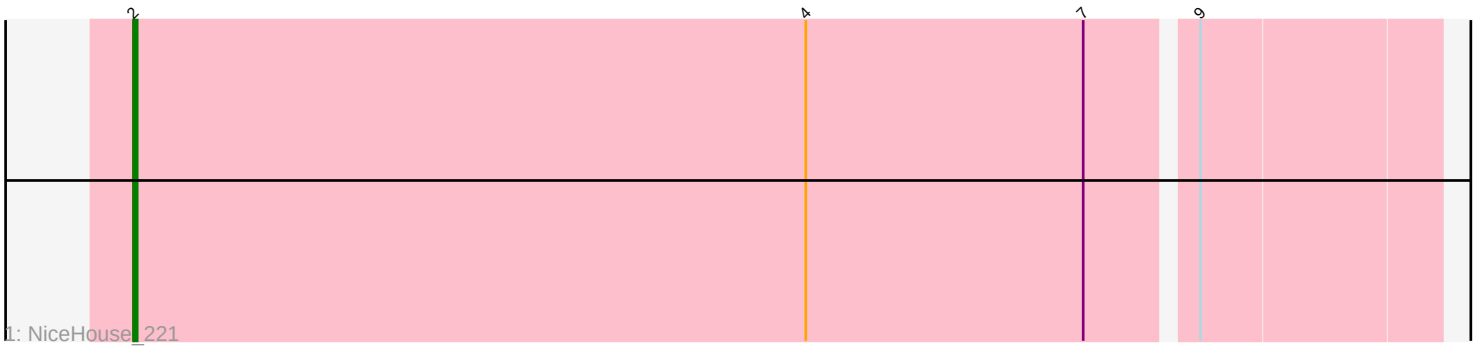
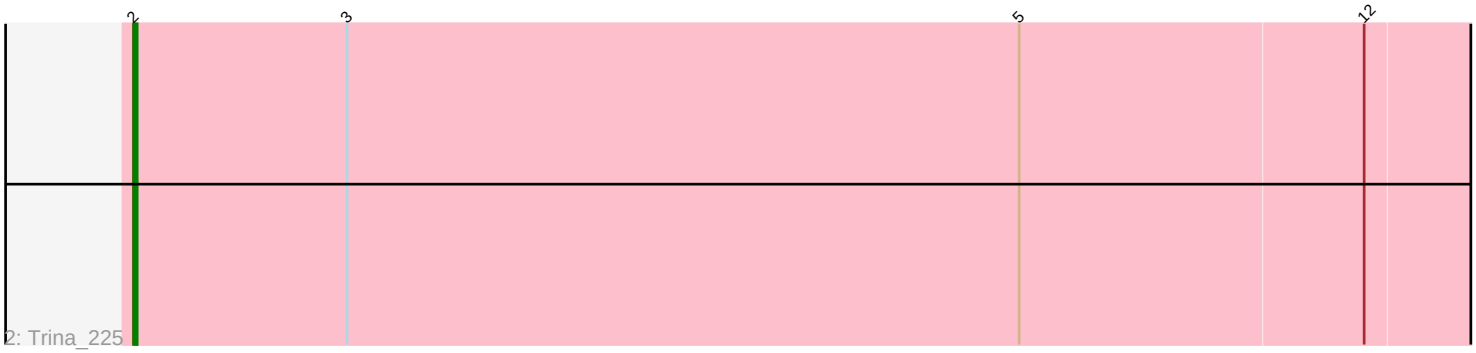
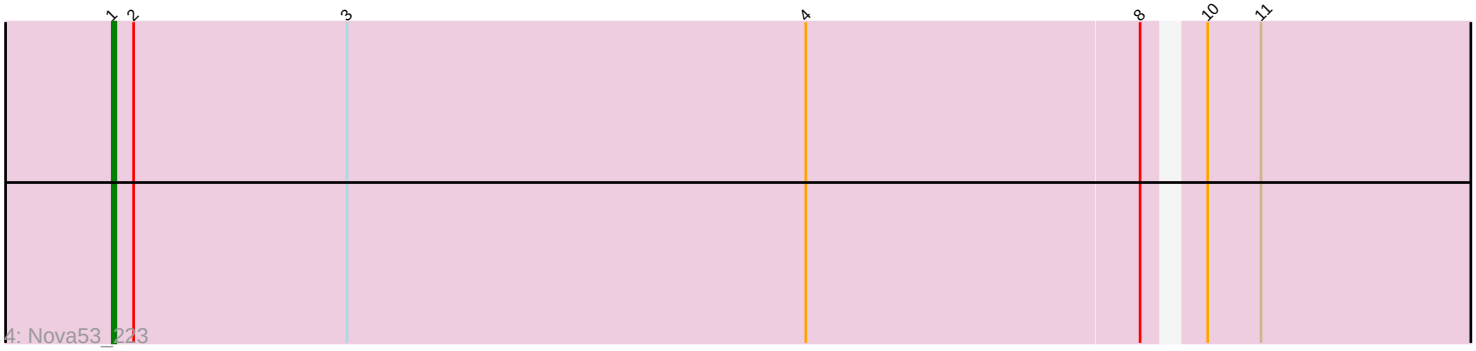
Pham 335746


2
4
7
9
1: NiceHouse_221


2
3
5
12
2: Trina_225


2
6
8
3: Francesca_212 + 1


1
2
3
4
8
10
11
4: Nova53_233

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 335746 Report

This analysis was run 11/14/26 on database version 672.

Pham number 335746 has 5 members, 0 are drafts.

Phages represented in each track:
• Track 1 : NiceHouse_221
• Track 2 : Trina_225
• Track 3 : Francesca_212, Dorin_211
• Track 4 : Nova53_223

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

The start number called the most often in the published annotations is 2, it was called in 4 of the 5 non-draft genes in the pham.

Genes that call this "Most Annotated" start:
• Dorin_211, Francesca_212, NiceHouse_221, Trina_225,

Genes that have the "Most Annotated" start but do not call it:
• Nova53_223,

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

Summary by start number:

Start 1:
• Found in 1 of 5 (20.0%) of genes in pham
• Manual Annotations of this start: 1 of 5
• Called 100.0% of time when present
• Phage (with cluster) where this start called: Nova53_223 (CG),

Start 2:
• Found in 5 of 5 (100.0%) of genes in pham
• Manual Annotations of this start: 4 of 5
• Called 80.0% of time when present
• Phage (with cluster) where this start called: Dorin_211 (CG), Francesca_212 (CG), NiceHouse_221 (CE), Trina_225 (CE),

Summary by clusters:

There are 2 clusters represented in this pham: CG, CE,

Info for manual annotations of cluster CE:
•Start number 2 was manually annotated 2 times for cluster CE.

Info for manual annotations of cluster CG:
•Start number 1 was manually annotated 1 time for cluster CG.
•Start number 2 was manually annotated 2 times for cluster CG.

Gene Information:

Gene: Dorin_211 Start: 112878, Stop: 113237, Start Num: 2
Candidate Starts for Dorin_211:
(Start: 2 @112878 has 4 MA's), (6, 113139), (8, 113160),

Gene: Francesca_212 Start: 113530, Stop: 113889, Start Num: 2
Candidate Starts for Francesca_212:
(Start: 2 @113530 has 4 MA's), (6, 113791), (8, 113812),

Gene: NiceHouse_221 Start: 119267, Stop: 119626, Start Num: 2
Candidate Starts for NiceHouse_221:
(Start: 2 @119267 has 4 MA's), (4, 119456), (7, 119534), (9, 119561),

Gene: Nova53_223 Start: 117635, Stop: 118009, Start Num: 1
Candidate Starts for Nova53_223:
(Start: 1 @117635 has 1 MA's), (Start: 2 @117641 has 4 MA's), (3, 117701), (4, 117830), (8, 117923), (10, 117935), (11, 117950),

Gene: Trina_225 Start: 118130, Stop: 118504, Start Num: 2
Candidate Starts for Trina_225:
(Start: 2 @118130 has 4 MA's), (3, 118190), (5, 118379), (12, 118475),