# Zoomed Pham 170593

	?≱	\
4. Parrie 20		
1: Berrie_32		
		( 8 <sup>0</sup>
2: Pumpkins_31		
		8,0
B: Cassia_31 + 1		
	<b>N.</b> 6	, o,o
4: Mudpuppy_31		
	<b>S</b>	^
5: ObiToo_33 + 1		
		,o 
6: VResidence_31		

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 170593 Report

This analysis was run 07/09/24 on database version 566.

Pham number 170593 has 8 members, 1 are drafts.

Phages represented in each track:

Track 1 : Berrie\_32

• Track 2 : Pumpkins\_31

Track 3 : Cassia\_31, TforTroy\_32

Track 4 : Mudpuppy\_31

Track 5 : ObiToo\_33, Crewmate\_34

• Track 6 : VResidence\_31

## 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 5 of the 7 non-draft genes in the pham.

Genes that call this "Most Annotated" start:

Berrie\_32, Cassia\_31, Mudpuppy\_31, Pumpkins\_31, TforTroy\_32, VResidence\_31,

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:

Crewmate\_34, ObiToo\_33,

## Summary by start number:

#### Start 4:

- Found in 6 of 8 (75.0%) of genes in pham
- Manual Annotations of this start: 5 of 7
- Called 100.0% of time when present
- Phage (with cluster) where this start called: Berrie\_32 (AZ1), Cassia\_31 (AZ1), Mudpuppy\_31 (AZ1), Pumpkins\_31 (AZ1), TforTroy\_32 (AZ1), VResidence\_31 (AZ1),

#### Start 5:

- Found in 2 of 8 (25.0%) of genes in pham
- Manual Annotations of this start: 2 of 7
- Called 100.0% of time when present

Phage (with cluster) where this start called: Crewmate\_34 (AZ1), ObiToo\_33 (AZ1),

## **Summary by clusters:**

There is one cluster represented in this pham: AZ1

Info for manual annotations of cluster AZ1:

- •Start number 4 was manually annotated 5 times for cluster AZ1.
- •Start number 5 was manually annotated 2 times for cluster AZ1.

#### Gene Information:

Gene: Berrie\_32 Start: 25131, Stop: 25277, Start Num: 4

Candidate Starts for Berrie 32:

(3, 25128), (Start: 4 @25131 has 5 MA's), (7, 25167), (9, 25230), (12, 25263),

Gene: Cassia\_31 Start: 24506, Stop: 24652, Start Num: 4

Candidate Starts for Cassia\_31:

(Start: 4 @24506 has 5 MA's), (7, 24542), (8, 24599), (10, 24608),

Gene: Crewmate\_34 Start: 24765, Stop: 24911, Start Num: 5

Candidate Starts for Crewmate 34:

(1, 24177), (Start: 5 @ 24765 has 2 MA's), (11, 24879),

Gene: Mudpuppy 31 Start: 23961, Stop: 24107, Start Num: 4

Candidate Starts for Mudpuppy\_31:

(2, 23766), (Start: 4 @ 23961 has 5 MA's), (6, 23970), (7, 23997), (9, 24060), (10, 24063),

Gene: ObiToo\_33 Start: 24505, Stop: 24651, Start Num: 5

Candidate Starts for ObiToo 33:

(1, 23917), (Start: 5 @24505 has 2 MA's), (11, 24619),

Gene: Pumpkins 31 Start: 24916, Stop: 25062, Start Num: 4

Candidate Starts for Pumpkins\_31:

(Start: 4 @ 24916 has 5 MA's), (7, 24952), (8, 25009), (10, 25018),

Gene: TforTroy\_32 Start: 24790, Stop: 24936, Start Num: 4

Candidate Starts for TforTroy 32:

(Start: 4 @24790 has 5 MA's), (7, 24826), (8, 24883), (10, 24892),

Gene: VResidence 31 Start: 24143, Stop: 24289, Start Num: 4

Candidate Starts for VResidence\_31:

(Start: 4 @ 24143 has 5 MA's), (10, 24245),