

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 7775 Report

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

Pham number 7775 has 8 members, 0 are drafts.

Phages represented in each track:

Track 1 : SparkleGoddess_243

• Track 2 : Karp_241, Belfort_242

Track 3 : Francob_249, Gilson_245

Track 4 : Comrade_238Track 5 : Kenrey_251Track 6 : Stigma_241

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

The start number called the most often in the published annotations is 6, it was called in 8 of the 8 non-draft genes in the pham.

Genes that call this "Most Annotated" start:

• Belfort_242, Comrade_238, Francob_249, Gilson_245, Karp_241, Kenrey_251, SparkleGoddess_243, Stigma_241,

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:

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Summary by start number:

Start 6:

- Found in 8 of 8 (100.0%) of genes in pham
- Manual Annotations of this start: 8 of 8
- Called 100.0% of time when present
- Phage (with cluster) where this start called: Belfort_242 (BK1), Comrade_238 (BK1), Francob_249 (BK1), Gilson_245 (BK1), Karp_241 (BK1), Kenrey_251 (BK1), SparkleGoddess_243 (BK1), Stigma_241 (BK1),

Summary by clusters:

There is one cluster represented in this pham: BK1

Info for manual annotations of cluster BK1:

•Start number 6 was manually annotated 8 times for cluster BK1.

Gene Information:

Gene: Belfort 242 Start: 116771, Stop: 116968, Start Num: 6

Candidate Starts for Belfort_242:

(3, 116702), (4, 116708), (5, 116720), (Start: 6 @116771 has 8 MA's), (10, 116819), (13, 116855),

Gene: Comrade_238 Start: 117058, Stop: 117255, Start Num: 6

Candidate Starts for Comrade 238:

(5, 117007), (Start: 6 @117058 has 8 MA's), (10, 117106), (13, 117142),

Gene: Francob 249 Start: 117384, Stop: 117581, Start Num: 6

Candidate Starts for Francob_249:

(1, 117297), (2, 117300), (3, 117315), (4, 117321), (5, 117333), (Start: 6 @117384 has 8 MA's), (10, 117432), (12, 117453), (13, 117468),

Gene: Gilson_245 Start: 116649, Stop: 116846, Start Num: 6

Candidate Starts for Gilson_245:

(1, 116562), (2, 116565), (3, 116580), (4, 116586), (5, 116598), (Start: 6 @116649 has 8 MA's), (10, 116697), (12, 116718), (13, 116733),

Gene: Karp_241 Start: 118369, Stop: 118566, Start Num: 6

Candidate Starts for Karp_241:

(3, 118300), (4, 118306), (5, 118318), (Start: 6 @118369 has 8 MA's), (10, 118417), (13, 118453),

Gene: Kenrey_251 Start: 117597, Stop: 117794, Start Num: 6

Candidate Starts for Kenrey 251:

(Start: 6 @ 117597 has 8 MA's), (7, 117630), (8, 117633), (9, 117639), (10, 117645), (11, 117651),

Gene: SparkleGoddess_243 Start: 117643, Stop: 117840, Start Num: 6

Candidate Starts for SparkleGoddess_243:

(1, 117556), (2, 117559), (3, 117574), (4, 117580), (5, 117592), (Start: 6 @117643 has 8 MA's), (10, 117691), (13, 117727),

Gene: Stigma 241 Start: 117605, Stop: 117802, Start Num: 6

Candidate Starts for Stigma_241:

(4, 117542), (5, 117554), (Start: 6 @117605 has 8 MA's), (10, 117653), (13, 117689),