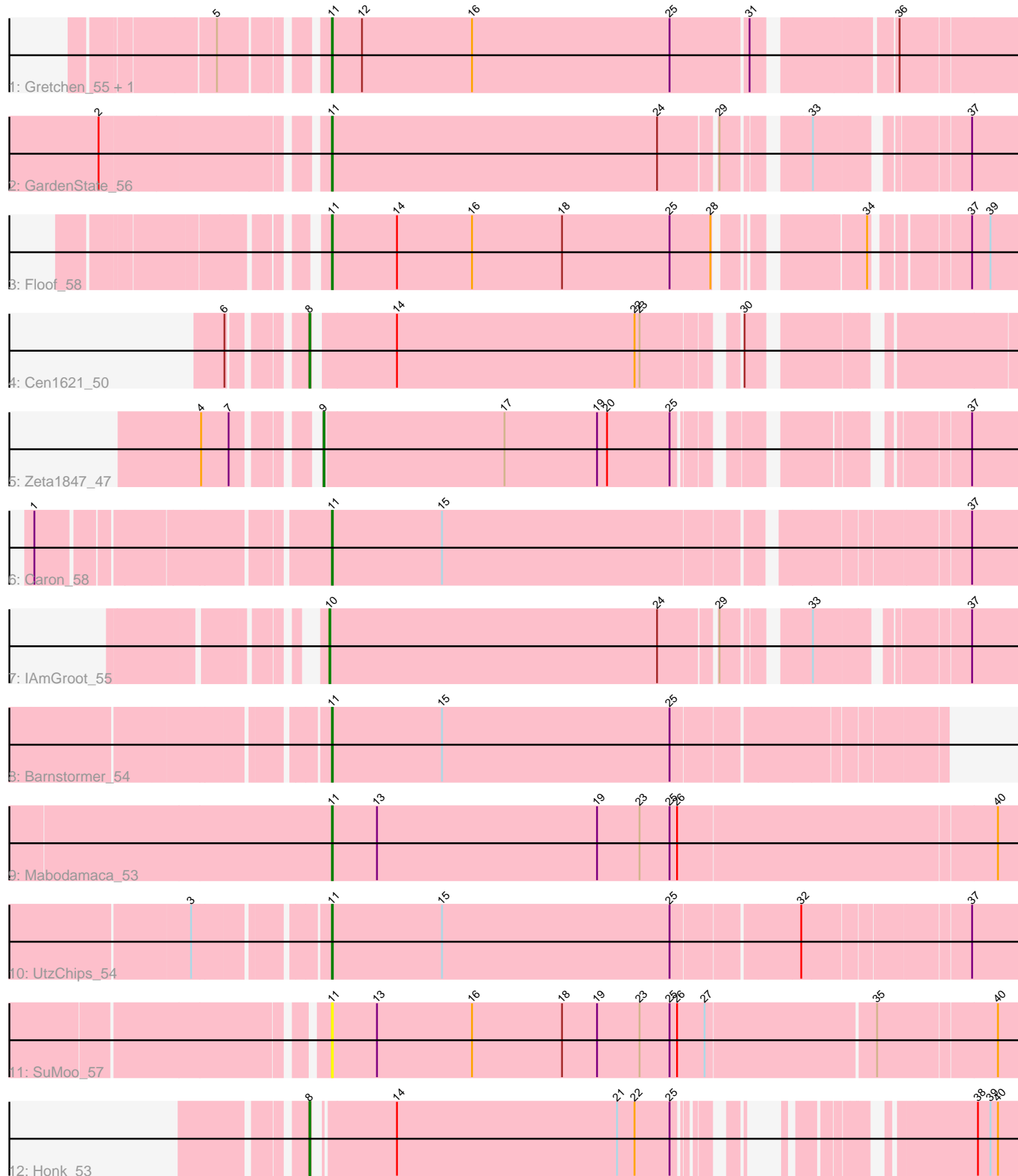


# Pham 32430



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 32430 Report

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

Pham number 32430 has 13 members, 1 are drafts.

Phages represented in each track:

- Track 1 : Gretchen\_55, Percival\_56
- Track 2 : GardenState\_56
- Track 3 : Floof\_58
- Track 4 : Cen1621\_50
- Track 5 : Zeta1847\_47
- Track 6 : Caron\_58
- Track 7 : IAmGroot\_55
- Track 8 : Barnstormer\_54
- Track 9 : Mabodamaca\_53
- Track 10 : UtzChips\_54
- Track 11 : SuMoo\_57
- Track 12 : Honk\_53

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

The start number called the most often in the published annotations is 11, it was called in 8 of the 12 non-draft genes in the pham.

Genes that call this "Most Annotated" start:

- Barnstormer\_54, Caron\_58, Floof\_58, GardenState\_56, Gretchen\_55, Mabodamaca\_53, Percival\_56, SuMoo\_57, UtzChips\_54,

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

- 

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

- Cen1621\_50, Honk\_53, IAmGroot\_55, Zeta1847\_47,

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

Start 8:

- Found in 2 of 13 ( 15.4% ) of genes in pham
- Manual Annotations of this start: 2 of 12
- Called 100.0% of time when present

- Phage (with cluster) where this start called: Cen1621\_50 (EH), Honk\_53 (EH),

Start 9:

- Found in 1 of 13 ( 7.7% ) of genes in pham
- Manual Annotations of this start: 1 of 12
- Called 100.0% of time when present
- Phage (with cluster) where this start called: Zeta1847\_47 (EH),

Start 10:

- Found in 1 of 13 ( 7.7% ) of genes in pham
- Manual Annotations of this start: 1 of 12
- Called 100.0% of time when present
- Phage (with cluster) where this start called: IAmGroot\_55 (EH),

Start 11:

- Found in 9 of 13 ( 69.2% ) of genes in pham
- Manual Annotations of this start: 8 of 12
- Called 100.0% of time when present
- Phage (with cluster) where this start called: Barnstormer\_54 (EH), Caron\_58 (EH), Floof\_58 (EH), GardenState\_56 (EH), Gretchen\_55 (EH), Mabodamaca\_53 (EH), Percival\_56 (EH), SuMoo\_57 (EH), UtzChips\_54 (EH),

### **Summary by clusters:**

There is one cluster represented in this pham: EH

Info for manual annotations of cluster EH:

- Start number 8 was manually annotated 2 times for cluster EH.
- Start number 9 was manually annotated 1 time for cluster EH.
- Start number 10 was manually annotated 1 time for cluster EH.
- Start number 11 was manually annotated 8 times for cluster EH.

### **Gene Information:**

Gene: Barnstormer\_54 Start: 35694, Stop: 36401, Start Num: 11

Candidate Starts for Barnstormer\_54:

(Start: 11 @35694 has 8 MA's), (15, 35826), (25, 36099),

Gene: Caron\_58 Start: 36781, Stop: 37575, Start Num: 11

Candidate Starts for Caron\_58:

(1, 36472), (Start: 11 @36781 has 8 MA's), (15, 36913), (37, 37498),

Gene: Cen1621\_50 Start: 34557, Stop: 35321, Start Num: 8

Candidate Starts for Cen1621\_50:

(6, 34485), (Start: 8 @34557 has 2 MA's), (14, 34647), (22, 34932), (23, 34938), (30, 35034),

Gene: Floof\_58 Start: 37269, Stop: 38048, Start Num: 11

Candidate Starts for Floof\_58:

(Start: 11 @37269 has 8 MA's), (14, 37347), (16, 37437), (18, 37545), (25, 37674), (28, 37722), (34, 37857), (37, 37956), (39, 37977),

Gene: GardenState\_56 Start: 36236, Stop: 36994, Start Num: 11

Candidate Starts for GardenState\_56:

(2, 35990), (Start: 11 @36236 has 8 MA's), (24, 36626), (29, 36689), (33, 36770), (37, 36929),

Gene: Gretchen\_55 Start: 37328, Stop: 38125, Start Num: 11

Candidate Starts for Gretchen\_55:

(5, 37223), (Start: 11 @37328 has 8 MA's), (12, 37364), (16, 37496), (25, 37733), (31, 37823), (36, 37967),

Gene: Honk\_53 Start: 34819, Stop: 35538, Start Num: 8

Candidate Starts for Honk\_53:

(Start: 8 @34819 has 2 MA's), (14, 34903), (21, 35167), (22, 35188), (25, 35230), (38, 35458), (39, 35473), (40, 35482),

Gene: IAmGroot\_55 Start: 36441, Stop: 37202, Start Num: 10

Candidate Starts for IAmGroot\_55:

(Start: 10 @36441 has 1 MA's), (24, 36834), (29, 36897), (33, 36978), (37, 37137),

Gene: Mabodamaca\_53 Start: 37633, Stop: 38448, Start Num: 11

Candidate Starts for Mabodamaca\_53:

(Start: 11 @37633 has 8 MA's), (13, 37687), (19, 37951), (23, 38002), (25, 38038), (26, 38047), (40, 38422),

Gene: Percival\_56 Start: 37164, Stop: 37961, Start Num: 11

Candidate Starts for Percival\_56:

(5, 37059), (Start: 11 @37164 has 8 MA's), (12, 37200), (16, 37332), (25, 37569), (31, 37659), (36, 37803),

Gene: SuMoo\_57 Start: 37124, Stop: 37933, Start Num: 11

Candidate Starts for SuMoo\_57:

(Start: 11 @37124 has 8 MA's), (13, 37178), (16, 37292), (18, 37400), (19, 37442), (23, 37493), (25, 37529), (26, 37538), (27, 37571), (35, 37769), (40, 37907),

Gene: UtzChips\_54 Start: 35679, Stop: 36491, Start Num: 11

Candidate Starts for UtzChips\_54:

(3, 35541), (Start: 11 @35679 has 8 MA's), (15, 35811), (25, 36084), (32, 36228), (37, 36414),

Gene: Zeta1847\_47 Start: 34179, Stop: 34964, Start Num: 9

Candidate Starts for Zeta1847\_47:

(4, 34071), (7, 34104), (Start: 9 @34179 has 1 MA's), (17, 34392), (19, 34503), (20, 34515), (25, 34590), (37, 34854),