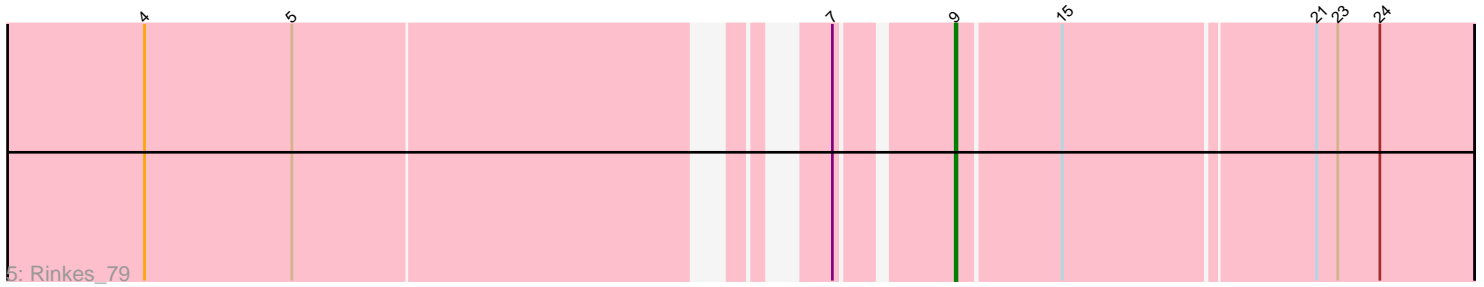
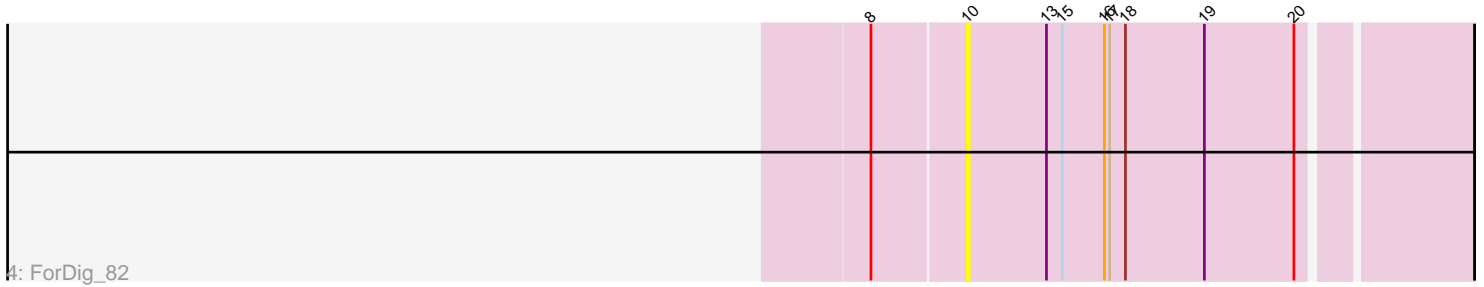
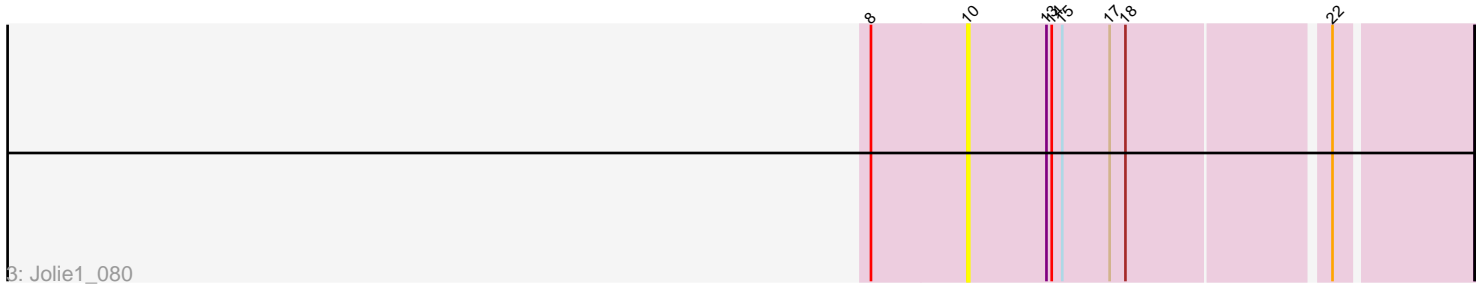
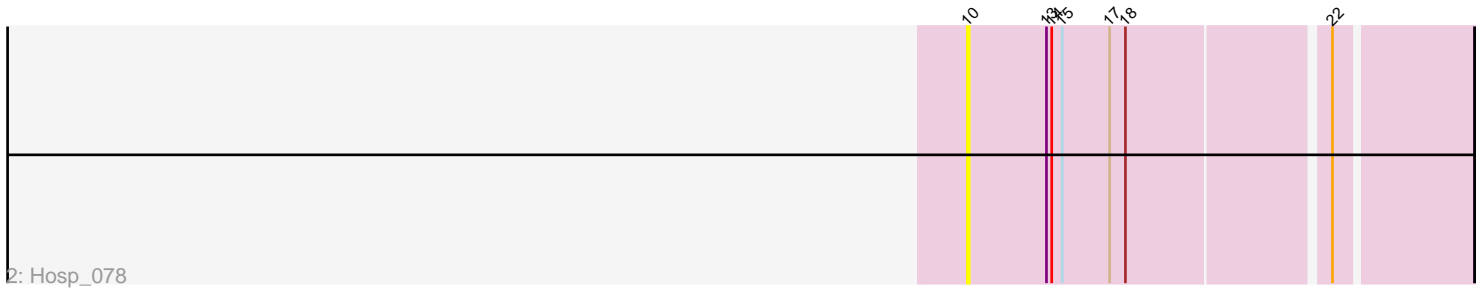
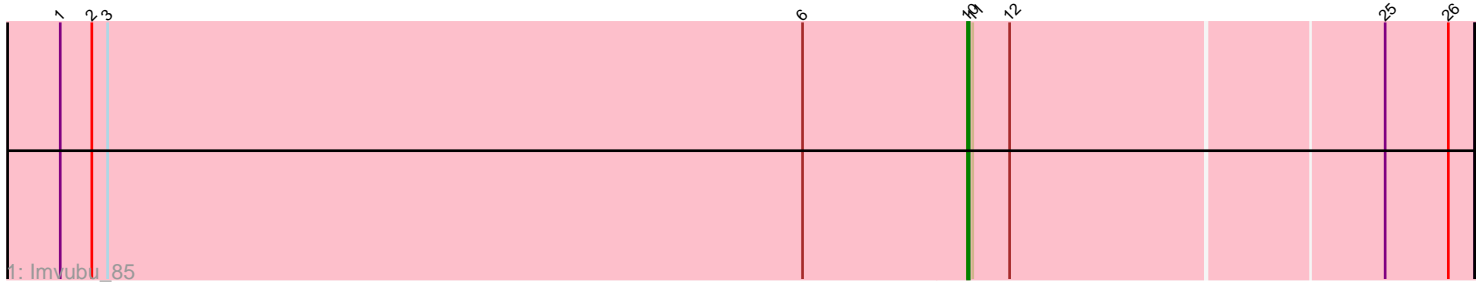


Pham 220565



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 220565 Report

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

Pham number 220565 has 5 members, 3 are drafts.

Phages represented in each track:

- Track 1 : Imvubu\_85
- Track 2 : Hosp\_078
- Track 3 : Jolie1\_080
- Track 4 : ForDig\_82
- Track 5 : Rinke\_79

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

The start number called the most often in the published annotations is 10, it was called in 1 of the 2 non-draft genes in the pham.

Genes that call this "Most Annotated" start:

- ForDig\_82, Hosp\_078, Imvubu\_85, Jolie1\_080,

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

- 

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

- Rinke\_79,

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

Start 9:

- Found in 1 of 5 ( 20.0% ) of genes in pham
- Manual Annotations of this start: 1 of 2
- Called 100.0% of time when present
- Phage (with cluster) where this start called: Rinke\_79 (B9),

Start 10:

- Found in 4 of 5 ( 80.0% ) of genes in pham
- Manual Annotations of this start: 1 of 2
- Called 100.0% of time when present
- Phage (with cluster) where this start called: ForDig\_82 (B6), Hosp\_078 (B6), Imvubu\_85 (B10), Jolie1\_080 (B6),

## Summary by clusters:

There are 3 clusters represented in this pham: B6, B9, B10,

Info for manual annotations of cluster B10:

- Start number 10 was manually annotated 1 time for cluster B10.

Info for manual annotations of cluster B9:

- Start number 9 was manually annotated 1 time for cluster B9.

## **Gene Information:**

Gene: ForDig\_82 Start: 63250, Stop: 62975, Start Num: 10

Candidate Starts for ForDig\_82:

(8, 63301), (Start: 10 @63250 has 1 MA's), (13, 63205), (15, 63196), (16, 63172), (17, 63169), (18, 63160), (19, 63115), (20, 63064),

Gene: Hosp\_078 Start: 61272, Stop: 61000, Start Num: 10

Candidate Starts for Hosp\_078:

(Start: 10 @61272 has 1 MA's), (13, 61227), (14, 61224), (15, 61218), (17, 61191), (18, 61182), (22, 61074),

Gene: Invubu\_85 Start: 64146, Stop: 63865, Start Num: 10

Candidate Starts for Invubu\_85:

(1, 64662), (2, 64644), (3, 64635), (6, 64239), (Start: 10 @64146 has 1 MA's), (11, 64143), (12, 64122), (25, 63915), (26, 63879),

Gene: Jolie1\_080 Start: 63299, Stop: 63027, Start Num: 10

Candidate Starts for Jolie1\_080:

(8, 63353), (Start: 10 @63299 has 1 MA's), (13, 63254), (14, 63251), (15, 63245), (17, 63218), (18, 63209), (22, 63101),

Gene: Rinkes\_79 Start: 64163, Stop: 63879, Start Num: 9

Candidate Starts for Rinkes\_79:

(4, 64562), (5, 64478), (7, 64220), (Start: 9 @64163 has 1 MA's), (15, 64106), (21, 63968), (23, 63956), (24, 63932),