

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 9617 Report

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

Pham number 9617 has 13 members, 6 are drafts.

Phages represented in each track:

• Track 1 : Gray\_122, Aloki\_117, Schomber\_124, Kabocha\_127, Chidiebere\_126, Pakusa\_119, Oogie\_122, Hanem\_124, ChisanaKitsune\_124

Track 2 : FlyingTortilla\_123, ScarletRaider\_123

Track 3 : DalanDe\_109Track 4 : UBSmoodge\_125

## Summary of Final Annotations (See graph section above for start numbers):

The start number called the most often in the published annotations is 3, it was called in 6 of the 7 non-draft genes in the pham.

Genes that call this "Most Annotated" start:

• Aloki\_117, Chidiebere\_126, ChisanaKitsune\_124, Gray\_122, Hanem\_124, Kabocha\_127, Oogie\_122, Pakusa\_119, Schomber\_124,

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

FlyingTortilla\_123, ScarletRaider\_123, UBSmoodge\_125,

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

DalanDe\_109,

# Summary by start number:

### Start 2:

- Found in 1 of 13 (7.7%) of genes in pham
- Manual Annotations of this start: 1 of 7
- Called 100.0% of time when present
- Phage (with cluster) where this start called: DalanDe 109 (DQ).

### Start 3:

- Found in 12 of 13 (92.3%) of genes in pham
- Manual Annotations of this start: 6 of 7
- Called 75.0% of time when present

• Phage (with cluster) where this start called: Aloki\_117 (DQ), Chidiebere\_126 (DQ), ChisanaKitsune\_124 (DQ), Gray\_122 (DQ), Hanem\_124 (DQ), Kabocha\_127 (DQ), Oogie\_122 (DQ), Pakusa\_119 (DQ), Schomber\_124 (DQ),

#### Start 5:

- Found in 3 of 13 (23.1%) of genes in pham
- No Manual Annotations of this start.
- Called 100.0% of time when present
- Phage (with cluster) where this start called: FlyingTortilla\_123 (DQ), ScarletRaider\_123 (DQ), UBSmoodge\_125 (DQ),

### Summary by clusters:

There is one cluster represented in this pham: DQ

Info for manual annotations of cluster DQ:

- •Start number 2 was manually annotated 1 time for cluster DQ.
- •Start number 3 was manually annotated 6 times for cluster DQ.

## Gene Information:

Gene: Aloki 117 Start: 87107, Stop: 87628, Start Num: 3

Candidate Starts for Aloki\_117:

(Start: 3 @87107 has 6 MA's), (4, 87134), (10, 87227), (11, 87398), (16, 87497),

Gene: Chidiebere\_126 Start: 89010, Stop: 89531, Start Num: 3

Candidate Starts for Chidiebere 126:

(Start: 3 @89010 has 6 MA's), (4, 89037), (10, 89130), (11, 89301), (16, 89400),

Gene: ChisanaKitsune 124 Start: 87253, Stop: 87774, Start Num: 3

Candidate Starts for ChisanaKitsune 124:

(Start: 3 @87253 has 6 MA's), (4, 87280), (10, 87373), (11, 87544), (16, 87643),

Gene: DalanDe\_109 Start: 85215, Stop: 85697, Start Num: 2

Candidate Starts for DalanDe 109:

(Start: 2 @85215 has 1 MA's), (6, 85257), (7, 85275), (8, 85344), (9, 85356), (11, 85530), (12, 85539), (13, 85566),

Gene: FlyingTortilla 123 Start: 91022, Stop: 91498, Start Num: 5

Candidate Starts for FlyingTortilla 123:

(1, 90986), (Start: 3 @ 90989 has 6 MA's), (4, 91016), (5, 91022), (6, 91028), (10, 91109),

Gene: Gray\_122 Start: 87566, Stop: 88087, Start Num: 3

Candidate Starts for Gray\_122:

(Start: 3 @87566 has 6 MA's), (4, 87593), (10, 87686), (11, 87857), (16, 87956),

Gene: Hanem 124 Start: 87107, Stop: 87628, Start Num: 3

Candidate Starts for Hanem 124:

(Start: 3 @87107 has 6 MA's), (4, 87134), (10, 87227), (11, 87398), (16, 87497),

Gene: Kabocha\_127 Start: 89823, Stop: 90323, Start Num: 3

Candidate Starts for Kabocha\_127:

(Start: 3 @89823 has 6 MA's), (4, 89850), (10, 89943), (11, 90114), (16, 90213),

Gene: Oogie\_122 Start: 89531, Stop: 90052, Start Num: 3

Candidate Starts for Oogie\_122:

(Start: 3 @ 89531 has 6 MA's), (4, 89558), (10, 89651), (11, 89822), (16, 89921),

Gene: Pakusa\_119 Start: 87035, Stop: 87556, Start Num: 3

Candidate Starts for Pakusa\_119:

(Start: 3 @87035 has 6 MA's), (4, 87062), (10, 87155), (11, 87326), (16, 87425),

Gene: ScarletRaider\_123 Start: 90239, Stop: 90691, Start Num: 5

Candidate Starts for ScarletRaider\_123:

(1, 90203), (Start: 3 @ 90206 has 6 MA's), (4, 90233), (5, 90239), (6, 90245), (10, 90326),

Gene: Schomber\_124 Start: 88211, Stop: 88732, Start Num: 3

Candidate Starts for Schomber 124:

(Start: 3 @88211 has 6 MA's), (4, 88238), (10, 88331), (11, 88502), (16, 88601),

Gene: UBSmoodge\_125 Start: 90837, Stop: 91292, Start Num: 5

Candidate Starts for UBSmoodge\_125:

(1, 90801), (Start: 3 @90804 has 6 MA's), (4, 90831), (5, 90837), (6, 90843), (10, 90924), (14, 91164),

(15, 91185), (16, 91194), (17, 91209),