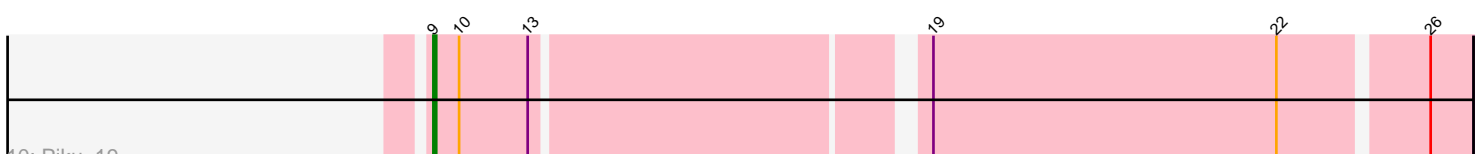
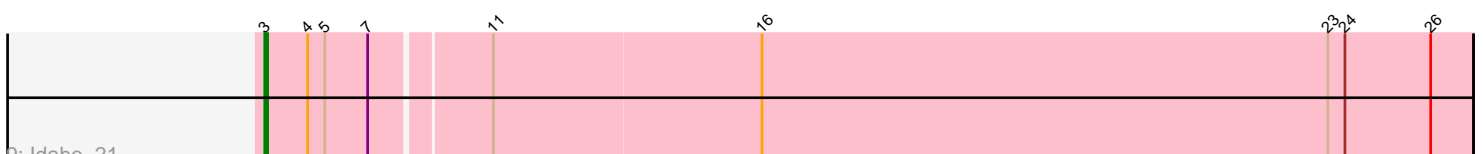
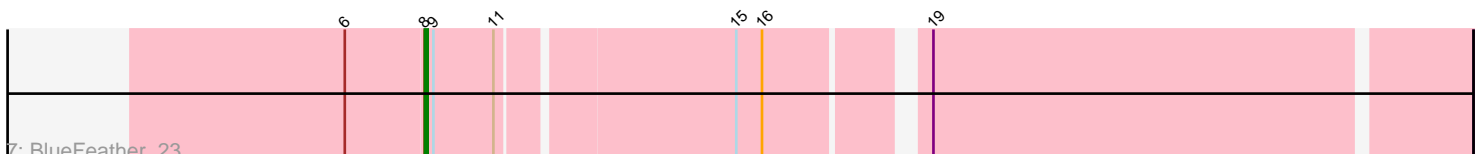
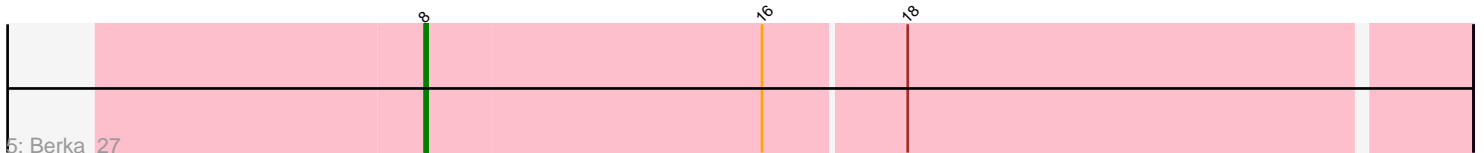
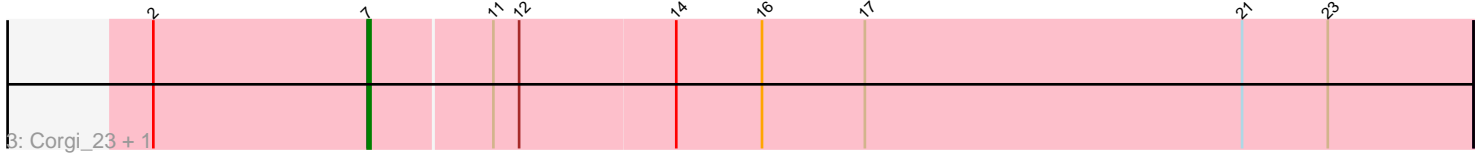
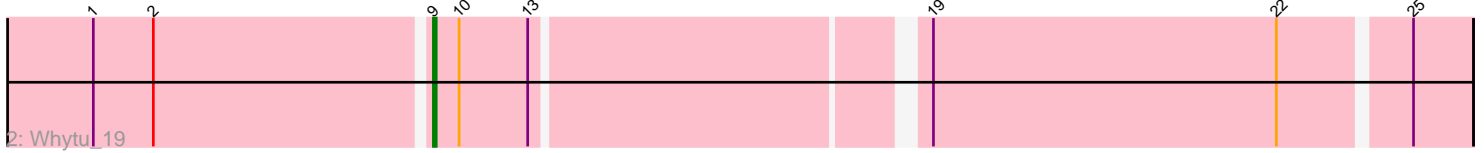
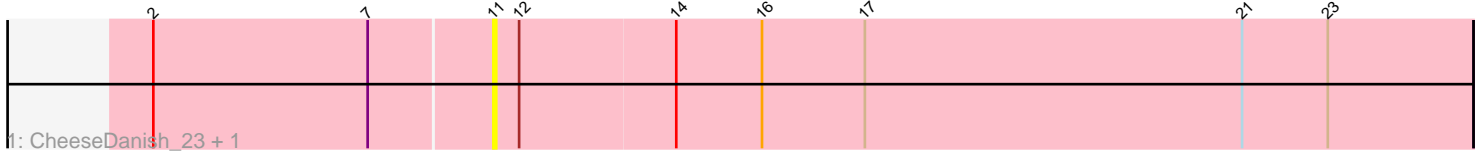


Pham 221970



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 221970 Report

This analysis was run 03/28/25 on database version 593.

Pham number 221970 has 12 members, 3 are drafts.

Phages represented in each track:

- Track 1 : CheeseDanish\_23, KNG13\_23
- Track 2 : Whytu\_19
- Track 3 : Corgi\_23, CabbageMan\_21
- Track 4 : Yavru\_20
- Track 5 : Berka\_27
- Track 6 : Utopia\_20
- Track 7 : BlueFeather\_23
- Track 8 : Noely\_21
- Track 9 : Idaho\_21
- Track 10 : Piku\_19

### ***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 3 of the 9 non-draft genes in the pham.

Genes that call this "Most Annotated" start:

- Piku\_19, Utopia\_20, Whytu\_19, Yavru\_20,

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

- BlueFeather\_23,

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

- Berka\_27, CabbageMan\_21, CheeseDanish\_23, Corgi\_23, Idaho\_21, KNG13\_23, Noely\_21,

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

Start 3:

- Found in 1 of 12 ( 8.3% ) of genes in pham
- Manual Annotations of this start: 1 of 9
- Called 100.0% of time when present
- Phage (with cluster) where this start called: Idaho\_21 (FE),

Start 7:

- Found in 6 of 12 ( 50.0% ) of genes in pham
- Manual Annotations of this start: 3 of 9
- Called 50.0% of time when present
- Phage (with cluster) where this start called: CabbageMan\_21 (FE), Corgi\_23 (FE), Noely\_21 (FE),

Start 8:

- Found in 2 of 12 ( 16.7% ) of genes in pham
- Manual Annotations of this start: 2 of 9
- Called 100.0% of time when present
- Phage (with cluster) where this start called: Berka\_27 (FE), BlueFeather\_23 (FE),

Start 9:

- Found in 5 of 12 ( 41.7% ) of genes in pham
- Manual Annotations of this start: 3 of 9
- Called 80.0% of time when present
- Phage (with cluster) where this start called: Piku\_19 (FE), Utopia\_20 (FE), Whytu\_19 (FE), Yavru\_20 (FE),

Start 11:

- Found in 7 of 12 ( 58.3% ) of genes in pham
- No Manual Annotations of this start.
- Called 28.6% of time when present
- Phage (with cluster) where this start called: CheeseDanish\_23 (FE), KNG13\_23 (FE),

**Summary by clusters:**

There is one cluster represented in this pham: FE

Info for manual annotations of cluster FE:

- Start number 3 was manually annotated 1 time for cluster FE.
- Start number 7 was manually annotated 3 times for cluster FE.
- Start number 8 was manually annotated 2 times for cluster FE.
- Start number 9 was manually annotated 3 times for cluster FE.

**Gene Information:**

Gene: Berka\_27 Start: 14718, Stop: 15074, Start Num: 8

Candidate Starts for Berka\_27:

(Start: 8 @14718 has 2 MA's), (16, 14835), (18, 14883),

Gene: BlueFeather\_23 Start: 15433, Stop: 15774, Start Num: 8

Candidate Starts for BlueFeather\_23:

(6, 15406), (Start: 8 @15433 has 2 MA's), (Start: 9 @15436 has 3 MA's), (11, 15457), (15, 15535), (16, 15544), (19, 15592),

Gene: CabbageMan\_21 Start: 14675, Stop: 15058, Start Num: 7

Candidate Starts for CabbageMan\_21:

(2, 14600), (Start: 7 @14675 has 3 MA's), (11, 14717), (12, 14726), (14, 14780), (16, 14810), (17, 14846), (21, 14978), (23, 15008),

Gene: CheeseDanish\_23 Start: 14876, Stop: 15217, Start Num: 11

Candidate Starts for CheeseDanish\_23:

(2, 14759), (Start: 7 @14834 has 3 MA's), (11, 14876), (12, 14885), (14, 14939), (16, 14969), (17, 15005), (21, 15137), (23, 15167),

Gene: Corgi\_23 Start: 14834, Stop: 15217, Start Num: 7

Candidate Starts for Corgi\_23:

(2, 14759), (Start: 7 @14834 has 3 MA's), (11, 14876), (12, 14885), (14, 14939), (16, 14969), (17, 15005), (21, 15137), (23, 15167),

Gene: Idaho\_21 Start: 14970, Stop: 15386, Start Num: 3

Candidate Starts for Idaho\_21:

(Start: 3 @14970 has 1 MA's), (4, 14985), (5, 14991), (Start: 7 @15006 has 3 MA's), (11, 15045), (16, 15138), (23, 15336), (24, 15342), (26, 15372),

Gene: KNG13\_23 Start: 14873, Stop: 15214, Start Num: 11

Candidate Starts for KNG13\_23:

(2, 14756), (Start: 7 @14831 has 3 MA's), (11, 14873), (12, 14882), (14, 14936), (16, 14966), (17, 15002), (21, 15134), (23, 15164),

Gene: Noely\_21 Start: 14189, Stop: 14569, Start Num: 7

Candidate Starts for Noely\_21:

(Start: 7 @14189 has 3 MA's), (11, 14231), (16, 14321), (20, 14435), (21, 14489), (26, 14555),

Gene: Piku\_19 Start: 14055, Stop: 14396, Start Num: 9

Candidate Starts for Piku\_19:

(Start: 9 @14055 has 3 MA's), (10, 14064), (13, 14088), (19, 14214), (22, 14334), (26, 14382),

Gene: Utopia\_20 Start: 14050, Stop: 14391, Start Num: 9

Candidate Starts for Utopia\_20:

(Start: 9 @14050 has 3 MA's), (10, 14059), (19, 14209), (22, 14329), (25, 14371),

Gene: Whytu\_19 Start: 14130, Stop: 14471, Start Num: 9

Candidate Starts for Whytu\_19:

(1, 14016), (2, 14037), (Start: 9 @14130 has 3 MA's), (10, 14139), (13, 14163), (19, 14289), (22, 14409), (25, 14451),

Gene: Yavru\_20 Start: 14060, Stop: 14401, Start Num: 9

Candidate Starts for Yavru\_20:

(Start: 9 @14060 has 3 MA's), (10, 14069), (19, 14219), (22, 14339),