



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 170290 Report

This analysis was run 07/09/24 on database version 566.

Pham number 170290 has 23 members, 1 are drafts.

Phages represented in each track:

- Track 1 : JF4_55, MK4_57, JF2_55, Isca_56, Phantastic_58
- Track 2 : Noella_56, Lambert1_56, Norbert_55, ResDef_57, Todacoro_56, QuinnKiro_55, Margo_56, Popcicle_56, Texage_55, Caviar_56, Hookmount_56, Veracruz_55, Pocahontas_56, Panamaxus_54, MA5_57
- Track 3 : Heathen_57, Scout_57, HelDan_57

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 22 of the 22 non-draft genes in the pham.

Genes that call this "Most Annotated" start:

- Caviar_56, Heathen_57, HelDan_57, Hookmount_56, Isca_56, JF2_55, JF4_55, Lambert1_56, MA5_57, MK4_57, Margo_56, Noella_56, Norbert_55, Panamaxus_54, Phantastic_58, Pocahontas_56, Popcicle_56, QuinnKiro_55, ResDef_57, Scout_57, Texage_55, Todacoro_56, Veracruz_55,

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 23 of 23 (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: Caviar_56 (A3), Heathen_57 (A3), HelDan_57 (A3), Hookmount_56 (A3), Isca_56 (A3), JF2_55 (A3), JF4_55 (A3), Lambert1_56 (A3), MA5_57 (A3), MK4_57 (A3), Margo_56 (A3), Noella_56 (A3), Norbert_55 (A3), Panamaxus_54 (A3), Phantastic_58 (A3), Pocahontas_56 (A3), Popcicle_56 (A3), QuinnKiro_55 (A3), ResDef_57 (A3), Scout_57 (A3), Texage_55

(A3), Todacoro_56 (A3), Veracruz_55 (A3),

Summary by clusters:

There is one cluster represented in this pham: A3

Info for manual annotations of cluster A3:

•Start number 3 was manually annotated 22 times for cluster A3.

Gene Information:

Gene: Caviar_56 Start: 37855, Stop: 37757, Start Num: 3

Candidate Starts for Caviar_56:

(Start: 3 @37855 has 22 MA's), (4, 37837),

Gene: Heathen_57 Start: 37581, Stop: 37489, Start Num: 3

Candidate Starts for Heathen_57:

(Start: 3 @37581 has 22 MA's), (4, 37563),

Gene: HelDan_57 Start: 37797, Stop: 37705, Start Num: 3

Candidate Starts for HelDan_57:

(Start: 3 @37797 has 22 MA's), (4, 37779),

Gene: Hookmount_56 Start: 37856, Stop: 37758, Start Num: 3

Candidate Starts for Hookmount_56:

(Start: 3 @37856 has 22 MA's), (4, 37838),

Gene: Isca_56 Start: 37366, Stop: 37271, Start Num: 3

Candidate Starts for Isca_56:

(1, 37609), (2, 37570), (Start: 3 @37366 has 22 MA's), (4, 37348),

Gene: JF2_55 Start: 36067, Stop: 35972, Start Num: 3

Candidate Starts for JF2_55:

(1, 36310), (2, 36271), (Start: 3 @36067 has 22 MA's), (4, 36049),

Gene: JF4_55 Start: 36067, Stop: 35972, Start Num: 3

Candidate Starts for JF4_55:

(1, 36310), (2, 36271), (Start: 3 @36067 has 22 MA's), (4, 36049),

Gene: Lambert1_56 Start: 37855, Stop: 37757, Start Num: 3

Candidate Starts for Lambert1_56:

(Start: 3 @37855 has 22 MA's), (4, 37837),

Gene: MA5_57 Start: 37267, Stop: 37175, Start Num: 3

Candidate Starts for MA5_57:

(Start: 3 @37267 has 22 MA's), (4, 37249),

Gene: MK4_57 Start: 37351, Stop: 37256, Start Num: 3

Candidate Starts for MK4_57:

(1, 37594), (2, 37555), (Start: 3 @37351 has 22 MA's), (4, 37333),

Gene: Margo_56 Start: 37881, Stop: 37783, Start Num: 3
Candidate Starts for Margo_56:
(Start: 3 @37881 has 22 MA's), (4, 37863),

Gene: Noella_56 Start: 37856, Stop: 37758, Start Num: 3
Candidate Starts for Noella_56:
(Start: 3 @37856 has 22 MA's), (4, 37838),

Gene: Norbert_55 Start: 37855, Stop: 37757, Start Num: 3
Candidate Starts for Norbert_55:
(Start: 3 @37855 has 22 MA's), (4, 37837),

Gene: Panamaxus_54 Start: 37855, Stop: 37757, Start Num: 3
Candidate Starts for Panamaxus_54:
(Start: 3 @37855 has 22 MA's), (4, 37837),

Gene: Phantastic_58 Start: 37528, Stop: 37433, Start Num: 3
Candidate Starts for Phantastic_58:
(1, 37771), (2, 37732), (Start: 3 @37528 has 22 MA's), (4, 37510),

Gene: Pocahontas_56 Start: 37852, Stop: 37754, Start Num: 3
Candidate Starts for Pocahontas_56:
(Start: 3 @37852 has 22 MA's), (4, 37834),

Gene: Popcicle_56 Start: 37852, Stop: 37754, Start Num: 3
Candidate Starts for Popcicle_56:
(Start: 3 @37852 has 22 MA's), (4, 37834),

Gene: QuinnKiro_55 Start: 37855, Stop: 37757, Start Num: 3
Candidate Starts for QuinnKiro_55:
(Start: 3 @37855 has 22 MA's), (4, 37837),

Gene: ResDef_57 Start: 37855, Stop: 37757, Start Num: 3
Candidate Starts for ResDef_57:
(Start: 3 @37855 has 22 MA's), (4, 37837),

Gene: Scout_57 Start: 37044, Stop: 36952, Start Num: 3
Candidate Starts for Scout_57:
(Start: 3 @37044 has 22 MA's), (4, 37026),

Gene: Texage_55 Start: 37856, Stop: 37758, Start Num: 3
Candidate Starts for Texage_55:
(Start: 3 @37856 has 22 MA's), (4, 37838),

Gene: Todacoro_56 Start: 37855, Stop: 37757, Start Num: 3
Candidate Starts for Todacoro_56:
(Start: 3 @37855 has 22 MA's), (4, 37837),

Gene: Veracruz_55 Start: 37855, Stop: 37757, Start Num: 3
Candidate Starts for Veracruz_55:
(Start: 3 @37855 has 22 MA's), (4, 37837),