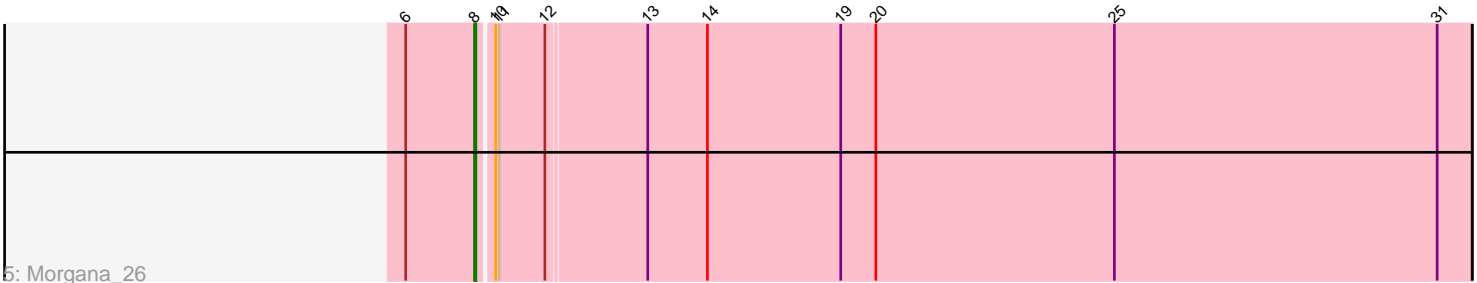
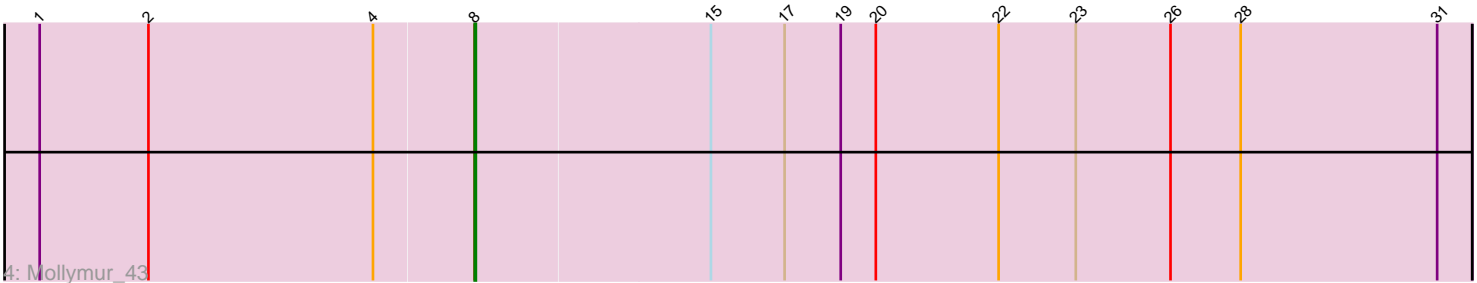
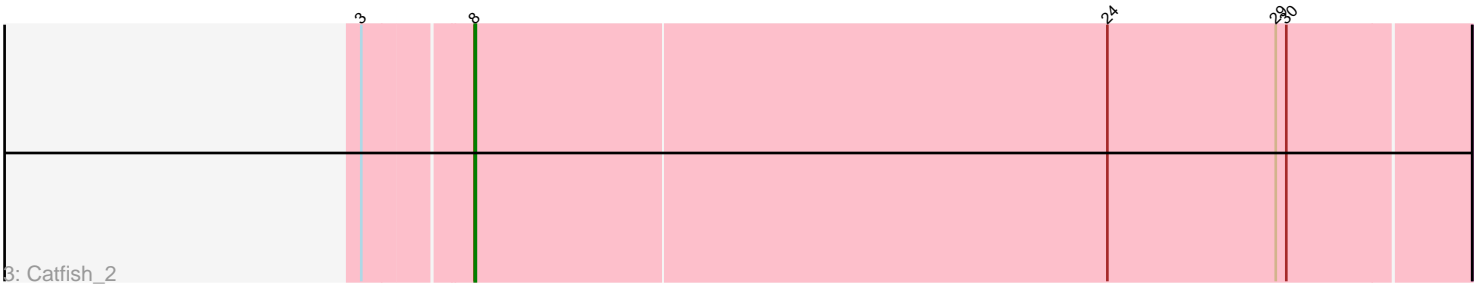
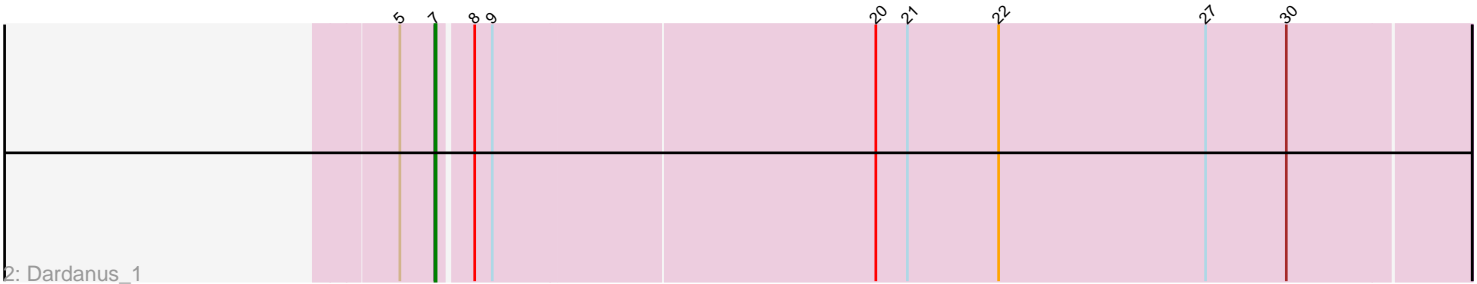
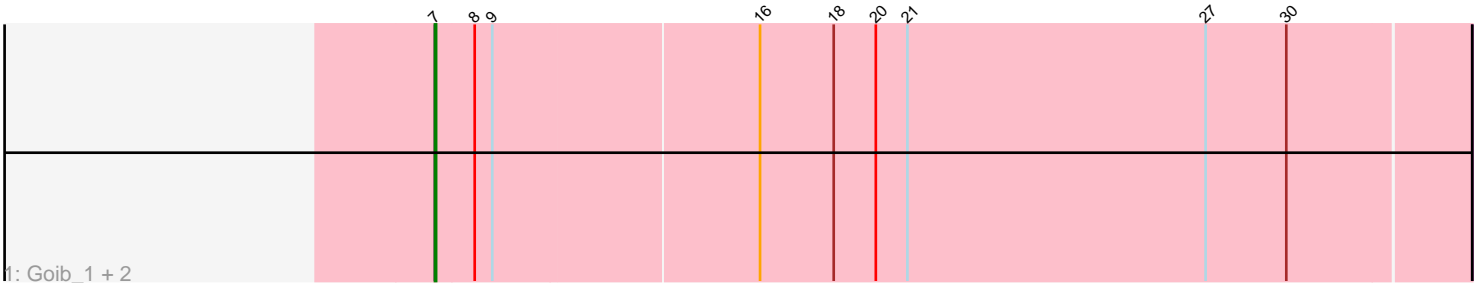


Pham 28256



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 28256 Report

This analysis was run 04/05/24 on database version 557.

Pham number 28256 has 7 members, 0 are drafts.

Phages represented in each track:

- Track 1 : Goib\_1, Vendetta\_1, Splinter\_1
- Track 2 : Dardanus\_1
- Track 3 : Catfish\_2
- Track 4 : Mollymur\_43
- Track 5 : Morgana\_26

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

The start number called the most often in the published annotations is 7, it was called in 4 of the 7 non-draft genes in the pham.

Genes that call this "Most Annotated" start:

- Dardanus\_1, Goib\_1, Splinter\_1, Vendetta\_1,

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

- 

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

- Catfish\_2, Mollymur\_43, Morgana\_26,

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

Start 7:

- Found in 4 of 7 ( 57.1% ) of genes in pham
- Manual Annotations of this start: 4 of 7
- Called 100.0% of time when present
- Phage (with cluster) where this start called: Dardanus\_1 (CU3), Goib\_1 (CU1), Splinter\_1 (CU1), Vendetta\_1 (CU1),

Start 8:

- Found in 7 of 7 ( 100.0% ) of genes in pham
- Manual Annotations of this start: 3 of 7
- Called 42.9% of time when present

- Phage (with cluster) where this start called: Catfish\_2 (CU5), Mollymur\_43 (DL), Morgana\_26 (DZ),

### **Summary by clusters:**

There are 5 clusters represented in this pham: DZ, DL, CU3, CU5, CU1,

Info for manual annotations of cluster CU1:

- Start number 7 was manually annotated 3 times for cluster CU1.

Info for manual annotations of cluster CU3:

- Start number 7 was manually annotated 1 time for cluster CU3.

Info for manual annotations of cluster CU5:

- Start number 8 was manually annotated 1 time for cluster CU5.

Info for manual annotations of cluster DL:

- Start number 8 was manually annotated 1 time for cluster DL.

Info for manual annotations of cluster DZ:

- Start number 8 was manually annotated 1 time for cluster DZ.

### **Gene Information:**

Gene: Catfish\_2 Start: 1219, Stop: 377, Start Num: 8

Candidate Starts for Catfish\_2:

(3, 1306), (Start: 8 @1219 has 3 MA's), (24, 682), (29, 538), (30, 529),

Gene: Dardanus\_1 Start: 1002, Stop: 136, Start Num: 7

Candidate Starts for Dardanus\_1:

(5, 1032), (Start: 7 @1002 has 4 MA's), (Start: 8 @975 has 3 MA's), (9, 960), (20, 639), (21, 612), (22, 534), (27, 357), (30, 288),

Gene: Goib\_1 Start: 1007, Stop: 135, Start Num: 7

Candidate Starts for Goib\_1:

(Start: 7 @1007 has 4 MA's), (Start: 8 @974 has 3 MA's), (9, 959), (16, 737), (18, 674), (20, 638), (21, 611), (27, 356), (30, 287),

Gene: Mollymur\_43 Start: 40248, Stop: 39400, Start Num: 8

Candidate Starts for Mollymur\_43:

(1, 40617), (2, 40524), (4, 40332), (Start: 8 @40248 has 3 MA's), (15, 40050), (17, 39987), (19, 39939), (20, 39909), (22, 39804), (23, 39738), (26, 39657), (28, 39597), (31, 39429),

Gene: Morgana\_26 Start: 24386, Stop: 23547, Start Num: 8

Candidate Starts for Morgana\_26:

(6, 24443), (Start: 8 @24386 has 3 MA's), (10, 24374), (11, 24371), (12, 24332), (13, 24251), (14, 24200), (19, 24086), (20, 24056), (25, 23852), (31, 23576),

Gene: Splinter\_1 Start: 1007, Stop: 135, Start Num: 7

Candidate Starts for Splinter\_1:

(Start: 7 @1007 has 4 MA's), (Start: 8 @974 has 3 MA's), (9, 959), (16, 737), (18, 674), (20, 638), (21, 611), (27, 356), (30, 287),

Gene: Vendetta\_1 Start: 1007, Stop: 135, Start Num: 7

Candidate Starts for Vendetta\_1:

(Start: 7 @1007 has 4 MA's), (Start: 8 @974 has 3 MA's), (9, 959), (16, 737), (18, 674), (20, 638), (21, 611), (27, 356), (30, 287),