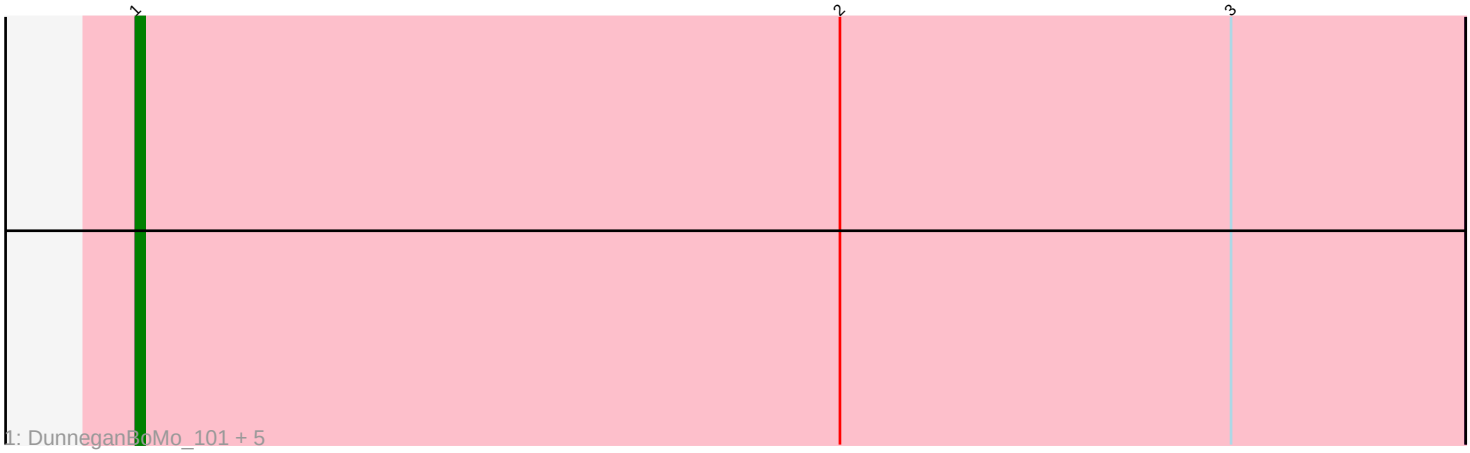
Pham 342856


1
2
3
1: DunneganBoMo_101 + 5


1
3
4
2: Atuin_106


1
3
3: ReginaGlobina_111

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 342856 Report

This analysis was run 11/14/26 on database version 672.

Pham number 342856 has 8 members, 2 are drafts.

Phages represented in each track:
• Track 1 : DunneganBoMo_101, BooTeria_102, Ellewin_100, Panchaali_103, WaddleDee_99, Emmetator_104
• Track 2 : Atuin_106
• Track 3 : ReginaGlobina_111

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:
• Atuin_106, BooTeria_102, DunneganBoMo_101, Ellewin_100, Emmetator_104, Panchaali_103, ReginaGlobina_111, WaddleDee_99,

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: Atuin_106 (FC), BooTeria_102 (FC), DunneganBoMo_101 (FC), Ellewin_100 (FC), Emmetator_104 (FC), Panchaali_103 (FC), ReginaGlobina_111 (FC), WaddleDee_99 (FC),

Summary by clusters:

There is one cluster represented in this pham: FC

Info for manual annotations of cluster FC:
•Start number 1 was manually annotated 6 times for cluster FC.

Gene Information:

Gene: Atuin_106 Start: 90868, Stop: 91020, Start Num: 1
Candidate Starts for Atuin_106:
(Start: 1 @90868 has 6 MA's), (3, 90994), (4, 91006),

Gene: BooTeria_102 Start: 87475, Stop: 87627, Start Num: 1
Candidate Starts for BooTeria_102:
(Start: 1 @87475 has 6 MA's), (2, 87556), (3, 87601),

Gene: DunneganBoMo_101 Start: 86744, Stop: 86896, Start Num: 1
Candidate Starts for DunneganBoMo_101:
(Start: 1 @86744 has 6 MA's), (2, 86825), (3, 86870),

Gene: Ellewin_100 Start: 86152, Stop: 86304, Start Num: 1
Candidate Starts for Ellewin_100:
(Start: 1 @86152 has 6 MA's), (2, 86233), (3, 86278),

Gene: Emmetator_104 Start: 86699, Stop: 86851, Start Num: 1
Candidate Starts for Emmetator_104:
(Start: 1 @86699 has 6 MA's), (2, 86780), (3, 86825),

Gene: Panchaali_103 Start: 87727, Stop: 87879, Start Num: 1
Candidate Starts for Panchaali_103:
(Start: 1 @87727 has 6 MA's), (2, 87808), (3, 87853),

Gene: ReginaGlobina_111 Start: 90670, Stop: 90822, Start Num: 1
Candidate Starts for ReginaGlobina_111:
(Start: 1 @90670 has 6 MA's), (3, 90796),

Gene: WaddleDee_99 Start: 85930, Stop: 86082, Start Num: 1
Candidate Starts for WaddleDee_99:
(Start: 1 @85930 has 6 MA's), (2, 86011), (3, 86056),