

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 293399 Report

This analysis was run 04/18/26 on database version 643.

Pham number 293399 has 9 members, 3 are drafts.

Phages represented in each track:

- Track 1 : Talos_9
- Track 2 : LuckySocke_8, Fudan_9, Alone3_9
- Track 3 : GiJojo_7
- Track 4 : Beleetus_8
- Track 5 : BRock_5
- Track 6 : Park1214_9
- Track 7 : Garnacho_7

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

Genes that call this "Most Annotated" start:

- Alone3_9, BRock_5, Beleetus_8, Fudan_9, Garnacho_7, GiJojo_7, LuckySocke_8, Park1214_9, Talos_9,

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 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: Alone3_9 (BS), BRock_5 (BS), Beleetus_8 (BS), Fudan_9 (BS), Garnacho_7 (BS), GiJojo_7 (BS), LuckySocke_8 (BS), Park1214_9 (BS), Talos_9 (BS),

Summary by clusters:

There is one cluster represented in this pham: BS

Info for manual annotations of cluster BS:

•Start number 1 was manually annotated 6 times for cluster BS.

Gene Information:

Gene: Alone3_9 Start: 3507, Stop: 3716, Start Num: 1

Candidate Starts for Alone3_9:

(Start: 1 @3507 has 6 MA's), (2, 3534), (3, 3549), (4, 3570), (7, 3624),

Gene: BRock_5 Start: 2508, Stop: 2729, Start Num: 1

Candidate Starts for BRock_5:

(Start: 1 @2508 has 6 MA's), (2, 2535), (5, 2613), (6, 2619), (9, 2637), (11, 2688), (13, 2724),

Gene: Beleetus_8 Start: 3177, Stop: 3386, Start Num: 1

Candidate Starts for Beleetus_8:

(Start: 1 @3177 has 6 MA's), (2, 3204), (7, 3294),

Gene: Fudan_9 Start: 3189, Stop: 3398, Start Num: 1

Candidate Starts for Fudan_9:

(Start: 1 @3189 has 6 MA's), (2, 3216), (3, 3231), (4, 3252), (7, 3306),

Gene: Garnacho_7 Start: 3161, Stop: 3370, Start Num: 1

Candidate Starts for Garnacho_7:

(Start: 1 @3161 has 6 MA's), (2, 3188), (3, 3203), (5, 3266), (7, 3278), (8, 3287), (10, 3317),

Gene: GiJojo_7 Start: 3108, Stop: 3317, Start Num: 1

Candidate Starts for GiJojo_7:

(Start: 1 @3108 has 6 MA's), (2, 3135), (4, 3171), (5, 3213), (7, 3225), (8, 3234), (12, 3300),

Gene: LuckySocke_8 Start: 3189, Stop: 3398, Start Num: 1

Candidate Starts for LuckySocke_8:

(Start: 1 @3189 has 6 MA's), (2, 3216), (3, 3231), (4, 3252), (7, 3306),

Gene: Park1214_9 Start: 3212, Stop: 3421, Start Num: 1

Candidate Starts for Park1214_9:

(Start: 1 @3212 has 6 MA's), (2, 3239), (5, 3317), (7, 3329),

Gene: Talos_9 Start: 3400, Stop: 3609, Start Num: 1

Candidate Starts for Talos_9:

(Start: 1 @3400 has 6 MA's), (2, 3427), (5, 3505), (7, 3517), (8, 3526),