

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 195851 Report

This analysis was run 12/09/24 on database version 580.

Pham number 195851 has 12 members, 1 are drafts.

Phages represented in each track:

• Track 1 : Tank 37, Wilde 37

• Track 2: Rizwana 35

Track 3: Beagle_46, Pointis_44, Pureglobe5_46

Track 4 : Odyssey395_47

Track 5 : MellowYellow_42, NyleyClemson_41

Track 6 : BruhMoment_42

Track 7 : AWGoat_35

Track 8 : SilentRX_37

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

The start number called the most often in the published annotations is 6, it was called in 6 of the 11 non-draft genes in the pham.

Genes that call this "Most Annotated" start:

 Beagle_46, BruhMoment_42, MellowYellow_42, NyleyClemson_41, Odyssey395_47, Pointis_44, Pureglobe5_46,

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

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

AWGoat_35, Rizwana_35, SilentRX_37, Tank_37, Wilde_37,

Summary by start number:

Start 6:

- Found in 7 of 12 (58.3%) of genes in pham
- Manual Annotations of this start: 6 of 11
- Called 100.0% of time when present
- Phage (with cluster) where this start called: Beagle_46 (AP2), BruhMoment_42 (AP3), MellowYellow_42 (AP2), NyleyClemson_41 (AP2), Odyssey395_47 (AP2), Pointis_44 (AP2), Pureglobe5_46 (AP2),

Start 7:

- Found in 2 of 12 (16.7%) of genes in pham
- Manual Annotations of this start: 2 of 11
- Called 100.0% of time when present
- Phage (with cluster) where this start called: AWGoat_35 (AP4), SilentRX_37 (AP4),

Start 8:

- Found in 3 of 12 (25.0%) of genes in pham
- Manual Annotations of this start: 1 of 11
- Called 33.3% of time when present
- Phage (with cluster) where this start called: Rizwana_35 (AP1),

Start 9:

- Found in 2 of 12 (16.7%) of genes in pham
- Manual Annotations of this start: 2 of 11
- Called 100.0% of time when present
- Phage (with cluster) where this start called: Tank_37 (AP1), Wilde_37 (AP1),

Summary by clusters:

There are 4 clusters represented in this pham: AP2, AP3, AP1, AP4,

Info for manual annotations of cluster AP1:

- •Start number 8 was manually annotated 1 time for cluster AP1.
- •Start number 9 was manually annotated 2 times for cluster AP1.

Info for manual annotations of cluster AP2:

•Start number 6 was manually annotated 5 times for cluster AP2.

Info for manual annotations of cluster AP3:

•Start number 6 was manually annotated 1 time for cluster AP3.

Info for manual annotations of cluster AP4:

•Start number 7 was manually annotated 2 times for cluster AP4.

Gene Information:

Gene: AWGoat 35 Start: 33444, Stop: 33779, Start Num: 7

Candidate Starts for AWGoat 35:

(Start: 7 @ 33444 has 2 MA's), (12, 33480), (17, 33549), (18, 33552), (19, 33561), (27, 33675), (32, 33729), (33, 33732), (34, 33741),

Gene: Beagle_46 Start: 34747, Stop: 35091, Start Num: 6

Candidate Starts for Beagle_46:

(3, 34687), (5, 34723), (Start: 6 @ 34747 has 6 MA's), (11, 34783), (16, 34855), (20, 34885), (25, 34930), (27, 34987), (28, 34990), (33, 35044),

Gene: BruhMoment 42 Start: 36111, Stop: 36455, Start Num: 6

Candidate Starts for BruhMoment 42:

(1, 35940), (2, 35952), (Start: 6 @ 36111 has 6 MA's), (18, 36228), (22, 36267), (23, 36270), (24, 36276), (27, 36351), (31, 36396), (33, 36408),

Gene: MellowYellow_42 Start: 33482, Stop: 33826, Start Num: 6

Candidate Starts for MellowYellow 42:

(3, 33422), (Start: 6 @33482 has 6 MA's), (11, 33518), (14, 33545), (16, 33590), (20, 33620), (25, 33665), (27, 33722), (28, 33725), (33, 33779),

Gene: NyleyClemson_41 Start: 33097, Stop: 33441, Start Num: 6

Candidate Starts for NyleyClemson 41:

(3, 33037), (Start: 6 @33097 has 6 MA's), (11, 33133), (14, 33160), (16, 33205), (20, 33235), (25, 33280), (27, 33337), (28, 33340), (33, 33394),

Gene: Odyssey395 47 Start: 34766, Stop: 35110, Start Num: 6

Candidate Starts for Odyssey395_47:

(3, 34706), (4, 34715), (5, 34742), (Start: 6 @34766 has 6 MA's), (11, 34802), (16, 34874), (20, 34904), (25, 34949), (27, 35006), (28, 35009), (33, 35063),

Gene: Pointis_44 Start: 34764, Stop: 35108, Start Num: 6

Candidate Starts for Pointis 44:

(3, 34704), (5, 34740), (Start: 6 @ 34764 has 6 MA's), (11, 34800), (16, 34872), (20, 34902), (25, 34947), (27, 35004), (28, 35007), (33, 35061),

Gene: Pureglobe5_46 Start: 34947, Stop: 35291, Start Num: 6

Candidate Starts for Pureglobe 546:

(3, 34887), (5, 34923), (Start: 6 @ 34947 has 6 MA's), (11, 34983), (16, 35055), (20, 35085), (25, 35130), (27, 35187), (28, 35190), (33, 35244),

Gene: Rizwana_35 Start: 34988, Stop: 35311, Start Num: 8

Candidate Starts for Rizwana 35:

(Start: 8 @34988 has 1 MA's), (10, 34997), (15, 35054), (21, 35120), (22, 35123), (26, 35162), (29, 35216), (30, 35222), (31, 35252),

Gene: SilentRX_37 Start: 34506, Stop: 34829, Start Num: 7

Candidate Starts for SilentRX 37:

(Start: 7 @34506 has 2 MA's), (13, 34551), (15, 34572), (17, 34599), (18, 34602), (19, 34611), (33, 34782), (34, 34791),

Gene: Tank_37 Start: 35024, Stop: 35344, Start Num: 9

Candidate Starts for Tank_37:

(Start: 8 @35021 has 1 MA's), (Start: 9 @35024 has 2 MA's), (15, 35087), (18, 35117), (26, 35195), (29, 35249), (31, 35285), (33, 35297),

Gene: Wilde_37 Start: 34866, Stop: 35186, Start Num: 9

Candidate Starts for Wilde 37:

(Start: 8 @34863 has 1 MA's), (Start: 9 @34866 has 2 MA's), (15, 34929), (18, 34959), (26, 35037), (29, 35091), (31, 35127), (33, 35139),