



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 284659 Report

This analysis was run 02/23/26 on database version 636.

Pham number 284659 has 4 members, 1 are drafts.

Phages represented in each track:

- Track 1 : Hereford_50, EvenBluerMoon_33, Aoka_34
- Track 2 : PrairieDogTown_34

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 2 of the 3 non-draft genes in the pham.

Genes that call this "Most Annotated" start:

- Aoka_34, EvenBluerMoon_33, Hereford_50,

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

- PrairieDogTown_34,

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

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Summary by start number:

Start 1:

- Found in 4 of 4 (100.0%) of genes in pham
- Manual Annotations of this start: 1 of 3
- Called 25.0% of time when present
- Phage (with cluster) where this start called: PrairieDogTown_34 (FO),

Start 2:

- Found in 4 of 4 (100.0%) of genes in pham
- Manual Annotations of this start: 2 of 3
- Called 75.0% of time when present
- Phage (with cluster) where this start called: Aoka_34 (FO), EvenBluerMoon_33 (FO), Hereford_50 (FO),

Summary by clusters:

There is one cluster represented in this pham: FO

Info for manual annotations of cluster FO:

- Start number 1 was manually annotated 1 time for cluster FO.
- Start number 2 was manually annotated 2 times for cluster FO.

Gene Information:

Gene: Aoka_34 Start: 26615, Stop: 27118, Start Num: 2

Candidate Starts for Aoka_34:

(Start: 1 @26579 has 1 MA's), (Start: 2 @26615 has 2 MA's),

Gene: EvenBluerMoon_33 Start: 26853, Stop: 27356, Start Num: 2

Candidate Starts for EvenBluerMoon_33:

(Start: 1 @26817 has 1 MA's), (Start: 2 @26853 has 2 MA's),

Gene: Hereford_50 Start: 26687, Stop: 27190, Start Num: 2

Candidate Starts for Hereford_50:

(Start: 1 @26651 has 1 MA's), (Start: 2 @26687 has 2 MA's),

Gene: PrairieDogTown_34 Start: 26819, Stop: 27358, Start Num: 1

Candidate Starts for PrairieDogTown_34:

(Start: 1 @26819 has 1 MA's), (Start: 2 @26855 has 2 MA's),