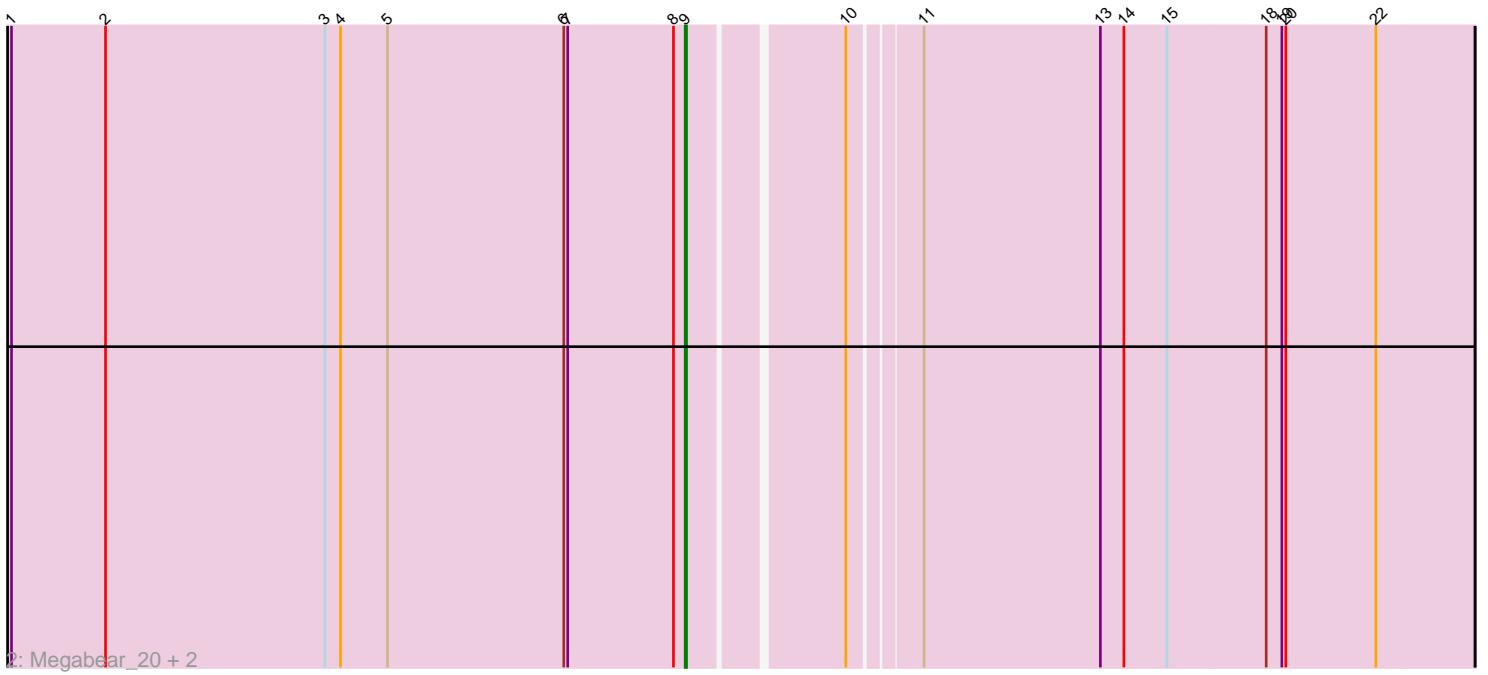
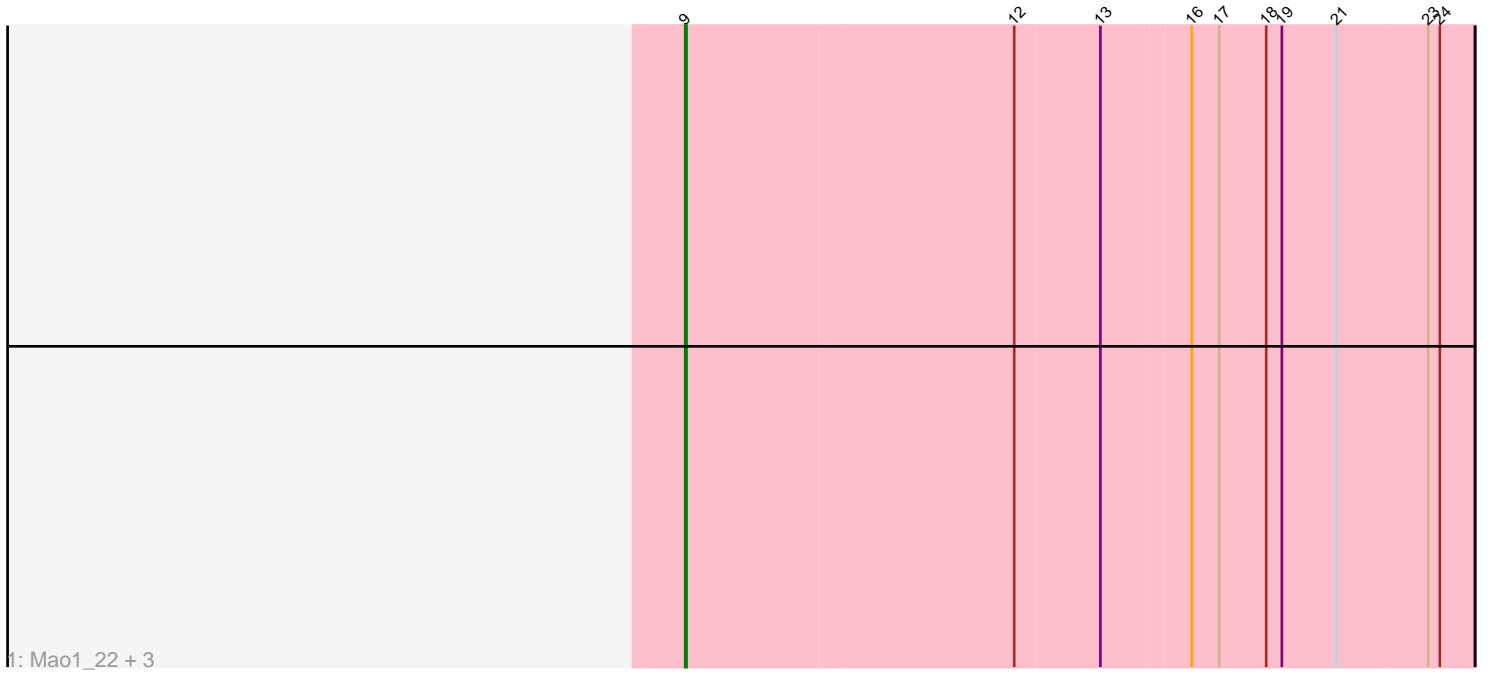


Pham 225209



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 225209 Report

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

Pham number 225209 has 7 members, 0 are drafts.

Phages represented in each track:

- Track 1 : Mao1_22, Sejanus_23, Dori_23, Mask_23
- Track 2 : Megabear_20, Cepens_20, Taptic_20

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 7 of the 7 non-draft genes in the pham.

Genes that call this "Most Annotated" start:

- Cepens_20, Dori_23, Mao1_22, Mask_23, Megabear_20, Sejanus_23, Taptic_20,

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

-

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

-

Summary by start number:

Start 9:

- Found in 7 of 7 (100.0%) of genes in pham
- Manual Annotations of this start: 7 of 7
- Called 100.0% of time when present
- Phage (with cluster) where this start called: Cepens_20 (W), Dori_23 (AD), Mao1_22 (AD), Mask_23 (AD), Megabear_20 (W), Sejanus_23 (AD), Taptic_20 (W),

Summary by clusters:

There are 2 clusters represented in this pham: AD, W,

Info for manual annotations of cluster AD:

- Start number 9 was manually annotated 4 times for cluster AD.

Info for manual annotations of cluster W:

- Start number 9 was manually annotated 3 times for cluster W.

Gene Information:

Gene: Cepens_20 Start: 18624, Stop: 19199, Start Num: 9

Candidate Starts for Cepens_20:

(1, 18108), (2, 18180), (3, 18348), (4, 18360), (5, 18396), (6, 18531), (7, 18534), (8, 18615), (Start: 9 @18624 has 7 MA's), (10, 18732), (11, 18780), (13, 18915), (14, 18933), (15, 18966), (18, 19041), (19, 19053), (20, 19056), (22, 19125),

Gene: Dori_23 Start: 16796, Stop: 17389, Start Num: 9

Candidate Starts for Dori_23:

(Start: 9 @16796 has 7 MA's), (12, 17045), (13, 17108), (16, 17174), (17, 17195), (18, 17231), (19, 17243), (21, 17285), (23, 17354), (24, 17363),

Gene: Mao1_22 Start: 17208, Stop: 17801, Start Num: 9

Candidate Starts for Mao1_22:

(Start: 9 @17208 has 7 MA's), (12, 17457), (13, 17520), (16, 17586), (17, 17607), (18, 17643), (19, 17655), (21, 17697), (23, 17766), (24, 17775),

Gene: Mask_23 Start: 17201, Stop: 17794, Start Num: 9

Candidate Starts for Mask_23:

(Start: 9 @17201 has 7 MA's), (12, 17450), (13, 17513), (16, 17579), (17, 17600), (18, 17636), (19, 17648), (21, 17690), (23, 17759), (24, 17768),

Gene: Megabear_20 Start: 18567, Stop: 19142, Start Num: 9

Candidate Starts for Megabear_20:

(1, 18051), (2, 18123), (3, 18291), (4, 18303), (5, 18339), (6, 18474), (7, 18477), (8, 18558), (Start: 9 @18567 has 7 MA's), (10, 18675), (11, 18723), (13, 18858), (14, 18876), (15, 18909), (18, 18984), (19, 18996), (20, 18999), (22, 19068),

Gene: Sejanus_23 Start: 18142, Stop: 18735, Start Num: 9

Candidate Starts for Sejanus_23:

(Start: 9 @18142 has 7 MA's), (12, 18391), (13, 18454), (16, 18520), (17, 18541), (18, 18577), (19, 18589), (21, 18631), (23, 18700), (24, 18709),

Gene: Taptic_20 Start: 18567, Stop: 19142, Start Num: 9

Candidate Starts for Taptic_20:

(1, 18051), (2, 18123), (3, 18291), (4, 18303), (5, 18339), (6, 18474), (7, 18477), (8, 18558), (Start: 9 @18567 has 7 MA's), (10, 18675), (11, 18723), (13, 18858), (14, 18876), (15, 18909), (18, 18984), (19, 18996), (20, 18999), (22, 19068),