



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 7708 Report

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

Pham number 7708 has 6 members, 0 are drafts.

Phages represented in each track:

- Track 1 : DmpstrDiver_181, BAKA_186, Minerva_180, Klein_184, Constella_176, Kalah2_177

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

The start number called the most often in the published annotations is 1, it was called in 6 of the 6 non-draft genes in the pham.

Genes that call this "Most Annotated" start:

- BAKA_186, Constella_176, DmpstrDiver_181, Kalah2_177, Klein_184, Minerva_180,

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

-

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

-

Summary by start number:

Start 1:

- Found in 6 of 6 (100.0%) of genes in pham
- Manual Annotations of this start: 6 of 6
- Called 100.0% of time when present
- Phage (with cluster) where this start called: BAKA_186 (J), Constella_176 (J), DmpstrDiver_181 (J), Kalah2_177 (J), Klein_184 (J), Minerva_180 (J),

Summary by clusters:

There is one cluster represented in this pham: J

Info for manual annotations of cluster J:

- Start number 1 was manually annotated 6 times for cluster J.

Gene Information:

Gene: BAKA_186 Start: 95077, Stop: 95187, Start Num: 1

Candidate Starts for BAKA_186:

(Start: 1 @95077 has 6 MA's), (2, 95107), (3, 95122),

Gene: Constella_176 Start: 92966, Stop: 93076, Start Num: 1

Candidate Starts for Constella_176:

(Start: 1 @92966 has 6 MA's), (2, 92996), (3, 93011),

Gene: DmpstrDiver_181 Start: 92899, Stop: 93009, Start Num: 1

Candidate Starts for DmpstrDiver_181:

(Start: 1 @92899 has 6 MA's), (2, 92929), (3, 92944),

Gene: Kalah2_177 Start: 94525, Stop: 94635, Start Num: 1

Candidate Starts for Kalah2_177:

(Start: 1 @94525 has 6 MA's), (2, 94555), (3, 94570),

Gene: Klein_184 Start: 92968, Stop: 93078, Start Num: 1

Candidate Starts for Klein_184:

(Start: 1 @92968 has 6 MA's), (2, 92998), (3, 93013),

Gene: Minerva_180 Start: 92605, Stop: 92715, Start Num: 1

Candidate Starts for Minerva_180:

(Start: 1 @92605 has 6 MA's), (2, 92635), (3, 92650),