



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 305612 Report

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

Pham number 305612 has 5 members, 2 are drafts.

Phages represented in each track:

- Track 1 : DreamEater_38
- Track 2 : TripleJ_42, Kukla_41, GUPitcher_39, PetriParty_43

Summary of Final Annotations (See graph section above for start numbers):

The start number called the most often in the published annotations is 6, it was called in 2 of the 3 non-draft genes in the pham.

Genes that call this "Most Annotated" start:

- GUPitcher_39, Kukla_41, PetriParty_43, TripleJ_42,

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

- DreamEater_38,

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

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Summary by start number:

Start 4:

- Found in 5 of 5 (100.0%) of genes in pham
- Manual Annotations of this start: 1 of 3
- Called 20.0% of time when present
- Phage (with cluster) where this start called: DreamEater_38 (FJ),

Start 6:

- Found in 5 of 5 (100.0%) of genes in pham
- Manual Annotations of this start: 2 of 3
- Called 80.0% of time when present
- Phage (with cluster) where this start called: GUPitcher_39 (FJ), Kukla_41 (FJ), PetriParty_43 (FJ), TripleJ_42 (FJ),

Summary by clusters:

There is one cluster represented in this pham: FJ

Info for manual annotations of cluster FJ:

- Start number 4 was manually annotated 1 time for cluster FJ.
- Start number 6 was manually annotated 2 times for cluster FJ.

Gene Information:

Gene: DreamEater_38 Start: 30245, Stop: 30595, Start Num: 4

Candidate Starts for DreamEater_38:

(1, 30191), (2, 30209), (3, 30218), (Start: 4 @30245 has 1 MA's), (5, 30251), (Start: 6 @30326 has 2 MA's), (7, 30449), (8, 30542), (9, 30551),

Gene: GUPitcher_39 Start: 30557, Stop: 30826, Start Num: 6

Candidate Starts for GUPitcher_39:

(1, 30422), (2, 30440), (3, 30449), (Start: 4 @30476 has 1 MA's), (5, 30482), (Start: 6 @30557 has 2 MA's), (7, 30680), (8, 30773), (9, 30782),

Gene: Kukla_41 Start: 30045, Stop: 30314, Start Num: 6

Candidate Starts for Kukla_41:

(1, 29910), (2, 29928), (3, 29937), (Start: 4 @29964 has 1 MA's), (5, 29970), (Start: 6 @30045 has 2 MA's), (7, 30168), (8, 30261), (9, 30270),

Gene: PetriParty_43 Start: 30282, Stop: 30551, Start Num: 6

Candidate Starts for PetriParty_43:

(1, 30147), (2, 30165), (3, 30174), (Start: 4 @30201 has 1 MA's), (5, 30207), (Start: 6 @30282 has 2 MA's), (7, 30405), (8, 30498), (9, 30507),

Gene: TripleJ_42 Start: 29886, Stop: 30155, Start Num: 6

Candidate Starts for TripleJ_42:

(1, 29751), (2, 29769), (3, 29778), (Start: 4 @29805 has 1 MA's), (5, 29811), (Start: 6 @29886 has 2 MA's), (7, 30009), (8, 30102), (9, 30111),