

1: Annalisa 1 + 9

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 87728 Report

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

Pham number 87728 has 10 members, 1 are drafts.

Phages represented in each track:

- Track 1 : Annalisa_1, MichaelScott_1, Easley_1, Clark_1, WinkNick_1, Beenie_1, Samman98_1, Thimann_1, Dolores_1, Sekhmet_1

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 9 of the 9 non-draft genes in the pham.

Genes that call this "Most Annotated" start:

- Annalisa_1, Beenie_1, Clark_1, Dolores_1, Easley_1, MichaelScott_1, Samman98_1, Sekhmet_1, Thimann_1, WinkNick_1,

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: 9 of 9
- Called 100.0% of time when present
- Phage (with cluster) where this start called: Annalisa_1 (CZ4), Beenie_1 (CZ4), Clark_1 (CZ4), Dolores_1 (CZ4), Easley_1 (CZ4), MichaelScott_1 (CZ4), Samman98_1 (CZ4), Sekhmet_1 (CZ4), Thimann_1 (CZ4), WinkNick_1 (CZ4),

Summary by clusters:

There is one cluster represented in this pham: CZ4

Info for manual annotations of cluster CZ4:

- Start number 1 was manually annotated 9 times for cluster CZ4.

Gene Information:

Gene: Annalisa_1 Start: 47, Stop: 517, Start Num: 1

Candidate Starts for Annalisa_1:

(Start: 1 @47 has 9 MA's), (2, 272), (3, 302), (4, 407), (5, 470),

Gene: Beenie_1 Start: 47, Stop: 517, Start Num: 1

Candidate Starts for Beenie_1:

(Start: 1 @47 has 9 MA's), (2, 272), (3, 302), (4, 407), (5, 470),

Gene: Clark_1 Start: 47, Stop: 517, Start Num: 1

Candidate Starts for Clark_1:

(Start: 1 @47 has 9 MA's), (2, 272), (3, 302), (4, 407), (5, 470),

Gene: Dolores_1 Start: 47, Stop: 517, Start Num: 1

Candidate Starts for Dolores_1:

(Start: 1 @47 has 9 MA's), (2, 272), (3, 302), (4, 407), (5, 470),

Gene: Easley_1 Start: 47, Stop: 517, Start Num: 1

Candidate Starts for Easley_1:

(Start: 1 @47 has 9 MA's), (2, 272), (3, 302), (4, 407), (5, 470),

Gene: MichaelScott_1 Start: 47, Stop: 517, Start Num: 1

Candidate Starts for MichaelScott_1:

(Start: 1 @47 has 9 MA's), (2, 272), (3, 302), (4, 407), (5, 470),

Gene: Samman98_1 Start: 47, Stop: 517, Start Num: 1

Candidate Starts for Samman98_1:

(Start: 1 @47 has 9 MA's), (2, 272), (3, 302), (4, 407), (5, 470),

Gene: Sekhmet_1 Start: 47, Stop: 517, Start Num: 1

Candidate Starts for Sekhmet_1:

(Start: 1 @47 has 9 MA's), (2, 272), (3, 302), (4, 407), (5, 470),

Gene: Thimann_1 Start: 47, Stop: 517, Start Num: 1

Candidate Starts for Thimann_1:

(Start: 1 @47 has 9 MA's), (2, 272), (3, 302), (4, 407), (5, 470),

Gene: WinkNick_1 Start: 47, Stop: 517, Start Num: 1

Candidate Starts for WinkNick_1:

(Start: 1 @47 has 9 MA's), (2, 272), (3, 302), (4, 407), (5, 470),