

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 168684 Report

This analysis was run 07/09/24 on database version 566.

Pham number 168684 has 10 members, 7 are drafts.

Phages represented in each track:

Track 1 : AinMach_14

Track 2 : Exile_14

Track 3: IttyBittyPiggy_15

Track 4 : Sue2 15

• Track 5 : ShakeItOph_15, MiniMommy_14

Track 6: B22_15, G4_15
Track 7: Doucette_15
Track 8: Coriander 16

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 3 non-draft genes in the pham.

Genes that call this "Most Annotated" start:

• B22_15, Doucette_15, G4_15,

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

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

• AinMach_14, Coriander_16, Exile_14, IttyBittyPiggy_15, MiniMommy_14, ShakeItOph_15, Sue2_15,

Summary by start number:

Start 2:

- Found in 3 of 10 (30.0%) of genes in pham
- Manual Annotations of this start: 3 of 3
- Called 100.0% of time when present
- Phage (with cluster) where this start called: B22_15 (BW), Doucette_15 (BW), G4_15 (BW),

Start 3:

- Found in 2 of 10 (20.0%) of genes in pham
- No Manual Annotations of this start.
- Called 100.0% of time when present
- Phage (with cluster) where this start called: AinMach_14 (AZ), IttyBittyPiggy_15 (AZ1),

Start 4:

- Found in 1 of 10 (10.0%) of genes in pham
- No Manual Annotations of this start.
- Called 100.0% of time when present
- Phage (with cluster) where this start called: Coriander_16 (DB),

Start 5:

- Found in 3 of 10 (30.0%) of genes in pham
- No Manual Annotations of this start.
- Called 100.0% of time when present
- Phage (with cluster) where this start called: Exile_14 (AZ), MiniMommy_14 (AZ4), ShakeltOph_15 (AZ4),

Start 9:

- Found in 5 of 10 (50.0%) of genes in pham
- No Manual Annotations of this start.
- Called 20.0% of time when present
- Phage (with cluster) where this start called: Sue2_15 (AZ1),

Summary by clusters:

There are 5 clusters represented in this pham: BW, DB, AZ, AZ4, AZ1,

Info for manual annotations of cluster BW:

•Start number 2 was manually annotated 3 times for cluster BW.

Gene Information:

Gene: AinMach 14 Start: 9534, Stop: 9839, Start Num: 3

Candidate Starts for AinMach 14:

(1, 9324), (3, 9534), (10, 9621), (20, 9783), (22, 9801),

Gene: B22 15 Start: 9378, Stop: 9746, Start Num: 2

Candidate Starts for B22_15:

(Start: 2 @9378 has 3 MA's), (9, 9498), (13, 9540), (15, 9555), (16, 9567), (24, 9717),

Gene: Coriander 16 Start: 10544, Stop: 10876, Start Num: 4

Candidate Starts for Coriander 16:

(4, 10544), (7, 10589), (8, 10595), (11, 10625), (12, 10643), (18, 10712), (19, 10772), (25, 10862),

Gene: Doucette_15 Start: 9437, Stop: 9805, Start Num: 2

Candidate Starts for Doucette_15:

(Start: 2 @9437 has 3 MA's), (6, 9521), (9, 9557), (13, 9599), (15, 9614), (16, 9626), (24, 9776),

Gene: Exile_14 Start: 10360, Stop: 10629, Start Num: 5

Candidate Starts for Exile_14:

(5, 10360), (9, 10411), (10, 10417), (23, 10618),

Gene: G4_15 Start: 9427, Stop: 9795, Start Num: 2

Candidate Starts for G4_15:

(Start: 2 @ 9427 has 3 MA's), (9, 9547), (13, 9589), (15, 9604), (16, 9616), (24, 9766),

Gene: IttyBittyPiggy_15 Start: 10410, Stop: 10706, Start Num: 3

Candidate Starts for IttyBittyPiggy_15:

(1, 10200), (3, 10410),

Gene: MiniMommy_14 Start: 8690, Stop: 8962, Start Num: 5

Candidate Starts for MiniMommy_14:

(5, 8690), (21, 8912),

Gene: ShakeltOph_15 Start: 8689, Stop: 8961, Start Num: 5

Candidate Starts for ShakeItOph_15:

(5, 8689), (21, 8911),

Gene: Sue2_15 Start: 10511, Stop: 10759, Start Num: 9

Candidate Starts for Sue2_15:

(1, 10220), (9, 10511), (14, 10556), (17, 10598),