

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 143116 Report

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

Pham number 143116 has 8 members, 7 are drafts.

Phages represented in each track:

• Track 1: DunneganBoMo 251

Track 2: Mimi 250, Bloom 249, Racecar 247

• Track 3 : DunneganBoMo\_253

Track 4 : DunneganBoMo\_252

• Track 5 : Talia1610\_247

Track 6 : SJReid\_276

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

The start number called the most often in the published annotations is 5, it was called in 1 of the 1 non-draft genes in the pham.

Genes that call this "Most Annotated" start:

Bloom\_249, Mimi\_250, Racecar\_247, Talia1610\_247,

Genes that have the "Most Annotated" start but do not call it:

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Genes that do not have the "Most Annotated" start:

• DunneganBoMo\_251, DunneganBoMo\_252, DunneganBoMo\_253, SJReid\_276,

# Summary by start number:

#### Start 2:

- Found in 1 of 8 (12.5%) of genes in pham
- No Manual Annotations of this start.
- Called 100.0% of time when present
- Phage (with cluster) where this start called: DunneganBoMo\_252 (FC),

### Start 5:

- Found in 4 of 8 (50.0%) of genes in pham
- Manual Annotations of this start: 1 of 1
- Called 100.0% of time when present

• Phage (with cluster) where this start called: Bloom\_249 (FC), Mimi\_250 (FC), Racecar\_247 (FC), Talia1610\_247 (FC),

## Start 6:

- Found in 2 of 8 (25.0%) of genes in pham
- No Manual Annotations of this start.
- Called 50.0% of time when present
- Phage (with cluster) where this start called: DunneganBoMo\_251 (FC),

## Start 11:

- Found in 1 of 8 (12.5%) of genes in pham
- No Manual Annotations of this start.
- Called 100.0% of time when present
- Phage (with cluster) where this start called: DunneganBoMo\_253 (FC),

#### Start 12:

- Found in 1 of 8 (12.5%) of genes in pham
- No Manual Annotations of this start.
- Called 100.0% of time when present
- Phage (with cluster) where this start called: SJReid\_276 (FC),

## Summary by clusters:

There is one cluster represented in this pham: FC

Info for manual annotations of cluster FC:

•Start number 5 was manually annotated 1 time for cluster FC.

### Gene Information:

Gene: Bloom 249 Start: 157954, Stop: 158643, Start Num: 5

Candidate Starts for Bloom 249:

(Start: 5 @ 157954 has 1 MA's), (13, 158056), (17, 158113), (24, 158197), (25, 158200), (32, 158410),

Gene: DunneganBoMo\_251 Start: 161249, Stop: 161899, Start Num: 6

Candidate Starts for DunneganBoMo 251:

(6, 161249), (10, 161267), (22, 161456), (27, 161558), (30, 161654), (33, 161708), (35, 161747),

Gene: DunneganBoMo 253 Start: 162842, Stop: 163462, Start Num: 11

Candidate Starts for DunneganBoMo\_253:

(9, 162806), (11, 162842), (15, 162917), (18, 162950), (36, 163379), (37, 163394), (38, 163454),

Gene: DunneganBoMo\_252 Start: 161908, Stop: 162771, Start Num: 2

Candidate Starts for DunneganBoMo\_252:

(1, 161782), (2, 161908), (3, 161992), (4, 161995), (27, 162430),

Gene: Mimi 250 Start: 157329, Stop: 158018, Start Num: 5

Candidate Starts for Mimi 250:

(Start: 5 @ 157329 has 1 MA's), (13, 157431), (17, 157488), (24, 157572), (25, 157575), (32, 157785),

Gene: Racecar\_247 Start: 157708, Stop: 158397, Start Num: 5

Candidate Starts for Racecar\_247:

(Start: 5 @ 157708 has 1 MA's), (13, 157810), (17, 157867), (24, 157951), (25, 157954), (32, 158164),

Gene: SJReid\_276 Start: 159658, Stop: 160269, Start Num: 12

Candidate Starts for SJReid\_276:

(6, 159604), (7, 159610), (8, 159616), (12, 159658), (14, 159736), (16, 159757), (19, 159781), (20, 159805), (21, 159811), (23, 159850), (26, 159949), (28, 159955), (29, 159958), (30, 160045), (31, 160060), (34, 160105),

Gene: Talia1610\_247 Start: 157983, Stop: 158684, Start Num: 5

Candidate Starts for Talia1610\_247:

(Start: 5 @ 157983 has 1 MA's), (13, 158085), (24, 158226), (25, 158229), (32, 158439),