I. Pad ioDiaboli, 325 ± 1		<u> </u>)	k 6	6 1	, ق	,0	, ,	v
1: Pauloliana 925.4.1									
1: Paulohiana 125 a 1									
S. BalloDishell, 325 ± 1									
v PolikiDishali 325 ± 1									
v BalkDisholi 325 ± 1									
t- PatioDiaholi 226 ± 1									
I. PalioDiaholi 325 + 1									
I: PalioDiaholi 335 + 1									
L. PattoDiaboli, 325 ± 1									
ty PathoDiaboli, 325 ± 1									
It Patholisholi 325 ± 1									
1: PatiloDiaholi 335 ± 1									
I: PallioDiaholi 325 ± 1									
t PalidaDiahaji 325 ± 1									
I: PaliloDiahali 325 ± 1									
t: PauloDiaholi 325 ± 1									
t: PalkloDiaholi 325 ± 1									
t: PaltoDiaholi 325 ± 1									
1: PalloDiaholi 325 ± 1									
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1- PalloDiaholi 325 ± 1									
1- PallloDiaholi 325 ± 1									
1- PaltloDiaholi 325 ± 1									
1- PallloDiaholi 325 ± 1									
1: PauloDiaholi 325 ± 1									
	1 · Do	uloDiaboli	325 ± 1						

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 13566 Report

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

Pham number 13566 has 2 members, 0 are drafts.

Phages represented in each track:

Track 1 : PauloDiaboli_325, A3Wally_324

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 2 of the 2 non-draft genes in the pham.

Genes that call this "Most Annotated" start:

• A3Wally_324, PauloDiaboli_325,

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 2 of 2 (100.0%) of genes in pham
- Manual Annotations of this start: 2 of 2
- Called 100.0% of time when present
- Phage (with cluster) where this start called: A3Wally_324 (GD1), PauloDiaboli_325 (GD1),

Summary by clusters:

There is one cluster represented in this pham: GD1

Info for manual annotations of cluster GD1:

•Start number 1 was manually annotated 2 times for cluster GD1.

Gene Information:

Gene: A3Wally_324 Start: 170287, Stop: 170625, Start Num: 1

Candidate Starts for A3Wally_324:

(Start: 1 @170287 has 2 MA's), (2, 170311), (3, 170356), (4, 170359), (5, 170380), (6, 170398), (7, 170470), (8, 170473), (9, 170512), (10, 170515), (11, 170578), (12, 170596),

Gene: PauloDiaboli_325 Start: 167800, Stop: 168132, Start Num: 1

Candidate Starts for PauloDiaboli_325:

(Start: 1 @167800 has 2 MA's), (2, 167818), (3, 167863), (4, 167866), (5, 167887), (6, 167905), (7, 167977), (8, 167980), (9, 168019), (10, 168022), (11, 168085), (12, 168103),