

A diagram showing a horizontal line divided into six segments by vertical lines. The segments are labeled 1 through 6 above the line. The entire area is shaded light pink.

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 6155 Report

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

Pham number 6155 has 8 members, 0 are drafts.

Phages represented in each track:

- Track 1 : Barco\_43, MintFen\_45, Archis\_48, BlingBling\_42, CaptainKirk2\_45, Melba\_44, Zarbodnamra\_44, Delian\_48

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

The start number called the most often in the published annotations is 2, it was called in 8 of the 8 non-draft genes in the pham.

Genes that call this "Most Annotated" start:

- Archis\_48, Barco\_43, BlingBling\_42, CaptainKirk2\_45, Delian\_48, Melba\_44, MintFen\_45, Zarbodnamra\_44,

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 2:

- 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: Archis\_48 (CV), Barco\_43 (CV), BlingBling\_42 (CV), CaptainKirk2\_45 (CV), Delian\_48 (CV), Melba\_44 (CV), MintFen\_45 (CV), Zarbodnamra\_44 (CV),

### **Summary by clusters:**

There is one cluster represented in this pham: CV

Info for manual annotations of cluster CV:

- Start number 2 was manually annotated 8 times for cluster CV.

**Gene Information:**

Gene: Archis\_48 Start: 33602, Stop: 34117, Start Num: 2

Candidate Starts for Archis\_48:

(1, 33389), (Start: 2 @33602 has 8 MA's), (3, 33722), (4, 33800), (5, 33896), (6, 33995),

Gene: Barco\_43 Start: 32652, Stop: 33167, Start Num: 2

Candidate Starts for Barco\_43:

(1, 32439), (Start: 2 @32652 has 8 MA's), (3, 32772), (4, 32850), (5, 32946), (6, 33045),

Gene: BlingBling\_42 Start: 33615, Stop: 34130, Start Num: 2

Candidate Starts for BlingBling\_42:

(1, 33402), (Start: 2 @33615 has 8 MA's), (3, 33735), (4, 33813), (5, 33909), (6, 34008),

Gene: CaptainKirk2\_45 Start: 32900, Stop: 33415, Start Num: 2

Candidate Starts for CaptainKirk2\_45:

(1, 32687), (Start: 2 @32900 has 8 MA's), (3, 33020), (4, 33098), (5, 33194), (6, 33293),

Gene: Delian\_48 Start: 32898, Stop: 33413, Start Num: 2

Candidate Starts for Delian\_48:

(1, 32685), (Start: 2 @32898 has 8 MA's), (3, 33018), (4, 33096), (5, 33192), (6, 33291),

Gene: Melba\_44 Start: 33440, Stop: 33955, Start Num: 2

Candidate Starts for Melba\_44:

(1, 33227), (Start: 2 @33440 has 8 MA's), (3, 33560), (4, 33638), (5, 33734), (6, 33833),

Gene: MintFen\_45 Start: 33514, Stop: 34029, Start Num: 2

Candidate Starts for MintFen\_45:

(1, 33301), (Start: 2 @33514 has 8 MA's), (3, 33634), (4, 33712), (5, 33808), (6, 33907),

Gene: Zarbodnamra\_44 Start: 32922, Stop: 33437, Start Num: 2

Candidate Starts for Zarbodnamra\_44:

(1, 32709), (Start: 2 @32922 has 8 MA's), (3, 33042), (4, 33120), (5, 33216), (6, 33315),