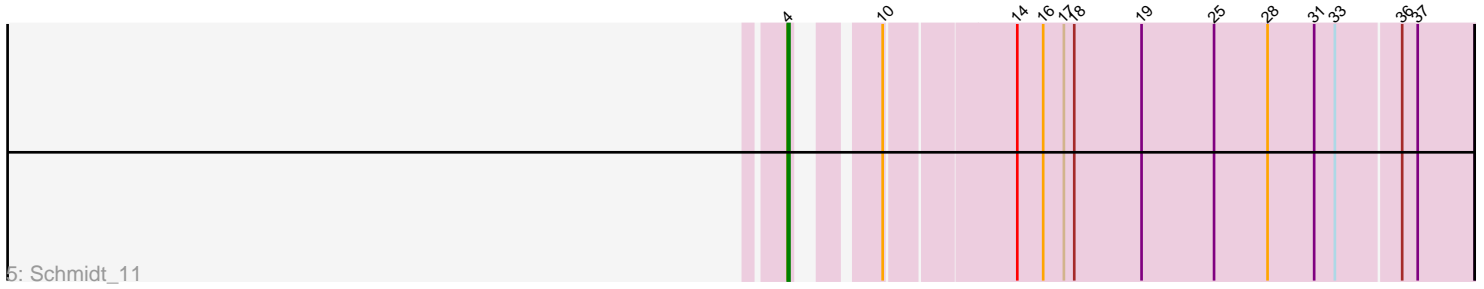
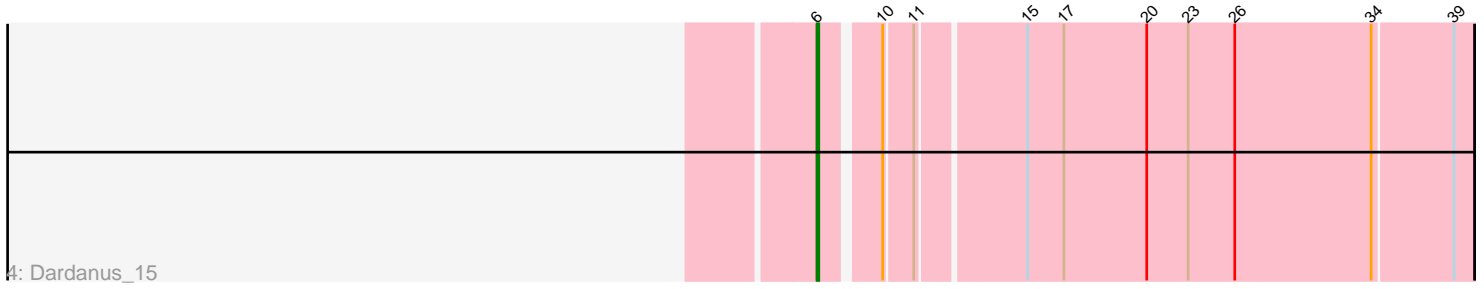
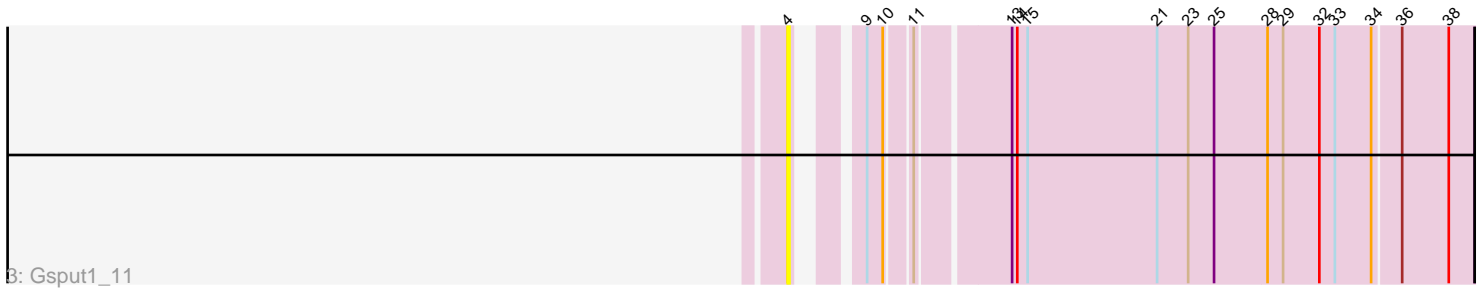
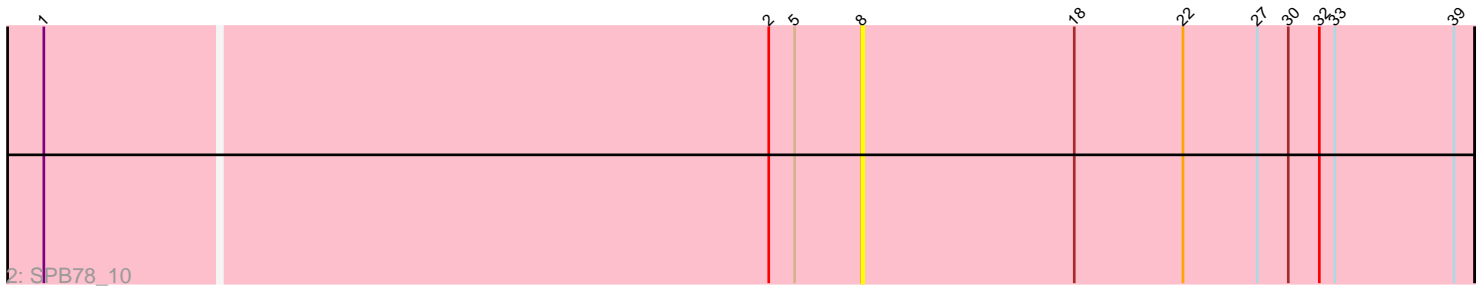
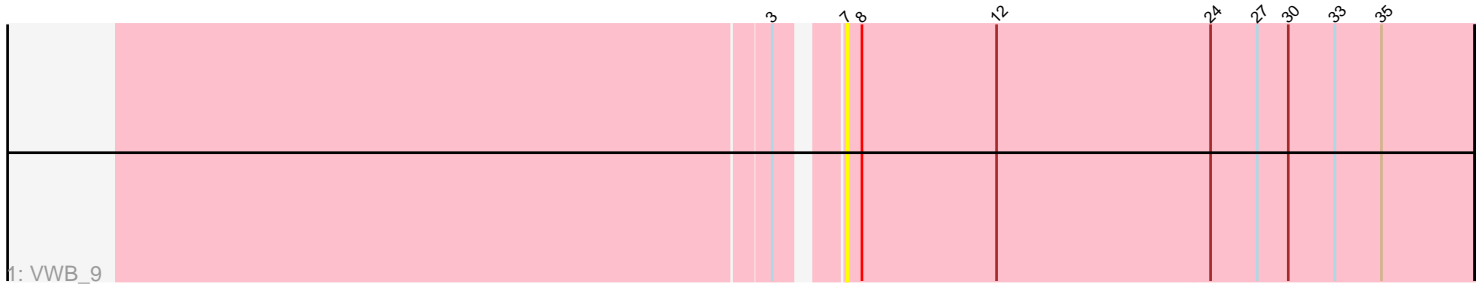


Pham 203728



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 203728 Report

This analysis was run 01/18/25 on database version 583.

WARNING: Pham size does not match number of genes in report. Either unphamerated genes have been added (by you) or starterator has removed genes due to invalid start codon.

Pham number 203728 has 5 members, 3 are drafts.

Phages represented in each track:

- Track 1 : VWB_9
- Track 2 : SPB78_10
- Track 3 : Gsput1_11
- Track 4 : Dardanus_15
- Track 5 : Schmidt_11

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

The start number called the most often in the published annotations is 4, it was called in 1 of the 2 non-draft genes in the pham.

Genes that call this "Most Annotated" start:

- Gsput1_11, Schmidt_11,

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

-

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

- Dardanus_15, SPB78_10, VWB_9,

Summary by start number:

Start 4:

- Found in 2 of 5 (40.0%) of genes in pham
- Manual Annotations of this start: 1 of 2
- Called 100.0% of time when present
- Phage (with cluster) where this start called: Gsput1_11 (CU2), Schmidt_11 (CU4),

Start 6:

- Found in 1 of 5 (20.0%) of genes in pham

- Manual Annotations of this start: 1 of 2
- Called 100.0% of time when present
- Phage (with cluster) where this start called: Dardanus_15 (CU3),

Start 7:

- Found in 1 of 5 (20.0%) of genes in pham
- No Manual Annotations of this start.
- Called 100.0% of time when present
- Phage (with cluster) where this start called: VWB_9 (BA),

Start 8:

- Found in 2 of 5 (40.0%) of genes in pham
- No Manual Annotations of this start.
- Called 50.0% of time when present
- Phage (with cluster) where this start called: SPB78_10 (BA),

Summary by clusters:

There are 4 clusters represented in this pham: CU4, CU3, BA, CU2,

Info for manual annotations of cluster CU3:

- Start number 6 was manually annotated 1 time for cluster CU3.

Info for manual annotations of cluster CU4:

- Start number 4 was manually annotated 1 time for cluster CU4.

Gene Information:

Gene: Dardanus_15 Start: 9541, Stop: 9897, Start Num: 6

Candidate Starts for Dardanus_15:

(Start: 6 @9541 has 1 MA's), (10, 9571), (11, 9586), (15, 9643), (17, 9664), (20, 9712), (23, 9736), (26, 9763), (34, 9841), (39, 9886),

Gene: Gspu1_11 Start: 7862, Stop: 8218, Start Num: 4

Candidate Starts for Gspu1_11:

(Start: 4 @7862 has 1 MA's), (9, 7886), (10, 7895), (11, 7907), (13, 7955), (14, 7958), (15, 7964), (21, 8039), (23, 8057), (25, 8072), (28, 8102), (29, 8111), (32, 8132), (33, 8141), (34, 8162), (36, 8177), (38, 8204),

Gene: SPB78_10 Start: 7392, Stop: 7745, Start Num: 8

Candidate Starts for SPB78_10:

(1, 6924), (2, 7338), (5, 7353), (8, 7392), (18, 7515), (22, 7578), (27, 7620), (30, 7638), (32, 7656), (33, 7665), (39, 7734),

Gene: Schmidt_11 Start: 7862, Stop: 8224, Start Num: 4

Candidate Starts for Schmidt_11:

(Start: 4 @7862 has 1 MA's), (10, 7895), (14, 7964), (16, 7979), (17, 7991), (18, 7997), (19, 8036), (25, 8078), (28, 8108), (31, 8135), (33, 8147), (36, 8183), (37, 8192),

Gene: VWB_9 Start: 7437, Stop: 7799, Start Num: 7

Candidate Starts for VWB_9:

(3, 7410), (7, 7437), (8, 7446), (12, 7524), (24, 7647), (27, 7674), (30, 7692), (33, 7719), (35, 7746),