



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 278882 Report

This analysis was run 02/07/26 on database version 634.

Pham number 278882 has 9 members, 3 are drafts.

Phages represented in each track:

- Track 1 : FruityLoops_77, UBSmoodge_80
- Track 2 : Schomber_74, Gray_76, Kabocha_77, Hanem_75
- Track 3 : ChisanaKitsune_73
- Track 4 : Lenoshki_76, Beted_76

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

The start number called the most often in the published annotations is 5, it was called in 6 of the 6 non-draft genes in the pham.

Genes that call this "Most Annotated" start:

- Beted_76, ChisanaKitsune_73, FruityLoops_77, Gray_76, Hanem_75, Kabocha_77, Lenoshki_76, Schomber_74, UBSmoodge_80,

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

-

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

-

Summary by start number:

Start 5:

- Found in 9 of 9 (100.0%) 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: Beted_76 (DQ), ChisanaKitsune_73 (DQ), FruityLoops_77 (DQ), Gray_76 (DQ), Hanem_75 (DQ), Kabocha_77 (DQ), Lenoshki_76 (DQ), Schomber_74 (DQ), UBSmoodge_80 (DQ),

Summary by clusters:

There is one cluster represented in this pham: DQ

Info for manual annotations of cluster DQ:

•Start number 5 was manually annotated 6 times for cluster DQ.

Gene Information:

Gene: Beted_76 Start: 62884, Stop: 63267, Start Num: 5

Candidate Starts for Beted_76:

(1, 62791), (3, 62818), (Start: 5 @62884 has 6 MA's), (8, 63067),

Gene: ChisanaKitsune_73 Start: 60428, Stop: 60832, Start Num: 5

Candidate Starts for ChisanaKitsune_73:

(1, 60335), (3, 60362), (Start: 5 @60428 has 6 MA's),

Gene: FruityLoops_77 Start: 64804, Stop: 65106, Start Num: 5

Candidate Starts for FruityLoops_77:

(2, 64732), (4, 64744), (Start: 5 @64804 has 6 MA's), (6, 64852), (7, 64894), (9, 65071),

Gene: Gray_76 Start: 61273, Stop: 61656, Start Num: 5

Candidate Starts for Gray_76:

(1, 61180), (3, 61207), (Start: 5 @61273 has 6 MA's),

Gene: Hanem_75 Start: 60758, Stop: 61162, Start Num: 5

Candidate Starts for Hanem_75:

(1, 60665), (3, 60692), (Start: 5 @60758 has 6 MA's),

Gene: Kabocha_77 Start: 62101, Stop: 62505, Start Num: 5

Candidate Starts for Kabocha_77:

(1, 62008), (3, 62035), (Start: 5 @62101 has 6 MA's),

Gene: Lenoshki_76 Start: 62884, Stop: 63267, Start Num: 5

Candidate Starts for Lenoshki_76:

(1, 62791), (3, 62818), (Start: 5 @62884 has 6 MA's), (8, 63067),

Gene: Schomber_74 Start: 60492, Stop: 60896, Start Num: 5

Candidate Starts for Schomber_74:

(1, 60399), (3, 60426), (Start: 5 @60492 has 6 MA's),

Gene: UBSmoodge_80 Start: 65247, Stop: 65549, Start Num: 5

Candidate Starts for UBSmoodge_80:

(2, 65175), (4, 65187), (Start: 5 @65247 has 6 MA's), (6, 65295), (7, 65337), (9, 65514),