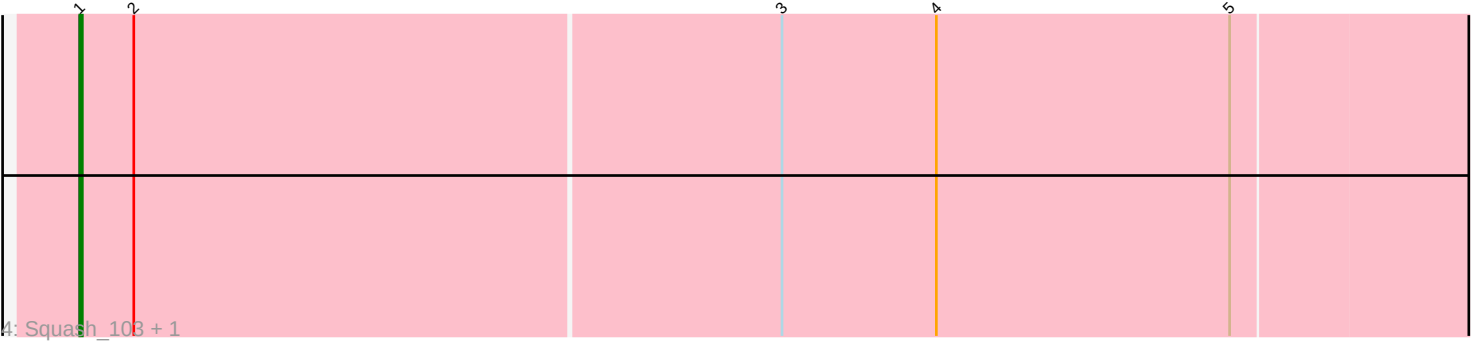
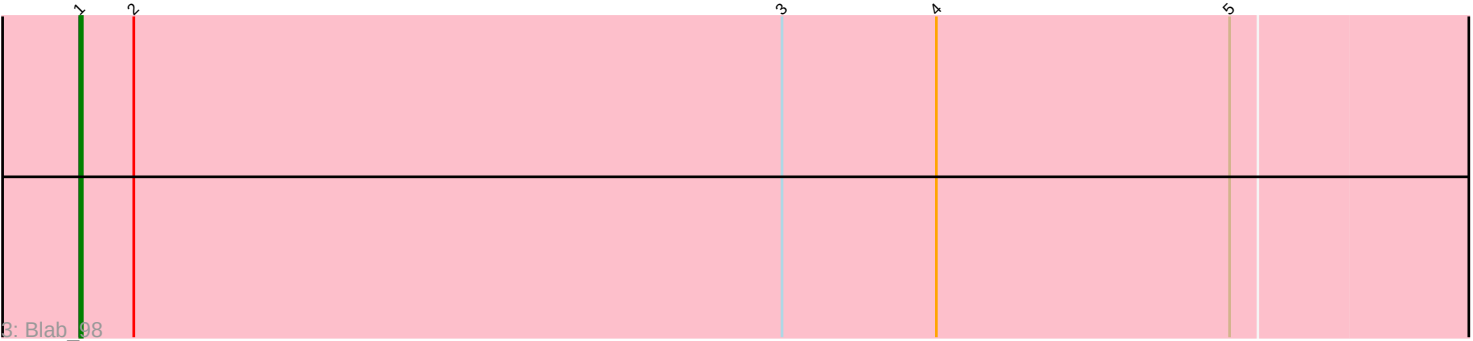
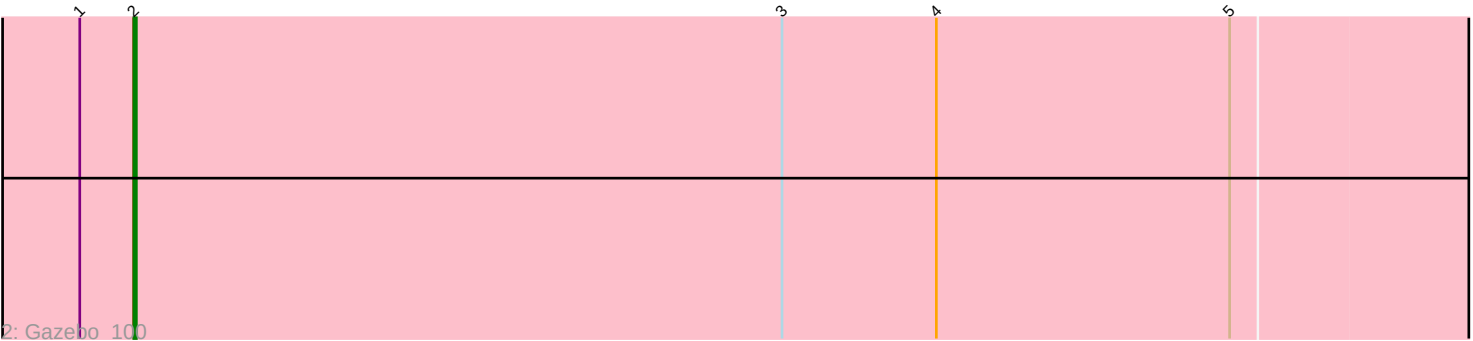
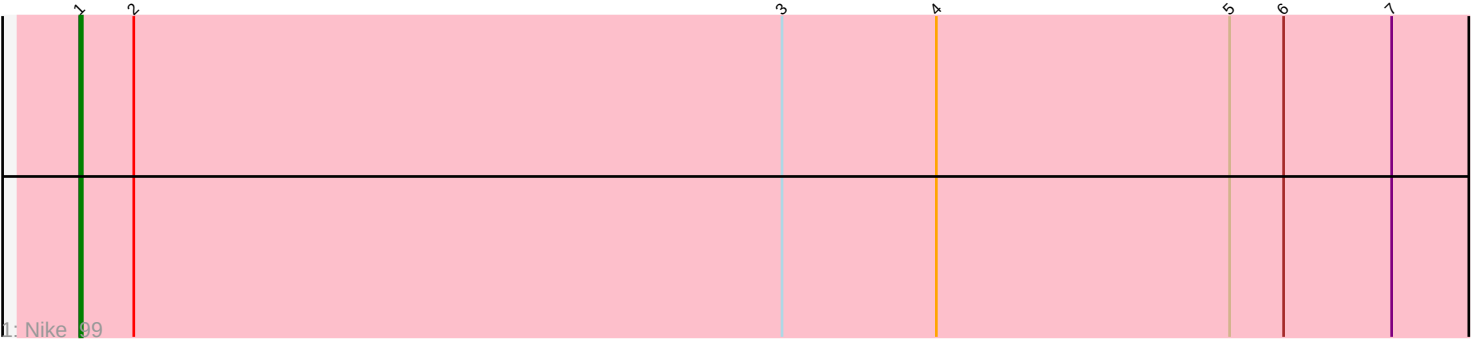




Note: Tracks are now grouped by subcluster and scaled. Switching in subcluster is indicated by changes in track color. Track scale is now set by default to display the region 30 bp upstream of start 1 to 30 bp downstream of the last possible start. If this default region is judged to be packed too tightly 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 331743 Report

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

Pham number 331743 has 5 members, 0 are drafts.

Phages represented in each track:

- Track 1 : Nike_99
- Track 2 : Gazebo_100
- Track 3 : Blab_98
- Track 4 : Squash_103, AluminumJesus_98

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

The start number called the most often in the published annotations is 1, it was called in 4 of the 5 non-draft genes in the pham.

Genes that call this "Most Annotated" start:

- AluminumJesus_98, Blab_98, Nike_99, Squash_103,

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

- Gazebo_100,

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

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Summary by start number:

Start 1:

- Found in 5 of 5 (100.0%) of genes in pham
- Manual Annotations of this start: 4 of 5
- Called 80.0% of time when present
- Phage (with cluster) where this start called: AluminumJesus_98 (EG), Blab_98 (EG), Nike_99 (EG), Squash_103 (EG),

Start 2:

- Found in 5 of 5 (100.0%) of genes in pham
- Manual Annotations of this start: 1 of 5
- Called 20.0% of time when present
- Phage (with cluster) where this start called: Gazebo_100 (EG),

Summary by clusters:

There is one cluster represented in this pham: EG

Info for manual annotations of cluster EG:

- Start number 1 was manually annotated 4 times for cluster EG.
- Start number 2 was manually annotated 1 time for cluster EG.

Gene Information:

Gene: AluminumJesus_98 Start: 59608, Stop: 59057, Start Num: 1

Candidate Starts for AluminumJesus_98:

(Start: 1 @59608 has 4 MA's), (Start: 2 @59587 has 1 MA's), (3, 59335), (4, 59275), (5, 59161),

Gene: Blab_98 Start: 60181, Stop: 59630, Start Num: 1

Candidate Starts for Blab_98:

(Start: 1 @60181 has 4 MA's), (Start: 2 @60160 has 1 MA's), (3, 59908), (4, 59848), (5, 59734),

Gene: Gazebo_100 Start: 60165, Stop: 59635, Start Num: 2

Candidate Starts for Gazebo_100:

(Start: 1 @60186 has 4 MA's), (Start: 2 @60165 has 1 MA's), (3, 59913), (4, 59853), (5, 59739),

Gene: Nike_99 Start: 60525, Stop: 59971, Start Num: 1

Candidate Starts for Nike_99:

(Start: 1 @60525 has 4 MA's), (Start: 2 @60504 has 1 MA's), (3, 60252), (4, 60192), (5, 60078), (6, 60057), (7, 60015),

Gene: Squash_103 Start: 60444, Stop: 59896, Start Num: 1

Candidate Starts for Squash_103:

(Start: 1 @60444 has 4 MA's), (Start: 2 @60423 has 1 MA's), (3, 60174), (4, 60114), (5, 60000),