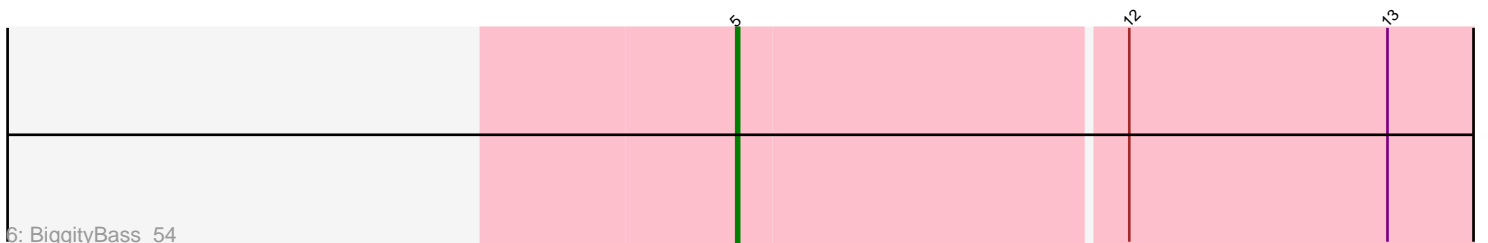
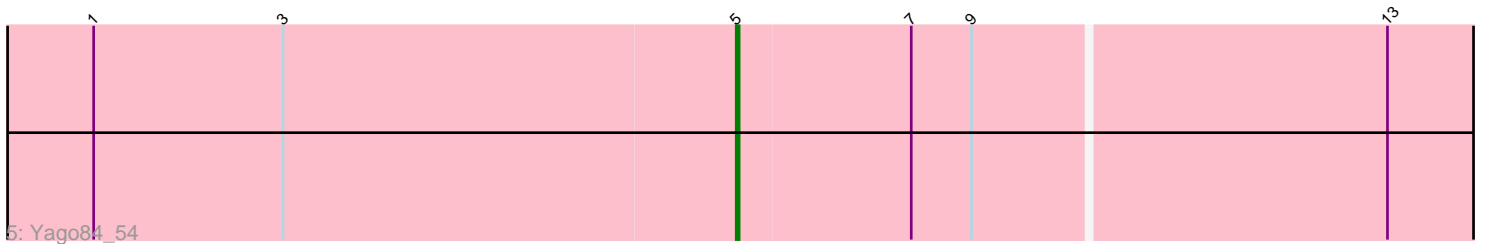
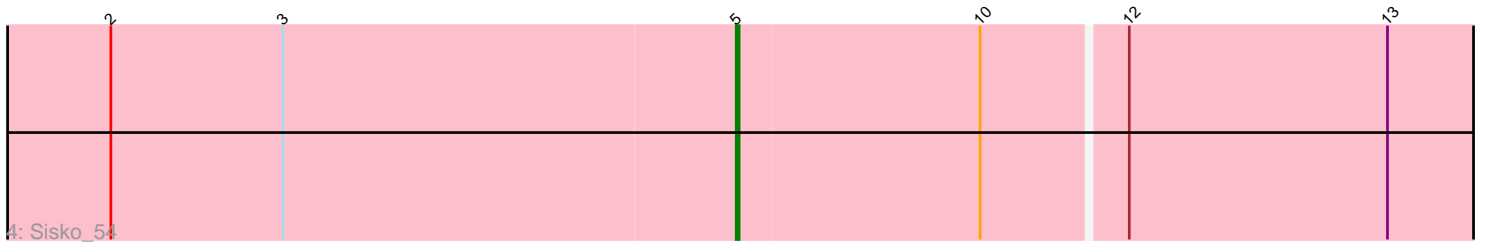
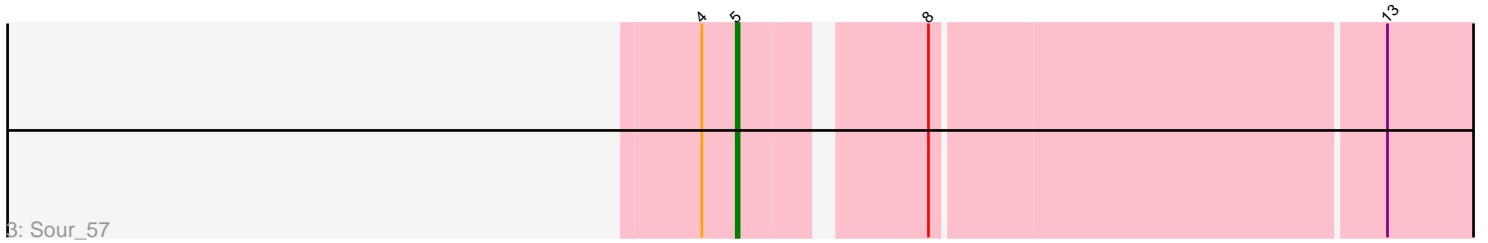
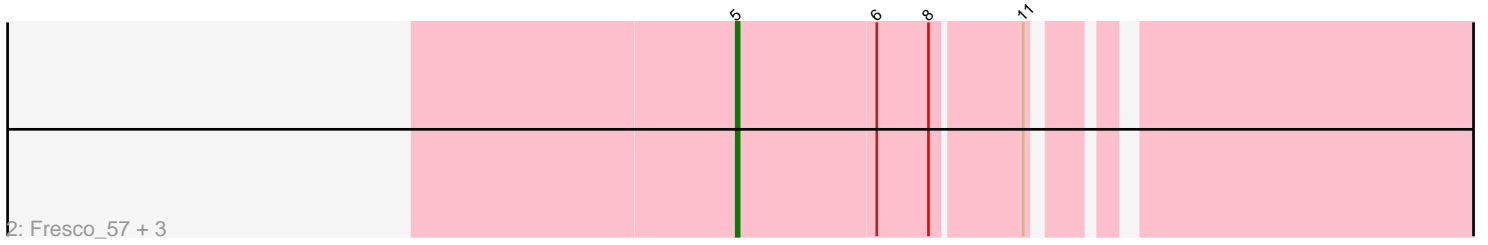
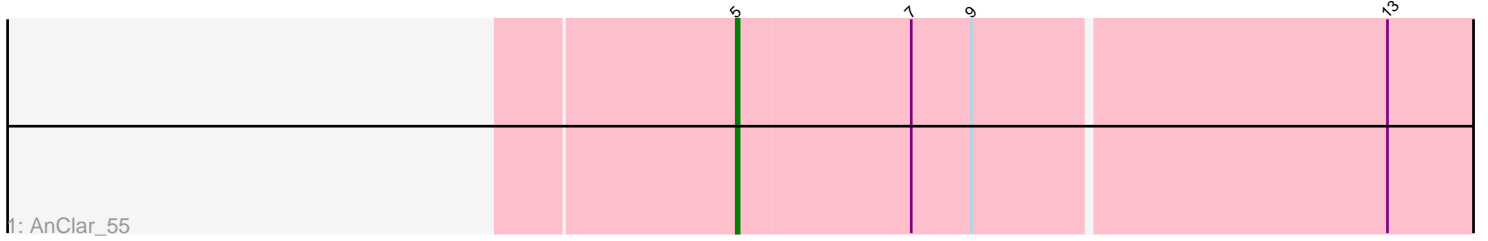


Pham 295364



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 295364 Report

This analysis was run 04/18/26 on database version 643.

Pham number 295364 has 9 members, 0 are drafts.

Phages represented in each track:

- Track 1 : AnClar_55
- Track 2 : Fresco_57, Axumite_57, Ligma_57, Shatter_57
- Track 3 : Sour_57
- Track 4 : Sisko_54
- Track 5 : Yago84_54
- Track 6 : BiggityBass_54

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

The start number called the most often in the published annotations is 5, it was called in 9 of the 9 non-draft genes in the pham.

Genes that call this "Most Annotated" start:

- AnClar_55, Axumite_57, BiggityBass_54, Fresco_57, Ligma_57, Shatter_57, Sisko_54, Sour_57, Yago84_54,

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 5:

- Found in 9 of 9 (100.0%) of genes in pham
- Manual Annotations of this start: 9 of 9
- Called 100.0% of time when present
- Phage (with cluster) where this start called: AnClar_55 (DR), Axumite_57 (DR), BiggityBass_54 (DR), Fresco_57 (DR), Ligma_57 (DR), Shatter_57 (DR), Sisko_54 (DR), Sour_57 (DR), Yago84_54 (DR),

Summary by clusters:

There is one cluster represented in this pham: DR

Info for manual annotations of cluster DR:

•Start number 5 was manually annotated 9 times for cluster DR.

Gene Information:

Gene: AnClar_55 Start: 49921, Stop: 49628, Start Num: 5

Candidate Starts for AnClar_55:

(Start: 5 @49921 has 9 MA's), (7, 49861), (9, 49840), (13, 49699),

Gene: Axumite_57 Start: 48183, Stop: 47899, Start Num: 5

Candidate Starts for Axumite_57:

(Start: 5 @48183 has 9 MA's), (6, 48135), (8, 48117), (11, 48087),

Gene: BiggityBass_54 Start: 49280, Stop: 48987, Start Num: 5

Candidate Starts for BiggityBass_54:

(Start: 5 @49280 has 9 MA's), (12, 49148), (13, 49058),

Gene: Fresco_57 Start: 48183, Stop: 47899, Start Num: 5

Candidate Starts for Fresco_57:

(Start: 5 @48183 has 9 MA's), (6, 48135), (8, 48117), (11, 48087),

Gene: Ligma_57 Start: 48183, Stop: 47899, Start Num: 5

Candidate Starts for Ligma_57:

(Start: 5 @48183 has 9 MA's), (6, 48135), (8, 48117), (11, 48087),

Gene: Shatter_57 Start: 48183, Stop: 47899, Start Num: 5

Candidate Starts for Shatter_57:

(Start: 5 @48183 has 9 MA's), (6, 48135), (8, 48117), (11, 48087),

Gene: Sisko_54 Start: 47905, Stop: 47612, Start Num: 5

Candidate Starts for Sisko_54:

(2, 48121), (3, 48061), (Start: 5 @47905 has 9 MA's), (10, 47821), (12, 47773), (13, 47683),

Gene: Sour_57 Start: 50612, Stop: 50331, Start Num: 5

Candidate Starts for Sour_57:

(4, 50624), (Start: 5 @50612 has 9 MA's), (8, 50555), (13, 50402),

Gene: Yago84_54 Start: 47998, Stop: 47705, Start Num: 5

Candidate Starts for Yago84_54:

(1, 48220), (3, 48154), (Start: 5 @47998 has 9 MA's), (7, 47938), (9, 47917), (13, 47776),