



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 3005 Report

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

Pham number 3005 has 26 members, 1 are drafts.

Phages represented in each track:

- Track 1 : Amao\_116, Marshmallow\_114, Dusk\_113, ChotaBhai\_116, BadStone\_113, Icee\_117, Gator\_114, Henry\_114, Phaja\_114, Cactus\_120, Tuco\_117, TeardropMSU\_112, Bask21\_116, ABCat\_113, IHOP\_114, Glexan\_115, Pharsalus\_112, Harella\_117, DrDrey\_116, Pat3\_113, Kostya\_116, Cookies\_113, Sotrice96\_117, Mindy\_115, SirDuracell\_116
- Track 2 : Inca\_112

### ***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 24 of the 25 non-draft genes in the pham.

Genes that call this "Most Annotated" start:

- ABCat\_113, Amao\_116, BadStone\_113, Bask21\_116, Cactus\_120, ChotaBhai\_116, Cookies\_113, DrDrey\_116, Dusk\_113, Gator\_114, Glexan\_115, Harella\_117, Henry\_114, IHOP\_114, Icee\_117, Kostya\_116, Marshmallow\_114, Mindy\_115, Pat3\_113, Phaja\_114, Pharsalus\_112, SirDuracell\_116, Sotrice96\_117, TeardropMSU\_112, Tuco\_117,

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

- Inca\_112,

Genes that do not have the "Most Annotated" start:

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### **Summary by start number:**

Start 1:

- Found in 26 of 26 ( 100.0% ) of genes in pham
- Manual Annotations of this start: 24 of 25
- Called 96.2% of time when present
- Phage (with cluster) where this start called: ABCat\_113 (E), Amao\_116 (E), BadStone\_113 (E), Bask21\_116 (E), Cactus\_120 (E), ChotaBhai\_116 (E), Cookies\_113 (E), DrDrey\_116 (E), Dusk\_113 (E), Gator\_114 (E), Glexan\_115 (E),

Harella\_117 (E), Henry\_114 (E), IHOP\_114 (E), Icee\_117 (E), Kostya\_116 (E), Marshmallow\_114 (E), Mindy\_115 (E), Pat3\_113 (E), Phaja\_114 (E), Pharsalus\_112 (E), SirDuracell\_116 (E), Sotrice96\_117 (E), TeardropMSU\_112 (E), Tuco\_117 (E),

Start 2:

- Found in 26 of 26 ( 100.0% ) of genes in pham
- Manual Annotations of this start: 1 of 25
- Called 3.8% of time when present
- Phage (with cluster) where this start called: Inca\_112 (E),

### **Summary by clusters:**

There is one cluster represented in this pham: E

Info for manual annotations of cluster E:

- Start number 1 was manually annotated 24 times for cluster E.
- Start number 2 was manually annotated 1 time for cluster E.

### **Gene Information:**

Gene: ABCat\_113 Start: 66247, Stop: 66456, Start Num: 1

Candidate Starts for ABCat\_113:

(Start: 1 @66247 has 24 MA's), (Start: 2 @66274 has 1 MA's),

Gene: Amao\_116 Start: 65810, Stop: 66019, Start Num: 1

Candidate Starts for Amao\_116:

(Start: 1 @65810 has 24 MA's), (Start: 2 @65837 has 1 MA's),

Gene: BadStone\_113 Start: 65795, Stop: 66004, Start Num: 1

Candidate Starts for BadStone\_113:

(Start: 1 @65795 has 24 MA's), (Start: 2 @65822 has 1 MA's),

Gene: Bask21\_116 Start: 64612, Stop: 64821, Start Num: 1

Candidate Starts for Bask21\_116:

(Start: 1 @64612 has 24 MA's), (Start: 2 @64639 has 1 MA's),

Gene: Cactus\_120 Start: 65155, Stop: 65364, Start Num: 1

Candidate Starts for Cactus\_120:

(Start: 1 @65155 has 24 MA's), (Start: 2 @65182 has 1 MA's),

Gene: ChotaBhai\_116 Start: 65554, Stop: 65763, Start Num: 1

Candidate Starts for ChotaBhai\_116:

(Start: 1 @65554 has 24 MA's), (Start: 2 @65581 has 1 MA's),

Gene: Cookies\_113 Start: 65540, Stop: 65749, Start Num: 1

Candidate Starts for Cookies\_113:

(Start: 1 @65540 has 24 MA's), (Start: 2 @65567 has 1 MA's),

Gene: DrDrey\_116 Start: 66933, Stop: 67142, Start Num: 1

Candidate Starts for DrDrey\_116:

(Start: 1 @66933 has 24 MA's), (Start: 2 @66960 has 1 MA's),

Gene: Dusk\_113 Start: 65455, Stop: 65664, Start Num: 1  
Candidate Starts for Dusk\_113:  
(Start: 1 @65455 has 24 MA's), (Start: 2 @65482 has 1 MA's),

Gene: Gator\_114 Start: 65507, Stop: 65716, Start Num: 1  
Candidate Starts for Gator\_114:  
(Start: 1 @65507 has 24 MA's), (Start: 2 @65534 has 1 MA's),

Gene: Glexan\_115 Start: 65657, Stop: 65866, Start Num: 1  
Candidate Starts for Glexan\_115:  
(Start: 1 @65657 has 24 MA's), (Start: 2 @65684 has 1 MA's),

Gene: Harella\_117 Start: 66173, Stop: 66382, Start Num: 1  
Candidate Starts for Harella\_117:  
(Start: 1 @66173 has 24 MA's), (Start: 2 @66200 has 1 MA's),

Gene: Henry\_114 Start: 65458, Stop: 65667, Start Num: 1  
Candidate Starts for Henry\_114:  
(Start: 1 @65458 has 24 MA's), (Start: 2 @65485 has 1 MA's),

Gene: IHOP\_114 Start: 65176, Stop: 65385, Start Num: 1  
Candidate Starts for IHOP\_114:  
(Start: 1 @65176 has 24 MA's), (Start: 2 @65203 has 1 MA's),

Gene: Icee\_117 Start: 65146, Stop: 65355, Start Num: 1  
Candidate Starts for Icee\_117:  
(Start: 1 @65146 has 24 MA's), (Start: 2 @65173 has 1 MA's),

Gene: Inca\_112 Start: 63382, Stop: 63564, Start Num: 2  
Candidate Starts for Inca\_112:  
(Start: 1 @63355 has 24 MA's), (Start: 2 @63382 has 1 MA's),

Gene: Kostya\_116 Start: 65278, Stop: 65487, Start Num: 1  
Candidate Starts for Kostya\_116:  
(Start: 1 @65278 has 24 MA's), (Start: 2 @65305 has 1 MA's),

Gene: Marshmallow\_114 Start: 65960, Stop: 66169, Start Num: 1  
Candidate Starts for Marshmallow\_114:  
(Start: 1 @65960 has 24 MA's), (Start: 2 @65987 has 1 MA's),

Gene: Mindy\_115 Start: 65119, Stop: 65328, Start Num: 1  
Candidate Starts for Mindy\_115:  
(Start: 1 @65119 has 24 MA's), (Start: 2 @65146 has 1 MA's),

Gene: Pat3\_113 Start: 64613, Stop: 64822, Start Num: 1  
Candidate Starts for Pat3\_113:  
(Start: 1 @64613 has 24 MA's), (Start: 2 @64640 has 1 MA's),

Gene: Phaja\_114 Start: 65181, Stop: 65390, Start Num: 1  
Candidate Starts for Phaja\_114:  
(Start: 1 @65181 has 24 MA's), (Start: 2 @65208 has 1 MA's),

Gene: Pharsalus\_112 Start: 65255, Stop: 65464, Start Num: 1  
Candidate Starts for Pharsalus\_112:  
(Start: 1 @65255 has 24 MA's), (Start: 2 @65282 has 1 MA's),

Gene: SirDuracell\_116 Start: 65112, Stop: 65321, Start Num: 1  
Candidate Starts for SirDuracell\_116:  
(Start: 1 @65112 has 24 MA's), (Start: 2 @65139 has 1 MA's),

Gene: Sotrice96\_117 Start: 65899, Stop: 66108, Start Num: 1  
Candidate Starts for Sotrice96\_117:  
(Start: 1 @65899 has 24 MA's), (Start: 2 @65926 has 1 MA's),

Gene: TeardropMSU\_112 Start: 65001, Stop: 65210, Start Num: 1  
Candidate Starts for TeardropMSU\_112:  
(Start: 1 @65001 has 24 MA's), (Start: 2 @65028 has 1 MA's),

Gene: Tuco\_117 Start: 67060, Stop: 67269, Start Num: 1  
Candidate Starts for Tuco\_117:  
(Start: 1 @67060 has 24 MA's), (Start: 2 @67087 has 1 MA's),