Pham 11779
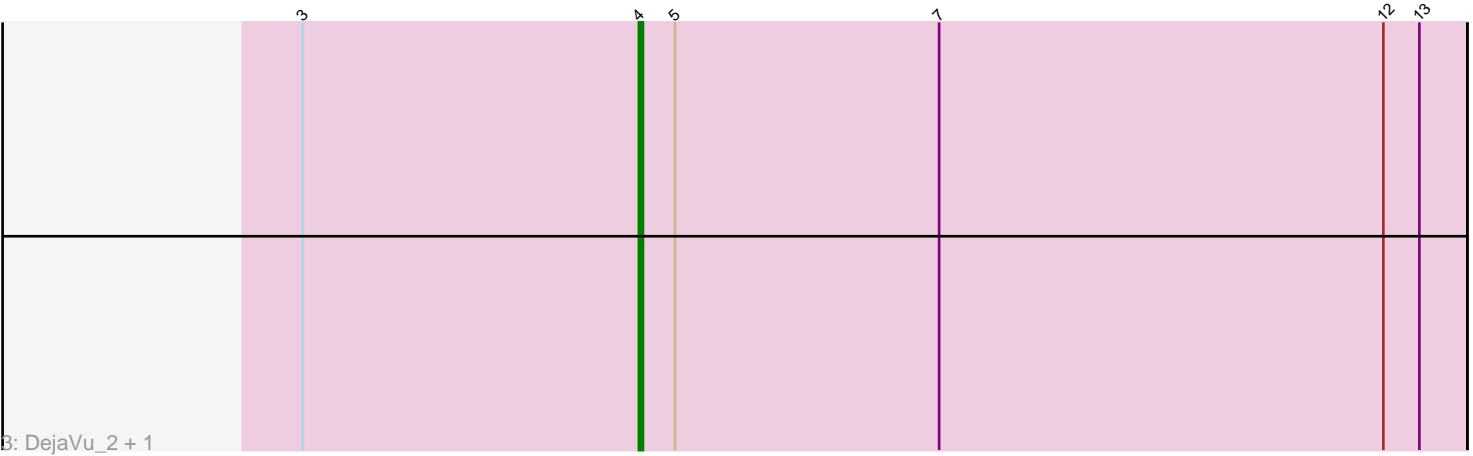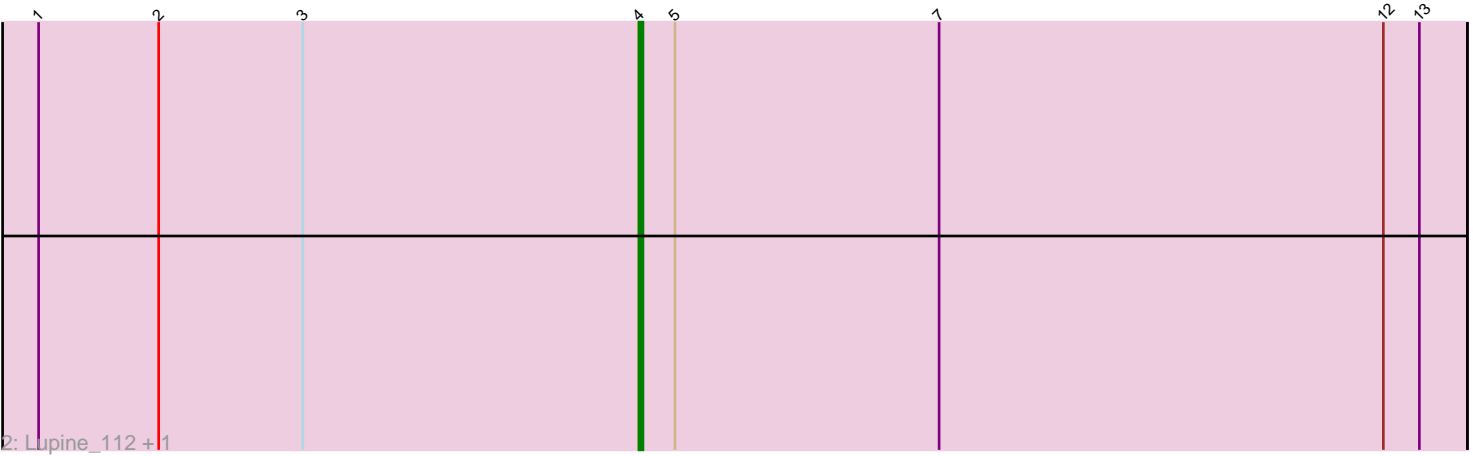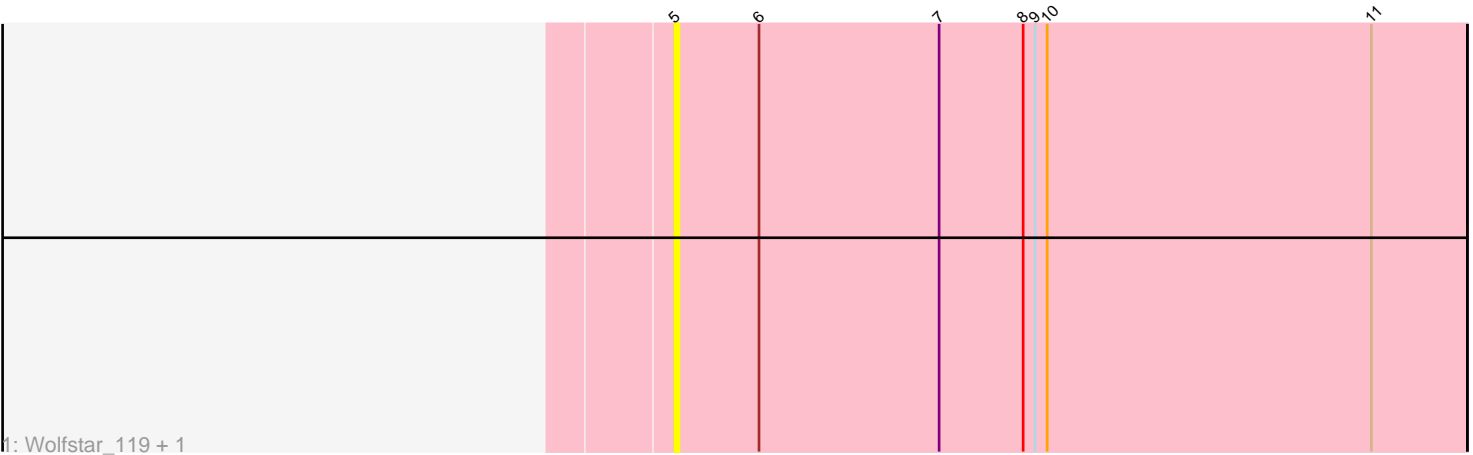

1: Wolfstar_119 + 1

2: Lupine_112 + 1

3: DejaVu_2 + 1

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 11779 Report

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

WARNING: Pham size does not match number of genes in report. Either unphamerated genes have been added (by you) or starterator has removed genes due to invalid start codon.

Pham number 11779 has 6 members, 2 are drafts.

Phages represented in each track:
• Track 1 : Wolfstar_119, Wolfstar_3
• Track 2 : Lupine_112, Lupine_2
• Track 3 : DejaVu_2, DejaVu_118

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

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

Genes that call this "Most Annotated" start:
• DejaVu_118, DejaVu_2, Lupine_112, Lupine_2,

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

Genes that do not have the "Most Annotated" start:
• Wolfstar_119, Wolfstar_3,

Summary by start number:

Start 4:
• Found in 4 of 6 (66.7%) of genes in pham
• Manual Annotations of this start: 4 of 4
• Called 100.0% of time when present
• Phage (with cluster) where this start called: DejaVu_118 (ED1), DejaVu_2 (ED1), Lupine_112 (ED1), Lupine_2 (ED1),

Start 5:
• Found in 6 of 6 (100.0%) of genes in pham
• No Manual Annotations of this start.

• Called 33.3% of time when present
• Phage (with cluster) where this start called: Wolfstar_119 (ED), Wolfstar_3 (ED),

Summary by clusters:

There are 2 clusters represented in this pham: ED, ED1,

Info for manual annotations of cluster ED1:
•Start number 4 was manually annotated 4 times for cluster ED1.

Gene Information:

Gene: DejaVu_2 Start: 754, Stop: 548, Start Num: 4
Candidate Starts for DejaVu_2:
(3, 838), (Start: 4 @754 has 4 MA's), (5, 745), (7, 679), (12, 568), (13, 559),

Gene: DejaVu_118 Start: 61134, Stop: 60928, Start Num: 4
Candidate Starts for DejaVu_118:
(3, 61218), (Start: 4 @61134 has 4 MA's), (5, 61125), (7, 61059), (12, 60948), (13, 60939),

Gene: Lupine_112 Start: 60015, Stop: 59809, Start Num: 4
Candidate Starts for Lupine_112:
(1, 60165), (2, 60135), (3, 60099), (Start: 4 @60015 has 4 MA's), (5, 60006), (7, 59940), (12, 59829), (13, 59820),

Gene: Lupine_2 Start: 742, Stop: 536, Start Num: 4
Candidate Starts for Lupine_2:
(1, 892), (2, 862), (3, 826), (Start: 4 @742 has 4 MA's), (5, 733), (7, 667), (12, 556), (13, 547),

Gene: Wolfstar_119 Start: 62235, Stop: 62038, Start Num: 5
Candidate Starts for Wolfstar_119:
(5, 62235), (6, 62214), (7, 62169), (8, 62148), (9, 62145), (10, 62142), (11, 62061),

Gene: Wolfstar_3 Start: 1094, Stop: 897, Start Num: 5
Candidate Starts for Wolfstar_3:
(5, 1094), (6, 1073), (7, 1028), (8, 1007), (9, 1004), (10, 1001), (11, 920),