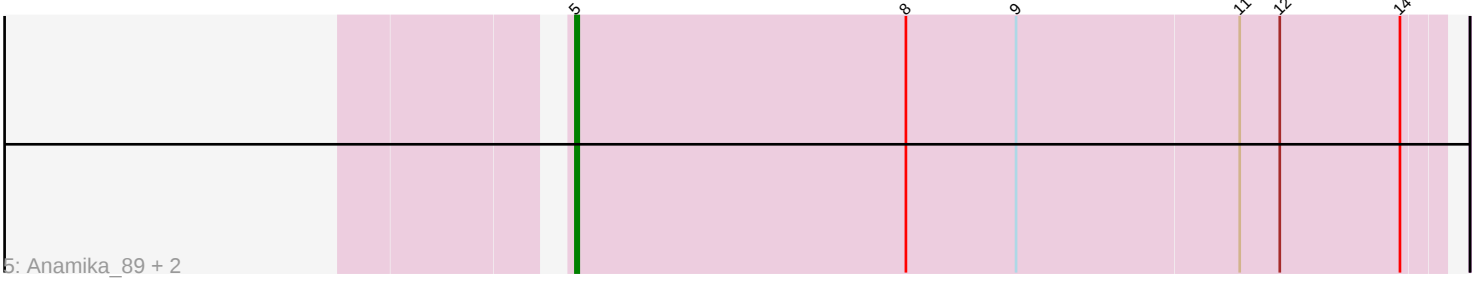
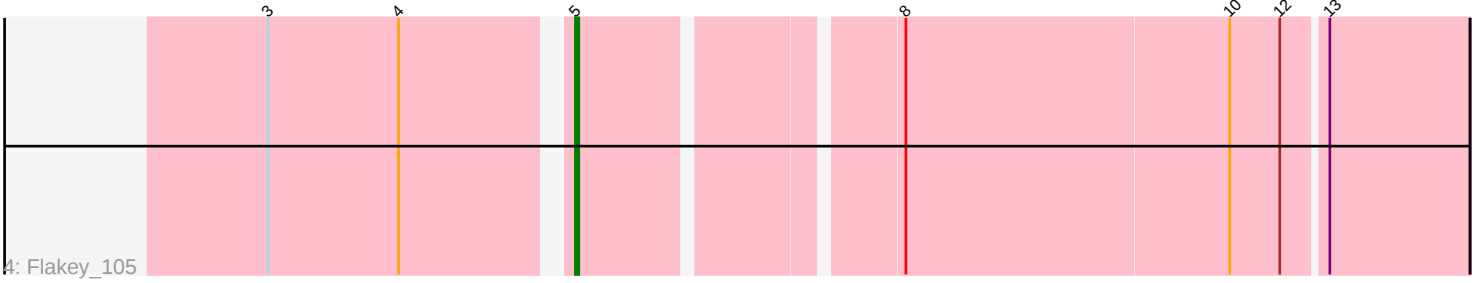
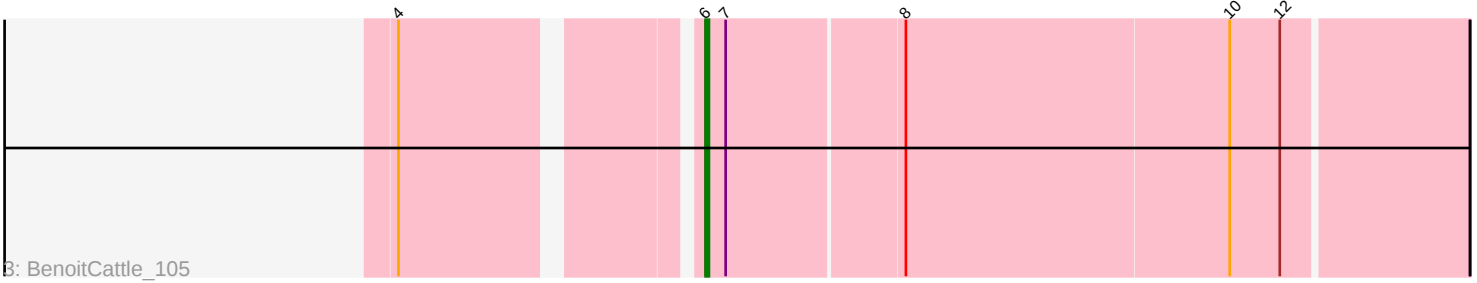
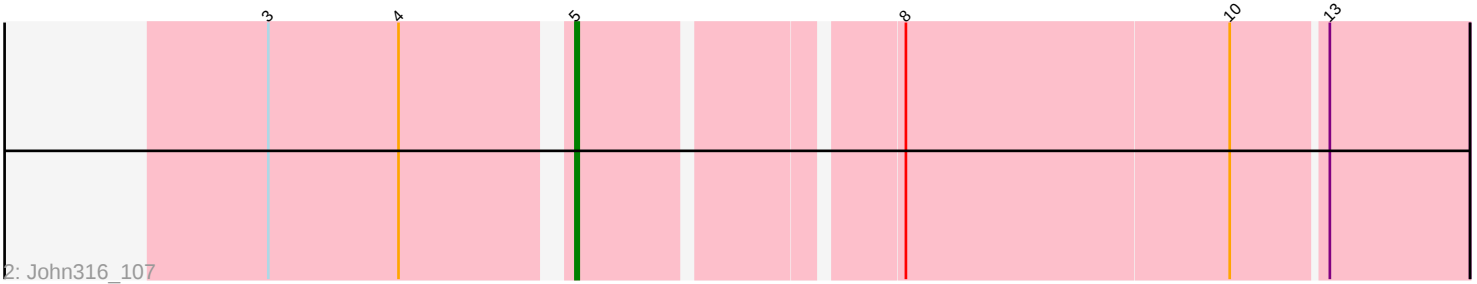
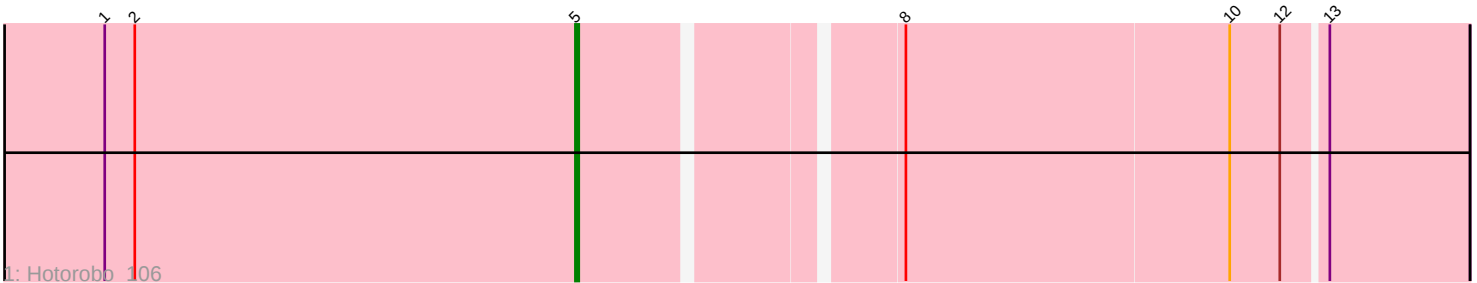


Pham 331616



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 331616 Report

This analysis was run 09/05/26 on database version 662.

Pham number 331616 has 7 members, 0 are drafts.

Phages represented in each track:

- Track 1 : Hotorobo_106
- Track 2 : John316_107
- Track 3 : BenoitCattle_105
- Track 4 : Flakey_105
- Track 5 : Anamika_89, Charianelly_90, Bianmat_90

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 6 of the 7 non-draft genes in the pham.

Genes that call this "Most Annotated" start:

- Anamika_89, Bianmat_90, Charianelly_90, Flakey_105, Hotorobo_106, John316_107,

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

-

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

- BenoitCattle_105,

Summary by start number:

Start 5:

- Found in 6 of 7 (85.7%) of genes in pham
- Manual Annotations of this start: 6 of 7
- Called 100.0% of time when present
- Phage (with cluster) where this start called: Anamika_89 (CS3), Bianmat_90 (CS3), Charianelly_90 (CS3), Flakey_105 (CS2), Hotorobo_106 (CS2), John316_107 (CS2),

Start 6:

- Found in 1 of 7 (14.3%) of genes in pham
- Manual Annotations of this start: 1 of 7
- Called 100.0% of time when present

- Phage (with cluster) where this start called: BenoitCattle_105 (CS2),

Summary by clusters:

There are 2 clusters represented in this pham: CS3, CS2,

Info for manual annotations of cluster CS2:

- Start number 5 was manually annotated 3 times for cluster CS2.
- Start number 6 was manually annotated 1 time for cluster CS2.

Info for manual annotations of cluster CS3:

- Start number 5 was manually annotated 3 times for cluster CS3.

Gene Information:

Gene: Anamika_89 Start: 70826, Stop: 70569, Start Num: 5

Candidate Starts for Anamika_89:

(Start: 5 @70826 has 6 MA's), (8, 70727), (9, 70694), (11, 70628), (12, 70616), (14, 70580),

Gene: BenoitCattle_105 Start: 73827, Stop: 73606, Start Num: 6

Candidate Starts for BenoitCattle_105:

(4, 73905), (Start: 6 @73827 has 1 MA's), (7, 73821), (8, 73770), (10, 73674), (12, 73659),

Gene: Bianmat_90 Start: 70838, Stop: 70581, Start Num: 5

Candidate Starts for Bianmat_90:

(Start: 5 @70838 has 6 MA's), (8, 70739), (9, 70706), (11, 70640), (12, 70628), (14, 70592),

Gene: Charianelly_90 Start: 70551, Stop: 70294, Start Num: 5

Candidate Starts for Charianelly_90:

(Start: 5 @70551 has 6 MA's), (8, 70452), (9, 70419), (11, 70353), (12, 70341), (14, 70305),

Gene: Flakey_105 Start: 74135, Stop: 73884, Start Num: 5

Candidate Starts for Flakey_105:

(3, 74219), (4, 74180), (Start: 5 @74135 has 6 MA's), (8, 74048), (10, 73952), (12, 73937), (13, 73925),

Gene: Hotorobo_106 Start: 74095, Stop: 73844, Start Num: 5

Candidate Starts for Hotorobo_106:

(1, 74236), (2, 74227), (Start: 5 @74095 has 6 MA's), (8, 74008), (10, 73912), (12, 73897), (13, 73885),

Gene: John316_107 Start: 74362, Stop: 74111, Start Num: 5

Candidate Starts for John316_107:

(3, 74446), (4, 74407), (Start: 5 @74362 has 6 MA's), (8, 74275), (10, 74179), (13, 74152),