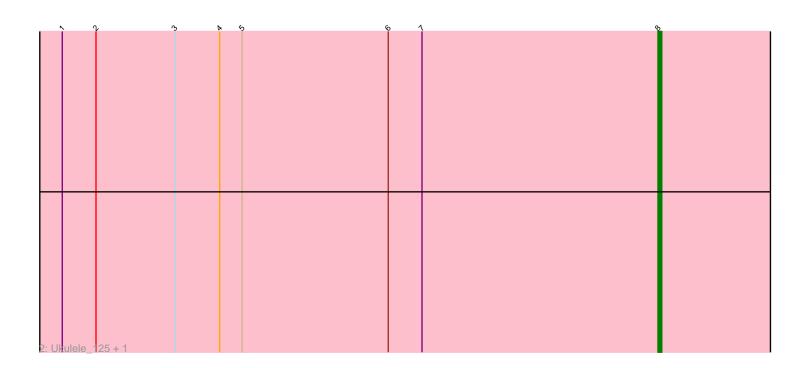
| 1: lcee_135 + 20  |                   | 9 |  |
|-------------------|-------------------|---|--|
| I: Icee_135 + 20  |                   |   |  |
| 1: Icee_135 + 20  |                   |   |  |
| 1: Icee_135 + 20  |                   |   |  |
| 1: Icee_135 + 20  |                   |   |  |
| I: Icee_135 + 20  |                   |   |  |
| 1: Icee_135 + 20  |                   |   |  |
| 1: Icee_135 + 20  |                   |   |  |
| 1: Icee_135 + 20  |                   |   |  |
| 1: lcee_135 + 20  |                   |   |  |
| 1: lcee_135 + 20  |                   |   |  |
| 1: lcee_135 + 20  |                   |   |  |
| 1: lcee_135 + 20  |                   |   |  |
| 1: Icee_135 + 20  |                   |   |  |
| 1: Icee_135 + 20  |                   |   |  |
| 1: Icee_135 + 20  |                   |   |  |
| 1: Icee_135 + 20  |                   |   |  |
| 1: Icee_135 + 20  |                   |   |  |
| 1: Icee_135 + 20  |                   |   |  |
| 1: Icee_135 + 20  |                   |   |  |
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| 1: Icee_135 + 20  |                   |   |  |
| 1: Icee_135 + 20  |                   |   |  |
| 1: Icee_135 + 20  |                   |   |  |
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| 1: Icee_135 + 20  |                   |   |  |
| 1: Icee_135 + 20  |                   |   |  |
| 1: Icee_135 + 20  |                   |   |  |
| 1: Icee_135 + 20  |                   |   |  |
| 1: Icee_135 + 20  |                   |   |  |
| 1: Icee_135 + 20  |                   |   |  |
| 1: Icee_135 + 20  |                   |   |  |
| 1: Icee_135 + 20  |                   |   |  |
| 1: Icee_135 + 20  |                   |   |  |
| 1: Icee_135 + 20  |                   |   |  |
| 1: Icee_135 + 20  |                   |   |  |
| 1: Icee_135 + 20  |                   |   |  |
| II. ICEE_155 + 20 | 1.1000 125   20   |   |  |
|                   | II. ICEE_133 + 20 |   |  |



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 106538 Report

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

Pham number 106538 has 23 members, 0 are drafts.

Phages represented in each track:

Track 1: Icee\_135, ABCat\_131, ChotaBhai\_134, NelitzaMV\_128, Phaja\_134, Thresher\_132, Murphy\_129, RiverMonster\_129, ShereKhan\_129, Bask21\_135, IHOP\_134, Tuco\_136, Nimrod\_131, Dusk\_131, BadStone\_131, Barbarian\_130, Maxxinista\_132, DrDrey\_134, TeardropMSU\_131, Mindy\_133, Simpliphy\_129
Track 2: Ukulele 125, Phaux 132

# 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 23 of the 23 non-draft genes in the pham.

Genes that call this "Most Annotated" start:

• ABCat\_131, BadStone\_131, Barbarian\_130, Bask21\_135, ChotaBhai\_134, DrDrey\_134, Dusk\_131, IHOP\_134, Icee\_135, Maxxinista\_132, Mindy\_133, Murphy\_129, NelitzaMV\_128, Nimrod\_131, Phaja\_134, Phaux\_132, RiverMonster\_129, ShereKhan\_129, Simpliphy\_129, TeardropMSU\_131, Thresher\_132, Tuco\_136, Ukulele\_125,

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 23 of 23 (100.0%) of genes in pham
- Manual Annotations of this start: 23 of 23
- Called 100.0% of time when present
- Phage (with cluster) where this start called: ABCat\_131 (E), BadStone\_131 (E), Barbarian\_130 (E), Bask21\_135 (E), ChotaBhai\_134 (E), DrDrey\_134 (E), Dusk\_131 (E), IHOP\_134 (E), Icee\_135 (E), Maxxinista\_132 (E), Mindy\_133 (E), Murphy\_129 (E), NelitzaMV\_128 (E), Nimrod\_131 (E), Phaja\_134 (E), Phaux\_132 (E),

RiverMonster\_129 (E), ShereKhan\_129 (E), Simpliphy\_129 (E), TeardropMSU\_131 (E), Thresher\_132 (E), Tuco\_136 (E), Ukulele\_125 (E),

# **Summary by clusters:**

There is one cluster represented in this pham: E

Info for manual annotations of cluster E:

•Start number 8 was manually annotated 23 times for cluster E.

### Gene Information:

Gene: ABCat\_131 Start: 71301, Stop: 71227, Start Num: 8

Candidate Starts for ABCat\_131: (Start: 8 @71301 has 23 MA's),

Gene: BadStone\_131 Start: 70849, Stop: 70775, Start Num: 8

Candidate Starts for BadStone\_131: (Start: 8 @70849 has 23 MA's),

Gene: Barbarian\_130 Start: 68606, Stop: 68532, Start Num: 8

Candidate Starts for Barbarian\_130: (Start: 8 @68606 has 23 MA's),

Gene: Bask21\_135 Start: 69772, Stop: 69698, Start Num: 8

Candidate Starts for Bask21\_135: (Start: 8 @69772 has 23 MA's),

Gene: ChotaBhai\_134 Start: 70608, Stop: 70534, Start Num: 8

Candidate Starts for ChotaBhai\_134: (Start: 8 @70608 has 23 MA's),

Gene: DrDrey 134 Start: 72082, Stop: 72008, Start Num: 8

Candidate Starts for DrDrey\_134: (Start: 8 @72082 has 23 MA's),

Gene: Dusk\_131 Start: 70509, Stop: 70435, Start Num: 8

Candidate Starts for Dusk\_131: (Start: 8 @70509 has 23 MA's),

Gene: IHOP 134 Start: 70325, Stop: 70251, Start Num: 8

Candidate Starts for IHOP\_134: (Start: 8 @70325 has 23 MA's),

Gene: Icee\_135 Start: 70201, Stop: 70127, Start Num: 8

Candidate Starts for Icee\_135: (Start: 8 @70201 has 23 MA's),

Gene: Maxxinista\_132 Start: 69783, Stop: 69709, Start Num: 8

Candidate Starts for Maxxinista\_132: (Start: 8 @69783 has 23 MA's),

Gene: Mindy\_133 Start: 70173, Stop: 70099, Start Num: 8

Candidate Starts for Mindy\_133: (Start: 8 @70173 has 23 MA's),

Gene: Murphy\_129 Start: 70157, Stop: 70083, Start Num: 8

Candidate Starts for Murphy\_129: (Start: 8 @70157 has 23 MA's),

Gene: NelitzaMV\_128 Start: 68480, Stop: 68406, Start Num: 8

Candidate Starts for NelitzaMV\_128: (Start: 8 @68480 has 23 MA's),

Gene: Nimrod\_131 Start: 70942, Stop: 70868, Start Num: 8

Candidate Starts for Nimrod\_131: (Start: 8 @70942 has 23 MA's),

Gene: Phaja\_134 Start: 70330, Stop: 70256, Start Num: 8

Candidate Starts for Phaja\_134: (Start: 8 @70330 has 23 MA's),

Gene: Phaux\_132 Start: 71097, Stop: 71023, Start Num: 8

Candidate Starts for Phaux\_132:

(1, 71256), (2, 71247), (3, 71226), (4, 71214), (5, 71208), (6, 71169), (7, 71160), (Start: 8 @71097 has

23 MA's),

Gene: RiverMonster\_129 Start: 70070, Stop: 69996, Start Num: 8

Candidate Starts for RiverMonster 129:

(Start: 8 @70070 has 23 MA's),

Gene: ShereKhan\_129 Start: 70829, Stop: 70755, Start Num: 8

Candidate Starts for ShereKhan\_129:

(Start: 8 @70829 has 23 MA's),

Gene: Simpliphy\_129 Start: 69641, Stop: 69567, Start Num: 8

Candidate Starts for Simpliphy\_129: (Start: 8 @69641 has 23 MA's),

Gene: TeardropMSU\_131 Start: 70055, Stop: 69981, Start Num: 8

Candidate Starts for TeardropMSU\_131:

(Start: 8 @70055 has 23 MA's),

Gene: Thresher\_132 Start: 70949, Stop: 70875, Start Num: 8

Candidate Starts for Thresher\_132: (Start: 8 @70949 has 23 MA's),

Gene: Tuco\_136 Start: 72114, Stop: 72040, Start Num: 8

Candidate Starts for Tuco\_136: (Start: 8 @72114 has 23 MA's),

Gene: Ukulele\_125 Start: 69450, Stop: 69376, Start Num: 8

Candidate Starts for Ukulele\_125:

(1, 69609), (2, 69600), (3, 69579), (4, 69567), (5, 69561), (6, 69522), (7, 69513), (Start: 8 @69450 has 23 MA's),