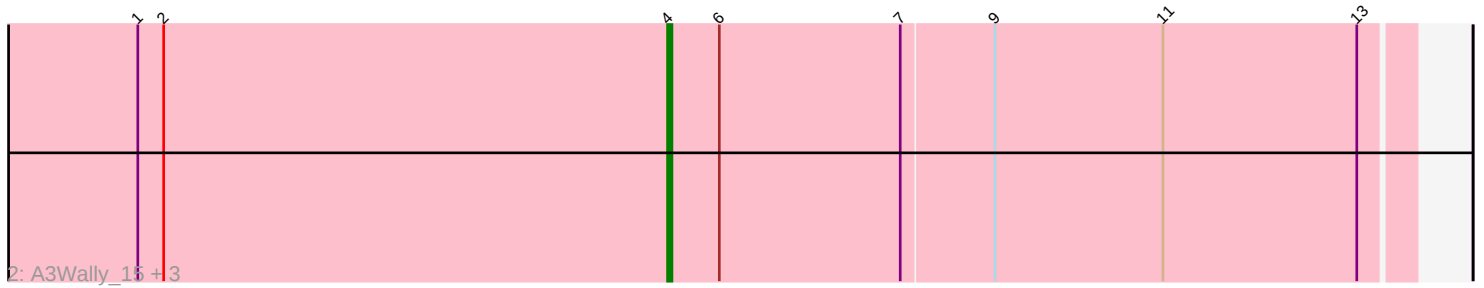
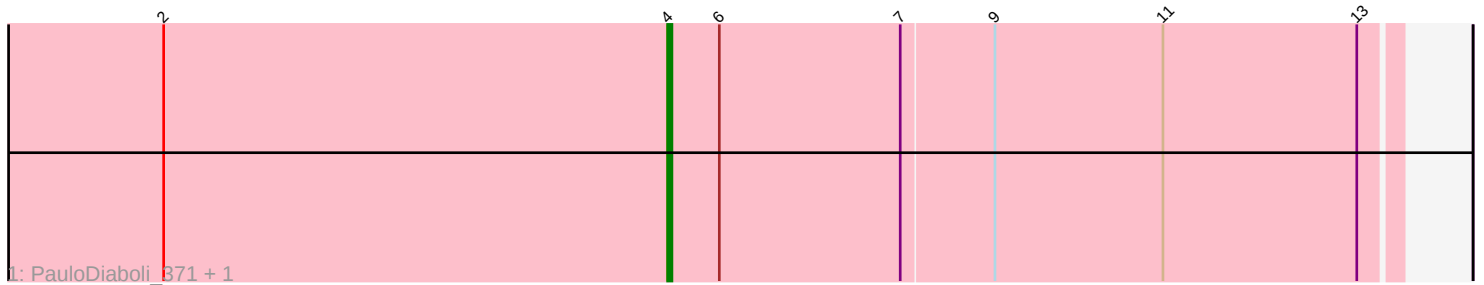


Pham 309658



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 309658 Report

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

Pham number 309658 has 12 members, 0 are drafts.

Phages represented in each track:

- Track 1 : PauloDiaboli_371, PauloDiaboli_16
- Track 2 : A3Wally_15, Dodo_362, Dodo_16, A3Wally_368
- Track 3 : Big4_341, Big4_15
- Track 4 : Zooman_13, Zooman_326
- Track 5 : Pumpernickel_320, Pumpernickel_19

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

The start number called the most often in the published annotations is 4, it was called in 8 of the 12 non-draft genes in the pham.

Genes that call this "Most Annotated" start:

- A3Wally_15, A3Wally_368, Big4_15, Big4_341, Dodo_16, Dodo_362, PauloDiaboli_16, PauloDiaboli_371,

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

-

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

- Pumpernickel_19, Pumpernickel_320, Zooman_13, Zooman_326,

Summary by start number:

Start 3:

- Found in 2 of 12 (16.7%) of genes in pham
- Manual Annotations of this start: 2 of 12
- Called 100.0% of time when present
- Phage (with cluster) where this start called: Zooman_13 (GD2), Zooman_326 (GD2),

Start 4:

- Found in 8 of 12 (66.7%) of genes in pham
- Manual Annotations of this start: 8 of 12
- Called 100.0% of time when present

- Phage (with cluster) where this start called: A3Wally_15 (GD1), A3Wally_368 (GD1), Big4_15 (GD2), Big4_341 (GD2), Dodo_16 (GD1), Dodo_362 (GD1), PauloDiaboli_16 (GD1), PauloDiaboli_371 (GD1),

Start 5:

- Found in 2 of 12 (16.7%) of genes in pham
- Manual Annotations of this start: 2 of 12
- Called 100.0% of time when present
- Phage (with cluster) where this start called: Pumpernickel_19 (GD4), Pumpernickel_320 (GD4),

Summary by clusters:

There are 3 clusters represented in this pham: GD1, GD2, GD4,

Info for manual annotations of cluster GD1:

- Start number 4 was manually annotated 6 times for cluster GD1.

Info for manual annotations of cluster GD2:

- Start number 3 was manually annotated 2 times for cluster GD2.
- Start number 4 was manually annotated 2 times for cluster GD2.

Info for manual annotations of cluster GD4:

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

Gene Information:

Gene: A3Wally_15 Start: 5373, Stop: 5543, Start Num: 4

Candidate Starts for A3Wally_15:

(1, 5250), (2, 5256), (Start: 4 @5373 has 8 MA's), (6, 5385), (7, 5427), (9, 5448), (11, 5487), (13, 5532),

Gene: A3Wally_368 Start: 184594, Stop: 184764, Start Num: 4

Candidate Starts for A3Wally_368:

(1, 184471), (2, 184477), (Start: 4 @184594 has 8 MA's), (6, 184606), (7, 184648), (9, 184669), (11, 184708), (13, 184753),

Gene: Big4_341 Start: 181671, Stop: 181856, Start Num: 4

Candidate Starts for Big4_341:

(Start: 4 @181671 has 8 MA's), (7, 181725), (8, 181731), (10, 181752), (11, 181785), (13, 181830),

Gene: Big4_15 Start: 6977, Stop: 7162, Start Num: 4

Candidate Starts for Big4_15:

(Start: 4 @6977 has 8 MA's), (7, 7031), (8, 7037), (10, 7058), (11, 7091), (13, 7136),

Gene: Dodo_362 Start: 183679, Stop: 183846, Start Num: 4

Candidate Starts for Dodo_362:

(1, 183556), (2, 183562), (Start: 4 @183679 has 8 MA's), (6, 183691), (7, 183733), (9, 183754), (11, 183793), (13, 183838),

Gene: Dodo_16 Start: 5479, Stop: 5646, Start Num: 4

Candidate Starts for Dodo_16:

(1, 5356), (2, 5362), (Start: 4 @5479 has 8 MA's), (6, 5491), (7, 5533), (9, 5554), (11, 5593), (13, 5638),

Gene: PauloDiaboli_371 Start: 182101, Stop: 182268, Start Num: 4

Candidate Starts for PauloDiaboli_371:

(2, 181984), (Start: 4 @182101 has 8 MA's), (6, 182113), (7, 182155), (9, 182176), (11, 182215), (13, 182260),

Gene: PauloDiaboli_16 Start: 5472, Stop: 5639, Start Num: 4

Candidate Starts for PauloDiaboli_16:

(2, 5355), (Start: 4 @5472 has 8 MA's), (6, 5484), (7, 5526), (9, 5547), (11, 5586), (13, 5631),

Gene: Pumpernickel_320 Start: 173874, Stop: 174047, Start Num: 5

Candidate Starts for Pumpernickel_320:

(Start: 5 @173874 has 2 MA's), (11, 173985), (12, 173994), (13, 174030),

Gene: Pumpernickel_19 Start: 7742, Stop: 7915, Start Num: 5

Candidate Starts for Pumpernickel_19:

(Start: 5 @7742 has 2 MA's), (11, 7853), (12, 7862), (13, 7898),

Gene: Zooman_13 Start: 6256, Stop: 6450, Start Num: 3

Candidate Starts for Zooman_13:

(Start: 3 @6256 has 2 MA's), (8, 6325), (11, 6379), (12, 6388), (13, 6424),

Gene: Zooman_326 Start: 181907, Stop: 182101, Start Num: 3

Candidate Starts for Zooman_326:

(Start: 3 @181907 has 2 MA's), (8, 181976), (11, 182030), (12, 182039), (13, 182075),