



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 194616 Report

This analysis was run 11/02/24 on database version 579.

Pham number 194616 has 8 members, 3 are drafts.

Phages represented in each track:

- Track 1 : HangryHippo_224, BlueOtter_224, Cursive_229, Scheme_229, Watermoore_225, Cross_225
- Track 2 : Gibbi_246
- Track 3 : Mugiwara_250

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 5 of the 5 non-draft genes in the pham.

Genes that call this "Most Annotated" start:

- BlueOtter_224, Cross_225, Cursive_229, HangryHippo_224, Mugiwara_250, Scheme_229, Watermoore_225,

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

- Gibbi_246,

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

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Summary by start number:

Start 2:

- Found in 2 of 8 (25.0%) of genes in pham
- No Manual Annotations of this start.
- Called 50.0% of time when present
- Phage (with cluster) where this start called: Gibbi_246 (BE2),

Start 3:

- Found in 8 of 8 (100.0%) of genes in pham
- Manual Annotations of this start: 5 of 5
- Called 87.5% of time when present
- Phage (with cluster) where this start called: BlueOtter_224 (BE1), Cross_225 (BE1), Cursive_229 (BE1), HangryHippo_224 (BE1), Mugiwara_250 (BE2), Scheme_229

(BE1), Watermoore_225 (BE1),

Summary by clusters:

There are 2 clusters represented in this pham: BE2, BE1,

Info for manual annotations of cluster BE1:

•Start number 3 was manually annotated 5 times for cluster BE1.

Gene Information:

Gene: BlueOtter_224 Start: 113515, Stop: 113667, Start Num: 3

Candidate Starts for BlueOtter_224:

(1, 113437), (Start: 3 @113515 has 5 MA's), (5, 113578), (7, 113659),

Gene: Cross_225 Start: 114160, Stop: 114312, Start Num: 3

Candidate Starts for Cross_225:

(1, 114082), (Start: 3 @114160 has 5 MA's), (5, 114223), (7, 114304),

Gene: Cursive_229 Start: 114267, Stop: 114419, Start Num: 3

Candidate Starts for Cursive_229:

(1, 114189), (Start: 3 @114267 has 5 MA's), (5, 114330), (7, 114411),

Gene: Gibbi_246 Start: 112069, Stop: 112251, Start Num: 2

Candidate Starts for Gibbi_246:

(1, 112021), (2, 112069), (Start: 3 @112099 has 5 MA's), (4, 112108), (6, 112165), (7, 112243),

Gene: HangryHippo_224 Start: 113515, Stop: 113667, Start Num: 3

Candidate Starts for HangryHippo_224:

(1, 113437), (Start: 3 @113515 has 5 MA's), (5, 113578), (7, 113659),

Gene: Mugiwara_250 Start: 114313, Stop: 114465, Start Num: 3

Candidate Starts for Mugiwara_250:

(1, 114235), (2, 114283), (Start: 3 @114313 has 5 MA's), (4, 114322), (6, 114379), (7, 114457),

Gene: Scheme_229 Start: 115451, Stop: 115603, Start Num: 3

Candidate Starts for Scheme_229:

(1, 115373), (Start: 3 @115451 has 5 MA's), (5, 115514), (7, 115595),

Gene: Watermoore_225 Start: 115158, Stop: 115310, Start Num: 3

Candidate Starts for Watermoore_225:

(1, 115080), (Start: 3 @115158 has 5 MA's), (5, 115221), (7, 115302),