

A table with a pink background, divided into two rows and four columns. The columns are separated by vertical lines of different colors: yellow, red, light blue, and orange. The top edge of the table has small black arrows pointing to the right above each column line. The text '1: Pemberton_79' is located in the bottom-left corner of the table.

1: Pemberton_79				

A table with a pink background, divided into two rows and four columns. The columns are separated by vertical lines of different colors: green, red, light blue, and orange. The top edge of the table has small black arrows pointing to the right above each column line. The text '2: BiPauneto_80' is located in the bottom-left corner of the table.

2: BiPauneto_80				

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 307470 Report

This analysis was run 06/27/26 on database version 652.

Pham number 307470 has 2 members, 1 are drafts.

Phages represented in each track:

- Track 1 : Pemberton_79
- Track 2 : BiPauneto_80

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:

- BiPauneto_80, Pemberton_79,

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 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: BiPauneto_80 (CR4), Pemberton_79 (CR4),

Summary by clusters:

There is one cluster represented in this pham: CR4

Info for manual annotations of cluster CR4:

- Start number 1 was manually annotated 1 time for cluster CR4.

Gene Information:

Gene: BiPauneto_80 Start: 57304, Stop: 57504, Start Num: 1

Candidate Starts for BiPauneto_80:

(Start: 1 @57304 has 1 MA's), (2, 57379), (3, 57415), (4, 57457),

Gene: Pemberton_79 Start: 55371, Stop: 55571, Start Num: 1

Candidate Starts for Pemberton_79:

(Start: 1 @55371 has 1 MA's), (2, 55446), (3, 55482), (4, 55524),