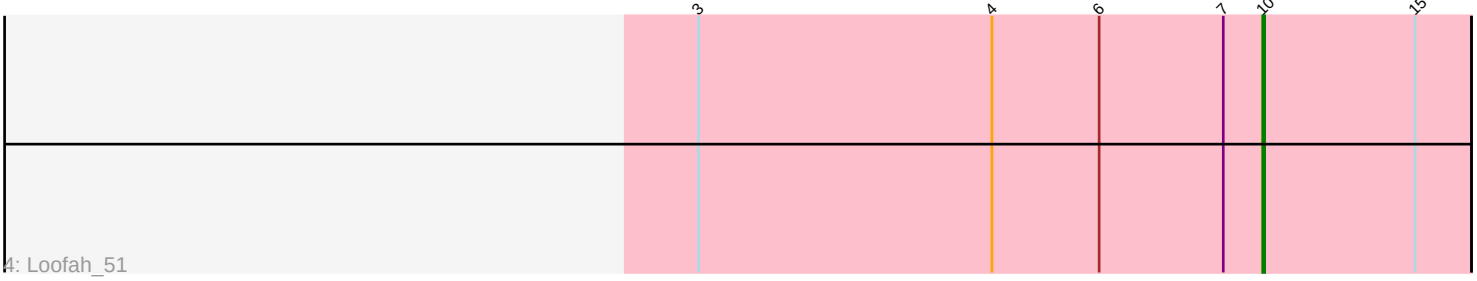
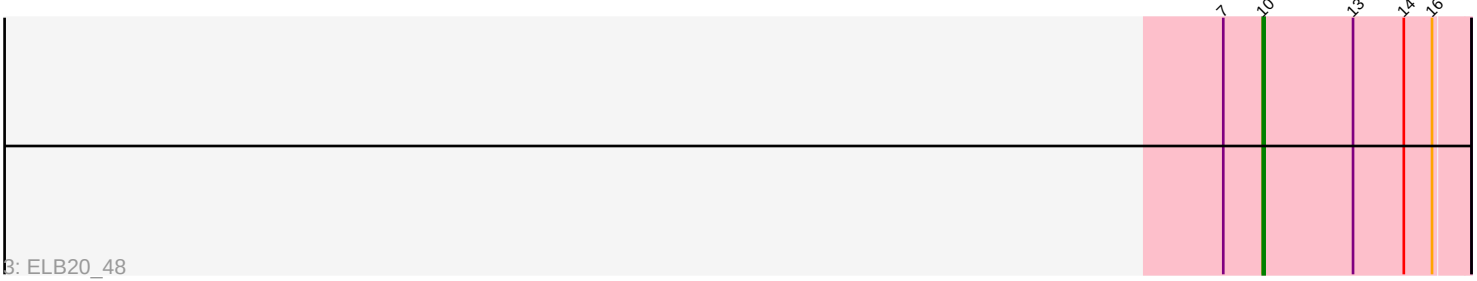
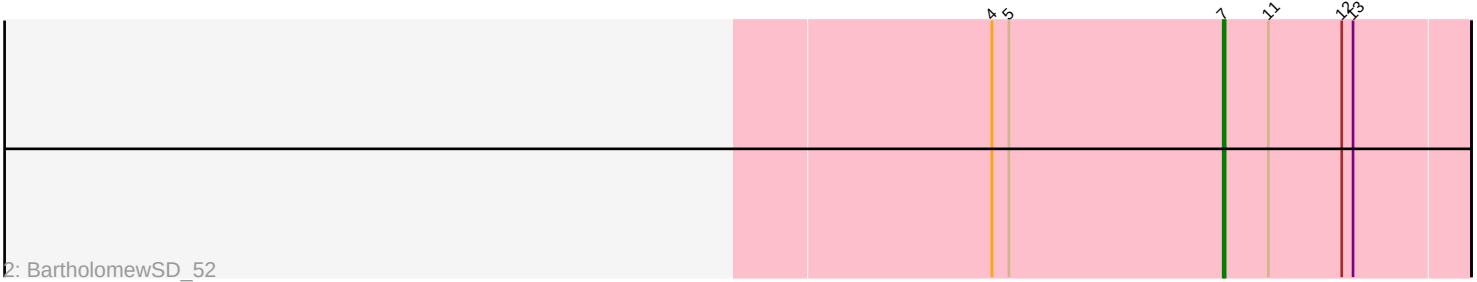


Pham 310052



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 310052 Report

This analysis was run 09/05/26 on database version 662.

Pham number 310052 has 7 members, 0 are drafts.

Phages represented in each track:

- Track 1 : Paedore_50
- Track 2 : BartholomewSD_52
- Track 3 : ELB20_48
- Track 4 : Loofah_51
- Track 5 : GreenWeasel_49, phiHau3_49, BroPlease_48

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 3 of the 7 non-draft genes in the pham.

Genes that call this "Most Annotated" start:

- BroPlease_48, GreenWeasel_49, phiHau3_49,

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

-

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

- BartholomewSD_52, ELB20_48, Loofah_51, Paedore_50,

Summary by start number:

Start 7:

- Found in 4 of 7 (57.1%) of genes in pham
- Manual Annotations of this start: 1 of 7
- Called 25.0% of time when present
- Phage (with cluster) where this start called: BartholomewSD_52 (BD2),

Start 8:

- Found in 3 of 7 (42.9%) of genes in pham
- Manual Annotations of this start: 3 of 7
- Called 100.0% of time when present
- Phage (with cluster) where this start called: BroPlease_48 (BD4), GreenWeasel_49 (BD4), phiHau3_49 (BD4),

Start 10:

- Found in 3 of 7 (42.9%) of genes in pham
- Manual Annotations of this start: 3 of 7
- Called 100.0% of time when present
- Phage (with cluster) where this start called: ELB20_48 (BD2), Loofah_51 (BD2), Paedore_50 (BD2),

Summary by clusters:

There are 2 clusters represented in this pham: BD4, BD2,

Info for manual annotations of cluster BD2:

- Start number 7 was manually annotated 1 time for cluster BD2.
- Start number 10 was manually annotated 3 times for cluster BD2.

Info for manual annotations of cluster BD4:

- Start number 8 was manually annotated 3 times for cluster BD4.

Gene Information:

Gene: BartholomewSD_52 Start: 35855, Stop: 35727, Start Num: 7

Candidate Starts for BartholomewSD_52:

(4, 35978), (5, 35969), (Start: 7 @35855 has 1 MA's), (11, 35831), (12, 35792), (13, 35786),

Gene: BroPlease_48 Start: 34002, Stop: 33886, Start Num: 8

Candidate Starts for BroPlease_48:

(Start: 8 @34002 has 3 MA's), (9, 33999),

Gene: ELB20_48 Start: 35704, Stop: 35597, Start Num: 10

Candidate Starts for ELB20_48:

(Start: 7 @35725 has 1 MA's), (Start: 10 @35704 has 3 MA's), (13, 35656), (14, 35629), (16, 35614),

Gene: GreenWeasel_49 Start: 34007, Stop: 33891, Start Num: 8

Candidate Starts for GreenWeasel_49:

(Start: 8 @34007 has 3 MA's), (9, 34004),

Gene: Loofah_51 Start: 35851, Stop: 35741, Start Num: 10

Candidate Starts for Loofah_51:

(3, 36151), (4, 35995), (6, 35938), (Start: 7 @35872 has 1 MA's), (Start: 10 @35851 has 3 MA's), (15, 35770),

Gene: Paedore_50 Start: 35836, Stop: 35726, Start Num: 10

Candidate Starts for Paedore_50:

(1, 36475), (2, 36304), (3, 36136), (4, 35980), (6, 35923), (Start: 7 @35857 has 1 MA's), (Start: 10 @35836 has 3 MA's), (15, 35755),

Gene: phiHau3_49 Start: 33970, Stop: 33854, Start Num: 8

Candidate Starts for phiHau3_49:

(Start: 8 @33970 has 3 MA's), (9, 33967),