

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 194642 Report

This analysis was run 11/02/24 on database version 579.

Pham number 194642 has 7 members, 1 are drafts.

Phages represented in each track:

Track 1 : CloverMinnie_57Track 2 : BobRoss2024_58

Track 3: AnarQue_58, Wooper_58, CaiB_58, MakoManhole_55

Track 4 : MossRose_57

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 6 of the 6 non-draft genes in the pham.

Genes that call this "Most Annotated" start:

AnarQue_58, CaiB_58, CloverMinnie_57, MakoManhole_55, MossRose_57, Wooper_58,

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:

BobRoss2024 58.

Summary by start number:

Start 1:

- Found in 1 of 7 (14.3%) of genes in pham
- No Manual Annotations of this start.
- Called 100.0% of time when present
- Phage (with cluster) where this start called: BobRoss2024_58 (DR),

Start 2:

- Found in 6 of 7 (85.7%) of genes in pham
- Manual Annotations of this start: 6 of 6
- Called 100.0% of time when present
- Phage (with cluster) where this start called: AnarQue_58 (DR), CaiB_58 (DR), CloverMinnie_57 (DR), MakoManhole_55 (DR), MossRose_57 (DR), Wooper_58

(DR),

Summary by clusters:

There is one cluster represented in this pham: DR

Info for manual annotations of cluster DR:

•Start number 2 was manually annotated 6 times for cluster DR.

Gene Information:

Gene: AnarQue_58 Start: 49672, Stop: 49292, Start Num: 2

Candidate Starts for AnarQue_58:

(Start: 2 @49672 has 6 MA's), (4, 49558), (6, 49516), (8, 49417), (9, 49411), (10, 49402), (11, 49342), (12, 49327),

Gene: BobRoss2024_58 Start: 48643, Stop: 48239, Start Num: 1

Candidate Starts for BobRoss2024_58:

(1, 48643), (3, 48583), (5, 48481), (6, 48463), (7, 48415), (10, 48352), (12, 48274), (13, 48259), (14, 48256), (15, 48253),

Gene: CaiB_58 Start: 50226, Stop: 49846, Start Num: 2

Candidate Starts for CaiB_58:

(Start: 2 @50226 has 6 MA's), (4, 50112), (6, 50070), (8, 49971), (9, 49965), (10, 49956), (11, 49896), (12, 49881),

Gene: CloverMinnie_57 Start: 49258, Stop: 48878, Start Num: 2

Candidate Starts for CloverMinnie_57:

(Start: 2 @49258 has 6 MA's), (4, 49144), (6, 49102), (11, 48928), (12, 48913), (15, 48892),

Gene: MakoManhole 55 Start: 48750, Stop: 48370, Start Num: 2

Candidate Starts for MakoManhole 55:

(Start: 2 @48750 has 6 MA's), (4, 48636), (6, 48594), (8, 48495), (9, 48489), (10, 48480), (11, 48420), (12, 48405),

Gene: MossRose 57 Start: 49303, Stop: 48923, Start Num: 2

Candidate Starts for MossRose 57:

(Start: 2 @49303 has 6 MA's), (4, 49189), (6, 49147), (8, 49048), (9, 49042), (10, 49033), (11, 48973), (12, 48958), (15, 48937),

Gene: Wooper 58 Start: 49644, Stop: 49264, Start Num: 2

Candidate Starts for Wooper_58:

(Start: 2 @49644 has 6 MA's), (4, 49530), (6, 49488), (8, 49389), (9, 49383), (10, 49374), (11, 49314), (12, 49299),