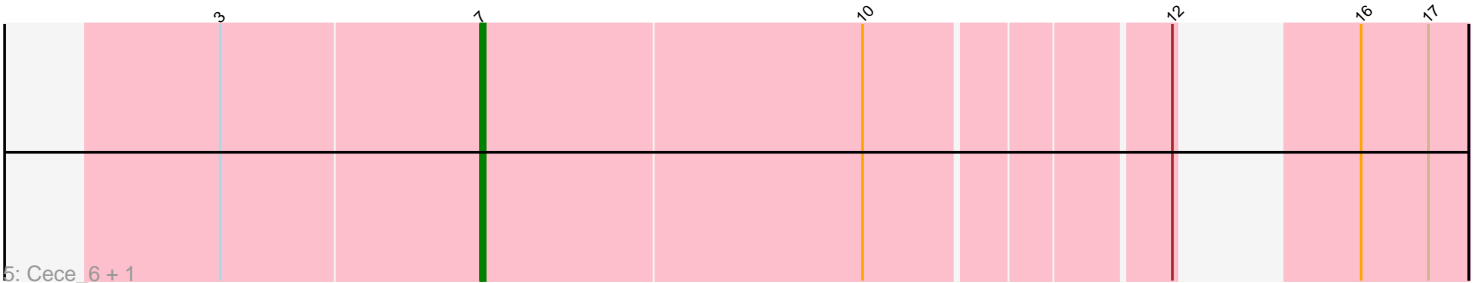
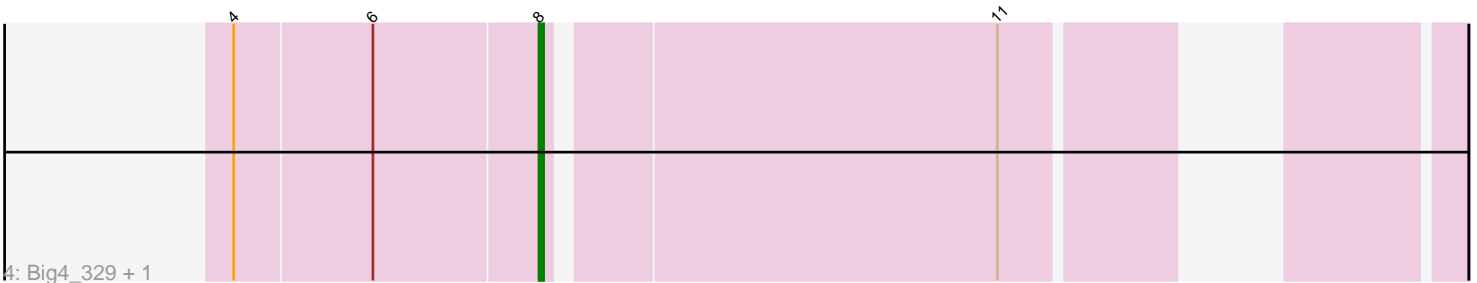
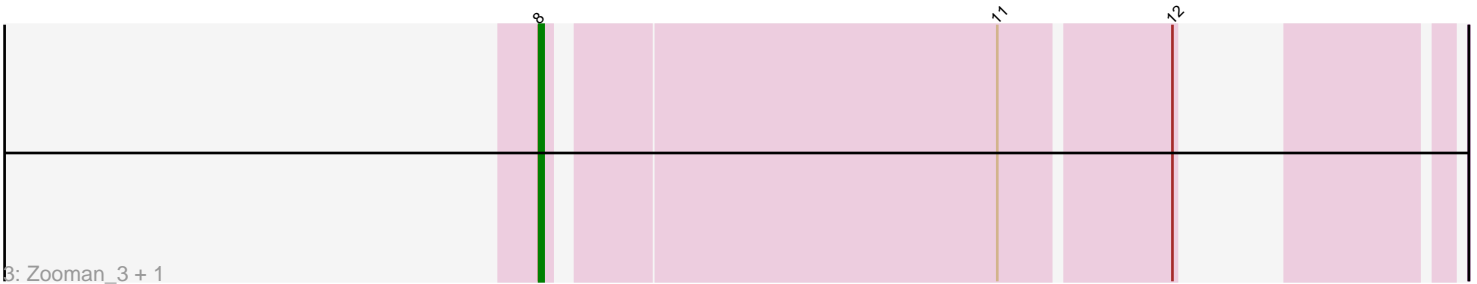


Pham 87513



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 87513 Report

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

Pham number 87513 has 10 members, 0 are drafts.

Phages represented in each track:

- Track 1 : PauloDiaboli\_358, PauloDiaboli\_3
- Track 2 : A3Wally\_357, A3Wally\_3
- Track 3 : Zooman\_3, Zooman\_316
- Track 4 : Big4\_329, Big4\_3
- Track 5 : Cece\_6, Cece\_308

### ***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\_3, A3Wally\_357, PauloDiaboli\_3, PauloDiaboli\_358,

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

- 

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

- Big4\_3, Big4\_329, Cece\_308, Cece\_6, Zooman\_3, Zooman\_316,

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

Start 7:

- 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\_308 (GD3), Cece\_6 (GD3),

Start 8:

- 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: Big4\_3 (GD2), Big4\_329 (GD2), Zooman\_3 (GD2), Zooman\_316 (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\_3 (GD1), A3Wally\_357 (GD1), PauloDiaboli\_3 (GD1), PauloDiaboli\_358 (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 8 was manually annotated 4 times for cluster GD2.

Info for manual annotations of cluster GD3:

- Start number 7 was manually annotated 2 times for cluster GD3.

### **Gene Information:**

Gene: A3Wally\_357 Start: 180321, Stop: 180515, Start Num: 9

Candidate Starts for A3Wally\_357:

(1, 180195), (2, 180216), (4, 180243), (5, 180264), (Start: 9 @180321 has 4 MA's), (11, 180411), (13, 180459), (14, 180462), (15, 180474),

Gene: A3Wally\_3 Start: 1100, Stop: 1294, Start Num: 9

Candidate Starts for A3Wally\_3:

(1, 974), (2, 995), (4, 1022), (5, 1043), (Start: 9 @1100 has 4 MA's), (11, 1190), (13, 1238), (14, 1241), (15, 1253),

Gene: Big4\_329 Start: 175815, Stop: 175985, Start Num: 8

Candidate Starts for Big4\_329:

(4, 175749), (6, 175779), (Start: 8 @175815 has 4 MA's), (11, 175911),

Gene: Big4\_3 Start: 1121, Stop: 1291, Start Num: 8

Candidate Starts for Big4\_3:

(4, 1055), (6, 1085), (Start: 8 @1121 has 4 MA's), (11, 1217),

Gene: Cece\_6 Start: 1742, Stop: 1930, Start Num: 7

Candidate Starts for Cece\_6:

(3, 1685), (Start: 7 @1742 has 2 MA's), (10, 1826), (12, 1889), (16, 1907), (17, 1922),

Gene: Cece\_308 Start: 170176, Stop: 170364, Start Num: 7

Candidate Starts for Cece\_308:

(3, 170119), (Start: 7 @170176 has 2 MA's), (10, 170260), (12, 170323), (16, 170341), (17, 170356),

Gene: PauloDiaboli\_358 Start: 177720, Stop: 177914, Start Num: 9

Candidate Starts for PauloDiaboli\_358:

(2, 177615), (4, 177642), (5, 177663), (Start: 9 @177720 has 4 MA's), (11, 177810), (13, 177858), (14, 177861), (15, 177873),

Gene: PauloDiaboli\_3 Start: 1091, Stop: 1285, Start Num: 9

Candidate Starts for PauloDiaboli\_3:

(2, 986), (4, 1013), (5, 1034), (Start: 9 @1091 has 4 MA's), (11, 1181), (13, 1229), (14, 1232), (15, 1244),

Gene: Zooman\_3 Start: 1121, Stop: 1288, Start Num: 8

Candidate Starts for Zooman\_3:

(Start: 8 @1121 has 4 MA's), (11, 1217), (12, 1253),

Gene: Zooman\_316 Start: 176772, Stop: 176939, Start Num: 8

Candidate Starts for Zooman\_316:

(Start: 8 @176772 has 4 MA's), (11, 176868), (12, 176904),