

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 214886 Report

This analysis was run 02/22/25 on database version 588.

Pham number 214886 has 10 members, 2 are drafts.

Phages represented in each track:

Track 1 : Yara_3Track 2 : Wilkos 3

Track 3: Rooney_4, Gibson_4

Track 4 : ShakeNBake_4, PHTowN_4, Lizz_4

Track 5 : Greenbelt_6Track 6 : Wentworth_4

Track 7 : Dryad_5

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

The start number called the most often in the published annotations is 6, it was called in 8 of the 8 non-draft genes in the pham.

Genes that call this "Most Annotated" start:

• Dryad_5, Gibson_4, Lizz_4, PHTowN_4, Rooney_4, ShakeNBake_4, Wentworth_4, Wilkos_3, Yara_3,

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

Greenbelt 6,

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

•

Summary by start number:

Start 3:

- Found in 1 of 10 (10.0%) of genes in pham
- No Manual Annotations of this start.
- Called 100.0% of time when present
- Phage (with cluster) where this start called: Greenbelt_6 (BN),

Start 6:

- Found in 10 of 10 (100.0%) of genes in pham
- Manual Annotations of this start: 8 of 8

- Called 90.0% of time when present
- Phage (with cluster) where this start called: Dryad_5 (BN), Gibson_4 (BN), Lizz_4 (BN), PHTowN_4 (BN), Rooney_4 (BN), ShakeNBake_4 (BN), Wentworth_4 (BN), Wilkos 3 (BN), Yara 3 (BN),

Summary by clusters:

There is one cluster represented in this pham: BN

Info for manual annotations of cluster BN:

Gene: Yara_3 Start: 1042, Stop: 1212, Start Num: 6

Start number 6 was manually annotated 8 times for cluster BN.

Gene Information: Gene: Dryad 5 Start: 1621, Stop: 1812, Start Num: 6 Candidate Starts for Dryad 5: (4, 1612), (Start: 6 @1621 has 8 MA's), (13, 1714), Gene: Gibson 4 Start: 1493, Stop: 1657, Start Num: 6 Candidate Starts for Gibson 4: (4, 1484), (5, 1487), (Start: 6 @1493 has 8 MA's), (7, 1499), (8, 1508), (11, 1544), (12, 1571), Gene: Greenbelt_6 Start: 1823, Stop: 1978, Start Num: 3 Candidate Starts for Greenbelt 6: (3, 1823), (Start: 6 @1838 has 8 MA's), (9, 1859), (10, 1880), (11, 1889), Gene: Lizz 4 Start: 1503, Stop: 1667, Start Num: 6 Candidate Starts for Lizz 4: (4, 1494), (5, 1497), (Start: 6 @ 1503 has 8 MA's), (7, 1509), (8, 1518), (11, 1554), Gene: PHTowN 4 Start: 1503, Stop: 1667, Start Num: 6 Candidate Starts for PHTowN 4: (4, 1494), (5, 1497), (Start: 6 @1503 has 8 MA's), (7, 1509), (8, 1518), (11, 1554), Gene: Rooney 4 Start: 1493, Stop: 1657, Start Num: 6 Candidate Starts for Rooney 4: (4, 1484), (5, 1487), (Start: 6 @ 1493 has 8 MA's), (7, 1499), (8, 1508), (11, 1544), (12, 1571), Gene: ShakeNBake 4 Start: 1503, Stop: 1667, Start Num: 6 Candidate Starts for ShakeNBake 4: (4, 1494), (5, 1497), (Start: 6 @1503 has 8 MA's), (7, 1509), (8, 1518), (11, 1554), Gene: Wentworth 4 Start: 1225, Stop: 1389, Start Num: 6 Candidate Starts for Wentworth 4: (1, 1174), (2, 1204), (4, 1216), (Start: 6 @1225 has 8 MA's), (7, 1231), (11, 1276), Gene: Wilkos 3 Start: 1485, Stop: 1649, Start Num: 6 Candidate Starts for Wilkos 3: (2, 1464), (4, 1476), (Start: 6 @1485 has 8 MA's), (8, 1500), (11, 1536), (12, 1563), (14, 1605), Candidate Starts for Yara_3: (Start: 6 @1042 has 8 MA's), (11, 1096), (13, 1138),