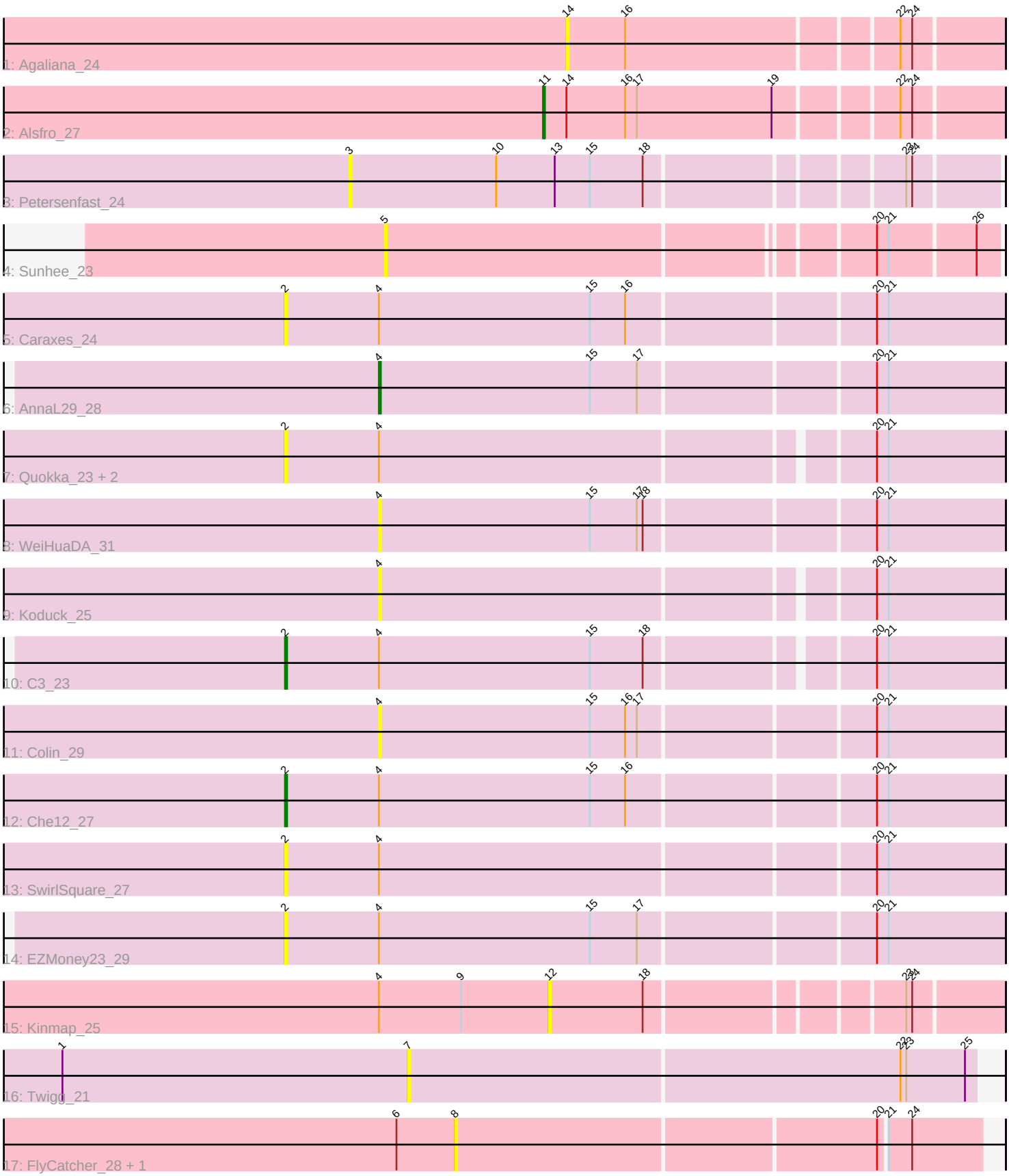
Pham 196841


1: Agaliana_24
2: Alsfro_27
3: Petersenfast_24
4: Sunhee_23
5: Caraxes_24
6: AnnaL29_28
7: Quokka_23 + 2
8: WeiHuaDA_31
9: Koduck_25
10: C3_23
11: Colin_29
12: Che12_27
13: SwirlSquare_27
14: EZMoney23_29
15: Kinmap_25
16: Twigg_21
17: FlyCatcher_28 + 1

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 196841 Report

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

Pham number 196841 has 20 members, 16 are drafts.

Phages represented in each track:
• Track 1 : Agaliana_24
• Track 2 : Alsfro_27
• Track 3 : Petersenfast_24
• Track 4 : Sunhee_23
• Track 5 : Caraxes_24
• Track 6 : AnnaL29_28
• Track 7 : Quokka_23, Bradman_24, MajorMajor_24
• Track 8 : WeiHuaDA_31
• Track 9 : Koduck_25
• Track 10 : C3_23
• Track 11 : Colin_29
• Track 12 : Che12_27
• Track 13 : SwirlSquare_27
• Track 14 : EZMoney23_29
• Track 15 : Kinmap_25
• Track 16 : Twigg_21
• Track 17 : FlyCatcher_28, Toro_27

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

Genes that call this "Most Annotated" start:
• Bradman_24, C3_23, Caraxes_24, Che12_27, EZMoney23_29, MajorMajor_24, Quokka_23, SwirlSquare_27,

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

Genes that do not have the "Most Annotated" start:
• Agaliana_24, Alsfro_27, AnnaL29_28, Colin_29, FlyCatcher_28, Kinmap_25, Koduck_25, Petersenfast_24, Sunhee_23, Toro_27, Twigg_21, WeiHuaDA_31,

Summary by start number:

Start 2:
• Found in 8 of 20 (40.0%) of genes in pham
• Manual Annotations of this start: 2 of 4
• Called 100.0% of time when present
• Phage (with cluster) where this start called: Bradman_24 (A2), C3_23 (A2), Caraxes_24 (A2), Che12_27 (A2), EZMoney23_29 (A2), MajorMajor_24 (A2), Quokka_23 (A2), SwirlSquare_27 (A2),

Start 3:
• Found in 1 of 20 (5.0%) of genes in pham
• No Manual Annotations of this start.
• Called 100.0% of time when present
• Phage (with cluster) where this start called: Petersenfast_24 (A11),

Start 4:
• Found in 13 of 20 (65.0%) of genes in pham
• Manual Annotations of this start: 1 of 4
• Called 30.8% of time when present
• Phage (with cluster) where this start called: AnnaL29_28 (A2), Colin_29 (A2), Koduck_25 (A2), WeiHuaDA_31 (A2),

Start 5:
• Found in 1 of 20 (5.0%) of genes in pham
• No Manual Annotations of this start.
• Called 100.0% of time when present
• Phage (with cluster) where this start called: Sunhee_23 (A14),

Start 7:
• Found in 1 of 20 (5.0%) of genes in pham
• No Manual Annotations of this start.
• Called 100.0% of time when present
• Phage (with cluster) where this start called: Twigg_21 (A5),

Start 8:
• Found in 2 of 20 (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: FlyCatcher_28 (A7), Toro_27 (A7),

Start 11:
• Found in 1 of 20 (5.0%) 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: Alsfro_27 (A1),

Start 12:
• Found in 1 of 20 (5.0%) of genes in pham
• No Manual Annotations of this start.
• Called 100.0% of time when present
• Phage (with cluster) where this start called: Kinmap_25 (A21),

Start 14:
• Found in 2 of 20 (10.0%) of genes in pham
• No Manual Annotations of this start.
• Called 50.0% of time when present
• Phage (with cluster) where this start called: Agaliana_24 (A1),

Summary by clusters:

There are 7 clusters represented in this pham: A21, A11, A14, A1, A2, A5, A7,

Info for manual annotations of cluster A1:
•Start number 11 was manually annotated 1 time for cluster A1.

Info for manual annotations of cluster A2:
•Start number 2 was manually annotated 2 times for cluster A2.
•Start number 4 was manually annotated 1 time for cluster A2.

Gene Information:

Gene: Agaliana_24 Start: 16414, Stop: 16626, Start Num: 14
Candidate Starts for Agaliana_24:
(14, 16414), (16, 16444), (22, 16576), (24, 16582),

Gene: Alsfro_27 Start: 16894, Stop: 17118, Start Num: 11
Candidate Starts for Alsfro_27:
(Start: 11 @16894 has 1 MA's), (14, 16906), (16, 16936), (17, 16942), (19, 17011), (22, 17068), (24, 17074),

Gene: AnnaL29_28 Start: 18178, Stop: 18489, Start Num: 4
Candidate Starts for AnnaL29_28:
(Start: 4 @18178 has 1 MA's), (15, 18286), (17, 18310), (20, 18424), (21, 18430),

Gene: Bradman_24 Start: 15185, Stop: 15538, Start Num: 2
Candidate Starts for Bradman_24:
(Start: 2 @15185 has 2 MA's), (Start: 4 @15233 has 1 MA's), (20, 15473), (21, 15479),

Gene: C3_23 Start: 16308, Stop: 16661, Start Num: 2
Candidate Starts for C3_23:
(Start: 2 @16308 has 2 MA's), (Start: 4 @16356 has 1 MA's), (15, 16464), (18, 16491), (20, 16596), (21, 16602),

Gene: Caraxes_24 Start: 14918, Stop: 15277, Start Num: 2
Candidate Starts for Caraxes_24:
(Start: 2 @14918 has 2 MA's), (Start: 4 @14966 has 1 MA's), (15, 15074), (16, 15092), (20, 15212), (21, 15218),

Gene: Che12_27 Start: 15178, Stop: 15537, Start Num: 2
Candidate Starts for Che12_27:
(Start: 2 @15178 has 2 MA's), (Start: 4 @15226 has 1 MA's), (15, 15334), (16, 15352), (20, 15472), (21, 15478),

Gene: Colin_29 Start: 18010, Stop: 18321, Start Num: 4
Candidate Starts for Colin_29:
(Start: 4 @18010 has 1 MA's), (15, 18118), (16, 18136), (17, 18142), (20, 18256), (21, 18262),

Gene: EZMoney23_29 Start: 18248, Stop: 18607, Start Num: 2
Candidate Starts for EZMoney23_29:
(Start: 2 @18248 has 2 MA's), (Start: 4 @18296 has 1 MA's), (15, 18404), (17, 18428), (20, 18542), (21, 18548),

Gene: FlyCatcher_28 Start: 17296, Stop: 17556, Start Num: 8
Candidate Starts for FlyCatcher_28:
(6, 17266), (8, 17296), (20, 17506), (21, 17509), (24, 17521),

Gene: Kinmap_25 Start: 16202, Stop: 16417, Start Num: 12
Candidate Starts for Kinmap_25:
(Start: 4 @16115 has 1 MA's), (9, 16157), (12, 16202), (18, 16250), (23, 16370), (24, 16373),

Gene: Koduck_25 Start: 15488, Stop: 15793, Start Num: 4
Candidate Starts for Koduck_25:
(Start: 4 @15488 has 1 MA's), (20, 15728), (21, 15734),

Gene: MajorMajor_24 Start: 15185, Stop: 15538, Start Num: 2
Candidate Starts for MajorMajor_24:
(Start: 2 @15185 has 2 MA's), (Start: 4 @15233 has 1 MA's), (20, 15473), (21, 15479),

Gene: Petersenfast_24 Start: 16853, Stop: 17167, Start Num: 3
Candidate Starts for Petersenfast_24:
(3, 16853), (10, 16928), (13, 16958), (15, 16976), (18, 17003), (23, 17123), (24, 17126),

Gene: Quokka_23 Start: 15185, Stop: 15538, Start Num: 2
Candidate Starts for Quokka_23:
(Start: 2 @15185 has 2 MA's), (Start: 4 @15233 has 1 MA's), (20, 15473), (21, 15479),

Gene: Sunhee_23 Start: 15505, Stop: 15801, Start Num: 5
Candidate Starts for Sunhee_23:
(5, 15505), (20, 15742), (21, 15748), (26, 15790),

Gene: SwirlSquare_27 Start: 15091, Stop: 15450, Start Num: 2
Candidate Starts for SwirlSquare_27:
(Start: 2 @15091 has 2 MA's), (Start: 4 @15139 has 1 MA's), (20, 15385), (21, 15391),

Gene: Toro_27 Start: 17296, Stop: 17556, Start Num: 8
Candidate Starts for Toro_27:
(6, 17266), (8, 17296), (20, 17506), (21, 17509), (24, 17521),

Gene: Twigg_21 Start: 14923, Stop: 15210, Start Num: 7
Candidate Starts for Twigg_21:
(1, 14746), (7, 14923), (22, 15172), (23, 15175), (25, 15205),

Gene: WeiHuaDA_31 Start: 18580, Stop: 18891, Start Num: 4
Candidate Starts for WeiHuaDA_31:
(Start: 4 @18580 has 1 MA's), (15, 18688), (17, 18712), (18, 18715), (20, 18826), (21, 18832),