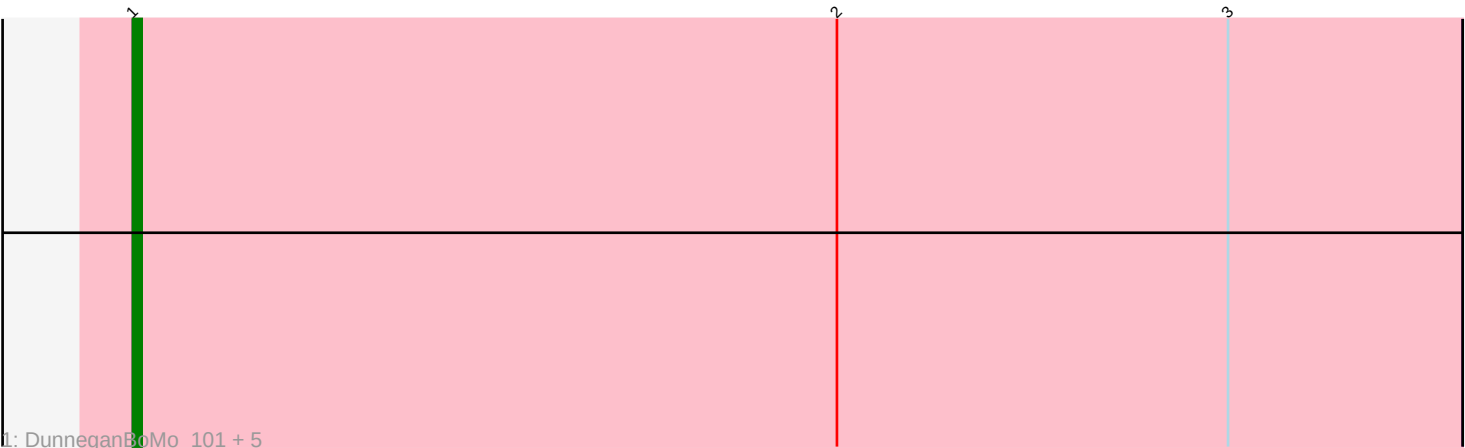




Note: Tracks are now grouped by subcluster and scaled. Switching in subcluster is indicated by changes in track color. Track scale is now set by default to display the region 30 bp upstream of start 1 to 30 bp downstream of the last possible start. If this default region is judged to be packed too tightly 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 329547 Report

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

Pham number 329547 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),