

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 192954 Report

This analysis was run 11/02/24 on database version 579.

WARNING: Pham size does not match number of genes in report. Either unphamerated genes have been added (by you) or starterator has removed genes due to invalid start codon.

Pham number 192954 has 16 members, 0 are drafts.

Phages represented in each track:

- Track 1 : Roman 8
- Track 2: Hortus1_11, OlinDD_11, Tandem_11, Alleb_12
- Track 3 : Pavlo 8
- Track 4: Pioneer3 9
- Track 5 : Hubbs_8
- Track 6: OlinDD_9, Hortus1_9, Alleb_10
- Track 7 : PhillyPhilly_9
- Track 8 : Platte_11
- Track 9 : DejaVu_9
- Track 10 : Pioneer3_11
- Track 11 : Lupine 8

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

The start number called the most often in the published annotations is 8, it was called in 6 of the 16 non-draft genes in the pham.

Genes that call this "Most Annotated" start:

• Alleb_12, Hortus1_11, OlinDD_11, Pioneer3_11, Platte_11, Tandem_11,

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

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

• Alleb_10, DejaVu_9, Hortus1_9, Hubbs_8, Lupine_8, OlinDD_9, Pavlo_8, PhillyPhilly_9, Pioneer3_9, Roman_8,

Summary by start number:

Start 3:

- Found in 4 of 16 (25.0%) of genes in pham
- Manual Annotations of this start: 1 of 16
- Called 25.0% of time when present
- Phage (with cluster) where this start called: PhillyPhilly_9 (ED1),

Start 6:

- Found in 6 of 16 (37.5%) of genes in pham
- Manual Annotations of this start: 5 of 16
- Called 83.3% of time when present
- Phage (with cluster) where this start called: DejaVu_9 (ED1), Hubbs_8 (ED1), Lupine_8 (ED1), Pavlo_8 (ED1), Roman_8 (ED1),

Start 7:

- Found in 4 of 16 (25.0%) of genes in pham
- Manual Annotations of this start: 4 of 16
- Called 100.0% of time when present
- Phage (with cluster) where this start called: Alleb_10 (ED1), Hortus1_9 (ED1), OlinDD_9 (ED1), Pioneer3_9 (ED1),

Start 8:

- Found in 6 of 16 (37.5%) of genes in pham
- Manual Annotations of this start: 6 of 16
- Called 100.0% of time when present
- Phage (with cluster) where this start called: Alleb_12 (ED1), Hortus1_11 (ED1), OlinDD_11 (ED1), Pioneer3_11 (ED1), Platte_11 (ED1), Tandem_11 (ED1),

Summary by clusters:

There is one cluster represented in this pham: ED1

Info for manual annotations of cluster ED1:

- •Start number 3 was manually annotated 1 time for cluster ED1.
- Start number 6 was manually annotated 5 times for cluster ED1.
- •Start number 7 was manually annotated 4 times for cluster ED1.
- •Start number 8 was manually annotated 6 times for cluster ED1.

Gene Information:

Gene: Alleb_12 Start: 4351, Stop: 4713, Start Num: 8

Candidate Starts for Alleb 12:

(4, 4258), (Start: 8 @4351 has 6 MA's), (9, 4405), (11, 4423), (16, 4555), (20, 4600), (25, 4696), (26, 4702),

Gene: Alleb_10 Start: 3636, Stop: 4010, Start Num: 7

Candidate Starts for Alleb 10:

(5, 3576), (Start: 7 @3636 has 4 MA's), (21, 3921), (22, 3936), (23, 3957), (26, 3999),

Gene: DejaVu 9 Start: 3447, Stop: 3833, Start Num: 6

Candidate Starts for DejaVu 9:

(Start: 6 @3447 has 5 MA's), (10, 3561), (12, 3579),

Gene: Hortus1_11 Start: 4305, Stop: 4667, Start Num: 8

Candidate Starts for Hortus1_11:

(4, 4212), (Start: 8 @4305 has 6 MA's), (9, 4359), (11, 4377), (16, 4509), (20, 4554), (25, 4650), (26, 4656),

Gene: Hortus1_9 Start: 3590, Stop: 3964, Start Num: 7

Candidate Starts for Hortus1_9:

(5, 3530), (Start: 7 @3590 has 4 MA's), (21, 3875), (22, 3890), (23, 3911), (26, 3953),

Gene: Hubbs_8 Start: 3631, Stop: 4017, Start Num: 6

Candidate Starts for Hubbs 8:

(Start: 3 @3574 has 1 MA's), (Start: 6 @3631 has 5 MA's), (10, 3745), (12, 3763), (15, 3826), (26, 4006),

Gene: Lupine_8 Start: 3428, Stop: 3817, Start Num: 6

Candidate Starts for Lupine_8:

(2, 3308), (Start: 3 @3371 has 1 MA's), (Start: 6 @3428 has 5 MA's), (12, 3560), (16, 3665), (20, 3707), (23, 3767), (26, 3806),

Gene: OlinDD_11 Start: 4304, Stop: 4666, Start Num: 8

Candidate Starts for OlinDD 11:

(4, 4211), (Start: 8 @4304 has 6 MA's), (9, 4358), (11, 4376), (16, 4508), (20, 4553), (25, 4649), (26, 4655).

Gene: OlinDD_9 Start: 3589, Stop: 3963, Start Num: 7

Candidate Starts for OlinDD_9:

(5, 3529), (Start: 7 @ 3589 has 4 MA's), (21, 3874), (22, 3889), (23, 3910), (26, 3952),

Gene: Pavlo_8 Start: 3702, Stop: 4091, Start Num: 6

Candidate Starts for Pavlo_8:

(1, 3465), (Start: 6 @ 3702 has 5 MA's), (16, 3939), (20, 3981), (23, 4041), (26, 4080),

Gene: PhillyPhilly_9 Start: 3491, Stop: 3940, Start Num: 3

Candidate Starts for PhillyPhilly 9:

(Start: 3 @3491 has 1 MA's), (Start: 6 @3548 has 5 MA's), (12, 3680), (16, 3785), (17, 3788), (19, 3809), (20, 3830), (26, 3929),

Gene: Pioneer3_9 Start: 3622, Stop: 3996, Start Num: 7

Candidate Starts for Pioneer3_9:

(5, 3562), (Start: 7 @3622 has 4 MA's), (21, 3907), (22, 3922), (23, 3943), (24, 3946), (26, 3985),

Gene: Pioneer3_11 Start: 4337, Stop: 4690, Start Num: 8

Candidate Starts for Pioneer3 11:

(4, 4244), (Start: 8 @4337 has 6 MA's), (13, 4418), (14, 4421), (16, 4532), (18, 4541), (20, 4577), (25, 4673), (26, 4679),

Gene: Platte_11 Start: 4138, Stop: 4491, Start Num: 8

Candidate Starts for Platte_11:

(4, 4045), (Start: 8 @4138 has 6 MA's), (13, 4219), (16, 4333), (18, 4342), (20, 4378), (25, 4474), (26, 4480),

Gene: Roman 8 Start: 3340, Stop: 3726, Start Num: 6

Candidate Starts for Roman_8:

(Start: 3 @3283 has 1 MA's), (Start: 6 @3340 has 5 MA's), (12, 3472), (15, 3535), (26, 3715),

Gene: Tandem_11 Start: 4242, Stop: 4604, Start Num: 8

Candidate Starts for Tandem_11:

(4, 4149), (Start: 8 @4242 has 6 MA's), (9, 4296), (11, 4314), (16, 4446), (20, 4491), (25, 4587), (26,

4593),