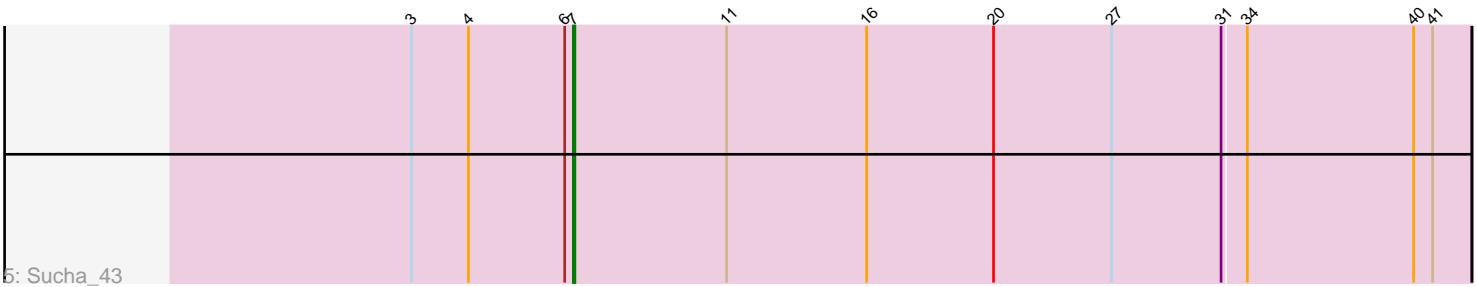
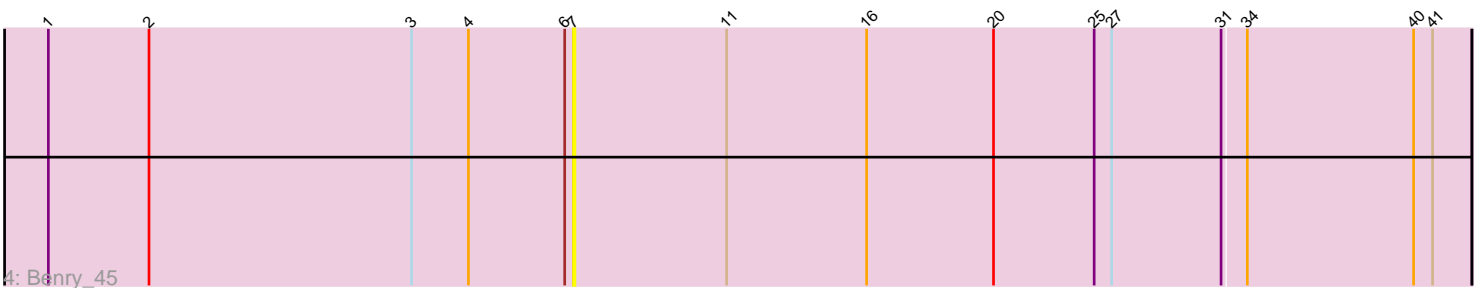
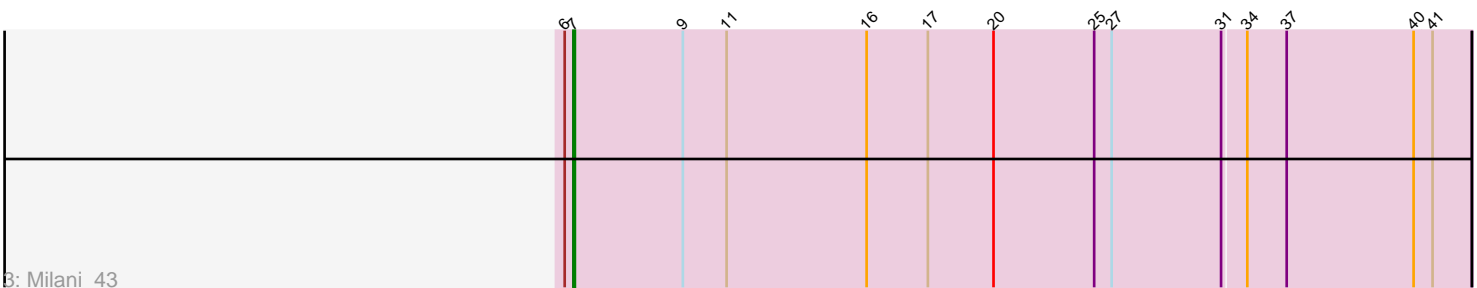
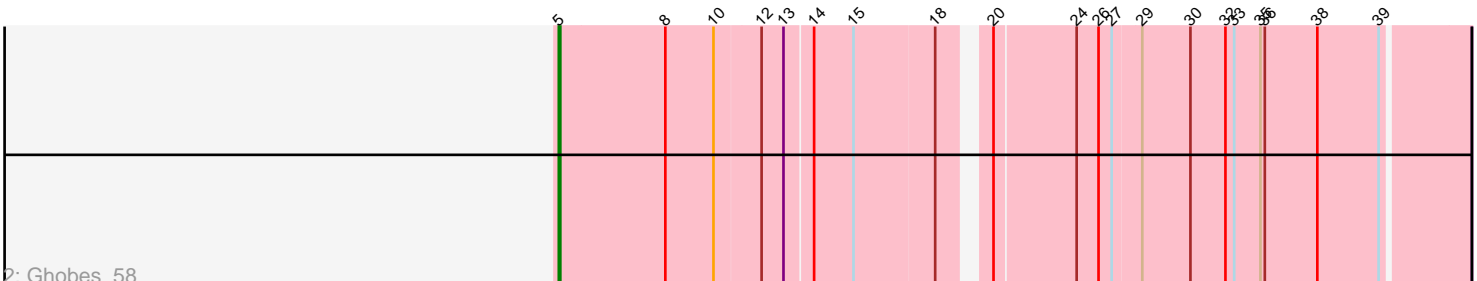
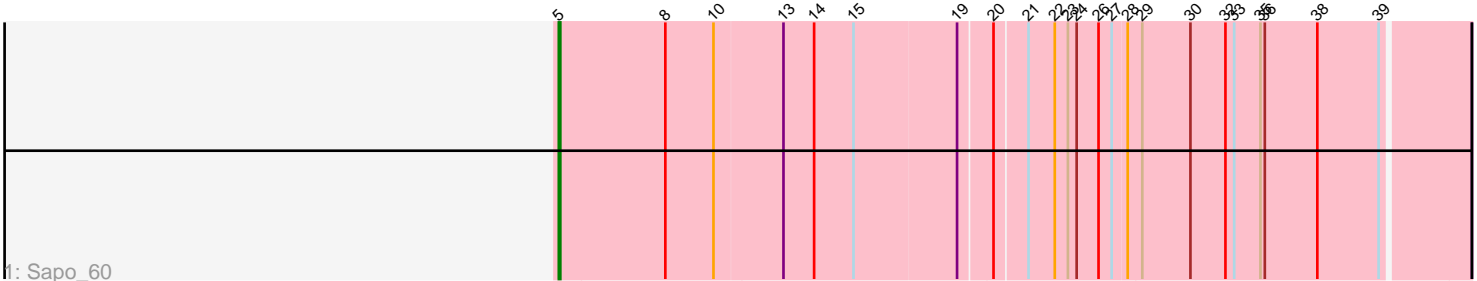


Pham 89120



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 89120 Report

This analysis was run 04/28/24 on database version 559.

Pham number 89120 has 5 members, 1 are drafts.

Phages represented in each track:

- Track 1 : Sapo_60
- Track 2 : Ghobes_58
- Track 3 : Milani_43
- Track 4 : Benry_45
- Track 5 : Sucha_43

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

The start number called the most often in the published annotations is 5, it was called in 2 of the 4 non-draft genes in the pham.

Genes that call this "Most Annotated" start:

- Ghobes_58, Sapo_60,

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

-

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

- Benry_45, Milani_43, Sucha_43,

Summary by start number:

Start 5:

- Found in 2 of 5 (40.0%) of genes in pham
- Manual Annotations of this start: 2 of 4
- Called 100.0% of time when present
- Phage (with cluster) where this start called: Ghobes_58 (DA), Sapo_60 (DA),

Start 7:

- Found in 3 of 5 (60.0%) of genes in pham
- Manual Annotations of this start: 2 of 4
- Called 100.0% of time when present
- Phage (with cluster) where this start called: Benry_45 (EJ), Milani_43 (EJ), Sucha_43 (EJ),

Summary by clusters:

There are 2 clusters represented in this pham: EJ, DA,

Info for manual annotations of cluster DA:

- Start number 5 was manually annotated 2 times for cluster DA.

Info for manual annotations of cluster EJ:

- Start number 7 was manually annotated 2 times for cluster EJ.

Gene Information:

Gene: Benry_45 Start: 27642, Stop: 28253, Start Num: 7

Candidate Starts for Benry_45:

(1, 27282), (2, 27351), (3, 27531), (4, 27570), (6, 27636), (Start: 7 @27642 has 2 MA's), (11, 27747), (16, 27843), (20, 27930), (25, 27999), (27, 28011), (31, 28086), (34, 28101), (40, 28215), (41, 28227),

Gene: Ghobes_58 Start: 44334, Stop: 44921, Start Num: 5

Candidate Starts for Ghobes_58:

(Start: 5 @44334 has 2 MA's), (8, 44406), (10, 44439), (12, 44469), (13, 44484), (14, 44502), (15, 44529), (18, 44583), (20, 44607), (24, 44661), (26, 44676), (27, 44685), (29, 44703), (30, 44736), (32, 44760), (33, 44766), (35, 44784), (36, 44787), (38, 44823), (39, 44865),

Gene: Milani_43 Start: 28064, Stop: 28675, Start Num: 7

Candidate Starts for Milani_43:

(6, 28058), (Start: 7 @28064 has 2 MA's), (9, 28139), (11, 28169), (16, 28265), (17, 28307), (20, 28352), (25, 28421), (27, 28433), (31, 28508), (34, 28523), (37, 28550), (40, 28637), (41, 28649),

Gene: Sapo_60 Start: 44598, Stop: 45200, Start Num: 5

Candidate Starts for Sapo_60:

(Start: 5 @44598 has 2 MA's), (8, 44670), (10, 44703), (13, 44748), (14, 44769), (15, 44796), (19, 44865), (20, 44886), (21, 44907), (22, 44925), (23, 44934), (24, 44940), (26, 44955), (27, 44964), (28, 44973), (29, 44982), (30, 45015), (32, 45039), (33, 45045), (35, 45063), (36, 45066), (38, 45102), (39, 45144),

Gene: Sucha_43 Start: 27259, Stop: 27870, Start Num: 7

Candidate Starts for Sucha_43:

(3, 27148), (4, 27187), (6, 27253), (Start: 7 @27259 has 2 MA's), (11, 27364), (16, 27460), (20, 27547), (27, 27628), (31, 27703), (34, 27718), (40, 27832), (41, 27844),