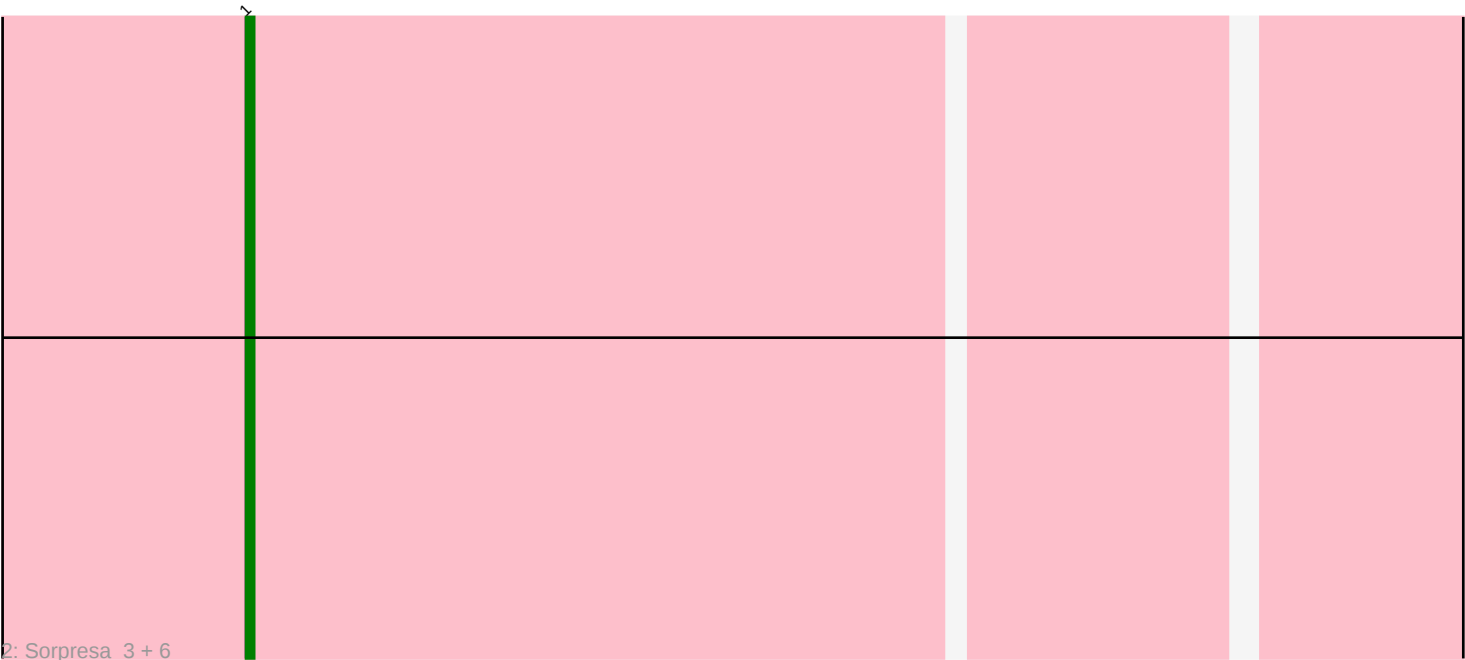




Note: Tracks are now grouped by subcluster and scaled. Switching in subcluster is indicated by changes in track color. Track scale is now set by default to display the region 30 bp upstream of start 1 to 30 bp downstream of the last possible start. If this default region is judged to be packed too tightly 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 323903 Report

This analysis was run 09/05/26 on database version 662.

Pham number 323903 has 10 members, 0 are drafts.

Phages represented in each track:

- Track 1 : Froghopper_4, Buttons_4, Manatee_4
- Track 2 : Sorpresa_3, Hermia_4, Pinto_4, SpikeBT_4, Noxious_4, AFIS_4, GageAP_5

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 10 of the 10 non-draft genes in the pham.

Genes that call this "Most Annotated" start:

- AFIS_4, Buttons_4, Froghopper_4, GageAP_5, Hermia_4, Manatee_4, Noxious_4, Pinto_4, Sorpresa_3, SpikeBT_4,

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 1:

- Found in 10 of 10 (100.0%) of genes in pham
- Manual Annotations of this start: 10 of 10
- Called 100.0% of time when present
- Phage (with cluster) where this start called: AFIS_4 (A1), Buttons_4 (A1), Froghopper_4 (A1), GageAP_5 (A1), Hermia_4 (A1), Manatee_4 (A1), Noxious_4 (A1), Pinto_4 (A1), Sorpresa_3 (A1), SpikeBT_4 (A1),

Summary by clusters:

There is one cluster represented in this pham: A1

Info for manual annotations of cluster A1:

- Start number 1 was manually annotated 10 times for cluster A1.

Gene Information:

Gene: AFIS_4 Start: 1662, Stop: 1808, Start Num: 1

Candidate Starts for AFIS_4:

(Start: 1 @1662 has 10 MA's),

Gene: Buttons_4 Start: 1561, Stop: 1740, Start Num: 1

Candidate Starts for Buttons_4:

(Start: 1 @1561 has 10 MA's), (2, 1681),

Gene: Froghopper_4 Start: 1562, Stop: 1741, Start Num: 1

Candidate Starts for Froghopper_4:

(Start: 1 @1562 has 10 MA's), (2, 1682),

Gene: GageAP_5 Start: 2005, Stop: 2151, Start Num: 1

Candidate Starts for GageAP_5:

(Start: 1 @2005 has 10 MA's),

Gene: Hermia_4 Start: 1563, Stop: 1709, Start Num: 1

Candidate Starts for Hermia_4:

(Start: 1 @1563 has 10 MA's),

Gene: Manatee_4 Start: 1566, Stop: 1745, Start Num: 1

Candidate Starts for Manatee_4:

(Start: 1 @1566 has 10 MA's), (2, 1686),

Gene: Noxious_4 Start: 1563, Stop: 1709, Start Num: 1

Candidate Starts for Noxious_4:

(Start: 1 @1563 has 10 MA's),

Gene: Pinto_4 Start: 1618, Stop: 1764, Start Num: 1

Candidate Starts for Pinto_4:

(Start: 1 @1618 has 10 MA's),

Gene: Sorpresa_3 Start: 1412, Stop: 1558, Start Num: 1

Candidate Starts for Sorpresa_3:

(Start: 1 @1412 has 10 MA's),

Gene: SpikeBT_4 Start: 1615, Stop: 1761, Start Num: 1

Candidate Starts for SpikeBT_4:

(Start: 1 @1615 has 10 MA's),