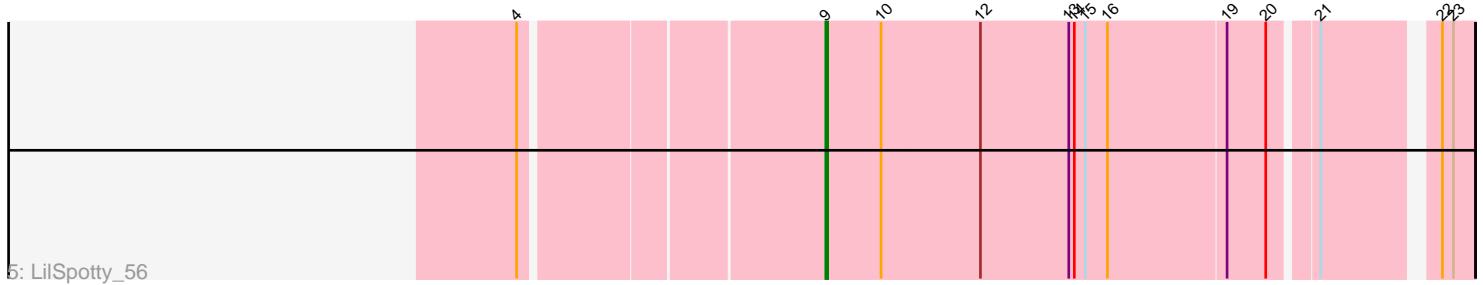
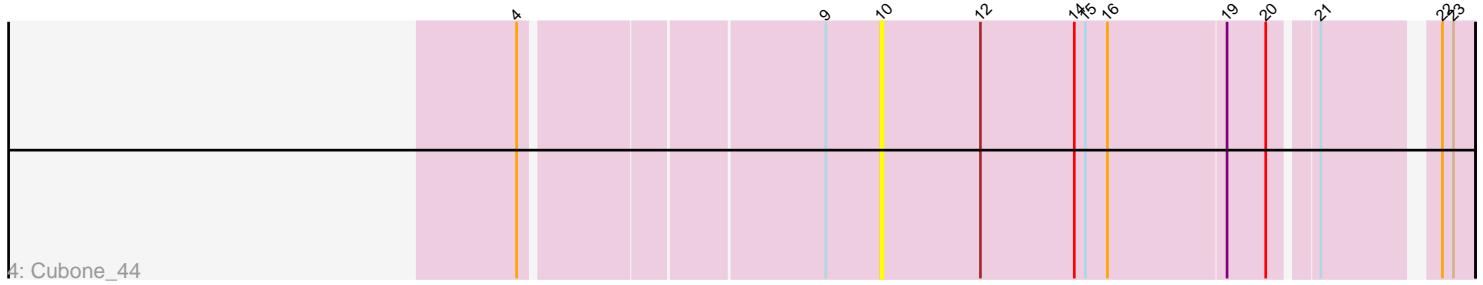
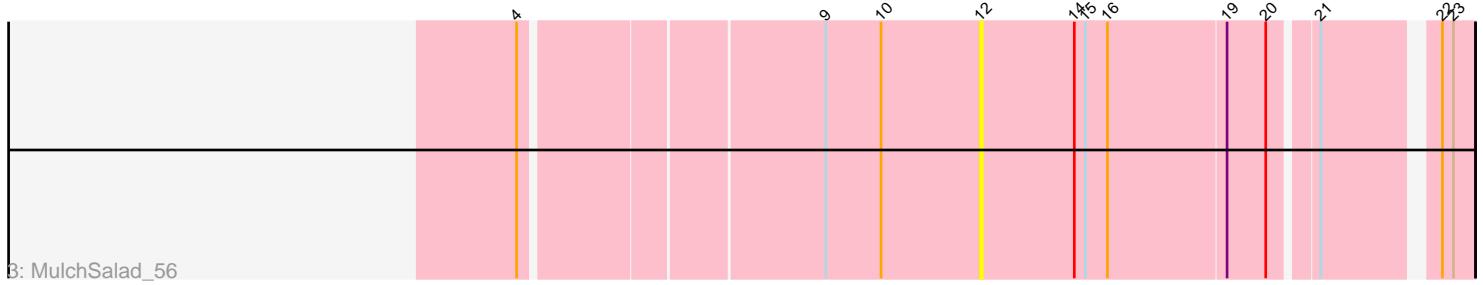
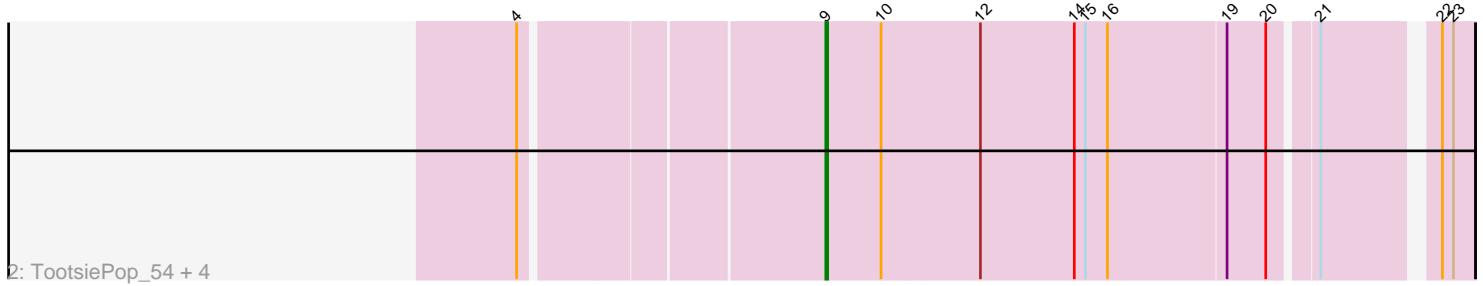
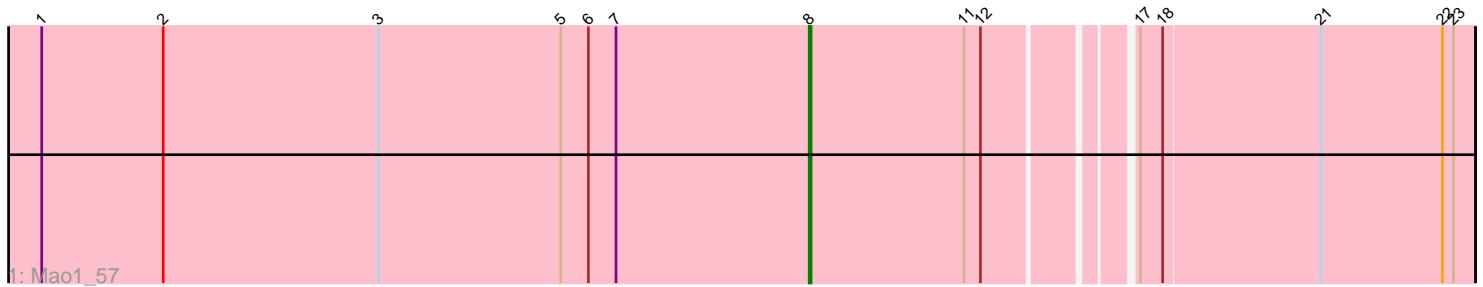


Pham 274946



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 274946 Report

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

Pham number 274946 has 9 members, 2 are drafts.

Phages represented in each track:

- Track 1 : Mao1_57
- Track 2 : TootsiePop_54, Misha28_54, Journey_41, Charlie_42, Awesomesauce_56
- Track 3 : MulchSalad_56
- Track 4 : Cubone_44
- Track 5 : LilSpotty_56

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

The start number called the most often in the published annotations is 9, it was called in 6 of the 7 non-draft genes in the pham.

Genes that call this "Most Annotated" start:

- Awesomesauce_56, Charlie_42, Journey_41, LilSpotty_56, Misha28_54, TootsiePop_54,

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

- Cubone_44, MulchSalad_56,

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

- Mao1_57,

Summary by start number:

Start 8:

- Found in 1 of 9 (11.1%) 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: Mao1_57 (AD),

Start 9:

- Found in 8 of 9 (88.9%) of genes in pham
- Manual Annotations of this start: 6 of 7
- Called 75.0% of time when present

- Phage (with cluster) where this start called: Awesomesauce_56 (F1), Charlie_42 (N), Journey_41 (N), LilSpotty_56 (singleton), Misha28_54 (F1), TootsiePop_54 (F1),

Start 10:

- Found in 8 of 9 (88.9%) of genes in pham
- No Manual Annotations of this start.
- Called 12.5% of time when present
- Phage (with cluster) where this start called: Cubone_44 (N),

Start 12:

- Found in 9 of 9 (100.0%) of genes in pham
- No Manual Annotations of this start.
- Called 11.1% of time when present
- Phage (with cluster) where this start called: MulchSalad_56 (F7),

Summary by clusters:

There are 5 clusters represented in this pham: F1, singleton, N, AD, F7,

Info for manual annotations of cluster AD:

- Start number 8 was manually annotated 1 time for cluster AD.

Info for manual annotations of cluster F1:

- Start number 9 was manually annotated 3 times for cluster F1.

Info for manual annotations of cluster N:

- Start number 9 was manually annotated 2 times for cluster N.

Gene Information:

Gene: Awesomesauce_56 Start: 37610, Stop: 37936, Start Num: 9

Candidate Starts for Awesomesauce_56:

(4, 37457), (Start: 9 @37610 has 6 MA's), (10, 37640), (12, 37691), (14, 37742), (15, 37748), (16, 37760), (19, 37823), (20, 37844), (21, 37865), (22, 37919), (23, 37925),

Gene: Charlie_42 Start: 29742, Stop: 30068, Start Num: 9

Candidate Starts for Charlie_42:

(4, 29589), (Start: 9 @29742 has 6 MA's), (10, 29772), (12, 29823), (14, 29874), (15, 29880), (16, 29892), (19, 29955), (20, 29976), (21, 29997), (22, 30051), (23, 30057),

Gene: Cubone_44 Start: 29794, Stop: 30090, Start Num: 10

Candidate Starts for Cubone_44:

(4, 29611), (Start: 9 @29764 has 6 MA's), (10, 29794), (12, 29845), (14, 29896), (15, 29902), (16, 29914), (19, 29977), (20, 29998), (21, 30019), (22, 30073), (23, 30079),

Gene: Journey_41 Start: 29742, Stop: 30068, Start Num: 9

Candidate Starts for Journey_41:

(4, 29589), (Start: 9 @29742 has 6 MA's), (10, 29772), (12, 29823), (14, 29874), (15, 29880), (16, 29892), (19, 29955), (20, 29976), (21, 29997), (22, 30051), (23, 30057),

Gene: LilSpotty_56 Start: 36924, Stop: 37250, Start Num: 9

Candidate Starts for LilSpotty_56:

(4, 36771), (Start: 9 @36924 has 6 MA's), (10, 36954), (12, 37005), (13, 37053), (14, 37056), (15, 37062), (16, 37074), (19, 37137), (20, 37158), (21, 37179), (22, 37233), (23, 37239),

Gene: Mao1_57 Start: 46089, Stop: 46430, Start Num: 8

Candidate Starts for Mao1_57:

(1, 45672), (2, 45738), (3, 45855), (5, 45954), (6, 45969), (7, 45984), (Start: 8 @46089 has 1 MA's), (11, 46173), (12, 46182), (17, 46251), (18, 46263), (21, 46347), (22, 46413), (23, 46419),

Gene: Misha28_54 Start: 37615, Stop: 37941, Start Num: 9

Candidate Starts for Misha28_54:

(4, 37462), (Start: 9 @37615 has 6 MA's), (10, 37645), (12, 37696), (14, 37747), (15, 37753), (16, 37765), (19, 37828), (20, 37849), (21, 37870), (22, 37924), (23, 37930),

Gene: MulchSalad_56 Start: 36879, Stop: 37124, Start Num: 12

Candidate Starts for MulchSalad_56:

(4, 36645), (Start: 9 @36798 has 6 MA's), (10, 36828), (12, 36879), (14, 36930), (15, 36936), (16, 36948), (19, 37011), (20, 37032), (21, 37053), (22, 37107), (23, 37113),

Gene: TootsiePop_54 Start: 37615, Stop: 37941, Start Num: 9

Candidate Starts for TootsiePop_54:

(4, 37462), (Start: 9 @37615 has 6 MA's), (10, 37645), (12, 37696), (14, 37747), (15, 37753), (16, 37765), (19, 37828), (20, 37849), (21, 37870), (22, 37924), (23, 37930),