



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 5175 Report

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

Pham number 5175 has 11 members, 0 are drafts.

Phages represented in each track:

- Track 1 : BigCheese_110, Winky_111, Breezona_111, Faith1_109, Netyap_109, Itos_111, Gabriela_111, Miley16_111, DrSeegs_109, BobsGarage_110, Zakai_112

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 11 of the 11 non-draft genes in the pham.

Genes that call this "Most Annotated" start:

- BigCheese_110, BobsGarage_110, Breezona_111, DrSeegs_109, Faith1_109, Gabriela_111, Itos_111, Miley16_111, Netyap_109, Winky_111, Zakai_112,

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 11 of 11 (100.0%) of genes in pham
- Manual Annotations of this start: 11 of 11
- Called 100.0% of time when present
- Phage (with cluster) where this start called: BigCheese_110 (L2), BobsGarage_110 (L2), Breezona_111 (L2), DrSeegs_109 (L2), Faith1_109 (L2), Gabriela_111 (L2), Itos_111 (L2), Miley16_111 (L2), Netyap_109 (L2), Winky_111 (L2), Zakai_112 (L2),

Summary by clusters:

There is one cluster represented in this pham: L2

Info for manual annotations of cluster L2:

- Start number 1 was manually annotated 11 times for cluster L2.

Gene Information:

Gene: BigCheese_110 Start: 64030, Stop: 64161, Start Num: 1

Candidate Starts for BigCheese_110:

(Start: 1 @64030 has 11 MA's), (2, 64087), (3, 64135),

Gene: BobsGarage_110 Start: 64030, Stop: 64161, Start Num: 1

Candidate Starts for BobsGarage_110:

(Start: 1 @64030 has 11 MA's), (2, 64087), (3, 64135),

Gene: Breezona_111 Start: 64365, Stop: 64496, Start Num: 1

Candidate Starts for Breezona_111:

(Start: 1 @64365 has 11 MA's), (2, 64422), (3, 64470),

Gene: DrSeegs_109 Start: 64365, Stop: 64496, Start Num: 1

Candidate Starts for DrSeegs_109:

(Start: 1 @64365 has 11 MA's), (2, 64422), (3, 64470),

Gene: Faith1_109 Start: 63673, Stop: 63804, Start Num: 1

Candidate Starts for Faith1_109:

(Start: 1 @63673 has 11 MA's), (2, 63730), (3, 63778),

Gene: Gabriela_111 Start: 63019, Stop: 63150, Start Num: 1

Candidate Starts for Gabriela_111:

(Start: 1 @63019 has 11 MA's), (2, 63076), (3, 63124),

Gene: Itos_111 Start: 62579, Stop: 62710, Start Num: 1

Candidate Starts for Itos_111:

(Start: 1 @62579 has 11 MA's), (2, 62636), (3, 62684),

Gene: Miley16_111 Start: 64365, Stop: 64496, Start Num: 1

Candidate Starts for Miley16_111:

(Start: 1 @64365 has 11 MA's), (2, 64422), (3, 64470),

Gene: Netyap_109 Start: 64079, Stop: 64210, Start Num: 1

Candidate Starts for Netyap_109:

(Start: 1 @64079 has 11 MA's), (2, 64136), (3, 64184),

Gene: Winky_111 Start: 64366, Stop: 64497, Start Num: 1

Candidate Starts for Winky_111:

(Start: 1 @64366 has 11 MA's), (2, 64423), (3, 64471),

Gene: Zakai_112 Start: 64076, Stop: 64207, Start Num: 1

Candidate Starts for Zakai_112:

(Start: 1 @64076 has 11 MA's), (2, 64133), (3, 64181),