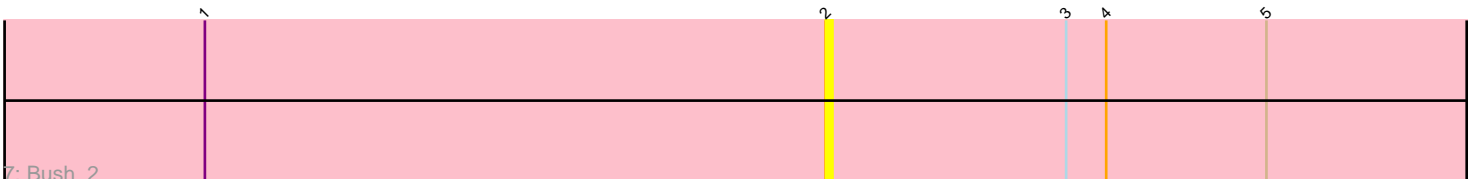


Pham 8570



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 8570 Report

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

Pham number 8570 has 9 members, 2 are drafts.

Phages represented in each track:

- Track 1 : Antuna_2
- Track 2 : Blett_2
- Track 3 : Dropshot_2, Appa_2
- Track 4 : Phonegingi_2
- Track 5 : MenE_2
- Track 6 : Pickles13_2
- Track 7 : Bush_2
- Track 8 : Warren_2

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

The start number called the most often in the published annotations is 2, it was called in 7 of the 7 non-draft genes in the pham.

Genes that call this "Most Annotated" start:

- Antuna_2, Appa_2, Blett_2, Bush_2, Dropshot_2, MenE_2, Phonegingi_2, Pickles13_2, Warren_2,

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 2:

- Found in 9 of 9 (100.0%) of genes in pham
- Manual Annotations of this start: 7 of 7
- Called 100.0% of time when present
- Phage (with cluster) where this start called: Antuna_2 (GA), Appa_2 (GA), Blett_2 (GA), Bush_2 (GA), Dropshot_2 (GA), MenE_2 (GA), Phonegingi_2 (GA), Pickles13_2 (GA), Warren_2 (GA),

Summary by clusters:

There is one cluster represented in this pham: GA

Info for manual annotations of cluster GA:

- Start number 2 was manually annotated 7 times for cluster GA.

Gene Information:

Gene: Antuna_2 Start: 501, Stop: 638, Start Num: 2

Candidate Starts for Antuna_2:

(1, 426), (Start: 2 @501 has 7 MA's), (3, 537), (4, 543), (5, 567),

Gene: Appa_2 Start: 519, Stop: 656, Start Num: 2

Candidate Starts for Appa_2:

(1, 426), (Start: 2 @519 has 7 MA's),

Gene: Blett_2 Start: 510, Stop: 647, Start Num: 2

Candidate Starts for Blett_2:

(1, 426), (Start: 2 @510 has 7 MA's), (4, 552),

Gene: Bush_2 Start: 519, Stop: 656, Start Num: 2

Candidate Starts for Bush_2:

(1, 426), (Start: 2 @519 has 7 MA's), (3, 555), (4, 561), (5, 585),

Gene: Dropshot_2 Start: 519, Stop: 656, Start Num: 2

Candidate Starts for Dropshot_2:

(1, 426), (Start: 2 @519 has 7 MA's),

Gene: MenE_2 Start: 519, Stop: 656, Start Num: 2

Candidate Starts for MenE_2:

(1, 426), (Start: 2 @519 has 7 MA's),

Gene: Phonegingi_2 Start: 519, Stop: 647, Start Num: 2

Candidate Starts for Phonegingi_2:

(1, 429), (Start: 2 @519 has 7 MA's), (3, 555), (4, 561),

Gene: Pickles13_2 Start: 510, Stop: 647, Start Num: 2

Candidate Starts for Pickles13_2:

(Start: 2 @510 has 7 MA's), (5, 576),

Gene: Warren_2 Start: 519, Stop: 656, Start Num: 2

Candidate Starts for Warren_2:

(1, 426), (Start: 2 @519 has 7 MA's), (5, 585),