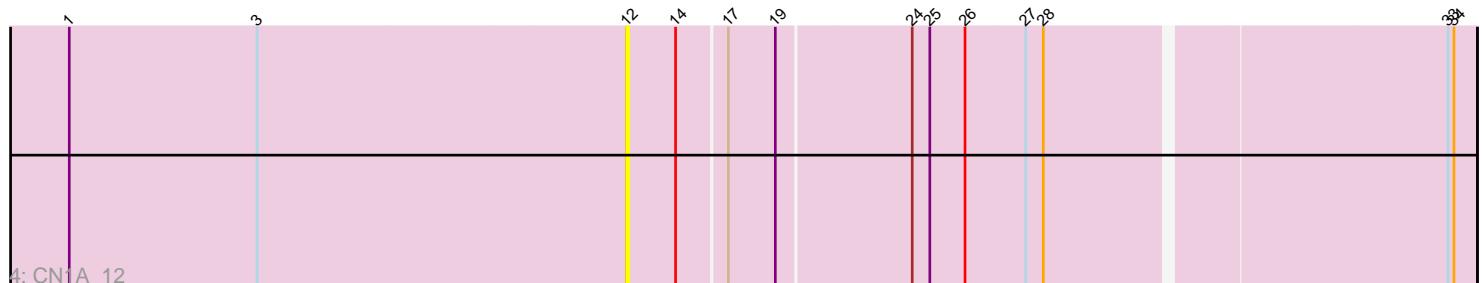
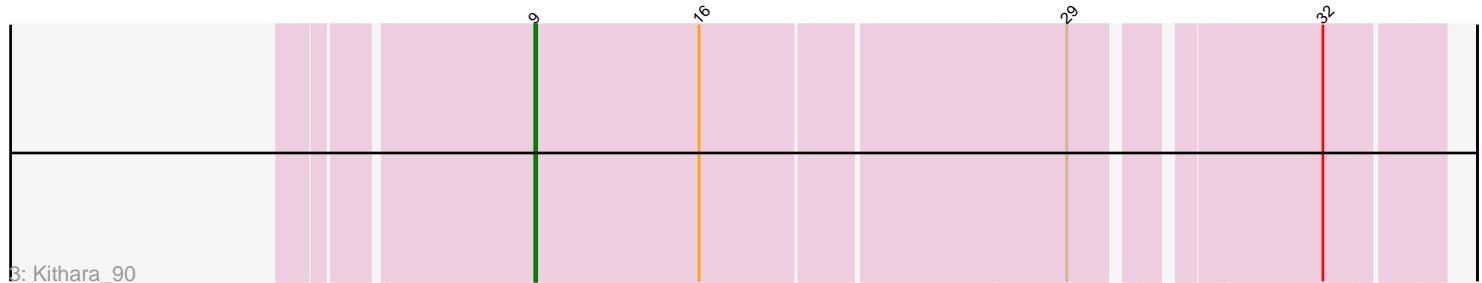
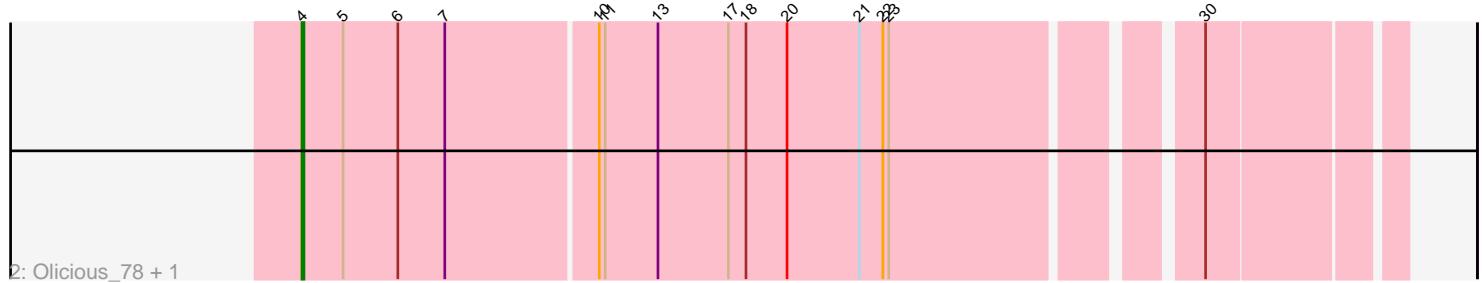
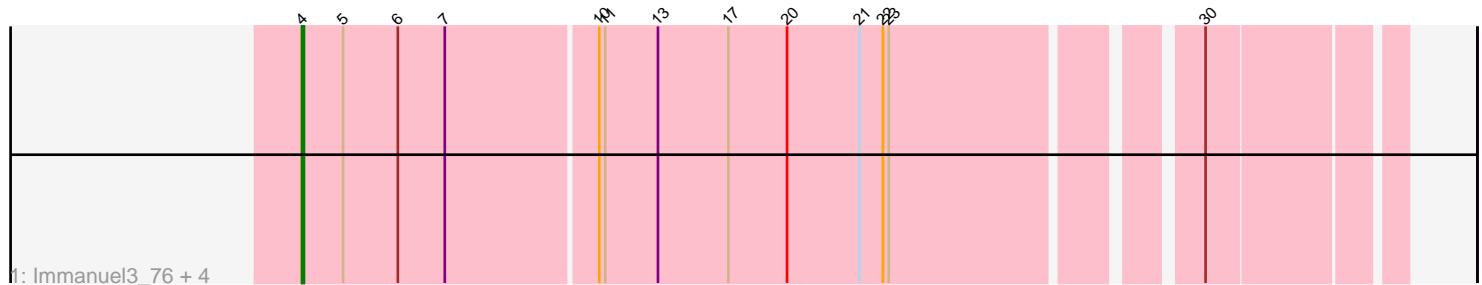


# Pham 281127



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 281127 Report

This analysis was run 02/07/26 on database version 634.

Pham number 281127 has 10 members, 1 are drafts.

Phages represented in each track:

- Track 1 : Immanuel3\_76, Percastrophe\_78, HaugeAnator\_78, ToriToki\_78, ZooBear\_78
- Track 2 : Olicious\_78, Romero\_78
- Track 3 : Kithara\_90
- Track 4 : CN1A\_12
- Track 5 : Yappy\_107

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

The start number called the most often in the published annotations is 4, it was called in 7 of the 9 non-draft genes in the pham.

Genes that call this "Most Annotated" start:

- HaugeAnator\_78, Immanuel3\_76, Olicious\_78, Percastrophe\_78, Romero\_78, ToriToki\_78, ZooBear\_78,

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

- 

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

- CN1A\_12, Kithara\_90, Yappy\_107,

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

Start 4:

- Found in 7 of 10 ( 70.0% ) of genes in pham
- Manual Annotations of this start: 7 of 9
- Called 100.0% of time when present
- Phage (with cluster) where this start called: HaugeAnator\_78 (BF), Immanuel3\_76 (BF), Olicious\_78 (BF), Percastrophe\_78 (BF), Romero\_78 (BF), ToriToki\_78 (BF), ZooBear\_78 (BF),

Start 8:

- Found in 1 of 10 ( 10.0% ) of genes in pham

- Manual Annotations of this start: 1 of 9
- Called 100.0% of time when present
- Phage (with cluster) where this start called: Yappy\_107 (singleton),

Start 9:

- Found in 1 of 10 ( 10.0% ) of genes in pham
- Manual Annotations of this start: 1 of 9
- Called 100.0% of time when present
- Phage (with cluster) where this start called: Kithara\_90 (singleton),

Start 12:

- Found in 1 of 10 ( 10.0% ) of genes in pham
- No Manual Annotations of this start.
- Called 100.0% of time when present
- Phage (with cluster) where this start called: CN1A\_12 (singleton),

### **Summary by clusters:**

There are 2 clusters represented in this pham: BF, singleton,

Info for manual annotations of cluster BF:

- Start number 4 was manually annotated 7 times for cluster BF.

### **Gene Information:**

Gene: CN1A\_12 Start: 5964, Stop: 6377, Start Num: 12

Candidate Starts for CN1A\_12:

(1, 5682), (3, 5778), (12, 5964), (14, 5988), (17, 6012), (19, 6036), (24, 6102), (25, 6111), (26, 6129), (27, 6159), (28, 6168), (33, 6363), (34, 6366),

Gene: HaugeAnator\_78 Start: 40177, Stop: 39656, Start Num: 4

Candidate Starts for HaugeAnator\_78:

(Start: 4 @40177 has 7 MA's), (5, 40156), (6, 40129), (7, 40105), (10, 40030), (11, 40027), (13, 40000), (17, 39964), (20, 39934), (21, 39898), (22, 39886), (23, 39883), (30, 39745),

Gene: Immanuel3\_76 Start: 40182, Stop: 39661, Start Num: 4

Candidate Starts for Immanuel3\_76:

(Start: 4 @40182 has 7 MA's), (5, 40161), (6, 40134), (7, 40110), (10, 40035), (11, 40032), (13, 40005), (17, 39969), (20, 39939), (21, 39903), (22, 39891), (23, 39888), (30, 39750),

Gene: Kithara\_90 Start: 59830, Stop: 59399, Start Num: 9

Candidate Starts for Kithara\_90:

(Start: 9 @59830 has 1 MA's), (16, 59746), (29, 59566), (32, 59455),

Gene: Olicious\_78 Start: 40180, Stop: 39659, Start Num: 4

Candidate Starts for Olicious\_78:

(Start: 4 @40180 has 7 MA's), (5, 40159), (6, 40132), (7, 40108), (10, 40033), (11, 40030), (13, 40003), (17, 39967), (18, 39958), (20, 39937), (21, 39901), (22, 39889), (23, 39886), (30, 39748),

Gene: Percastrophe\_78 Start: 40112, Stop: 39591, Start Num: 4

Candidate Starts for Percastrophe\_78:

(Start: 4 @40112 has 7 MA's), (5, 40091), (6, 40064), (7, 40040), (10, 39965), (11, 39962), (13, 39935), (17, 39899), (20, 39869), (21, 39833), (22, 39821), (23, 39818), (30, 39680),

Gene: Romero\_78 Start: 40173, Stop: 39652, Start Num: 4

Candidate Starts for Romero\_78:

(Start: 4 @40173 has 7 MA's), (5, 40152), (6, 40125), (7, 40101), (10, 40026), (11, 40023), (13, 39996), (17, 39960), (18, 39951), (20, 39930), (21, 39894), (22, 39882), (23, 39879), (30, 39741),

Gene: ToriToki\_78 Start: 40176, Stop: 39655, Start Num: 4

Candidate Starts for ToriToki\_78:

(Start: 4 @40176 has 7 MA's), (5, 40155), (6, 40128), (7, 40104), (10, 40029), (11, 40026), (13, 39999), (17, 39963), (20, 39933), (21, 39897), (22, 39885), (23, 39882), (30, 39744),

Gene: Yappy\_107 Start: 55269, Stop: 54802, Start Num: 8

Candidate Starts for Yappy\_107:

(2, 55377), (Start: 8 @55269 has 1 MA's), (15, 55164), (31, 54900),

Gene: ZooBear\_78 Start: 40177, Stop: 39656, Start Num: 4

Candidate Starts for ZooBear\_78:

(Start: 4 @40177 has 7 MA's), (5, 40156), (6, 40129), (7, 40105), (10, 40030), (11, 40027), (13, 40000), (17, 39964), (20, 39934), (21, 39898), (22, 39886), (23, 39883), (30, 39745),