Pham 346564


4
6
7
9
10
14
1: Faja_33
5
12
13
2: Pipp_32
1
5
8
11
15
17
3: KappaFarmDelta_34 + 1
5
15
17
4: CheeseSpider_35
2
3
6
16
5: Rooter_25

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 346564 Report

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

Pham number 346564 has 6 members, 0 are drafts.

Phages represented in each track:
• Track 1 : Faja_33
• Track 2 : Pipp_32
• Track 3 : KappaFarmDelta_34, GEazy_31
• Track 4 : CheeseSpider_35
• Track 5 : Rooter_25

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

The start number called the most often in the published annotations is 5, it was called in 4 of the 6 non-draft genes in the pham.

Genes that call this "Most Annotated" start:
• CheeseSpider_35, GEazy_31, KappaFarmDelta_34, Pipp_32,

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

Genes that do not have the "Most Annotated" start:
• Faja_33, Rooter_25,

Summary by start number:

Start 3:
• Found in 1 of 6 (16.7%) of genes in pham
• Manual Annotations of this start: 1 of 6
• Called 100.0% of time when present
• Phage (with cluster) where this start called: Rooter_25 (FB),

Start 5:
• Found in 4 of 6 (66.7%) of genes in pham
• Manual Annotations of this start: 4 of 6
• Called 100.0% of time when present
• Phage (with cluster) where this start called: CheeseSpider_35 (CV), GEazy_31 (DB), KappaFarmDelta_34 (CV), Pipp_32 (CV),

Start 7:
• Found in 1 of 6 (16.7%) of genes in pham
• Manual Annotations of this start: 1 of 6
• Called 100.0% of time when present
• Phage (with cluster) where this start called: Faja_33 (AY),

Summary by clusters:

There are 4 clusters represented in this pham: AY, FB, DB, CV,

Info for manual annotations of cluster AY:
•Start number 7 was manually annotated 1 time for cluster AY.

Info for manual annotations of cluster CV:
•Start number 5 was manually annotated 3 times for cluster CV.

Info for manual annotations of cluster DB:
•Start number 5 was manually annotated 1 time for cluster DB.

Info for manual annotations of cluster FB:
•Start number 3 was manually annotated 1 time for cluster FB.

Gene Information:

Gene: CheeseSpider_35 Start: 27464, Stop: 27237, Start Num: 5
Candidate Starts for CheeseSpider_35:
(Start: 5 @27464 has 4 MA's), (15, 27308), (17, 27278),

Gene: Faja_33 Start: 24279, Stop: 24067, Start Num: 7
Candidate Starts for Faja_33:
(4, 24294), (6, 24288), (Start: 7 @24279 has 1 MA's), (9, 24243), (10, 24228), (14, 24132),

Gene: GEazy_31 Start: 26375, Stop: 26142, Start Num: 5
Candidate Starts for GEazy_31:
(1, 26429), (Start: 5 @26375 has 4 MA's), (8, 26354), (11, 26261), (15, 26219), (17, 26189),

Gene: KappaFarmDelta_34 Start: 27349, Stop: 27116, Start Num: 5
Candidate Starts for KappaFarmDelta_34:
(1, 27403), (Start: 5 @27349 has 4 MA's), (8, 27328), (11, 27235), (15, 27193), (17, 27163),

Gene: Pipp_32 Start: 27488, Stop: 27261, Start Num: 5
Candidate Starts for Pipp_32:
(Start: 5 @27488 has 4 MA's), (12, 27359), (13, 27356),

Gene: Rooter_25 Start: 20212, Stop: 19976, Start Num: 3
Candidate Starts for Rooter_25:
(2, 20215), (Start: 3 @20212 has 1 MA's), (6, 20203), (16, 20020),