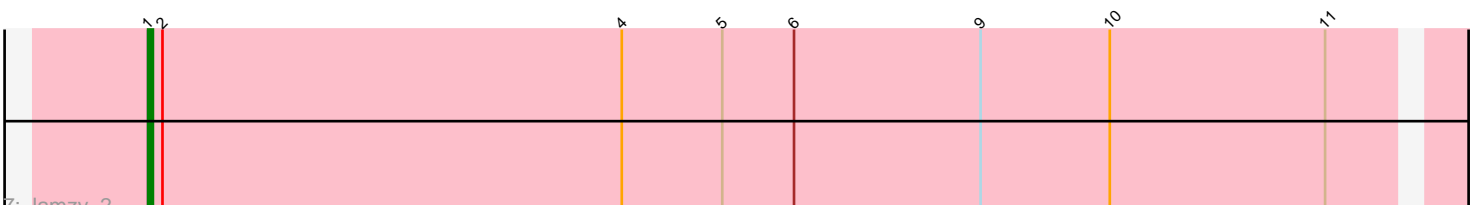
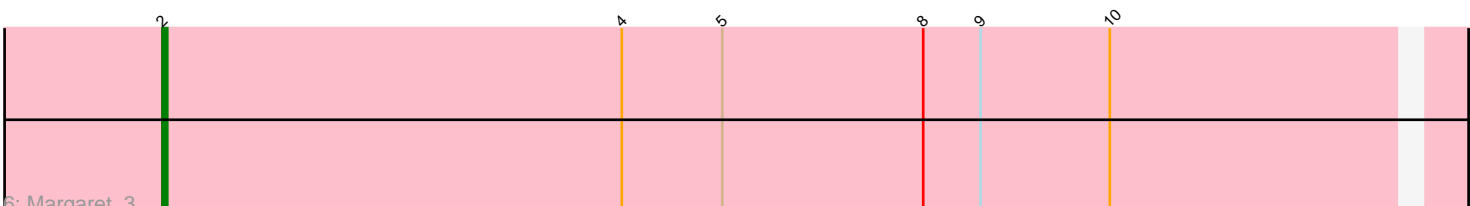
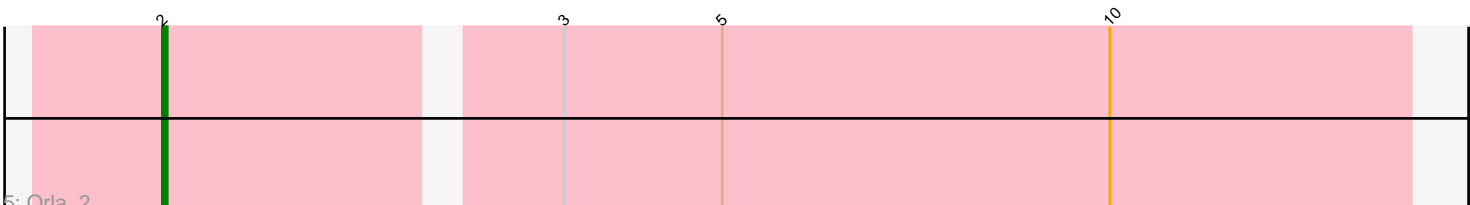
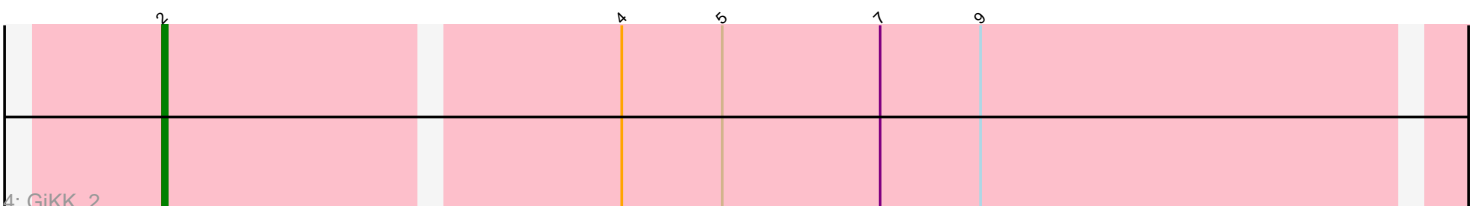
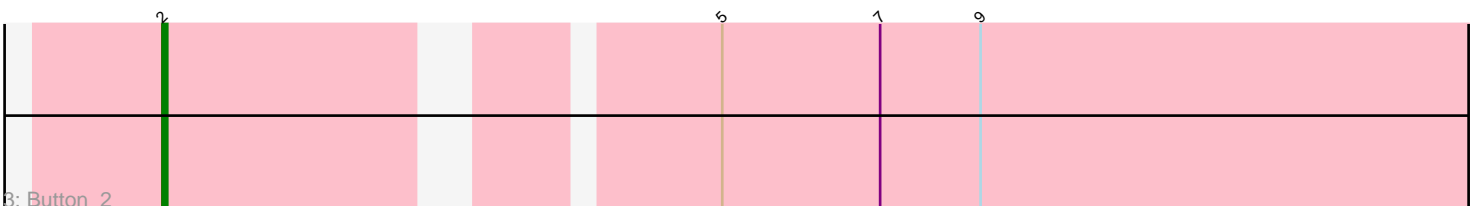
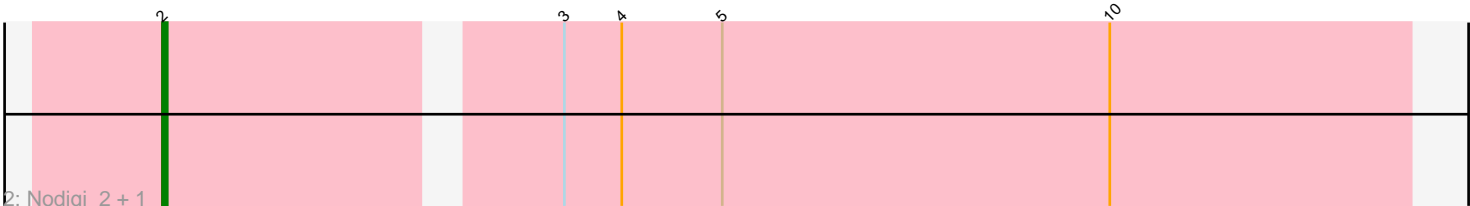
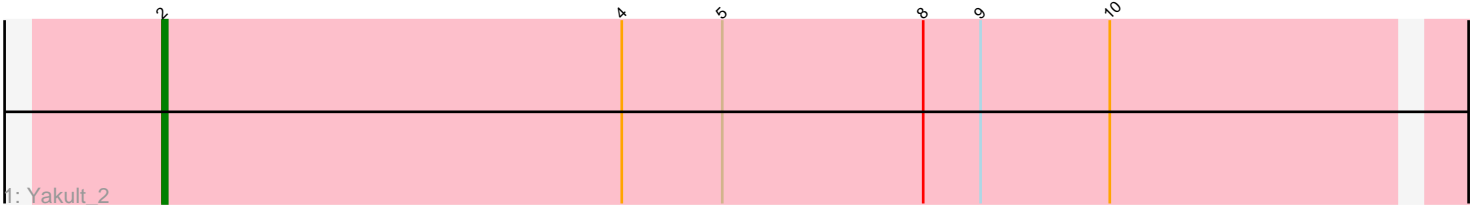


Pham 7785



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 7785 Report

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

Pham number 7785 has 8 members, 0 are drafts.

Phages represented in each track:

- Track 1 : Yakult_2
- Track 2 : Nodigi_2, Hexbug_2
- Track 3 : Button_2
- Track 4 : GiKK_2
- Track 5 : Orla_2
- Track 6 : Margaret_3
- Track 7 : Jamzy_2

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 7 of the 8 non-draft genes in the pham.

Genes that call this "Most Annotated" start:

- Button_2, GiKK_2, Hexbug_2, Margaret_3, Nodigi_2, Orla_2, Yakult_2,

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

- Jamzy_2,

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

•

Summary by start number:

Start 1:

- Found in 1 of 8 (12.5%) of genes in pham
- Manual Annotations of this start: 1 of 8
- Called 100.0% of time when present
- Phage (with cluster) where this start called: Jamzy_2 (CT),

Start 2:

- Found in 8 of 8 (100.0%) of genes in pham
- Manual Annotations of this start: 7 of 8
- Called 87.5% of time when present

- Phage (with cluster) where this start called: Button_2 (CT), GiKK_2 (CT), Hexbug_2 (CT), Margaret_3 (CT), Nodigi_2 (CT), Orla_2 (CT), Yakult_2 (CT),

Summary by clusters:

There is one cluster represented in this pham: CT

Info for manual annotations of cluster CT:

- Start number 1 was manually annotated 1 time for cluster CT.
- Start number 2 was manually annotated 7 times for cluster CT.

Gene Information:

Gene: Button_2 Start: 505, Stop: 762, Start Num: 2

Candidate Starts for Button_2:

(Start: 2 @505 has 7 MA's), (5, 604), (7, 637), (9, 658),

Gene: GiKK_2 Start: 504, Stop: 767, Start Num: 2

Candidate Starts for GiKK_2:

(Start: 2 @504 has 7 MA's), (4, 594), (5, 615), (7, 648), (9, 669),

Gene: Hexbug_2 Start: 505, Stop: 756, Start Num: 2

Candidate Starts for Hexbug_2:

(Start: 2 @505 has 7 MA's), (3, 580), (4, 592), (5, 613), (10, 694),

Gene: Jamzy_2 Start: 501, Stop: 773, Start Num: 1

Candidate Starts for Jamzy_2:

(Start: 1 @501 has 1 MA's), (Start: 2 @504 has 7 MA's), (4, 600), (5, 621), (6, 636), (9, 675), (10, 702), (11, 747),

Gene: Margaret_3 Start: 1128, Stop: 1397, Start Num: 2

Candidate Starts for Margaret_3:

(Start: 2 @1128 has 7 MA's), (4, 1224), (5, 1245), (8, 1287), (9, 1299), (10, 1326),

Gene: Nodigi_2 Start: 505, Stop: 756, Start Num: 2

Candidate Starts for Nodigi_2:

(Start: 2 @505 has 7 MA's), (3, 580), (4, 592), (5, 613), (10, 694),

Gene: Orla_2 Start: 504, Stop: 755, Start Num: 2

Candidate Starts for Orla_2:

(Start: 2 @504 has 7 MA's), (3, 579), (5, 612), (10, 693),

Gene: Yakult_2 Start: 503, Stop: 772, Start Num: 2

Candidate Starts for Yakult_2:

(Start: 2 @503 has 7 MA's), (4, 599), (5, 620), (8, 662), (9, 674), (10, 701),