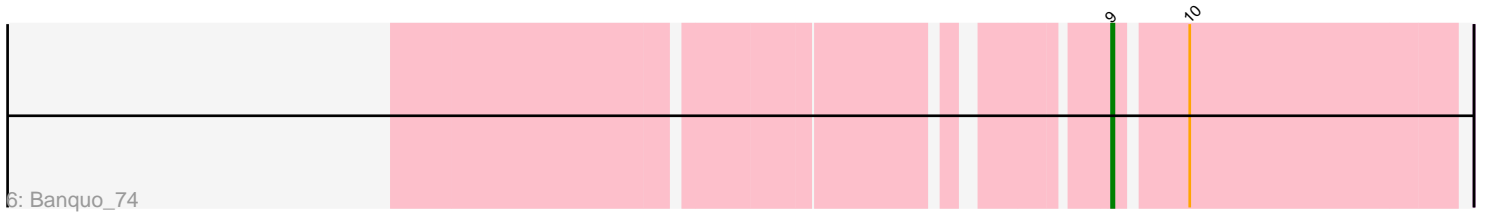
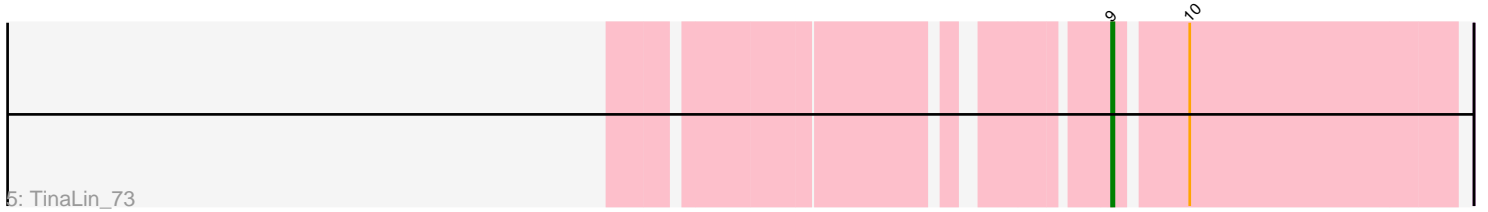
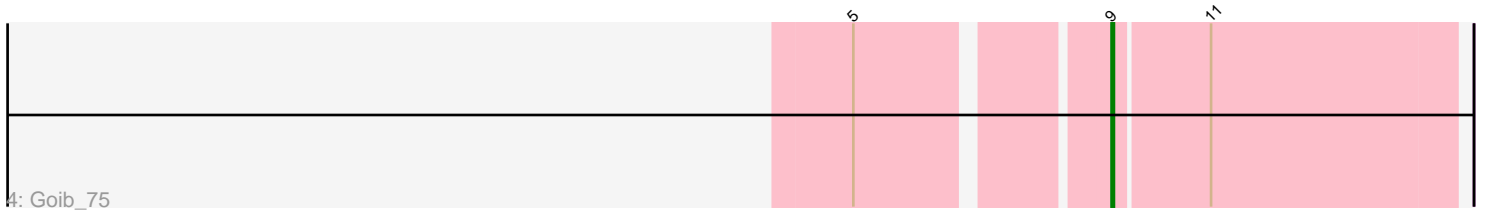
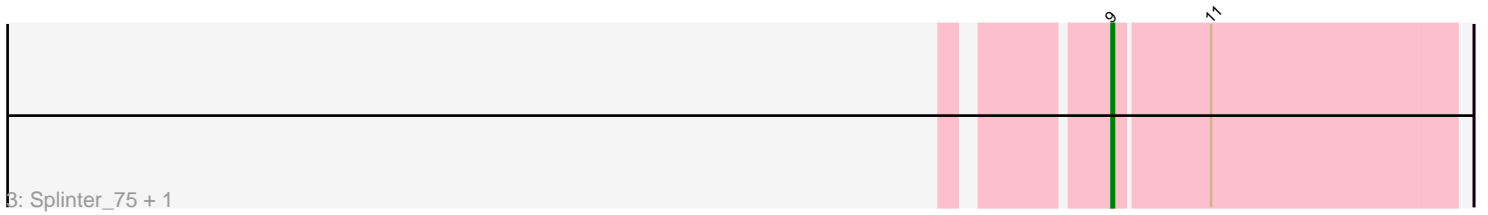
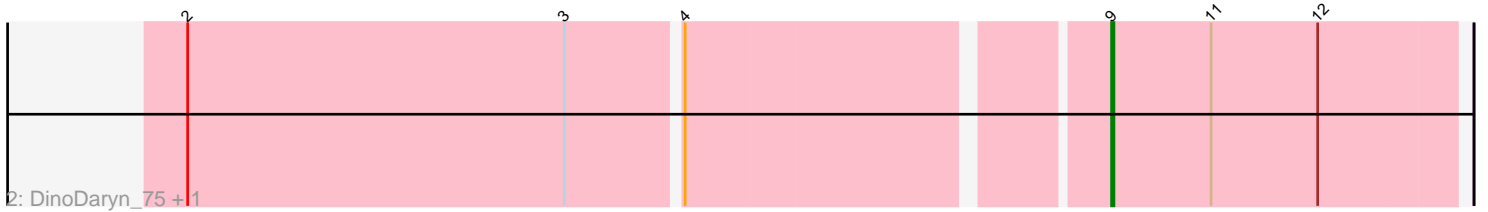
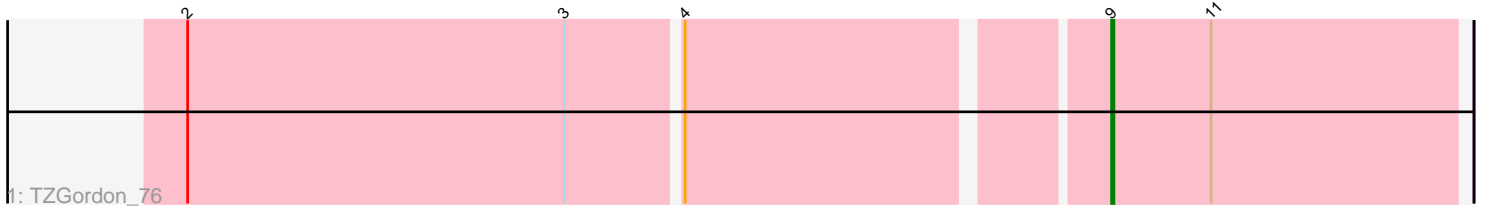


Pham 6400



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 6400 Report

This analysis was run 07/09/24 on database version 566.

Pham number 6400 has 9 members, 0 are drafts.

Phages represented in each track:

- Track 1 : TZGordon_76
- Track 2 : DinoDaryn_75, Huff_75
- Track 3 : Splinter_75, Vendetta_75
- Track 4 : Goib_75
- Track 5 : TinaLin_73
- Track 6 : Banquo_74
- Track 7 : Schmidt_75

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

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

Genes that call this "Most Annotated" start:

- Banquo_74, DinoDaryn_75, Goib_75, Huff_75, Schmidt_75, Splinter_75, TZGordon_76, TinaLin_73, Vendetta_75,

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 9:

- 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: Banquo_74 (CU1), DinoDaryn_75 (CU1), Goib_75 (CU1), Huff_75 (CU1), Schmidt_75 (CU4), Splinter_75 (CU1), TZGordon_76 (CU1), TinaLin_73 (CU1), Vendetta_75 (CU1),

Summary by clusters:

There are 2 clusters represented in this pham: CU4, CU1,

Info for manual annotations of cluster CU1:

•Start number 9 was manually annotated 8 times for cluster CU1.

Info for manual annotations of cluster CU4:

•Start number 9 was manually annotated 1 time for cluster CU4.

Gene Information:

Gene: Banquo_74 Start: 42885, Stop: 43022, Start Num: 9

Candidate Starts for Banquo_74:

(Start: 9 @42885 has 9 MA's), (10, 42912),

Gene: DinoDaryn_75 Start: 43179, Stop: 43322, Start Num: 9

Candidate Starts for DinoDaryn_75:

(2, 42813), (3, 42972), (4, 43017), (Start: 9 @43179 has 9 MA's), (11, 43221), (12, 43266),

Gene: Goib_75 Start: 44282, Stop: 44422, Start Num: 9

Candidate Starts for Goib_75:

(5, 44189), (Start: 9 @44282 has 9 MA's), (11, 44321),

Gene: Huffy_75 Start: 43179, Stop: 43322, Start Num: 9

Candidate Starts for Huffy_75:

(2, 42813), (3, 42972), (4, 43017), (Start: 9 @43179 has 9 MA's), (11, 43221), (12, 43266),

Gene: Schmidt_75 Start: 42708, Stop: 42860, Start Num: 9

Candidate Starts for Schmidt_75:

(1, 42273), (6, 42618), (7, 42657), (8, 42702), (Start: 9 @42708 has 9 MA's), (11, 42750), (13, 42837), (14, 42840),

Gene: Splinter_75 Start: 44266, Stop: 44406, Start Num: 9

Candidate Starts for Splinter_75:

(Start: 9 @44266 has 9 MA's), (11, 44305),

Gene: TZGordon_76 Start: 43146, Stop: 43289, Start Num: 9

Candidate Starts for TZGordon_76:

(2, 42780), (3, 42939), (4, 42984), (Start: 9 @43146 has 9 MA's), (11, 43188),

Gene: TinaLin_73 Start: 42583, Stop: 42720, Start Num: 9

Candidate Starts for TinaLin_73:

(Start: 9 @42583 has 9 MA's), (10, 42610),

Gene: Vendetta_75 Start: 44266, Stop: 44406, Start Num: 9

Candidate Starts for Vendetta_75:

(Start: 9 @44266 has 9 MA's), (11, 44305),