



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 142968 Report

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

Pham number 142968 has 15 members, 3 are drafts.

Phages represented in each track:

- Track 1 : Abscondus\_35, Dusty\_34, Culver\_36
- Track 2 : Aphelion\_36, Lozinak\_36, Bachita\_38, Smoothie\_37, Cucurbita\_38, PhinkBoden\_36, Toniann\_36, Norvs\_37, Engineer\_37
- Track 3 : ClubL\_36, Miskis\_38
- Track 4 : WilliamBoone\_36

### ***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\_35, Aphelion\_36, Bachita\_38, ClubL\_36, Cucurbita\_38, Culver\_36, Dusty\_34, Engineer\_37, Lozinak\_36, Miskis\_38, Norvs\_37, PhinkBoden\_36, Smoothie\_37, Toniann\_36, WilliamBoone\_36,

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\_35 (CQ), Aphelion\_36 (CQ1), Bachita\_38 (CQ1), ClubL\_36 (CQ1), Cucurbita\_38 (CQ1), Culver\_36 (CQ1), Dusty\_34 (CQ), Engineer\_37 (CQ1), Lozinak\_36 (CQ1), Miskis\_38 (CQ), Norvs\_37 (CQ), PhinkBoden\_36 (CQ1), Smoothie\_37 (CQ1), Toniann\_36 (CQ1), WilliamBoone\_36 (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\_35 Start: 15180, Stop: 15494, Start Num: 1

Candidate Starts for Abscondus\_35:

(Start: 1 @15180 has 12 MA's), (2, 15201), (3, 15207), (4, 15297), (5, 15318), (6, 15321),

Gene: Aphelion\_36 Start: 15425, Stop: 15763, Start Num: 1

Candidate Starts for Aphelion\_36:

(Start: 1 @15425 has 12 MA's), (4, 15542), (5, 15563), (6, 15566), (7, 15617), (8, 15641), (9, 15653), (10, 15665),

Gene: Bachita\_38 Start: 15858, Stop: 16196, Start Num: 1

Candidate Starts for Bachita\_38:

(Start: 1 @15858 has 12 MA's), (4, 15975), (5, 15996), (6, 15999), (7, 16050), (8, 16074), (9, 16086), (10, 16098),

Gene: ClubL\_36 Start: 15359, Stop: 15685, Start Num: 1

Candidate Starts for ClubL\_36:

(Start: 1 @15359 has 12 MA's), (4, 15476), (5, 15497), (6, 15500), (7, 15551), (8, 15575), (9, 15587),

Gene: Cucurbita\_38 Start: 16717, Stop: 17055, Start Num: 1

Candidate Starts for Cucurbita\_38:

(Start: 1 @16717 has 12 MA's), (4, 16834), (5, 16855), (6, 16858), (7, 16909), (8, 16933), (9, 16945), (10, 16957),

Gene: Culver\_36 Start: 15180, Stop: 15494, Start Num: 1

Candidate Starts for Culver\_36:

(Start: 1 @15180 has 12 MA's), (2, 15201), (3, 15207), (4, 15297), (5, 15318), (6, 15321),

Gene: Dusty\_34 Start: 15180, Stop: 15494, Start Num: 1

Candidate Starts for Dusty\_34:

(Start: 1 @15180 has 12 MA's), (2, 15201), (3, 15207), (4, 15297), (5, 15318), (6, 15321),

Gene: Engineer\_37 Start: 15373, Stop: 15711, Start Num: 1

Candidate Starts for Engineer\_37:

(Start: 1 @15373 has 12 MA's), (4, 15490), (5, 15511), (6, 15514), (7, 15565), (8, 15589), (9, 15601), (10, 15613),

Gene: Lozinak\_36 Start: 15428, Stop: 15766, Start Num: 1

Candidate Starts for Lozinak\_36:

(Start: 1 @15428 has 12 MA's), (4, 15545), (5, 15566), (6, 15569), (7, 15620), (8, 15644), (9, 15656), (10, 15668),

Gene: Miskis\_38 Start: 15203, Stop: 15529, Start Num: 1

Candidate Starts for Miskis\_38:

(Start: 1 @15203 has 12 MA's), (4, 15320), (5, 15341), (6, 15344), (7, 15395), (8, 15419), (9, 15431),

Gene: Norvs\_37 Start: 15430, Stop: 15768, Start Num: 1

Candidate Starts for Norvs\_37:

(Start: 1 @15430 has 12 MA's), (4, 15547), (5, 15568), (6, 15571), (7, 15622), (8, 15646), (9, 15658), (10, 15670),

Gene: PhinkBoden\_36 Start: 15811, Stop: 16149, Start Num: 1

Candidate Starts for PhinkBoden\_36:

(Start: 1 @15811 has 12 MA's), (4, 15928), (5, 15949), (6, 15952), (7, 16003), (8, 16027), (9, 16039), (10, 16051),

Gene: Smoothie\_37 Start: 15428, Stop: 15766, Start Num: 1

Candidate Starts for Smoothie\_37:

(Start: 1 @15428 has 12 MA's), (4, 15545), (5, 15566), (6, 15569), (7, 15620), (8, 15644), (9, 15656), (10, 15668),

Gene: Toniann\_36 Start: 15373, Stop: 15711, Start Num: 1

Candidate Starts for Toniann\_36:

(Start: 1 @15373 has 12 MA's), (4, 15490), (5, 15511), (6, 15514), (7, 15565), (8, 15589), (9, 15601), (10, 15613),

Gene: WilliamBoone\_36 Start: 14761, Stop: 15075, Start Num: 1

Candidate Starts for WilliamBoone\_36:

(Start: 1 @14761 has 12 MA's), (2, 14782), (5, 14899),