Pham 6382

1: A3Wally_34 + 3

2: Zooman_35 + 1

3: Big4_38 + 1

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 6382 Report

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

Pham number 6382 has 8 members, 0 are drafts.

Phages represented in each track:
• Track 1 : A3Wally_34, PauloDiaboli_389, PauloDiaboli_34, A3Wally_387
• Track 2 : Zooman_35, Zooman_348
• Track 3 : Big4_38, Big4_364

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

The start number called the most often in the published annotations is 1, it was called in 8 of the 8 non-draft genes in the pham.

Genes that call this "Most Annotated" start:
• A3Wally_34, A3Wally_387, Big4_364, Big4_38, PauloDiaboli_34, PauloDiaboli_389, Zooman_348, Zooman_35,

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

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

Summary by start number:

Start 1:
• Found in 8 of 8 (100.0%) of genes in pham
• Manual Annotations of this start: 8 of 8
• Called 100.0% of time when present
• Phage (with cluster) where this start called: A3Wally_34 (GD1), A3Wally_387 (GD1), Big4_364 (GD2), Big4_38 (GD2), PauloDiaboli_34 (GD1), PauloDiaboli_389 (GD1), Zooman_348 (GD2), Zooman_35 (GD2),

Summary by clusters:

There are 2 clusters represented in this pham: GD1, GD2,

Info for manual annotations of cluster GD1:

•Start number 1 was manually annotated 4 times for cluster GD1.

Info for manual annotations of cluster GD2:
•Start number 1 was manually annotated 4 times for cluster GD2.

Gene Information:

Gene: A3Wally_34 Start: 11828, Stop: 11938, Start Num: 1
Candidate Starts for A3Wally_34:
(Start: 1 @11828 has 8 MA's), (3, 11912), (4, 11918),

Gene: A3Wally_387 Start: 191049, Stop: 191159, Start Num: 1
Candidate Starts for A3Wally_387:
(Start: 1 @191049 has 8 MA's), (3, 191133), (4, 191139),

Gene: Big4_38 Start: 14704, Stop: 14823, Start Num: 1
Candidate Starts for Big4_38:
(Start: 1 @14704 has 8 MA's), (2, 14764),

Gene: Big4_364 Start: 189398, Stop: 189517, Start Num: 1
Candidate Starts for Big4_364:
(Start: 1 @189398 has 8 MA's), (2, 189458),

Gene: PauloDiaboli_389 Start: 188297, Stop: 188407, Start Num: 1
Candidate Starts for PauloDiaboli_389:
(Start: 1 @188297 has 8 MA's), (3, 188381), (4, 188387),

Gene: PauloDiaboli_34 Start: 11668, Stop: 11778, Start Num: 1
Candidate Starts for PauloDiaboli_34:
(Start: 1 @11668 has 8 MA's), (3, 11752), (4, 11758),

Gene: Zooman_35 Start: 14155, Stop: 14274, Start Num: 1
Candidate Starts for Zooman_35:
(Start: 1 @14155 has 8 MA's),

Gene: Zooman_348 Start: 189806, Stop: 189925, Start Num: 1
Candidate Starts for Zooman_348:
(Start: 1 @189806 has 8 MA's),