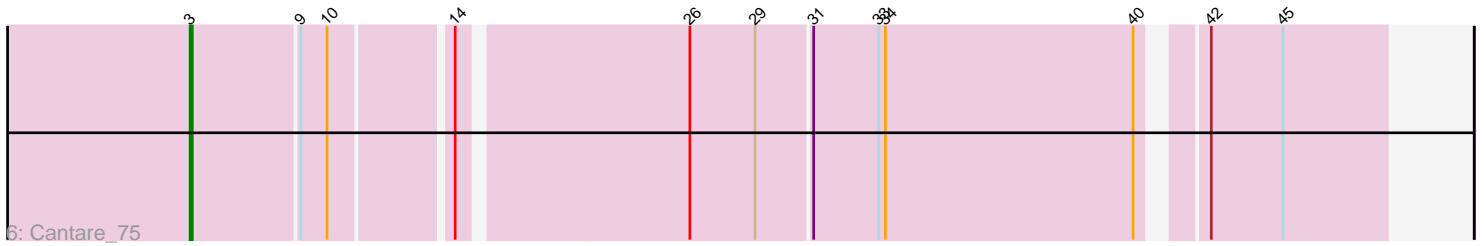
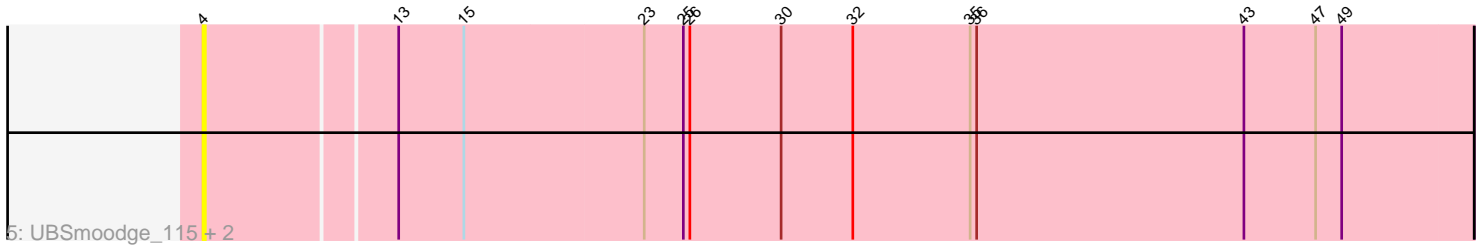
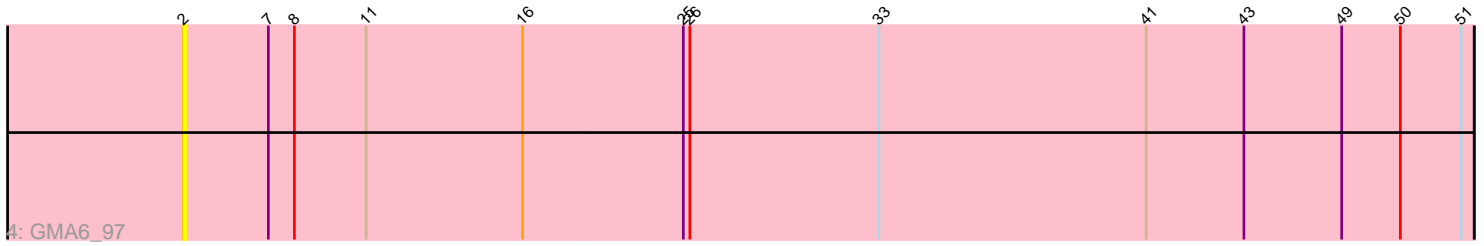
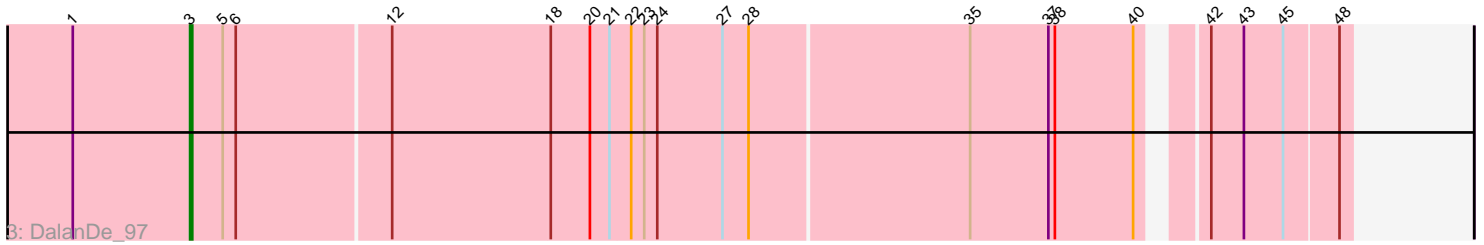
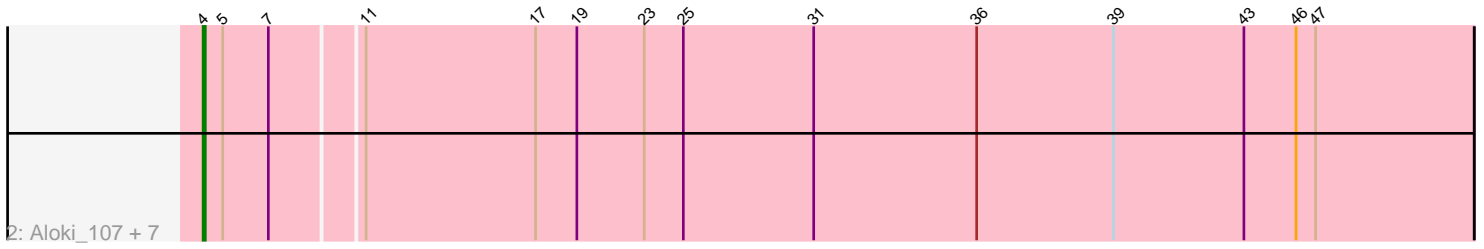
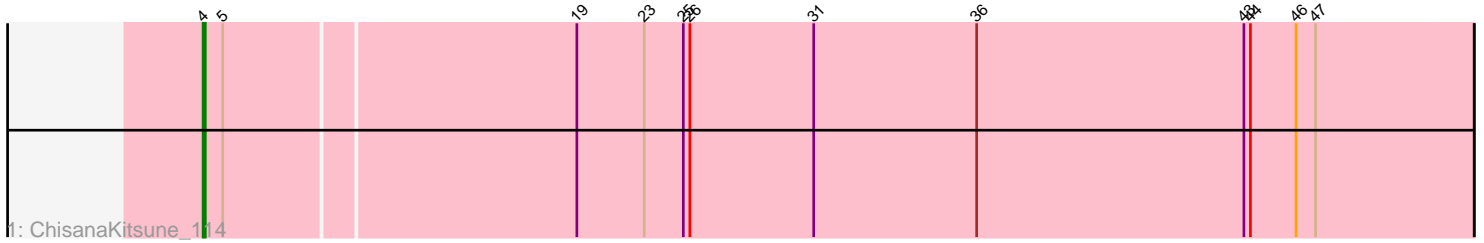


# Pham 159486



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 159486 Report

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

Pham number 159486 has 15 members, 7 are drafts.

Phages represented in each track:

- Track 1 : ChisanaKitsune\_114
- Track 2 : Alok\_107, Gray\_112, Chidiebere\_115, Pakusa\_109, Oogie\_110, Hanem\_114, Kabocha\_116, Schomber\_113
- Track 3 : DalanDe\_97
- Track 4 : GMA6\_97
- Track 5 : UBSmoodge\_115, FlyingTortilla\_113, ScarletRaider\_113
- Track 6 : Cantare\_75

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

The start number called the most often in the published annotations is 4, it was called in 6 of the 8 non-draft genes in the pham.

Genes that call this "Most Annotated" start:

- Alok\_107, Chidiebere\_115, ChisanaKitsune\_114, FlyingTortilla\_113, Gray\_112, Hanem\_114, Kabocha\_116, Oogie\_110, Pakusa\_109, ScarletRaider\_113, Schomber\_113, UBSmoodge\_115,

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

- 

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

- Cantare\_75, DalanDe\_97, GMA6\_97,

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

Start 2:

- Found in 1 of 15 ( 6.7% ) of genes in pham
- No Manual Annotations of this start.
- Called 100.0% of time when present
- Phage (with cluster) where this start called: GMA6\_97 (DQ),

Start 3:

- Found in 2 of 15 ( 13.3% ) of genes in pham

- Manual Annotations of this start: 2 of 8
- Called 100.0% of time when present
- Phage (with cluster) where this start called: Cantare\_75 (singleton), DalanDe\_97 (DQ),

Start 4:

- Found in 12 of 15 ( 80.0% ) of genes in pham
- Manual Annotations of this start: 6 of 8
- Called 100.0% of time when present
- Phage (with cluster) where this start called: Aloki\_107 (DQ), Chidiebere\_115 (DQ), ChisanaKitsune\_114 (DQ), FlyingTortilla\_113 (DQ), Gray\_112 (DQ), Hanem\_114 (DQ), Kabocha\_116 (DQ), Oogie\_110 (DQ), Pakusa\_109 (DQ), ScarletRaider\_113 (DQ), Schomber\_113 (DQ), UBSmoodge\_115 (DQ),

### Summary by clusters:

There are 2 clusters represented in this pham: singleton, DQ,

Info for manual annotations of cluster DQ:

- Start number 3 was manually annotated 1 time for cluster DQ.
- Start number 4 was manually annotated 6 times for cluster DQ.

### Gene Information:

Gene: Aloki\_107 Start: 78726, Stop: 79301, Start Num: 4

Candidate Starts for Aloki\_107:

(Start: 4 @78726 has 6 MA's), (5, 78735), (7, 78756), (11, 78795), (17, 78873), (19, 78891), (23, 78921), (25, 78939), (31, 78999), (36, 79074), (39, 79137), (43, 79197), (46, 79221), (47, 79230),

Gene: Cantare\_75 Start: 61854, Stop: 62366, Start Num: 3

Candidate Starts for Cantare\_75:

(Start: 3 @61854 has 2 MA's), (9, 61902), (10, 61914), (14, 61965), (26, 62064), (29, 62094), (31, 62118), (33, 62148), (34, 62151), (40, 62265), (42, 62286), (45, 62319),

Gene: Chidiebere\_115 Start: 79856, Stop: 80431, Start Num: 4

Candidate Starts for Chidiebere\_115:

(Start: 4 @79856 has 6 MA's), (5, 79865), (7, 79886), (11, 79925), (17, 80003), (19, 80021), (23, 80051), (25, 80069), (31, 80129), (36, 80204), (39, 80267), (43, 80327), (46, 80351), (47, 80360),

Gene: ChisanaKitsune\_114 Start: 78857, Stop: 79432, Start Num: 4

Candidate Starts for ChisanaKitsune\_114:

(Start: 4 @78857 has 6 MA's), (5, 78866), (19, 79022), (23, 79052), (25, 79070), (26, 79073), (31, 79130), (36, 79205), (43, 79328), (44, 79331), (46, 79352), (47, 79361),

Gene: DalanDe\_97 Start: 77200, Stop: 77709, Start Num: 3

Candidate Starts for DalanDe\_97:

(1, 77146), (Start: 3 @77200 has 2 MA's), (5, 77215), (6, 77221), (12, 77290), (18, 77362), (20, 77380), (21, 77389), (22, 77398), (23, 77404), (24, 77410), (27, 77440), (28, 77452), (35, 77551), (37, 77587), (38, 77590), (40, 77626), (42, 77647), (43, 77662), (45, 77680), (48, 77704),

Gene: FlyingTortilla\_113 Start: 82708, Stop: 83283, Start Num: 4

Candidate Starts for FlyingTortilla\_113:

(Start: 4 @82708 has 6 MA's), (13, 82792), (15, 82822), (23, 82903), (25, 82921), (26, 82924), (30, 82966), (32, 82999), (35, 83053), (36, 83056), (43, 83179), (47, 83212), (49, 83224),

Gene: GMA6\_97 Start: 70563, Stop: 71153, Start Num: 2

Candidate Starts for GMA6\_97:

(2, 70563), (7, 70602), (8, 70614), (11, 70647), (16, 70719), (25, 70791), (26, 70794), (33, 70881), (41, 71004), (43, 71049), (49, 71094), (50, 71121), (51, 71148),

Gene: Gray\_112 Start: 79189, Stop: 79764, Start Num: 4

Candidate Starts for Gray\_112:

(Start: 4 @79189 has 6 MA's), (5, 79198), (7, 79219), (11, 79258), (17, 79336), (19, 79354), (23, 79384), (25, 79402), (31, 79462), (36, 79537), (39, 79600), (43, 79660), (46, 79684), (47, 79693),

Gene: Hanem\_114 Start: 78726, Stop: 79301, Start Num: 4

Candidate Starts for Hanem\_114:

(Start: 4 @78726 has 6 MA's), (5, 78735), (7, 78756), (11, 78795), (17, 78873), (19, 78891), (23, 78921), (25, 78939), (31, 78999), (36, 79074), (39, 79137), (43, 79197), (46, 79221), (47, 79230),

Gene: Kabocha\_116 Start: 80669, Stop: 81244, Start Num: 4

Candidate Starts for Kabocha\_116:

(Start: 4 @80669 has 6 MA's), (5, 80678), (7, 80699), (11, 80738), (17, 80816), (19, 80834), (23, 80864), (25, 80882), (31, 80942), (36, 81017), (39, 81080), (43, 81140), (46, 81164), (47, 81173),

Gene: Oogie\_110 Start: 80689, Stop: 81264, Start Num: 4

Candidate Starts for Oogie\_110:

(Start: 4 @80689 has 6 MA's), (5, 80698), (7, 80719), (11, 80758), (17, 80836), (19, 80854), (23, 80884), (25, 80902), (31, 80962), (36, 81037), (39, 81100), (43, 81160), (46, 81184), (47, 81193),

Gene: Pakusa\_109 Start: 78654, Stop: 79229, Start Num: 4

Candidate Starts for Pakusa\_109:

(Start: 4 @78654 has 6 MA's), (5, 78663), (7, 78684), (11, 78723), (17, 78801), (19, 78819), (23, 78849), (25, 78867), (31, 78927), (36, 79002), (39, 79065), (43, 79125), (46, 79149), (47, 79158),

Gene: ScarletRaider\_113 Start: 81925, Stop: 82500, Start Num: 4

Candidate Starts for ScarletRaider\_113:

(Start: 4 @81925 has 6 MA's), (13, 82009), (15, 82039), (23, 82120), (25, 82138), (26, 82141), (30, 82183), (32, 82216), (35, 82270), (36, 82273), (43, 82396), (47, 82429), (49, 82441),

Gene: Schomber\_113 Start: 79057, Stop: 79632, Start Num: 4

Candidate Starts for Schomber\_113:

(Start: 4 @79057 has 6 MA's), (5, 79066), (7, 79087), (11, 79126), (17, 79204), (19, 79222), (23, 79252), (25, 79270), (31, 79330), (36, 79405), (39, 79468), (43, 79528), (46, 79552), (47, 79561),

Gene: UBSmoodge\_115 Start: 82495, Stop: 83070, Start Num: 4

Candidate Starts for UBSmoodge\_115:

(Start: 4 @82495 has 6 MA's), (13, 82579), (15, 82609), (23, 82690), (25, 82708), (26, 82711), (30, 82753), (32, 82786), (35, 82840), (36, 82843), (43, 82966), (47, 82999), (49, 83011),