



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 205832 Report

This analysis was run 02/22/25 on database version 588.

Pham number 205832 has 8 members, 0 are drafts.

Phages represented in each track:

- Track 1 : Megabear_55, Cepens_55, GodPhather_56, Argie_56, Taptic_55, Jeon_55, D12_54, PR_55

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 8 of the 8 non-draft genes in the pham.

Genes that call this "Most Annotated" start:

- Argie_56, Cepens_55, D12_54, GodPhather_56, Jeon_55, Megabear_55, PR_55, Taptic_55,

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: 8 of 8
- Called 100.0% of time when present
- Phage (with cluster) where this start called: Argie_56 (W), Cepens_55 (W), D12_54 (W), GodPhather_56 (W), Jeon_55 (W), Megabear_55 (W), PR_55 (W), Taptic_55 (W),

Summary by clusters:

There is one cluster represented in this pham: W

Info for manual annotations of cluster W:

- Start number 1 was manually annotated 8 times for cluster W.

Gene Information:

Gene: Argie_56 Start: 45292, Stop: 45573, Start Num: 1

Candidate Starts for Argie_56:

(Start: 1 @45292 has 8 MA's), (2, 45352), (3, 45409), (4, 45481), (5, 45553),

Gene: Cepens_55 Start: 45007, Stop: 45288, Start Num: 1

Candidate Starts for Cepens_55:

(Start: 1 @45007 has 8 MA's), (2, 45067), (3, 45124), (4, 45196), (5, 45268),

Gene: D12_54 Start: 45555, Stop: 45836, Start Num: 1

Candidate Starts for D12_54:

(Start: 1 @45555 has 8 MA's), (2, 45615), (3, 45672), (4, 45744), (5, 45816),

Gene: GodPhather_56 Start: 45795, Stop: 46076, Start Num: 1

Candidate Starts for GodPhather_56:

(Start: 1 @45795 has 8 MA's), (2, 45855), (3, 45912), (4, 45984), (5, 46056),

Gene: Jeon_55 Start: 45537, Stop: 45818, Start Num: 1

Candidate Starts for Jeon_55:

(Start: 1 @45537 has 8 MA's), (2, 45597), (3, 45654), (4, 45726), (5, 45798),

Gene: Megabear_55 Start: 44928, Stop: 45209, Start Num: 1

Candidate Starts for Megabear_55:

(Start: 1 @44928 has 8 MA's), (2, 44988), (3, 45045), (4, 45117), (5, 45189),

Gene: PR_55 Start: 45532, Stop: 45813, Start Num: 1

Candidate Starts for PR_55:

(Start: 1 @45532 has 8 MA's), (2, 45592), (3, 45649), (4, 45721), (5, 45793),

Gene: Taptic_55 Start: 44928, Stop: 45209, Start Num: 1

Candidate Starts for Taptic_55:

(Start: 1 @44928 has 8 MA's), (2, 44988), (3, 45045), (4, 45117), (5, 45189),