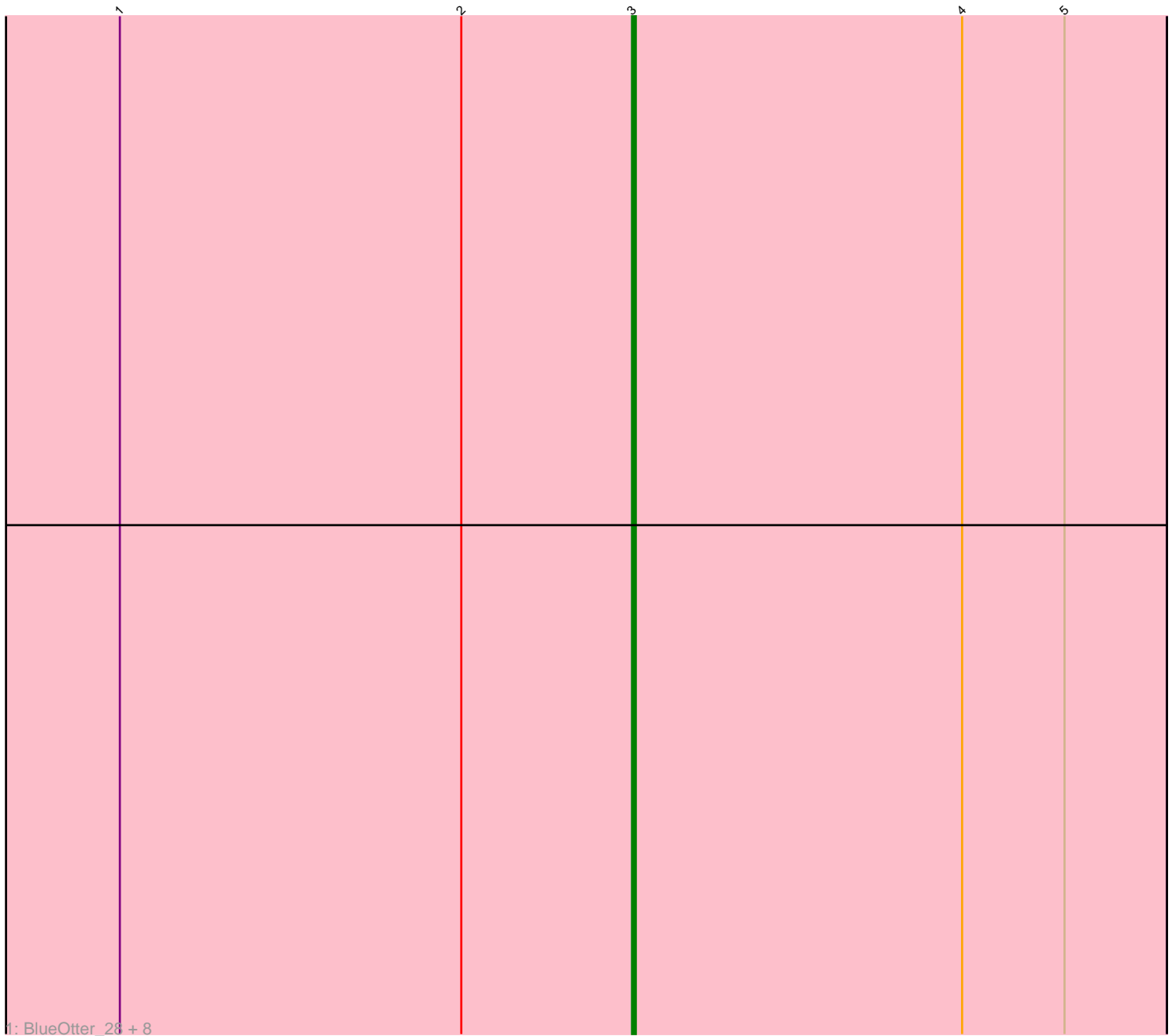


Pham 30960



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 30960 Report

This analysis was run 04/05/24 on database version 557.

Pham number 30960 has 9 members, 5 are drafts.

Phages represented in each track:

- Track 1 : BlueOtter_28, PacManQ_28, Cross_26, Larnav_31, HangryHippo_28, Lululemon_28, Leo04_26, Cursive_24, Watermoore_26

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 4 of the 4 non-draft genes in the pham.

Genes that call this "Most Annotated" start:

- BlueOtter_28, Cross_26, Cursive_24, HangryHippo_28, Larnav_31, Leo04_26, Lululemon_28, PacManQ_28, Watermoore_26,

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 9 of 9 (100.0%) of genes in pham
- Manual Annotations of this start: 4 of 4
- Called 100.0% of time when present
- Phage (with cluster) where this start called: BlueOtter_28 (BE1), Cross_26 (BE1), Cursive_24 (BE1), HangryHippo_28 (BE1), Larnav_31 (BE1), Leo04_26 (BE1), Lululemon_28 (BE1), PacManQ_28 (BE1), Watermoore_26 (BE1),

Summary by clusters:

There is one cluster represented in this pham: BE1

Info for manual annotations of cluster BE1:

- Start number 3 was manually annotated 4 times for cluster BE1.

Gene Information:

Gene: BlueOtter_28 Start: 12246, Stop: 12106, Start Num: 3

Candidate Starts for BlueOtter_28:

(1, 12381), (2, 12291), (Start: 3 @12246 has 4 MA's), (4, 12159), (5, 12132),

Gene: Cross_26 Start: 12247, Stop: 12107, Start Num: 3

Candidate Starts for Cross_26:

(1, 12382), (2, 12292), (Start: 3 @12247 has 4 MA's), (4, 12160), (5, 12133),

Gene: Cursive_24 Start: 11064, Stop: 10924, Start Num: 3

Candidate Starts for Cursive_24:

(1, 11199), (2, 11109), (Start: 3 @11064 has 4 MA's), (4, 10977), (5, 10950),

Gene: HangryHippo_28 Start: 12246, Stop: 12106, Start Num: 3

Candidate Starts for HangryHippo_28:

(1, 12381), (2, 12291), (Start: 3 @12246 has 4 MA's), (4, 12159), (5, 12132),

Gene: Larnav_31 Start: 12205, Stop: 12065, Start Num: 3

Candidate Starts for Larnav_31:

(1, 12340), (2, 12250), (Start: 3 @12205 has 4 MA's), (4, 12118), (5, 12091),

Gene: Leo04_26 Start: 12245, Stop: 12105, Start Num: 3

Candidate Starts for Leo04_26:

(1, 12380), (2, 12290), (Start: 3 @12245 has 4 MA's), (4, 12158), (5, 12131),

Gene: Lululemon_28 Start: 11626, Stop: 11486, Start Num: 3

Candidate Starts for Lululemon_28:

(1, 11761), (2, 11671), (Start: 3 @11626 has 4 MA's), (4, 11539), (5, 11512),

Gene: PacManQ_28 Start: 11626, Stop: 11486, Start Num: 3

Candidate Starts for PacManQ_28:

(1, 11761), (2, 11671), (Start: 3 @11626 has 4 MA's), (4, 11539), (5, 11512),

Gene: Watermoore_26 Start: 12247, Stop: 12107, Start Num: 3

Candidate Starts for Watermoore_26:

(1, 12382), (2, 12292), (Start: 3 @12247 has 4 MA's), (4, 12160), (5, 12133),