Pham 348374


2
6
21
1: Jflix2_59
2
5
6
11
13
17
21
24
2: REQ1_29
2
6
7
21
25
3: SFuller_58
2
6
13
21
4: Madraxi_62
1
8
9
14
15
16
18
22
23
5: BigAnti_49
3
4
10
12
19
20
6: Jace_74

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 348374 Report

This analysis was run 11/14/26 on database version 672.

Pham number 348374 has 6 members, 2 are drafts.

Phages represented in each track:
• Track 1 : Jflix2_59
• Track 2 : REQ1_29
• Track 3 : SFuller_58
• Track 4 : Madraxi_62
• Track 5 : BigAnti_49
• Track 6 : Jace_74

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 3 of the 4 non-draft genes in the pham.

Genes that call this "Most Annotated" start:
• Jflix2_59, Madraxi_62, REQ1_29, SFuller_58,

Genes that have the "Most Annotated" start but do not call it:
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Genes that do not have the "Most Annotated" start:
• BigAnti_49, Jace_74,

Summary by start number:

Start 1:
• Found in 1 of 6 (16.7%) of genes in pham
• No Manual Annotations of this start.
• Called 100.0% of time when present
• Phage (with cluster) where this start called: BigAnti_49 (singleton),

Start 2:
• Found in 4 of 6 (66.7%) 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: Jflix2_59 (CF), Madraxi_62 (CF), REQ1_29 (CF), SFuller_58 (CF),

Start 3:
• Found in 1 of 6 (16.7%) 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: Jace_74 (singleton),

Summary by clusters:

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

Info for manual annotations of cluster CF:
•Start number 2 was manually annotated 3 times for cluster CF.

Gene Information:

Gene: BigAnti_49 Start: 44134, Stop: 44700, Start Num: 1
Candidate Starts for BigAnti_49:
(1, 44134), (8, 44365), (9, 44371), (14, 44473), (15, 44476), (16, 44515), (18, 44554), (22, 44632), (23, 44680),

Gene: Jace_74 Start: 43206, Stop: 43682, Start Num: 3
Candidate Starts for Jace_74:
(Start: 3 @43206 has 1 MA's), (4, 43215), (10, 43401), (12, 43419), (19, 43557), (20, 43563),

Gene: Jflix2_59 Start: 40098, Stop: 40586, Start Num: 2
Candidate Starts for Jflix2_59:
(Start: 2 @40098 has 3 MA's), (6, 40209), (21, 40482),

Gene: Madraxi_62 Start: 42601, Stop: 43107, Start Num: 2
Candidate Starts for Madraxi_62:
(Start: 2 @42601 has 3 MA's), (6, 42712), (13, 42844), (21, 42982),

Gene: REQ1_29 Start: 11628, Stop: 12131, Start Num: 2
Candidate Starts for REQ1_29:
(Start: 2 @11628 has 3 MA's), (5, 11727), (6, 11739), (11, 11850), (13, 11871), (17, 11946), (21, 12009), (24, 12090),

Gene: SFuller_58 Start: 40290, Stop: 40778, Start Num: 2
Candidate Starts for SFuller_58:
(Start: 2 @40290 has 3 MA's), (6, 40401), (7, 40425), (21, 40674), (25, 40761),