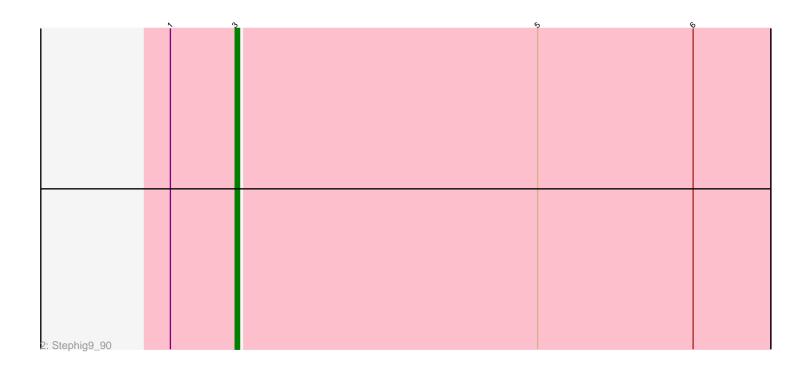
	<b>D</b>	×	•	
1: Roary_92 + 6				



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 106977 Report

This analysis was run 04/05/24 on database version 557.

Pham number 106977 has 8 members, 0 are drafts.

Phages represented in each track:

• Track 1: Roary\_92, Dixon\_89, Phillis\_87, NearlyHeadless\_91, Danforth\_91,

Expelliarmus\_87, Smeadley\_90

Track 2 : Stephig9\_90

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

The start number called the most often in the published annotations is 2, it was called in 7 of the 8 non-draft genes in the pham.

Genes that call this "Most Annotated" start:

• Danforth\_91, Dixon\_89, Expelliarmus\_87, NearlyHeadless\_91, Phillis\_87, Roary\_92, Smeadley\_90,

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

•

Genes that do not have the "Most Annotated" start:

Stephig9\_90,

### Summary by start number:

#### Start 2:

- Found in 7 of 8 (87.5%) of genes in pham
- Manual Annotations of this start: 7 of 8
- Called 100.0% of time when present
- Phage (with cluster) where this start called: Danforth\_91 (A8), Dixon\_89 (A8), Expelliarmus\_87 (A8), NearlyHeadless\_91 (A8), Phillis\_87 (A8), Roary\_92 (A8), Smeadley\_90 (A8),

#### Start 3:

- Found in 1 of 8 (12.5%) of genes in pham
- Manual Annotations of this start: 1 of 8
- Called 100.0% of time when present
- Phage (with cluster) where this start called: Stephig9 90 (A8).

## Summary by clusters:

There is one cluster represented in this pham: A8

Info for manual annotations of cluster A8:

- •Start number 2 was manually annotated 7 times for cluster A8.
- •Start number 3 was manually annotated 1 time for cluster A8.

#### Gene Information:

Gene: Danforth\_91 Start: 47917, Stop: 47792, Start Num: 2

Candidate Starts for Danforth\_91:

(Start: 2 @47917 has 7 MA's), (4, 47887), (5, 47845), (6, 47809),

Gene: Dixon\_89 Start: 47322, Stop: 47197, Start Num: 2

Candidate Starts for Dixon\_89:

(Start: 2 @47322 has 7 MA's), (4, 47292), (5, 47250), (6, 47214),

Gene: Expelliarmus\_87 Start: 47536, Stop: 47411, Start Num: 2

Candidate Starts for Expelliarmus\_87:

(Start: 2 @ 47536 has 7 MA's), (4, 47506), (5, 47464), (6, 47428),

Gene: NearlyHeadless\_91 Start: 47654, Stop: 47529, Start Num: 2

Candidate Starts for NearlyHeadless 91:

(Start: 2 @ 47654 has 7 MA's), (4, 47624), (5, 47582), (6, 47546),

Gene: Phillis\_87 Start: 46667, Stop: 46542, Start Num: 2

Candidate Starts for Phillis\_87:

(Start: 2 @ 46667 has 7 MA's), (4, 46637), (5, 46595), (6, 46559),

Gene: Roary\_92 Start: 47794, Stop: 47669, Start Num: 2

Candidate Starts for Roary 92:

(Start: 2 @ 47794 has 7 MA's), (4, 47764), (5, 47722), (6, 47686),

Gene: Smeadley\_90 Start: 47658, Stop: 47533, Start Num: 2

Candidate Starts for Smeadley 90:

(Start: 2 @ 47658 has 7 MA's), (4, 47628), (5, 47586), (6, 47550),

Gene: Stephig9\_90 Start: 47819, Stop: 47697, Start Num: 3

Candidate Starts for Stephig9 90:

(1, 47834), (Start: 3 @ 47819 has 1 MA's), (5, 47750), (6, 47714),