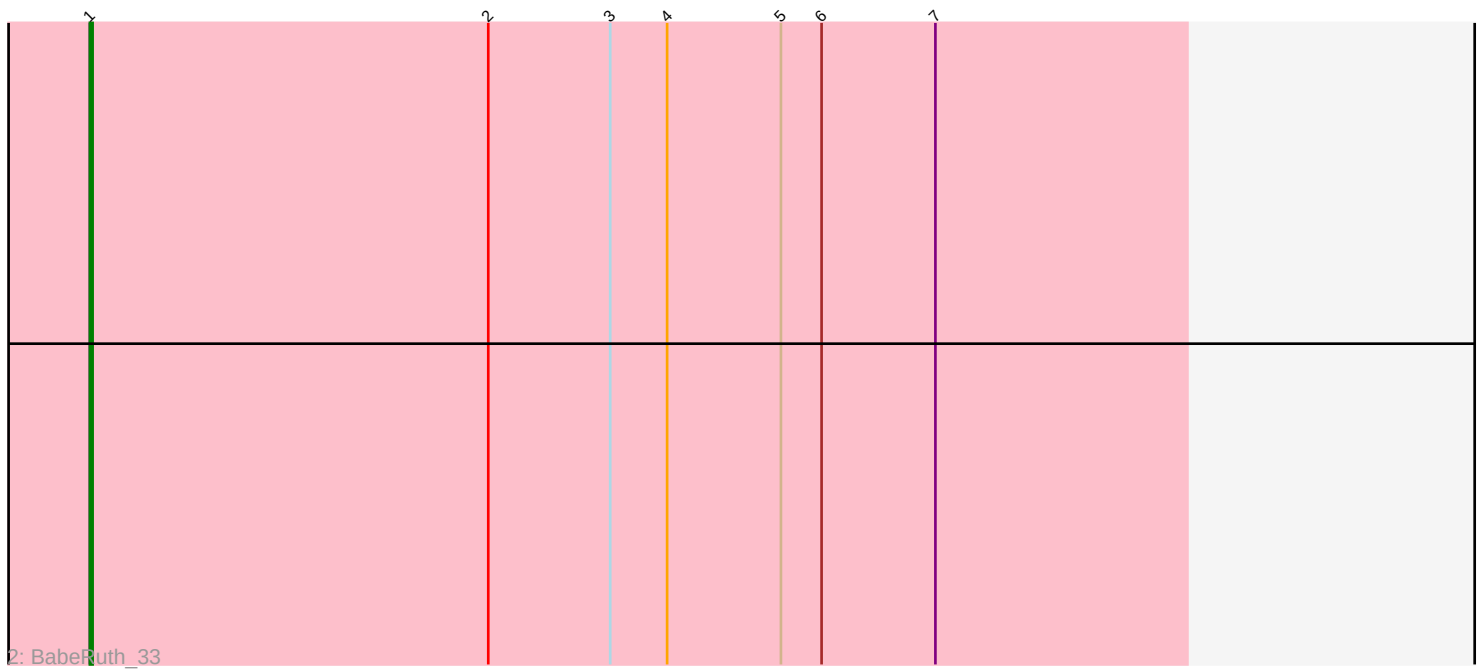
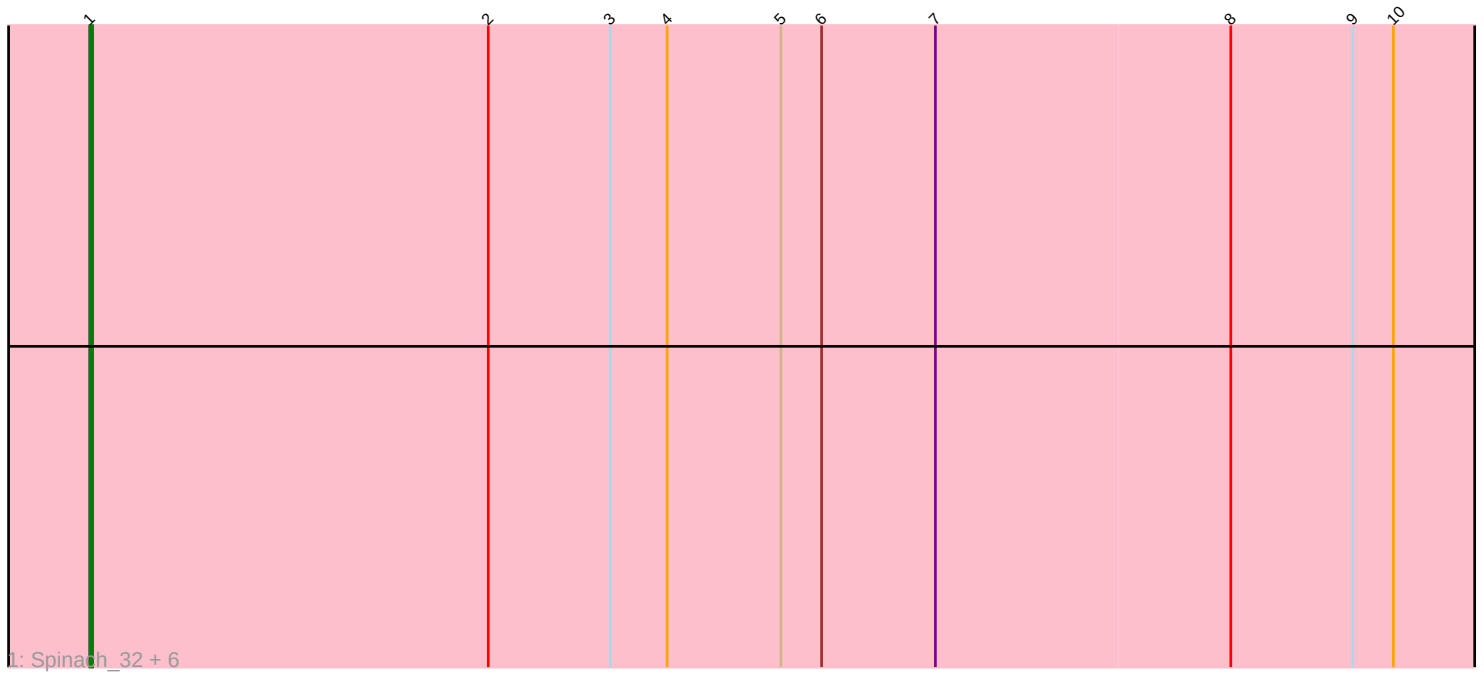


Pham 309892



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 309892 Report

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

Pham number 309892 has 8 members, 0 are drafts.

Phages represented in each track:

- Track 1 : Spinach_32, ShrimpFriedEgg_32, Purgamenstris_32, PhancyPhin_32, Hanako_32, Nенаe_32, Redi_32
- Track 2 : BabeRuth_33

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

The start number called the most often in the published annotations is 1, it was called in 8 of the 8 non-draft genes in the pham.

Genes that call this "Most Annotated" start:

- BabeRuth_33, Hanako_32, Nенаe_32, PhancyPhin_32, Purgamenstris_32, Redi_32, ShrimpFriedEgg_32, Spinach_32,

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 1:

- Found in 8 of 8 (100.0%) of genes in pham
- Manual Annotations of this start: 8 of 8
- Called 100.0% of time when present
- Phage (with cluster) where this start called: BabeRuth_33 (N), Hanako_32 (N), Nенаe_32 (N), PhancyPhin_32 (N), Purgamenstris_32 (N), Redi_32 (N), ShrimpFriedEgg_32 (N), Spinach_32 (N),

Summary by clusters:

There is one cluster represented in this pham: N

Info for manual annotations of cluster N:

- Start number 1 was manually annotated 8 times for cluster N.

Gene Information:

Gene: BabeRuth_33 Start: 26820, Stop: 26416, Start Num: 1

Candidate Starts for BabeRuth_33:

(Start: 1 @26820 has 8 MA's), (2, 26673), (3, 26628), (4, 26607), (5, 26565), (6, 26550), (7, 26508),

Gene: Hanako_32 Start: 26819, Stop: 26262, Start Num: 1

Candidate Starts for Hanako_32:

(Start: 1 @26819 has 8 MA's), (2, 26672), (3, 26627), (4, 26606), (5, 26564), (6, 26549), (7, 26507), (8, 26399), (9, 26354), (10, 26339),

Gene: Nенаe_32 Start: 26819, Stop: 26262, Start Num: 1

Candidate Starts for Nенаe_32:

(Start: 1 @26819 has 8 MA's), (2, 26672), (3, 26627), (4, 26606), (5, 26564), (6, 26549), (7, 26507), (8, 26399), (9, 26354), (10, 26339),

Gene: PhancyPhin_32 Start: 26816, Stop: 26259, Start Num: 1

Candidate Starts for PhancyPhin_32:

(Start: 1 @26816 has 8 MA's), (2, 26669), (3, 26624), (4, 26603), (5, 26561), (6, 26546), (7, 26504), (8, 26396), (9, 26351), (10, 26336),

Gene: Purgamenstris_32 Start: 26819, Stop: 26262, Start Num: 1

Candidate Starts for Purgamenstris_32:

(Start: 1 @26819 has 8 MA's), (2, 26672), (3, 26627), (4, 26606), (5, 26564), (6, 26549), (7, 26507), (8, 26399), (9, 26354), (10, 26339),

Gene: Redi_32 Start: 26819, Stop: 26262, Start Num: 1

Candidate Starts for Redi_32:

(Start: 1 @26819 has 8 MA's), (2, 26672), (3, 26627), (4, 26606), (5, 26564), (6, 26549), (7, 26507), (8, 26399), (9, 26354), (10, 26339),

Gene: ShrimpFriedEgg_32 Start: 26819, Stop: 26262, Start Num: 1

Candidate Starts for ShrimpFriedEgg_32:

(Start: 1 @26819 has 8 MA's), (2, 26672), (3, 26627), (4, 26606), (5, 26564), (6, 26549), (7, 26507), (8, 26399), (9, 26354), (10, 26339),

Gene: Spinach_32 Start: 26819, Stop: 26262, Start Num: 1

Candidate Starts for Spinach_32:

(Start: 1 @26819 has 8 MA's), (2, 26672), (3, 26627), (4, 26606), (5, 26564), (6, 26549), (7, 26507), (8, 26399), (9, 26354), (10, 26339),