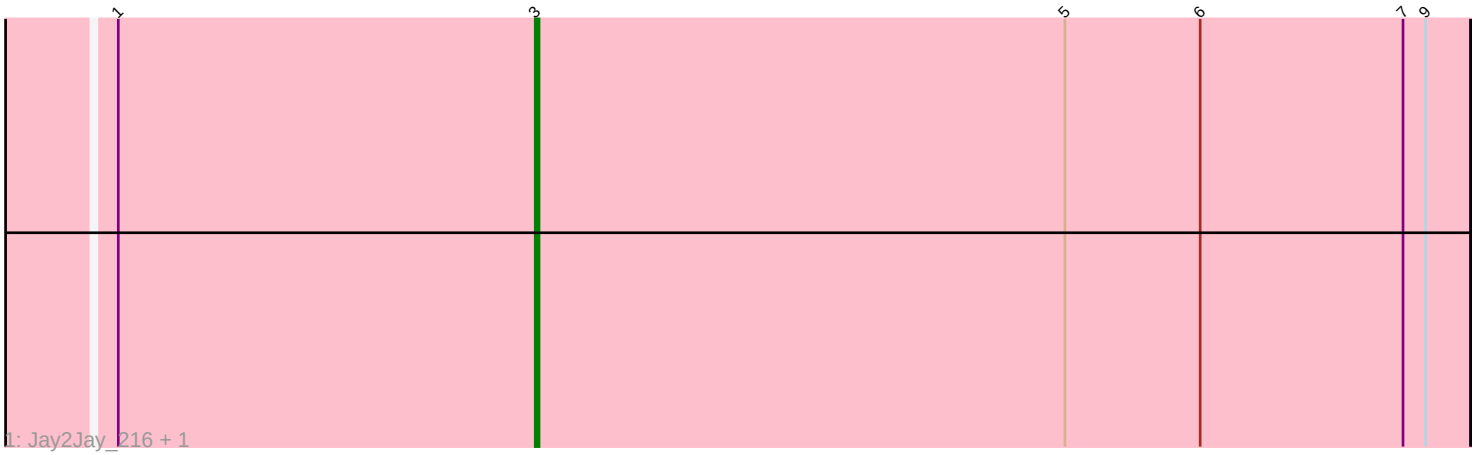
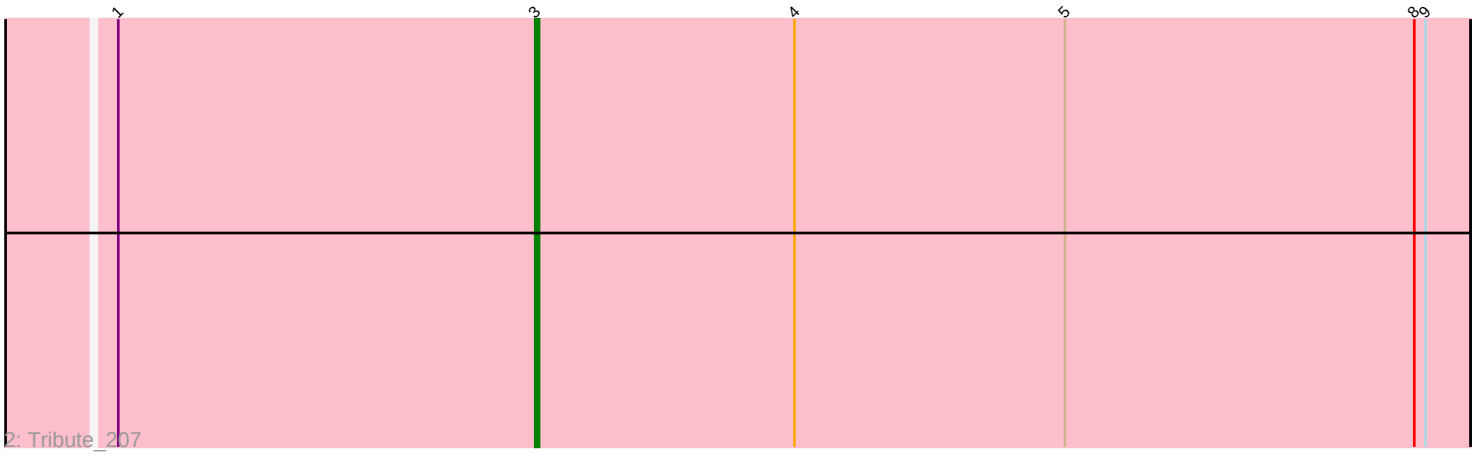
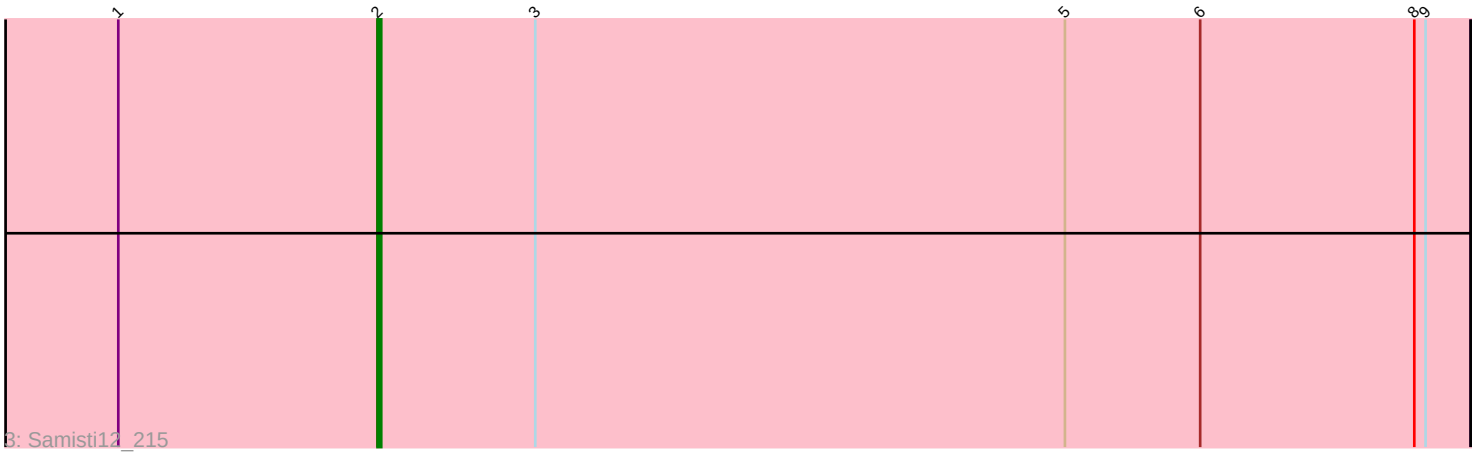
Pham 346648


1
3
5
6
7
9
1: Jay2Jay_216 + 1


1
3
4
5
8
9
2: Tribute_207


1
2
3
5
6
8
9
3: Samisti13_215

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 346648 Report

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

Pham number 346648 has 4 members, 0 are drafts.

Phages represented in each track:
• Track 1 : Jay2Jay_216, Warpy_214
• Track 2 : Tribute_207
• Track 3 : Samisti12_215

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 3 of the 4 non-draft genes in the pham.

Genes that call this "Most Annotated" start:
• Jay2Jay_216, Tribute_207, Warpy_214,

Genes that have the "Most Annotated" start but do not call it:
• Samisti12_215,

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

Summary by start number:

Start 2:
• Found in 1 of 4 (25.0%) of genes in pham
• Manual Annotations of this start: 1 of 4
• Called 100.0% of time when present
• Phage (with cluster) where this start called: Samisti12_215 (BE1),

Start 3:
• Found in 4 of 4 (100.0%) of genes in pham
• Manual Annotations of this start: 3 of 4
• Called 75.0% of time when present
• Phage (with cluster) where this start called: Jay2Jay_216 (BE1), Tribute_207 (BE1), Warpy_214 (BE1),

Summary by clusters:

There is one cluster represented in this pham: BE1

Info for manual annotations of cluster BE1:
•Start number 2 was manually annotated 1 time for cluster BE1.
•Start number 3 was manually annotated 3 times for cluster BE1.

Gene Information:

Gene: Jay2Jay_216 Start: 107222, Stop: 107470, Start Num: 3
Candidate Starts for Jay2Jay_216:
(1, 107111), (Start: 3 @107222 has 3 MA's), (5, 107363), (6, 107399), (7, 107453), (9, 107459),

Gene: Samisti12_215 Start: 108521, Stop: 108811, Start Num: 2
Candidate Starts for Samisti12_215:
(1, 108452), (Start: 2 @108521 has 1 MA's), (Start: 3 @108563 has 3 MA's), (5, 108704), (6, 108740), (8, 108797), (9, 108800),

Gene: Tribute_207 Start: 107557, Stop: 107805, Start Num: 3
Candidate Starts for Tribute_207:
(1, 107446), (Start: 3 @107557 has 3 MA's), (4, 107626), (5, 107698), (8, 107791), (9, 107794),

Gene: Warpy_214 Start: 106660, Stop: 106908, Start Num: 3
Candidate Starts for Warpy_214:
(1, 106549), (Start: 3 @106660 has 3 MA's), (5, 106801), (6, 106837), (7, 106891), (9, 106897),