



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 161007 Report

This analysis was run 05/04/24 on database version 560.

Pham number 161007 has 7 members, 0 are drafts.

Phages represented in each track:

- Track 1 : TunaTartare_201
- Track 2 : Belfort_206, Karp_202, Comrade_203
- Track 3 : Sham_193
- Track 4 : SparkleGoddess_206, Stigma_205

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

The start number called the most often in the published annotations is 5, it was called in 5 of the 7 non-draft genes in the pham.

Genes that call this "Most Annotated" start:

- Belfort_206, Comrade_203, Karp_202, SparkleGoddess_206, Stigma_205,

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

-

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

- Sham_193, TunaTartare_201,

Summary by start number:

Start 4:

- Found in 2 of 7 (28.6%) of genes in pham
- Manual Annotations of this start: 2 of 7
- Called 100.0% of time when present
- Phage (with cluster) where this start called: Sham_193 (BK1), TunaTartare_201 (BK1),

Start 5:

- Found in 5 of 7 (71.4%) of genes in pham
- Manual Annotations of this start: 5 of 7
- Called 100.0% of time when present
- Phage (with cluster) where this start called: Belfort_206 (BK1), Comrade_203 (BK1), Karp_202 (BK1), SparkleGoddess_206 (BK1), Stigma_205 (BK1),

Summary by clusters:

There is one cluster represented in this pham: BK1

Info for manual annotations of cluster BK1:

- Start number 4 was manually annotated 2 times for cluster BK1.
- Start number 5 was manually annotated 5 times for cluster BK1.

Gene Information:

Gene: Belfort_206 Start: 103754, Stop: 104002, Start Num: 5

Candidate Starts for Belfort_206:

(3, 103730), (Start: 5 @103754 has 5 MA's), (6, 103784), (7, 103796), (8, 103826), (11, 103880), (13, 103925), (14, 103934),

Gene: Comrade_203 Start: 103955, Stop: 104203, Start Num: 5

Candidate Starts for Comrade_203:

(3, 103931), (Start: 5 @103955 has 5 MA's), (6, 103985), (7, 103997), (8, 104027), (11, 104081), (13, 104126), (14, 104135),

Gene: Karp_202 Start: 104121, Stop: 104369, Start Num: 5

Candidate Starts for Karp_202:

(3, 104097), (Start: 5 @104121 has 5 MA's), (6, 104151), (7, 104163), (8, 104193), (11, 104247), (13, 104292), (14, 104301),

Gene: Sham_193 Start: 102724, Stop: 102972, Start Num: 4

Candidate Starts for Sham_193:

(Start: 4 @102724 has 2 MA's), (8, 102796), (9, 102805), (10, 102841), (15, 102919), (16, 102922),

Gene: SparkleGoddess_206 Start: 104150, Stop: 104398, Start Num: 5

Candidate Starts for SparkleGoddess_206:

(1, 104054), (2, 104072), (3, 104126), (Start: 5 @104150 has 5 MA's), (6, 104180), (7, 104192), (8, 104222), (11, 104276), (13, 104321), (14, 104330),

Gene: Stigma_205 Start: 104388, Stop: 104636, Start Num: 5

Candidate Starts for Stigma_205:

(1, 104292), (2, 104310), (3, 104364), (Start: 5 @104388 has 5 MA's), (6, 104418), (7, 104430), (8, 104460), (11, 104514), (13, 104559), (14, 104568),

Gene: TunaTartare_201 Start: 104839, Stop: 105087, Start Num: 4

Candidate Starts for TunaTartare_201:

(Start: 4 @104839 has 2 MA's), (8, 104911), (9, 104920), (12, 104977), (16, 105037), (17, 105064),