



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 295008 Report

This analysis was run 04/18/26 on database version 643.

Pham number 295008 has 27 members, 5 are drafts.

Phages represented in each track:

- Track 1 : Platte\_5, Tandem\_5, Platte\_115, Tandem\_116
- Track 2 : Welcome\_7, Necrophoxinus\_123, RunningBrook\_122, HollowPurple\_123, Casablanacas\_5, Deschain\_6, Deschain\_120, StevieWelch\_7, Necrophoxinus\_7, DustyDino\_125, DustyDino\_7, SteakFry\_6, Casablanacas\_120, Musetta\_120, Yuma\_120, RunningBrook\_6, Welcome\_124, Musetta\_7, HollowPurple\_7, StevieWelch\_125, Yuma\_7
- Track 3 : ASegato\_7, ASegato\_121

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

The start number called the most often in the published annotations is 2, it was called in 22 of the 22 non-draft genes in the pham.

Genes that call this "Most Annotated" start:

- ASegato\_121, ASegato\_7, Casablanacas\_120, Casablanacas\_5, Deschain\_120, Deschain\_6, DustyDino\_125, DustyDino\_7, HollowPurple\_123, HollowPurple\_7, Musetta\_120, Musetta\_7, Necrophoxinus\_123, Necrophoxinus\_7, Platte\_115, Platte\_5, RunningBrook\_122, RunningBrook\_6, SteakFry\_6, StevieWelch\_125, StevieWelch\_7, Tandem\_116, Tandem\_5, Welcome\_124, Welcome\_7, Yuma\_120, Yuma\_7,

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 2:

- Found in 27 of 27 ( 100.0% ) of genes in pham
- Manual Annotations of this start: 22 of 22
- Called 100.0% of time when present

- Phage (with cluster) where this start called: ASegato\_121 (ED2), ASegato\_7 (ED2), Casablanacas\_120 (ED2), Casablanacas\_5 (ED2), Deschain\_120 (ED2), Deschain\_6 (ED2), DustyDino\_125 (ED2), DustyDino\_7 (ED2), HollowPurple\_123 (ED2), HollowPurple\_7 (ED2), Musetta\_120 (ED2), Musetta\_7 (ED2), Necrophoxinus\_123 (ED2), Necrophoxinus\_7 (ED2), Platte\_115 (ED1), Platte\_5 (ED1), RunningBrook\_122 (ED2), RunningBrook\_6 (ED2), SteakFry\_6 (ED2), StevieWelch\_125 (ED2), StevieWelch\_7 (ED2), Tandem\_116 (ED1), Tandem\_5 (ED1), Welcome\_124 (ED2), Welcome\_7 (ED2), Yuma\_120 (ED2), Yuma\_7 (ED2),

### **Summary by clusters:**

There are 2 clusters represented in this pham: ED2, ED1,

Info for manual annotations of cluster ED1:

- Start number 2 was manually annotated 4 times for cluster ED1.

Info for manual annotations of cluster ED2:

- Start number 2 was manually annotated 18 times for cluster ED2.

### **Gene Information:**

Gene: ASegato\_7 Start: 2656, Stop: 2270, Start Num: 2

Candidate Starts for ASegato\_7:

(Start: 2 @2656 has 22 MA's), (3, 2599), (4, 2482), (5, 2461), (7, 2341), (8, 2308),

Gene: ASegato\_121 Start: 62105, Stop: 61719, Start Num: 2

Candidate Starts for ASegato\_121:

(Start: 2 @62105 has 22 MA's), (3, 62048), (4, 61931), (5, 61910), (7, 61790), (8, 61757),

Gene: Casablanacas\_5 Start: 2450, Stop: 2064, Start Num: 2

Candidate Starts for Casablanacas\_5:

(Start: 2 @2450 has 22 MA's), (3, 2393), (4, 2276), (7, 2135), (8, 2102),

Gene: Casablanacas\_120 Start: 61467, Stop: 61081, Start Num: 2

Candidate Starts for Casablanacas\_120:

(Start: 2 @61467 has 22 MA's), (3, 61410), (4, 61293), (7, 61152), (8, 61119),

Gene: Deschain\_6 Start: 2556, Stop: 2170, Start Num: 2

Candidate Starts for Deschain\_6:

(Start: 2 @2556 has 22 MA's), (3, 2499), (4, 2382), (7, 2241), (8, 2208),

Gene: Deschain\_120 Start: 62137, Stop: 61751, Start Num: 2

Candidate Starts for Deschain\_120:

(Start: 2 @62137 has 22 MA's), (3, 62080), (4, 61963), (7, 61822), (8, 61789),

Gene: DustyDino\_125 Start: 62782, Stop: 62396, Start Num: 2

Candidate Starts for DustyDino\_125:

(Start: 2 @62782 has 22 MA's), (3, 62725), (4, 62608), (7, 62467), (8, 62434),

Gene: DustyDino\_7 Start: 2692, Stop: 2306, Start Num: 2

Candidate Starts for DustyDino\_7:

(Start: 2 @2692 has 22 MA's), (3, 2635), (4, 2518), (7, 2377), (8, 2344),

Gene: HollowPurple\_123 Start: 62727, Stop: 62341, Start Num: 2

Candidate Starts for HollowPurple\_123:

(Start: 2 @62727 has 22 MA's), (3, 62670), (4, 62553), (7, 62412), (8, 62379),

Gene: HollowPurple\_7 Start: 2654, Stop: 2268, Start Num: 2

Candidate Starts for HollowPurple\_7:

(Start: 2 @2654 has 22 MA's), (3, 2597), (4, 2480), (7, 2339), (8, 2306),

Gene: Musetta\_120 Start: 62463, Stop: 62077, Start Num: 2

Candidate Starts for Musetta\_120:

(Start: 2 @62463 has 22 MA's), (3, 62406), (4, 62289), (7, 62148), (8, 62115),

Gene: Musetta\_7 Start: 2668, Stop: 2282, Start Num: 2

Candidate Starts for Musetta\_7:

(Start: 2 @2668 has 22 MA's), (3, 2611), (4, 2494), (7, 2353), (8, 2320),

Gene: Necrophoxinus\_123 Start: 62723, Stop: 62337, Start Num: 2

Candidate Starts for Necrophoxinus\_123:

(Start: 2 @62723 has 22 MA's), (3, 62666), (4, 62549), (7, 62408), (8, 62375),

Gene: Necrophoxinus\_7 Start: 2480, Stop: 2094, Start Num: 2

Candidate Starts for Necrophoxinus\_7:

(Start: 2 @2480 has 22 MA's), (3, 2423), (4, 2306), (7, 2165), (8, 2132),

Gene: Platte\_5 Start: 2255, Stop: 1875, Start Num: 2

Candidate Starts for Platte\_5:

(1, 2285), (Start: 2 @2255 has 22 MA's), (4, 2084), (6, 1973), (8, 1913), (9, 1907),

Gene: Platte\_115 Start: 61803, Stop: 61423, Start Num: 2

Candidate Starts for Platte\_115:

(1, 61833), (Start: 2 @61803 has 22 MA's), (4, 61632), (6, 61521), (8, 61461), (9, 61455),

Gene: RunningBrook\_122 Start: 62782, Stop: 62396, Start Num: 2

Candidate Starts for RunningBrook\_122:

(Start: 2 @62782 has 22 MA's), (3, 62725), (4, 62608), (7, 62467), (8, 62434),

Gene: RunningBrook\_6 Start: 2692, Stop: 2306, Start Num: 2

Candidate Starts for RunningBrook\_6:

(Start: 2 @2692 has 22 MA's), (3, 2635), (4, 2518), (7, 2377), (8, 2344),

Gene: SteakFry\_6 Start: 2654, Stop: 2268, Start Num: 2

Candidate Starts for SteakFry\_6:

(Start: 2 @2654 has 22 MA's), (3, 2597), (4, 2480), (7, 2339), (8, 2306),

Gene: StevieWelch\_7 Start: 2553, Stop: 2167, Start Num: 2

Candidate Starts for StevieWelch\_7:

(Start: 2 @2553 has 22 MA's), (3, 2496), (4, 2379), (7, 2238), (8, 2205),

Gene: StevieWelch\_125 Start: 62799, Stop: 62413, Start Num: 2

Candidate Starts for StevieWelch\_125:

(Start: 2 @62799 has 22 MA's), (3, 62742), (4, 62625), (7, 62484), (8, 62451),

Gene: Tandem\_5 Start: 2360, Stop: 1980, Start Num: 2

Candidate Starts for Tandem\_5:

(1, 2390), (Start: 2 @2360 has 22 MA's), (4, 2189), (6, 2078), (8, 2018), (9, 2012),

Gene: Tandem\_116 Start: 62203, Stop: 61823, Start Num: 2

Candidate Starts for Tandem\_116:

(1, 62233), (Start: 2 @62203 has 22 MA's), (4, 62032), (6, 61921), (8, 61861), (9, 61855),

Gene: Welcome\_7 Start: 2667, Stop: 2281, Start Num: 2

Candidate Starts for Welcome\_7:

(Start: 2 @2667 has 22 MA's), (3, 2610), (4, 2493), (7, 2352), (8, 2319),

Gene: Welcome\_124 Start: 62811, Stop: 62425, Start Num: 2

Candidate Starts for Welcome\_124:

(Start: 2 @62811 has 22 MA's), (3, 62754), (4, 62637), (7, 62496), (8, 62463),

Gene: Yuma\_120 Start: 61614, Stop: 61228, Start Num: 2

Candidate Starts for Yuma\_120:

(Start: 2 @61614 has 22 MA's), (3, 61557), (4, 61440), (7, 61299), (8, 61266),

Gene: Yuma\_7 Start: 2563, Stop: 2177, Start Num: 2

Candidate Starts for Yuma\_7:

(Start: 2 @2563 has 22 MA's), (3, 2506), (4, 2389), (7, 2248), (8, 2215),