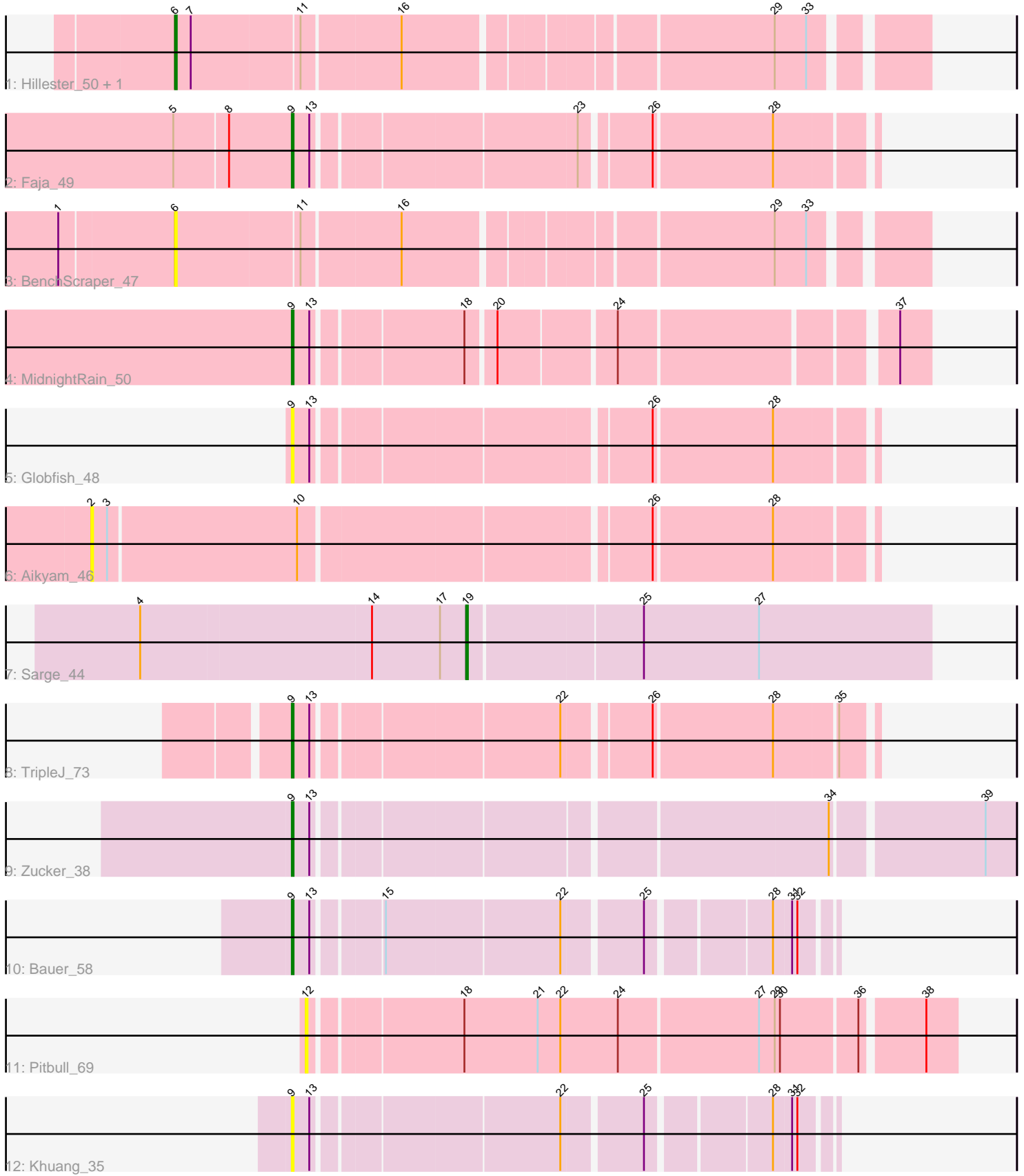


Pham 156781



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 156781 Report

This analysis was run 04/12/24 on database version 558.

Pham number 156781 has 13 members, 6 are drafts.

Phages represented in each track:

- Track 1 : Hillester\_50, RadFad\_50
- Track 2 : Faja\_49
- Track 3 : BenchScraper\_47
- Track 4 : MidnightRain\_50
- Track 5 : Globfish\_48
- Track 6 : Aikyam\_46
- Track 7 : Sarge\_44
- Track 8 : TripleJ\_73
- Track 9 : Zucker\_38
- Track 10 : Bauer\_58
- Track 11 : Pitbull\_69
- Track 12 : Khuang\_35

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

The start number called the most often in the published annotations is 9, it was called in 5 of the 7 non-draft genes in the pham.

Genes that call this "Most Annotated" start:

- Bauer\_58, Faja\_49, Globfish\_48, Khuang\_35, MidnightRain\_50, TripleJ\_73, Zucker\_38,

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

- 

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

- Aikyam\_46, BenchScraper\_47, Hillester\_50, Pitbull\_69, RadFad\_50, Sarge\_44,

### **Summary by start number:**

Start 2:

- Found in 1 of 13 ( 7.7% ) of genes in pham
- No Manual Annotations of this start.
- Called 100.0% of time when present

- Phage (with cluster) where this start called: Aikyam\_46 (AY),

Start 6:

- Found in 3 of 13 ( 23.1% ) of genes in pham
- Manual Annotations of this start: 1 of 7
- Called 100.0% of time when present
- Phage (with cluster) where this start called: BenchScraper\_47 (AY), Hillester\_50 (AY), RadFad\_50 (AY),

Start 9:

- Found in 7 of 13 ( 53.8% ) of genes in pham
- Manual Annotations of this start: 5 of 7
- Called 100.0% of time when present
- Phage (with cluster) where this start called: Bauer\_58 (FN), Faja\_49 (AY), Globfish\_48 (AY), Khuang\_35 (UNK), MidnightRain\_50 (AY), TripleJ\_73 (FJ), Zucker\_38 (FN),

Start 12:

- Found in 1 of 13 ( 7.7% ) of genes in pham
- No Manual Annotations of this start.
- Called 100.0% of time when present
- Phage (with cluster) where this start called: Pitbull\_69 (FQ),

Start 19:

- Found in 1 of 13 ( 7.7% ) of genes in pham
- Manual Annotations of this start: 1 of 7
- Called 100.0% of time when present
- Phage (with cluster) where this start called: Sarge\_44 (FB),

### **Summary by clusters:**

There are 6 clusters represented in this pham: FQ, FB, AY, FJ, UNK, FN,

Info for manual annotations of cluster AY:

- Start number 6 was manually annotated 1 time for cluster AY.
- Start number 9 was manually annotated 2 times for cluster AY.

Info for manual annotations of cluster FB:

- Start number 19 was manually annotated 1 time for cluster FB.

Info for manual annotations of cluster FJ:

- Start number 9 was manually annotated 1 time for cluster FJ.

Info for manual annotations of cluster FN:

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

### **Gene Information:**

Gene: Aikyam\_46 Start: 27868, Stop: 28284, Start Num: 2

Candidate Starts for Aikyam\_46:  
(2, 27868), (3, 27877), (10, 27982), (26, 28168), (28, 28234),

Gene: Bauer\_58 Start: 32955, Stop: 33227, Start Num: 9  
Candidate Starts for Bauer\_58:  
(Start: 9 @32955 has 5 MA's), (13, 32964), (15, 32997), (22, 33093), (25, 33135), (28, 33198), (31, 33207), (32, 33210),

Gene: BenchScraper\_47 Start: 29609, Stop: 30001, Start Num: 6  
Candidate Starts for BenchScraper\_47:  
(1, 29546), (Start: 6 @29609 has 1 MA's), (11, 29678), (16, 29732), (29, 29927), (33, 29945),

Gene: Faja\_49 Start: 31772, Stop: 32071, Start Num: 9  
Candidate Starts for Faja\_49:  
(5, 31706), (8, 31736), (Start: 9 @31772 has 5 MA's), (13, 31781), (23, 31919), (26, 31955), (28, 32021),

Gene: Globfish\_48 Start: 30153, Stop: 30452, Start Num: 9  
Candidate Starts for Globfish\_48:  
(Start: 9 @30153 has 5 MA's), (13, 30162), (26, 30336), (28, 30402),

Gene: Hillester\_50 Start: 30882, Stop: 31274, Start Num: 6  
Candidate Starts for Hillester\_50:  
(Start: 6 @30882 has 1 MA's), (7, 30891), (11, 30951), (16, 31005), (29, 31200), (33, 31218),

Gene: Khuang\_35 Start: 24689, Stop: 24417, Start Num: 9  
Candidate Starts for Khuang\_35:  
(Start: 9 @24689 has 5 MA's), (13, 24680), (22, 24551), (25, 24509), (28, 24446), (31, 24437), (32, 24434),

Gene: MidnightRain\_50 Start: 31295, Stop: 31621, Start Num: 9  
Candidate Starts for MidnightRain\_50:  
(Start: 9 @31295 has 5 MA's), (13, 31304), (18, 31382), (20, 31397), (24, 31460), (37, 31604),

Gene: Pitbull\_69 Start: 37588, Stop: 37938, Start Num: 12  
Candidate Starts for Pitbull\_69:  
(12, 37588), (18, 37669), (21, 37711), (22, 37723), (24, 37756), (27, 37834), (29, 37843), (30, 37846), (36, 37888), (38, 37921),

Gene: RadFad\_50 Start: 30882, Stop: 31274, Start Num: 6  
Candidate Starts for RadFad\_50:  
(Start: 6 @30882 has 1 MA's), (7, 30891), (11, 30951), (16, 31005), (29, 31200), (33, 31218),

Gene: Sarge\_44 Start: 27015, Stop: 27275, Start Num: 19  
Candidate Starts for Sarge\_44:  
(4, 26832), (14, 26961), (17, 27000), (Start: 19 @27015 has 1 MA's), (25, 27111), (27, 27177),

Gene: TripleJ\_73 Start: 41236, Stop: 41535, Start Num: 9  
Candidate Starts for TripleJ\_73:  
(Start: 9 @41236 has 5 MA's), (13, 41245), (22, 41374), (26, 41419), (28, 41485), (35, 41518),

Gene: Zucker\_38 Start: 28357, Stop: 28737, Start Num: 9  
Candidate Starts for Zucker\_38:

(Start: 9 @28357 has 5 MA's), (13, 28366), (34, 28639), (39, 28720),