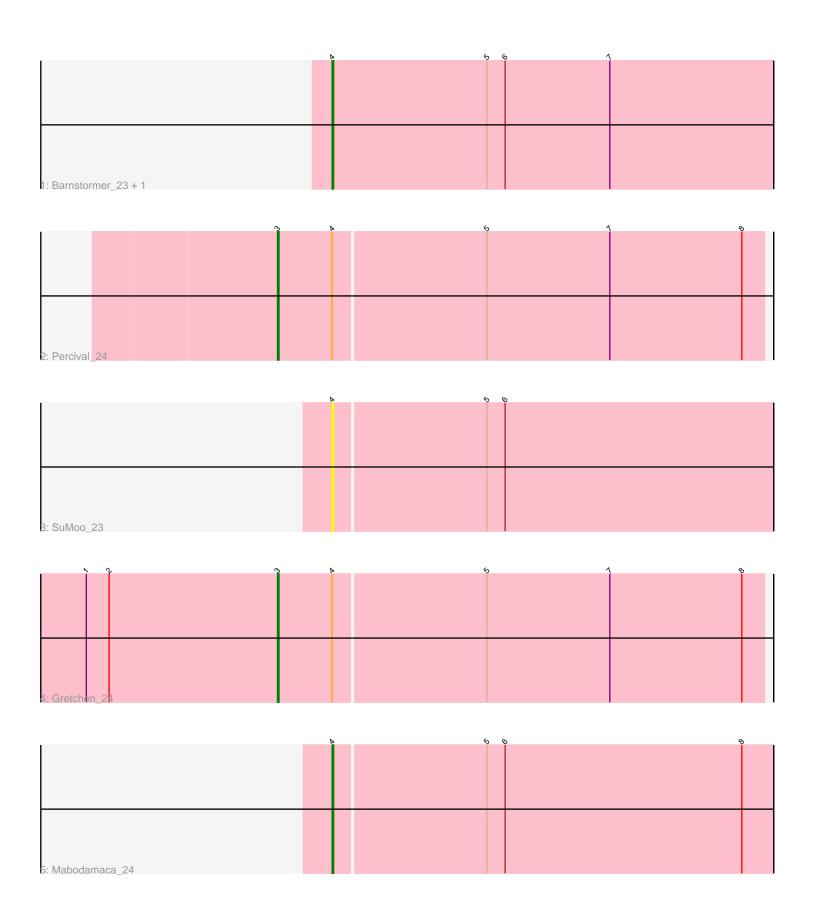
Pham 15412



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 15412 Report

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

Pham number 15412 has 6 members, 1 are drafts.

Phages represented in each track:

Track 1: Barnstormer_23, UtzChips_23

Track 2 : Percival_24
Track 3 : SuMoo_23
Track 4 : Gretchen_24
Track 5 : Mabodamaca 24

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 3 of the 5 non-draft genes in the pham.

Genes that call this "Most Annotated" start:

Barnstormer_23, Mabodamaca_24, SuMoo_23, UtzChips_23,

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

Gretchen_24, Percival_24,

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

Summary by start number:

Start 3:

- Found in 2 of 6 (33.3%) of genes in pham
- Manual Annotations of this start: 2 of 5
- Called 100.0% of time when present
- Phage (with cluster) where this start called: Gretchen_24 (EH), Percival_24 (EH),

Start 4:

- Found in 6 of 6 (100.0%) of genes in pham
- Manual Annotations of this start: 3 of 5
- Called 66.7% of time when present
- Phage (with cluster) where this start called: Barnstormer_23 (EH), Mabodamaca_24 (EH), SuMoo 23 (EH), UtzChips 23 (EH),

Summary by clusters:

There is one cluster represented in this pham: EH

Info for manual annotations of cluster EH:

- •Start number 3 was manually annotated 2 times for cluster EH.
- •Start number 4 was manually annotated 3 times for cluster EH.

Gene Information:

Gene: Barnstormer_23 Start: 19301, Stop: 19591, Start Num: 4

Candidate Starts for Barnstormer_23:

(Start: 4 @ 19301 has 3 MA's), (5, 19403), (6, 19415), (7, 19484),

Gene: Gretchen_24 Start: 20393, Stop: 20710, Start Num: 3

Candidate Starts for Gretchen_24:

(1, 20267), (2, 20282), (Start: 3 @20393 has 2 MA's), (Start: 4 @20429 has 3 MA's), (5, 20528), (7,

20609), (8, 20696),

Gene: Mabodamaca_24 Start: 19658, Stop: 19945, Start Num: 4

Candidate Starts for Mabodamaca 24:

(Start: 4 @ 19658 has 3 MA's), (5, 19757), (6, 19769), (8, 19925),

Gene: Percival_24 Start: 20225, Stop: 20542, Start Num: 3

Candidate Starts for Percival_24:

(Start: 3 @20225 has 2 MA's), (Start: 4 @20261 has 3 MA's), (5, 20360), (7, 20441), (8, 20528),

Gene: SuMoo_23 Start: 18614, Stop: 18901, Start Num: 4

Candidate Starts for SuMoo 23:

(Start: 4 @18614 has 3 MA's), (5, 18713), (6, 18725),

Gene: UtzChips 23 Start: 19287, Stop: 19577, Start Num: 4

Candidate Starts for UtzChips_23:

(Start: 4 @ 19287 has 3 MA's), (5, 19389), (6, 19401), (7, 19470),