

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 165450 Report

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

Pham number 165450 has 9 members, 1 are drafts.

Phages represented in each track:

Track 1 : Rooney_91, Gibson_91

Track 2 : Wilkos_90Track 3 : Yara 89

• Track 4 : ShakeNBake 90

• Track 5 : Lizz 90, PHTowN 90

Track 6 : Dryad_93Track 7 : Wentworth 90

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

The start number called the most often in the published annotations is 1, it was called in 7 of the 8 non-draft genes in the pham.

Genes that call this "Most Annotated" start:

• Dryad_93, Gibson_91, Lizz_90, PHTowN_90, Rooney_91, ShakeNBake_90, Wentworth_90, Wilkos_90,

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

• Yara_89,

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

Summary by start number:

Start 1:

- Found in 8 of 9 (88.9%) 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: Dryad_93 (BN), Gibson_91 (BN), Lizz_90 (BN), PHTowN_90 (BN), Rooney_91 (BN), ShakeNBake_90 (BN), Wentworth_90 (BN), Wilkos_90 (BN),

Start 2:

- Found in 1 of 9 (11.1%) 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: Yara_89 (BN),

Summary by clusters:

There is one cluster represented in this pham: BN

Info for manual annotations of cluster BN:

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

Gene Information:

Gene: Dryad 93 Start: 59282, Stop: 58686, Start Num: 1

Candidate Starts for Dryad_93:

(Start: 1 @59282 has 7 MA's), (3, 59171), (4, 59144), (5, 59132), (6, 58970), (11, 58910), (15, 58784), (16, 58769), (17, 58730),

Gene: Gibson_91 Start: 59251, Stop: 58649, Start Num: 1

Candidate Starts for Gibson_91:

(Start: 1 @59251 has 7 MA's), (5, 59101), (11, 58873), (13, 58846), (14, 58813), (16, 58732), (17, 58693),

Gene: Lizz_90 Start: 59245, Stop: 58643, Start Num: 1

Candidate Starts for Lizz_90:

(Start: 1 @59245 has 7 MA's), (3, 59131), (5, 59092), (11, 58867), (13, 58840), (14, 58807), (16, 58726), (17, 58687),

Gene: PHTowN 90 Start: 59243, Stop: 58641, Start Num: 1

Candidate Starts for PHTowN 90:

(Start: 1 @59243 has 7 MA's), (3, 59129), (5, 59090), (11, 58865), (13, 58838), (14, 58805), (16, 58724), (17, 58685),

Gene: Rooney_91 Start: 59259, Stop: 58657, Start Num: 1

Candidate Starts for Roonev 91:

(Start: 1 @59259 has 7 MA's), (5, 59109), (11, 58881), (13, 58854), (14, 58821), (16, 58740), (17, 58701),

Gene: ShakeNBake 90 Start: 59266, Stop: 58664, Start Num: 1

Candidate Starts for ShakeNBake_90:

(Start: 1 @59266 has 7 MA's), (3, 59152), (5, 59113), (11, 58888), (13, 58861), (14, 58828), (16, 58747), (17, 58708),

Gene: Wentworth 90 Start: 58648, Stop: 58046, Start Num: 1

Candidate Starts for Wentworth 90:

(Start: 1 @58648 has 7 MA's), (3, 58534), (5, 58495), (6, 58321), (16, 58129), (17, 58090),

Gene: Wilkos_90 Start: 58442, Stop: 57849, Start Num: 1

Candidate Starts for Wilkos_90:

(Start: 1 @58442 has 7 MA's), (3, 58328), (4, 58301), (5, 58289), (6, 58115), (7, 58088), (9, 58073), (10, 58067), (14, 58004), (18, 57881),

Gene: Yara_89 Start: 59333, Stop: 58701, Start Num: 2

Candidate Starts for Yara_89:

(Start: 2 @59333 has 1 MA's), (5, 59186), (8, 58949), (12, 58913), (13, 58898), (14, 58865), (16,

58784), (17, 58745),