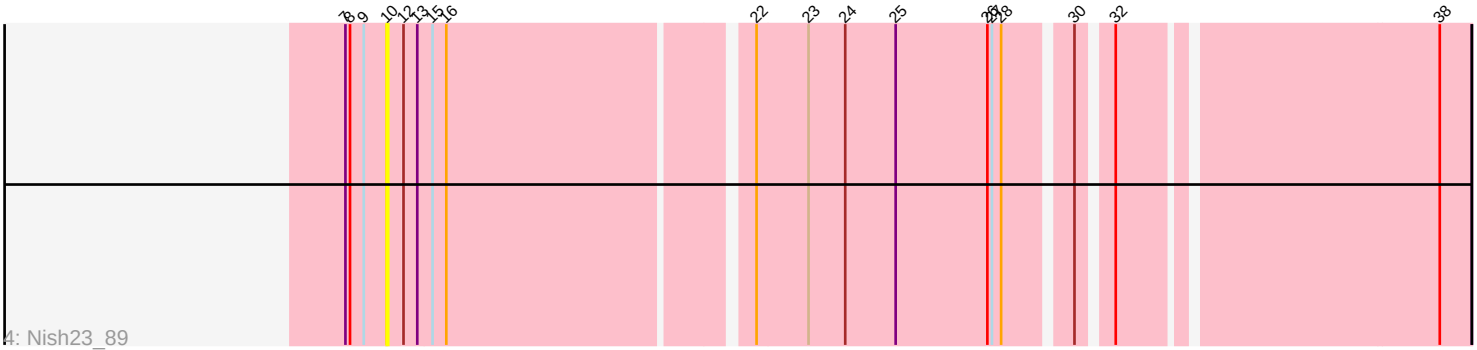
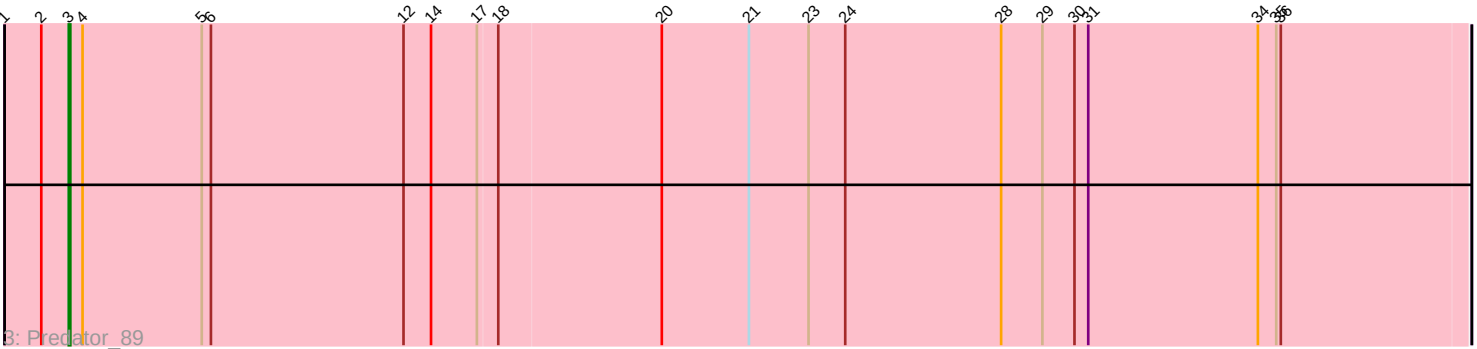
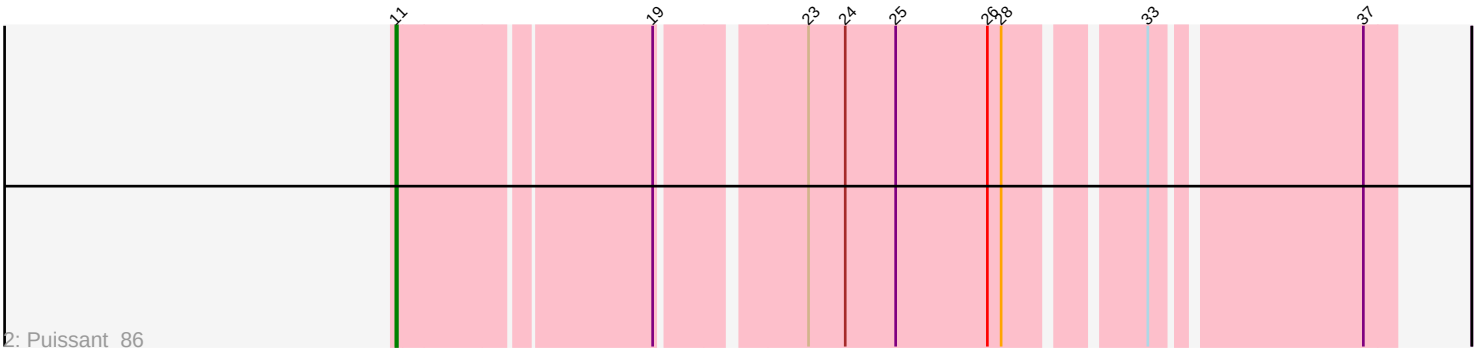
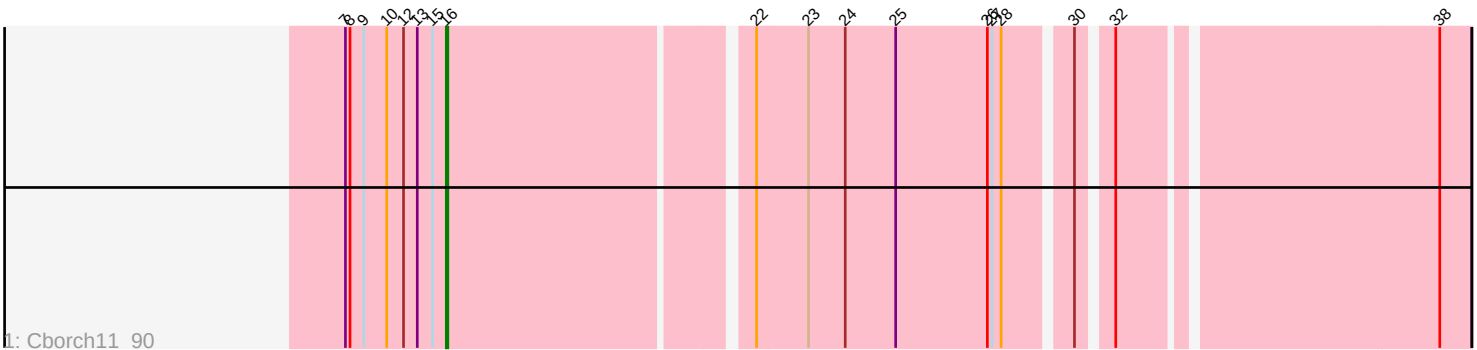


Pham 310280



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 310280 Report

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

Pham number 310280 has 4 members, 1 are drafts.

Phages represented in each track:

- Track 1 : Cborch11_90
- Track 2 : Puissant_86
- Track 3 : Predator_89
- Track 4 : Nish23_89

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

The start number called the most often in the published annotations is 11, it was called in 1 of the 3 non-draft genes in the pham.

Genes that call this "Most Annotated" start:

- Puissant_86,

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

-

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

- Cborch11_90, Nish23_89, Predator_89,

Summary by start number:

Start 3:

- Found in 1 of 4 (25.0%) of genes in pham
- Manual Annotations of this start: 1 of 3
- Called 100.0% of time when present
- Phage (with cluster) where this start called: Predator_89 (H1),

Start 10:

- Found in 2 of 4 (50.0%) of genes in pham
- No Manual Annotations of this start.
- Called 50.0% of time when present
- Phage (with cluster) where this start called: Nish23_89 (H1),

Start 11:

- Found in 1 of 4 (25.0%) of genes in pham
- Manual Annotations of this start: 1 of 3
- Called 100.0% of time when present
- Phage (with cluster) where this start called: Puissant_86 (H1),

Start 16:

- Found in 2 of 4 (50.0%) of genes in pham
- Manual Annotations of this start: 1 of 3
- Called 50.0% of time when present
- Phage (with cluster) where this start called: Cborch11_90 (H1),

Summary by clusters:

There is one cluster represented in this pham: H1

Info for manual annotations of cluster H1:

- Start number 3 was manually annotated 1 time for cluster H1.
- Start number 11 was manually annotated 1 time for cluster H1.
- Start number 16 was manually annotated 1 time for cluster H1.

Gene Information:

Gene: Cborch11_90 Start: 63152, Stop: 63769, Start Num: 16

Candidate Starts for Cborch11_90:

(7, 63089), (8, 63092), (9, 63101), (10, 63116), (12, 63125), (13, 63134), (15, 63143), (Start: 16 @63152 has 1 MA's), (22, 63338), (23, 63371), (24, 63395), (25, 63428), (26, 63488), (27, 63491), (28, 63497), (30, 63536), (32, 63554), (38, 63749),

Gene: Nish23_89 Start: 63116, Stop: 63769, Start Num: 10

Candidate Starts for Nish23_89:

(7, 63089), (8, 63092), (9, 63101), (10, 63116), (12, 63125), (13, 63134), (15, 63143), (Start: 16 @63152 has 1 MA's), (22, 63338), (23, 63371), (24, 63395), (25, 63428), (26, 63488), (27, 63491), (28, 63497), (30, 63536), (32, 63554), (38, 63749),

Gene: Predator_89 Start: 64539, Stop: 65447, Start Num: 3

Candidate Starts for Predator_89:

(1, 64497), (2, 64521), (Start: 3 @64539 has 1 MA's), (4, 64548), (5, 64626), (6, 64632), (12, 64758), (14, 64776), (17, 64806), (18, 64818), (20, 64923), (21, 64980), (23, 65019), (24, 65043), (28, 65145), (29, 65172), (30, 65193), (31, 65202), (34, 65313), (35, 65325), (36, 65328),

Gene: Puissant_86 Start: 64050, Stop: 64640, Start Num: 11

Candidate Starts for Puissant_86:

(Start: 11 @64050 has 1 MA's), (19, 64206), (23, 64290), (24, 64314), (25, 64347), (26, 64407), (28, 64416), (33, 64494), (37, 64620),