Pham 200750

		<del>م</del> <sup>ري</sup>	20		
1: Dryad_84					
	6 1	onR <sup>h</sup> n <sup>3</sup>	, <i>6</i> ,870		
2: Rooney_82					
		R 121×	15 VA 25		
B: Rooney_68 + 1					
×γ 3 ≽	6	ond is	৻ড়৻ড়৻ঢ়		
4: Glbson_82					
	6	ark ro	.80		
5: Lizz_80 + 2					
		<del>ر</del> ۍ	20		
6: Greenbelt_124					
		NO NO	<b>1</b> 80		
7: Wilkos_82					
		<i>\</i> <sup>⊙</sup>	20		Ŷ
B: Yara_78					
		<u>ب</u> ې	1	Ŷ	
9: Dryad_72					
		arth right	1 <sup>0</sup>		

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 200750 Report

This analysis was run 01/18/25 on database version 583.

Pham number 200750 has 13 members, 2 are drafts.

Phages represented in each track:

- Track 1 : Dryad\_84
- Track 2 : Rooney\_82
- Track 3 : Rooney\_68, Gibson\_68
- Track 4 : Gibson\_82
- Track 5 : Lizz\_80, PHTowN\_80, ShakeNBake\_80
- Track 6 : Greenbelt\_124
- Track 7 : Wilkos\_82
- Track 8 : Yara\_78
- Track 9 : Dryad\_72
- Track 10 : Wentworth\_79

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

The start number called the most often in the published annotations is 13, it was called in 9 of the 11 non-draft genes in the pham.

Genes that call this "Most Annotated" start: • Dryad\_72, Dryad\_84, Gibson\_82, Greenbelt\_124, Lizz\_80, PHTowN\_80, Rooney\_82, ShakeNBake\_80, Wentworth\_79, Wilkos\_82, Yara\_78,

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

Genes that do not have the "Most Annotated" start: • Gibson\_68, Rooney\_68,

## Summary by start number:

Start 13:

- Found in 11 of 13 (84.6%) of genes in pham
- Manual Annotation's of this start: 9 of 11
- Called 100.0% of time when present
- Phage (with cluster) where this start called: Dryad\_72 (BN), Dryad\_84 (BN),

Gibson\_82 (BN), Greenbelt\_124 (BN), Lizz\_80 (BN), PHTowN\_80 (BN), Rooney\_82

(BN), ShakeNBake\_80 (BN), Wentworth\_79 (BN), Wilkos\_82 (BN), Yara\_78 (BN),

Start 14:

- Found in 2 of 13 (15.4%) of genes in pham
- Manual Annotations of this start: 2 of 11
- Called 100.0% of time when present
- Phage (with cluster) where this start called: Gibson\_68 (BN), Rooney\_68 (BN),

### Summary by clusters:

There is one cluster represented in this pham: BN

Info for manual annotations of cluster BN:Start number 13 was manually annotated 9 times for cluster BN.Start number 14 was manually annotated 2 times for cluster BN.

### Gene Information:

Gene: Dryad\_84 Start: 55433, Stop: 54966, Start Num: 13 Candidate Starts for Dryad\_84: (8, 55472), (Start: 13 @55433 has 9 MA's), (20, 55280),

Gene: Dryad\_72 Start: 48943, Stop: 48380, Start Num: 13 Candidate Starts for Dryad\_72: (Start: 13 @48943 has 9 MA's), (17, 48808), (22, 48613),

Gene: Gibson\_82 Start: 55230, Stop: 54763, Start Num: 13 Candidate Starts for Gibson\_82: (1, 55767), (2, 55743), (3, 55698), (4, 55476), (5, 55404), (9, 55266), (10, 55260), (11, 55254), (Start: 13 @55230 has 9 MA's), (16, 55101), (18, 55089), (20, 55077),

Gene: Gibson\_68 Start: 48192, Stop: 47644, Start Num: 14 Candidate Starts for Gibson\_68: (10, 48231), (11, 48225), (12, 48207), (Start: 14 @48192 has 2 MA's), (15, 48087), (16, 48072), (17, 48066), (21, 48015),

Gene: Greenbelt\_124 Start: 51064, Stop: 50546, Start Num: 13 Candidate Starts for Greenbelt\_124: (Start: 13 @51064 has 9 MA's), (20, 50911),

Gene: Lizz\_80 Start: 54885, Stop: 54421, Start Num: 13 Candidate Starts for Lizz\_80: (6, 55032), (9, 54921), (10, 54915), (11, 54909), (Start: 13 @54885 has 9 MA's), (19, 54738), (20, 54732),

Gene: PHTowN\_80 Start: 54883, Stop: 54419, Start Num: 13 Candidate Starts for PHTowN\_80: (6, 55030), (9, 54919), (10, 54913), (11, 54907), (Start: 13 @54883 has 9 MA's), (19, 54736), (20, 54730),

Gene: Rooney\_82 Start: 55236, Stop: 54769, Start Num: 13

Candidate Starts for Rooney\_82: (6, 55383), (7, 55344), (9, 55272), (10, 55266), (11, 55260), (Start: 13 @55236 has 9 MA's), (16, 55107), (18, 55095), (20, 55083),

Gene: Rooney\_68 Start: 48168, Stop: 47620, Start Num: 14 Candidate Starts for Rooney\_68: (10, 48207), (11, 48201), (12, 48183), (Start: 14 @48168 has 2 MA's), (15, 48063), (16, 48048), (17, 48042), (21, 47991),

Gene: ShakeNBake\_80 Start: 54910, Stop: 54446, Start Num: 13 Candidate Starts for ShakeNBake\_80: (6, 55057), (9, 54946), (10, 54940), (11, 54934), (Start: 13 @54910 has 9 MA's), (19, 54763), (20, 54757),

Gene: Wentworth\_79 Start: 54336, Stop: 53863, Start Num: 13 Candidate Starts for Wentworth\_79: (9, 54372), (10, 54366), (11, 54360), (Start: 13 @54336 has 9 MA's), (20, 54183),

Gene: Wilkos\_82 Start: 54813, Stop: 54361, Start Num: 13 Candidate Starts for Wilkos\_82: (10, 54843), (Start: 13 @54813 has 9 MA's), (19, 54666), (20, 54660),

Gene: Yara\_78 Start: 53975, Stop: 53490, Start Num: 13 Candidate Starts for Yara\_78: (Start: 13 @53975 has 9 MA's), (20, 53825), (23, 53498),