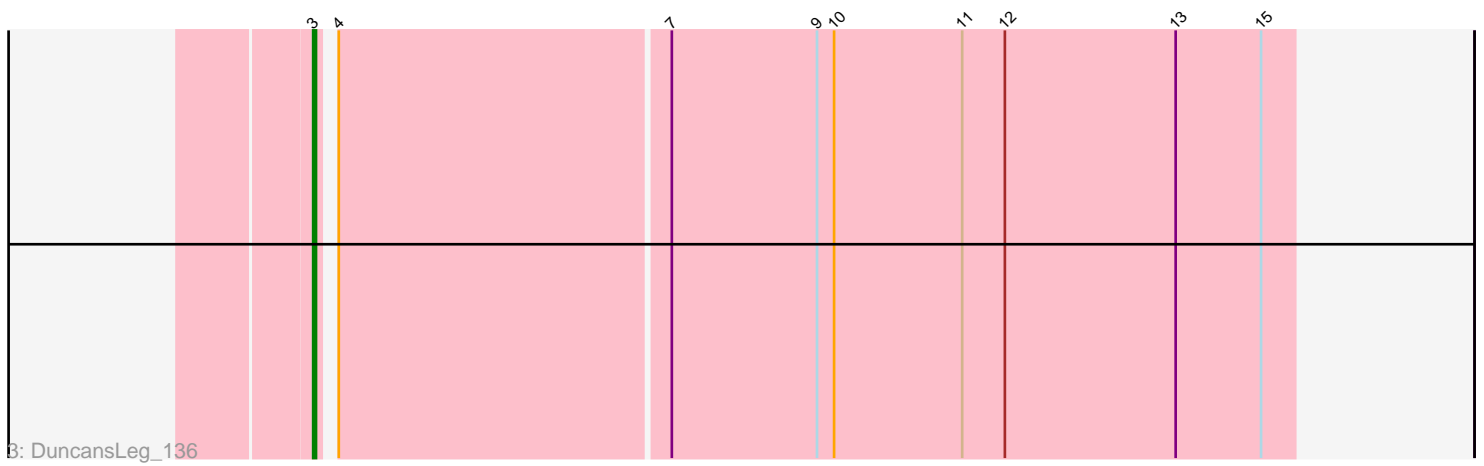
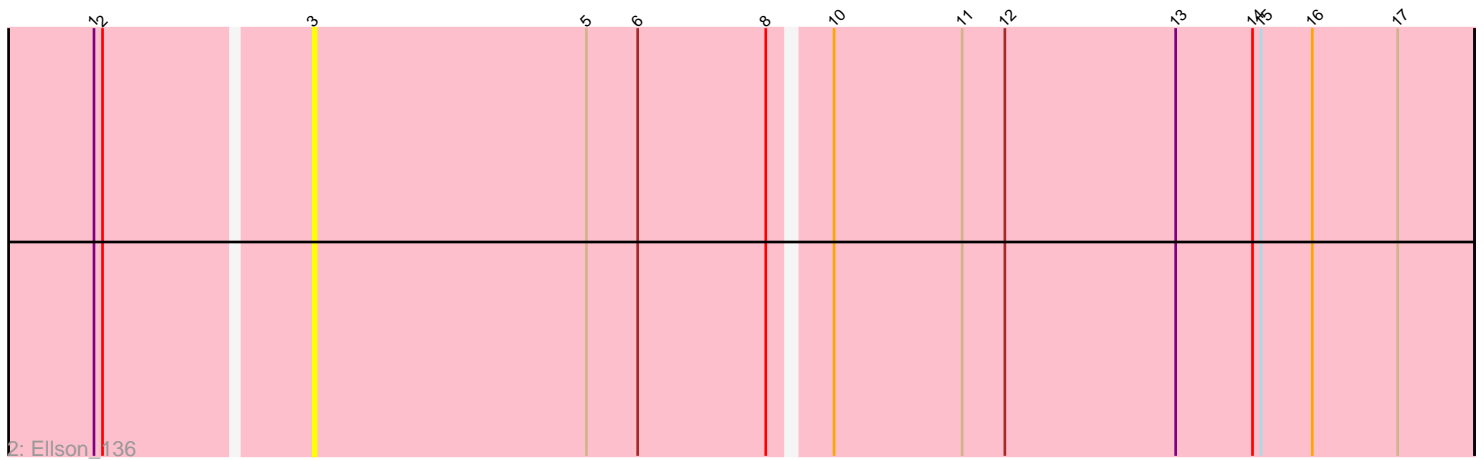
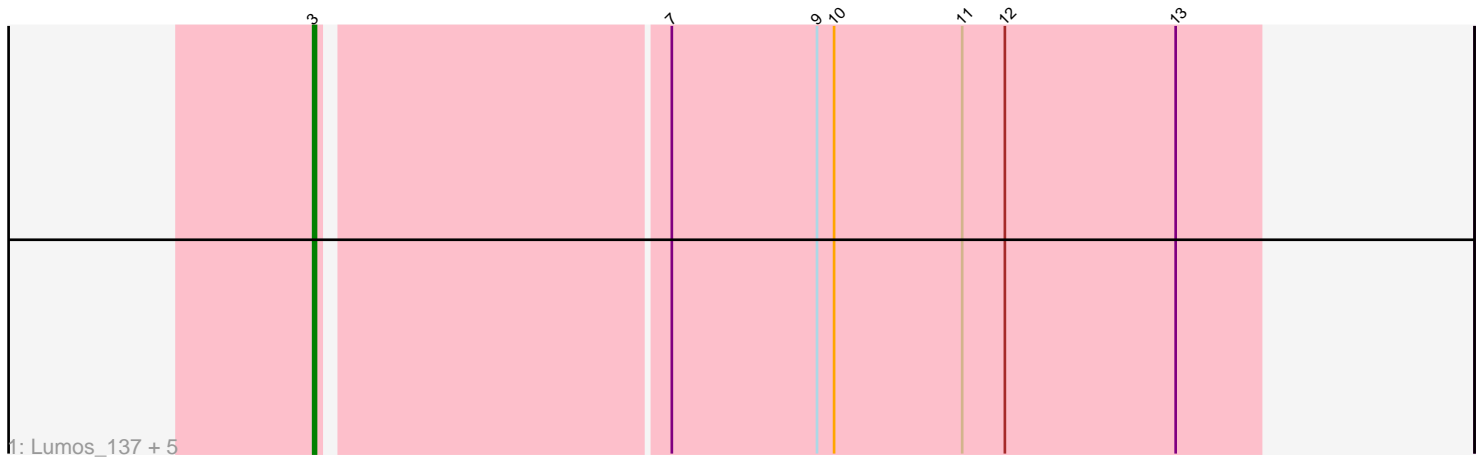


Pham 88130



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 88130 Report

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

Pham number 88130 has 8 members, 2 are drafts.

Phages represented in each track:

- Track 1 : Lumos_137, Jubie_138, Snenia_137, MsGreen_138, Jobypre_140, Clautastrophe_130
- Track 2 : Ellson_136
- Track 3 : DuncansLeg_136

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

The start number called the most often in the published annotations is 3, it was called in 6 of the 6 non-draft genes in the pham.

Genes that call this "Most Annotated" start:

- Clautastrophe_130, DuncansLeg_136, Ellson_136, Jobypre_140, Jubie_138, Lumos_137, MsGreen_138, Snenia_137,

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 3:

- Found in 8 of 8 (100.0%) of genes in pham
- Manual Annotations of this start: 6 of 6
- Called 100.0% of time when present
- Phage (with cluster) where this start called: Clautastrophe_130 (L3), DuncansLeg_136 (L3), Ellson_136 (L3), Jobypre_140 (L3), Jubie_138 (L3), Lumos_137 (L3), MsGreen_138 (L3), Snenia_137 (L3),

Summary by clusters:

There is one cluster represented in this pham: L3

Info for manual annotations of cluster L3:

- Start number 3 was manually annotated 6 times for cluster L3.

Gene Information:

Gene: Clautastrophe_130 Start: 70899, Stop: 70576, Start Num: 3

Candidate Starts for Clautastrophe_130:

(Start: 3 @70899 has 6 MA's), (7, 70782), (9, 70731), (10, 70725), (11, 70680), (12, 70665), (13, 70605),

Gene: DuncansLeg_136 Start: 74439, Stop: 74104, Start Num: 3

Candidate Starts for DuncansLeg_136:

(Start: 3 @74439 has 6 MA's), (4, 74436), (7, 74322), (9, 74271), (10, 74265), (11, 74220), (12, 74205), (13, 74145), (15, 74115),

Gene: Ellson_136 Start: 74808, Stop: 74407, Start Num: 3

Candidate Starts for Ellson_136:

(1, 74880), (2, 74877), (Start: 3 @74808 has 6 MA's), (5, 74712), (6, 74694), (8, 74649), (10, 74631), (11, 74586), (12, 74571), (13, 74511), (14, 74484), (15, 74481), (16, 74463), (17, 74433),

Gene: Jobypre_140 Start: 74463, Stop: 74140, Start Num: 3

Candidate Starts for Jobypre_140:

(Start: 3 @74463 has 6 MA's), (7, 74346), (9, 74295), (10, 74289), (11, 74244), (12, 74229), (13, 74169),

Gene: Jubie_138 Start: 74626, Stop: 74303, Start Num: 3

Candidate Starts for Jubie_138:

(Start: 3 @74626 has 6 MA's), (7, 74509), (9, 74458), (10, 74452), (11, 74407), (12, 74392), (13, 74332),

Gene: Lumos_137 Start: 74430, Stop: 74107, Start Num: 3

Candidate Starts for Lumos_137:

(Start: 3 @74430 has 6 MA's), (7, 74313), (9, 74262), (10, 74256), (11, 74211), (12, 74196), (13, 74136),

Gene: MsGreen_138 Start: 74463, Stop: 74140, Start Num: 3

Candidate Starts for MsGreen_138:

(Start: 3 @74463 has 6 MA's), (7, 74346), (9, 74295), (10, 74289), (11, 74244), (12, 74229), (13, 74169),

Gene: Snenia_137 Start: 74469, Stop: 74146, Start Num: 3

Candidate Starts for Snenia_137:

(Start: 3 @74469 has 6 MA's), (7, 74352), (9, 74301), (10, 74295), (11, 74250), (12, 74235), (13, 74175),