

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 171768 Report

This analysis was run 07/10/24 on database version 566.

Pham number 171768 has 21 members, 4 are drafts.

Phages represented in each track:

- Track 1 : Anatole_31
- Track 2 : B3_31
- Track 3 : G4_32
- Track 4 : B22_32, Doucette_34
- Track 5 : E6_31, E1_31
- Track 6 : Eileen_37, Judy_42, GlobiWarming_41, RootBeer_34, Bridgette_41,
- ChuckDuck_42, Peas_38, Constance_41
- Track 7 : Kumotta_33, MargaretKali_33
- Track 8 : BrayBeast_36
- Track 9 : BlackSpider_39
- Track 10 : 32HC_49, Rem711_48

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

The start number called the most often in the published annotations is 4, it was called in 9 of the 17 non-draft genes in the pham.

Genes that call this "Most Annotated" start:

• BlackSpider_39, BrayBeast_36, Bridgette_41, ChuckDuck_42, Constance_41, Eileen_37, GlobiWarming_41, Judy_42, Kumotta_33, MargaretKali_33, Peas_38, RootBeer_34,

Genes that have the "Most Annotated" start but do not call it: • 32HC_49, Rem711_48,

Genes that do not have the "Most Annotated" start: • Anatole_31, B22_32, B3_31, Doucette_34, E1_31, E6_31, G4_32,

Summary by start number:

Start 3:

- Found in 3 of 21 (14.3%) of genes in pham
- Manual Annotations of this start: 1 of 17
- Called 33.3% of time when present

• Phage (with cluster) where this start called: Anatole_31 (BV),

Start 4:

- Found in 14 of 21 (66.7%) of genes in pham
- Manual Annotations of this start: 9 of 17
- Called 85.7% of time when present

• Phage (with cluster) where this start called: BlackSpider_39 (FN), BrayBeast_36 (FB), Bridgette_41 (FA), ChuckDuck_42 (FA), Constance_41 (FA), Eileen_37 (FA), GlobiWarming_41 (FA), Judy_42 (FA), Kumotta_33 (FB), MargaretKali_33 (FB), Peas_38 (FA), RootBeer_34 (FA),

Start 5:

- Found in 4 of 21 (19.0%) of genes in pham
- Manual Annotations of this start: 1 of 17
- Called 25.0% of time when present
- Phage (with cluster) where this start called: B3_31 (BV),

Start 7:

- Found in 3 of 21 (14.3%) of genes in pham
- Manual Annotations of this start: 2 of 17
- Called 66.7% of time when present
- Phage (with cluster) where this start called: E1_31 (BV), E6_31 (BW),

Start 8:

- Found in 6 of 21 (28.6%) of genes in pham
- Manual Annotations of this start: 3 of 17
- Called 66.7% of time when present
- Phage (with cluster) where this start called: 32HC_49 (Z), B22_32 (BW), Doucette 34 (BW), Rem711 48 (Z),

Start 11:

- Found in 4 of 21 (19.0%) of genes in pham
- Manual Annotations of this start: 1 of 17
- Called 25.0% of time when present
- Phage (with cluster) where this start called: G4_32 (BW),

Summary by clusters:

There are 6 clusters represented in this pham: FB, FA, BV, BW, Z, FN,

Info for manual annotations of cluster BV:

- •Start number 3 was manually annotated 1 time for cluster BV.
- •Start number 5 was manually annotated 1 time for cluster BV.
- •Start number 7 was manually annotated 1 time for cluster BV.

Info for manual annotations of cluster BW:

- •Start number 7 was manually annotated 1 time for cluster BW.
- •Start number 8 was manually annotated 2 times for cluster BW.
- •Start number 11 was manually annotated 1 time for cluster BW.

Info for manual annotations of cluster FA:

•Start number 4 was manually annotated 6 times for cluster FA.

Info for manual annotations of cluster FB: •Start number 4 was manually annotated 3 times for cluster FB.

Info for manual annotations of cluster Z: •Start number 8 was manually annotated 1 time for cluster Z.

Gene Information:

Gene: 32HC_49 Start: 34204, Stop: 34452, Start Num: 8 Candidate Starts for 32HC_49: (Start: 4 @34177 has 9 MA's), (Start: 8 @34204 has 3 MA's), (9, 34213), (12, 34267), (13, 34282), (14, 34288), (20, 34363),

Gene: Anatole_31 Start: 23401, Stop: 23664, Start Num: 3 Candidate Starts for Anatole_31: (Start: 3 @23401 has 1 MA's), (6, 23413), (Start: 7 @23422 has 2 MA's), (10, 23440), (15, 23509), (16, 23521), (17, 23527), (21, 23590), (25, 23632),

Gene: B22_32 Start: 23397, Stop: 23636, Start Num: 8 Candidate Starts for B22_32: (1, 23325), (2, 23346), (Start: 5 @23376 has 1 MA's), (Start: 8 @23397 has 3 MA's), (Start: 11 @23433 has 1 MA's), (18, 23502), (19, 23526), (24, 23592),

Gene: B3_31 Start: 23506, Stop: 23766, Start Num: 5 Candidate Starts for B3_31: (Start: 5 @23506 has 1 MA's), (Start: 8 @23527 has 3 MA's), (Start: 11 @23563 has 1 MA's), (18, 23632), (19, 23656), (24, 23722),

Gene: BlackSpider_39 Start: 28835, Stop: 29107, Start Num: 4 Candidate Starts for BlackSpider_39: (Start: 4 @28835 has 9 MA's), (22, 29033), (26, 29084),

Gene: BrayBeast_36 Start: 26166, Stop: 26420, Start Num: 4 Candidate Starts for BrayBeast_36: (Start: 4 @26166 has 9 MA's), (23, 26370),

Gene: Bridgette_41 Start: 28919, Stop: 29173, Start Num: 4 Candidate Starts for Bridgette_41: (Start: 4 @28919 has 9 MA's), (13, 29003), (14, 29009),

Gene: ChuckDuck_42 Start: 28742, Stop: 28996, Start Num: 4 Candidate Starts for ChuckDuck_42: (Start: 4 @28742 has 9 MA's), (13, 28826), (14, 28832),

Gene: Constance_41 Start: 29071, Stop: 29325, Start Num: 4 Candidate Starts for Constance_41: (Start: 4 @29071 has 9 MA's), (13, 29155), (14, 29161),

Gene: Doucette_34 Start: 24811, Stop: 25050, Start Num: 8 Candidate Starts for Doucette_34: (1, 24739), (2, 24760), (Start: 5 @24790 has 1 MA's), (Start: 8 @24811 has 3 MA's), (Start: 11 @24847 has 1 MA's), (18, 24916), (19, 24940), (24, 25006),

Gene: E1_31 Start: 23422, Stop: 23664, Start Num: 7 Candidate Starts for E1_31: (Start: 3 @23401 has 1 MA's), (6, 23413), (Start: 7 @23422 has 2 MA's), (10, 23440), (15, 23509), (16, 23521), (17, 23527), (21, 23590), (25, 23632),

Gene: E6_31 Start: 23144, Stop: 23386, Start Num: 7 Candidate Starts for E6_31: (Start: 3 @23123 has 1 MA's), (6, 23135), (Start: 7 @23144 has 2 MA's), (10, 23162), (15, 23231), (16, 23243), (17, 23249), (21, 23312), (25, 23354),

Gene: Eileen_37 Start: 27149, Stop: 27403, Start Num: 4 Candidate Starts for Eileen_37: (Start: 4 @27149 has 9 MA's), (13, 27233), (14, 27239),

Gene: G4_32 Start: 24466, Stop: 24669, Start Num: 11 Candidate Starts for G4_32: (Start: 5 @24409 has 1 MA's), (Start: 8 @24430 has 3 MA's), (Start: 11 @24466 has 1 MA's), (18, 24535), (19, 24559), (24, 24625),

Gene: GlobiWarming_41 Start: 28325, Stop: 28579, Start Num: 4 Candidate Starts for GlobiWarming_41: (Start: 4 @28325 has 9 MA's), (13, 28409), (14, 28415),

Gene: Judy_42 Start: 29237, Stop: 29491, Start Num: 4 Candidate Starts for Judy_42: (Start: 4 @29237 has 9 MA's), (13, 29321), (14, 29327),

Gene: Kumotta_33 Start: 26166, Stop: 26420, Start Num: 4 Candidate Starts for Kumotta_33: (Start: 4 @26166 has 9 MA's), (14, 26256), (27, 26400), (28, 26409),

Gene: MargaretKali_33 Start: 25801, Stop: 26055, Start Num: 4 Candidate Starts for MargaretKali_33: (Start: 4 @25801 has 9 MA's), (14, 25891), (27, 26035), (28, 26044),

Gene: Peas_38 Start: 29296, Stop: 29550, Start Num: 4 Candidate Starts for Peas_38: (Start: 4 @29296 has 9 MA's), (13, 29380), (14, 29386),

Gene: Rem711_48 Start: 34188, Stop: 34436, Start Num: 8 Candidate Starts for Rem711_48: (Start: 4 @34161 has 9 MA's), (Start: 8 @34188 has 3 MA's), (9, 34197), (12, 34251), (13, 34266), (14, 34272), (20, 34347),

Gene: RootBeer_34 Start: 25199, Stop: 25453, Start Num: 4 Candidate Starts for RootBeer_34: (Start: 4 @25199 has 9 MA's), (13, 25283), (14, 25289),