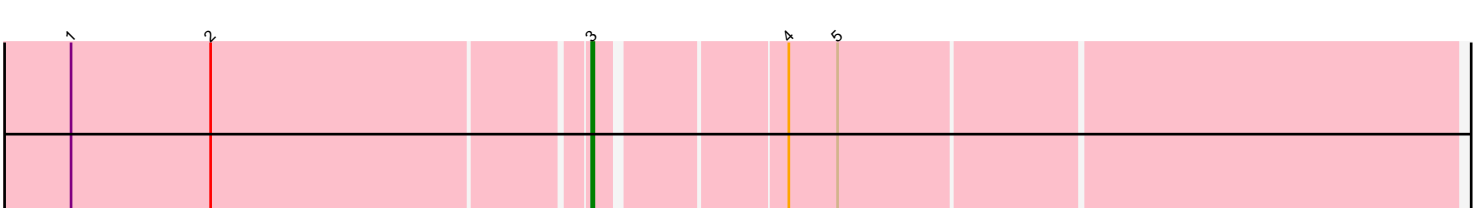
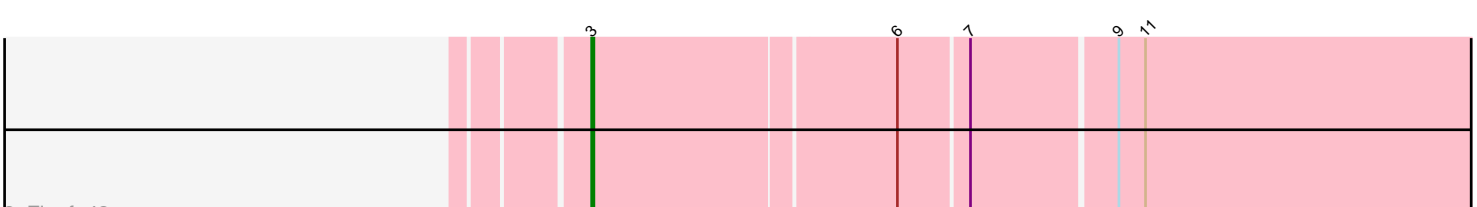
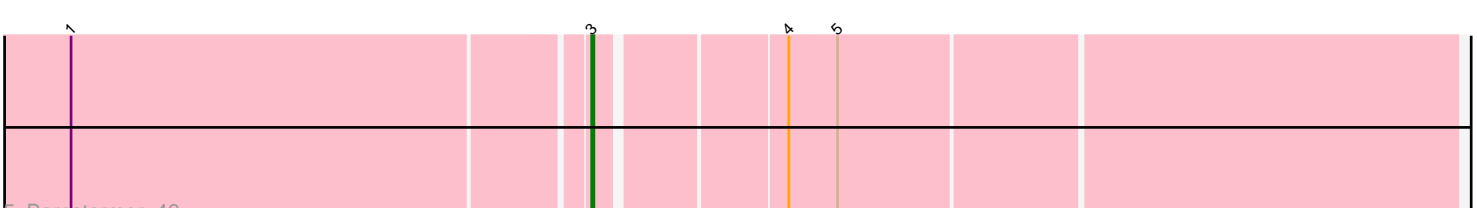
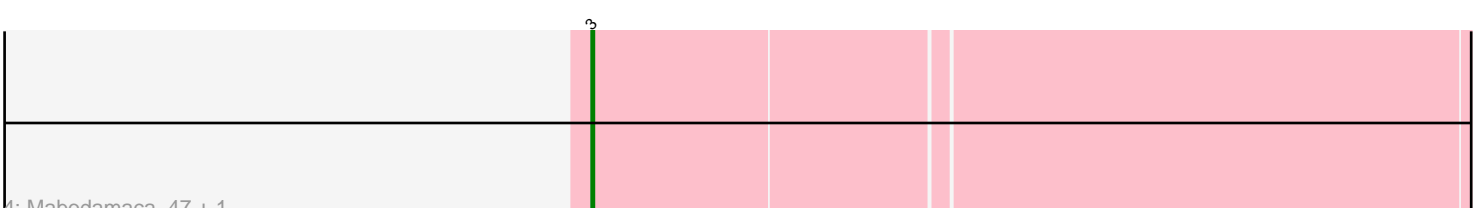
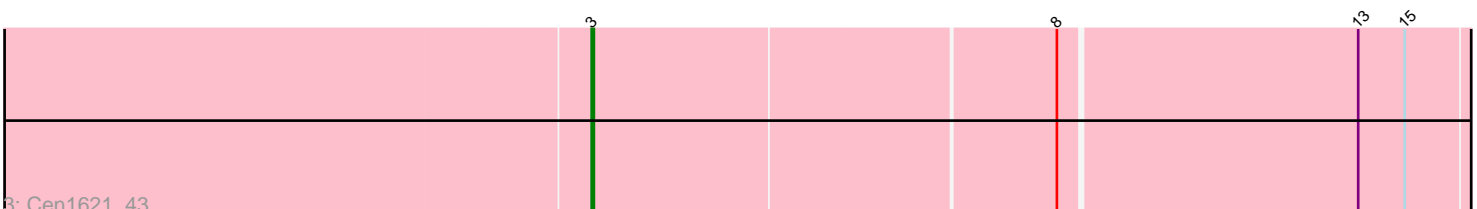
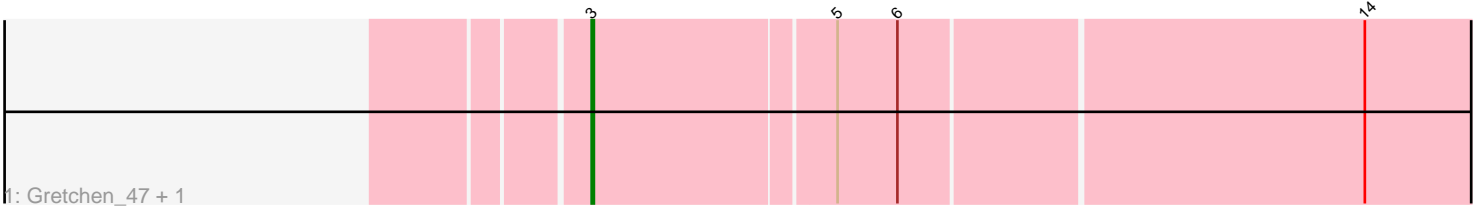


Pham 7405



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 7405 Report

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

Pham number 7405 has 10 members, 1 are drafts.

Phages represented in each track:

- Track 1 : Gretchen\_47, Percival\_48
- Track 2 : GardenState\_47, IAmGroot\_47
- Track 3 : Cen1621\_43
- Track 4 : Mabodamaca\_47, SuMoo\_49
- Track 5 : Barnstormer\_46
- Track 6 : Floof\_48
- Track 7 : UtzChips\_46

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

The start number called the most often in the published annotations is 3, it was called in 9 of the 9 non-draft genes in the pham.

Genes that call this "Most Annotated" start:

- Barnstormer\_46, Cen1621\_43, Floof\_48, GardenState\_47, Gretchen\_47, IAmGroot\_47, Mabodamaca\_47, Percival\_48, SuMoo\_49, UtzChips\_46,

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 3:

- Found in 10 of 10 ( 100.0% ) of genes in pham
- Manual Annotations of this start: 9 of 9
- Called 100.0% of time when present
- Phage (with cluster) where this start called: Barnstormer\_46 (EH), Cen1621\_43 (EH), Floof\_48 (EH), GardenState\_47 (EH), Gretchen\_47 (EH), IAmGroot\_47 (EH), Mabodamaca\_47 (EH), Percival\_48 (EH), SuMoo\_49 (EH), UtzChips\_46 (EH),

### **Summary by clusters:**

There is one cluster represented in this pham: EH

Info for manual annotations of cluster EH:

- Start number 3 was manually annotated 9 times for cluster EH.

**Gene Information:**

Gene: Barnstormer\_46 Start: 30765, Stop: 31136, Start Num: 3

Candidate Starts for Barnstormer\_46:

(1, 30540), (Start: 3 @30765 has 9 MA's), (4, 30843), (5, 30864),

Gene: Cen1621\_43 Start: 29290, Stop: 29697, Start Num: 3

Candidate Starts for Cen1621\_43:

(Start: 3 @29290 has 9 MA's), (8, 29494), (13, 29626), (15, 29647),

Gene: Floof\_48 Start: 31609, Stop: 32013, Start Num: 3

Candidate Starts for Floof\_48:

(Start: 3 @31609 has 9 MA's), (6, 31741), (7, 31771), (9, 31834), (11, 31846),

Gene: GardenState\_47 Start: 31002, Stop: 31415, Start Num: 3

Candidate Starts for GardenState\_47:

(Start: 3 @31002 has 9 MA's), (10, 31239), (12, 31335),

Gene: Gretchen\_47 Start: 32225, Stop: 32629, Start Num: 3

Candidate Starts for Gretchen\_47:

(Start: 3 @32225 has 9 MA's), (5, 32330), (6, 32357), (14, 32561),

Gene: IAmGroot\_47 Start: 31469, Stop: 31882, Start Num: 3

Candidate Starts for IAmGroot\_47:

(Start: 3 @31469 has 9 MA's), (10, 31706), (12, 31802),

Gene: Mabodamaca\_47 Start: 32040, Stop: 32444, Start Num: 3

Candidate Starts for Mabodamaca\_47:

(Start: 3 @32040 has 9 MA's),

Gene: Percival\_48 Start: 32060, Stop: 32464, Start Num: 3

Candidate Starts for Percival\_48:

(Start: 3 @32060 has 9 MA's), (5, 32165), (6, 32192), (14, 32396),

Gene: SuMoo\_49 Start: 31703, Stop: 32107, Start Num: 3

Candidate Starts for SuMoo\_49:

(Start: 3 @31703 has 9 MA's),

Gene: UtzChips\_46 Start: 30751, Stop: 31122, Start Num: 3

Candidate Starts for UtzChips\_46:

(1, 30526), (2, 30589), (Start: 3 @30751 has 9 MA's), (4, 30829), (5, 30850),