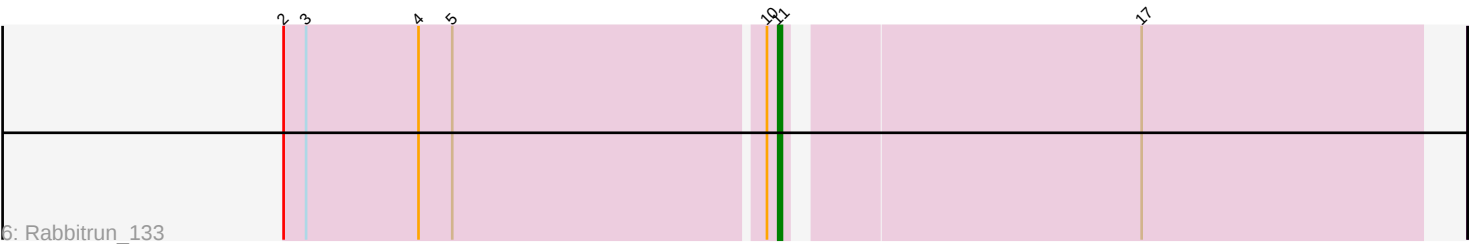
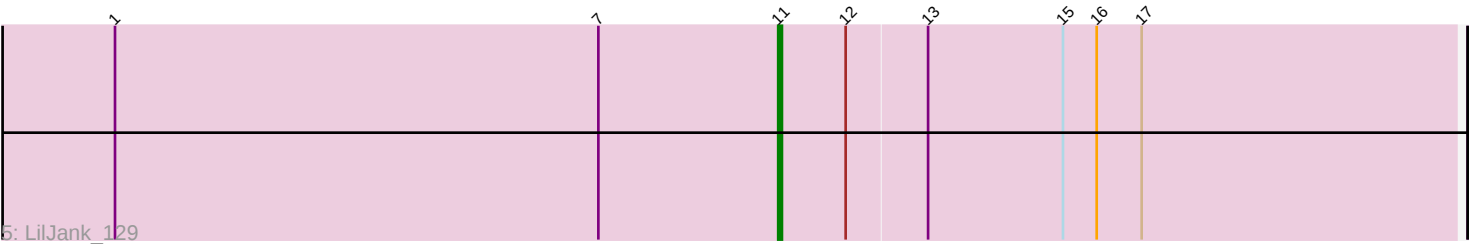
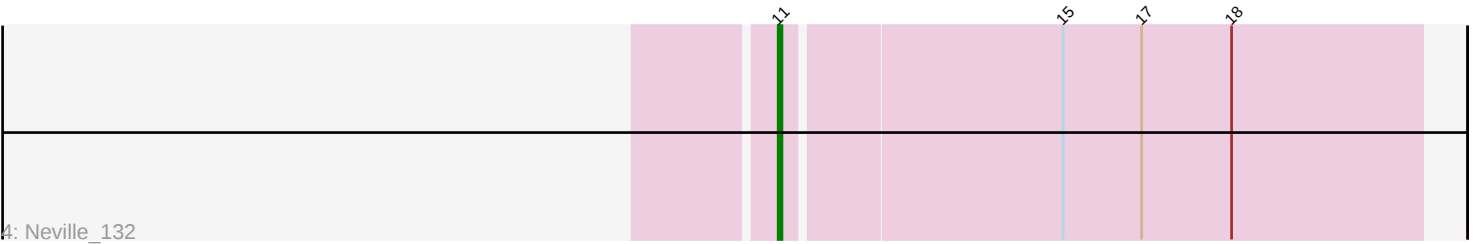
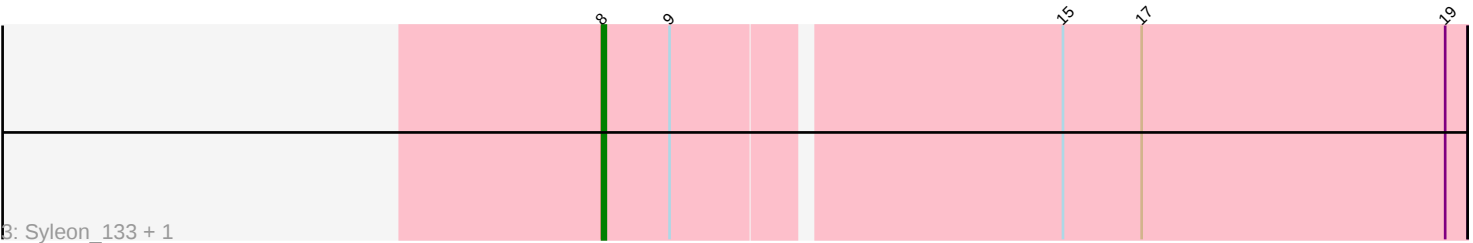
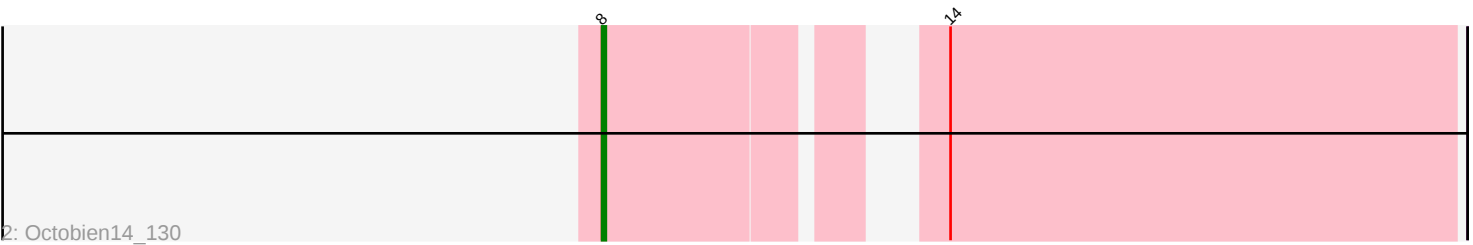
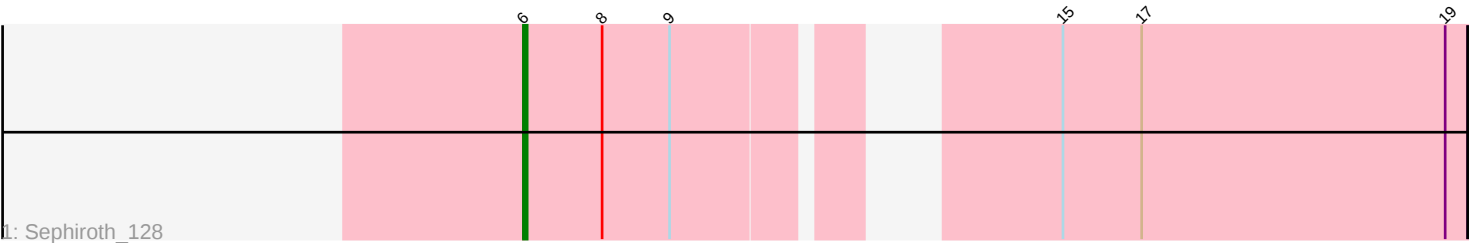


Pham 331593



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 331593 Report

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

Pham number 331593 has 7 members, 0 are drafts.

Phages represented in each track:

- Track 1 : Sephiroth_128
- Track 2 : Octobien14_130
- Track 3 : Syleon_133, Kudefre_134
- Track 4 : Neville_132
- Track 5 : LilJank_129
- Track 6 : Rabbitrun_133

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

The start number called the most often in the published annotations is 11, it was called in 3 of the 7 non-draft genes in the pham.

Genes that call this "Most Annotated" start:

- LilJank_129, Neville_132, Rabbitrun_133,

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

-

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

- Kudefre_134, Octobien14_130, Sephiroth_128, Syleon_133,

Summary by start number:

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: Sephiroth_128 (DU1),

Start 8:

- Found in 4 of 7 (57.1%) of genes in pham
- Manual Annotations of this start: 3 of 7
- Called 75.0% of time when present

- Phage (with cluster) where this start called: Kudefre_134 (DU1), Octobien14_130 (DU1), Syleon_133 (DU1),

Start 11:

- Found in 3 of 7 (42.9%) of genes in pham
- Manual Annotations of this start: 3 of 7
- Called 100.0% of time when present
- Phage (with cluster) where this start called: LilJank_129 (DU2), Neville_132 (DU2), Rabbitrun_133 (DU2),

Summary by clusters:

There are 2 clusters represented in this pham: DU1, DU2,

Info for manual annotations of cluster DU1:

- Start number 6 was manually annotated 1 time for cluster DU1.
- Start number 8 was manually annotated 3 times for cluster DU1.

Info for manual annotations of cluster DU2:

- Start number 11 was manually annotated 3 times for cluster DU2.

Gene Information:

Gene: Kudefre_134 Start: 71195, Stop: 70971, Start Num: 8

Candidate Starts for Kudefre_134:

(Start: 8 @71195 has 3 MA's), (9, 71177), (15, 71078), (17, 71057), (19, 70976),

Gene: LilJank_129 Start: 72754, Stop: 72575, Start Num: 11

Candidate Starts for LilJank_129:

(1, 72931), (7, 72802), (Start: 11 @72754 has 3 MA's), (12, 72736), (13, 72715), (15, 72679), (16, 72670), (17, 72658),

Gene: Neville_132 Start: 71846, Stop: 71679, Start Num: 11

Candidate Starts for Neville_132:

(Start: 11 @71846 has 3 MA's), (15, 71774), (17, 71753), (18, 71729),

Gene: Octobien14_130 Start: 69603, Stop: 69397, Start Num: 8

Candidate Starts for Octobien14_130:

(Start: 8 @69603 has 3 MA's), (14, 69531),

Gene: Rabbitrun_133 Start: 73473, Stop: 73309, Start Num: 11

Candidate Starts for Rabbitrun_133:

(2, 73602), (3, 73596), (4, 73566), (5, 73557), (10, 73476), (Start: 11 @73473 has 3 MA's), (17, 73383),

Gene: Sephiroth_128 Start: 70772, Stop: 70548, Start Num: 6

Candidate Starts for Sephiroth_128:

(Start: 6 @70772 has 1 MA's), (Start: 8 @70751 has 3 MA's), (9, 70733), (15, 70655), (17, 70634), (19, 70553),

Gene: Syleon_133 Start: 71523, Stop: 71299, Start Num: 8

Candidate Starts for Syleon_133:

(Start: 8 @71523 has 3 MA's), (9, 71505), (15, 71406), (17, 71385), (19, 71304),