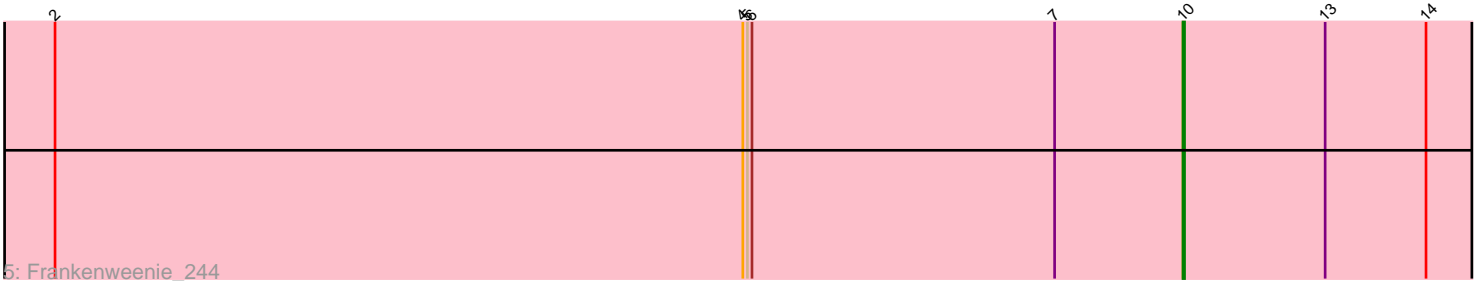
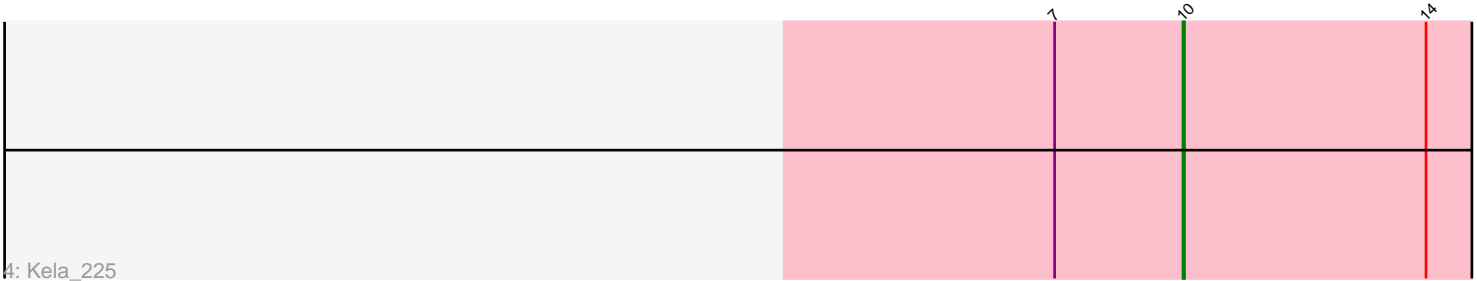
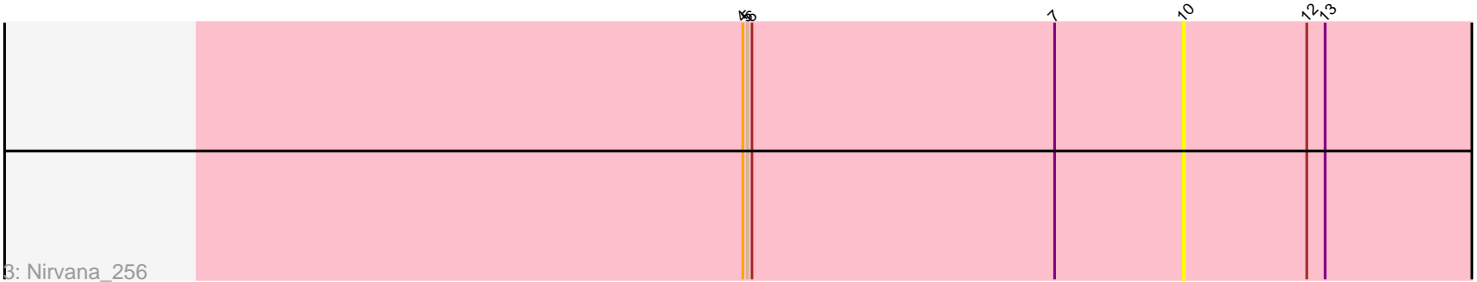
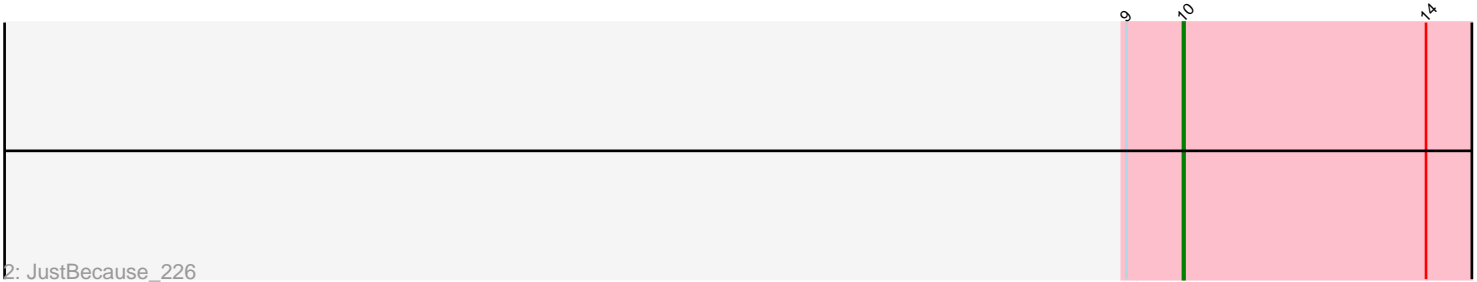
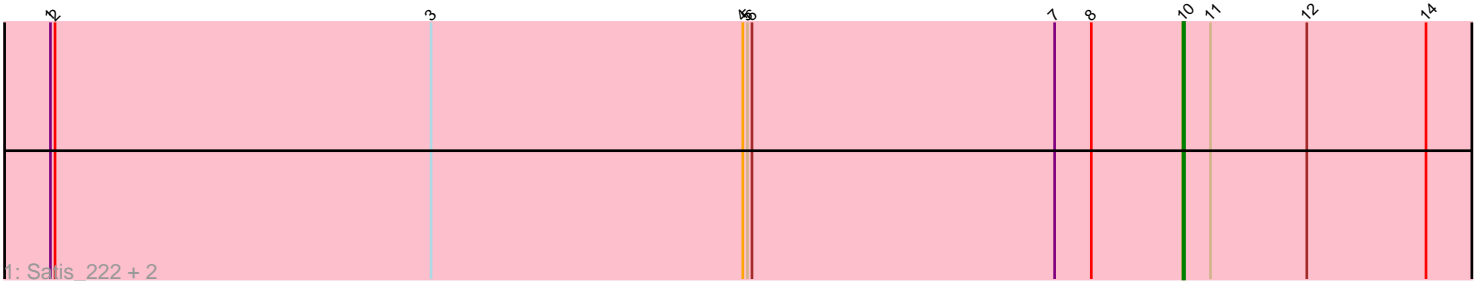


Pham 7552



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 7552 Report

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

Pham number 7552 has 7 members, 1 are drafts.

Phages represented in each track:

- Track 1 : Satis_222, Kradal_224, EhyElimayoE_225
- Track 2 : JustBecause_226
- Track 3 : Nirvana_256
- Track 4 : Kela_225
- Track 5 : Frankenweenie_244

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

The start number called the most often in the published annotations is 10, it was called in 6 of the 6 non-draft genes in the pham.

Genes that call this "Most Annotated" start:

- EhyElimayoE_225, Frankenweenie_244, JustBecause_226, Kela_225, Kradal_224, Nirvana_256, Satis_222,

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 10:

- Found in 7 of 7 (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: EhyElimayoE_225 (BM), Frankenweenie_244 (BM), JustBecause_226 (BM), Kela_225 (BM), Kradal_224 (BM), Nirvana_256 (BM), Satis_222 (BM),

Summary by clusters:

There is one cluster represented in this pham: BM

Info for manual annotations of cluster BM:

- Start number 10 was manually annotated 6 times for cluster BM.

Gene Information:

Gene: EhyElimayoE_225 Start: 134319, Stop: 134582, Start Num: 10

Candidate Starts for EhyElimayoE_225:

(1, 133578), (2, 133581), (3, 133827), (4, 134031), (5, 134034), (6, 134037), (7, 134235), (8, 134259),
(Start: 10 @134319 has 6 MA's), (11, 134337), (12, 134400), (14, 134478),

Gene: Frankenweenie_244 Start: 145531, Stop: 145797, Start Num: 10

Candidate Starts for Frankenweenie_244:

(2, 144793), (4, 145243), (5, 145246), (6, 145249), (7, 145447), (Start: 10 @145531 has 6 MA's), (13, 145624), (14, 145690),

Gene: JustBecause_226 Start: 132496, Stop: 132759, Start Num: 10

Candidate Starts for JustBecause_226:

(9, 132460), (Start: 10 @132496 has 6 MA's), (14, 132655),

Gene: Kela_225 Start: 133079, Stop: 133342, Start Num: 10

Candidate Starts for Kela_225:

(7, 132995), (Start: 10 @133079 has 6 MA's), (14, 133238),

Gene: Kradal_224 Start: 134316, Stop: 134579, Start Num: 10

Candidate Starts for Kradal_224:

(1, 133575), (2, 133578), (3, 133824), (4, 134028), (5, 134031), (6, 134034), (7, 134232), (8, 134256),
(Start: 10 @134316 has 6 MA's), (11, 134334), (12, 134397), (14, 134475),

Gene: Nirvana_256 Start: 144385, Stop: 144651, Start Num: 10

Candidate Starts for Nirvana_256:

(4, 144097), (5, 144100), (6, 144103), (7, 144301), (Start: 10 @144385 has 6 MA's), (12, 144466), (13, 144478),

Gene: Satis_222 Start: 134653, Stop: 134916, Start Num: 10

Candidate Starts for Satis_222:

(1, 133912), (2, 133915), (3, 134161), (4, 134365), (5, 134368), (6, 134371), (7, 134569), (8, 134593),
(Start: 10 @134653 has 6 MA's), (11, 134671), (12, 134734), (14, 134812),