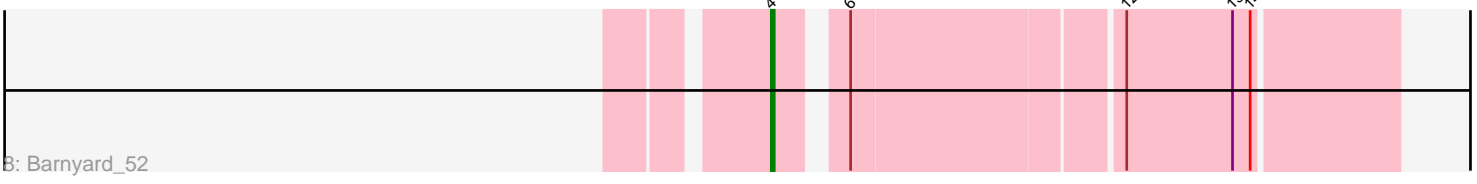
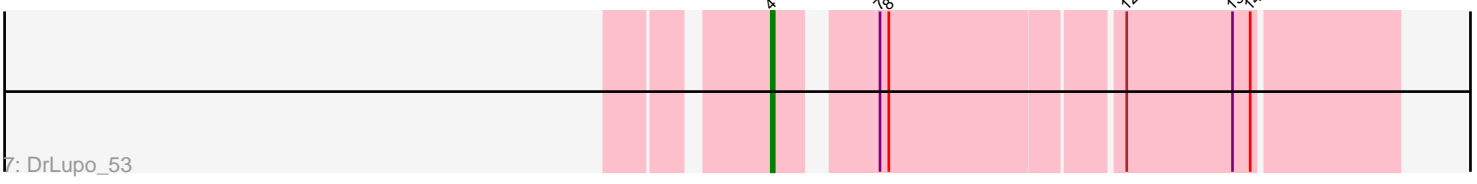
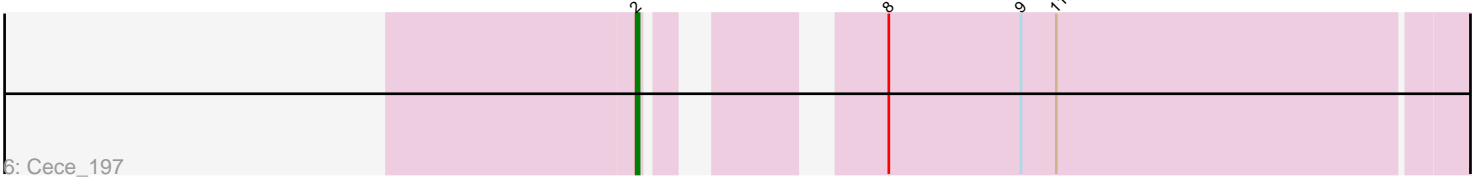
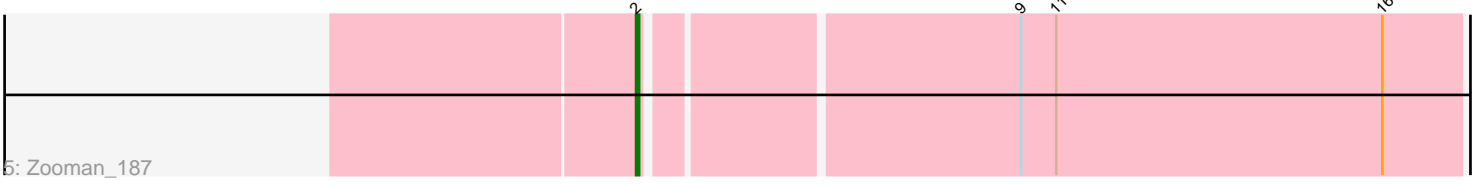
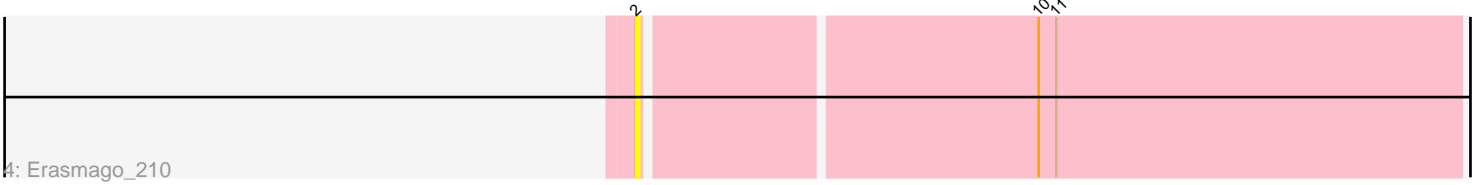
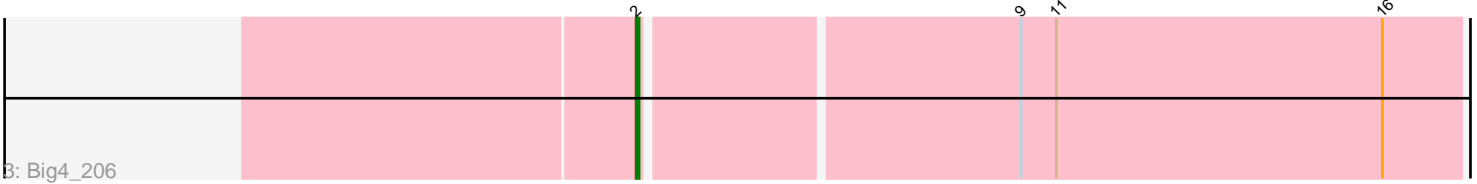
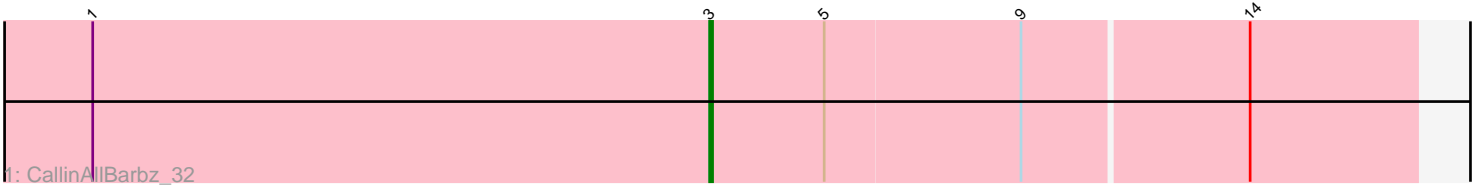


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Note: Tracks are now grouped by subcluster and scaled. Switching in subcluster is indicated by changes in track color. Track scale is now set by default to display the region 30 bp upstream of start 1 to 30 bp downstream of the last possible start. 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 278840 Report

This analysis was run 02/07/26 on database version 634.

Pham number 278840 has 10 members, 2 are drafts.

Phages represented in each track:

- Track 1 : CallinAllBarbz\_32
- Track 2 : A3Wally\_221, PauloDiaboli\_221, Dodo\_220
- Track 3 : Big4\_206
- Track 4 : Erasmago\_210
- Track 5 : Zooman\_187
- Track 6 : Cece\_197
- Track 7 : DrLupo\_53
- Track 8 : Barnyard\_52

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

The start number called the most often in the published annotations is 2, it was called in 5 of the 8 non-draft genes in the pham.

Genes that call this "Most Annotated" start:

- A3Wally\_221, Big4\_206, Cece\_197, Dodo\_220, Erasmago\_210, PauloDiaboli\_221, Zooman\_187,

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

- 

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

- Barnyard\_52, CallinAllBarbz\_32, DrLupo\_53,

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

Start 2:

- Found in 7 of 10 ( 70.0% ) of genes in pham
- Manual Annotations of this start: 5 of 8
- Called 100.0% of time when present
- Phage (with cluster) where this start called: A3Wally\_221 (GD1), Big4\_206 (GD2), Cece\_197 (GD3), Dodo\_220 (GD1), Erasmago\_210 (GD2), PauloDiaboli\_221 (GD1), Zooman\_187 (GD2),

#### Start 3:

- Found in 1 of 10 ( 10.0% ) of genes in pham
- Manual Annotations of this start: 1 of 8
- Called 100.0% of time when present
- Phage (with cluster) where this start called: CallinAllBarbz\_32 (FP),

#### Start 4:

- Found in 2 of 10 ( 20.0% ) 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: Barnyard\_52 (H2), DrLupo\_53 (H2),

### **Summary by clusters:**

There are 5 clusters represented in this pham: H2, GD1, GD2, GD3, FP,

Info for manual annotations of cluster FP:

- Start number 3 was manually annotated 1 time for cluster FP.

Info for manual annotations of cluster GD1:

- Start number 2 was manually annotated 2 times for cluster GD1.

Info for manual annotations of cluster GD2:

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

Info for manual annotations of cluster GD3:

- Start number 2 was manually annotated 1 time for cluster GD3.

Info for manual annotations of cluster H2:

- Start number 4 was manually annotated 2 times for cluster H2.

### **Gene Information:**

Gene: A3Wally\_221 Start: 119347, Stop: 119087, Start Num: 2

Candidate Starts for A3Wally\_221:

(Start: 2 @119347 has 5 MA's), (9, 119239), (11, 119227), (15, 119140),

Gene: Barnyard\_52 Start: 39091, Stop: 39285, Start Num: 4

Candidate Starts for Barnyard\_52:

(Start: 4 @39091 has 2 MA's), (6, 39109), (12, 39196), (13, 39232), (14, 39238),

Gene: Big4\_206 Start: 115772, Stop: 115500, Start Num: 2

Candidate Starts for Big4\_206:

(Start: 2 @115772 has 5 MA's), (9, 115649), (11, 115637), (16, 115526),

Gene: CallinAllBarbz\_32 Start: 24070, Stop: 24306, Start Num: 3

Candidate Starts for CallinAllBarbz\_32:

(1, 23860), (Start: 3 @24070 has 1 MA's), (5, 24109), (9, 24175), (14, 24250),

Gene: Cece\_197 Start: 120223, Stop: 119972, Start Num: 2

Candidate Starts for Cece\_197:

(Start: 2 @120223 has 5 MA's), (8, 120166), (9, 120121), (11, 120109),

Gene: Dodo\_220 Start: 119149, Stop: 118889, Start Num: 2

Candidate Starts for Dodo\_220:

(Start: 2 @119149 has 5 MA's), (9, 119041), (11, 119029), (15, 118942),

Gene: DrLupo\_53 Start: 39502, Stop: 39696, Start Num: 4

Candidate Starts for DrLupo\_53:

(Start: 4 @39502 has 2 MA's), (7, 39529), (8, 39532), (12, 39607), (13, 39643), (14, 39649),

Gene: Erasmago\_210 Start: 114028, Stop: 113756, Start Num: 2

Candidate Starts for Erasmago\_210:

(Start: 2 @114028 has 5 MA's), (10, 113899), (11, 113893),

Gene: PauloDiaboli\_221 Start: 117560, Stop: 117300, Start Num: 2

Candidate Starts for PauloDiaboli\_221:

(Start: 2 @117560 has 5 MA's), (9, 117452), (11, 117440), (15, 117353),

Gene: Zooman\_187 Start: 111993, Stop: 111724, Start Num: 2

Candidate Starts for Zooman\_187:

(Start: 2 @111993 has 5 MA's), (9, 111873), (11, 111861), (16, 111750),