



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 107059 Report

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

Pham number 107059 has 7 members, 0 are drafts.

Phages represented in each track:

 Track 1: KaiHaiDragon_55, Quhwah_58, Teamocil_57, Piperis_55, Phorgeous_54, ClearAsMud_58

Track 2 : EarickHC_55

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

Genes that call this "Most Annotated" start:

 ClearAsMud_58, KaiHaiDragon_55, Phorgeous_54, Piperis_55, Quhwah_58, Teamocil_57,

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

EarickHC_55,

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

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Summary by start number:

Start 1:

- Found in 7 of 7 (100.0%) of genes in pham
- Manual Annotations of this start: 6 of 7
- Called 85.7% of time when present
- Phage (with cluster) where this start called: ClearAsMud_58 (EC), KaiHaiDragon_55 (EC), Phorgeous_54 (EC), Piperis_55 (EC), Quhwah_58 (EC), Teamocil_57 (EC),

Start 3:

- Found in 7 of 7 (100.0%) of genes in pham
- Manual Annotations of this start: 1 of 7
- Called 14.3% of time when present
- Phage (with cluster) where this start called: EarickHC_55 (EC),

Summary by clusters:

There is one cluster represented in this pham: EC

Info for manual annotations of cluster EC:

- •Start number 1 was manually annotated 6 times for cluster EC.
- •Start number 3 was manually annotated 1 time for cluster EC.

Gene Information:

Gene: ClearAsMud 58 Start: 36822, Stop: 37004, Start Num: 1

Candidate Starts for ClearAsMud_58:

(Start: 1 @36822 has 6 MA's), (2, 36843), (Start: 3 @36873 has 1 MA's),

Gene: EarickHC 55 Start: 36293, Stop: 36424, Start Num: 3

Candidate Starts for EarickHC 55:

(Start: 1 @36242 has 6 MA's), (2, 36263), (Start: 3 @36293 has 1 MA's),

Gene: KaiHaiDragon_55 Start: 36242, Stop: 36424, Start Num: 1

Candidate Starts for KaiHaiDragon_55:

(Start: 1 @36242 has 6 MA's), (2, 36263), (Start: 3 @36293 has 1 MA's),

Gene: Phorgeous_54 Start: 36171, Stop: 36353, Start Num: 1

Candidate Starts for Phorgeous 54:

(Start: 1 @ 36171 has 6 MA's), (2, 36192), (Start: 3 @ 36222 has 1 MA's),

Gene: Piperis_55 Start: 36294, Stop: 36476, Start Num: 1

Candidate Starts for Piperis_55:

(Start: 1 @36294 has 6 MA's), (2, 36315), (Start: 3 @36345 has 1 MA's),

Gene: Quhwah 58 Start: 36952, Stop: 37134, Start Num: 1

Candidate Starts for Quhwah 58:

(Start: 1 @ 36952 has 6 MA's), (2, 36973), (Start: 3 @ 37003 has 1 MA's),

Gene: Teamocil_57 Start: 36421, Stop: 36603, Start Num: 1

Candidate Starts for Teamocil_57:

(Start: 1 @36421 has 6 MA's), (2, 36442), (Start: 3 @36472 has 1 MA's),