1: Lonel/Bol, 74 + 1		<u>,</u>	ე	<u>ه</u> 5	<i>ۆ</i> ،	18	9 10	,
1 Longle (Sc). 74 + 1								
1 Jone W. Rol. 74 + 1								
1 Longly RG1 74 + 1								
1 Jone Rei 74+1								
N. Longi-Rai, 7.4 + 1								
It Jonew Roi, 74+1								
I' Lonek-Boi, 74+1								
t'i nnekBai 74+1								
t: Lone/Poi 74 + 1								
1: Lone/PBi 74 + 1								
1: Lone//Roi 74 + 1								
1: Lone VR01 74 + 1								
1: Lone vBri 74+1								
1: LonelyBai 74+1								
1: Lone (Boi, 74 + 1								
1: Lone WBoi, 74 + 1								
1: LonelyBoi, 74 + 1								
1: Lone /Boi 74 + 1								
1: Lone WBgi 74 + 1								
1: Lone vBoi 74 + 1								
1: LoneivBoi: 74 + 1								
1: LonelyBoi, 74 + 1								
1: Lone WBoi 74 + 1								
1: Lone vBoi: 74 + 1								
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1: Lone vBoi 74 + 1								
	1: Lone	VBoi 74 + 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 90028 Report

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

Pham number 90028 has 2 members, 0 are drafts.

Phages represented in each track:

Track 1 : LonelyBoi_74, Confidence_75

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

The start number called the most often in the published annotations is 4, it was called in 2 of the 2 non-draft genes in the pham.

Genes that call this "Most Annotated" start:

• Confidence_75, LonelyBoi_74,

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 4:

- 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: Confidence_75 (CY1), LonelyBoi_74 (CY),

Summary by clusters:

There are 2 clusters represented in this pham: CY, CY1,

Info for manual annotations of cluster CY:

•Start number 4 was manually annotated 1 time for cluster CY.

Info for manual annotations of cluster CY1:

Start number 4 was manually annotated 1 time for cluster CY1.

Gene Information:

Gene: Confidence_75 Start: 48360, Stop: 48758, Start Num: 4

Candidate Starts for Confidence_75:

(1, 48129), (2, 48246), (3, 48327), (Start: 4 @48360 has 2 MA's), (5, 48414), (6, 48507), (7, 48519), (8, 48522), (9, 48636), (10, 48660),

Gene: LonelyBoi_74 Start: 51507, Stop: 51905, Start Num: 4

Candidate Starts for LonelyBoi_74:

 $(1,\,51276),\,(2,\,51393),\,(3,\,51474),\,(Start:\,4\,\,@\,51507\,\,has\,\,2\,\,MA's),\,(5,\,51561),\,(6,\,51654),\,(7,\,51666),\,(8,\,61654),\,(1,\,10066),\,(1,\,1$

51669), (9, 51783), (10, 51807),