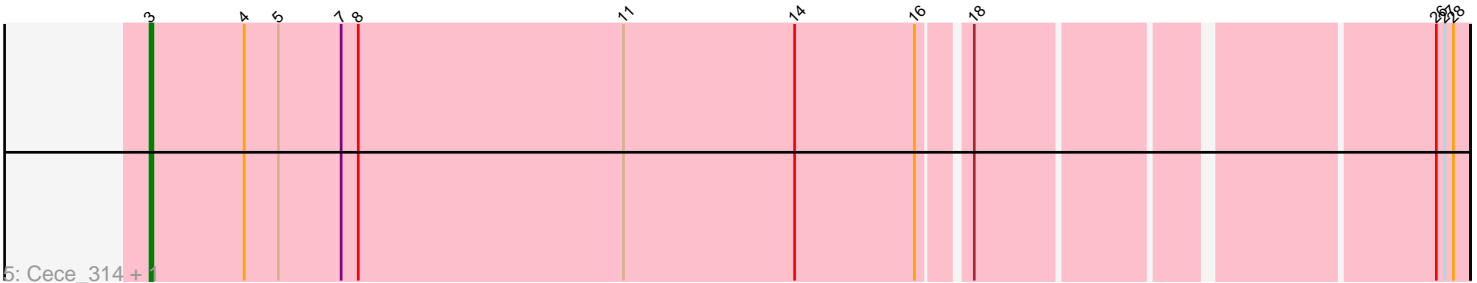
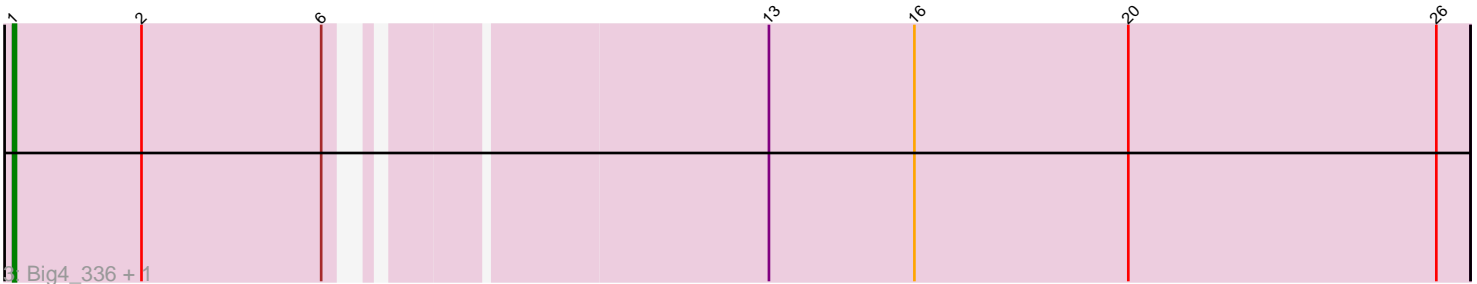
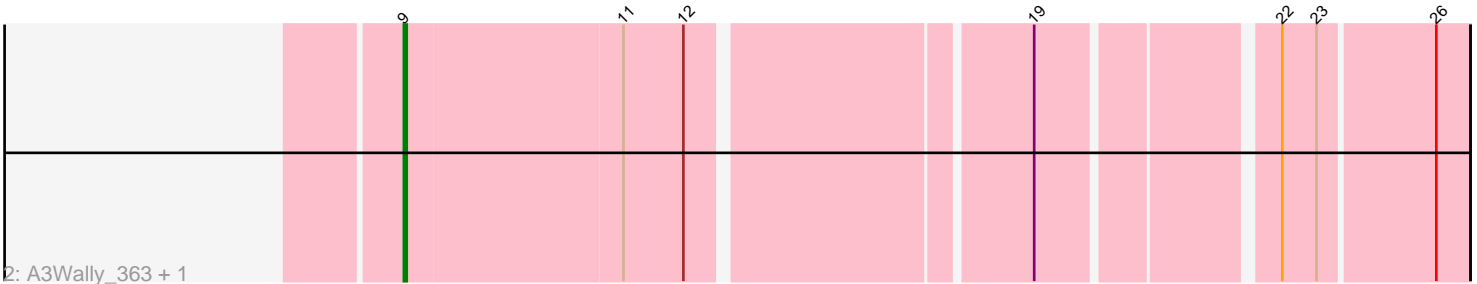
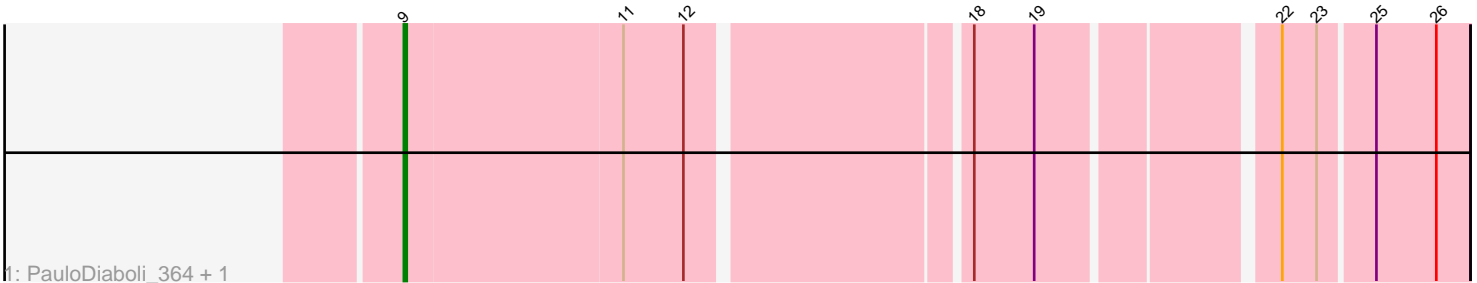


Pham 5699



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 5699 Report

This analysis was run 04/28/24 on database version 559.

Pham number 5699 has 10 members, 0 are drafts.

Phages represented in each track:

- Track 1 : PauloDiaboli\_364, PauloDiaboli\_9
- Track 2 : A3Wally\_363, A3Wally\_9
- Track 3 : Big4\_336, Big4\_10
- Track 4 : Zooman\_321, Zooman\_8
- Track 5 : Cece\_314, Cece\_12

### ***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 4 of the 10 non-draft genes in the pham.

Genes that call this "Most Annotated" start:

- A3Wally\_363, A3Wally\_9, PauloDiaboli\_364, PauloDiaboli\_9,

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

- 

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

- Big4\_10, Big4\_336, Cece\_12, Cece\_314, Zooman\_321, Zooman\_8,

### **Summary by start number:**

Start 1:

- Found in 2 of 10 ( 20.0% ) of genes in pham
- Manual Annotations of this start: 2 of 10
- Called 100.0% of time when present
- Phage (with cluster) where this start called: Big4\_10 (GD2), Big4\_336 (GD2),

Start 3:

- Found in 2 of 10 ( 20.0% ) of genes in pham
- Manual Annotations of this start: 2 of 10
- Called 100.0% of time when present
- Phage (with cluster) where this start called: Cece\_12 (GD3), Cece\_314 (GD3),

Start 6:

- Found in 4 of 10 ( 40.0% ) of genes in pham
- Manual Annotations of this start: 2 of 10
- Called 50.0% of time when present
- Phage (with cluster) where this start called: Zooman\_321 (GD2), Zooman\_8 (GD2),

Start 9:

- Found in 4 of 10 ( 40.0% ) of genes in pham
- Manual Annotations of this start: 4 of 10
- Called 100.0% of time when present
- Phage (with cluster) where this start called: A3Wally\_363 (GD1), A3Wally\_9 (GD1), PauloDiaboli\_364 (GD1), PauloDiaboli\_9 (GD1),

### **Summary by clusters:**

There are 3 clusters represented in this pham: GD1, GD2, GD3,

Info for manual annotations of cluster GD1:

- Start number 9 was manually annotated 4 times for cluster GD1.

Info for manual annotations of cluster GD2:

- Start number 1 was manually annotated 2 times for cluster GD2.
- Start number 6 was manually annotated 2 times for cluster GD2.

Info for manual annotations of cluster GD3:

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

### **Gene Information:**

Gene: A3Wally\_363 Start: 182572, Stop: 182916, Start Num: 9

Candidate Starts for A3Wally\_363:

(Start: 9 @182572 has 4 MA's), (11, 182647), (12, 182668), (19, 182779), (22, 182854), (23, 182866), (26, 182905),

Gene: A3Wally\_9 Start: 3351, Stop: 3695, Start Num: 9

Candidate Starts for A3Wally\_9:

(Start: 9 @3351 has 4 MA's), (11, 3426), (12, 3447), (19, 3558), (22, 3633), (23, 3645), (26, 3684),

Gene: Big4\_336 Start: 178974, Stop: 179462, Start Num: 1

Candidate Starts for Big4\_336:

(Start: 1 @178974 has 2 MA's), (2, 179019), (Start: 6 @179082 has 2 MA's), (13, 179217), (16, 179268), (20, 179343), (26, 179451),

Gene: Big4\_10 Start: 4280, Stop: 4768, Start Num: 1

Candidate Starts for Big4\_10:

(Start: 1 @4280 has 2 MA's), (2, 4325), (Start: 6 @4388 has 2 MA's), (13, 4523), (16, 4574), (20, 4649), (26, 4757),

Gene: Cece\_314 Start: 172354, Stop: 172794, Start Num: 3

Candidate Starts for Cece\_314:

(Start: 3 @172354 has 2 MA's), (4, 172387), (5, 172399), (7, 172420), (8, 172426), (11, 172519), (14, 172579), (16, 172621), (18, 172636), (26, 172783), (27, 172786), (28, 172789),

Gene: Cece\_12 Start: 3920, Stop: 4360, Start Num: 3

Candidate Starts for Cece\_12:

(Start: 3 @3920 has 2 MA's), (4, 3953), (5, 3965), (7, 3986), (8, 3992), (11, 4085), (14, 4145), (16, 4187), (18, 4202), (26, 4349), (27, 4352), (28, 4355),

Gene: PauloDiaboli\_364 Start: 179815, Stop: 180159, Start Num: 9

Candidate Starts for PauloDiaboli\_364:

(Start: 9 @179815 has 4 MA's), (11, 179890), (12, 179911), (18, 180001), (19, 180022), (22, 180097), (23, 180109), (25, 180127), (26, 180148),

Gene: PauloDiaboli\_9 Start: 3186, Stop: 3530, Start Num: 9

Candidate Starts for PauloDiaboli\_9:

(Start: 9 @3186 has 4 MA's), (11, 3261), (12, 3282), (18, 3372), (19, 3393), (22, 3468), (23, 3480), (25, 3498), (26, 3519),

Gene: Zooman\_321 Start: 179227, Stop: 179628, Start Num: 6

Candidate Starts for Zooman\_321:

(Start: 6 @179227 has 2 MA's), (10, 179266), (15, 179416), (17, 179440), (21, 179548), (22, 179563), (24, 179590), (26, 179617),

Gene: Zooman\_8 Start: 3576, Stop: 3977, Start Num: 6

Candidate Starts for Zooman\_8:

(Start: 6 @3576 has 2 MA's), (10, 3615), (15, 3765), (17, 3789), (21, 3897), (22, 3912), (24, 3939), (26, 3966),