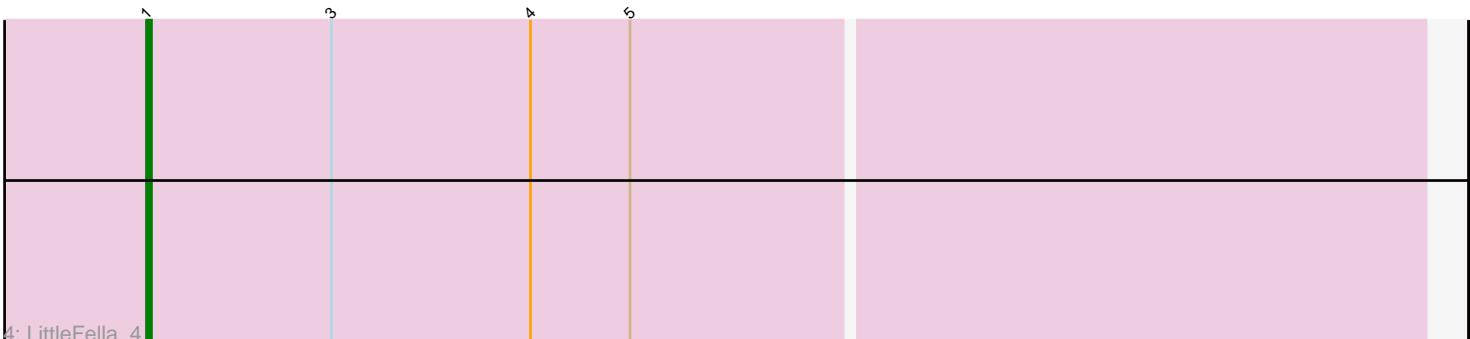
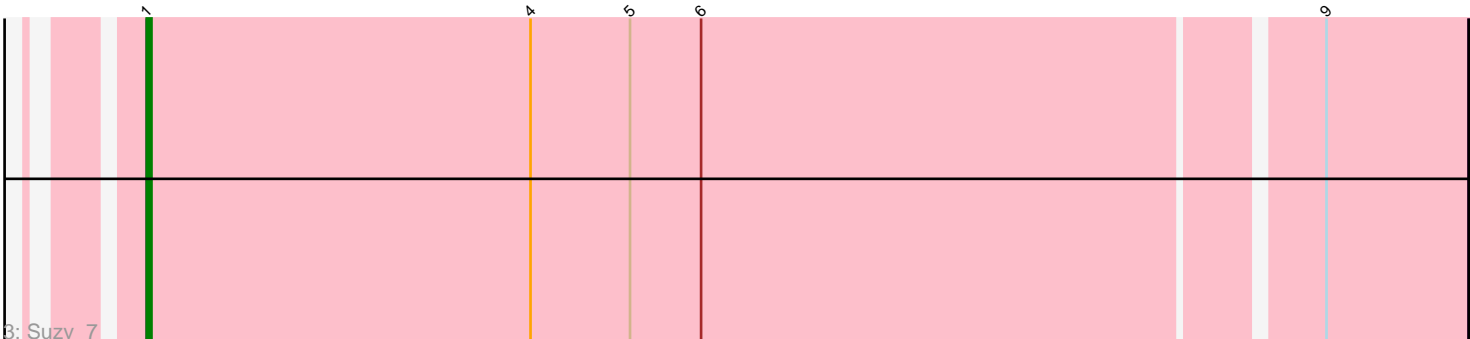
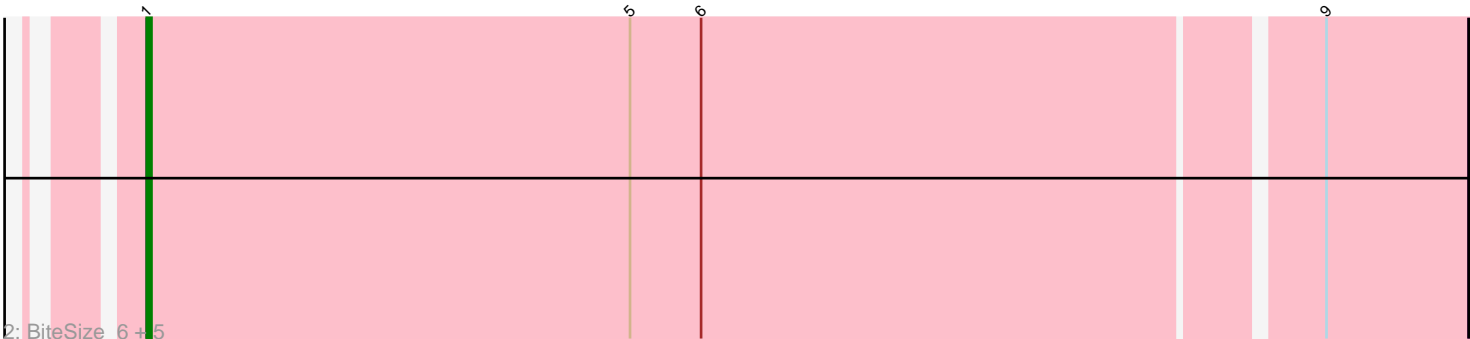
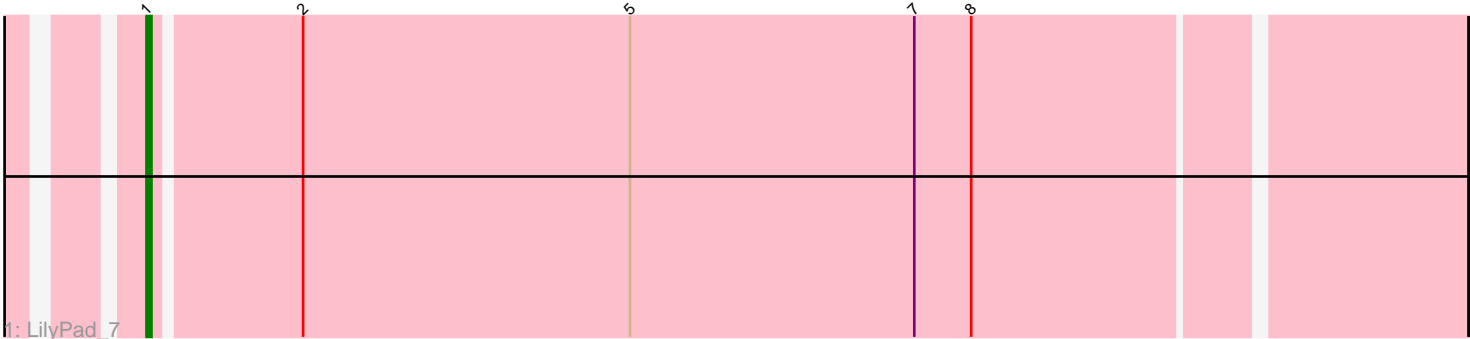


Pham 5873



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 5873 Report

This analysis was run 04/05/24 on database version 557.

Pham number 5873 has 9 members, 0 are drafts.

Phages represented in each track:

- Track 1 : LilyPad_7
- Track 2 : BiteSize_6, Beyoncage_6, Djokovic_6, Sienna_6, Madi_6, Terapin_6
- Track 3 : Suzy_7
- Track 4 : LittleFella_4

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

The start number called the most often in the published annotations is 1, it was called in 9 of the 9 non-draft genes in the pham.

Genes that call this "Most Annotated" start:

- Beyoncage_6, BiteSize_6, Djokovic_6, LilyPad_7, LittleFella_4, Madi_6, Sienna_6, Suzy_7, Terapin_6,

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 1:

- Found in 9 of 9 (100.0%) of genes in pham
- Manual Annotations of this start: 9 of 9
- Called 100.0% of time when present
- Phage (with cluster) where this start called: Beyoncage_6 (DG1), BiteSize_6 (DG1), Djokovic_6 (DG1), LilyPad_7 (DG1), LittleFella_4 (DG2), Madi_6 (DG1), Sienna_6 (DG1), Suzy_7 (DG1), Terapin_6 (DG1),

Summary by clusters:

There are 2 clusters represented in this pham: DG2, DG1,

Info for manual annotations of cluster DG1:

- Start number 1 was manually annotated 8 times for cluster DG1.

Info for manual annotations of cluster DG2:

- Start number 1 was manually annotated 1 time for cluster DG2.

Gene Information:

Gene: Beyoncage_6 Start: 4196, Stop: 4468, Start Num: 1

Candidate Starts for Beyoncage_6:

(Start: 1 @4196 has 9 MA's), (5, 4298), (6, 4313), (9, 4439),

Gene: BiteSize_6 Start: 4196, Stop: 4468, Start Num: 1

Candidate Starts for BiteSize_6:

(Start: 1 @4196 has 9 MA's), (5, 4298), (6, 4313), (9, 4439),

Gene: Djokovic_6 Start: 4196, Stop: 4468, Start Num: 1

Candidate Starts for Djokovic_6:

(Start: 1 @4196 has 9 MA's), (5, 4298), (6, 4313), (9, 4439),

Gene: LilyPad_7 Start: 4957, Stop: 5226, Start Num: 1

Candidate Starts for LilyPad_7:

(Start: 1 @4957 has 9 MA's), (2, 4987), (5, 5056), (7, 5116), (8, 5128),

Gene: LittleFella_4 Start: 4143, Stop: 4409, Start Num: 1

Candidate Starts for LittleFella_4:

(Start: 1 @4143 has 9 MA's), (3, 4182), (4, 4224), (5, 4245),

Gene: Madi_6 Start: 4196, Stop: 4468, Start Num: 1

Candidate Starts for Madi_6:

(Start: 1 @4196 has 9 MA's), (5, 4298), (6, 4313), (9, 4439),

Gene: Sienna_6 Start: 4196, Stop: 4468, Start Num: 1

Candidate Starts for Sienna_6:

(Start: 1 @4196 has 9 MA's), (5, 4298), (6, 4313), (9, 4439),

Gene: Suzy_7 Start: 5468, Stop: 5740, Start Num: 1

Candidate Starts for Suzy_7:

(Start: 1 @5468 has 9 MA's), (4, 5549), (5, 5570), (6, 5585), (9, 5711),

Gene: Terapin_6 Start: 4196, Stop: 4468, Start Num: 1

Candidate Starts for Terapin_6:

(Start: 1 @4196 has 9 MA's), (5, 4298), (6, 4313), (9, 4439),