Pham 6886


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 6886 Report

This analysis was run 04/05/24 on database version 557.
Pham number 6886 has 8 members, 1 are drafts.
Phages represented in each track:

- Track 1 : PHTowN_41, ShakeNBake_41, Lizz_41
- Track 2 : Rooney $\overline{4} 1$
- Track 3 : Dryad_45
- Track 4 : Wentworth 42
- Track 5 : Wilkos 42
- Track 6 : Gibson_41


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

The start number called the most often in the published annotations is 7, it was called in 5 of the 7 non-draft genes in the pham.

Genes that call this "Most Annotated" start:

- Dryad_45, Lizz_41, PHTowN_41, Rooney_41, ShakeNBake_41,

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

- Gibson_41, Wentworth_42, Wilkos_42,

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

## Summary by start number:

Start 4:

- Found in 5 of 8 ( 62.5\% ) of genes in pham
- Manual Annotations of this start: 1 of 7
- Called $40.0 \%$ of time when present
- Phage (with cluster) where this start called: Wentworth_42 (BN), Wilkos_42 (BN),

Start 5:

- Found in 3 of 8 ( $37.5 \%$ ) of genes in pham
- Manual Annotations of this start: 1 of 7
- Called $33.3 \%$ of time when present
- Phage (with cluster) where this start called: Gibson_41 (BN),

Start 7:

- Found in 8 of 8 ( $100.0 \%$ ) of genes in pham
- Manual Annotations of this start: 5 of 7
- Called $62.5 \%$ of time when present
- Phage (with cluster) where this start called: Dryad_45 (BN), Lizz_41 (BN),

PHTowN_41 (BN), Rooney_41 (BN), ShakeNBake_41 (BN),

## Summary by clusters:

There is one cluster represented in this pham: BN
Info for manual annotations of cluster BN:

- Start number 4 was manually annotated 1 time for cluster BN.
- Start number 5 was manually annotated 1 time for cluster BN.
-Start number 7 was manually annotated 5 times for cluster BN.


## Gene Information:

Gene: Dryad_45 Start: 32679, Stop: 32527, Start Num: 7
Candidate Starts for Dryad_45:
(1, 32772), (3, 32745), (Start: 5 @32715 has 1 MA's), (Start: 7 @32679 has 5 MA's), (9, 32592),
Gene: Gibson_41 Start: 32230, Stop: 32036, Start Num: 5
Candidate Starts for Gibson_41:
(Start: 5 @32230 has 1 MA's), (Start: 7 @32188 has 5 MA's), (8, 32137),
Gene: Lizz_41 Start: 32002, Stop: 31850, Start Num: 7
Candidate Starts for Lizz_41:
(1, 32101), (Start: 4 @32050 has 1 MA's), (6, 32041), (Start: 7 @32002 has 5 MA's), (8, 31951), (9, 31915),

Gene: PHTowN_41 Start: 32000, Stop: 31848, Start Num: 7
Candidate Starts for PHTowN_41:
(1, 32099), (Start: 4 @32048 has 1 MA's), (6, 32039), (Start: 7 @32000 has 5 MA's), (8, 31949), (9, 31913),

Gene: Rooney_41 Start: 32185, Stop: 32033, Start Num: 7
Candidate Starts for Rooney_41:
(Start: 5 @32227 has 1 MA's), (Start: 7 @32185 has 5 MA's), (8, 32134),
Gene: ShakeNBake_41 Start: 32018, Stop: 31866, Start Num: 7
Candidate Starts for ShakeNBake_41:
(1, 32117), (Start: 4 @32066 has 1 MA's), (6, 32057), (Start: 7 @32018 has 5 MA's), (8, 31967), (9, 31931),

Gene: Wentworth_42 Start: 32535, Stop: 32335, Start Num: 4
Candidate Starts for Wentworth_42:
(3, 32559), (Start: 4 @32535 has 1 MA's), (6, 32526), (Start: 7 @32487 has 5 MA's), (10, 32352),
Gene: Wilkos_42 Start: 32356, Stop: 32156, Start Num: 4

Candidate Starts for Wilkos_42:
(2, 32398), (3, 32380), (Start: 4 @32356 has 1 MA's), (Start: 7 @32308 has 5 MA's), (9, 32221),

