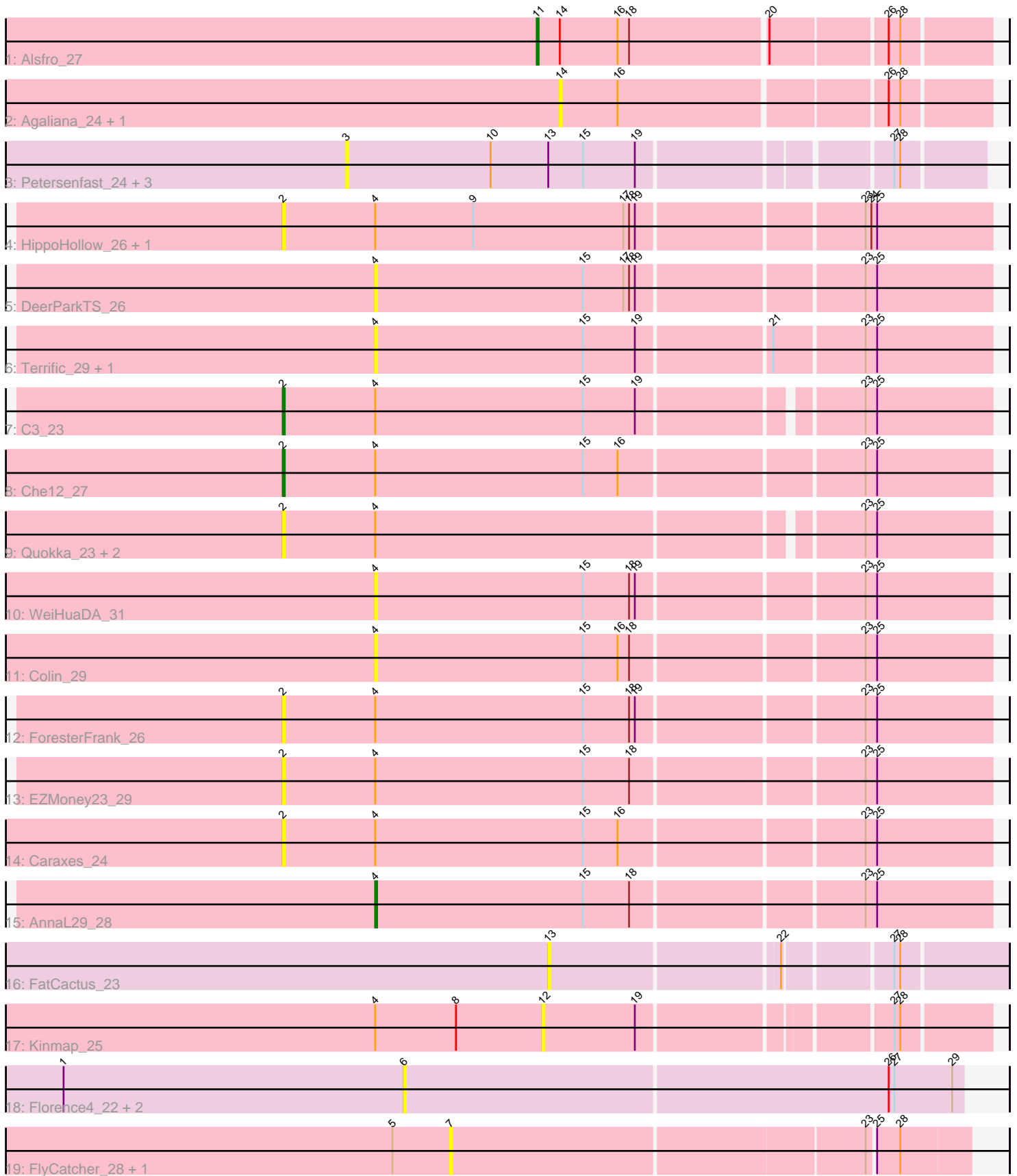


Pham 311876



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 311876 Report

This analysis was run 06/27/26 on database version 652.

Pham number 311876 has 30 members, 26 are drafts.

Phages represented in each track:

- Track 1 : Alsfro\_27
- Track 2 : Agaliana\_24, ChristmasHams\_24
- Track 3 : Petersenfast\_24, Saskia\_24, Wolpertinger\_24, Kimba\_25
- Track 4 : HippoHollow\_26, LionsBait\_26
- Track 5 : DeerParkTS\_26
- Track 6 : Terrific\_29, Gratitude\_26
- Track 7 : C3\_23
- Track 8 : Che12\_27
- Track 9 : Quokka\_23, Bradman\_24, MajorMajor\_24
- Track 10 : WeiHuaDA\_31
- Track 11 : Colin\_29
- Track 12 : ForesterFrank\_26
- Track 13 : EZMoney23\_29
- Track 14 : Caraxes\_24
- Track 15 : AnnaL29\_28
- Track 16 : FatCactus\_23
- Track 17 : Kinmap\_25
- Track 18 : Florence4\_22, Prinashe11\_22, Florence1\_21
- Track 19 : 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, ForesterFrank\_26, HippoHollow\_26, LionsBait\_26, MajorMajor\_24, Quokka\_23,

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, ChristmasHams\_24, Colin\_29, DeerParkTS\_26, FatCactus\_23, Florence1\_21, Florence4\_22, FlyCatcher\_28, Gratitude\_26, Kimba\_25, Kinmap\_25, Petersenfast\_24, Prinashe11\_22, Saskia\_24, Terrific\_29, Toro\_27, WeiHuaDA\_31, Wolpertinger\_24,

### Summary by start number:

#### Start 2:

- Found in 10 of 30 ( 33.3% ) 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), ForesterFrank\_26 (A2), HippoHollow\_26 (A2), LionsBait\_26 (A2), MajorMajor\_24 (A2), Quokka\_23 (A2),

#### Start 3:

- Found in 4 of 30 ( 13.3% ) of genes in pham
- No Manual Annotations of this start.
- Called 100.0% of time when present
- Phage (with cluster) where this start called: Kimba\_25 (A11), Petersenfast\_24 (A11), Saskia\_24 (A11), Wolpertinger\_24 (A11),

#### Start 4:

- Found in 17 of 30 ( 56.7% ) of genes in pham
- Manual Annotations of this start: 1 of 4
- Called 35.3% of time when present
- Phage (with cluster) where this start called: AnnaL29\_28 (A2), Colin\_29 (A2), DeerParkTS\_26 (A2), Gratitude\_26 (A2), Terrific\_29 (A2), WeiHuaDA\_31 (A2),

#### Start 6:

- Found in 3 of 30 ( 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: Florence1\_21 (A5), Florence4\_22 (A5), Prinashe11\_22 (A5),

#### Start 7:

- Found in 2 of 30 ( 6.7% ) 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 30 ( 3.3% ) 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: Alsfro\_27 (A1),

#### Start 12:

- Found in 1 of 30 ( 3.3% ) 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 13:

- Found in 5 of 30 ( 16.7% ) of genes in pham
- No Manual Annotations of this start.
- Called 20.0% of time when present
- Phage (with cluster) where this start called: FatCactus\_23 (A20),

Start 14:

- Found in 3 of 30 ( 10.0% ) of genes in pham
- No Manual Annotations of this start.
- Called 66.7% of time when present
- Phage (with cluster) where this start called: Agaliana\_24 (A1), ChristmasHams\_24 (A1),

**Summary by clusters:**

There are 7 clusters represented in this pham: A20, A21, A11, 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), (26, 16576), (28, 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), (18, 16942), (20, 17011), (26, 17068), (28, 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), (18, 18310), (23, 18424), (25, 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), (23, 15473), (25, 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), (19, 16491), (23, 16596), (25, 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), (23, 15212), (25, 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), (23, 15472), (25, 15478),

Gene: ChristmasHams\_24 Start: 16173, Stop: 16385, Start Num: 14

Candidate Starts for ChristmasHams\_24:

(14, 16173), (16, 16203), (26, 16335), (28, 16341),

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), (18, 18142), (23, 18256), (25, 18262),

Gene: DeerParkTS\_26 Start: 16386, Stop: 16697, Start Num: 4

Candidate Starts for DeerParkTS\_26:

(Start: 4 @16386 has 1 MA's), (15, 16494), (17, 16515), (18, 16518), (19, 16521), (23, 16632), (25, 16638),

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), (18, 18428), (23, 18542), (25, 18548),

Gene: FatCactus\_23 Start: 14466, Stop: 14687, Start Num: 13

Candidate Starts for FatCactus\_23:

(13, 14466), (22, 14580), (27, 14631), (28, 14634),

Gene: Florence1\_21 Start: 14740, Stop: 15027, Start Num: 6

Candidate Starts for Florence1\_21:

(1, 14563), (6, 14740), (26, 14989), (27, 14992), (29, 15022),

Gene: Florence4\_22 Start: 14740, Stop: 15027, Start Num: 6

Candidate Starts for Florence4\_22:

(1, 14563), (6, 14740), (26, 14989), (27, 14992), (29, 15022),

Gene: FlyCatcher\_28 Start: 17296, Stop: 17556, Start Num: 7

Candidate Starts for FlyCatcher\_28:

(5, 17266), (7, 17296), (23, 17506), (25, 17509), (28, 17521),

Gene: ForesterFrank\_26 Start: 17788, Stop: 18147, Start Num: 2

Candidate Starts for ForesterFrank\_26:

(Start: 2 @17788 has 2 MA's), (Start: 4 @17836 has 1 MA's), (15, 17944), (18, 17968), (19, 17971), (23, 18082), (25, 18088),

Gene: Gratitude\_26 Start: 16814, Stop: 17125, Start Num: 4

Candidate Starts for Gratitude\_26:

(Start: 4 @16814 has 1 MA's), (15, 16922), (19, 16949), (21, 17015), (23, 17060), (25, 17066),

Gene: HippoHollow\_26 Start: 18036, Stop: 18395, Start Num: 2

Candidate Starts for HippoHollow\_26:

(Start: 2 @18036 has 2 MA's), (Start: 4 @18084 has 1 MA's), (9, 18135), (17, 18213), (18, 18216), (19, 18219), (23, 18330), (24, 18333), (25, 18336),

Gene: Kimba\_25 Start: 16908, Stop: 17222, Start Num: 3

Candidate Starts for Kimba\_25:

(3, 16908), (10, 16983), (13, 17013), (15, 17031), (19, 17058), (27, 17178), (28, 17181),

Gene: Kinmap\_25 Start: 16202, Stop: 16417, Start Num: 12

Candidate Starts for Kinmap\_25:

(Start: 4 @16115 has 1 MA's), (8, 16157), (12, 16202), (19, 16250), (27, 16370), (28, 16373),

Gene: LionsBait\_26 Start: 18030, Stop: 18389, Start Num: 2

Candidate Starts for LionsBait\_26:

(Start: 2 @18030 has 2 MA's), (Start: 4 @18078 has 1 MA's), (9, 18129), (17, 18207), (18, 18210), (19, 18213), (23, 18324), (24, 18327), (25, 18330),

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), (23, 15473), (25, 15479),

Gene: Petersenfast\_24 Start: 16853, Stop: 17167, Start Num: 3

Candidate Starts for Petersenfast\_24:

(3, 16853), (10, 16928), (13, 16958), (15, 16976), (19, 17003), (27, 17123), (28, 17126),

Gene: Prinashe11\_22 Start: 14920, Stop: 15207, Start Num: 6

Candidate Starts for Prinashe11\_22:

(1, 14743), (6, 14920), (26, 15169), (27, 15172), (29, 15202),

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), (23, 15473), (25, 15479),

Gene: Saskia\_24 Start: 16850, Stop: 17164, Start Num: 3

Candidate Starts for Saskia\_24:

(3, 16850), (10, 16925), (13, 16955), (15, 16973), (19, 17000), (27, 17120), (28, 17123),

Gene: Terrific\_29 Start: 16826, Stop: 17137, Start Num: 4

Candidate Starts for Terrific\_29:

(Start: 4 @16826 has 1 MA's), (15, 16934), (19, 16961), (21, 17027), (23, 17072), (25, 17078),

Gene: Toro\_27 Start: 17296, Stop: 17556, Start Num: 7

Candidate Starts for Toro\_27:

(5, 17266), (7, 17296), (23, 17506), (25, 17509), (28, 17521),

Gene: WeiHuaDA\_31 Start: 18580, Stop: 18891, Start Num: 4

Candidate Starts for WeiHuaDA\_31:

(Start: 4 @18580 has 1 MA's), (15, 18688), (18, 18712), (19, 18715), (23, 18826), (25, 18832),

Gene: Wolpertinger\_24 Start: 16853, Stop: 17167, Start Num: 3

Candidate Starts for Wolpertinger\_24:

(3, 16853), (10, 16928), (13, 16958), (15, 16976), (19, 17003), (27, 17123), (28, 17126),