



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 156872 Report

This analysis was run 04/12/24 on database version 558.

Pham number 156872 has 9 members, 0 are drafts.

Phages represented in each track:

- Track 1 : D29_43.1, D32_46, Naji_46, StarStuff_46
- Track 2 : Kerberos_46, DBQu4n_46, Tomathan_46
- Track 3 : Duplo_46, Pomar16_46

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

The start number called the most often in the published annotations is 3, it was called in 6 of the 9 non-draft genes in the pham.

Genes that call this "Most Annotated" start:

- D29_43.1, D32_46, Duplo_46, Naji_46, Pomar16_46, StarStuff_46,

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

- DBQu4n_46, Kerberos_46, Tomathan_46,

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

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Summary by start number:

Start 2:

- Found in 5 of 9 (55.6%) of genes in pham
- Manual Annotations of this start: 3 of 9
- Called 60.0% of time when present
- Phage (with cluster) where this start called: DBQu4n_46 (A2), Kerberos_46 (A2), Tomathan_46 (A2),

Start 3:

- Found in 9 of 9 (100.0%) of genes in pham
- Manual Annotations of this start: 6 of 9
- Called 66.7% of time when present
- Phage (with cluster) where this start called: D29_43.1 (A2), D32_46 (A2), Duplo_46 (A2), Naji_46 (A2), Pomar16_46 (A2), StarStuff_46 (A2),

Summary by clusters:

There is one cluster represented in this pham: A2

Info for manual annotations of cluster A2:

- Start number 2 was manually annotated 3 times for cluster A2.
- Start number 3 was manually annotated 6 times for cluster A2.

Gene Information:

Gene: D29_43.1 Start: 29319, Stop: 29179, Start Num: 3

Candidate Starts for D29_43.1:

(1, 29373), (Start: 3 @29319 has 6 MA's), (4, 29283), (5, 29265), (6, 29235), (7, 29226),

Gene: D32_46 Start: 29319, Stop: 29179, Start Num: 3

Candidate Starts for D32_46:

(1, 29373), (Start: 3 @29319 has 6 MA's), (4, 29283), (5, 29265), (6, 29235), (7, 29226),

Gene: DBQu4n_46 Start: 29346, Stop: 29179, Start Num: 2

Candidate Starts for DBQu4n_46:

(1, 29373), (Start: 2 @29346 has 3 MA's), (Start: 3 @29319 has 6 MA's), (4, 29283), (6, 29235), (7, 29226),

Gene: Duplo_46 Start: 29357, Stop: 29217, Start Num: 3

Candidate Starts for Duplo_46:

(1, 29411), (Start: 2 @29384 has 3 MA's), (Start: 3 @29357 has 6 MA's), (4, 29321), (6, 29273), (7, 29264),

Gene: Kerberos_46 Start: 29346, Stop: 29179, Start Num: 2

Candidate Starts for Kerberos_46:

(1, 29373), (Start: 2 @29346 has 3 MA's), (Start: 3 @29319 has 6 MA's), (4, 29283), (6, 29235), (7, 29226),

Gene: Naji_46 Start: 29319, Stop: 29179, Start Num: 3

Candidate Starts for Naji_46:

(1, 29373), (Start: 3 @29319 has 6 MA's), (4, 29283), (5, 29265), (6, 29235), (7, 29226),

Gene: Pomar16_46 Start: 29360, Stop: 29220, Start Num: 3

Candidate Starts for Pomar16_46:

(1, 29414), (Start: 2 @29387 has 3 MA's), (Start: 3 @29360 has 6 MA's), (4, 29324), (6, 29276), (7, 29267),

Gene: StarStuff_46 Start: 29322, Stop: 29182, Start Num: 3

Candidate Starts for StarStuff_46:

(1, 29376), (Start: 3 @29322 has 6 MA's), (4, 29286), (5, 29268), (6, 29238), (7, 29229),

Gene: Tomathan_46 Start: 29346, Stop: 29179, Start Num: 2

Candidate Starts for Tomathan_46:

(1, 29373), (Start: 2 @29346 has 3 MA's), (Start: 3 @29319 has 6 MA's), (4, 29283), (6, 29235), (7, 29226),