



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 7304 Report

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

Pham number 7304 has 6 members, 0 are drafts.

Phages represented in each track:

- Track 1 : Big4_34, Big4_360
- Track 2 : Zooman_31, Zooman_344
- Track 3 : Cece_337, Cece_35

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

Genes that call this "Most Annotated" start:

- Big4_34, Big4_360, Zooman_31, Zooman_344,

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

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

- Cece_337, Cece_35,

Summary by start number:

Start 2:

- Found in 2 of 6 (33.3%) of genes in pham
- Manual Annotations of this start: 2 of 6
- Called 100.0% of time when present
- Phage (with cluster) where this start called: Cece_337 (GD3), Cece_35 (GD3),

Start 3:

- Found in 4 of 6 (66.7%) of genes in pham
- Manual Annotations of this start: 4 of 6
- Called 100.0% of time when present
- Phage (with cluster) where this start called: Big4_34 (GD2), Big4_360 (GD2), Zooman_31 (GD2), Zooman_344 (GD2),

Summary by clusters:

There are 2 clusters represented in this pham: GD2, GD3,

Info for manual annotations of cluster GD2:

- Start number 3 was manually annotated 4 times for cluster GD2.

Info for manual annotations of cluster GD3:

- Start number 2 was manually annotated 2 times for cluster GD3.

Gene Information:

Gene: Big4_34 Start: 13506, Stop: 13796, Start Num: 3

Candidate Starts for Big4_34:

(Start: 3 @13506 has 4 MA's), (6, 13626), (7, 13668), (8, 13692), (9, 13698), (10, 13725),

Gene: Big4_360 Start: 188200, Stop: 188490, Start Num: 3

Candidate Starts for Big4_360:

(Start: 3 @188200 has 4 MA's), (6, 188320), (7, 188362), (8, 188386), (9, 188392), (10, 188419),

Gene: Cece_337 Start: 182048, Stop: 182347, Start Num: 2

Candidate Starts for Cece_337:

(1, 182033), (Start: 2 @182048 has 2 MA's), (4, 182102), (5, 182117), (7, 182219), (9, 182249), (12, 182318),

Gene: Cece_35 Start: 13614, Stop: 13913, Start Num: 2

Candidate Starts for Cece_35:

(1, 13599), (Start: 2 @13614 has 2 MA's), (4, 13668), (5, 13683), (7, 13785), (9, 13815), (12, 13884),

Gene: Zooman_31 Start: 12945, Stop: 13241, Start Num: 3

Candidate Starts for Zooman_31:

(Start: 3 @12945 has 4 MA's), (7, 13113), (8, 13137), (10, 13170), (11, 13200),

Gene: Zooman_344 Start: 188596, Stop: 188892, Start Num: 3

Candidate Starts for Zooman_344:

(Start: 3 @188596 has 4 MA's), (7, 188764), (8, 188788), (10, 188821), (11, 188851),