



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 281065 Report

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

Pham number 281065 has 12 members, 0 are drafts.

Phages represented in each track:

- Track 1 : Benczkowski14_77, Katyusha_75, Niagara_76, Teech_77, Hollow_77
- Track 2 : Kvothe_75
- Track 3 : Tredge_76, Teatealatte_77
- Track 4 : ASerpRocky_71
- Track 5 : Vitaenoi_74, Philon9_74
- Track 6 : Demosthenes_71

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 12 of the 12 non-draft genes in the pham.

Genes that call this "Most Annotated" start:

- ASerpRocky_71, Benczkowski14_77, Demosthenes_71, Hollow_77, Katyusha_75, Kvothe_75, Niagara_76, Philon9_74, Teatealatte_77, Teech_77, Tredge_76, Vitaenoi_74,

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 2:

- Found in 12 of 12 (100.0%) of genes in pham
- Manual Annotations of this start: 12 of 12
- Called 100.0% of time when present
- Phage (with cluster) where this start called: ASerpRocky_71 (CS4), Benczkowski14_77 (CS4), Demosthenes_71 (CS4), Hollow_77 (CS4), Katyusha_75 (CS4), Kvothe_75 (CS4), Niagara_76 (CS4), Philon9_74 (CS4), Teatealatte_77 (CS4), Teech_77 (CS4), Tredge_76 (CS4), Vitaenoi_74 (CS4),

Summary by clusters:

There is one cluster represented in this pham: CS4

Info for manual annotations of cluster CS4:

•Start number 2 was manually annotated 12 times for cluster CS4.

Gene Information:

Gene: ASerpRocky_71 Start: 61030, Stop: 60557, Start Num: 2

Candidate Starts for ASerpRocky_71:

(Start: 2 @61030 has 12 MA's), (11, 60739), (12, 60733), (15, 60610), (18, 60568),

Gene: Benczkowski14_77 Start: 62275, Stop: 61802, Start Num: 2

Candidate Starts for Benczkowski14_77:

(Start: 2 @62275 has 12 MA's), (3, 62224), (5, 62128), (8, 62041), (9, 62038), (10, 62023), (11, 61984), (12, 61978), (13, 61945), (14, 61906), (16, 61852),

Gene: Demosthenes_71 Start: 61052, Stop: 60579, Start Num: 2

Candidate Starts for Demosthenes_71:

(Start: 2 @61052 has 12 MA's), (10, 60800), (11, 60761), (12, 60755), (13, 60722), (14, 60683), (16, 60629),

Gene: Hollow_77 Start: 62444, Stop: 61971, Start Num: 2

Candidate Starts for Hollow_77:

(Start: 2 @62444 has 12 MA's), (3, 62393), (5, 62297), (8, 62210), (9, 62207), (10, 62192), (11, 62153), (12, 62147), (13, 62114), (14, 62075), (16, 62021),

Gene: Katyusha_75 Start: 62275, Stop: 61802, Start Num: 2

Candidate Starts for Katyusha_75:

(Start: 2 @62275 has 12 MA's), (3, 62224), (5, 62128), (8, 62041), (9, 62038), (10, 62023), (11, 61984), (12, 61978), (13, 61945), (14, 61906), (16, 61852),

Gene: Kvothe_75 Start: 62117, Stop: 61644, Start Num: 2

Candidate Starts for Kvothe_75:

(1, 62180), (Start: 2 @62117 has 12 MA's), (3, 62066), (8, 61883), (9, 61880), (10, 61865), (11, 61826), (12, 61820), (13, 61787), (14, 61748), (16, 61694),

Gene: Niagara_76 Start: 62208, Stop: 61735, Start Num: 2

Candidate Starts for Niagara_76:

(Start: 2 @62208 has 12 MA's), (3, 62157), (5, 62061), (8, 61974), (9, 61971), (10, 61956), (11, 61917), (12, 61911), (13, 61878), (14, 61839), (16, 61785),

Gene: Philon9_74 Start: 62676, Stop: 62200, Start Num: 2

Candidate Starts for Philon9_74:

(1, 62739), (Start: 2 @62676 has 12 MA's), (4, 62610), (6, 62526), (7, 62508), (8, 62442), (16, 62250), (17, 62241), (18, 62211),

Gene: Teatealatte_77 Start: 62105, Stop: 61632, Start Num: 2

Candidate Starts for Teatealatte_77:

(Start: 2 @62105 has 12 MA's), (3, 62054), (8, 61871), (9, 61868), (10, 61853), (11, 61814), (12, 61808), (13, 61775), (14, 61736), (16, 61682),

Gene: Teech_77 Start: 61924, Stop: 61451, Start Num: 2

Candidate Starts for Teech_77:

(Start: 2 @61924 has 12 MA's), (3, 61873), (5, 61777), (8, 61690), (9, 61687), (10, 61672), (11, 61633), (12, 61627), (13, 61594), (14, 61555), (16, 61501),

Gene: Tredge_76 Start: 62105, Stop: 61632, Start Num: 2

Candidate Starts for Tredge_76:

(Start: 2 @62105 has 12 MA's), (3, 62054), (8, 61871), (9, 61868), (10, 61853), (11, 61814), (12, 61808), (13, 61775), (14, 61736), (16, 61682),

Gene: Vitaenoi_74 Start: 62660, Stop: 62184, Start Num: 2

Candidate Starts for Vitaenoi_74:

(1, 62723), (Start: 2 @62660 has 12 MA's), (4, 62594), (6, 62510), (7, 62492), (8, 62426), (16, 62234), (17, 62225), (18, 62195),