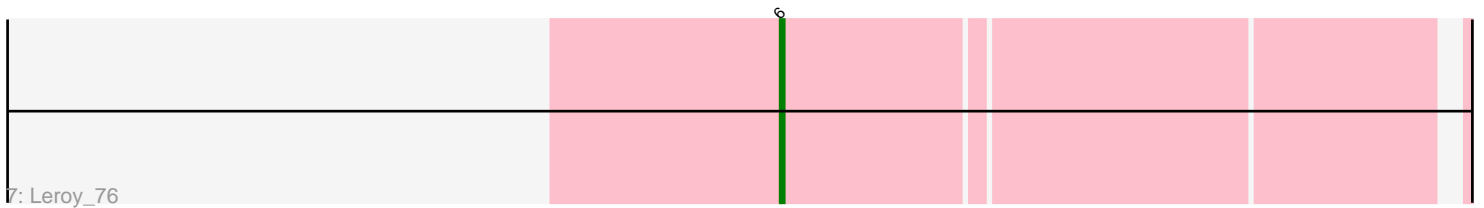
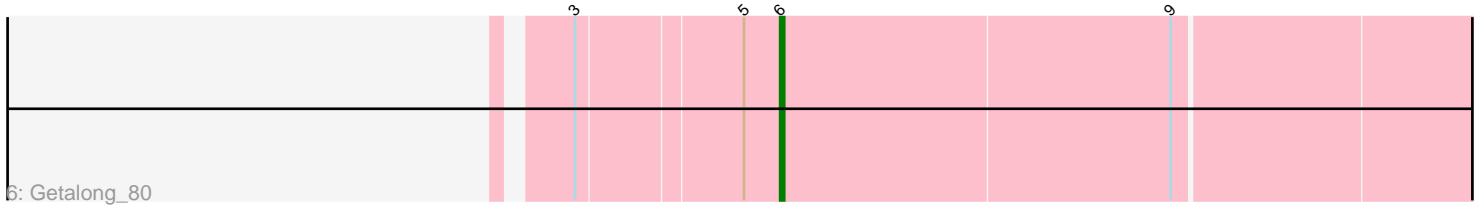
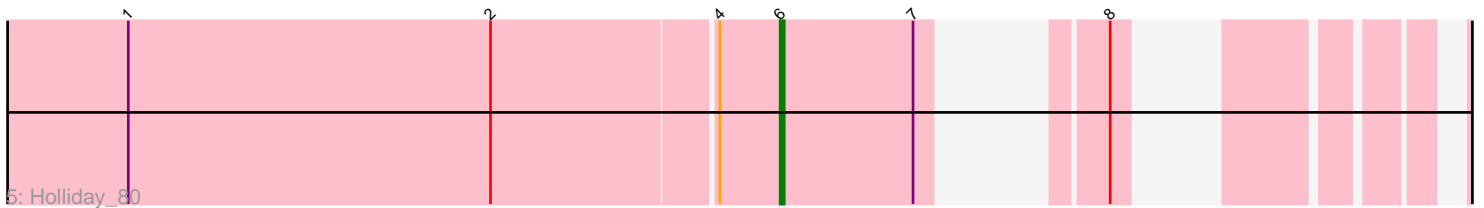
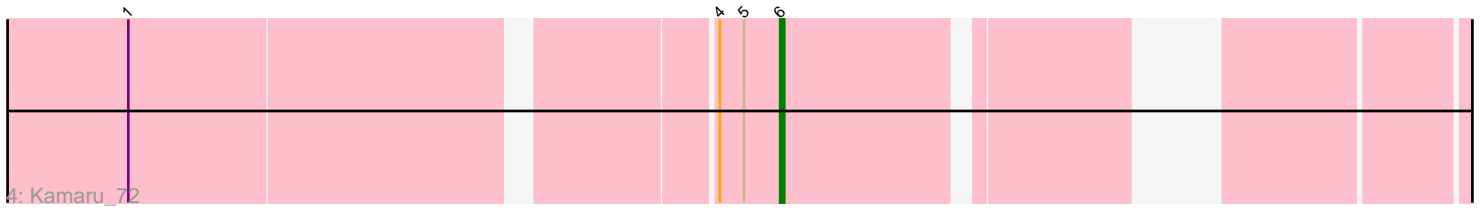
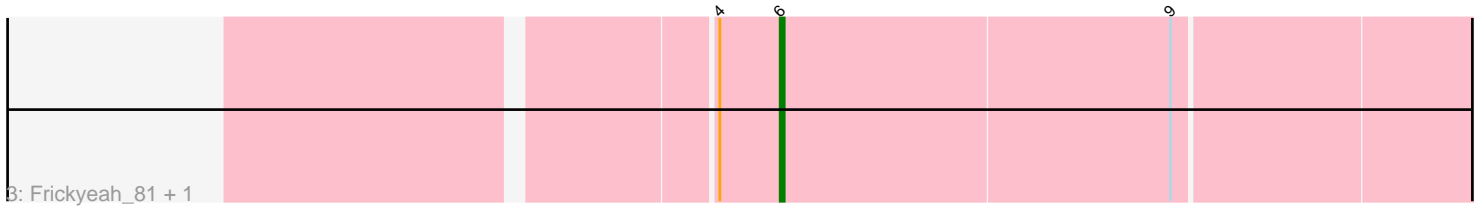
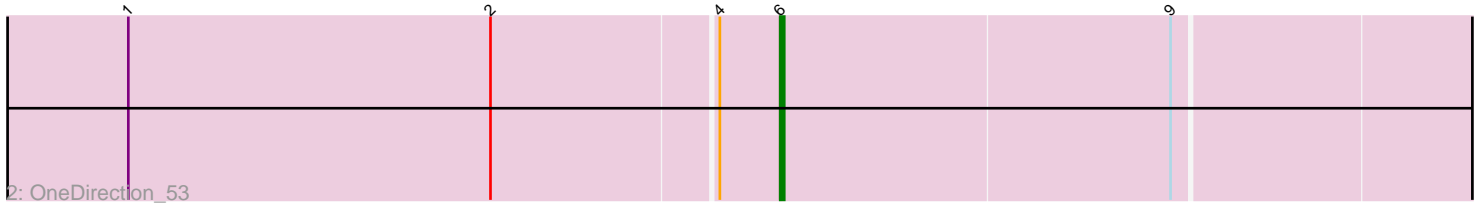
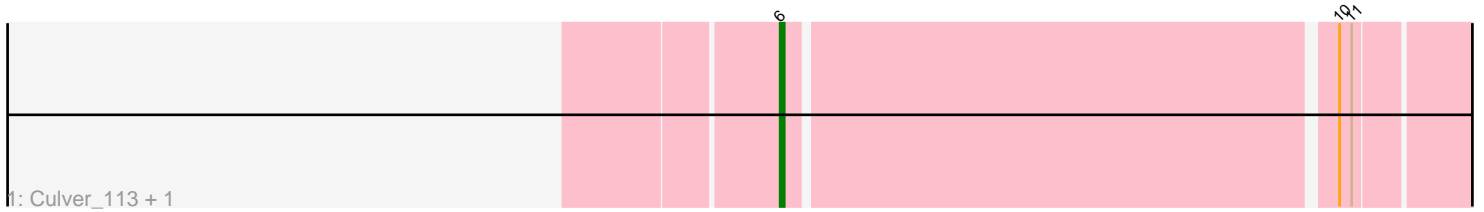


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Note: Tracks are now grouped by subcluster and scaled. Switching in subcluster is indicated by changes in track color. Track scale is now set by default to display the region 30 bp upstream of start 1 to 30 bp downstream of the last possible start. 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 225128 Report

This analysis was run 03/28/25 on database version 593.

Pham number 225128 has 9 members, 1 are drafts.

Phages represented in each track:

- Track 1 : Culver\_113, Abscondus\_112
- Track 2 : OneDirection\_53
- Track 3 : Frickyeah\_81, BENtherdunthat\_79
- Track 4 : Kamaru\_72
- Track 5 : Holliday\_80
- Track 6 : Getalong\_80
- Track 7 : Leroy\_76

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

The start number called the most often in the published annotations is 6, it was called in 8 of the 8 non-draft genes in the pham.

Genes that call this "Most Annotated" start:

- Abscondus\_112, BENtherdunthat\_79, Culver\_113, Frickyeah\_81, Getalong\_80, Holliday\_80, Kamaru\_72, Leroy\_76, OneDirection\_53,

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 6:

- 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: Abscondus\_112 (CQ1), BENtherdunthat\_79 (DN1), Culver\_113 (CQ1), Frickyeah\_81 (DN1), Getalong\_80 (DN1), Holliday\_80 (DN1), Kamaru\_72 (DN1), Leroy\_76 (DN1), OneDirection\_53 (CZ6),

## Summary by clusters:

There are 3 clusters represented in this pham: CZ6, CQ1, DN1,

Info for manual annotations of cluster CQ1:

- Start number 6 was manually annotated 2 times for cluster CQ1.

Info for manual annotations of cluster CZ6:

- Start number 6 was manually annotated 1 time for cluster CZ6.

Info for manual annotations of cluster DN1:

- Start number 6 was manually annotated 5 times for cluster DN1.

## Gene Information:

Gene: Abscondus\_112 Start: 65388, Stop: 65648, Start Num: 6

Candidate Starts for Abscondus\_112:

(Start: 6 @65388 has 8 MA's), (10, 65520), (11, 65523),

Gene: BENtherdunthat\_79 Start: 44974, Stop: 45240, Start Num: 6

Candidate Starts for BENtherdunthat\_79:

(4, 44959), (Start: 6 @44974 has 8 MA's), (9, 45070),

Gene: Culver\_113 Start: 64860, Stop: 65120, Start Num: 6

Candidate Starts for Culver\_113:

(Start: 6 @64860 has 8 MA's), (10, 64992), (11, 64995),

Gene: Frickyeah\_81 Start: 45105, Stop: 45371, Start Num: 6

Candidate Starts for Frickyeah\_81:

(4, 45090), (Start: 6 @45105 has 8 MA's), (9, 45201),

Gene: Getalong\_80 Start: 46400, Stop: 46666, Start Num: 6

Candidate Starts for Getalong\_80:

(3, 46352), (5, 46391), (Start: 6 @46400 has 8 MA's), (9, 46496),

Gene: Holliday\_80 Start: 48011, Stop: 48211, Start Num: 6

Candidate Starts for Holliday\_80:

(1, 47852), (2, 47942), (4, 47996), (Start: 6 @48011 has 8 MA's), (7, 48044), (8, 48062),

Gene: Kamaru\_72 Start: 43594, Stop: 43830, Start Num: 6

Candidate Starts for Kamaru\_72:

(1, 43444), (4, 43579), (5, 43585), (Start: 6 @43594 has 8 MA's),

Gene: Leroy\_76 Start: 44184, Stop: 44435, Start Num: 6

Candidate Starts for Leroy\_76:

(Start: 6 @44184 has 8 MA's),

Gene: OneDirection\_53 Start: 33352, Stop: 33618, Start Num: 6

Candidate Starts for OneDirection\_53:

(1, 33193), (2, 33283), (4, 33337), (Start: 6 @33352 has 8 MA's), (9, 33448),