

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 7657 Report

This analysis was run 04/05/24 on database version 557.

Pham number 7657 has 9 members, 3 are drafts.

Phages represented in each track:

Track 1 : Baee\_47, Acadian\_48, Rich\_46

Track 2 : Suigeneris\_48, Donny\_48

Track 3 : Nangang\_47, DigitDog\_47

Track 4 : Serendipitous\_47

Track 5 : EastView9101\_48

# 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 6 of the 6 non-draft genes in the pham.

Genes that call this "Most Annotated" start:

Acadian\_48, Baee\_47, Donny\_48, Rich\_46, Serendipitous\_47, Suigeneris\_48,

Genes that have the "Most Annotated" start but do not call it:

DigitDog\_47, EastView9101\_48, Nangang\_47,

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: 6 of 6
- Called 66.7% of time when present
- Phage (with cluster) where this start called: Acadian\_48 (B5), Baee\_47 (B5), Donny\_48 (B5), Rich\_46 (B5), Serendipitous\_47 (B5), Suigeneris\_48 (B5),

### Start 2:

- Found in 9 of 9 ( 100.0% ) of genes in pham
- No Manual Annotations of this start.
- Called 33.3% of time when present

• Phage (with cluster) where this start called: DigitDog\_47 (B5), EastView9101\_48 (B5), Nangang\_47 (B5),

### **Summary by clusters:**

There is one cluster represented in this pham: B5

Info for manual annotations of cluster B5:

•Start number 1 was manually annotated 6 times for cluster B5.

### Gene Information:

Gene: Acadian\_48 Start: 42592, Stop: 42729, Start Num: 1

Candidate Starts for Acadian 48:

(Start: 1 @ 42592 has 6 MA's), (2, 42610), (3, 42619), (4, 42634),

Gene: Baee\_47 Start: 42476, Stop: 42613, Start Num: 1

Candidate Starts for Baee 47:

(Start: 1 @ 42476 has 6 MA's), (2, 42494), (3, 42503), (4, 42518),

Gene: DigitDog\_47 Start: 42310, Stop: 42432, Start Num: 2

Candidate Starts for DigitDog 47:

(Start: 1 @42292 has 6 MA's), (2, 42310),

Gene: Donny\_48 Start: 42597, Stop: 42734, Start Num: 1

Candidate Starts for Donny\_48:

(Start: 1 @42597 has 6 MA's), (2, 42615),

Gene: EastView9101\_48 Start: 42265, Stop: 42384, Start Num: 2

Candidate Starts for EastView9101\_48:

(Start: 1 @42247 has 6 MA's), (2, 42265), (3, 42274), (4, 42289),

Gene: Nangang 47 Start: 42675, Stop: 42797, Start Num: 2

Candidate Starts for Nangang\_47:

(Start: 1 @ 42657 has 6 MA's), (2, 42675),

Gene: Rich\_46 Start: 42464, Stop: 42601, Start Num: 1

Candidate Starts for Rich 46:

(Start: 1 @ 42464 has 6 MA's), (2, 42482), (3, 42491), (4, 42506),

Gene: Serendipitous 47 Start: 42407, Stop: 42547, Start Num: 1

Candidate Starts for Serendipitous\_47:

(Start: 1 @42407 has 6 MA's), (2, 42425), (4, 42449),

Gene: Suigeneris\_48 Start: 42597, Stop: 42734, Start Num: 1

Candidate Starts for Suigeneris\_48: (Start: 1 @42597 has 6 MA's), (2, 42615),