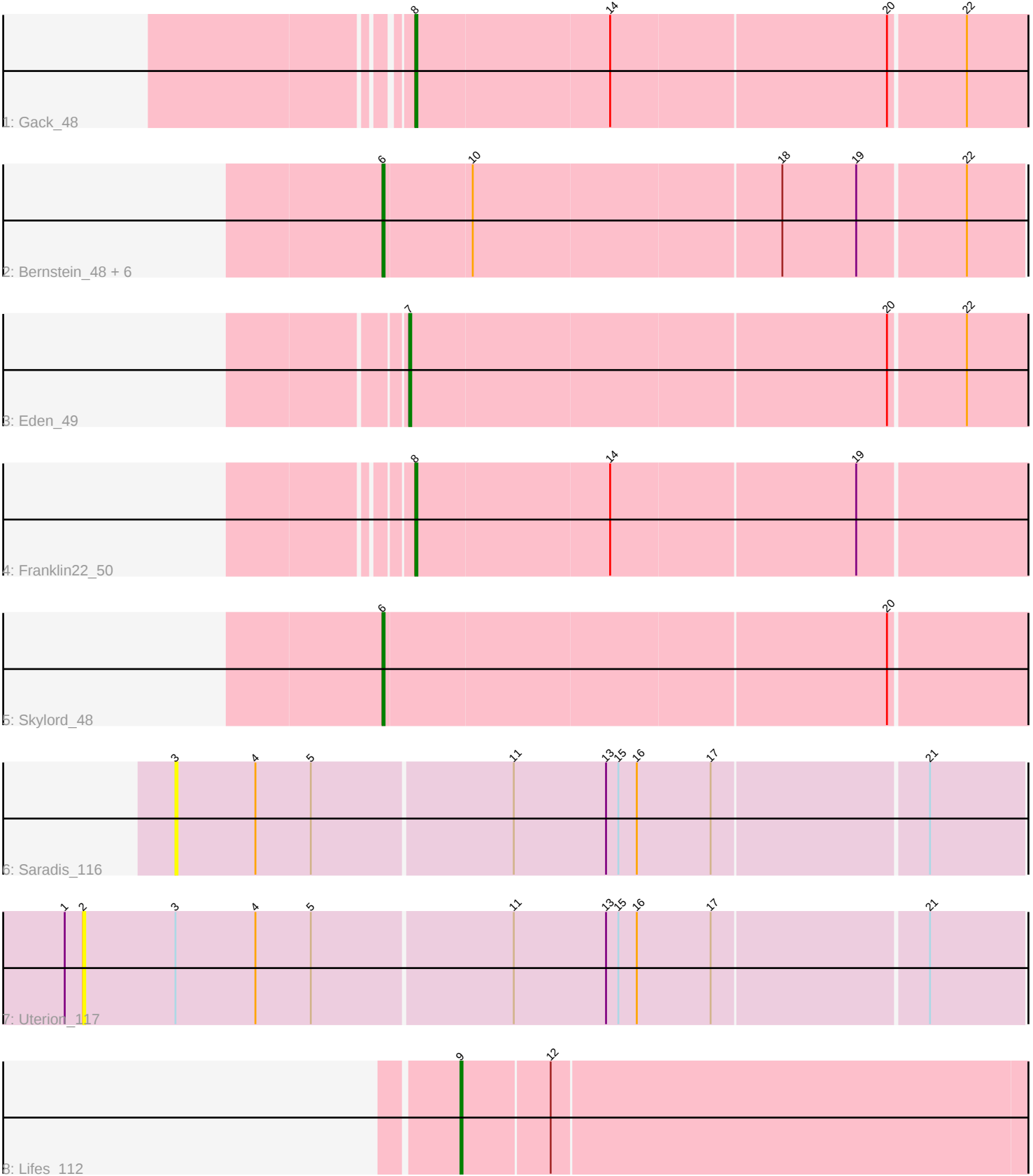


Pham 309536



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 309536 Report

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

Pham number 309536 has 14 members, 2 are drafts.

Phages represented in each track:

- Track 1 : Gack\_48
- Track 2 : Bernstein\_48, Rollins\_48, Clayda5\_50, Coltrane\_48, Brahms\_48, Vitas\_48, Armstrong\_48
- Track 3 : Eden\_49
- Track 4 : Franklin22\_50
- Track 5 : Skylord\_48
- Track 6 : Saradis\_116
- Track 7 : Uterion\_117
- Track 8 : Lifes\_112

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

The start number called the most often in the published annotations is 6, it was called in 8 of the 12 non-draft genes in the pham.

Genes that call this "Most Annotated" start:

- Armstrong\_48, Bernstein\_48, Brahms\_48, Clayda5\_50, Coltrane\_48, Rollins\_48, Skylord\_48, Vitas\_48,

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

- 

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

- Eden\_49, Franklin22\_50, Gack\_48, Lifes\_112, Saradis\_116, Uterion\_117,

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

Start 2:

- Found in 1 of 14 ( 7.1% ) of genes in pham
- No Manual Annotations of this start.
- Called 100.0% of time when present
- Phage (with cluster) where this start called: Uterion\_117 (ED1),

Start 3:

- Found in 2 of 14 ( 14.3% ) of genes in pham
- No Manual Annotations of this start.
- Called 50.0% of time when present
- Phage (with cluster) where this start called: Saradis\_116 (ED1),

Start 6:

- Found in 8 of 14 ( 57.1% ) 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: Armstrong\_48 (EB), Bernstein\_48 (EB), Brahms\_48 (EB), Clayda5\_50 (EB), Coltrane\_48 (EB), Rollins\_48 (EB), Skylord\_48 (EB), Vitas\_48 (EB),

Start 7:

- Found in 1 of 14 ( 7.1% ) 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: Eden\_49 (EB),

Start 8:

- Found in 2 of 14 ( 14.3% ) 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: Franklin22\_50 (EB), Gack\_48 (EB),

Start 9:

- Found in 1 of 14 ( 7.1% ) 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: Lifes\_112 (GB),

### **Summary by clusters:**

There are 3 clusters represented in this pham: ED1, GB, EB,

Info for manual annotations of cluster EB:

- Start number 6 was manually annotated 8 times for cluster EB.
- Start number 7 was manually annotated 1 time for cluster EB.
- Start number 8 was manually annotated 2 times for cluster EB.

Info for manual annotations of cluster GB:

- Start number 9 was manually annotated 1 time for cluster GB.

### **Gene Information:**

Gene: Armstrong\_48 Start: 32505, Stop: 32819, Start Num: 6

Candidate Starts for Armstrong\_48:

(Start: 6 @32505 has 8 MA's), (10, 32547), (18, 32694), (19, 32730), (22, 32781),

Gene: Bernstein\_48 Start: 32503, Stop: 32817, Start Num: 6

Candidate Starts for Bernstein\_48:

(Start: 6 @32503 has 8 MA's), (10, 32545), (18, 32692), (19, 32728), (22, 32779),

Gene: Brahms\_48 Start: 32423, Stop: 32737, Start Num: 6

Candidate Starts for Brahms\_48:

(Start: 6 @32423 has 8 MA's), (10, 32465), (18, 32612), (19, 32648), (22, 32699),

Gene: Clayda5\_50 Start: 32488, Stop: 32802, Start Num: 6

Candidate Starts for Clayda5\_50:

(Start: 6 @32488 has 8 MA's), (10, 32530), (18, 32677), (19, 32713), (22, 32764),

Gene: Coltrane\_48 Start: 32423, Stop: 32737, Start Num: 6

Candidate Starts for Coltrane\_48:

(Start: 6 @32423 has 8 MA's), (10, 32465), (18, 32612), (19, 32648), (22, 32699),

Gene: Eden\_49 Start: 33518, Stop: 33847, Start Num: 7

Candidate Starts for Eden\_49:

(Start: 7 @33518 has 1 MA's), (20, 33746), (22, 33782),

Gene: Franklin22\_50 Start: 33066, Stop: 33392, Start Num: 8

Candidate Starts for Franklin22\_50:

(Start: 8 @33066 has 2 MA's), (14, 33159), (19, 33276),

Gene: Gack\_48 Start: 33118, Stop: 33444, Start Num: 8

Candidate Starts for Gack\_48:

(Start: 8 @33118 has 2 MA's), (14, 33211), (20, 33343), (22, 33379),

Gene: Lifes\_112 Start: 55394, Stop: 55107, Start Num: 9

Candidate Starts for Lifes\_112:

(Start: 9 @55394 has 1 MA's), (12, 55352),

Gene: Rollins\_48 Start: 32503, Stop: 32817, Start Num: 6

Candidate Starts for Rollins\_48:

(Start: 6 @32503 has 8 MA's), (10, 32545), (18, 32692), (19, 32728), (22, 32779),

Gene: Saradis\_116 Start: 59097, Stop: 58678, Start Num: 3

Candidate Starts for Saradis\_116:

(3, 59097), (4, 59058), (5, 59031), (11, 58935), (13, 58890), (15, 58884), (16, 58875), (17, 58839), (21, 58737),

Gene: Skylord\_48 Start: 32434, Stop: 32772, Start Num: 6

Candidate Starts for Skylord\_48:

(Start: 6 @32434 has 8 MA's), (20, 32674),

Gene: Uterion\_117 Start: 59102, Stop: 58638, Start Num: 2

Candidate Starts for Uterion\_117:

(1, 59111), (2, 59102), (3, 59057), (4, 59018), (5, 58991), (11, 58895), (13, 58850), (15, 58844), (16, 58835), (17, 58799), (21, 58697),

Gene: Vitas\_48 Start: 32494, Stop: 32808, Start Num: 6

Candidate Starts for Vitas\_48:

(Start: 6 @32494 has 8 MA's), (10, 32536), (18, 32683), (19, 32719), (22, 32770),