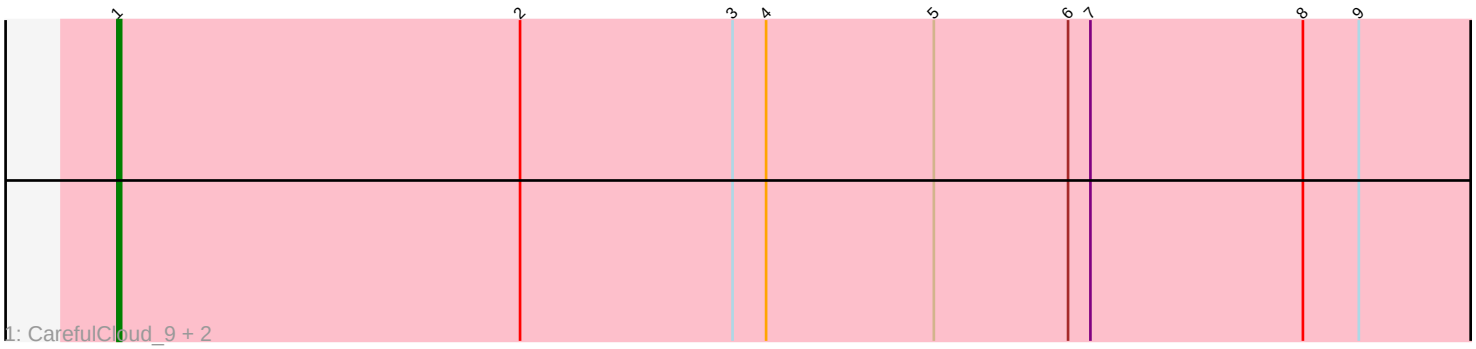
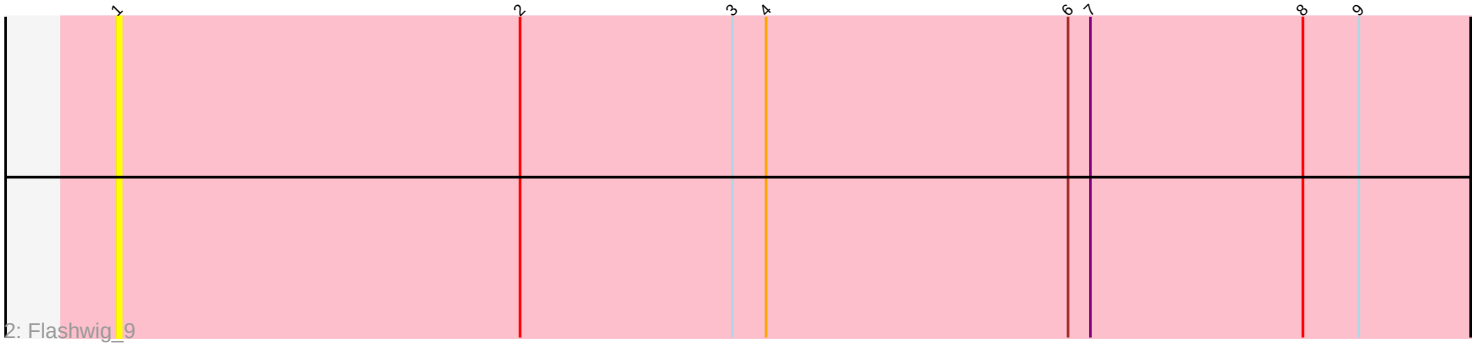
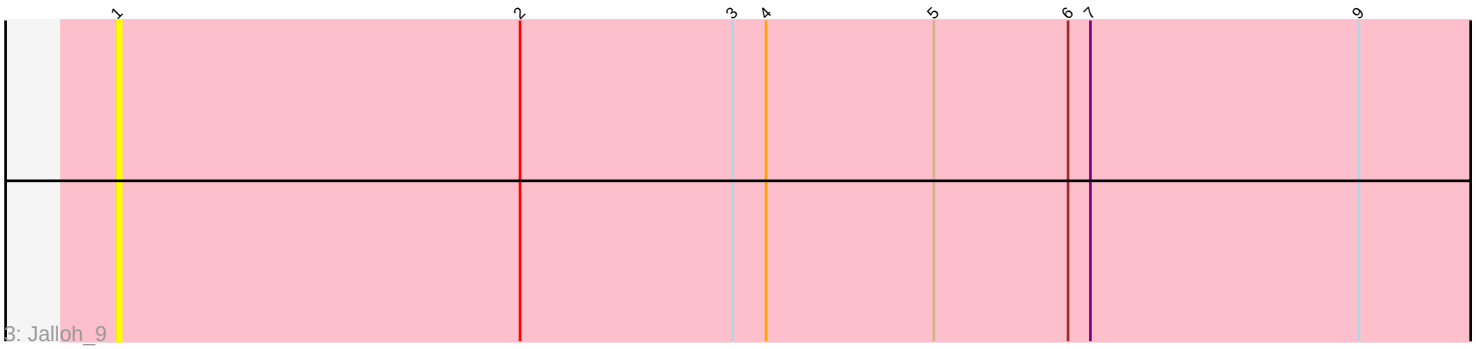
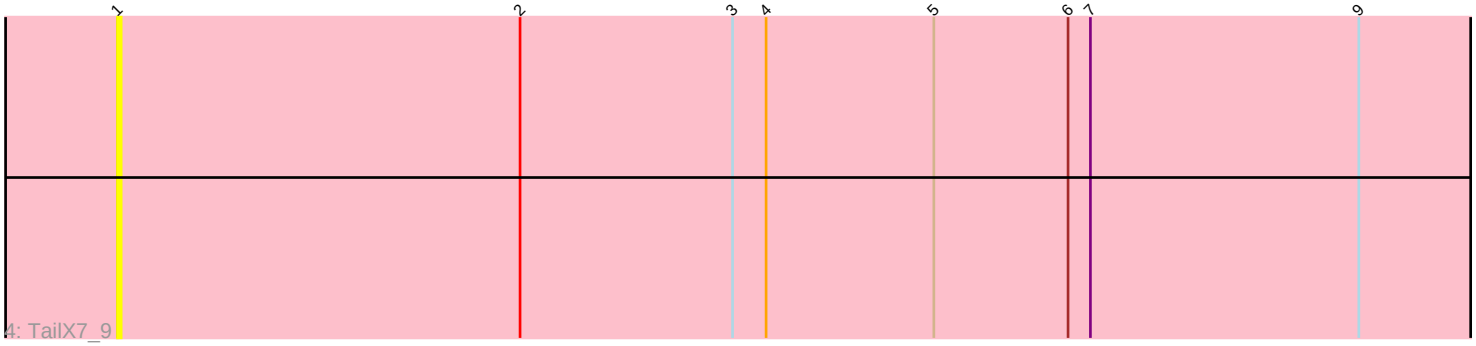
Pham 348371


1
2
3
4
5
6
7
8
9
1: CarefulCloud_9 + 2


1
2
3
4
6
7
8
9
2: Flashwig_9


1
2
3
4
5
6
7
9
3: Jalloh_9


1
2
3
4
5
6
7
9
4: TailX7_9

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 348371 Report

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

Pham number 348371 has 6 members, 5 are drafts.

Phages represented in each track:
• Track 1 : CarefulCloud_9, HorseRider_9, CyranoPS_9
• Track 2 : Flashwig_9
• Track 3 : Jalloh_9
• Track 4 : TailX7_9

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

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

Genes that call this "Most Annotated" start:
• CarefulCloud_9, CyranoPS_9, Flashwig_9, HorseRider_9, Jalloh_9, TailX7_9,

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 1:
• Found in 6 of 6 (100.0%) of genes in pham
• Manual Annotations of this start: 1 of 1
• Called 100.0% of time when present
• Phage (with cluster) where this start called: CarefulCloud_9 (ER), CyranoPS_9 (ER), Flashwig_9 (ER), HorseRider_9 (ER), Jalloh_9 (ER), TailX7_9 (ER),

Summary by clusters:

There is one cluster represented in this pham: ER

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

Gene Information:

Gene: CarefulCloud_9 Start: 5960, Stop: 6328, Start Num: 1
Candidate Starts for CarefulCloud_9:
(Start: 1 @5960 has 1 MA's), (2, 6068), (3, 6125), (4, 6134), (5, 6179), (6, 6215), (7, 6221), (8, 6278), (9, 6293),

Gene: CyranoPS_9 Start: 5958, Stop: 6326, Start Num: 1
Candidate Starts for CyranoPS_9:
(Start: 1 @5958 has 1 MA's), (2, 6066), (3, 6123), (4, 6132), (5, 6177), (6, 6213), (7, 6219), (8, 6276), (9, 6291),

Gene: Flashwig_9 Start: 5960, Stop: 6328, Start Num: 1
Candidate Starts for Flashwig_9:
(Start: 1 @5960 has 1 MA's), (2, 6068), (3, 6125), (4, 6134), (6, 6215), (7, 6221), (8, 6278), (9, 6293),

Gene: HorseRider_9 Start: 5960, Stop: 6328, Start Num: 1
Candidate Starts for HorseRider_9:
(Start: 1 @5960 has 1 MA's), (2, 6068), (3, 6125), (4, 6134), (5, 6179), (6, 6215), (7, 6221), (8, 6278), (9, 6293),

Gene: Jalloh_9 Start: 5968, Stop: 6336, Start Num: 1
Candidate Starts for Jalloh_9:
(Start: 1 @5968 has 1 MA's), (2, 6076), (3, 6133), (4, 6142), (5, 6187), (6, 6223), (7, 6229), (9, 6301),

Gene: TailX7_9 Start: 5957, Stop: 6325, Start Num: 1
Candidate Starts for TailX7_9:
(Start: 1 @5957 has 1 MA's), (2, 6065), (3, 6122), (4, 6131), (5, 6176), (6, 6212), (7, 6218), (9, 6290),