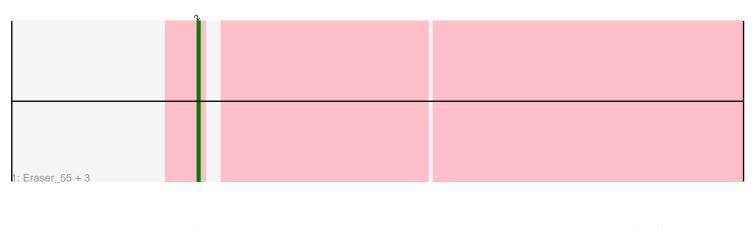
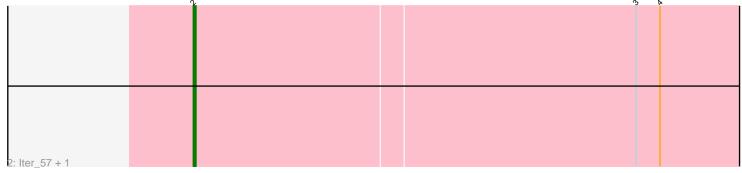
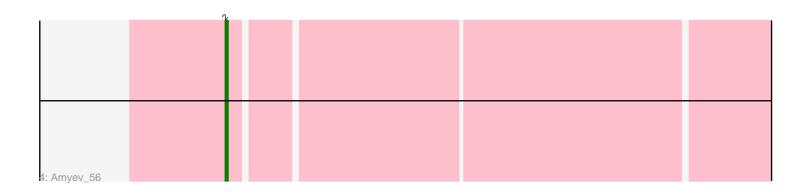
Pham 87784





	N	< l>	γ		
2	Fian_55				
Ο.	nan_55				



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 87784 Report

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

Pham number 87784 has 8 members, 1 are drafts.

Phages represented in each track:

- Track 1 : Eraser_55, Asa16_55, London_55, Niobe_55
- Track 2 : Iter_57, Ascela_57
- Track 3 : Tian_55
- Track 4 : Amyev_56

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

The start number called the most often in the published annotations is 2, it was called in 7 of the 7 non-draft genes in the pham.

Genes that call this "Most Annotated" start: • Amyev_56, Asa16_55, Ascela_57, Eraser_55, Iter_57, London_55, Niobe_55, Tian_55,

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

•

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

Summary by start number:

Start 2:

- Found in 8 of 8 (100.0%) of genes in pham
- Manual Annotations of this start: 7 of 7
- Called 100.0% of time when present

• Phage (with cluster) where this start called: Amyev_56 (AZ1), Asa16_55 (AZ1), Ascela_57 (AZ1), Eraser_55 (AZ1), Iter_57 (AZ1), London_55 (AZ1), Niobe_55 (AZ1), Tian_55 (AZ1),

Summary by clusters:

There is one cluster represented in this pham: AZ1

Info for manual annotations of cluster AZ1: •Start number 2 was manually annotated 7 times for cluster AZ1.

Gene Information:

Gene: Amyev_56 Start: 40609, Stop: 40833, Start Num: 2 Candidate Starts for Amyev_56: (Start: 2 @40609 has 7 MA's),

Gene: Asa16_55 Start: 39306, Stop: 39530, Start Num: 2 Candidate Starts for Asa16_55: (Start: 2 @39306 has 7 MA's),

Gene: Ascela_57 Start: 38762, Stop: 38992, Start Num: 2 Candidate Starts for Ascela_57: (Start: 2 @38762 has 7 MA's), (3, 38927), (4, 38936),

Gene: Eraser_55 Start: 39313, Stop: 39537, Start Num: 2 Candidate Starts for Eraser_55: (Start: 2 @39313 has 7 MA's),

Gene: Iter_57 Start: 38754, Stop: 38984, Start Num: 2 Candidate Starts for Iter_57: (Start: 2 @38754 has 7 MA's), (3, 38919), (4, 38928),

Gene: London_55 Start: 39304, Stop: 39528, Start Num: 2 Candidate Starts for London_55: (Start: 2 @39304 has 7 MA's),

Gene: Niobe_55 Start: 39307, Stop: 39531, Start Num: 2 Candidate Starts for Niobe_55: (Start: 2 @39307 has 7 MA's),

Gene: Tian_55 Start: 40608, Stop: 40832, Start Num: 2 Candidate Starts for Tian_55: (1, 40569), (Start: 2 @40608 has 7 MA's),