



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 203665 Report

This analysis was run 03/07/25 on database version 590.

Pham number 203665 has 8 members, 7 are drafts.

Phages represented in each track:

- Track 1 : LeoJr_204, Atuin_195
- Track 2 : DunneganBoMo_201, WaddleDee_200, Ellewin_199, KSunshine22_198
- Track 3 : ReginaGlobina_207
- Track 4 : Panchaali_207

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 1 of the 1 non-draft genes in the pham.

Genes that call this "Most Annotated" start:

- Atuin_195, DunneganBoMo_201, Ellewin_199, KSunshine22_198, LeoJr_204, Panchaali_207, ReginaGlobina_207, WaddleDee_200,

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 8 of 8 (100.0%) of genes in pham
- Manual Annotations of this start: 1 of 1
- Called 100.0% of time when present
- Phage (with cluster) where this start called: Atuin_195 (FC), DunneganBoMo_201 (FC), Ellewin_199 (FC), KSunshine22_198 (FC), LeoJr_204 (FC), Panchaali_207 (FC), ReginaGlobina_207 (FC), WaddleDee_200 (FC),

Summary by clusters:

There is one cluster represented in this pham: FC

Info for manual annotations of cluster FC:

•Start number 1 was manually annotated 1 time for cluster FC.

Gene Information:

Gene: Atuin_195 Start: 138894, Stop: 139280, Start Num: 1

Candidate Starts for Atuin_195:

(Start: 1 @138894 has 1 MA's), (3, 138999), (4, 139050), (5, 139071), (6, 139074), (7, 139083),

Gene: DunneganBoMo_201 Start: 144050, Stop: 144415, Start Num: 1

Candidate Starts for DunneganBoMo_201:

(Start: 1 @144050 has 1 MA's), (2, 144077), (5, 144218),

Gene: Ellewin_199 Start: 143316, Stop: 143681, Start Num: 1

Candidate Starts for Ellewin_199:

(Start: 1 @143316 has 1 MA's), (2, 143343), (5, 143484),

Gene: KSunshine22_198 Start: 142727, Stop: 143092, Start Num: 1

Candidate Starts for KSunshine22_198:

(Start: 1 @142727 has 1 MA's), (2, 142754), (5, 142895),

Gene: LeoJr_204 Start: 139068, Stop: 139454, Start Num: 1

Candidate Starts for LeoJr_204:

(Start: 1 @139068 has 1 MA's), (3, 139173), (4, 139224), (5, 139245), (6, 139248), (7, 139257),

Gene: Panchaali_207 Start: 144812, Stop: 145174, Start Num: 1

Candidate Starts for Panchaali_207:

(Start: 1 @144812 has 1 MA's), (5, 144977), (8, 145109),

Gene: ReginaGlobina_207 Start: 140351, Stop: 140737, Start Num: 1

Candidate Starts for ReginaGlobina_207:

(Start: 1 @140351 has 1 MA's), (3, 140456), (4, 140507), (5, 140528), (6, 140531), (7, 140540),

Gene: WaddleDee_200 Start: 142511, Stop: 142876, Start Num: 1

Candidate Starts for WaddleDee_200:

(Start: 1 @142511 has 1 MA's), (2, 142538), (5, 142679),