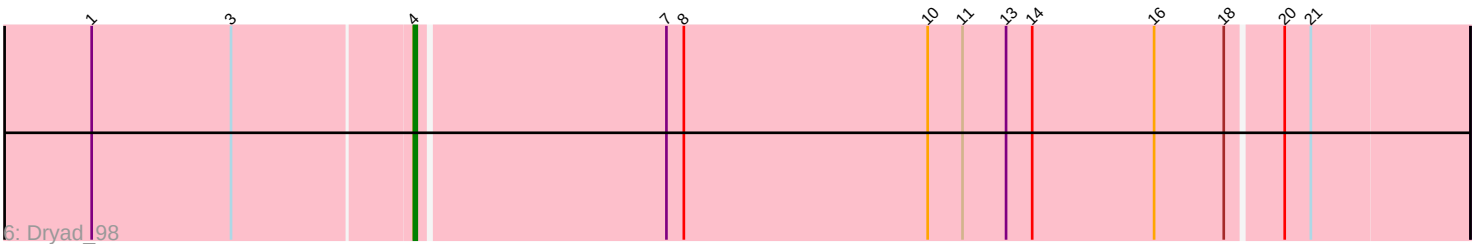
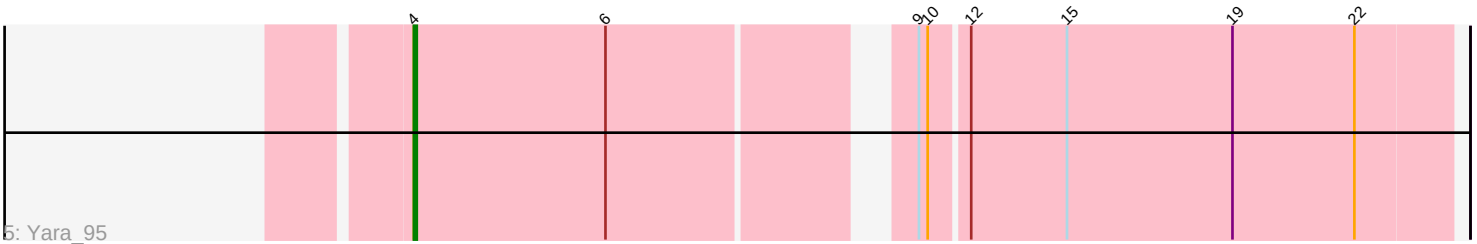
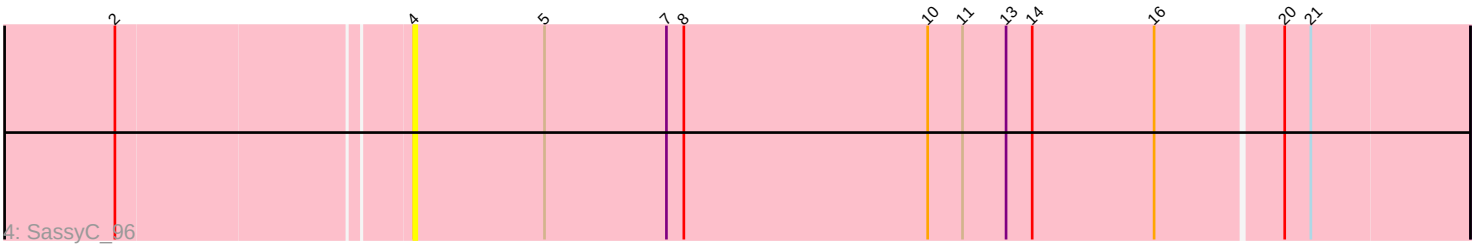
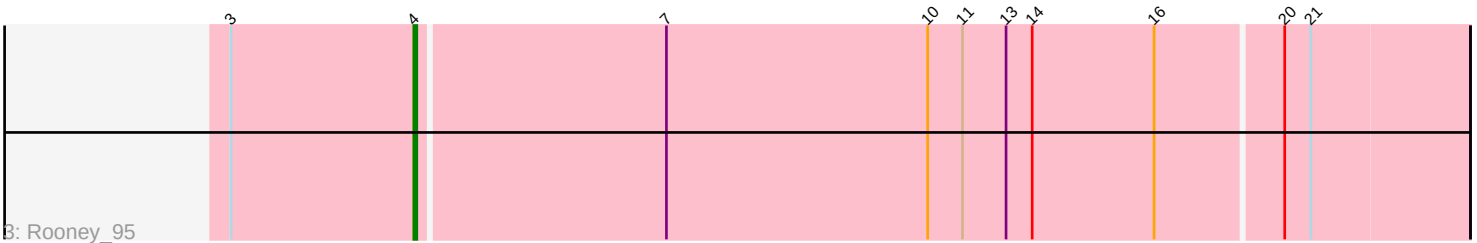
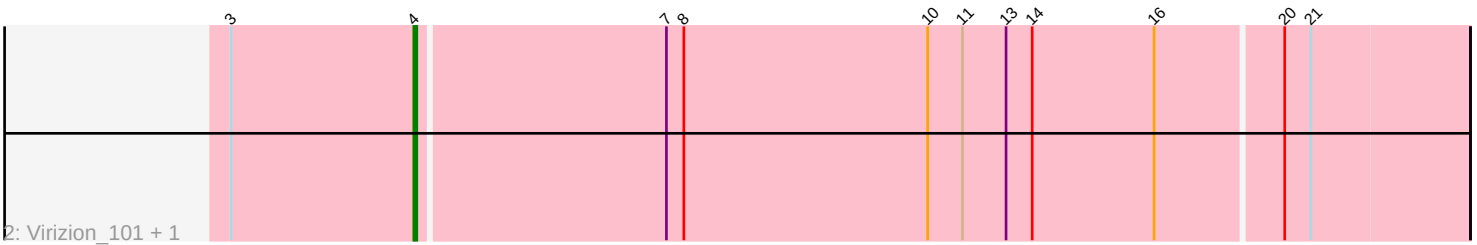
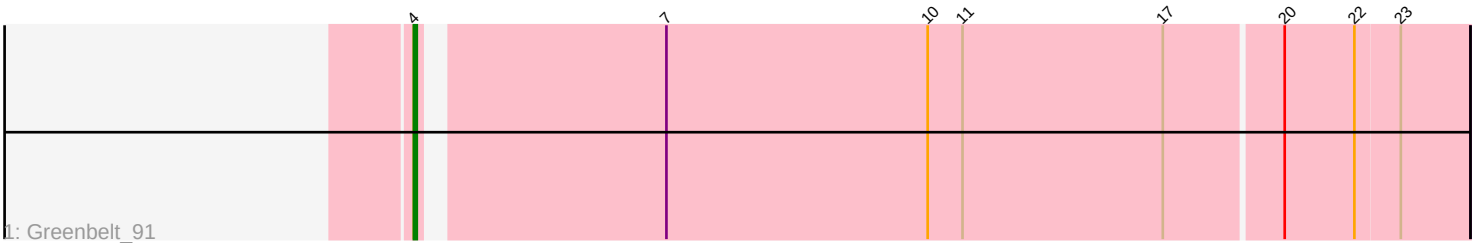


Pham 309979



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 309979 Report

This analysis was run 09/05/26 on database version 662.

Pham number 309979 has 7 members, 2 are drafts.

Phages represented in each track:

- Track 1 : Greenbelt_91
- Track 2 : Virizion_101, Gibson_95
- Track 3 : Rooney_95
- Track 4 : SassyC_96
- Track 5 : Yara_95
- Track 6 : Dryad_98

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 5 of the 5 non-draft genes in the pham.

Genes that call this "Most Annotated" start:

- Dryad_98, Gibson_95, Greenbelt_91, Rooney_95, SassyC_96, Virizion_101, Yara_95,

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 4:

- Found in 7 of 7 (100.0%) of genes in pham
- Manual Annotations of this start: 5 of 5
- Called 100.0% of time when present
- Phage (with cluster) where this start called: Dryad_98 (BN), Gibson_95 (BN), Greenbelt_91 (BN), Rooney_95 (BN), SassyC_96 (BN), Virizion_101 (BN), Yara_95 (BN),

Summary by clusters:

There is one cluster represented in this pham: BN

Info for manual annotations of cluster BN:

•Start number 4 was manually annotated 5 times for cluster BN.

Gene Information:

Gene: Dryad_98 Start: 61804, Stop: 61448, Start Num: 4

Candidate Starts for Dryad_98:

(1, 61912), (3, 61864), (Start: 4 @61804 has 5 MA's), (7, 61720), (8, 61714), (10, 61630), (11, 61618), (13, 61603), (14, 61594), (16, 61552), (18, 61528), (20, 61510), (21, 61501),

Gene: Gibson_95 Start: 61610, Stop: 61254, Start Num: 4

Candidate Starts for Gibson_95:

(3, 61673), (Start: 4 @61610 has 5 MA's), (7, 61526), (8, 61520), (10, 61436), (11, 61424), (13, 61409), (14, 61400), (16, 61358), (20, 61316), (21, 61307),

Gene: Greenbelt_91 Start: 57682, Stop: 57332, Start Num: 4

Candidate Starts for Greenbelt_91:

(Start: 4 @57682 has 5 MA's), (7, 57604), (10, 57514), (11, 57502), (17, 57433), (20, 57394), (22, 57370), (23, 57355),

Gene: Rooney_95 Start: 61624, Stop: 61268, Start Num: 4

Candidate Starts for Rooney_95:

(3, 61687), (Start: 4 @61624 has 5 MA's), (7, 61540), (10, 61450), (11, 61438), (13, 61423), (14, 61414), (16, 61372), (20, 61330), (21, 61321),

Gene: SassyC_96 Start: 60685, Stop: 60326, Start Num: 4

Candidate Starts for SassyC_96:

(2, 60781), (Start: 4 @60685 has 5 MA's), (5, 60640), (7, 60598), (8, 60592), (10, 60508), (11, 60496), (13, 60481), (14, 60472), (16, 60430), (20, 60388), (21, 60379),

Gene: Virizion_101 Start: 61824, Stop: 61468, Start Num: 4

Candidate Starts for Virizion_101:

(3, 61887), (Start: 4 @61824 has 5 MA's), (7, 61740), (8, 61734), (10, 61650), (11, 61638), (13, 61623), (14, 61614), (16, 61572), (20, 61530), (21, 61521),

Gene: Yara_95 Start: 61554, Stop: 61219, Start Num: 4

Candidate Starts for Yara_95:

(Start: 4 @61554 has 5 MA's), (6, 61488), (9, 61398), (10, 61395), (12, 61383), (15, 61350), (19, 61293), (22, 61251),