

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 6512 Report

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

Pham number 6512 has 9 members, 1 are drafts.

Phages represented in each track:

Track 1: PHTowN\_52, Rooney\_52, Gibson\_52

Track 2 : ShakeNBake\_52, Lizz\_52

Track 3 : Wilkos\_55Track 4 : Dryad\_57Track 5 : Yara\_47

• Track 6 : Wentworth\_53

## 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 8 of the 8 non-draft genes in the pham.

Genes that call this "Most Annotated" start:

• Dryad\_57, Gibson\_52, Lizz\_52, PHTowN\_52, Rooney\_52, ShakeNBake\_52, Wentworth\_53, Wilkos\_55, Yara\_47,

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:

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# Summary by start number:

#### Start 3:

- Found in 9 of 9 (100.0%) of genes in pham
- Manual Annotations of this start: 8 of 8
- Called 100.0% of time when present
- Phage (with cluster) where this start called: Dryad\_57 (BN), Gibson\_52 (BN), Lizz\_52 (BN), PHTowN\_52 (BN), Rooney\_52 (BN), ShakeNBake\_52 (BN), Wentworth\_53 (BN), Wilkos\_55 (BN), Yara\_47 (BN),

## Summary by clusters:

There is one cluster represented in this pham: BN

Info for manual annotations of cluster BN:

•Start number 3 was manually annotated 8 times for cluster BN.

### Gene Information:

Gene: Dryad 57 Start: 40277, Stop: 40014, Start Num: 3

Candidate Starts for Dryad\_57:

(1, 40565), (2, 40481), (Start: 3 @40277 has 8 MA's), (5, 40241), (8, 40208), (10, 40166), (12, 40130), (18, 40055), (19, 40034),

Gene: Gibson\_52 Start: 39314, Stop: 39051, Start Num: 3

Candidate Starts for Gibson 52:

(Start: 3 @39314 has 8 MA's), (6, 39275), (8, 39245), (10, 39203), (12, 39167), (13, 39152), (16, 39125), (18, 39092), (19, 39071),

Gene: Lizz\_52 Start: 39124, Stop: 38861, Start Num: 3

Candidate Starts for Lizz 52:

(Start: 3 @39124 has 8 MA's), (7, 39073), (8, 39055), (10, 39013), (12, 38977), (17, 38929), (18, 38902), (19, 38881),

Gene: PHTowN\_52 Start: 39122, Stop: 38859, Start Num: 3

Candidate Starts for PHTowN 52:

(Start: 3 @39122 has 8 MA's), (6, 39083), (8, 39053), (10, 39011), (12, 38975), (13, 38960), (16, 38933), (18, 38900), (19, 38879),

Gene: Rooney\_52 Start: 39311, Stop: 39048, Start Num: 3

Candidate Starts for Rooney\_52:

(Start: 3 @39311 has 8 MA's), (6, 39272), (8, 39242), (10, 39200), (12, 39164), (13, 39149), (16, 39122), (18, 39089), (19, 39068),

Gene: ShakeNBake 52 Start: 39140, Stop: 38877, Start Num: 3

Candidate Starts for ShakeNBake\_52:

(Start: 3 @39140 has 8 MA's), (7, 39089), (8, 39071), (10, 39029), (12, 38993), (17, 38945), (18, 38918), (19, 38897),

Gene: Wentworth 53 Start: 39540, Stop: 39277, Start Num: 3

Candidate Starts for Wentworth 53:

(Start: 3 @39540 has 8 MA's), (6, 39501), (8, 39471), (10, 39429), (12, 39393), (13, 39378), (16, 39351), (18, 39318), (19, 39297),

Gene: Wilkos\_55 Start: 39396, Stop: 39133, Start Num: 3

Candidate Starts for Wilkos\_55:

(Start: 3 @39396 has 8 MA's), (5, 39360), (6, 39357), (8, 39327), (10, 39285), (12, 39249), (13, 39234), (18, 39174), (19, 39153),

Gene: Yara 47 Start: 37858, Stop: 37589, Start Num: 3

Candidate Starts for Yara 47:

(Start: 3 @37858 has 8 MA's), (4, 37846), (9, 37759), (10, 37744), (11, 37714), (14, 37681), (15, 37672),