



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 189270 Report

This analysis was run 01/18/25 on database version 583.

Pham number 189270 has 2 members, 1 are drafts.

Phages represented in each track:

Track 1 : Nova53_213Track 2 : Dorin_199

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

The start number called the most often in the published annotations is 12, it was called in 1 of the 1 non-draft genes in the pham.

Genes that call this "Most Annotated" start:

Dorin_199, Nova53_213,

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 12:

- Found in 2 of 2 (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: Dorin 199 (CG), Nova53 213 (CG).

Summary by clusters:

There is one cluster represented in this pham: CG

Info for manual annotations of cluster CG:

Start number 12 was manually annotated 1 time for cluster CG.

Gene Information:

Gene: Dorin_199 Start: 109238, Stop: 109510, Start Num: 12

Candidate Starts for Dorin_199:

(Start: 12 @109238 has 1 MA's), (13, 109352), (14, 109358), (15, 109412), (16, 109418), (18, 109445), (19, 109469),

Gene: Nova53_213 Start: 114115, Stop: 114387, Start Num: 12

Candidate Starts for Nova53_213:

(1, 113917), (2, 113920), (3, 113953), (4, 113959), (5, 113995), (6, 114001), (7, 114007), (8, 114016), (9, 114028), (10, 114043), (11, 114079), (Start: 12 @114115 has 1 MA's), (13, 114229), (16, 114295), (17, 114301), (18, 114322), (19, 114346),