Pham 9220

1: MacGully_101

2: Dalilpop_93

3: GTE5_78

4: Flapper_94

5: GRU1_80
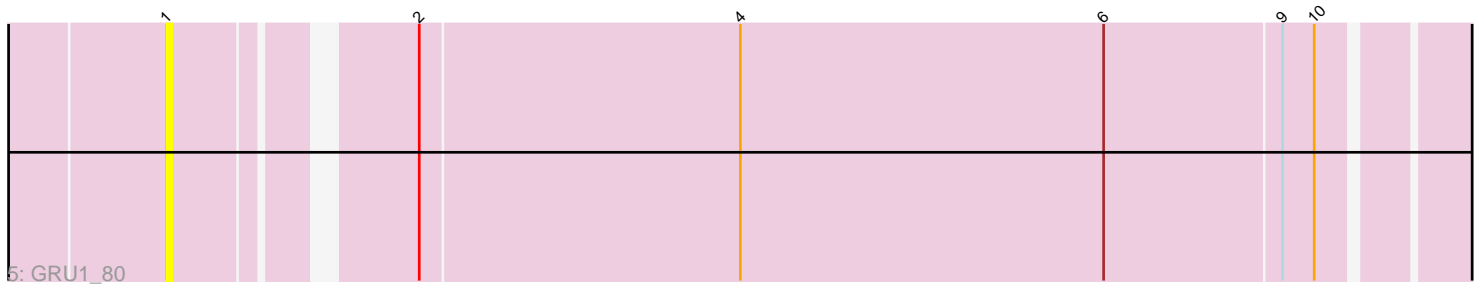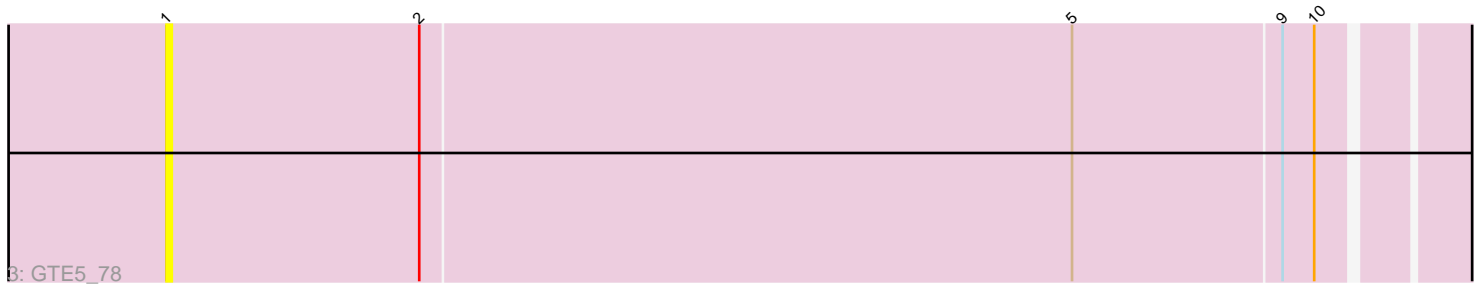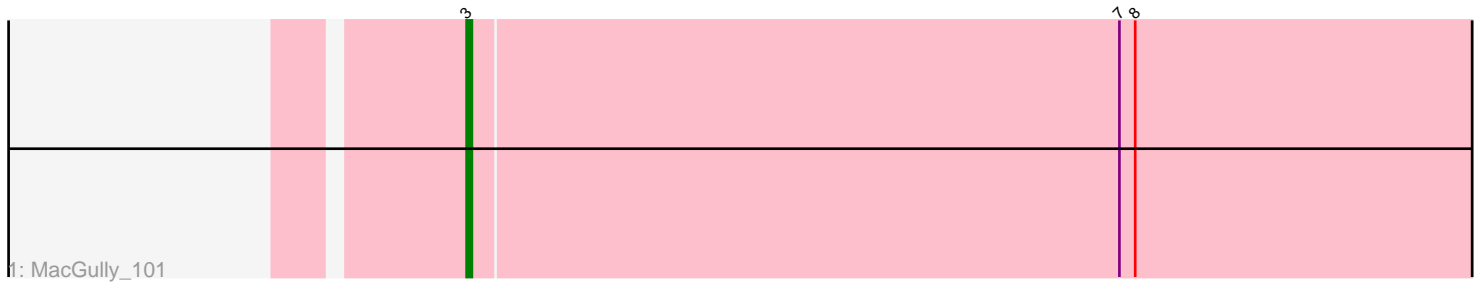
Note: Tracks are now grouped by subcluster and scaled. Switching in subcluster is indicated by changes in track color. Track scale is now set by default to display the region 30 bp upstream of start 1 to 30 bp downstream of the last possible start. If this default region is judged to be packed too tightly 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 9220 Report

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

Pham number 9220 has 5 members, 2 are drafts.

Phages represented in each track:
• Track 1 : MacGully_101
• Track 2 : Dalilpop_93
• Track 3 : GTE5_78
• Track 4 : Flapper_94
• Track 5 : GRU1_80

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 2 of the 3 non-draft genes in the pham.

Genes that call this "Most Annotated" start:
• Dalilpop_93, Flapper_94, GRU1_80, GTE5_78,

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

Genes that do not have the "Most Annotated" start:
• MacGully_101,

Summary by start number:

Start 1:
• Found in 4 of 5 (80.0%) of genes in pham
• Manual Annotations of this start: 2 of 3
• Called 100.0% of time when present
• Phage (with cluster) where this start called: Dalilpop_93 (CR1), Flapper_94 (CR1), GRU1_80 (CR1), GTE5_78 (CR1),

Start 3:
• Found in 1 of 5 (20.0%) of genes in pham
• Manual Annotations of this start: 1 of 3
• Called 100.0% of time when present
• Phage (with cluster) where this start called: MacGully_101 (CR),

Summary by clusters:

There are 2 clusters represented in this pham: CR, CR1,

Info for manual annotations of cluster CR:
•Start number 3 was manually annotated 1 time for cluster CR.

Info for manual annotations of cluster CR1:
•Start number 1 was manually annotated 2 times for cluster CR1.

Gene Information:
Gene: Dalilpop_93 Start: 65317, Stop: 65595, Start Num: 1
Candidate Starts for Dalilpop_93:
(Start: 1 @65317 has 2 MA's), (2, 65365), (5, 65488), (6, 65494), (9, 65527), (10, 65533),

Gene: Flapper_94 Start: 65831, Stop: 66103, Start Num: 1
Candidate Starts for Flapper_94:
(Start: 1 @65831 has 2 MA's), (2, 65873), (5, 65996), (6, 66002),

Gene: GRU1_80 Start: 57406, Stop: 57675, Start Num: 1
Candidate Starts for GRU1_80:
(Start: 1 @57406 has 2 MA's), (2, 57445), (4, 57505), (6, 57574), (9, 57607), (10, 57613),

Gene: GTE5_78 Start: 57674, Stop: 57952, Start Num: 1
Candidate Starts for GTE5_78:
(Start: 1 @57674 has 2 MA's), (2, 57722), (5, 57845), (9, 57884), (10, 57890),

Gene: MacGully_101 Start: 65485, Stop: 65712, Start Num: 3
Candidate Starts for MacGully_101:
(Start: 3 @65485 has 1 MA's), (7, 65608), (8, 65611),