



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 105370 Report

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

Pham number 105370 has 8 members, 0 are drafts.

Phages represented in each track:

- Track 1 : Confidence_32
- Track 2 : Bunnybear_30
- Track 3 : Kita_35
- Track 4 : Jambalaya_38
- Track 5 : Malisha_38, Spooky_37
- Track 6 : BENtherdunthat_37
- Track 7 : Eyre_29

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

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

Genes that call this "Most Annotated" start:

- BENtherdunthat_37, Bunnybear_30, Confidence_32, Eyre_29, Jambalaya_38, Kita_35, Malisha_38, Spooky_37,

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 8:

- 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: BENtherdunthat_37 (DN1), Bunnybear_30 (CZ), Confidence_32 (CY1), Eyre_29 (singleton), Jambalaya_38 (DC1), Kita_35 (CZ1), Malisha_38 (DN), Spooky_37 (DN2),

Summary by clusters:

There are 8 clusters represented in this pham: DN, singleton, CZ1, CY1, CZ, DN1, DN2, DC1,

Info for manual annotations of cluster CY1:

- Start number 8 was manually annotated 1 time for cluster CY1.

Info for manual annotations of cluster CZ:

- Start number 8 was manually annotated 1 time for cluster CZ.

Info for manual annotations of cluster CZ1:

- Start number 8 was manually annotated 1 time for cluster CZ1.

Info for manual annotations of cluster DC1:

- Start number 8 was manually annotated 1 time for cluster DC1.

Info for manual annotations of cluster DN:

- Start number 8 was manually annotated 1 time for cluster DN.

Info for manual annotations of cluster DN1:

- Start number 8 was manually annotated 1 time for cluster DN1.

Info for manual annotations of cluster DN2:

- Start number 8 was manually annotated 1 time for cluster DN2.

Gene Information:

Gene: BENtherdunthat_37 Start: 28409, Stop: 28489, Start Num: 8

Candidate Starts for BENtherdunthat_37:

(1, 28253), (Start: 8 @28409 has 8 MA's), (9, 28442),

Gene: Bunnybear_30 Start: 27670, Stop: 27744, Start Num: 8

Candidate Starts for Bunnybear_30:

(4, 27583), (Start: 8 @27670 has 8 MA's),

Gene: Confidence_32 Start: 27302, Stop: 27379, Start Num: 8

Candidate Starts for Confidence_32:

(Start: 8 @27302 has 8 MA's),

Gene: Eyre_29 Start: 24447, Stop: 24527, Start Num: 8

Candidate Starts for Eyre_29:

(2, 24321), (3, 24336), (6, 24429), (Start: 8 @24447 has 8 MA's),

Gene: Jambalaya_38 Start: 33394, Stop: 33465, Start Num: 8

Candidate Starts for Jambalaya_38:

(5, 33352), (7, 33379), (Start: 8 @33394 has 8 MA's), (10, 33448),

Gene: Kita_35 Start: 29024, Stop: 29104, Start Num: 8

Candidate Starts for Kita_35:

(Start: 8 @29024 has 8 MA's),

Gene: Malisha_38 Start: 29395, Stop: 29481, Start Num: 8
Candidate Starts for Malisha_38:
(Start: 8 @29395 has 8 MA's), (11, 29452),

Gene: Spooky_37 Start: 29041, Stop: 29127, Start Num: 8
Candidate Starts for Spooky_37:
(Start: 8 @29041 has 8 MA's), (11, 29098),