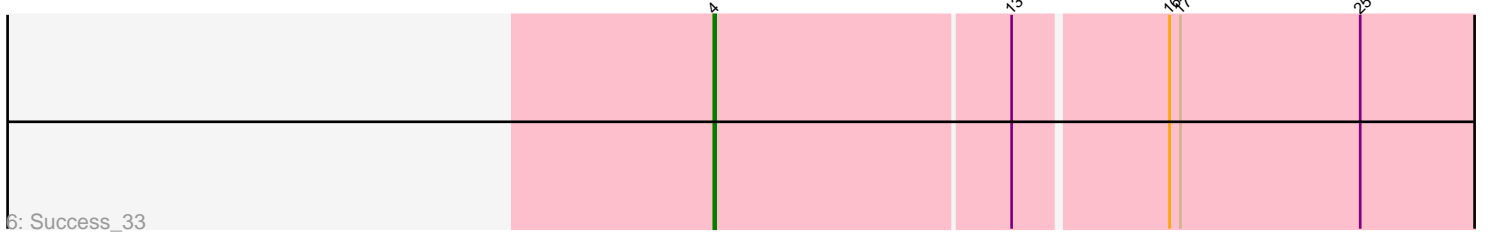
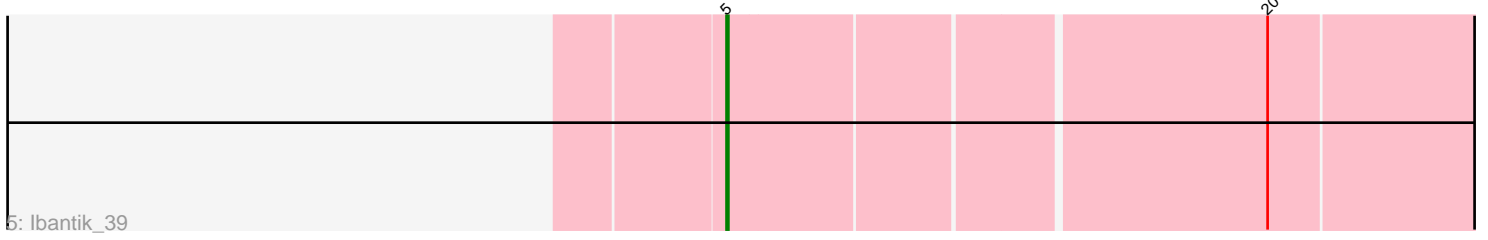
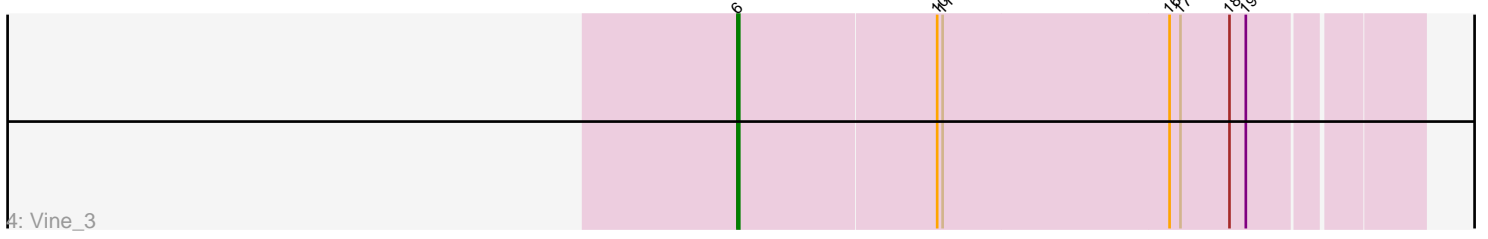
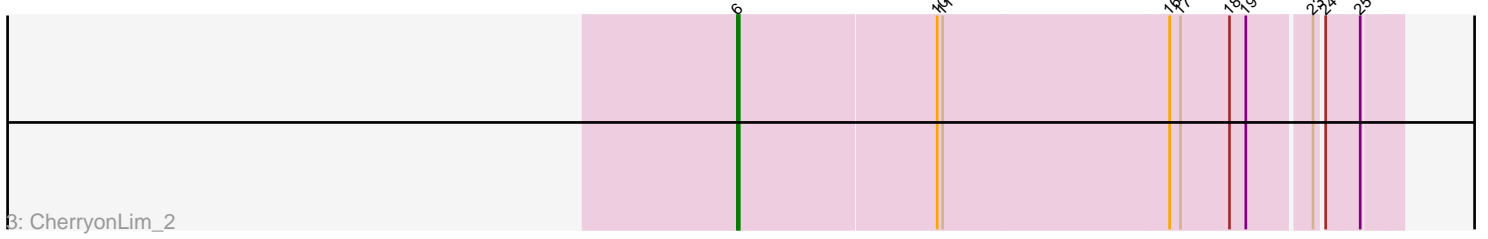
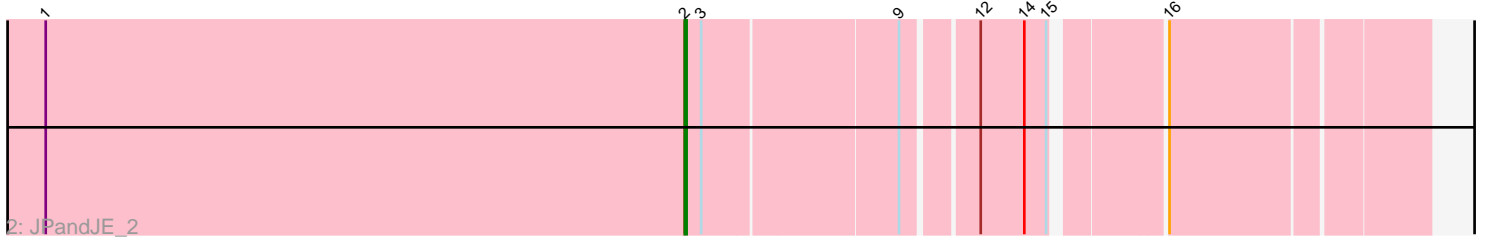
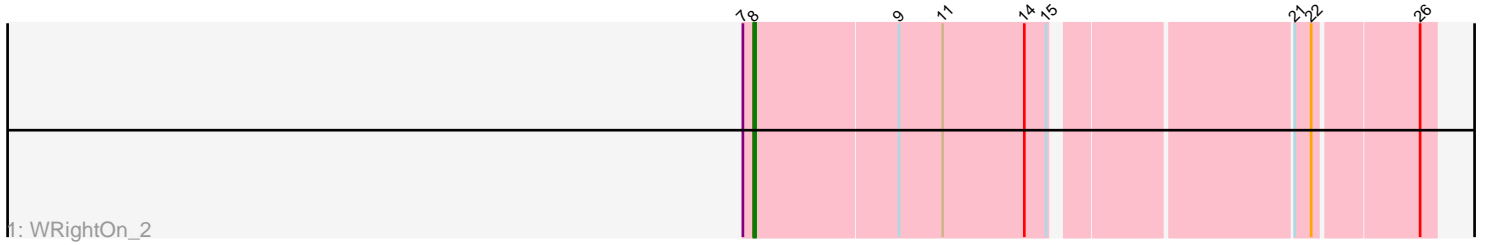


Pham 197211



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 197211 Report

This analysis was run 12/09/24 on database version 580.

Pham number 197211 has 6 members, 0 are drafts.

Phages represented in each track:

- Track 1 : WRightOn_2
- Track 2 : JPandJE_2
- Track 3 : CherryonLim_2
- Track 4 : Vine_3
- Track 5 : lbantik_39
- Track 6 : Success_33

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

Genes that call this "Most Annotated" start:

- CherryonLim_2, Vine_3,

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

-

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

- lbantik_39, JPandJE_2, Success_33, WRightOn_2,

Summary by start number:

Start 2:

- Found in 1 of 6 (16.7%) of genes in pham
- Manual Annotations of this start: 1 of 6
- Called 100.0% of time when present
- Phage (with cluster) where this start called: JPandJE_2 (BF),

Start 4:

- Found in 1 of 6 (16.7%) of genes in pham
- Manual Annotations of this start: 1 of 6
- Called 100.0% of time when present
- Phage (with cluster) where this start called: Success_33 (singleton),

Start 5:

- Found in 1 of 6 (16.7%) of genes in pham
- Manual Annotations of this start: 1 of 6
- Called 100.0% of time when present
- Phage (with cluster) where this start called: Ibantik_39 (singleton),

Start 6:

- Found in 2 of 6 (33.3%) of genes in pham
- Manual Annotations of this start: 2 of 6
- Called 100.0% of time when present
- Phage (with cluster) where this start called: CherryonLim_2 (CT), Vine_3 (CT),

Start 8:

- Found in 1 of 6 (16.7%) of genes in pham
- Manual Annotations of this start: 1 of 6
- Called 100.0% of time when present
- Phage (with cluster) where this start called: WRightOn_2 (BF),

Summary by clusters:

There are 3 clusters represented in this pham: BF, singleton, CT,

Info for manual annotations of cluster BF:

- Start number 2 was manually annotated 1 time for cluster BF.
- Start number 8 was manually annotated 1 time for cluster BF.

Info for manual annotations of cluster CT:

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

Gene Information:

Gene: CherryonLim_2 Start: 640, Stop: 990, Start Num: 6

Candidate Starts for CherryonLim_2:

(Start: 6 @640 has 2 MA's), (10, 748), (11, 751), (16, 874), (17, 880), (18, 907), (19, 916), (23, 949), (24, 952), (25, 970),

Gene: Ibantik_39 Start: 18281, Stop: 17859, Start Num: 5

Candidate Starts for Ibantik_39:

(Start: 5 @18281 has 1 MA's), (20, 17999),

Gene: JPandJE_2 Start: 2649, Stop: 3017, Start Num: 2

Candidate Starts for JPandJE_2:

(1, 2298), (Start: 2 @2649 has 1 MA's), (3, 2658), (9, 2760), (12, 2796), (14, 2820), (15, 2832), (16, 2883),

Gene: Success_33 Start: 20053, Stop: 19616, Start Num: 4

Candidate Starts for Success_33:

(Start: 4 @20053 has 1 MA's), (13, 19894), (16, 19813), (17, 19807), (25, 19708),

Gene: Vine_3 Start: 1265, Stop: 1630, Start Num: 6

Candidate Starts for Vine_3:

(Start: 6 @1265 has 2 MA's), (10, 1373), (11, 1376), (16, 1499), (17, 1505), (18, 1532), (19, 1541),

Gene: WRightOn_2 Start: 2629, Stop: 2976, Start Num: 8

Candidate Starts for WRightOn_2:

(7, 2626), (Start: 8 @2629 has 1 MA's), (9, 2707), (11, 2731), (14, 2776), (15, 2788), (21, 2905), (22, 2914), (26, 2968),