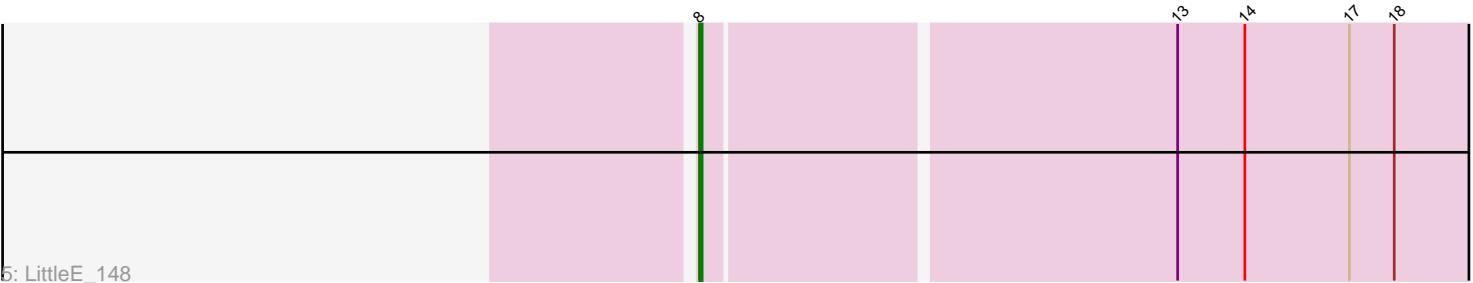
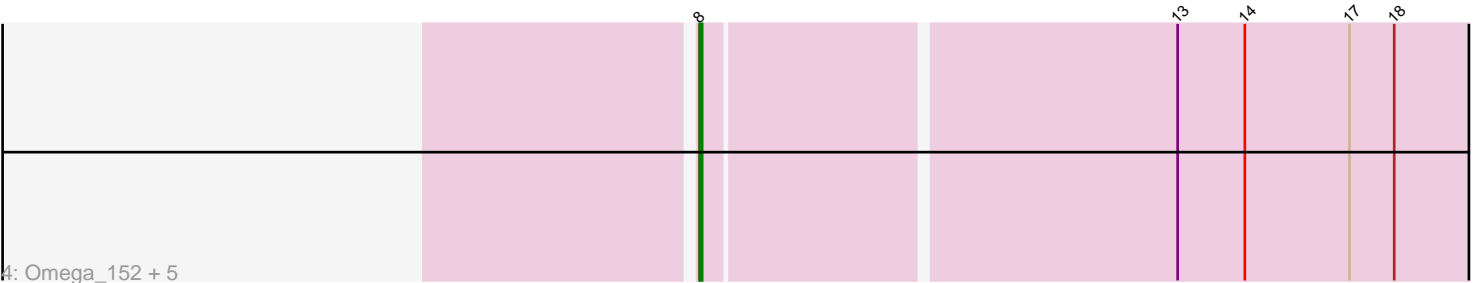
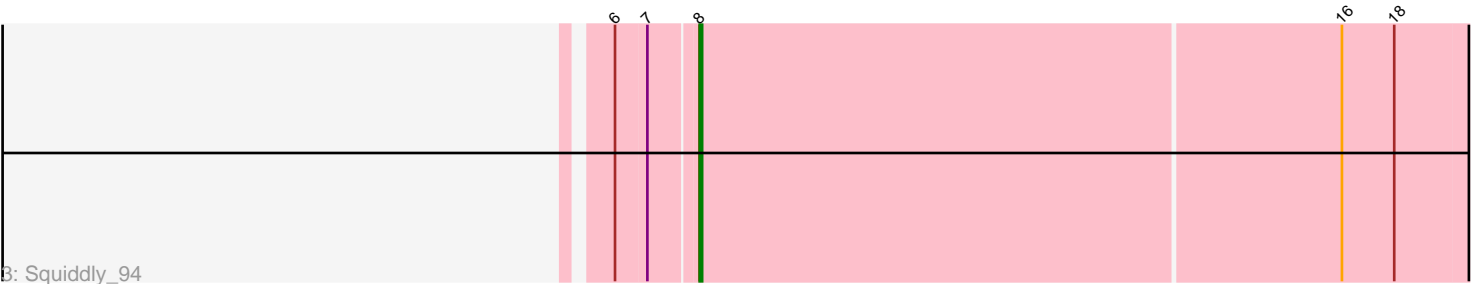
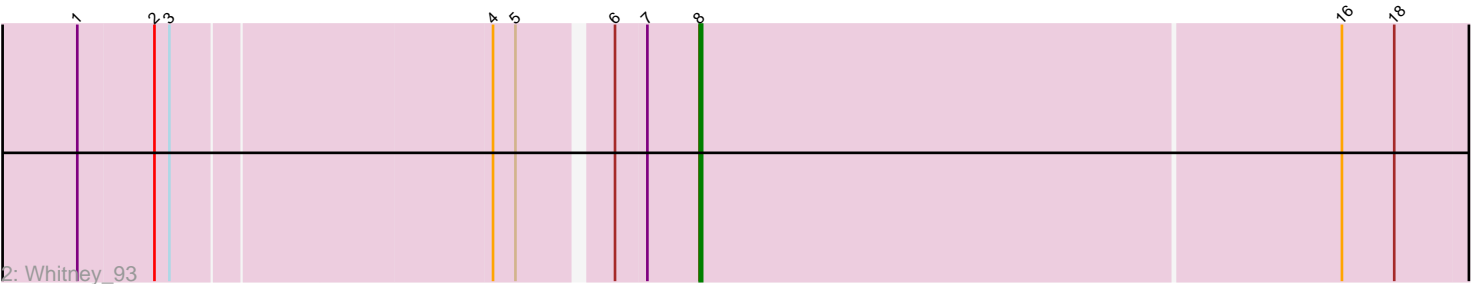
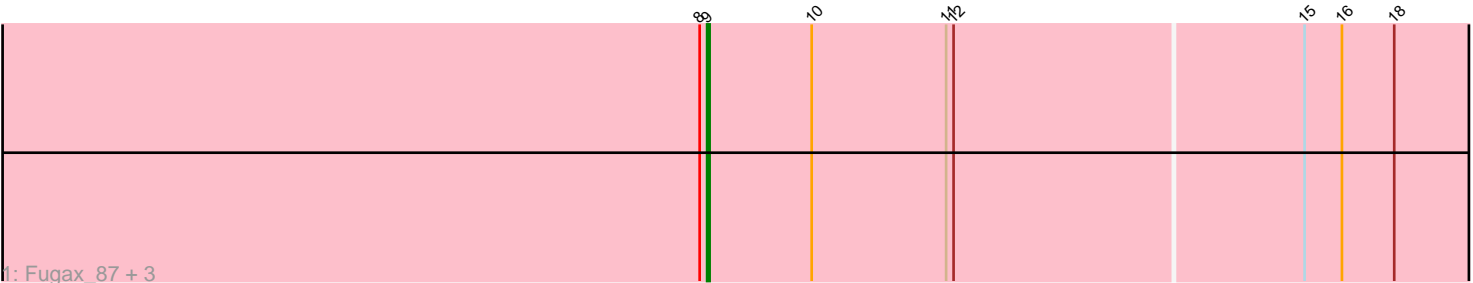


Pham 87182



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 87182 Report

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

Pham number 87182 has 13 members, 0 are drafts.

Phages represented in each track:

- Track 1 : Fugax\_87, ClamChowder\_86, Danyall\_84, Barb\_87
- Track 2 : Whitney\_93
- Track 3 : Squiddly\_94
- Track 4 : Omega\_152, Courthouse\_140, Hannaconda\_138, Squint\_139, Ariel\_144, Superphikiman\_142
- Track 5 : LittleE\_148

### ***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 9 of the 13 non-draft genes in the pham.

Genes that call this "Most Annotated" start:

- Ariel\_144, Courthouse\_140, Hannaconda\_138, LittleE\_148, Omega\_152, Squiddly\_94, Squint\_139, Superphikiman\_142, Whitney\_93,

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

- Barb\_87, ClamChowder\_86, Danyall\_84, Fugax\_87,

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

•

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

Start 8:

- Found in 13 of 13 ( 100.0% ) of genes in pham
- Manual Annotations of this start: 9 of 13
- Called 69.2% of time when present
- Phage (with cluster) where this start called: Ariel\_144 (J), Courthouse\_140 (J), Hannaconda\_138 (J), LittleE\_148 (J), Omega\_152 (J), Squiddly\_94 (DN2), Squint\_139 (J), Superphikiman\_142 (J), Whitney\_93 (DN1),

Start 9:

- Found in 4 of 13 ( 30.8% ) of genes in pham

- Manual Annotations of this start: 4 of 13
- Called 100.0% of time when present
- Phage (with cluster) where this start called: Barb\_87 (DC1), ClamChowder\_86 (DC1), Danyall\_84 (DC1), Fugax\_87 (DC1),

### **Summary by clusters:**

There are 4 clusters represented in this pham: DN2, J, DN1, DC1,

Info for manual annotations of cluster DC1:

- Start number 9 was manually annotated 4 times for cluster DC1.

Info for manual annotations of cluster DN1:

- Start number 8 was manually annotated 1 time for cluster DN1.

Info for manual annotations of cluster DN2:

- Start number 8 was manually annotated 1 time for cluster DN2.

Info for manual annotations of cluster J:

- Start number 8 was manually annotated 7 times for cluster J.

### **Gene Information:**

Gene: Ariel\_144 Start: 73552, Stop: 73857, Start Num: 8

Candidate Starts for Ariel\_144:

(Start: 8 @73552 has 9 MA's), (13, 73735), (14, 73762), (17, 73804), (18, 73822),

Gene: Barb\_87 Start: 55893, Stop: 55591, Start Num: 9

Candidate Starts for Barb\_87:

(Start: 8 @55896 has 9 MA's), (Start: 9 @55893 has 4 MA's), (10, 55851), (11, 55797), (12, 55794), (15, 55656), (16, 55641), (18, 55620),

Gene: ClamChowder\_86 Start: 55893, Stop: 55591, Start Num: 9

Candidate Starts for ClamChowder\_86:

(Start: 8 @55896 has 9 MA's), (Start: 9 @55893 has 4 MA's), (10, 55851), (11, 55797), (12, 55794), (15, 55656), (16, 55641), (18, 55620),

Gene: Courthouse\_140 Start: 73410, Stop: 73715, Start Num: 8

Candidate Starts for Courthouse\_140:

(Start: 8 @73410 has 9 MA's), (13, 73593), (14, 73620), (17, 73662), (18, 73680),

Gene: Danyall\_84 Start: 55074, Stop: 54772, Start Num: 9

Candidate Starts for Danyall\_84:

(Start: 8 @55077 has 9 MA's), (Start: 9 @55074 has 4 MA's), (10, 55032), (11, 54978), (12, 54975), (15, 54837), (16, 54822), (18, 54801),

Gene: Fugax\_87 Start: 55885, Stop: 55583, Start Num: 9

Candidate Starts for Fugax\_87:

(Start: 8 @55888 has 9 MA's), (Start: 9 @55885 has 4 MA's), (10, 55843), (11, 55789), (12, 55786), (15, 55648), (16, 55633), (18, 55612),

Gene: Hannaconda\_138 Start: 74159, Stop: 74464, Start Num: 8  
Candidate Starts for Hannaconda\_138:  
(Start: 8 @74159 has 9 MA's), (13, 74342), (14, 74369), (17, 74411), (18, 74429),

Gene: LittleE\_148 Start: 76801, Stop: 77106, Start Num: 8  
Candidate Starts for LittleE\_148:  
(Start: 8 @76801 has 9 MA's), (13, 76984), (14, 77011), (17, 77053), (18, 77071),

Gene: Omega\_152 Start: 77879, Stop: 78184, Start Num: 8  
Candidate Starts for Omega\_152:  
(Start: 8 @77879 has 9 MA's), (13, 78062), (14, 78089), (17, 78131), (18, 78149),

Gene: Squiddly\_94 Start: 54160, Stop: 54468, Start Num: 8  
Candidate Starts for Squiddly\_94:  
(6, 54130), (7, 54142), (Start: 8 @54160 has 9 MA's), (16, 54415), (18, 54436),

Gene: Squint\_139 Start: 73950, Stop: 74255, Start Num: 8  
Candidate Starts for Squint\_139:  
(Start: 8 @73950 has 9 MA's), (13, 74133), (14, 74160), (17, 74202), (18, 74220),

Gene: Superphikiman\_142 Start: 73692, Stop: 73997, Start Num: 8  
Candidate Starts for Superphikiman\_142:  
(Start: 8 @73692 has 9 MA's), (13, 73875), (14, 73902), (17, 73944), (18, 73962),

Gene: Whitney\_93 Start: 52482, Stop: 52790, Start Num: 8  
Candidate Starts for Whitney\_93:  
(1, 52248), (2, 52278), (3, 52284), (4, 52407), (5, 52416), (6, 52449), (7, 52461), (Start: 8 @52482 has 9 MA's), (16, 52737), (18, 52758),