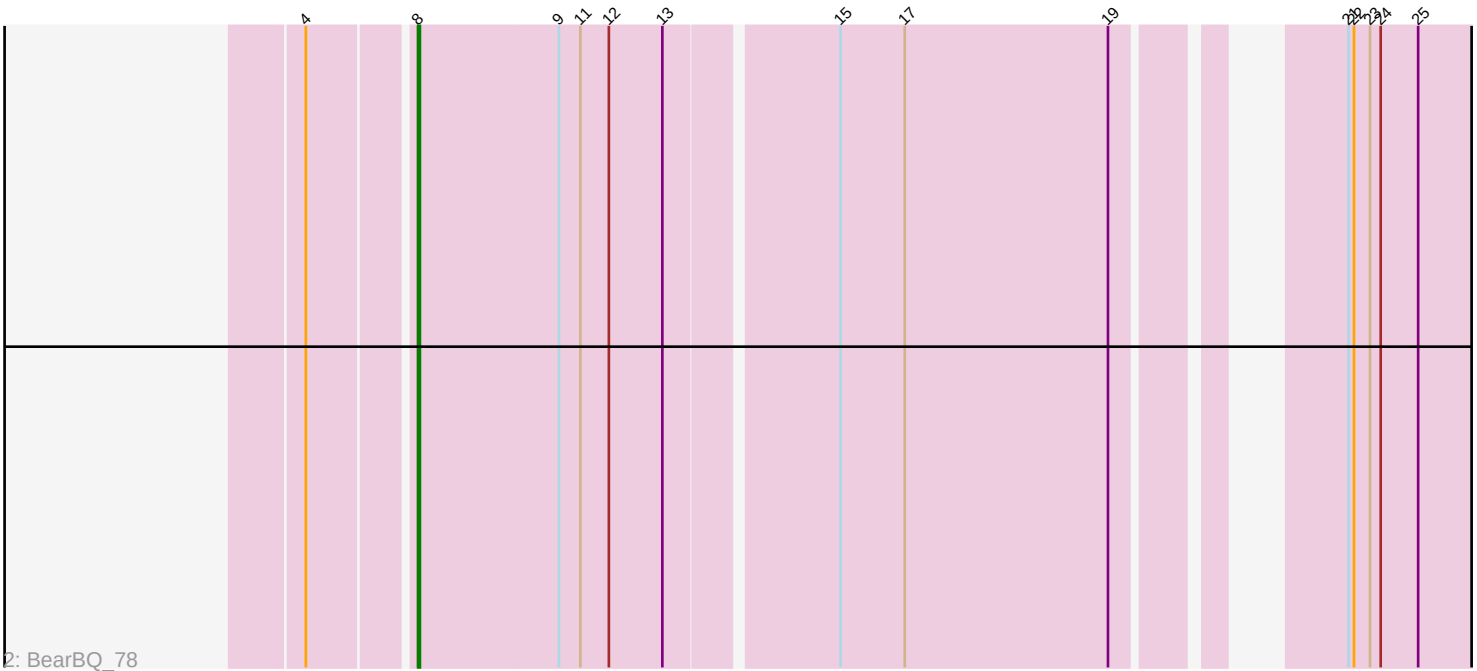
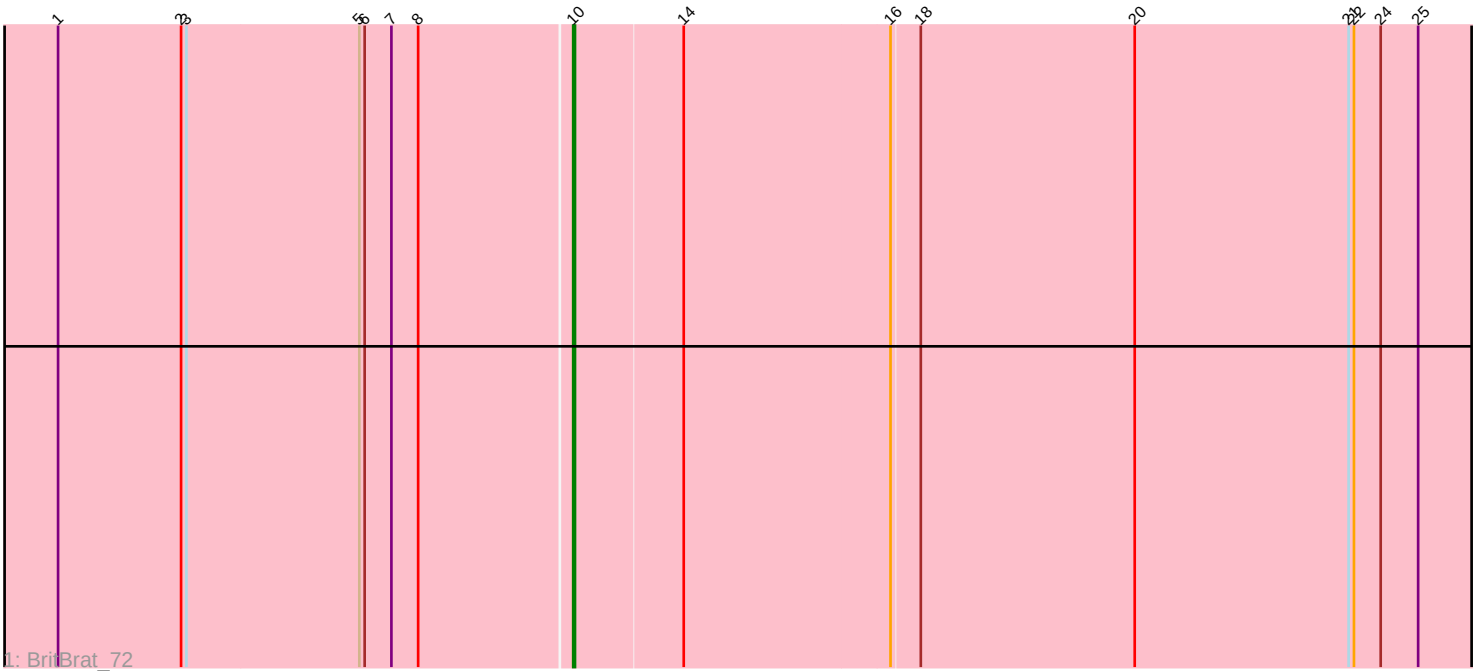




Note: Tracks are now grouped by subcluster and scaled. Switching in subcluster is indicated by changes in track color. Track scale is now set by default to display the region 30 bp upstream of start 1 to 30 bp downstream of the last possible start. If this default region is judged to be packed too tightly 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 292088 Report

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

Pham number 292088 has 2 members, 0 are drafts.

Phages represented in each track:

- Track 1 : BritBrat_72
- Track 2 : BearBQ_78

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 1 of the 2 non-draft genes in the pham.

Genes that call this "Most Annotated" start:

- BritBrat_72,

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

-

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

- BearBQ_78,

Summary by start number:

Start 8:

- Found in 2 of 2 (100.0%) of genes in pham
- Manual Annotations of this start: 1 of 2
- Called 50.0% of time when present
- Phage (with cluster) where this start called: BearBQ_78 (DN),

Start 10:

- Found in 1 of 2 (50.0%) of genes in pham
- Manual Annotations of this start: 1 of 2
- Called 100.0% of time when present
- Phage (with cluster) where this start called: BritBrat_72 (CY2),

Summary by clusters:

There are 2 clusters represented in this pham: DN, CY2,

Info for manual annotations of cluster CY2:

- Start number 10 was manually annotated 1 time for cluster CY2.

Info for manual annotations of cluster DN:

- Start number 8 was manually annotated 1 time for cluster DN.

Gene Information:

Gene: BearBQ_78 Start: 46707, Stop: 47315, Start Num: 8

Candidate Starts for BearBQ_78:

(4, 46653), (Start: 8 @46707 has 1 MA's), (9, 46785), (11, 46797), (12, 46812), (13, 46842), (15, 46932), (17, 46968), (19, 47082), (21, 47169), (22, 47172), (23, 47181), (24, 47187), (25, 47208),

Gene: BritBrat_72 Start: 45785, Stop: 46360, Start Num: 10

Candidate Starts for BritBrat_72:

(1, 45500), (2, 45569), (3, 45572), (5, 45668), (6, 45671), (7, 45686), (Start: 8 @45701 has 1 MA's), (Start: 10 @45785 has 1 MA's), (14, 45845), (16, 45959), (18, 45974), (20, 46094), (21, 46214), (22, 46217), (24, 46232), (25, 46253),