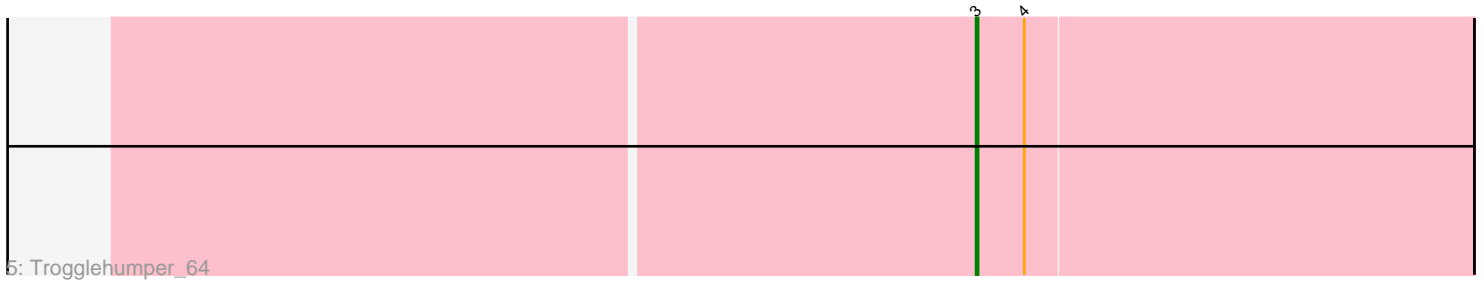
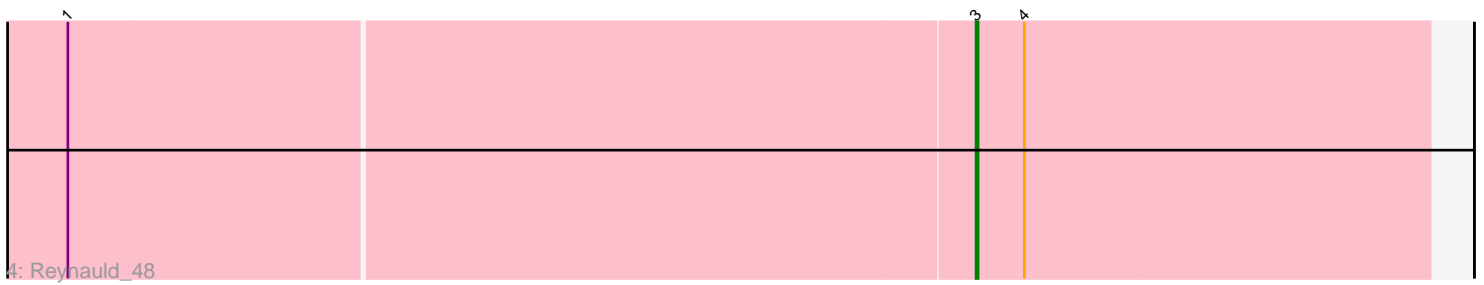
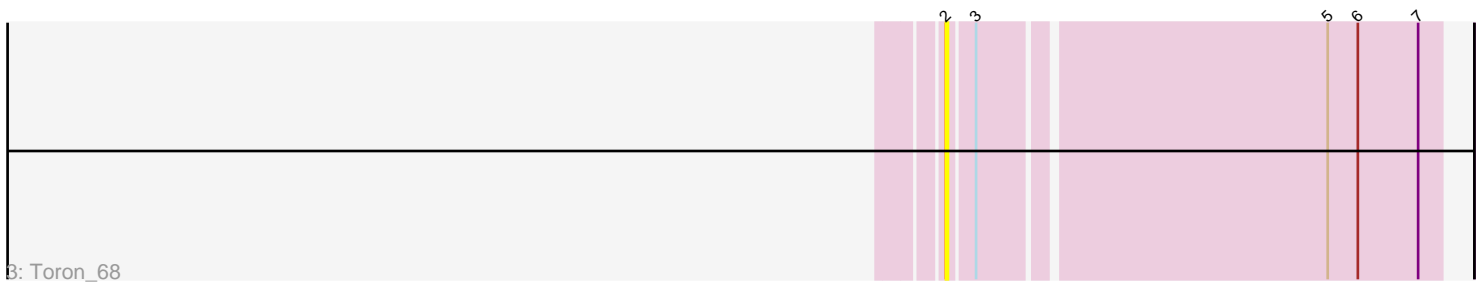
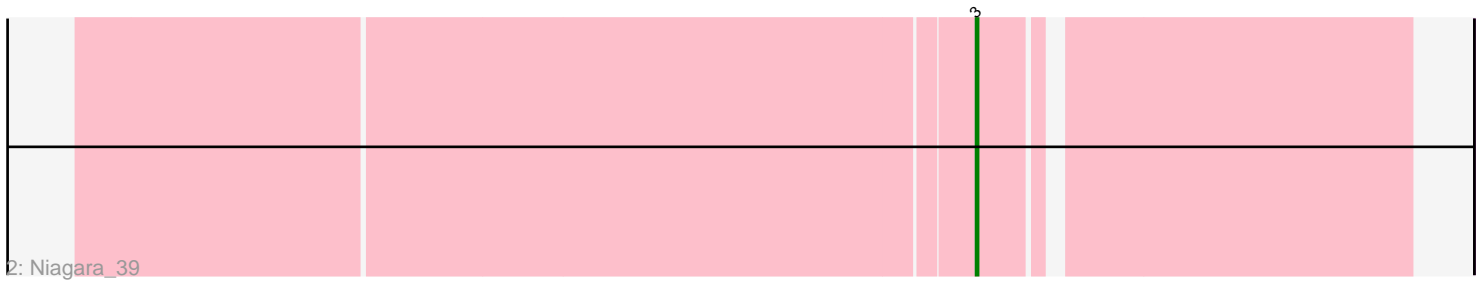


Pham 216705



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 216705 Report

This analysis was run 02/22/25 on database version 588.

Pham number 216705 has 13 members, 2 are drafts.

Phages represented in each track:

- Track 1 : Benczkowski14_39, Kvothe_39, Tredge_40, Demosthenes_39, Teatealatte_40, Katyusha_39, Hollow_40, ASerpRocky_40, Teech_40
- Track 2 : Niagara_39
- Track 3 : Toron_68
- Track 4 : Reynauld_48
- Track 5 : Trogglehumper_64

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 11 of the 11 non-draft genes in the pham.

Genes that call this "Most Annotated" start:

- ASerpRocky_40, Benczkowski14_39, Demosthenes_39, Hollow_40, Katyusha_39, Kvothe_39, Niagara_39, Reynauld_48, Teatealatte_40, Teech_40, Tredge_40, Trogglehumper_64,

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

- Toron_68,

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

-

Summary by start number:

Start 2:

- Found in 1 of 13 (7.7%) of genes in pham
- No Manual Annotations of this start.
- Called 100.0% of time when present
- Phage (with cluster) where this start called: Toron_68 (F6),

Start 3:

- Found in 13 of 13 (100.0%) of genes in pham
- Manual Annotations of this start: 11 of 11

- Called 92.3% of time when present
- Phage (with cluster) where this start called: ASerpRocky_40 (CS4), Benczkowski14_39 (CS4), Demosthenes_39 (CS4), Hollow_40 (CS4), Katyusha_39 (CS4), Kvothe_39 (CS4), Niagara_39 (CS4), Reynauld_48 (singleton), Teatealatte_40 (CS4), Teech_40 (CS4), Tredge_40 (CS4), Trogglehumper_64 (singleton),

Summary by clusters:

There are 3 clusters represented in this pham: CS4, singleton, F6,

Info for manual annotations of cluster CS4:

- Start number 3 was manually annotated 9 times for cluster CS4.

Gene Information:

Gene: ASerpRocky_40 Start: 44241, Stop: 44041, Start Num: 3

Candidate Starts for ASerpRocky_40:

(Start: 3 @44241 has 11 MA's),

Gene: Benczkowski14_39 Start: 44274, Stop: 44074, Start Num: 3

Candidate Starts for Benczkowski14_39:

(Start: 3 @44274 has 11 MA's),

Gene: Demosthenes_39 Start: 44214, Stop: 44014, Start Num: 3

Candidate Starts for Demosthenes_39:

(Start: 3 @44214 has 11 MA's),

Gene: Hollow_40 Start: 44620, Stop: 44420, Start Num: 3

Candidate Starts for Hollow_40:

(Start: 3 @44620 has 11 MA's),

Gene: Katyusha_39 Start: 44274, Stop: 44074, Start Num: 3

Candidate Starts for Katyusha_39:

(Start: 3 @44274 has 11 MA's),

Gene: Kvothe_39 Start: 44173, Stop: 43973, Start Num: 3

Candidate Starts for Kvothe_39:

(Start: 3 @44173 has 11 MA's),

Gene: Niagara_39 Start: 44231, Stop: 44031, Start Num: 3

Candidate Starts for Niagara_39:

(Start: 3 @44231 has 11 MA's),

Gene: Reynauld_48 Start: 45364, Stop: 45140, Start Num: 3

Candidate Starts for Reynauld_48:

(1, 45808), (Start: 3 @45364 has 11 MA's), (4, 45340),

Gene: Teatealatte_40 Start: 44176, Stop: 43976, Start Num: 3

Candidate Starts for Teatealatte_40:

(Start: 3 @44176 has 11 MA's),

Gene: Teech_40 Start: 43977, Stop: 43777, Start Num: 3
Candidate Starts for Teech_40:
(Start: 3 @43977 has 11 MA's),

Gene: Toron_68 Start: 42444, Stop: 42677, Start Num: 2
Candidate Starts for Toron_68:
(2, 42444), (Start: 3 @42456 has 11 MA's), (5, 42621), (6, 42636), (7, 42666),

Gene: Tredge_40 Start: 44176, Stop: 43976, Start Num: 3
Candidate Starts for Tredge_40:
(Start: 3 @44176 has 11 MA's),

Gene: Trogglehumper_64 Start: 53677, Stop: 53435, Start Num: 3
Candidate Starts for Trogglehumper_64:
(Start: 3 @53677 has 11 MA's), (4, 53653),