

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 171984 Report

This analysis was run 07/10/24 on database version 566.

Pham number 171984 has 9 members, 0 are drafts.

Phages represented in each track:

- Track 1 : Peregrin_19
- Track 2 : Grayson_26
- Track 3 : Grayson_21
- Track 4 : Peregrin_20
- Track 5 : Peregrin 24
- Track 6 : Weasels 2_18
- Track 7 : Weasels2 22
- Track 8 : Grayson_20
- Track 9 : Success 26

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

The start number called the most often in the published annotations is 9, it was called in 5 of the 9 non-draft genes in the pham.

Genes that call this "Most Annotated" start:

• Grayson_20, Grayson_21, Peregrin_19, Peregrin_20, Weasels2_18,

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

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

• Grayson_26, Peregrin_24, Success_26, Weasels2_22,

Summary by start number:

Start 3:

- Found in 1 of 9 (11.1%) of genes in pham
- Manual Annotations of this start: 1 of 9
- Called 100.0% of time when present
- Phage (with cluster) where this start called: Success_26 (singleton),

Start 8:

• Found in 3 of 9 (33.3%) of genes in pham

- Manual Annotations of this start: 3 of 9
- Called 100.0% of time when present
- Phage (with cluster) where this start called: Grayson_26 (CB), Peregrin_24 (CB), Weasels2_22 (CB),

Start 9:

- Found in 5 of 9 (55.6%) of genes in pham
- Manual Annotations of this start: 5 of 9
- Called 100.0% of time when present
- Phage (with cluster) where this start called: Grayson_20 (CB), Grayson_21 (CB), Peregrin_19 (CB), Peregrin_20 (CB), Weasels2_18 (CB),

Summary by clusters:

There are 2 clusters represented in this pham: CB, singleton,

Info for manual annotations of cluster CB:

- •Start number 8 was manually annotated 3 times for cluster CB.
- •Start number 9 was manually annotated 5 times for cluster CB.

Gene Information:

Gene: Grayson_26 Start: 7449, Stop: 7075, Start Num: 8

Candidate Starts for Grayson_26:

(Start: 8 @7449 has 3 MA's), (21, 7281), (23, 7254), (26, 7209), (30, 7140),

Gene: Grayson_21 Start: 6068, Stop: 5700, Start Num: 9

Candidate Starts for Grayson_21:

(4, 6098), (5, 6092), (7, 6086), (Start: 9 @6068 has 5 MA's), (19, 5924), (21, 5909), (33, 5717),

Gene: Grayson 20 Start: 5707, Stop: 5357, Start Num: 9

Candidate Starts for Grayson 20:

(6, 5728), (Start: 9 @5707 has 5 MA's), (28, 5467),

Gene: Peregrin_19 Start: 5539, Stop: 5195, Start Num: 9

Candidate Starts for Peregrin_19: (Start: 9 @5539 has 5 MA's), (28, 5299),

Gene: Peregrin 20 Start: 5906, Stop: 5532, Start Num: 9

Candidate Starts for Peregrin_20:

(Start: 9 @ 5906 has 5 MA's), (19, 5762), (21, 5747), (33, 5555), (34, 5552),

Gene: Peregrin_24 Start: 6762, Stop: 6388, Start Num: 8

Candidate Starts for Peregrin_24:

(Start: 8 @ 6762 has 3 MA's), (12, 6708), (18, 6633), (20, 6597), (21, 6594), (25, 6534),

Gene: Success 26 Start: 17594, Stop: 17208, Start Num: 3

Candidate Starts for Success 26:

(1, 17702), (2, 17618), (Start: 3 @ 17594 has 1 MA's), (11, 17534), (13, 17492), (17, 17456), (24, 17376), (20, 17373)

17366), (29, 17279), (30, 17273),

Gene: Weasels2_18 Start: 5542, Stop: 5186, Start Num: 9

Candidate Starts for Weasels2_18:

(Start: 9 @5542 has 5 MA's), (14, 5473), (15, 5455), (31, 5224), (32, 5209), (35, 5200),

Gene: Weasels2_22 Start: 6969, Stop: 6571, Start Num: 8

Candidate Starts for Weasels2_22:

(Start: 8 @6969 has 3 MA's), (10, 6942), (16, 6867), (22, 6786), (27, 6723), (28, 6690), (30, 6657),