



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 87628 Report

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

Pham number 87628 has 9 members, 0 are drafts.

Phages represented in each track:

- Track 1 : Madi_17, Sienna_17, Beyoncage_17, Terapin_17, BiteSize_17, Djokovic_17
- Track 2 : LilyPad_18
- Track 3 : Suzy_18
- Track 4 : LittleFella_19

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

Genes that call this "Most Annotated" start:

- Beyoncage_17, BiteSize_17, Djokovic_17, Madi_17, Sienna_17, Suzy_18, Terapin_17,

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:

- LilyPad_18, LittleFella_19,

Summary by start number:

Start 2:

- Found in 7 of 9 (77.8%) of genes in pham
- Manual Annotations of this start: 7 of 9
- Called 100.0% of time when present
- Phage (with cluster) where this start called: Beyoncage_17 (DG1), BiteSize_17 (DG1), Djokovic_17 (DG1), Madi_17 (DG1), Sienna_17 (DG1), Suzy_18 (DG1), Terapin_17 (DG1),

Start 3:

- Found in 1 of 9 (11.1%) of genes in pham
- Manual Annotations of this start: 1 of 9

- Called 100.0% of time when present
- Phage (with cluster) where this start called: LittleFella_19 (DG2),

Start 4:

- Found in 1 of 9 (11.1%) of genes in pham
- Manual Annotations of this start: 1 of 9
- Called 100.0% of time when present
- Phage (with cluster) where this start called: LilyPad_18 (DG1),

Summary by clusters:

There are 2 clusters represented in this pham: DG2, DG1,

Info for manual annotations of cluster DG1:

- Start number 2 was manually annotated 7 times for cluster DG1.
- Start number 4 was manually annotated 1 time for cluster DG1.

Info for manual annotations of cluster DG2:

- Start number 3 was manually annotated 1 time for cluster DG2.

Gene Information:

Gene: Beyoncage_17 Start: 10309, Stop: 10662, Start Num: 2

Candidate Starts for Beyoncage_17:

(1, 10306), (Start: 2 @10309 has 7 MA's), (6, 10441), (10, 10594),

Gene: BiteSize_17 Start: 10309, Stop: 10662, Start Num: 2

Candidate Starts for BiteSize_17:

(1, 10306), (Start: 2 @10309 has 7 MA's), (6, 10441), (10, 10594),

Gene: Djokovic_17 Start: 10309, Stop: 10662, Start Num: 2

Candidate Starts for Djokovic_17:

(1, 10306), (Start: 2 @10309 has 7 MA's), (6, 10441), (10, 10594),

Gene: LilyPad_18 Start: 11024, Stop: 11326, Start Num: 4

Candidate Starts for LilyPad_18:

(Start: 4 @11024 has 1 MA's), (8, 11129), (10, 11258),

Gene: LittleFella_19 Start: 12893, Stop: 13201, Start Num: 3

Candidate Starts for LittleFella_19:

(Start: 3 @12893 has 1 MA's), (5, 12953), (7, 12986), (9, 13091), (10, 13133), (11, 13151),

Gene: Madi_17 Start: 10309, Stop: 10662, Start Num: 2

Candidate Starts for Madi_17:

(1, 10306), (Start: 2 @10309 has 7 MA's), (6, 10441), (10, 10594),

Gene: Sienna_17 Start: 10309, Stop: 10662, Start Num: 2

Candidate Starts for Sienna_17:

(1, 10306), (Start: 2 @10309 has 7 MA's), (6, 10441), (10, 10594),

Gene: Suzy_18 Start: 11536, Stop: 11889, Start Num: 2

Candidate Starts for Suzy_18:

(Start: 2 @11536 has 7 MA's), (10, 11821),

Gene: Terapin_17 Start: 10309, Stop: 10662, Start Num: 2

Candidate Starts for Terapin_17:

(1, 10306), (Start: 2 @10309 has 7 MA's), (6, 10441), (10, 10594),