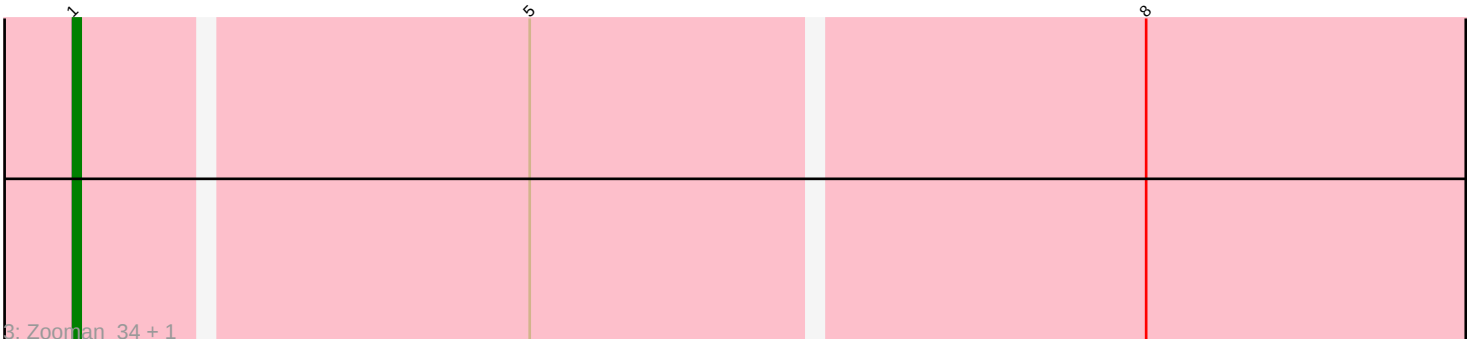
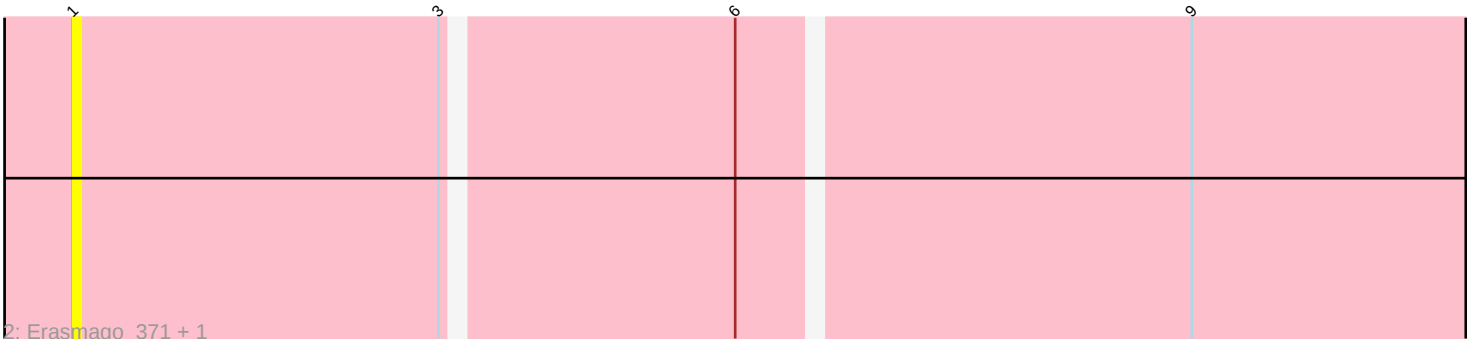
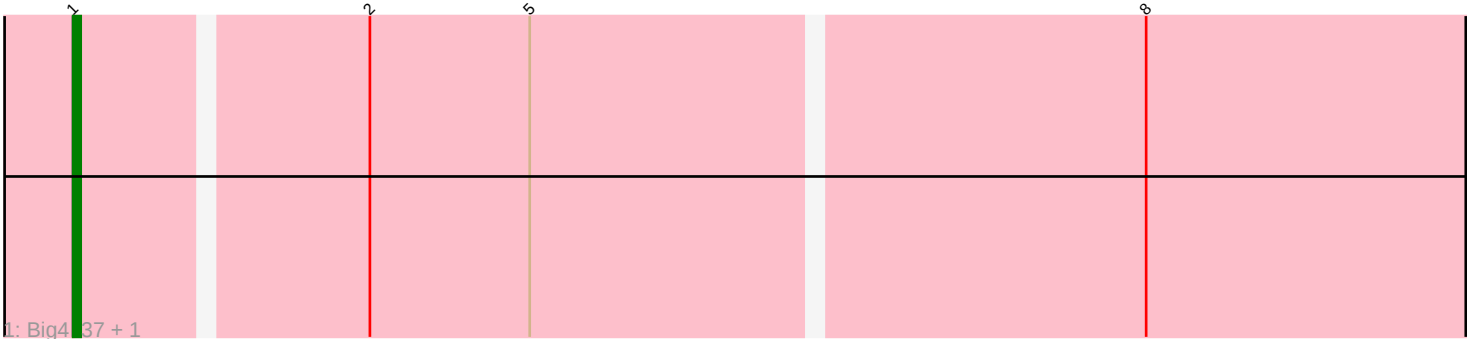




Note: Tracks are now grouped by subcluster and scaled. Switching in subcluster is indicated by changes in track color. Track scale is now set by default to display the region 30 bp upstream of start 1 to 30 bp downstream of the last possible start. If this default region is judged to be packed too tightly 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 278906 Report

This analysis was run 02/07/26 on database version 634.

Pham number 278906 has 8 members, 2 are drafts.

Phages represented in each track:

- Track 1 : Big4_37, Big4_363
- Track 2 : Erasmago_371, Erasmago_33
- Track 3 : Zooman_34, Zooman_347
- Track 4 : Pumpernickel_343, Pumpernickel_42

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 6 of the 6 non-draft genes in the pham.

Genes that call this "Most Annotated" start:

- Big4_363, Big4_37, Erasmago_33, Erasmago_371, Pumpernickel_343, Pumpernickel_42, Zooman_34, Zooman_347,

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: 6 of 6
- Called 100.0% of time when present
- Phage (with cluster) where this start called: Big4_363 (GD2), Big4_37 (GD2), Erasmago_33 (GD2), Erasmago_371 (GD2), Pumpernickel_343 (GD4), Pumpernickel_42 (GD4), Zooman_34 (GD2), Zooman_347 (GD2),

Summary by clusters:

There are 2 clusters represented in this pham: GD2, GD4,

Info for manual annotations of cluster GD2:

- Start number 1 was manually annotated 4 times for cluster GD2.

Info for manual annotations of cluster GD4:

- Start number 1 was manually annotated 2 times for cluster GD4.

Gene Information:

Gene: Big4_37 Start: 14515, Stop: 14694, Start Num: 1

Candidate Starts for Big4_37:

(Start: 1 @14515 has 6 MA's), (2, 14551), (5, 14572), (8, 14650),

Gene: Big4_363 Start: 189209, Stop: 189388, Start Num: 1

Candidate Starts for Big4_363:

(Start: 1 @189209 has 6 MA's), (2, 189245), (5, 189266), (8, 189344),

Gene: Erasmago_371 Start: 187039, Stop: 187218, Start Num: 1

Candidate Starts for Erasmago_371:

(Start: 1 @187039 has 6 MA's), (3, 187087), (6, 187123), (9, 187180),

Gene: Erasmago_33 Start: 12530, Stop: 12709, Start Num: 1

Candidate Starts for Erasmago_33:

(Start: 1 @12530 has 6 MA's), (3, 12578), (6, 12614), (9, 12671),

Gene: Pumpernickel_343 Start: 182268, Stop: 182447, Start Num: 1

Candidate Starts for Pumpernickel_343:

(Start: 1 @182268 has 6 MA's), (4, 182316), (7, 182397), (8, 182406), (10, 182415),

Gene: Pumpernickel_42 Start: 16136, Stop: 16315, Start Num: 1

Candidate Starts for Pumpernickel_42:

(Start: 1 @16136 has 6 MA's), (4, 16184), (7, 16265), (8, 16274), (10, 16283),

Gene: Zooman_34 Start: 13966, Stop: 14145, Start Num: 1

Candidate Starts for Zooman_34:

(Start: 1 @13966 has 6 MA's), (5, 14023), (8, 14101),

Gene: Zooman_347 Start: 189617, Stop: 189796, Start Num: 1

Candidate Starts for Zooman_347:

(Start: 1 @189617 has 6 MA's), (5, 189674), (8, 189752),