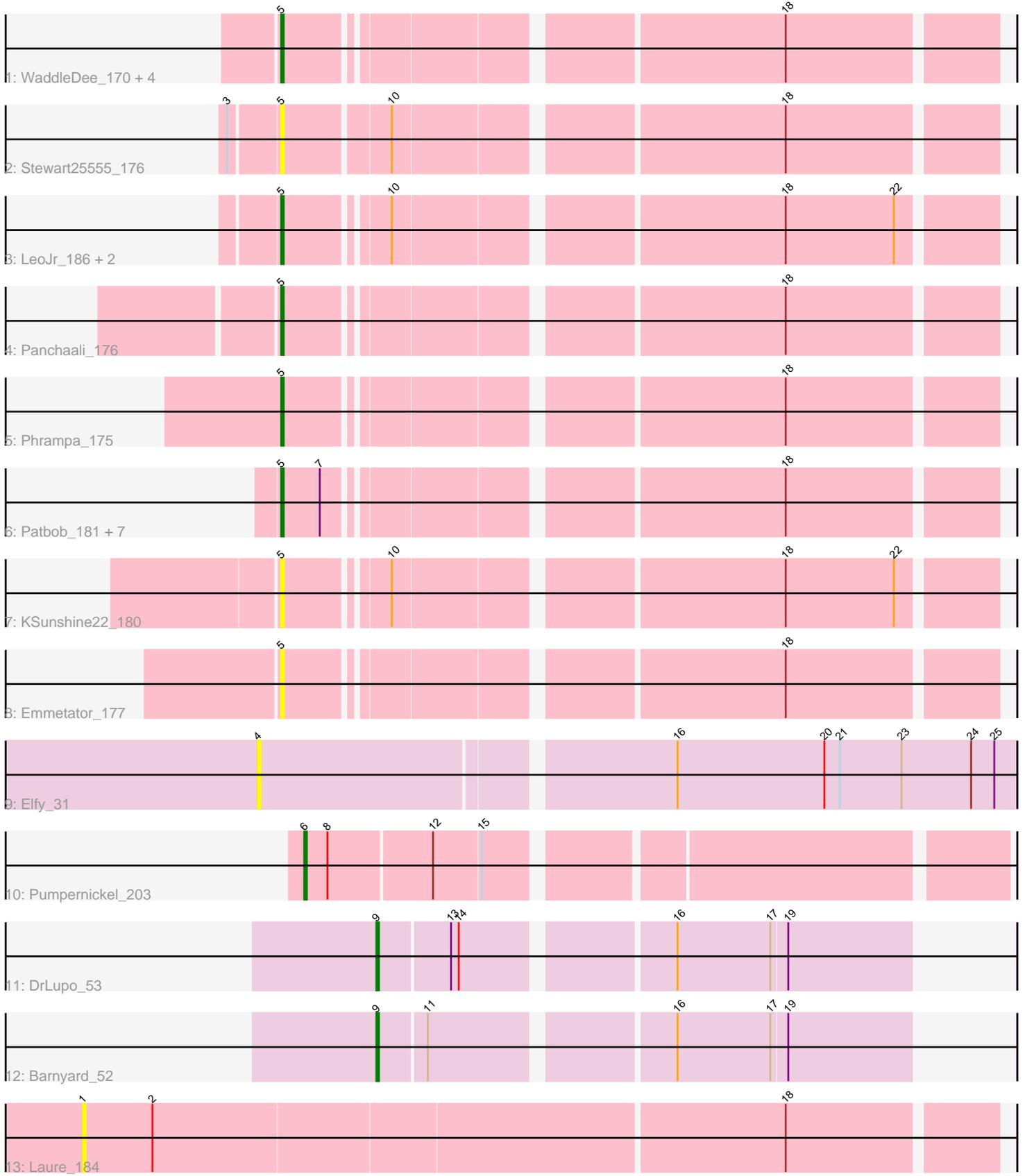


Pham 291455



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 291455 Report

This analysis was run 03/28/26 on database version 641.

Pham number 291455 has 26 members, 13 are drafts.

Phages represented in each track:

- Track 1 : WaddleDee\_170, BooTeria\_182, DunneganBoMo\_174, Ellewin\_180, Artu\_177
- Track 2 : Stewart25555\_176
- Track 3 : LeoJr\_186, ReginaGlobina\_190, Atuin\_177
- Track 4 : Panchaali\_176
- Track 5 : Phrampa\_175
- Track 6 : Patbob\_181, GoldenEssence\_170, Bloom\_187, FloraSnap32\_180, Racecar\_184, Talia1610\_184, Mimi\_183, FrostedClock\_186
- Track 7 : KSunshine22\_180
- Track 8 : Emmetator\_177
- Track 9 : Elfy\_31
- Track 10 : Pumpernickel\_203
- Track 11 : DrLupo\_53
- Track 12 : Barnyard\_52
- Track 13 : Laure\_184

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

The start number called the most often in the published annotations is 5, it was called in 10 of the 13 non-draft genes in the pham.

Genes that call this "Most Annotated" start:

- Artu\_177, Atuin\_177, Bloom\_187, BooTeria\_182, DunneganBoMo\_174, Ellewin\_180, Emmetator\_177, FloraSnap32\_180, FrostedClock\_186, GoldenEssence\_170, KSunshine22\_180, LeoJr\_186, Mimi\_183, Panchaali\_176, Patbob\_181, Phrampa\_175, Racecar\_184, ReginaGlobina\_190, Stewart25555\_176, Talia1610\_184, WaddleDee\_170,

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

- 

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

- Barnyard\_52, DrLupo\_53, Elfy\_31, Laure\_184, Pumpernickel\_203,

## Summary by start number:

### Start 1:

- Found in 1 of 26 ( 3.8% ) of genes in pham
- No Manual Annotations of this start.
- Called 100.0% of time when present
- Phage (with cluster) where this start called: Laure\_184 (UNK),

### Start 4:

- Found in 1 of 26 ( 3.8% ) of genes in pham
- No Manual Annotations of this start.
- Called 100.0% of time when present
- Phage (with cluster) where this start called: Elfy\_31 (FP),

### Start 5:

- Found in 21 of 26 ( 80.8% ) of genes in pham
- Manual Annotations of this start: 10 of 13
- Called 100.0% of time when present
- Phage (with cluster) where this start called: Artu\_177 (FC), Atuin\_177 (FC), Bloom\_187 (FC), BooTeria\_182 (FC), DunneganBoMo\_174 (FC), Ellewin\_180 (FC), Emmetator\_177 (FC), FloraSnap32\_180 (FC), FrostedClock\_186 (FC), GoldenEssence\_170 (FC), KSunshine22\_180 (FC), LeoJr\_186 (FC), Mimi\_183 (FC), Panchaali\_176 (FC), Patbob\_181 (FC), Phrampa\_175 (FC), Racecar\_184 (FC), ReginaGlobina\_190 (FC), Stewart25555\_176 (FC), Talia1610\_184 (FC), WaddleDee\_170 (FC),

### Start 6:

- Found in 1 of 26 ( 3.8% ) of genes in pham
- Manual Annotations of this start: 1 of 13
- Called 100.0% of time when present
- Phage (with cluster) where this start called: Pumpernickel\_203 (GD4),

### Start 9:

- Found in 2 of 26 ( 7.7% ) of genes in pham
- Manual Annotations of this start: 2 of 13
- Called 100.0% of time when present
- Phage (with cluster) where this start called: Barnyard\_52 (H2), DrLupo\_53 (H2),

## Summary by clusters:

There are 5 clusters represented in this pham: H2, FP, UNK, FC, GD4,

Info for manual annotations of cluster FC:

- Start number 5 was manually annotated 10 times for cluster FC.

Info for manual annotations of cluster GD4:

- Start number 6 was manually annotated 1 time for cluster GD4.

Info for manual annotations of cluster H2:

- Start number 9 was manually annotated 2 times for cluster H2.

**Gene Information:**

Gene: Artu\_177 Start: 120610, Stop: 120864, Start Num: 5

Candidate Starts for Artu\_177:

(Start: 5 @120610 has 10 MA's), (18, 120787),

Gene: Atuin\_177 Start: 122575, Stop: 122829, Start Num: 5

Candidate Starts for Atuin\_177:

(Start: 5 @122575 has 10 MA's), (10, 122611), (18, 122752), (22, 122794),

Gene: Barnyard\_52 Start: 39091, Stop: 39285, Start Num: 9

Candidate Starts for Barnyard\_52:

(Start: 9 @39091 has 2 MA's), (11, 39109), (16, 39196), (17, 39232), (19, 39238),

Gene: Bloom\_187 Start: 122149, Stop: 122403, Start Num: 5

Candidate Starts for Bloom\_187:

(Start: 5 @122149 has 10 MA's), (7, 122164), (18, 122326),

Gene: BooTeria\_182 Start: 120032, Stop: 120286, Start Num: 5

Candidate Starts for BooTeria\_182:

(Start: 5 @120032 has 10 MA's), (18, 120209),

Gene: DrLupo\_53 Start: 39502, Stop: 39696, Start Num: 9

Candidate Starts for DrLupo\_53:

(Start: 9 @39502 has 2 MA's), (13, 39529), (14, 39532), (16, 39607), (17, 39643), (19, 39649),

Gene: DunneganBoMo\_174 Start: 119214, Stop: 119468, Start Num: 5

Candidate Starts for DunneganBoMo\_174:

(Start: 5 @119214 has 10 MA's), (18, 119391),

Gene: Elfy\_31 Start: 23896, Stop: 24180, Start Num: 4

Candidate Starts for Elfy\_31:

(4, 23896), (16, 24049), (20, 24106), (21, 24112), (23, 24136), (24, 24163), (25, 24172),

Gene: Ellewin\_180 Start: 119388, Stop: 119642, Start Num: 5

Candidate Starts for Ellewin\_180:

(Start: 5 @119388 has 10 MA's), (18, 119565),

Gene: Emmetator\_177 Start: 119534, Stop: 119788, Start Num: 5

Candidate Starts for Emmetator\_177:

(Start: 5 @119534 has 10 MA's), (18, 119711),

Gene: FloraSnap32\_180 Start: 120480, Stop: 120734, Start Num: 5

Candidate Starts for FloraSnap32\_180:

(Start: 5 @120480 has 10 MA's), (7, 120495), (18, 120657),

Gene: FrostedClock\_186 Start: 122724, Stop: 122978, Start Num: 5

Candidate Starts for FrostedClock\_186:

(Start: 5 @122724 has 10 MA's), (7, 122739), (18, 122901),

Gene: GoldenEssence\_170 Start: 115722, Stop: 115976, Start Num: 5

Candidate Starts for GoldenEssence\_170:  
(Start: 5 @115722 has 10 MA's), (7, 115737), (18, 115899),

Gene: KSunshine22\_180 Start: 121028, Stop: 121282, Start Num: 5  
Candidate Starts for KSunshine22\_180:  
(Start: 5 @121028 has 10 MA's), (10, 121064), (18, 121205), (22, 121247),

Gene: Laure\_184 Start: 113724, Stop: 114068, Start Num: 1  
Candidate Starts for Laure\_184:  
(1, 113724), (2, 113751), (18, 113991),

Gene: LeoJr\_186 Start: 123129, Stop: 123383, Start Num: 5  
Candidate Starts for LeoJr\_186:  
(Start: 5 @123129 has 10 MA's), (10, 123165), (18, 123306), (22, 123348),

Gene: Mimi\_183 Start: 121776, Stop: 122030, Start Num: 5  
Candidate Starts for Mimi\_183:  
(Start: 5 @121776 has 10 MA's), (7, 121791), (18, 121953),

Gene: Panchaali\_176 Start: 120271, Stop: 120525, Start Num: 5  
Candidate Starts for Panchaali\_176:  
(Start: 5 @120271 has 10 MA's), (18, 120448),

Gene: Patbob\_181 Start: 122338, Stop: 122592, Start Num: 5  
Candidate Starts for Patbob\_181:  
(Start: 5 @122338 has 10 MA's), (7, 122353), (18, 122515),

Gene: Phrampa\_175 Start: 123930, Stop: 124184, Start Num: 5  
Candidate Starts for Phrampa\_175:  
(Start: 5 @123930 has 10 MA's), (18, 124107),

Gene: Pumpernickel\_203 Start: 117028, Stop: 116777, Start Num: 6  
Candidate Starts for Pumpernickel\_203:  
(Start: 6 @117028 has 1 MA's), (8, 117019), (12, 116980), (15, 116962),

Gene: Racecar\_184 Start: 122742, Stop: 122996, Start Num: 5  
Candidate Starts for Racecar\_184:  
(Start: 5 @122742 has 10 MA's), (7, 122757), (18, 122919),

Gene: ReginaGlobina\_190 Start: 124398, Stop: 124652, Start Num: 5  
Candidate Starts for ReginaGlobina\_190:  
(Start: 5 @124398 has 10 MA's), (10, 124434), (18, 124575), (22, 124617),

Gene: Stewart25555\_176 Start: 121983, Stop: 122240, Start Num: 5  
Candidate Starts for Stewart25555\_176:  
(3, 121965), (Start: 5 @121983 has 10 MA's), (10, 122022), (18, 122163),

Gene: Talia1610\_184 Start: 122159, Stop: 122413, Start Num: 5  
Candidate Starts for Talia1610\_184:  
(Start: 5 @122159 has 10 MA's), (7, 122174), (18, 122336),

Gene: WaddleDee\_170 Start: 118487, Stop: 118741, Start Num: 5  
Candidate Starts for WaddleDee\_170:

(Start: 5 @118487 has 10 MA's), (18, 118664),