Pham 136201

	,	1
1: Talia1610_56 + 2		
	ę)
2: Mimi_58 + 1		
	,	
B: DunneganBoMo_44		
b. Durineganbolvio_44		'
3	•	
4: SJReid_55		
5: Atuin_44		

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 136201 Report

This analysis was run 04/28/24 on database version 559.

Pham number 136201 has 8 members, 7 are drafts.

Phages represented in each track:

Track 1: Talia1610_56, Patbob_48, Bloom_57

• Track 2 : Mimi_58, Racecar_53

Track 3 : DunneganBoMo_44Track 4 : SJReid 55

• Track 5 : Atuin_44

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

The start number called the most often in the published annotations is 3, it was called in 1 of the 1 non-draft genes in the pham.

Genes that call this "Most Annotated" start:

• Atuin_44, Bloom_57, DunneganBoMo_44, Mimi_58, Patbob_48, Racecar_53, SJReid_55, Talia1610_56,

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:

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Summary by start number:

Start 3:

- Found in 8 of 8 (100.0%) 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: Atuin_44 (FC), Bloom_57 (FC), DunneganBoMo_44 (FC), Mimi_58 (FC), Patbob_48 (FC), Racecar_53 (FC), SJReid_55 (FC), Talia1610_56 (FC),

Summary by clusters:

There is one cluster represented in this pham: FC

Info for manual annotations of cluster FC:

•Start number 3 was manually annotated 1 time for cluster FC.

Gene Information:

Gene: Atuin_44 Start: 18462, Stop: 18800, Start Num: 3

Candidate Starts for Atuin_44:

(Start: 3 @18462 has 1 MA's), (5, 18528),

Gene: Bloom_57 Start: 23011, Stop: 23370, Start Num: 3

Candidate Starts for Bloom_57:

(Start: 3 @23011 has 1 MA's), (4, 23062), (5, 23077), (7, 23083),

Gene: DunneganBoMo 44 Start: 17038, Stop: 17346, Start Num: 3

Candidate Starts for DunneganBoMo_44:

(1, 17005), (2, 17011), (Start: 3 @17038 has 1 MA's), (5, 17104), (6, 17107),

Gene: Mimi 58 Start: 22124, Stop: 22483, Start Num: 3

Candidate Starts for Mimi 58:

(Start: 3 @22124 has 1 MA's), (5, 22190), (7, 22196),

Gene: Patbob_48 Start: 21496, Stop: 21855, Start Num: 3

Candidate Starts for Patbob 48:

(Start: 3 @ 21496 has 1 MA's), (4, 21547), (5, 21562), (7, 21568),

Gene: Racecar_53 Start: 22777, Stop: 23136, Start Num: 3

Candidate Starts for Racecar_53:

(Start: 3 @22777 has 1 MA's), (5, 22843), (7, 22849),

Gene: SJReid 55 Start: 22818, Stop: 23138, Start Num: 3

Candidate Starts for SJReid 55:

(Start: 3 @22818 has 1 MA's), (5, 22884),

Gene: Talia1610_56 Start: 22142, Stop: 22501, Start Num: 3

Candidate Starts for Talia1610_56:

(Start: 3 @ 22142 has 1 MA's), (4, 22193), (5, 22208), (7, 22214),