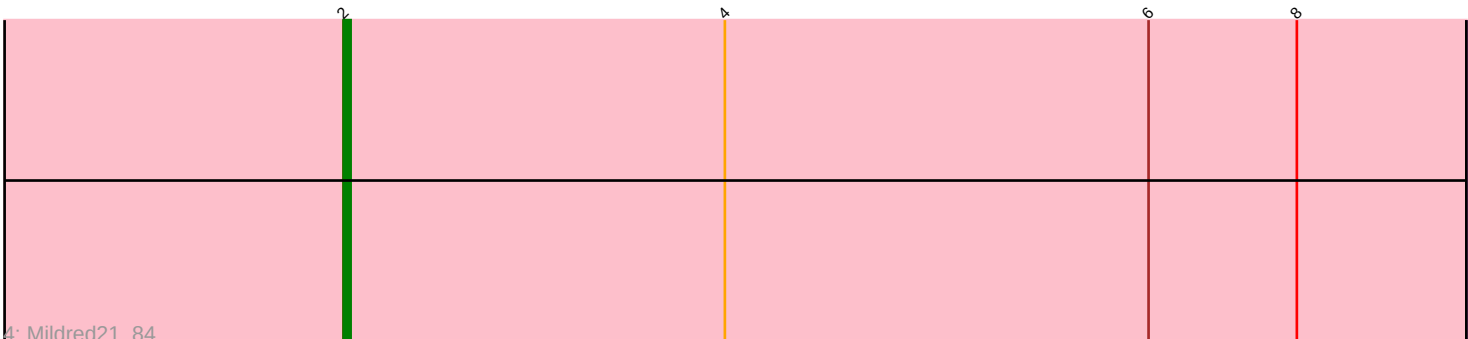
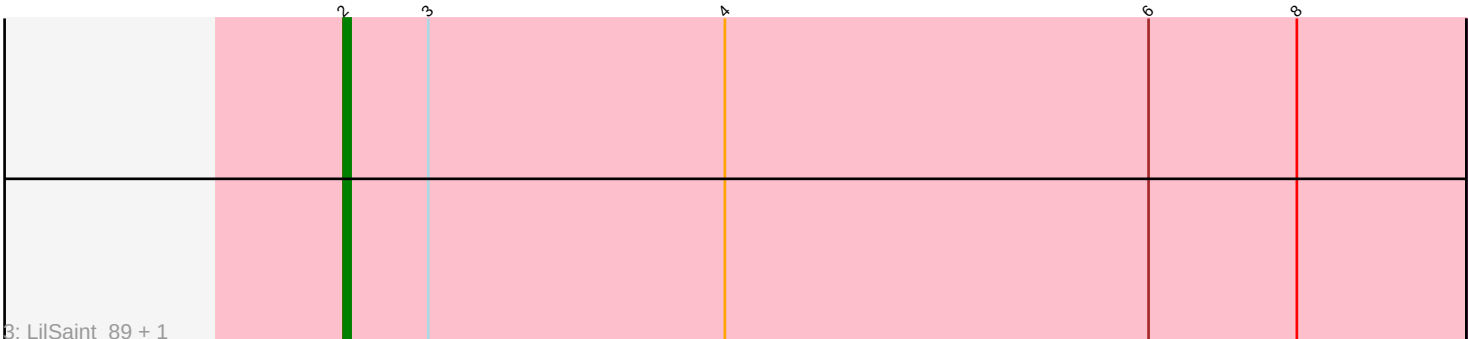
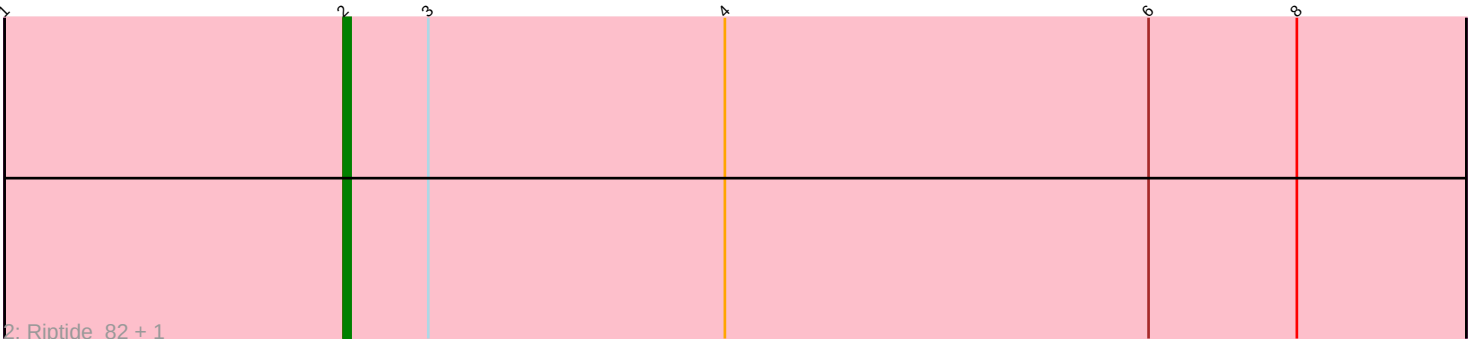
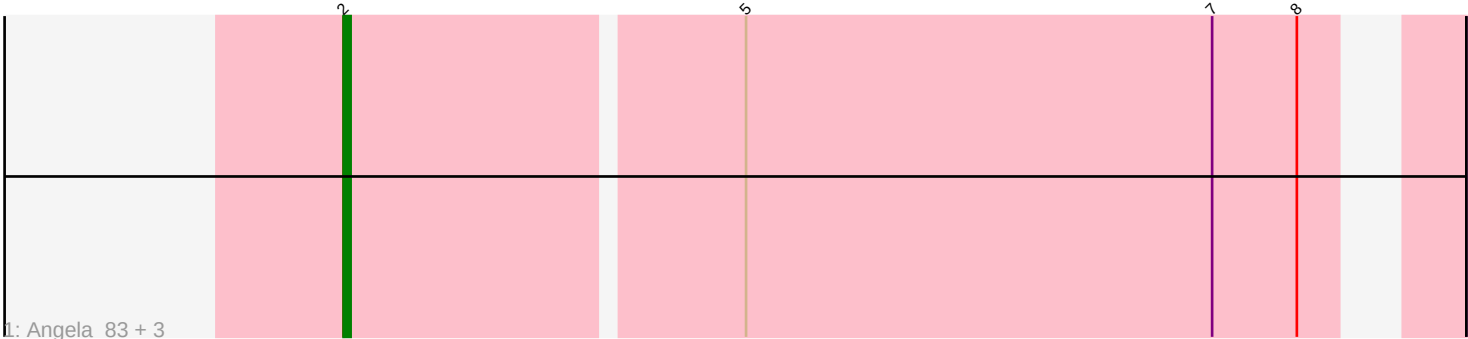




Note: Tracks are now grouped by subcluster and scaled. Switching in subcluster is indicated by changes in track color. Track scale is now set by default to display the region 30 bp upstream of start 1 to 30 bp downstream of the last possible start. If this default region is judged to be packed too tightly 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 281159 Report

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

Pham number 281159 has 9 members, 1 are drafts.

Phages represented in each track:

- Track 1 : Angela_83, LilMartin_83, Marsus_84, MulchMansion_83
- Track 2 : Riptide_82, Anedea_85
- Track 3 : LilSaint_89, Bmoc_84
- Track 4 : Mildred21_84

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 8 of the 8 non-draft genes in the pham.

Genes that call this "Most Annotated" start:

- Anedea_85, Angela_83, Bmoc_84, LilMartin_83, LilSaint_89, Marsus_84, Mildred21_84, MulchMansion_83, Riptide_82,

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

-

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

-

Summary by start number:

Start 2:

- Found in 9 of 9 (100.0%) of genes in pham
- Manual Annotations of this start: 8 of 8
- Called 100.0% of time when present
- Phage (with cluster) where this start called: Anedea_85 (BE1), Angela_83 (BE1), Bmoc_84 (BE1), LilMartin_83 (BE1), LilSaint_89 (BE1), Marsus_84 (BE1), Mildred21_84 (BE1), MulchMansion_83 (BE1), Riptide_82 (BE1),

Summary by clusters:

There is one cluster represented in this pham: BE1

Info for manual annotations of cluster BE1:

•Start number 2 was manually annotated 8 times for cluster BE1.

Gene Information:

Gene: Anedea_85 Start: 63484, Stop: 63642, Start Num: 2

Candidate Starts for Anedea_85:

(1, 63436), (Start: 2 @63484 has 8 MA's), (3, 63496), (4, 63538), (6, 63598), (8, 63619),

Gene: Angela_83 Start: 62975, Stop: 63121, Start Num: 2

Candidate Starts for Angela_83:

(Start: 2 @62975 has 8 MA's), (5, 63029), (7, 63095), (8, 63107),

Gene: Bmoc_84 Start: 63322, Stop: 63480, Start Num: 2

Candidate Starts for Bmoc_84:

(Start: 2 @63322 has 8 MA's), (3, 63334), (4, 63376), (6, 63436), (8, 63457),

Gene: LilMartin_83 Start: 62923, Stop: 63069, Start Num: 2

Candidate Starts for LilMartin_83:

(Start: 2 @62923 has 8 MA's), (5, 62977), (7, 63043), (8, 63055),

Gene: LilSaint_89 Start: 63157, Stop: 63315, Start Num: 2

Candidate Starts for LilSaint_89:

(Start: 2 @63157 has 8 MA's), (3, 63169), (4, 63211), (6, 63271), (8, 63292),

Gene: Marsus_84 Start: 63172, Stop: 63318, Start Num: 2

Candidate Starts for Marsus_84:

(Start: 2 @63172 has 8 MA's), (5, 63226), (7, 63292), (8, 63304),

Gene: Mildred21_84 Start: 63128, Stop: 63286, Start Num: 2

Candidate Starts for Mildred21_84:

(Start: 2 @63128 has 8 MA's), (4, 63182), (6, 63242), (8, 63263),

Gene: MulchMansion_83 Start: 62923, Stop: 63069, Start Num: 2

Candidate Starts for MulchMansion_83:

(Start: 2 @62923 has 8 MA's), (5, 62977), (7, 63043), (8, 63055),

Gene: Riptide_82 Start: 62701, Stop: 62859, Start Num: 2

Candidate Starts for Riptide_82:

(1, 62653), (Start: 2 @62701 has 8 MA's), (3, 62713), (4, 62755), (6, 62815), (8, 62836),