Pham 225289

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 225289 Report

This analysis was run 03/28/25 on database version 593.

Pham number 225289 has 5 members, 4 are drafts.

Phages represented in each track:
• Track 1 : McMater_45, Dartin_45, Richo_45
• Track 2 : YoureAdopted_42
• Track 3 : Doughnut_46

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

The start number called the most often in the published annotations is 7, it was called in 1 of the 1 non-draft genes in the pham.

Genes that call this "Most Annotated" start:
• Doughnut_46, YoureAdopted_42,

Genes that have the "Most Annotated" start but do not call it:
• Dartin_45, McMater_45, Richo_45,

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

Summary by start number:

Start 2:
• Found in 3 of 5 (60.0%) of genes in pham
• No Manual Annotations of this start.
• Called 100.0% of time when present
• Phage (with cluster) where this start called: Dartin_45 (K1), McMater_45 (K1), Richo_45 (K1),

Start 7:
• Found in 5 of 5 (100.0%) of genes in pham
• Manual Annotations of this start: 1 of 1
• Called 40.0% of time when present
• Phage (with cluster) where this start called: Doughnut_46 (K2), YoureAdopted_42 (K1),

Summary by clusters:

There are 2 clusters represented in this pham: K2, K1,

Info for manual annotations of cluster K1:
•Start number 7 was manually annotated 1 time for cluster K1.

Gene Information:
Gene: Dartin_45 Start: 35648, Stop: 35256, Start Num: 2
Candidate Starts for Dartin_45:
(2, 35648), (6, 35564), (Start: 7 @35549 has 1 MA's), (10, 35411), (12, 35387), (15, 35324), (16, 35300), (18, 35267),

Gene: Doughnut_46 Start: 36568, Stop: 36287, Start Num: 7
Candidate Starts for Doughnut_46:
(1, 36718), (3, 36664), (4, 36628), (5, 36613), (Start: 7 @36568 has 1 MA's), (8, 36508), (9, 36448), (11, 36418), (13, 36370),

Gene: McMater_45 Start: 35648, Stop: 35256, Start Num: 2
Candidate Starts for McMater_45:
(2, 35648), (6, 35564), (Start: 7 @35549 has 1 MA's), (10, 35411), (12, 35387), (15, 35324), (16, 35300), (18, 35267),

Gene: Richo_45 Start: 35648, Stop: 35256, Start Num: 2
Candidate Starts for Richo_45:
(2, 35648), (6, 35564), (Start: 7 @35549 has 1 MA's), (10, 35411), (12, 35387), (15, 35324), (16, 35300), (18, 35267),

Gene: YoureAdopted_42 Start: 35085, Stop: 34804, Start Num: 7
Candidate Starts for YoureAdopted_42:
(Start: 7 @35085 has 1 MA's), (14, 34878), (17, 34830),