



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 107022 Report

This analysis was run 04/05/24 on database version 557.

Pham number 107022 has 7 members, 0 are drafts.

Phages represented in each track:

- Track 1 : Panchino_36, Andies_37, Snekmaggedon_37, Jamie19_37, SpongeBob_37, Shweta_37, Phrann_41

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 7 non-draft genes in the pham.

Genes that call this "Most Annotated" start:

- Andies_37, Jamie19_37, Panchino_36, Phrann_41, Shweta_37, Snekmaggedon_37, SpongeBob_37,

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 1:

- Found in 7 of 7 (100.0%) of genes in pham
- Manual Annotations of this start: 7 of 7
- Called 100.0% of time when present
- Phage (with cluster) where this start called: Andies_37 (N), Jamie19_37 (N), Panchino_36 (N), Phrann_41 (N), Shweta_37 (N), Snekmaggedon_37 (N), SpongeBob_37 (N),

Summary by clusters:

There is one cluster represented in this pham: N

Info for manual annotations of cluster N:

- Start number 1 was manually annotated 7 times for cluster N.

Gene Information:

Gene: Andies_37 Start: 28992, Stop: 29165, Start Num: 1

Candidate Starts for Andies_37:

(Start: 1 @28992 has 7 MA's), (2, 29004), (3, 29007), (4, 29025), (5, 29034), (6, 29046), (7, 29073), (8, 29097), (9, 29115), (10, 29121), (11, 29142),

Gene: Jamie19_37 Start: 28873, Stop: 29046, Start Num: 1

Candidate Starts for Jamie19_37:

(Start: 1 @28873 has 7 MA's), (2, 28885), (3, 28888), (4, 28906), (5, 28915), (6, 28927), (7, 28954), (8, 28978), (9, 28996), (10, 29002), (11, 29023),

Gene: Panchino_36 Start: 30219, Stop: 30392, Start Num: 1

Candidate Starts for Panchino_36:

(Start: 1 @30219 has 7 MA's), (2, 30231), (3, 30234), (4, 30252), (5, 30261), (6, 30273), (7, 30300), (8, 30324), (9, 30342), (10, 30348), (11, 30369),

Gene: Phrann_41 Start: 30899, Stop: 31072, Start Num: 1

Candidate Starts for Phrann_41:

(Start: 1 @30899 has 7 MA's), (2, 30911), (3, 30914), (4, 30932), (5, 30941), (6, 30953), (7, 30980), (8, 31004), (9, 31022), (10, 31028), (11, 31049),

Gene: Shweta_37 Start: 29003, Stop: 29176, Start Num: 1

Candidate Starts for Shweta_37:

(Start: 1 @29003 has 7 MA's), (2, 29015), (3, 29018), (4, 29036), (5, 29045), (6, 29057), (7, 29084), (8, 29108), (9, 29126), (10, 29132), (11, 29153),

Gene: Snekmaggedon_37 Start: 28873, Stop: 29046, Start Num: 1

Candidate Starts for Snekmaggedon_37:

(Start: 1 @28873 has 7 MA's), (2, 28885), (3, 28888), (4, 28906), (5, 28915), (6, 28927), (7, 28954), (8, 28978), (9, 28996), (10, 29002), (11, 29023),

Gene: SpongeBob_37 Start: 28873, Stop: 29046, Start Num: 1

Candidate Starts for SpongeBob_37:

(Start: 1 @28873 has 7 MA's), (2, 28885), (3, 28888), (4, 28906), (5, 28915), (6, 28927), (7, 28954), (8, 28978), (9, 28996), (10, 29002), (11, 29023),