Pham 88296

1: Fresco_55 + 3

2: Mariokart_53 + 1

3: LittleMunchkin_55

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 88296 Report

This analysis was run 04/05/24 on database version 557.

Pham number 88296 has 7 members, 3 are drafts.

Phages represented in each track:
• Track 1 : Fresco_55, Axumite_55, Ligma_55, Shatter_55
• Track 2 : Mariokart_53, CharlottesWeb_53
• Track 3 : LittleMunchkin_55

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

The start number called the most often in the published annotations is 4, it was called in 3 of the 4 non-draft genes in the pham.

Genes that call this "Most Annotated" start:
• Axumite_55, Fresco_55, Ligma_55, LittleMunchkin_55, Shatter_55,

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

Genes that do not have the "Most Annotated" start:
• CharlottesWeb_53, Mariokart_53,

Summary by start number:

Start 4:
• Found in 5 of 7 (71.4%) of genes in pham
• Manual Annotations of this start: 3 of 4
• Called 100.0% of time when present
• Phage (with cluster) where this start called: Axumite_55 (DR), Fresco_55 (DR), Ligma_55 (DR), LittleMunchkin_55 (DR), Shatter_55 (DR),

Start 5:
• Found in 2 of 7 (28.6%) of genes in pham
• Manual Annotations of this start: 1 of 4
• Called 100.0% of time when present
• Phage (with cluster) where this start called: CharlottesWeb_53 (DR), Mariokart_53 (DR),

Summary by clusters:

There is one cluster represented in this pham: DR

Info for manual annotations of cluster DR:
•Start number 4 was manually annotated 3 times for cluster DR.
•Start number 5 was manually annotated 1 time for cluster DR.

Gene Information:
Gene: Axumite_55 Start: 47477, Stop: 47031, Start Num: 4
Candidate Starts for Axumite_55:
(Start: 4 @47477 has 3 MA's), (6, 47447), (8, 47426), (15, 47204),

Gene: CharlottesWeb_53 Start: 46544, Stop: 46098, Start Num: 5
Candidate Starts for CharlottesWeb_53:
(1, 46616), (3, 46601), (Start: 5 @46544 has 1 MA's), (6, 46514), (8, 46493), (12, 46322), (13, 46295),
(15, 46265), (18, 46109),

Gene: Fresco_55 Start: 47477, Stop: 47031, Start Num: 4
Candidate Starts for Fresco_55:
(Start: 4 @47477 has 3 MA's), (6, 47447), (8, 47426), (15, 47204),

Gene: Ligma_55 Start: 47477, Stop: 47031, Start Num: 4
Candidate Starts for Ligma_55:
(Start: 4 @47477 has 3 MA's), (6, 47447), (8, 47426), (15, 47204),

Gene: LittleMunchkin_55 Start: 49817, Stop: 49383, Start Num: 4
Candidate Starts for LittleMunchkin_55:
(2, 49877), (Start: 4 @49817 has 3 MA's), (7, 49784), (9, 49748), (10, 49706), (11, 49631), (14,
49562), (16, 49523), (17, 49397),

Gene: Mariokart_53 Start: 46830, Stop: 46384, Start Num: 5
Candidate Starts for Mariokart_53:
(1, 46902), (3, 46887), (Start: 5 @46830 has 1 MA's), (6, 46800), (8, 46779), (12, 46608), (13, 46581),
(15, 46551), (18, 46395),

Gene: Shatter_55 Start: 47477, Stop: 47031, Start Num: 4
Candidate Starts for Shatter_55:
(Start: 4 @47477 has 3 MA's), (6, 47447), (8, 47426), (15, 47204),