



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 106839 Report

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

Pham number 106839 has 10 members, 0 are drafts.

Phages represented in each track:

- Track 1 : ILeeKay\_81, Eyeball\_84, BaconJack\_86, Briton15\_86, Seabiscuit\_84, McGuire\_83, Sibs6\_85, Doom\_76, STLscum\_91
- Track 2 : BeesKnees\_84

### ***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:

- BaconJack\_86, BeesKnees\_84, Briton15\_86, Doom\_76, Eyeball\_84, ILeeKay\_81, McGuire\_83, STLscum\_91, Seabiscuit\_84, Sibs6\_85,

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: BaconJack\_86 (A1), BeesKnees\_84 (A1), Briton15\_86 (A1), Doom\_76 (A1), Eyeball\_84 (A1), ILeeKay\_81 (A1), McGuire\_83 (A1), STLscum\_91 (A1), Seabiscuit\_84 (A1), Sibs6\_85 (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: BaconJack\_86 Start: 50344, Stop: 50150, Start Num: 1

Candidate Starts for BaconJack\_86:

(Start: 1 @50344 has 10 MA's), (2, 50332), (3, 50239), (4, 50227),

Gene: BeesKnees\_84 Start: 47280, Stop: 47086, Start Num: 1

Candidate Starts for BeesKnees\_84:

(Start: 1 @47280 has 10 MA's), (2, 47268), (3, 47175), (4, 47163), (5, 47094),

Gene: Briton15\_86 Start: 49423, Stop: 49229, Start Num: 1

Candidate Starts for Briton15\_86:

(Start: 1 @49423 has 10 MA's), (2, 49411), (3, 49318), (4, 49306),

Gene: Doom\_76 Start: 47395, Stop: 47201, Start Num: 1

Candidate Starts for Doom\_76:

(Start: 1 @47395 has 10 MA's), (2, 47383), (3, 47290), (4, 47278),

Gene: Eyeball\_84 Start: 48830, Stop: 48636, Start Num: 1

Candidate Starts for Eyeball\_84:

(Start: 1 @48830 has 10 MA's), (2, 48818), (3, 48725), (4, 48713),

Gene: ILeeKay\_81 Start: 46786, Stop: 46592, Start Num: 1

Candidate Starts for ILeeKay\_81:

(Start: 1 @46786 has 10 MA's), (2, 46774), (3, 46681), (4, 46669),

Gene: McGuire\_83 Start: 47428, Stop: 47234, Start Num: 1

Candidate Starts for McGuire\_83:

(Start: 1 @47428 has 10 MA's), (2, 47416), (3, 47323), (4, 47311),

Gene: STLscum\_91 Start: 49511, Stop: 49317, Start Num: 1

Candidate Starts for STLscum\_91:

(Start: 1 @49511 has 10 MA's), (2, 49499), (3, 49406), (4, 49394),

Gene: Seabiscuit\_84 Start: 47662, Stop: 47468, Start Num: 1

Candidate Starts for Seabiscuit\_84:

(Start: 1 @47662 has 10 MA's), (2, 47650), (3, 47557), (4, 47545),

Gene: Sibs6\_85 Start: 46259, Stop: 46065, Start Num: 1

Candidate Starts for Sibs6\_85:

(Start: 1 @46259 has 10 MA's), (2, 46247), (3, 46154), (4, 46142),