



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 28047 Report

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

Pham number 28047 has 9 members, 2 are drafts.

Phages represented in each track:

- Track 1 : Cborch11_48, Oaker_46, Puissant_48, Megatron06_50, Thumb_47, Konstantine_50, Phreeze_45, Beckerton_46
- Track 2 : Damien_47

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 7 of the 7 non-draft genes in the pham.

Genes that call this "Most Annotated" start:

- Beckerton_46, Cborch11_48, Damien_47, Konstantine_50, Megatron06_50, Oaker_46, Phreeze_45, Puissant_48, Thumb_47,

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 9 of 9 (100.0%) of genes in pham
- Manual Annotations of this start: 7 of 7
- Called 100.0% of time when present
- Phage (with cluster) where this start called: Beckerton_46 (H1), Cborch11_48 (H1), Damien_47 (H1), Konstantine_50 (H1), Megatron06_50 (H1), Oaker_46 (H1), Phreeze_45 (H1), Puissant_48 (H1), Thumb_47 (H1),

Summary by clusters:

There is one cluster represented in this pham: H1

Info for manual annotations of cluster H1:

- Start number 1 was manually annotated 7 times for cluster H1.

Gene Information:

Gene: Beckerton_46 Start: 40343, Stop: 40528, Start Num: 1

Candidate Starts for Beckerton_46:

(Start: 1 @40343 has 7 MA's), (2, 40397), (3, 40418), (4, 40427),

Gene: Cborch11_48 Start: 39859, Stop: 40044, Start Num: 1

Candidate Starts for Cborch11_48:

(Start: 1 @39859 has 7 MA's), (2, 39913), (3, 39934), (4, 39943),

Gene: Damien_47 Start: 39822, Stop: 40007, Start Num: 1

Candidate Starts for Damien_47:

(Start: 1 @39822 has 7 MA's), (3, 39897), (4, 39906),

Gene: Konstantine_50 Start: 40653, Stop: 40838, Start Num: 1

Candidate Starts for Konstantine_50:

(Start: 1 @40653 has 7 MA's), (2, 40707), (3, 40728), (4, 40737),

Gene: Megatron06_50 Start: 40397, Stop: 40582, Start Num: 1

Candidate Starts for Megatron06_50:

(Start: 1 @40397 has 7 MA's), (2, 40451), (3, 40472), (4, 40481),

Gene: Oaker_46 Start: 40112, Stop: 40297, Start Num: 1

Candidate Starts for Oaker_46:

(Start: 1 @40112 has 7 MA's), (2, 40166), (3, 40187), (4, 40196),

Gene: Phreeze_45 Start: 39419, Stop: 39604, Start Num: 1

Candidate Starts for Phreeze_45:

(Start: 1 @39419 has 7 MA's), (2, 39473), (3, 39494), (4, 39503),

Gene: Puissant_48 Start: 39933, Stop: 40118, Start Num: 1

Candidate Starts for Puissant_48:

(Start: 1 @39933 has 7 MA's), (2, 39987), (3, 40008), (4, 40017),

Gene: Thumb_47 Start: 39854, Stop: 40039, Start Num: 1

Candidate Starts for Thumb_47:

(Start: 1 @39854 has 7 MA's), (2, 39908), (3, 39929), (4, 39938),