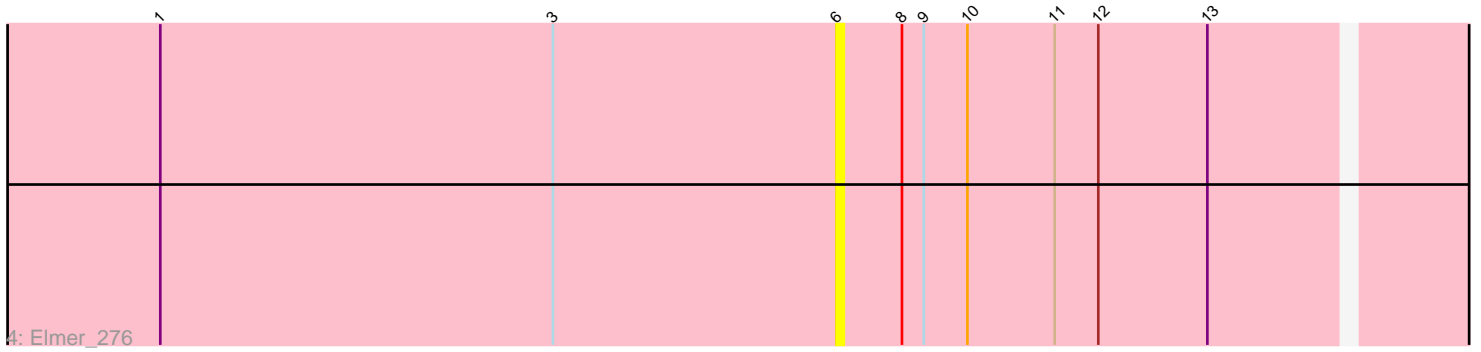
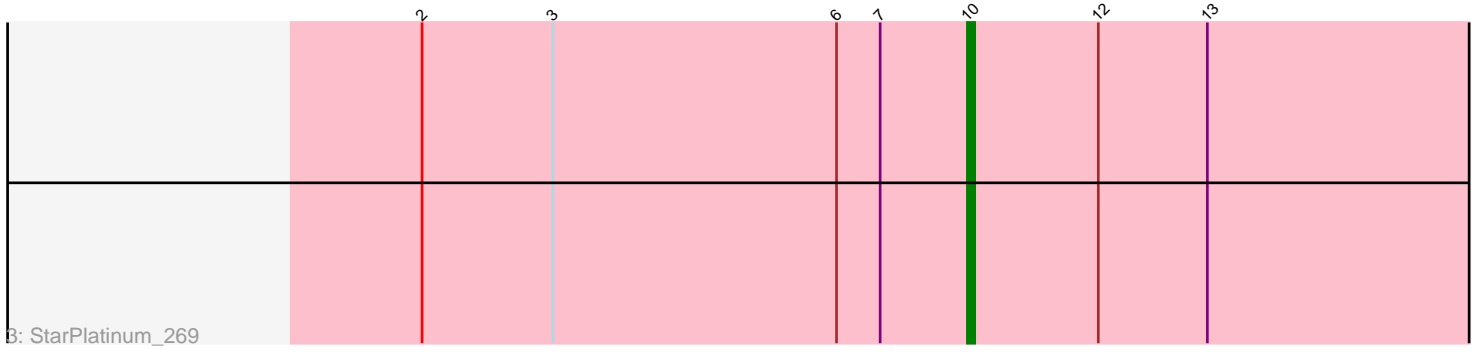
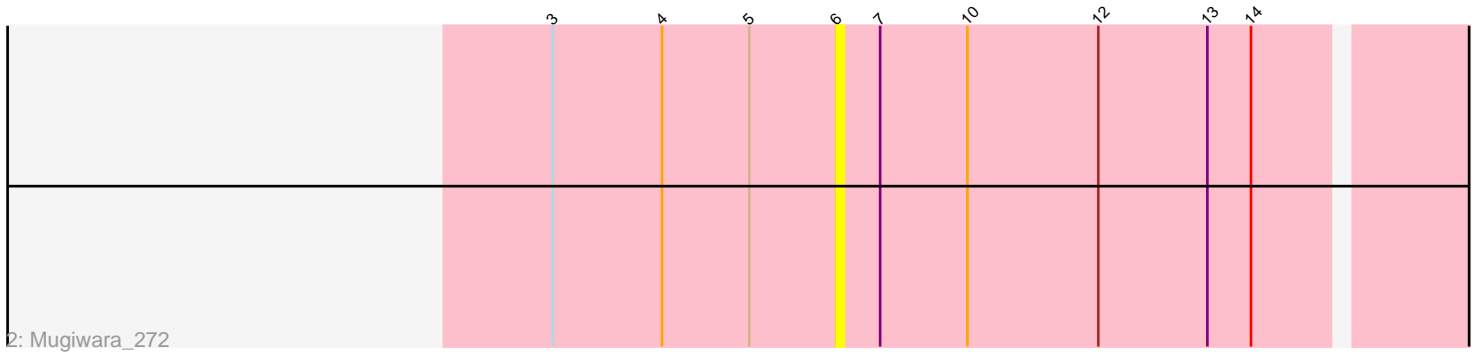
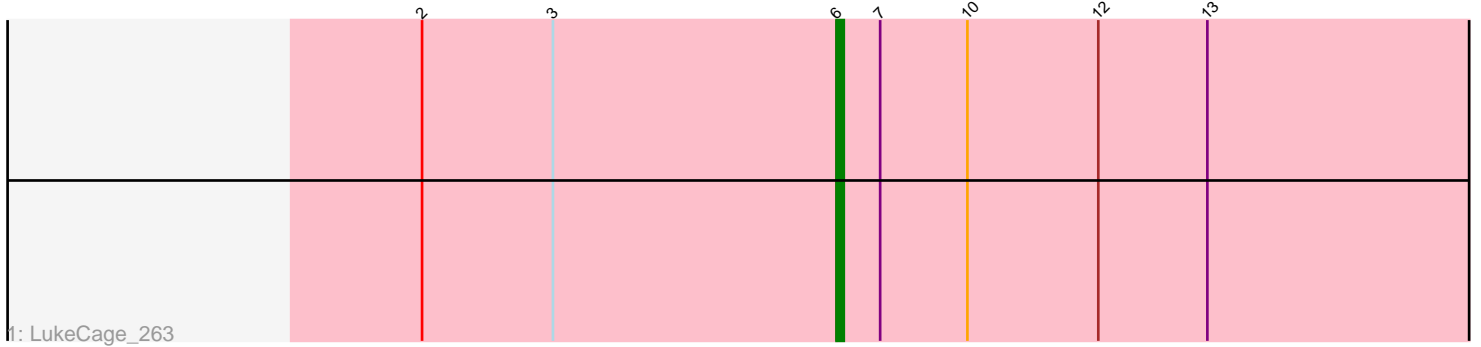


Pham 164320



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 164320 Report

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

Pham number 164320 has 4 members, 2 are drafts.

Phages represented in each track:

- Track 1 : LukeCage_263
- Track 2 : Mugiwara_272
- Track 3 : StarPlatinum_269
- Track 4 : Elmer_276

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

The start number called the most often in the published annotations is 6, it was called in 1 of the 2 non-draft genes in the pham.

Genes that call this "Most Annotated" start:

- Elmer_276, LukeCage_263, Mugiwara_272,

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

- StarPlatinum_269,

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

-

Summary by start number:

Start 6:

- Found in 4 of 4 (100.0%) of genes in pham
- Manual Annotations of this start: 1 of 2
- Called 75.0% of time when present
- Phage (with cluster) where this start called: Elmer_276 (BE2), LukeCage_263 (BE2), Mugiwara_272 (BE2),

Start 10:

- Found in 4 of 4 (100.0%) of genes in pham
- Manual Annotations of this start: 1 of 2
- Called 25.0% of time when present
- Phage (with cluster) where this start called: StarPlatinum_269 (BE2),

Summary by clusters:

There is one cluster represented in this pham: BE2

Info for manual annotations of cluster BE2:

- Start number 6 was manually annotated 1 time for cluster BE2.
- Start number 10 was manually annotated 1 time for cluster BE2.

Gene Information:

Gene: Elmer_276 Start: 121832, Stop: 121954, Start Num: 6

Candidate Starts for Elmer_276:

(1, 121739), (3, 121793), (Start: 6 @121832 has 1 MA's), (8, 121841), (9, 121844), (Start: 10 @121850 has 1 MA's), (11, 121862), (12, 121868), (13, 121883),

Gene: LukeCage_263 Start: 120379, Stop: 120528, Start Num: 6

Candidate Starts for LukeCage_263:

(2, 120322), (3, 120340), (Start: 6 @120379 has 1 MA's), (7, 120385), (Start: 10 @120397 has 1 MA's), (12, 120415), (13, 120430),

Gene: Mugiwara_272 Start: 120861, Stop: 120980, Start Num: 6

Candidate Starts for Mugiwara_272:

(3, 120822), (4, 120837), (5, 120849), (Start: 6 @120861 has 1 MA's), (7, 120867), (Start: 10 @120879 has 1 MA's), (12, 120897), (13, 120912), (14, 120918),

Gene: StarPlatinum_269 Start: 121173, Stop: 121304, Start Num: 10

Candidate Starts for StarPlatinum_269:

(2, 121098), (3, 121116), (Start: 6 @121155 has 1 MA's), (7, 121161), (Start: 10 @121173 has 1 MA's), (12, 121191), (13, 121206),