

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 213317 Report

This analysis was run 02/22/25 on database version 588.

Pham number 213317 has 7 members, 3 are drafts.

Phages represented in each track:

Track 1 : Phaila_85

Track 2 : GantcherGoblin_80

Track 3 : Zeina_84Track 4 : Argan 82

• Track 5 : KevinMinion 80

Track 6 : Uzumaki_81Track 7 : Navi1117 87

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 3 of the 4 non-draft genes in the pham.

Genes that call this "Most Annotated" start:

• GantcherGoblin_80, Navi1117_87, Uzumaki_81, Zeina_84,

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

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

Argan_82, KevinMinion_80, Phaila_85,

Summary by start number:

Start 1:

- Found in 2 of 7 (28.6%) of genes in pham
- No Manual Annotations of this start.
- Called 100.0% of time when present
- Phage (with cluster) where this start called: KevinMinion_80 (AU6), Phaila_85 (AU6),

Start 2:

- Found in 1 of 7 (14.3%) of genes in pham
- Manual Annotations of this start: 1 of 4
- Called 100.0% of time when present

Phage (with cluster) where this start called: Argan_82 (AU6),

Start 3:

- Found in 4 of 7 (57.1%) of genes in pham
- Manual Annotations of this start: 3 of 4
- Called 100.0% of time when present
- Phage (with cluster) where this start called: GantcherGoblin_80 (AU6), Navi1117_87 (AU6), Uzumaki_81 (AU6), Zeina_84 (AU6),

Summary by clusters:

There is one cluster represented in this pham: AU6

Info for manual annotations of cluster AU6:

- •Start number 2 was manually annotated 1 time for cluster AU6.
- •Start number 3 was manually annotated 3 times for cluster AU6.

Gene Information:

Gene: Argan 82 Start: 48814, Stop: 49053, Start Num: 2

Candidate Starts for Argan_82:

(Start: 2 @ 48814 has 1 MA's), (6, 48874), (8, 48886), (11, 48937), (14, 48979), (15, 48988),

Gene: GantcherGoblin 80 Start: 48628, Stop: 48867, Start Num: 3

Candidate Starts for GantcherGoblin 80:

(Start: 3 @48628 has 3 MA's), (5, 48664), (6, 48688), (7, 48691), (8, 48700), (11, 48751), (14, 48793), (15, 48802),

Gene: KevinMinion_80 Start: 49418, Stop: 49684, Start Num: 1

Candidate Starts for KevinMinion 80:

(1, 49418), (6, 49505), (8, 49517), (10, 49565), (11, 49568), (13, 49604), (14, 49610), (15, 49619),

Gene: Navi1117 87 Start: 50186, Stop: 50425, Start Num: 3

Candidate Starts for Navi1117_87:

(Start: 3 @50186 has 3 MA's), (4, 50219), (5, 50222), (6, 50246), (8, 50258), (10, 50306), (11, 50309), (12, 50333), (14, 50351), (15, 50360),

Gene: Phaila 85 Start: 49012, Stop: 49278, Start Num: 1

Candidate Starts for Phaila 85:

(1, 49012), (6, 49099), (8, 49111), (11, 49162), (12, 49186), (14, 49204), (15, 49213),

Gene: Uzumaki_81 Start: 48596, Stop: 48835, Start Num: 3

Candidate Starts for Uzumaki_81:

(Start: 3 @48596 has 3 MA's), (4, 48629), (5, 48632), (6, 48656), (7, 48659), (8, 48668), (11, 48719), (14, 48761),

Gene: Zeina 84 Start: 48968, Stop: 49207, Start Num: 3

Candidate Starts for Zeina 84:

(Start: 3 @48968 has 3 MA's), (4, 49001), (5, 49004), (6, 49028), (7, 49031), (8, 49040), (9, 49058), (10, 49088), (11, 49091), (14, 49133), (15, 49142),