

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 205875 Report

This analysis was run 02/22/25 on database version 588.

Pham number 205875 has 8 members, 1 are drafts.

Phages represented in each track:

Track 1 : Daddyjeff_61Track 2 : Pippin15_62

Track 3 : AddiRose_60

Track 4 : Sweetclover_59

• Track 5 : Yotsuba 61

• Track 6 : Serenabean_60

Track 7 : Eevee_61Track 8 : JoyLin_61

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

The start number called the most often in the published annotations is 8, it was called in 4 of the 7 non-draft genes in the pham.

Genes that call this "Most Annotated" start:

Eevee_61, JoyLin_61, Pippin15_62, Yotsuba_61,

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

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

AddiRose_60, Daddyjeff_61, Serenabean_60, Sweetclover_59,

Summary by start number:

Start 7:

- Found in 4 of 8 (50.0%) of genes in pham
- Manual Annotations of this start: 3 of 7
- Called 100.0% of time when present
- Phage (with cluster) where this start called: AddiRose_60 (JA), Daddyjeff_61 (JA), Serenabean_60 (JA), Sweetclover_59 (JA),

Start 8:

• Found in 4 of 8 (50.0%) of genes in pham

- Manual Annotations of this start: 4 of 7
- Called 100.0% of time when present
- Phage (with cluster) where this start called: Eevee_61 (JA), JoyLin_61 (JA), Pippin15_62 (JA), Yotsuba_61 (JA),

Summary by clusters:

There is one cluster represented in this pham: JA

Info for manual annotations of cluster JA:

- •Start number 7 was manually annotated 3 times for cluster JA.
- Start number 8 was manually annotated 4 times for cluster JA.

Gene Information:

Gene: AddiRose 60 Start: 40973, Stop: 41491, Start Num: 7

Candidate Starts for AddiRose_60:

(Start: 7 @40973 has 3 MA's), (9, 41051), (10, 41090), (11, 41138), (12, 41189), (18, 41306), (20, 41351), (22, 41408),

Gene: Daddyjeff_61 Start: 41194, Stop: 41703, Start Num: 7

Candidate Starts for Daddyjeff_61:

(5, 40825), (Start: 7 @41194 has 3 MA's), (9, 41272), (10, 41311), (11, 41359), (12, 41410), (13, 41452), (20, 41572),

Gene: Eevee_61 Start: 44507, Stop: 45004, Start Num: 8

Candidate Starts for Eevee_61:

(6, 44414), (Start: 8 @44507 has 4 MA's), (12, 44717), (15, 44783), (16, 44795), (21, 44912), (23, 44936), (24, 44957),

Gene: JoyLin 61 Start: 44627, Stop: 45124, Start Num: 8

Candidate Starts for JoyLin 61:

(6, 44534), (Start: 8 @44627 has 4 MA's), (12, 44837), (15, 44903), (16, 44915), (19, 44972), (21, 45032), (24, 45077),

Gene: Pippin15 62 Start: 41542, Stop: 42039, Start Num: 8

Candidate Starts for Pippin15 62:

(6, 41446), (Start: 8 @41542 has 4 MA's), (12, 41752), (14, 41797), (16, 41830), (19, 41887), (21, 41947), (23, 41971), (24, 41992),

Gene: Serenabean 60 Start: 41143, Stop: 41652, Start Num: 7

Candidate Starts for Serenabean 60:

(1, 40474), (2, 40588), (3, 40669), (4, 40690), (Start: 7 @41143 has 3 MA's), (9, 41221), (10, 41260), (11, 41308), (12, 41359), (13, 41401), (20, 41521),

Gene: Sweetclover_59 Start: 40126, Stop: 40638, Start Num: 7

Candidate Starts for Sweetclover 59:

(6, 40039), (Start: 7 @ 40126 has 3 MA's), (10, 40243), (12, 40342), (17, 40435),

Gene: Yotsuba 61 Start: 44645, Stop: 45142, Start Num: 8

Candidate Starts for Yotsuba_61:

(6, 44549), (Start: 8 @44645 has 4 MA's), (12, 44855), (15, 44921), (16, 44933), (21, 45050), (23, 45074), (24, 45095),