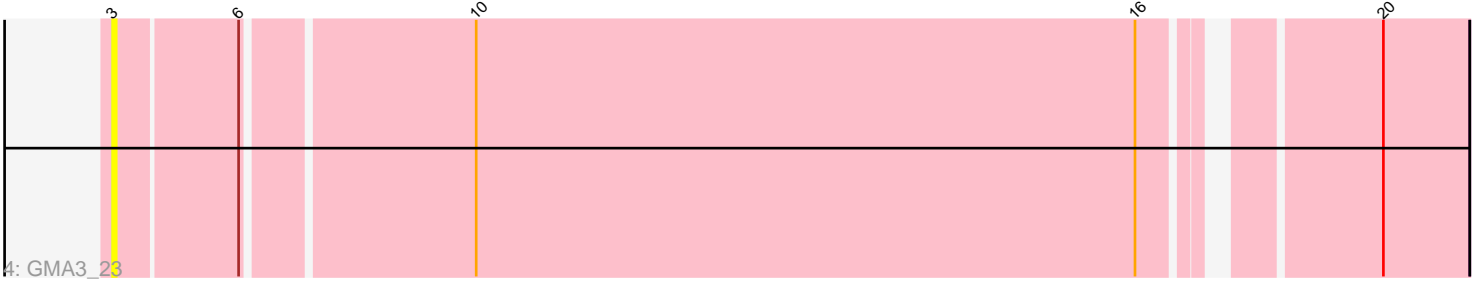
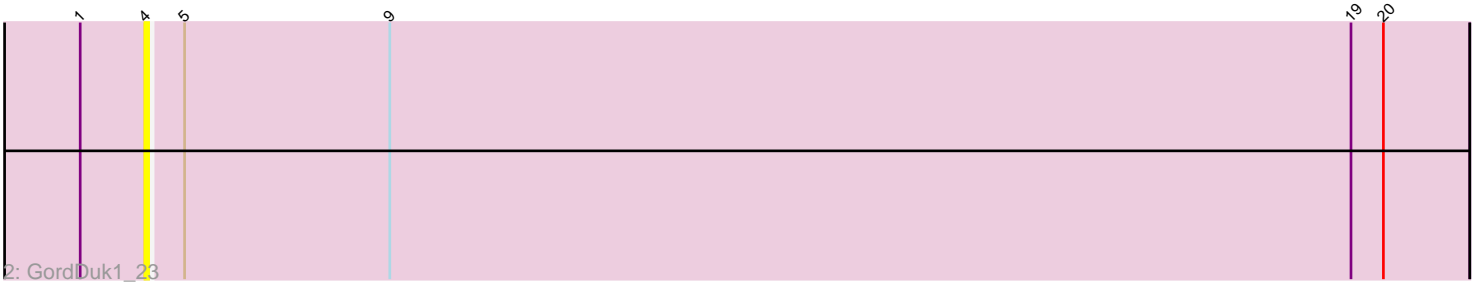
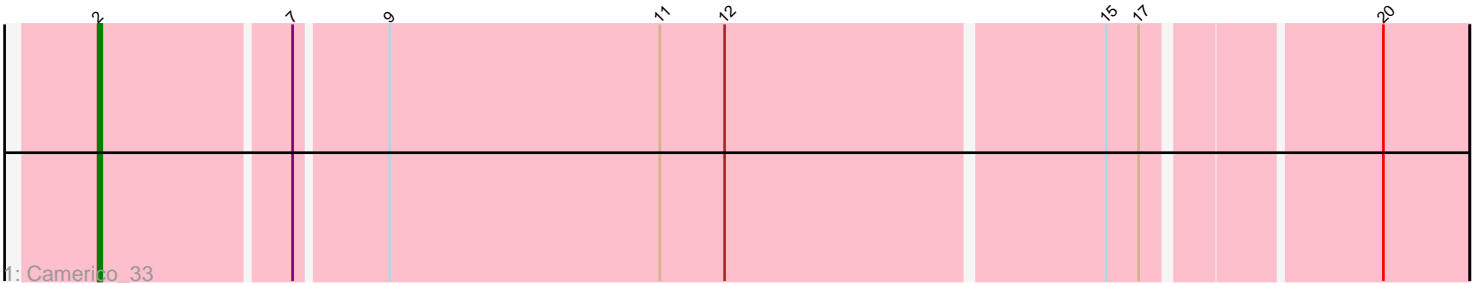


Pham 7347



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 7347 Report

This analysis was run 04/28/24 on database version 559.

Pham number 7347 has 6 members, 4 are drafts.

Phages represented in each track:

- Track 1 : Camerico_33
- Track 2 : GordDuk1_23
- Track 3 : GordTnk2_23, Gmala1_23
- Track 4 : GMA3_23
- Track 5 : Jumbo_29

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

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

Genes that call this "Most Annotated" start:

- Camerico_33,

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

-

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

- GMA3_23, Gmala1_23, GordDuk1_23, GordTnk2_23, Jumbo_29,

Summary by start number:

Start 2:

- Found in 1 of 6 (16.7%) 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: Camerico_33 (DF),

Start 3:

- Found in 2 of 6 (33.3%) 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: GMA3_23 (DF2), Jumbo_29 (DF3),

Start 4:

- Found in 3 of 6 (50.0%) of genes in pham
- No Manual Annotations of this start.
- Called 100.0% of time when present
- Phage (with cluster) where this start called: Gmala1_23 (DF1), GordDuk1_23 (DF1), GordTnk2_23 (DF1),

Summary by clusters:

There are 4 clusters represented in this pham: DF, DF1, DF3, DF2,

Info for manual annotations of cluster DF:

- Start number 2 was manually annotated 1 time for cluster DF.

Info for manual annotations of cluster DF3:

- Start number 3 was manually annotated 1 time for cluster DF3.

Gene Information:

Gene: Camerico_33 Start: 40298, Stop: 40660, Start Num: 2

Candidate Starts for Camerico_33:

(Start: 2 @40298 has 1 MA's), (7, 40349), (9, 40373), (11, 40448), (12, 40466), (15, 40568), (17, 40577), (20, 40637),

Gene: GMA3_23 Start: 34586, Stop: 34939, Start Num: 3

Candidate Starts for GMA3_23:

(Start: 3 @34586 has 1 MA's), (6, 34619), (10, 34679), (16, 34862), (20, 34916),

Gene: Gmala1_23 Start: 32972, Stop: 33337, Start Num: 4

Candidate Starts for Gmala1_23:

(4, 32972), (5, 32981), (9, 33038), (19, 33305), (20, 33314),

Gene: GordDuk1_23 Start: 32912, Stop: 33277, Start Num: 4

Candidate Starts for GordDuk1_23:

(1, 32894), (4, 32912), (5, 32921), (9, 32978), (19, 33245), (20, 33254),

Gene: GordTnk2_23 Start: 32879, Stop: 33244, Start Num: 4

Candidate Starts for GordTnk2_23:

(4, 32879), (5, 32888), (9, 32945), (19, 33212), (20, 33221),

Gene: Jumbo_29 Start: 38914, Stop: 39270, Start Num: 3

Candidate Starts for Jumbo_29:

(Start: 3 @38914 has 1 MA's), (6, 38947), (8, 38977), (13, 39112), (14, 39142), (18, 39202),