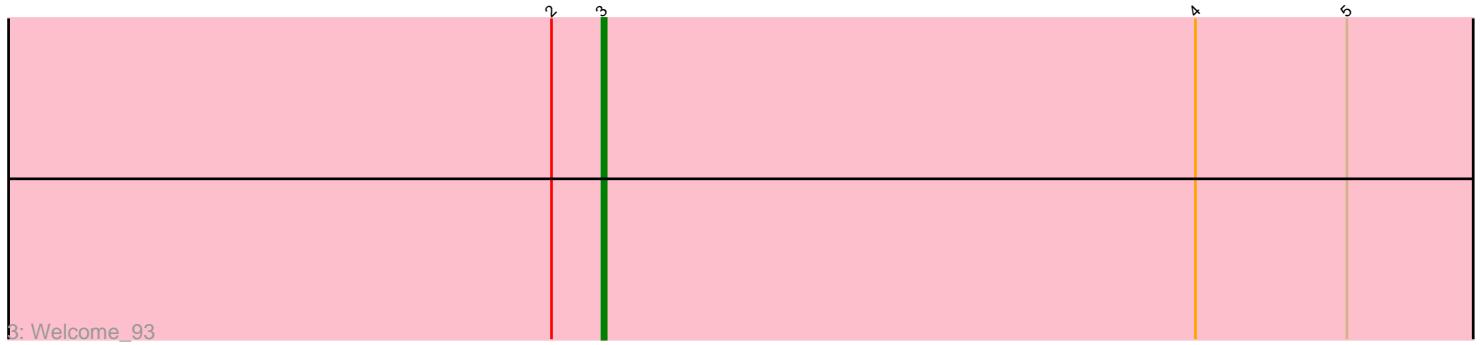
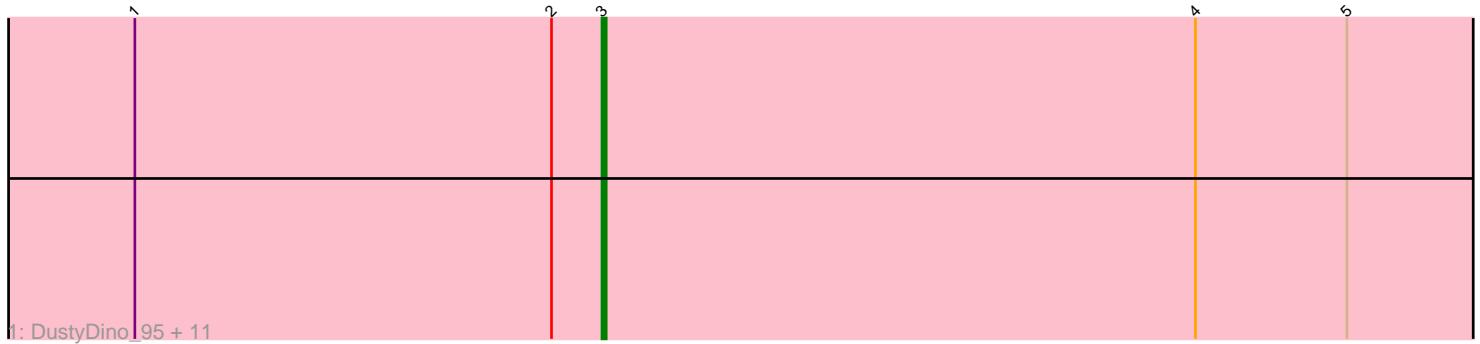


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Note: Tracks are now grouped by subcluster and scaled. Switching in subcluster is indicated by changes in track color. Track scale is now set by default to display the region 30 bp upstream of start 1 to 30 bp downstream of the last possible start. 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 276659 Report

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

Pham number 276659 has 16 members, 5 are drafts.

Phages represented in each track:

- Track 1 : DustyDino\_95, HollowPurple\_92, Lyell\_91, ASegato\_89, Yuma\_90, RunningBrook\_93, Fork\_87, Deschain\_92, Casablancas\_92, Erenyeager\_91, Necrophoxinus\_93, SteakFry\_90
- Track 2 : Musetta\_90, Shroomer\_94
- Track 3 : Welcome\_93
- Track 4 : Issa7\_90

### ***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 11 of the 11 non-draft genes in the pham.

Genes that call this "Most Annotated" start:

- ASegato\_89, Casablancas\_92, Deschain\_92, DustyDino\_95, Erenyeager\_91, Fork\_87, HollowPurple\_92, Lyell\_91, Musetta\_90, Necrophoxinus\_93, RunningBrook\_93, Shroomer\_94, SteakFry\_90, Welcome\_93, Yuma\_90,

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

- Issa7\_90,

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

- 

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

Start 2:

- Found in 16 of 16 ( 100.0% ) of genes in pham
- No Manual Annotations of this start.
- Called 6.2% of time when present
- Phage (with cluster) where this start called: Issa7\_90 (ED2),

Start 3:

- Found in 16 of 16 ( 100.0% ) of genes in pham
- Manual Annotations of this start: 11 of 11

- Called 93.8% of time when present
- Phage (with cluster) where this start called: ASegato\_89 (ED2), Casablancas\_92 (ED2), Deschain\_92 (ED2), DustyDino\_95 (ED2), Erenyeager\_91 (ED2), Fork\_87 (ED2), HollowPurple\_92 (ED2), Lyell\_91 (ED2), Musetta\_90 (ED2), Necrophoxinus\_93 (ED2), RunningBrook\_93 (ED2), Shroomer\_94 (ED2), SteakFry\_90 (ED2), Welcome\_93 (ED2), Yuma\_90 (ED2),

### **Summary by clusters:**

There is one cluster represented in this pham: ED2

Info for manual annotations of cluster ED2:

- Start number 3 was manually annotated 11 times for cluster ED2.

### **Gene Information:**

Gene: ASegato\_89 Start: 51352, Stop: 51128, Start Num: 3

Candidate Starts for ASegato\_89:

(1, 51463), (2, 51364), (Start: 3 @51352 has 11 MA's), (4, 51211), (5, 51175),

Gene: Casablancas\_92 Start: 51268, Stop: 51044, Start Num: 3

Candidate Starts for Casablancas\_92:

(1, 51379), (2, 51280), (Start: 3 @51268 has 11 MA's), (4, 51127), (5, 51091),

Gene: Deschain\_92 Start: 51932, Stop: 51708, Start Num: 3

Candidate Starts for Deschain\_92:

(1, 52043), (2, 51944), (Start: 3 @51932 has 11 MA's), (4, 51791), (5, 51755),

Gene: DustyDino\_95 Start: 52491, Stop: 52267, Start Num: 3

Candidate Starts for DustyDino\_95:

(1, 52602), (2, 52503), (Start: 3 @52491 has 11 MA's), (4, 52350), (5, 52314),

Gene: Erenyeager\_91 Start: 51280, Stop: 51056, Start Num: 3

Candidate Starts for Erenyeager\_91:

(1, 51391), (2, 51292), (Start: 3 @51280 has 11 MA's), (4, 51139), (5, 51103),

Gene: Fork\_87 Start: 51230, Stop: 51006, Start Num: 3

Candidate Starts for Fork\_87:

(1, 51341), (2, 51242), (Start: 3 @51230 has 11 MA's), (4, 51089), (5, 51053),

Gene: HollowPurple\_92 Start: 51789, Stop: 51565, Start Num: 3

Candidate Starts for HollowPurple\_92:

(1, 51900), (2, 51801), (Start: 3 @51789 has 11 MA's), (4, 51648), (5, 51612),

Gene: Issa7\_90 Start: 51257, Stop: 51021, Start Num: 2

Candidate Starts for Issa7\_90:

(1, 51356), (2, 51257), (Start: 3 @51245 has 11 MA's), (4, 51104), (5, 51068),

Gene: Lyell\_91 Start: 51441, Stop: 51217, Start Num: 3

Candidate Starts for Lyell\_91:

(1, 51552), (2, 51453), (Start: 3 @51441 has 11 MA's), (4, 51300), (5, 51264),

Gene: Musetta\_90 Start: 51780, Stop: 51556, Start Num: 3

Candidate Starts for Musetta\_90:

(2, 51792), (Start: 3 @51780 has 11 MA's), (4, 51639), (5, 51603),

Gene: Necrophoxinus\_93 Start: 52127, Stop: 51903, Start Num: 3

Candidate Starts for Necrophoxinus\_93:

(1, 52238), (2, 52139), (Start: 3 @52127 has 11 MA's), (4, 51986), (5, 51950),

Gene: RunningBrook\_93 Start: 52491, Stop: 52267, Start Num: 3

Candidate Starts for RunningBrook\_93:

(1, 52602), (2, 52503), (Start: 3 @52491 has 11 MA's), (4, 52350), (5, 52314),

Gene: Shroomer\_94 Start: 51731, Stop: 51507, Start Num: 3

Candidate Starts for Shroomer\_94:

(2, 51743), (Start: 3 @51731 has 11 MA's), (4, 51590), (5, 51554),

Gene: SteakFry\_90 Start: 51789, Stop: 51565, Start Num: 3

Candidate Starts for SteakFry\_90:

(1, 51900), (2, 51801), (Start: 3 @51789 has 11 MA's), (4, 51648), (5, 51612),

Gene: Welcome\_93 Start: 51943, Stop: 51719, Start Num: 3

Candidate Starts for Welcome\_93:

(2, 51955), (Start: 3 @51943 has 11 MA's), (4, 51802), (5, 51766),

Gene: Yuma\_90 Start: 51452, Stop: 51228, Start Num: 3

Candidate Starts for Yuma\_90:

(1, 51563), (2, 51464), (Start: 3 @51452 has 11 MA's), (4, 51311), (5, 51275),