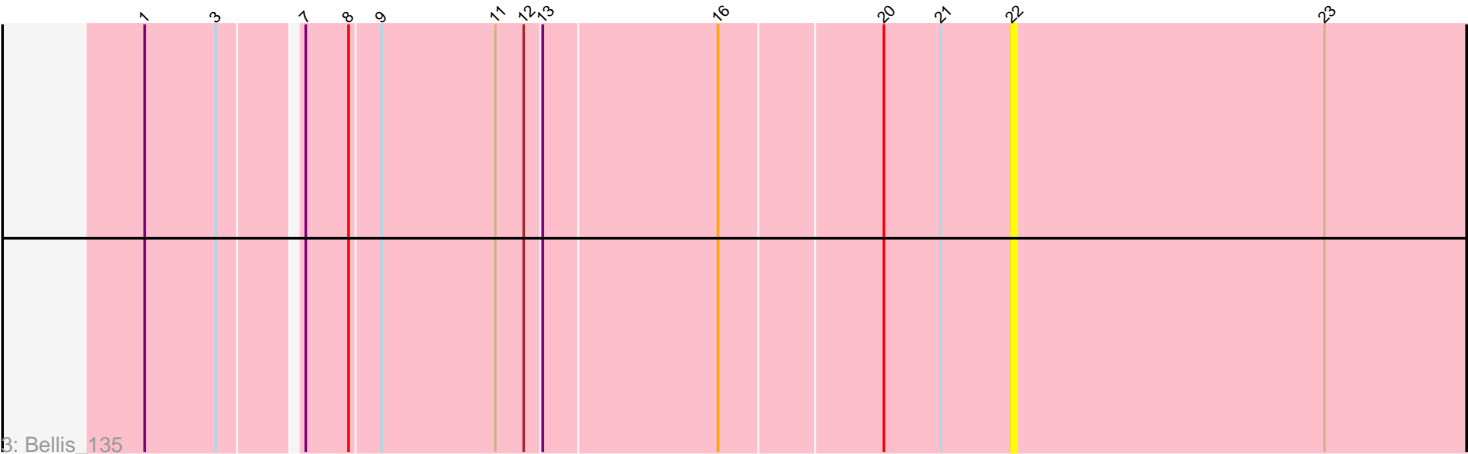
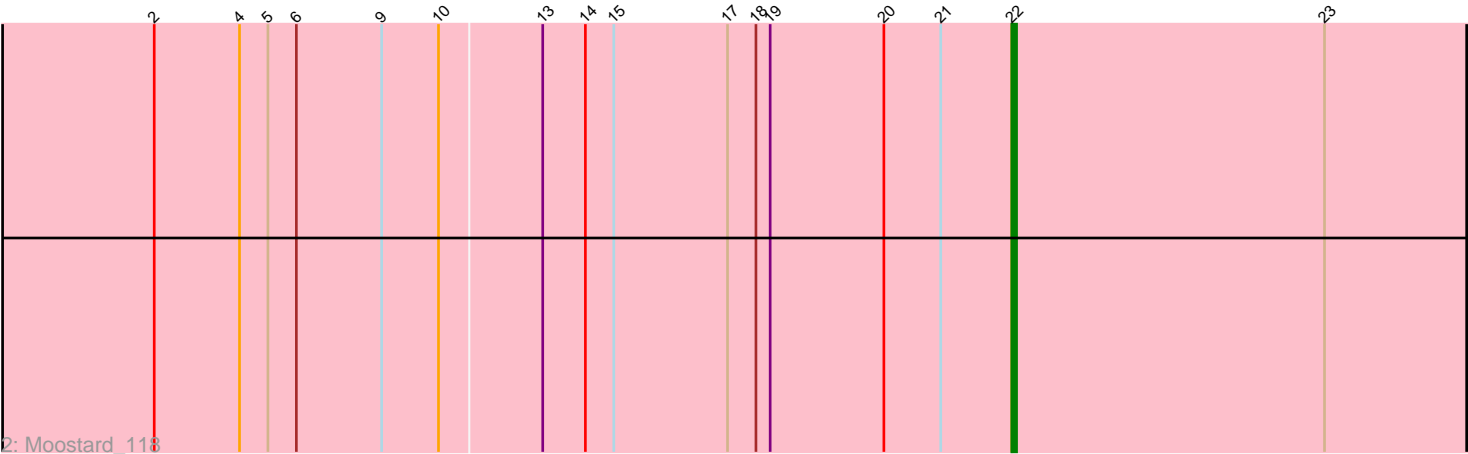
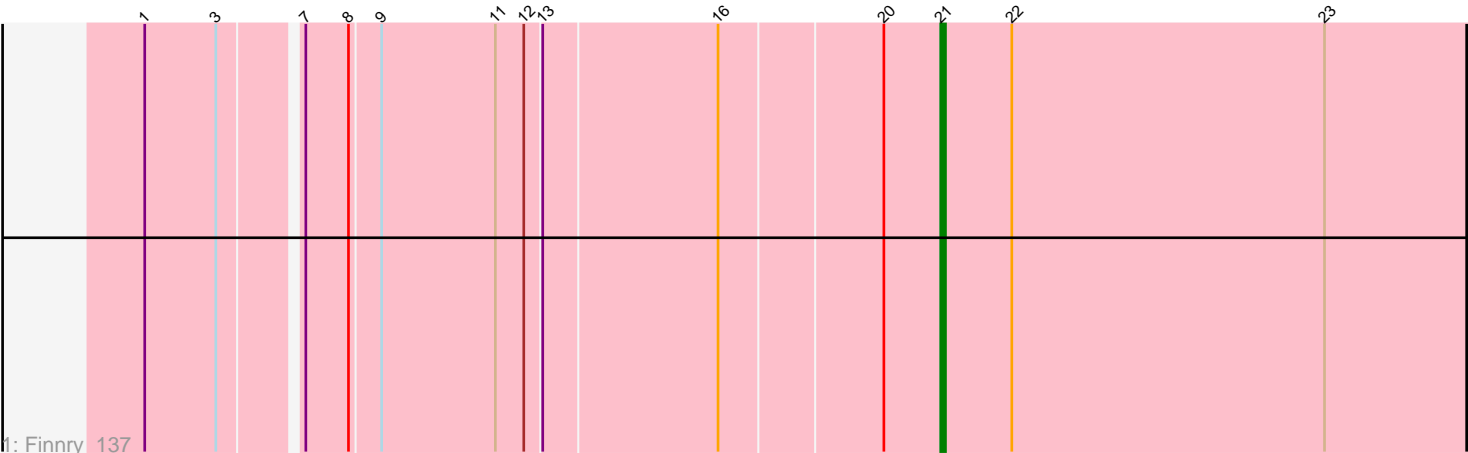


Pham 109395



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 109395 Report

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

Pham number 109395 has 3 members, 1 are drafts.

Phages represented in each track:

- Track 1 : Finnry_137
- Track 2 : Moostard_118
- Track 3 : Bellis_135

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

The start number called the most often in the published annotations is 21, it was called in 1 of the 2 non-draft genes in the pham.

Genes that call this "Most Annotated" start:

- Finnry_137,

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

- Bellis_135, Moostard_118,

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

-

Summary by start number:

Start 21:

- Found in 3 of 3 (100.0%) of genes in pham
- Manual Annotations of this start: 1 of 2
- Called 33.3% of time when present
- Phage (with cluster) where this start called: Finnry_137 (L3),

Start 22:

- Found in 3 of 3 (100.0%) of genes in pham
- Manual Annotations of this start: 1 of 2
- Called 66.7% of time when present
- Phage (with cluster) where this start called: Bellis_135 (L3), Moostard_118 (L3),

Summary by clusters:

There is one cluster represented in this pham: L3

Info for manual annotations of cluster L3:

- Start number 21 was manually annotated 1 time for cluster L3.
- Start number 22 was manually annotated 1 time for cluster L3.

Gene Information:

Gene: Bellis_135 Start: 74302, Stop: 74189, Start Num: 22

Candidate Starts for Bellis_135:

(1, 74476), (3, 74461), (7, 74446), (8, 74437), (9, 74431), (11, 74407), (12, 74401), (13, 74398), (16, 74362), (20, 74329), (Start: 21 @74317 has 1 MA's), (Start: 22 @74302 has 1 MA's), (23, 74236),

Gene: Finnry_137 Start: 74774, Stop: 74646, Start Num: 21

Candidate Starts for Finnry_137:

(1, 74933), (3, 74918), (7, 74903), (8, 74894), (9, 74888), (11, 74864), (12, 74858), (13, 74855), (16, 74819), (20, 74786), (Start: 21 @74774 has 1 MA's), (Start: 22 @74759 has 1 MA's), (23, 74693),

Gene: Moostard_118 Start: 68644, Stop: 68531, Start Num: 22

Candidate Starts for Moostard_118:

(2, 68824), (4, 68806), (5, 68800), (6, 68794), (9, 68776), (10, 68764), (13, 68743), (14, 68734), (15, 68728), (17, 68704), (18, 68698), (19, 68695), (20, 68671), (Start: 21 @68659 has 1 MA's), (Start: 22 @68644 has 1 MA's), (23, 68578),