

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 106893 Report

This analysis was run 04/05/24 on database version 557.

Pham number 106893 has 9 members, 1 are drafts.

Phages represented in each track:

• Track 1 : Redno2 230

Track 2 : Ejimix_230, Dove_232

• Track 3 : StevieRay_55, Phegasus_55, Bunnies_56, Majeke_55, Mangethe_55, Zilizebeth 55

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 8 of the 8 non-draft genes in the pham.

Genes that call this "Most Annotated" start:

• Bunnies_56, Dove_232, Ejimix_230, Majeke_55, Mangethe_55, Phegasus_55, Redno2_230, StevieRay_55, Zilizebeth_55,

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

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

Summary by start number:

Start 2:

- Found in 9 of 9 (100.0%) of genes in pham
- Manual Annotations of this start: 8 of 8
- Called 100.0% of time when present
- Phage (with cluster) where this start called: Bunnies_56 (P1), Dove_232 (J),
 Ejimix_230 (J), Majeke_55 (P1), Mangethe_55 (P1), Phegasus_55 (P1), Redno2_230 (J), StevieRay_55 (P1), Zilizebeth_55 (P1),

Summary by clusters:

There are 2 clusters represented in this pham: J, P1,

Info for manual annotations of cluster J:

•Start number 2 was manually annotated 2 times for cluster J.

Info for manual annotations of cluster P1:

•Start number 2 was manually annotated 6 times for cluster P1.

Gene Information:

Gene: Bunnies 56 Start: 37483, Stop: 37310, Start Num: 2

Candidate Starts for Bunnies 56:

(1, 37525), (Start: 2 @37483 has 8 MA's), (3, 37471), (5, 37435), (6, 37426), (7, 37417), (8, 37393), (11, 37330),

Gene: Dove_232 Start: 107312, Stop: 107151, Start Num: 2

Candidate Starts for Dove 232:

(Start: 2 @107312 has 8 MA's), (8, 107225), (9, 107222), (10, 107201),

Gene: Ejimix_230 Start: 110258, Stop: 110097, Start Num: 2

Candidate Starts for Ejimix 230:

(Start: 2 @110258 has 8 MA's), (8, 110171), (9, 110168), (10, 110147),

Gene: Majeke 55 Start: 37272, Stop: 37099, Start Num: 2

Candidate Starts for Majeke_55:

(1, 37314), (Start: 2 @37272 has 8 MA's), (3, 37260), (5, 37224), (6, 37215), (7, 37206), (8, 37182), (11, 37119),

Gene: Mangethe_55 Start: 37272, Stop: 37099, Start Num: 2

Candidate Starts for Mangethe_55:

(1, 37314), (Start: 2 @37272 has 8 MA's), (3, 37260), (5, 37224), (6, 37215), (7, 37206), (8, 37182), (11, 37119),

Gene: Phegasus 55 Start: 37239, Stop: 37066, Start Num: 2

Candidate Starts for Phegasus 55:

(1, 37281), (Start: 2 @37239 has 8 MA's), (3, 37227), (5, 37191), (6, 37182), (7, 37173), (8, 37149), (11, 37086),

Gene: Redno2 230 Start: 107309, Stop: 107145, Start Num: 2

Candidate Starts for Redno2 230:

(Start: 2 @ 107309 has 8 MA's), (4, 107261), (8, 107222), (9, 107219), (10, 107198), (12, 107153),

Gene: StevieRay 55 Start: 37435, Stop: 37262, Start Num: 2

Candidate Starts for StevieRay_55:

(1, 37477), (Start: 2 @37435 has 8 MA's), (3, 37423), (5, 37387), (6, 37378), (7, 37369), (8, 37345), (11, 37282),

Gene: Zilizebeth_55 Start: 37263, Stop: 37090, Start Num: 2

Candidate Starts for Zilizebeth 55:

(1, 37305), (Start: 2 @37263 has 8 MA's), (3, 37251), (5, 37215), (6, 37206), (7, 37197), (8, 37173), (11, 37110),