



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 5082 Report

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

Pham number 5082 has 15 members, 3 are drafts.

Phages represented in each track:

- Track 1 : Miskis\_98, Aphelion\_101, Dusty\_97, Abscondus\_99
- Track 2 : Engineer\_101, Smoothie\_101, Cucurbita\_101, Toniann\_102, Lozinak\_100, PhinkBoden\_99, Bachita\_102
- Track 3 : Culver\_102, ClubL\_101, Norvs\_102
- Track 4 : WilliamBoone\_102

### ***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 12 of the 12 non-draft genes in the pham.

Genes that call this "Most Annotated" start:

- Abscondus\_99, Aphelion\_101, Bachita\_102, ClubL\_101, Cucurbita\_101, Culver\_102, Dusty\_97, Engineer\_101, Lozinak\_100, Miskis\_98, Norvs\_102, PhinkBoden\_99, Smoothie\_101, Toniann\_102, WilliamBoone\_102,

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 15 of 15 ( 100.0% ) of genes in pham
- Manual Annotations of this start: 12 of 12
- Called 100.0% of time when present
- Phage (with cluster) where this start called: Abscondus\_99 (CQ), Aphelion\_101 (CQ1), Bachita\_102 (CQ1), ClubL\_101 (CQ1), Cucurbita\_101 (CQ1), Culver\_102 (CQ1), Dusty\_97 (CQ), Engineer\_101 (CQ1), Lozinak\_100 (CQ1), Miskis\_98 (CQ), Norvs\_102 (CQ), PhinkBoden\_99 (CQ1), Smoothie\_101 (CQ1), Toniann\_102 (CQ1), WilliamBoone\_102 (CQ1),

## Summary by clusters:

There are 2 clusters represented in this pham: CQ1, CQ,

Info for manual annotations of cluster CQ:

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

Info for manual annotations of cluster CQ1:

- Start number 1 was manually annotated 11 times for cluster CQ1.

## Gene Information:

Gene: Abscondus\_99 Start: 59444, Stop: 59635, Start Num: 1

Candidate Starts for Abscondus\_99:

(Start: 1 @59444 has 12 MA's), (2, 59471),

Gene: Aphelion\_101 Start: 60997, Stop: 61188, Start Num: 1

Candidate Starts for Aphelion\_101:

(Start: 1 @60997 has 12 MA's), (2, 61024),

Gene: Bachita\_102 Start: 60655, Stop: 60846, Start Num: 1

Candidate Starts for Bachita\_102:

(Start: 1 @60655 has 12 MA's), (3, 60700),

Gene: ClubL\_101 Start: 59587, Stop: 59775, Start Num: 1

Candidate Starts for ClubL\_101:

(Start: 1 @59587 has 12 MA's), (4, 59650), (5, 59701),

Gene: Cucurbita\_101 Start: 61162, Stop: 61353, Start Num: 1

Candidate Starts for Cucurbita\_101:

(Start: 1 @61162 has 12 MA's), (3, 61207),

Gene: Culver\_102 Start: 59487, Stop: 59675, Start Num: 1

Candidate Starts for Culver\_102:

(Start: 1 @59487 has 12 MA's), (4, 59550), (5, 59601),

Gene: Dusty\_97 Start: 59503, Stop: 59694, Start Num: 1

Candidate Starts for Dusty\_97:

(Start: 1 @59503 has 12 MA's), (2, 59530),

Gene: Engineer\_101 Start: 60628, Stop: 60819, Start Num: 1

Candidate Starts for Engineer\_101:

(Start: 1 @60628 has 12 MA's), (3, 60673),

Gene: Lozinak\_100 Start: 60466, Stop: 60657, Start Num: 1

Candidate Starts for Lozinak\_100:

(Start: 1 @60466 has 12 MA's), (3, 60511),

Gene: Miskis\_98 Start: 59166, Stop: 59357, Start Num: 1

Candidate Starts for Miskis\_98:

(Start: 1 @59166 has 12 MA's), (2, 59193),

Gene: Norvs\_102 Start: 60495, Stop: 60683, Start Num: 1  
Candidate Starts for Norvs\_102:  
(Start: 1 @60495 has 12 MA's), (4, 60558), (5, 60609),

Gene: PhinkBoden\_99 Start: 60256, Stop: 60447, Start Num: 1  
Candidate Starts for PhinkBoden\_99:  
(Start: 1 @60256 has 12 MA's), (3, 60301),

Gene: Smoothie\_101 Start: 60466, Stop: 60657, Start Num: 1  
Candidate Starts for Smoothie\_101:  
(Start: 1 @60466 has 12 MA's), (3, 60511),

Gene: Toniann\_102 Start: 60435, Stop: 60626, Start Num: 1  
Candidate Starts for Toniann\_102:  
(Start: 1 @60435 has 12 MA's), (3, 60480),

Gene: WilliamBoone\_102 Start: 59204, Stop: 59395, Start Num: 1  
Candidate Starts for WilliamBoone\_102:  
(Start: 1 @59204 has 12 MA's), (3, 59249), (6, 59345),