



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 304021 Report

This analysis was run 06/08/26 on database version 649.

Pham number 304021 has 6 members, 0 are drafts.

Phages represented in each track:

- Track 1 : IchabodCrane_116, MindFlayer_117, KentuckyRacer_122, Rikishi_119
- Track 2 : JimJam_122, Spilled_122

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 4 of the 6 non-draft genes in the pham.

Genes that call this "Most Annotated" start:

- IchabodCrane_116, KentuckyRacer_122, MindFlayer_117, Rikishi_119,

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

- JimJam_122, Spilled_122,

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

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Summary by start number:

Start 1:

- Found in 6 of 6 (100.0%) of genes in pham
- Manual Annotations of this start: 4 of 6
- Called 66.7% of time when present
- Phage (with cluster) where this start called: IchabodCrane_116 (BE2), KentuckyRacer_122 (BE2), MindFlayer_117 (BE2), Rikishi_119 (BE2),

Start 2:

- Found in 6 of 6 (100.0%) of genes in pham
- Manual Annotations of this start: 2 of 6
- Called 33.3% of time when present
- Phage (with cluster) where this start called: JimJam_122 (BE2), Spilled_122 (BE2),

Summary by clusters:

There is one cluster represented in this pham: BE2

Info for manual annotations of cluster BE2:

- Start number 1 was manually annotated 4 times for cluster BE2.
- Start number 2 was manually annotated 2 times for cluster BE2.

Gene Information:

Gene: IchabodCrane_116 Start: 77536, Stop: 77793, Start Num: 1

Candidate Starts for IchabodCrane_116:

(Start: 1 @77536 has 4 MA's), (Start: 2 @77593 has 2 MA's), (3, 77638), (4, 77683), (5, 77767),

Gene: JimJam_122 Start: 78415, Stop: 78615, Start Num: 2

Candidate Starts for JimJam_122:

(Start: 1 @78358 has 4 MA's), (Start: 2 @78415 has 2 MA's), (3, 78460), (4, 78505), (5, 78589),

Gene: KentuckyRacer_122 Start: 78222, Stop: 78479, Start Num: 1

Candidate Starts for KentuckyRacer_122:

(Start: 1 @78222 has 4 MA's), (Start: 2 @78279 has 2 MA's), (3, 78324), (4, 78369), (5, 78453),

Gene: MindFlayer_117 Start: 77443, Stop: 77700, Start Num: 1

Candidate Starts for MindFlayer_117:

(Start: 1 @77443 has 4 MA's), (Start: 2 @77500 has 2 MA's), (3, 77545), (4, 77590), (5, 77674),

Gene: Rikishi_119 Start: 77741, Stop: 77998, Start Num: 1

Candidate Starts for Rikishi_119:

(Start: 1 @77741 has 4 MA's), (Start: 2 @77798 has 2 MA's), (3, 77843), (4, 77888), (5, 77972),

Gene: Spilled_122 Start: 77964, Stop: 78164, Start Num: 2

Candidate Starts for Spilled_122:

(Start: 1 @77907 has 4 MA's), (Start: 2 @77964 has 2 MA's), (3, 78009), (4, 78054), (5, 78138),