



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 310132 Report

This analysis was run 09/05/26 on database version 662.

Pham number 310132 has 6 members, 0 are drafts.

Phages represented in each track:

- Track 1 : PauloDiaboli_265
- Track 2 : PauloDiaboli_264, Dodo_259
- Track 3 : A3Wally_265
- Track 4 : Dodo_260
- Track 5 : A3Wally_264

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

The start number called the most often in the published annotations is 3, it was called in 2 of the 6 non-draft genes in the pham.

Genes that call this "Most Annotated" start:

- Dodo_259, PauloDiaboli_264,

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

- A3Wally_264,

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

- A3Wally_265, Dodo_260, PauloDiaboli_265,

Summary by start number:

Start 1:

- Found in 3 of 6 (50.0%) of genes in pham
- Manual Annotations of this start: 2 of 6
- Called 66.7% of time when present
- Phage (with cluster) where this start called: A3Wally_265 (GD1), PauloDiaboli_265 (GD1),

Start 2:

- Found in 2 of 6 (33.3%) of genes in pham
- Manual Annotations of this start: 1 of 6
- Called 50.0% of time when present
- Phage (with cluster) where this start called: Dodo_260 (GD1),

Start 3:

- Found in 3 of 6 (50.0%) of genes in pham
- Manual Annotations of this start: 2 of 6
- Called 66.7% of time when present
- Phage (with cluster) where this start called: Dodo_259 (GD1), PauloDiaboli_264 (GD1),

Start 4:

- Found in 3 of 6 (50.0%) of genes in pham
- Manual Annotations of this start: 1 of 6
- Called 33.3% of time when present
- Phage (with cluster) where this start called: A3Wally_264 (GD1),

Summary by clusters:

There is one cluster represented in this pham: GD1

Info for manual annotations of cluster GD1:

- Start number 1 was manually annotated 2 times for cluster GD1.
- Start number 2 was manually annotated 1 time for cluster GD1.
- Start number 3 was manually annotated 2 times for cluster GD1.
- Start number 4 was manually annotated 1 time for cluster GD1.

Gene Information:

Gene: A3Wally_265 Start: 145405, Stop: 145743, Start Num: 1

Candidate Starts for A3Wally_265:

(Start: 1 @145405 has 2 MA's), (5, 145474), (6, 145480), (7, 145483), (9, 145540), (10, 145561), (12, 145597), (13, 145603), (15, 145633), (17, 145666),

Gene: A3Wally_264 Start: 145136, Stop: 145411, Start Num: 4

Candidate Starts for A3Wally_264:

(Start: 3 @145130 has 2 MA's), (Start: 4 @145136 has 1 MA's), (11, 145265), (13, 145292), (14, 145304), (16, 145328),

Gene: Dodo_259 Start: 144817, Stop: 145098, Start Num: 3

Candidate Starts for Dodo_259:

(Start: 3 @144817 has 2 MA's), (Start: 4 @144823 has 1 MA's), (11, 144952), (13, 144979), (14, 144991),

Gene: Dodo_260 Start: 145098, Stop: 145430, Start Num: 2

Candidate Starts for Dodo_260:

(Start: 1 @145092 has 2 MA's), (Start: 2 @145098 has 1 MA's), (8, 145182), (12, 145284), (13, 145290), (15, 145320), (17, 145353),

Gene: PauloDiaboli_265 Start: 142607, Stop: 142945, Start Num: 1

Candidate Starts for PauloDiaboli_265:

(Start: 1 @142607 has 2 MA's), (Start: 2 @142613 has 1 MA's), (8, 142697), (12, 142799), (13, 142805), (15, 142835), (17, 142868),

Gene: PauloDiaboli_264 Start: 142332, Stop: 142613, Start Num: 3

Candidate Starts for PauloDiaboli_264:

(Start: 3 @142332 has 2 MA's), (Start: 4 @142338 has 1 MA's), (11, 142467), (13, 142494), (14, 142506),