



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 278661 Report

This analysis was run 02/07/26 on database version 634.

Pham number 278661 has 17 members, 3 are drafts.

Phages represented in each track:

- Track 1 : Momos\_4, Ellison17\_4, Prophecy\_4, Mimi16\_4, Grotle\_4
- Track 2 : Brave\_4, Squall\_4, Penne\_4, Fairywren\_4
- Track 3 : Trapezoid\_4, Reje\_4
- Track 4 : Razzleberry\_4
- Track 5 : Thatch\_4, Mantle\_4
- Track 6 : Transit\_4
- Track 7 : Pize\_4
- Track 8 : Ayka\_4

### ***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 14 of the 14 non-draft genes in the pham.

Genes that call this "Most Annotated" start:

- Ayka\_4, Brave\_4, Ellison17\_4, Fairywren\_4, Grotle\_4, Mantle\_4, Mimi16\_4, Momos\_4, Penne\_4, Pize\_4, Prophecy\_4, Razzleberry\_4, Reje\_4, Squall\_4, Thatch\_4, Transit\_4, Trapezoid\_4,

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 3:

- Found in 17 of 17 ( 100.0% ) of genes in pham
- Manual Annotations of this start: 14 of 14
- Called 100.0% of time when present
- Phage (with cluster) where this start called: Ayka\_4 (UNK), Brave\_4 (JB1), Ellison17\_4 (JB1), Fairywren\_4 (JB1), Grotle\_4 (JB1), Mantle\_4 (JB2), Mimi16\_4 (JB1), Momos\_4 (JB1), Penne\_4 (JB1), Pize\_4 (UNK), Prophecy\_4 (JB1),

Razzleberry\_4 (JB1), Reje\_4 (UNK), Squall\_4 (JB1), Thatch\_4 (JB2), Transit\_4 (JB2), Trapezoid\_4 (JB1),

### **Summary by clusters:**

There are 3 clusters represented in this pham: UNK, JB2, JB1,

Info for manual annotations of cluster JB1:

- Start number 3 was manually annotated 11 times for cluster JB1.

Info for manual annotations of cluster JB2:

- Start number 3 was manually annotated 3 times for cluster JB2.

### **Gene Information:**

Gene: Ayka\_4 Start: 3953, Stop: 4189, Start Num: 3

Candidate Starts for Ayka\_4:

(Start: 3 @3953 has 14 MA's), (4, 4061), (5, 4103),

Gene: Brave\_4 Start: 3867, Stop: 4109, Start Num: 3

Candidate Starts for Brave\_4:

(Start: 3 @3867 has 14 MA's), (7, 4050),

Gene: Ellison17\_4 Start: 3732, Stop: 3974, Start Num: 3

Candidate Starts for Ellison17\_4:

(Start: 3 @3732 has 14 MA's), (9, 3927),

Gene: Fairywren\_4 Start: 3836, Stop: 4078, Start Num: 3

Candidate Starts for Fairywren\_4:

(Start: 3 @3836 has 14 MA's), (7, 4019),

Gene: Grotle\_4 Start: 3691, Stop: 3933, Start Num: 3

Candidate Starts for Grotle\_4:

(Start: 3 @3691 has 14 MA's), (9, 3886),

Gene: Mantle\_4 Start: 4363, Stop: 4614, Start Num: 3

Candidate Starts for Mantle\_4:

(2, 4339), (Start: 3 @4363 has 14 MA's),

Gene: Mimi16\_4 Start: 3742, Stop: 3984, Start Num: 3

Candidate Starts for Mimi16\_4:

(Start: 3 @3742 has 14 MA's), (9, 3937),

Gene: Momos\_4 Start: 3732, Stop: 3974, Start Num: 3

Candidate Starts for Momos\_4:

(Start: 3 @3732 has 14 MA's), (9, 3927),

Gene: Penne\_4 Start: 3870, Stop: 4112, Start Num: 3

Candidate Starts for Penne\_4:

(Start: 3 @3870 has 14 MA's), (7, 4053),

Gene: Pize\_4 Start: 3685, Stop: 3936, Start Num: 3  
Candidate Starts for Pize\_4:  
(Start: 3 @3685 has 14 MA's), (8, 3880),

Gene: Prophecy\_4 Start: 3742, Stop: 3984, Start Num: 3  
Candidate Starts for Prophecy\_4:  
(Start: 3 @3742 has 14 MA's), (9, 3937),

Gene: Razzleberry\_4 Start: 3717, Stop: 3950, Start Num: 3  
Candidate Starts for Razzleberry\_4:  
(Start: 3 @3717 has 14 MA's), (6, 3888), (9, 3906),

Gene: Reje\_4 Start: 3729, Stop: 3959, Start Num: 3  
Candidate Starts for Reje\_4:  
(Start: 3 @3729 has 14 MA's), (5, 3873), (10, 3915),

Gene: Squall\_4 Start: 3837, Stop: 4079, Start Num: 3  
Candidate Starts for Squall\_4:  
(Start: 3 @3837 has 14 MA's), (7, 4020),

Gene: Thatch\_4 Start: 4160, Stop: 4396, Start Num: 3  
Candidate Starts for Thatch\_4:  
(2, 4136), (Start: 3 @4160 has 14 MA's),

Gene: Transit\_4 Start: 4101, Stop: 4337, Start Num: 3  
Candidate Starts for Transit\_4:  
(1, 3996), (2, 4077), (Start: 3 @4101 has 14 MA's),

Gene: Trapezoid\_4 Start: 3762, Stop: 3992, Start Num: 3  
Candidate Starts for Trapezoid\_4:  
(Start: 3 @3762 has 14 MA's), (5, 3906), (10, 3948),