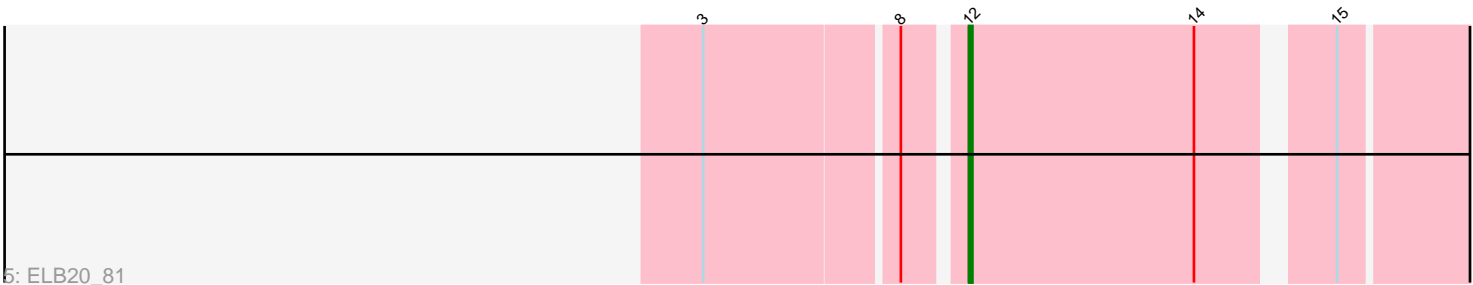
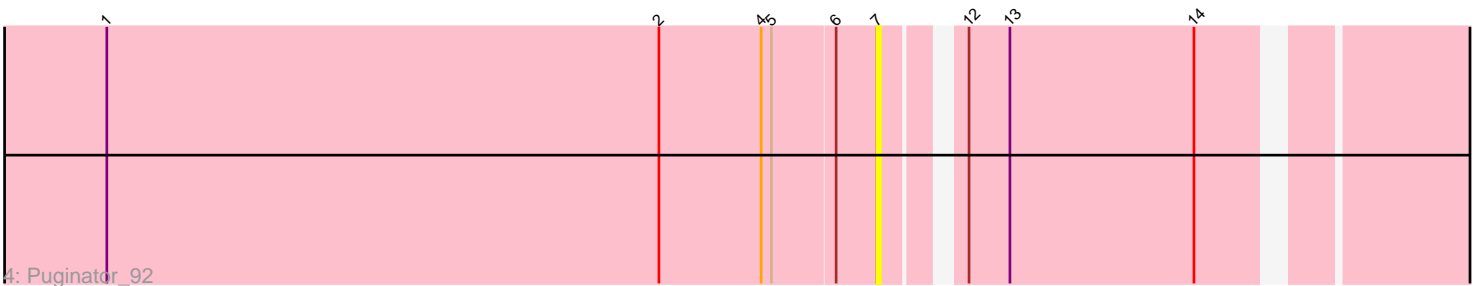
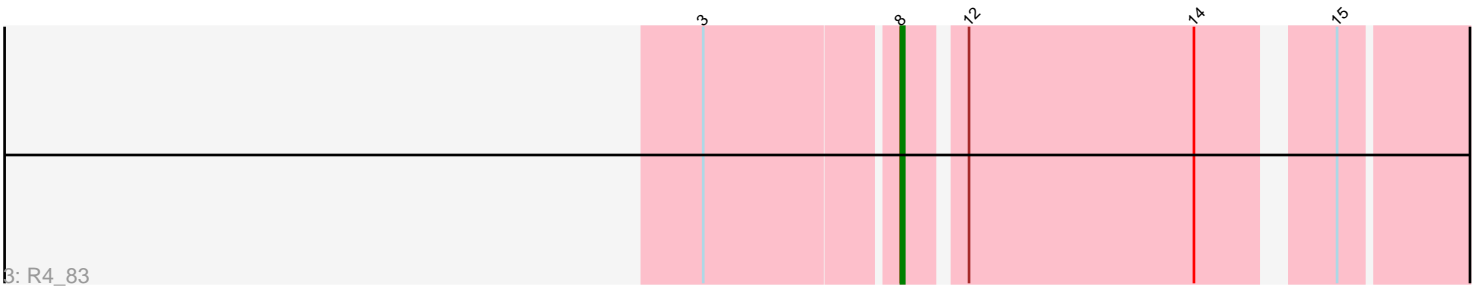
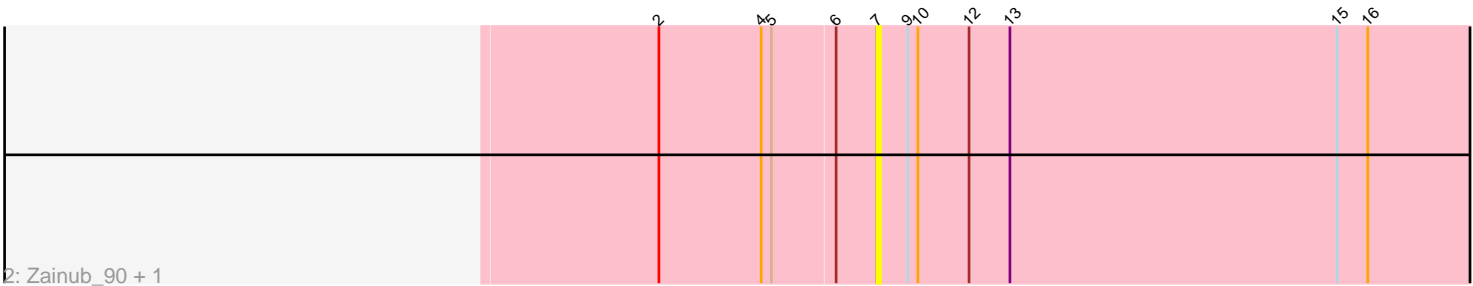
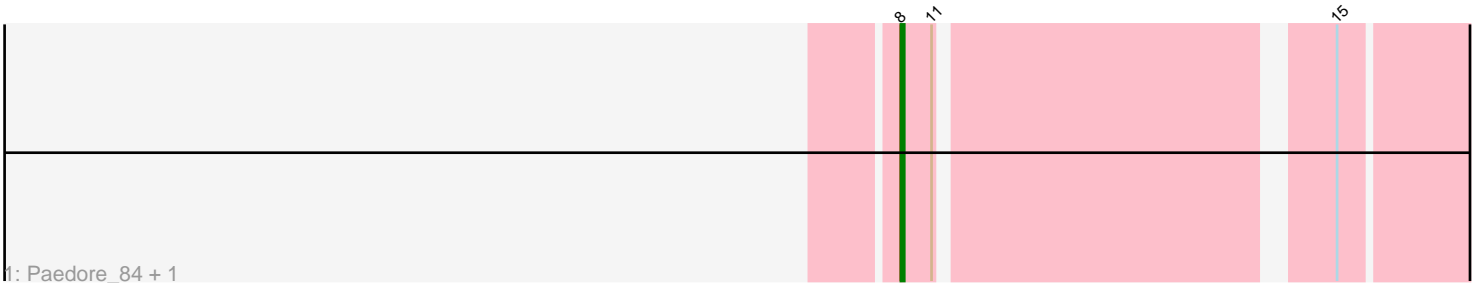


Pham 107356



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 107356 Report

This analysis was run 04/28/24 on database version 559.

Pham number 107356 has 7 members, 4 are drafts.

Phages represented in each track:

- Track 1 : Paedore\_84, Loofah\_87
- Track 2 : Zainub\_90, Paolo\_86
- Track 3 : R4\_83
- Track 4 : Puginator\_92
- Track 5 : ELB20\_81

### ***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 2 of the 3 non-draft genes in the pham.

Genes that call this "Most Annotated" start:

- Loofah\_87, Paedore\_84, R4\_83,

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

- ELB20\_81,

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

- Paolo\_86, Puginator\_92, Zainub\_90,

### **Summary by start number:**

Start 7:

- Found in 3 of 7 ( 42.9% ) of genes in pham
- No Manual Annotations of this start.
- Called 100.0% of time when present
- Phage (with cluster) where this start called: Paolo\_86 (BD2), Puginator\_92 (BD2), Zainub\_90 (BD2),

Start 8:

- Found in 4 of 7 ( 57.1% ) of genes in pham
- Manual Annotations of this start: 2 of 3
- Called 75.0% of time when present

- Phage (with cluster) where this start called: Loofah\_87 (BD2), Paedore\_84 (BD2), R4\_83 (BD2),

Start 12:

- Found in 5 of 7 ( 71.4% ) of genes in pham
- Manual Annotations of this start: 1 of 3
- Called 20.0% of time when present
- Phage (with cluster) where this start called: ELB20\_81 (BD2),

### **Summary by clusters:**

There is one cluster represented in this pham: BD2

Info for manual annotations of cluster BD2:

- Start number 8 was manually annotated 2 times for cluster BD2.
- Start number 12 was manually annotated 1 time for cluster BD2.

### **Gene Information:**

Gene: ELB20\_81 Start: 50160, Stop: 49957, Start Num: 12

Candidate Starts for ELB20\_81:

(3, 50229), (Start: 8 @50175 has 2 MA's), (Start: 12 @50160 has 1 MA's), (14, 50094), (15, 50061),

Gene: Loofah\_87 Start: 51412, Stop: 51194, Start Num: 8

Candidate Starts for Loofah\_87:

(Start: 8 @51412 has 2 MA's), (11, 51403), (15, 51298),

Gene: Paedore\_84 Start: 51581, Stop: 51363, Start Num: 8

Candidate Starts for Paedore\_84:

(Start: 8 @51581 has 2 MA's), (11, 51572), (15, 51467),

Gene: Paolo\_86 Start: 50004, Stop: 49774, Start Num: 7

Candidate Starts for Paolo\_86:

(2, 50067), (4, 50037), (5, 50034), (6, 50016), (7, 50004), (9, 49995), (10, 49992), (Start: 12 @49977 has 1 MA's), (13, 49965), (15, 49869), (16, 49860),

Gene: Puginator\_92 Start: 50053, Stop: 49844, Start Num: 7

Candidate Starts for Puginator\_92:

(1, 50278), (2, 50116), (4, 50086), (5, 50083), (6, 50065), (7, 50053), (Start: 12 @50035 has 1 MA's), (13, 50023), (14, 49969),

Gene: R4\_83 Start: 50097, Stop: 49879, Start Num: 8

Candidate Starts for R4\_83:

(3, 50151), (Start: 8 @50097 has 2 MA's), (Start: 12 @50082 has 1 MA's), (14, 50016), (15, 49983),

Gene: Zainub\_90 Start: 50184, Stop: 49954, Start Num: 7

Candidate Starts for Zainub\_90:

(2, 50247), (4, 50217), (5, 50214), (6, 50196), (7, 50184), (9, 50175), (10, 50172), (Start: 12 @50157 has 1 MA's), (13, 50145), (15, 50049), (16, 50040),