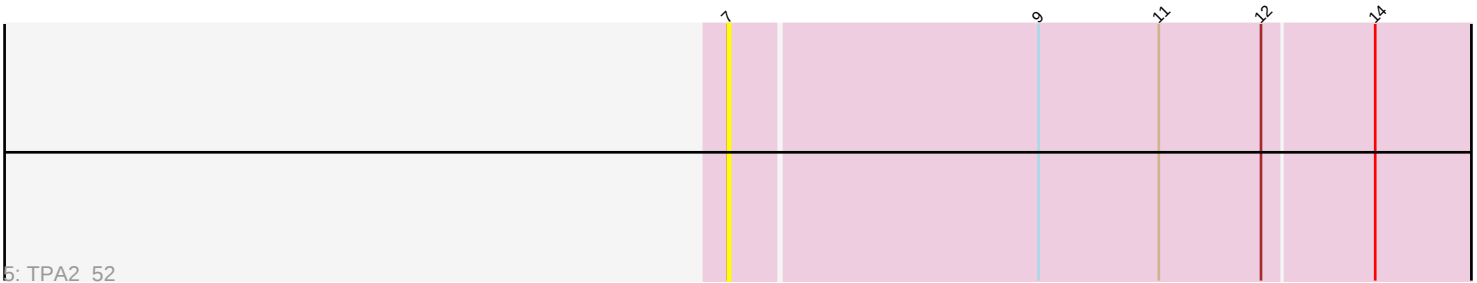
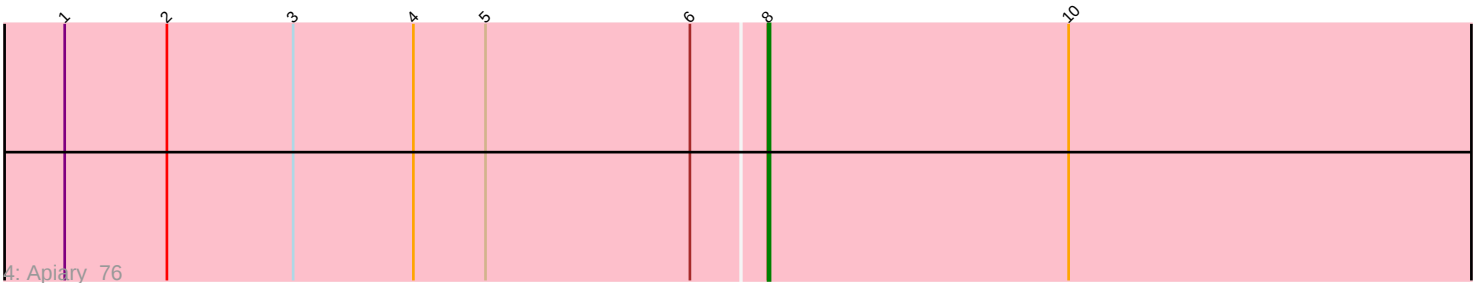
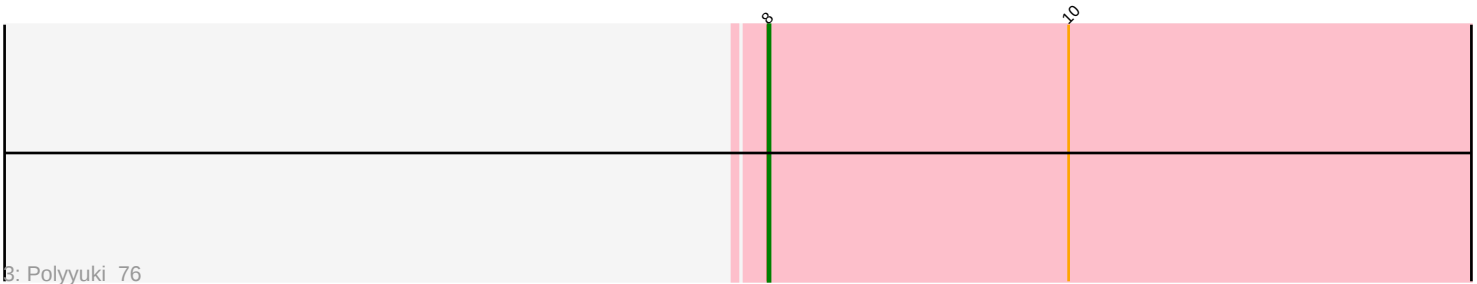
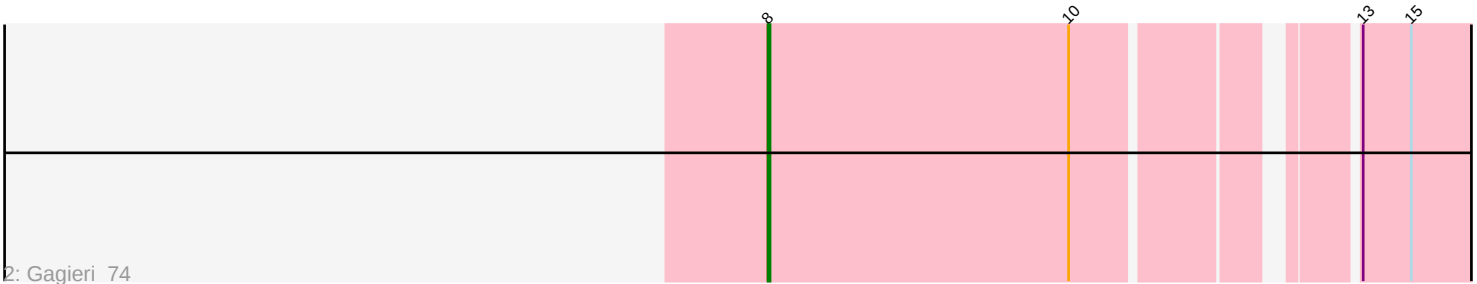
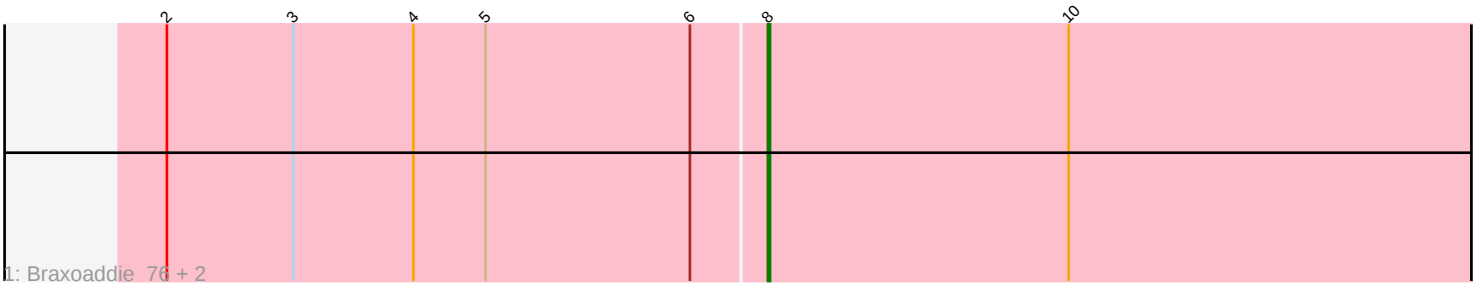


Pham 291855



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 291855 Report

This analysis was run 03/28/26 on database version 641.

Pham number 291855 has 7 members, 1 are drafts.

Phages represented in each track:

- Track 1 : Braxoaddie_76, Maselop_76, CoffeeBean_76
- Track 2 : Gagieri_74
- Track 3 : Polygyuki_76
- Track 4 : Apiary_76
- Track 5 : TPA2_52

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

The start number called the most often in the published annotations is 8, it was called in 6 of the 6 non-draft genes in the pham.

Genes that call this "Most Annotated" start:

- Apiary_76, Braxoaddie_76, CoffeeBean_76, Gagieri_74, Maselop_76, Polygyuki_76,

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

-

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

- TPA2_52,

Summary by start number:

Start 7:

- Found in 1 of 7 (14.3%) of genes in pham
- No Manual Annotations of this start.
- Called 100.0% of time when present
- Phage (with cluster) where this start called: TPA2_52 (singleton),

Start 8:

- Found in 6 of 7 (85.7%) 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: Apiary_76 (CR6), Braxoaddie_76 (CR6), CoffeeBean_76 (CR6), Gagieri_74 (CR6), Maselop_76 (CR6), Polygyuki_76 (CR6),

Summary by clusters:

There are 2 clusters represented in this pham: singleton, CR6,

Info for manual annotations of cluster CR6:

•Start number 8 was manually annotated 6 times for cluster CR6.

Gene Information:

Gene: Apiary_76 Start: 55270, Stop: 54890, Start Num: 8

Candidate Starts for Apiary_76:

(1, 55618), (2, 55567), (3, 55504), (4, 55444), (5, 55408), (6, 55306), (Start: 8 @55270 has 6 MA's), (10, 55120),

Gene: Braxoaddie_76 Start: 55255, Stop: 54875, Start Num: 8

Candidate Starts for Braxoaddie_76:

(2, 55552), (3, 55489), (4, 55429), (5, 55393), (6, 55291), (Start: 8 @55255 has 6 MA's), (10, 55105),

Gene: CoffeeBean_76 Start: 55213, Stop: 54833, Start Num: 8

Candidate Starts for CoffeeBean_76:

(2, 55510), (3, 55447), (4, 55387), (5, 55351), (6, 55249), (Start: 8 @55213 has 6 MA's), (10, 55063),

Gene: Gagieri_74 Start: 55043, Stop: 54696, Start Num: 8

Candidate Starts for Gagieri_74:

(Start: 8 @55043 has 6 MA's), (10, 54893), (13, 54776), (15, 54752),

Gene: Maselop_76 Start: 55289, Stop: 54909, Start Num: 8

Candidate Starts for Maselop_76:

(2, 55586), (3, 55523), (4, 55463), (5, 55427), (6, 55325), (Start: 8 @55289 has 6 MA's), (10, 55139),

Gene: Polyuyki_76 Start: 55281, Stop: 54901, Start Num: 8

Candidate Starts for Polyuyki_76:

(Start: 8 @55281 has 6 MA's), (10, 55131),

Gene: TPA2_52 Start: 39328, Stop: 38936, Start Num: 7

Candidate Starts for TPA2_52:

(7, 39328), (9, 39178), (11, 39118), (12, 39067), (14, 39013),