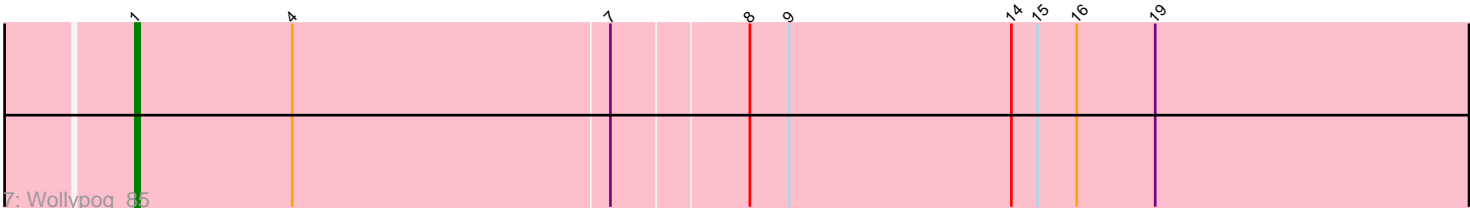
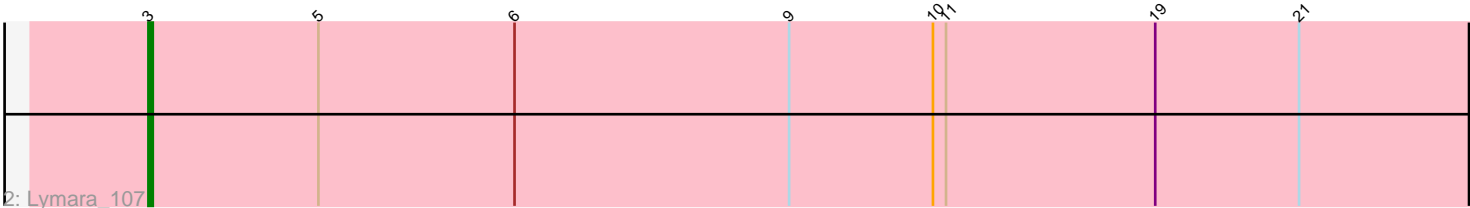
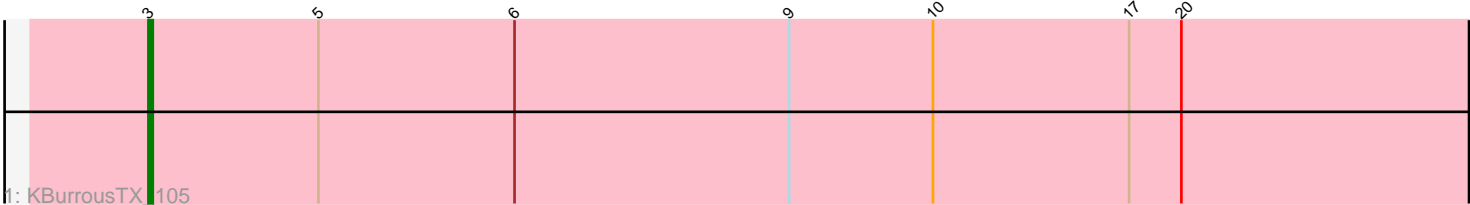


Pham 7037



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 7037 Report

This analysis was run 04/05/24 on database version 557.

Pham number 7037 has 7 members, 1 are drafts.

Phages represented in each track:

- Track 1 : KBurrousTX\_105
- Track 2 : Lymara\_107
- Track 3 : Colucci\_112
- Track 4 : ArV1\_096
- Track 5 : DrYang\_103
- Track 6 : Isolde\_97
- Track 7 : Wollypog\_85

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

The start number called the most often in the published annotations is 3, it was called in 4 of the 6 non-draft genes in the pham.

Genes that call this "Most Annotated" start:

- ArV1\_096, Colucci\_112, DrYang\_103, KBurrousTX\_105, Lymara\_107,

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

- 

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

- Isolde\_97, Wollypog\_85,

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

Start 1:

- Found in 1 of 7 ( 14.3% ) 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: Wollypog\_85 (singleton),

Start 2:

- Found in 1 of 7 ( 14.3% ) 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: Isolde\_97 (AY),

Start 3:

- Found in 5 of 7 ( 71.4% ) of genes in pham
- Manual Annotations of this start: 4 of 6
- Called 100.0% of time when present
- Phage (with cluster) where this start called: ArV1\_096 (AR), Colucci\_112 (AR), DrYang\_103 (AR), KBurrousTX\_105 (AR), Lymara\_107 (AR),

### **Summary by clusters:**

There are 3 clusters represented in this pham: AY, singleton, AR,

Info for manual annotations of cluster AR:

- Start number 3 was manually annotated 4 times for cluster AR.

Info for manual annotations of cluster AY:

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

### **Gene Information:**

Gene: ArV1\_096 Start: 68521, Stop: 68817, Start Num: 3

Candidate Starts for ArV1\_096:

(Start: 3 @68521 has 4 MA's), (6, 68599), (10, 68695), (12, 68701), (19, 68746), (22, 68788),

Gene: Colucci\_112 Start: 69441, Stop: 69737, Start Num: 3

Candidate Starts for Colucci\_112:

(Start: 3 @69441 has 4 MA's), (6, 69519), (10, 69615), (12, 69621), (13, 69630), (16, 69648), (19, 69666),

Gene: DrYang\_103 Start: 67786, Stop: 68094, Start Num: 3

Candidate Starts for DrYang\_103:

(Start: 3 @67786 has 4 MA's), (6, 67870), (10, 67966), (17, 68011), (20, 68023),

Gene: Isolde\_97 Start: 52510, Stop: 52809, Start Num: 2

Candidate Starts for Isolde\_97:

(Start: 2 @52510 has 1 MA's), (6, 52591), (18, 52735), (19, 52738),

Gene: KBurrousTX\_105 Start: 69717, Stop: 70025, Start Num: 3

Candidate Starts for KBurrousTX\_105:

(Start: 3 @69717 has 4 MA's), (5, 69756), (6, 69801), (9, 69864), (10, 69897), (17, 69942), (20, 69954),

Gene: Lymara\_107 Start: 67501, Stop: 67809, Start Num: 3

Candidate Starts for Lymara\_107:

(Start: 3 @67501 has 4 MA's), (5, 67540), (6, 67585), (9, 67648), (10, 67681), (11, 67684), (19, 67732), (21, 67765),

Gene: Wollypog\_85 Start: 57428, Stop: 57730, Start Num: 1

Candidate Starts for Wollypog\_85:

(Start: 1 @57428 has 1 MA's), (4, 57464), (7, 57536), (8, 57566), (9, 57575), (14, 57626), (15, 57632),  
(16, 57641), (19, 57659),