<b>\</b>	<u> </u>	6	·	. 4	. 9		<b>S</b>	~ ~	Ç5	m/ko	10 1
1: Hitter_31											
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o o	S &	6	ę	) ^	, q	5		, , , , , , , , , , , , , , , , , , ,	ζ,	M <sub>P</sub>	,6 ,1 
2: Archis_32											
		6		, 1	٩	5	, ,0	^	\ <sub>0</sub>	M <sub>R</sub> O	,6
3: Pollux_30 + 3											
		6	્	, 1	q	5	<b>S</b>	,^	, <sub>0</sub>	₩ <sub>p</sub>	<i>√</i> 0
4.5.11											
4: Bradis <mark>sa_30</mark>											
		6	ę	,	q	5	, ,	,	, o	"ke	, o
5: Hedwig_32											
, 		6		)	·	<b>S</b>	,0	Λ.	<i>√</i> 6	1,50	<b>,</b> ©
6: Lucky10_27											

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 87682 Report

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

Pham number 87682 has 9 members, 0 are drafts.

Phages represented in each track:

Track 1 : Hitter\_31Track 2 : Archis\_32

Track 3: Pollux\_30, Begonia\_32, Malachai\_32, Floral\_30

Track 4 : Bradissa\_30Track 5 : Hedwig\_32Track 6 : Lucky10 27

## 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 9 of the 9 non-draft genes in the pham.

Genes that call this "Most Annotated" start:

 Archis\_32, Begonia\_32, Bradissa\_30, Floral\_30, Hedwig\_32, Hitter\_31, Lucky10\_27, Malachai\_32, Pollux\_30,

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 3:

- Found in 9 of 9 (100.0%) of genes in pham
- Manual Annotations of this start: 9 of 9
- Called 100.0% of time when present
- Phage (with cluster) where this start called: Archis\_32 (CV), Begonia\_32 (CV), Bradissa\_30 (CY1), Floral\_30 (CY1), Hedwig\_32 (DB), Hitter\_31 (CV), Lucky10\_27 (DH), Malachai\_32 (CV), Pollux\_30 (CY1),

### Summary by clusters:

There are 4 clusters represented in this pham: CY1, DB, CV, DH,

Info for manual annotations of cluster CV:

•Start number 3 was manually annotated 4 times for cluster CV.

Info for manual annotations of cluster CY1:

•Start number 3 was manually annotated 3 times for cluster CY1.

Info for manual annotations of cluster DB:

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

Info for manual annotations of cluster DH:

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

### Gene Information:

Gene: Archis\_32 Start: 27504, Stop: 28439, Start Num: 3

Candidate Starts for Archis 32:

(Start: 3 @27504 has 9 MA's), (4, 27630), (5, 27672), (6, 27777), (7, 27894), (8, 27981), (9, 28098), (11, 28182), (12, 28224), (13, 28257), (14, 28320), (15, 28323), (16, 28374), (17, 28398),

Gene: Begonia\_32 Start: 27738, Stop: 28667, Start Num: 3

Candidate Starts for Begonia\_32:

(Start: 3 @27738 has 9 MA's), (5, 27900), (6, 28005), (7, 28122), (8, 28209), (9, 28326), (10, 28392), (11, 28410), (13, 28485), (14, 28548), (15, 28551), (16, 28602),

Gene: Bradissa\_30 Start: 26559, Stop: 27488, Start Num: 3

Candidate Starts for Bradissa 30:

(Start: 3 @26559 has 9 MA's), (5, 26721), (6, 26826), (7, 26943), (8, 27030), (9, 27147), (11, 27231), (13, 27306), (14, 27369), (15, 27372), (16, 27423),

Gene: Floral 30 Start: 26412, Stop: 27341, Start Num: 3

Candidate Starts for Floral 30:

(Start: 3 @26412 has 9 MA's), (5, 26574), (6, 26679), (7, 26796), (8, 26883), (9, 27000), (10, 27066), (11, 27084), (13, 27159), (14, 27222), (15, 27225), (16, 27276),

Gene: Hedwig 32 Start: 27130, Stop: 28059, Start Num: 3

Candidate Starts for Hedwig\_32:

(2, 27076), (Start: 3 @27130 has 9 MA's), (5, 27292), (6, 27397), (7, 27514), (8, 27601), (9, 27718), (10, 27784), (11, 27802), (13, 27877), (14, 27940), (15, 27943), (16, 27994),

Gene: Hitter\_31 Start: 27134, Stop: 28063, Start Num: 3

Candidate Starts for Hitter 31:

(1, 27008), (Start: 3 @27134 has 9 MA's), (4, 27254), (5, 27296), (6, 27401), (7, 27518), (8, 27605), (9, 27722), (11, 27806), (12, 27848), (13, 27881), (14, 27944), (15, 27947), (16, 27998), (17, 28022),

Gene: Lucky10 27 Start: 23886, Stop: 24815, Start Num: 3

Candidate Starts for Luckv10 27:

(Start: 3 @23886 has 9 MA's), (5, 24048), (6, 24153), (7, 24270), (8, 24357), (9, 24474), (10, 24540), (11, 24558), (13, 24633), (15, 24699), (16, 24750),

Gene: Malachai\_32 Start: 27738, Stop: 28667, Start Num: 3

Candidate Starts for Malachai\_32:

(Start: 3 @27738 has 9 MA's), (5, 27900), (6, 28005), (7, 28122), (8, 28209), (9, 28326), (10, 28392), (11, 28410), (13, 28485), (14, 28548), (15, 28551), (16, 28602),

Gene: Pollux\_30 Start: 26412, Stop: 27341, Start Num: 3

Candidate Starts for Pollux\_30:

(Start: 3 @26412 has 9 MA's), (5, 26574), (6, 26679), (7, 26796), (8, 26883), (9, 27000), (10, 27066), (11, 27084), (13, 27159), (14, 27222), (15, 27225), (16, 27276),