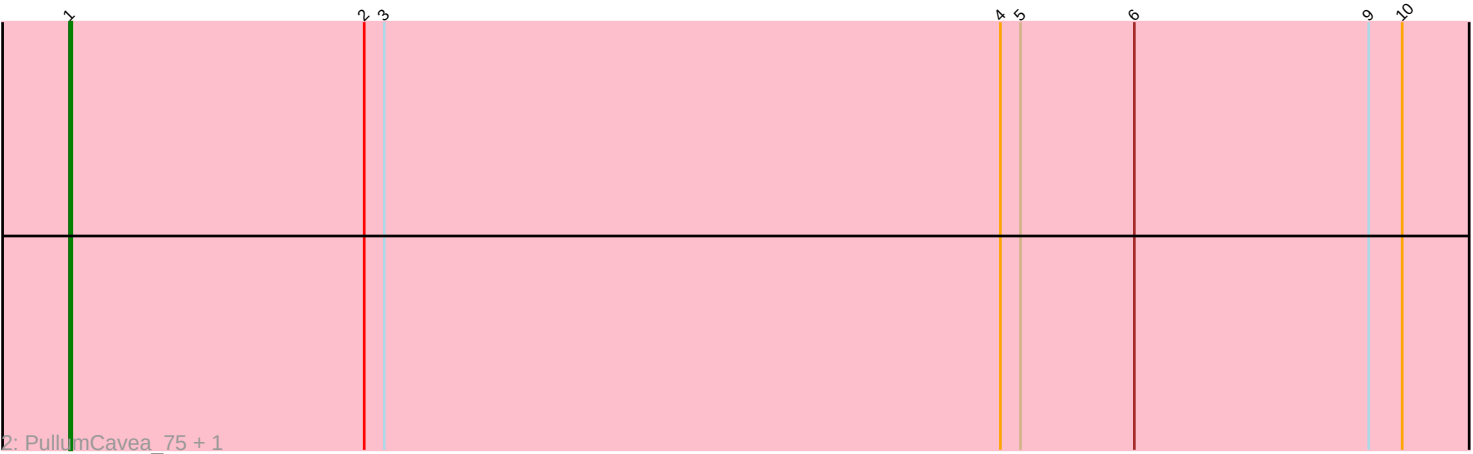




Note: Tracks are now grouped by subcluster and scaled. Switching in subcluster is indicated by changes in track color. Track scale is now set by default to display the region 30 bp upstream of start 1 to 30 bp downstream of the last possible start. If this default region is judged to be packed too tightly 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 281166 Report

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

Pham number 281166 has 9 members, 0 are drafts.

Phages represented in each track:

- Track 1 : Twister6_77, Salvador_76, Togo_76, SmokingBunny_77, Evamon_76
- Track 2 : PullumCavea_75, Phlop_74
- Track 3 : VanDeWege_78, Arri_77

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 9 of the 9 non-draft genes in the pham.

Genes that call this "Most Annotated" start:

- Arri_77, Evamon_76, Phlop_74, PullumCavea_75, Salvador_76, SmokingBunny_77, Togo_76, Twister6_77, VanDeWege_78,

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: 9 of 9
- Called 100.0% of time when present
- Phage (with cluster) where this start called: Arri_77 (DC1), Evamon_76 (DC1), Phlop_74 (DC1), PullumCavea_75 (DC1), Salvador_76 (DC1), SmokingBunny_77 (DC1), Togo_76 (DC1), Twister6_77 (DC1), VanDeWege_78 (DC1),

Summary by clusters:

There is one cluster represented in this pham: DC1

Info for manual annotations of cluster DC1:

- Start number 1 was manually annotated 9 times for cluster DC1.

Gene Information:

Gene: Arri_77 Start: 51472, Stop: 52107, Start Num: 1

Candidate Starts for Arri_77:

(Start: 1 @51472 has 9 MA's), (2, 51604), (3, 51613), (4, 51889), (6, 51949), (7, 52000), (8, 52003),

Gene: Evamon_76 Start: 51207, Stop: 51866, Start Num: 1

Candidate Starts for Evamon_76:

(Start: 1 @51207 has 9 MA's), (2, 51339), (3, 51348), (4, 51624), (5, 51633), (6, 51684), (9, 51789), (10, 51804),

Gene: Phlop_74 Start: 51091, Stop: 51750, Start Num: 1

Candidate Starts for Phlop_74:

(Start: 1 @51091 has 9 MA's), (2, 51223), (3, 51232), (4, 51508), (5, 51517), (6, 51568), (9, 51673), (10, 51688),

Gene: PullumCavea_75 Start: 51268, Stop: 51927, Start Num: 1

Candidate Starts for PullumCavea_75:

(Start: 1 @51268 has 9 MA's), (2, 51400), (3, 51409), (4, 51685), (5, 51694), (6, 51745), (9, 51850), (10, 51865),

Gene: Salvador_76 Start: 51205, Stop: 51864, Start Num: 1

Candidate Starts for Salvador_76:

(Start: 1 @51205 has 9 MA's), (2, 51337), (3, 51346), (4, 51622), (5, 51631), (6, 51682), (9, 51787), (10, 51802),

Gene: SmokingBunny_77 Start: 52201, Stop: 52860, Start Num: 1

Candidate Starts for SmokingBunny_77:

(Start: 1 @52201 has 9 MA's), (2, 52333), (3, 52342), (4, 52618), (5, 52627), (6, 52678), (9, 52783), (10, 52798),

Gene: Togo_76 Start: 52201, Stop: 52860, Start Num: 1

Candidate Starts for Togo_76:

(Start: 1 @52201 has 9 MA's), (2, 52333), (3, 52342), (4, 52618), (5, 52627), (6, 52678), (9, 52783), (10, 52798),

Gene: Twister6_77 Start: 50550, Stop: 51209, Start Num: 1

Candidate Starts for Twister6_77:

(Start: 1 @50550 has 9 MA's), (2, 50682), (3, 50691), (4, 50967), (5, 50976), (6, 51027), (9, 51132), (10, 51147),

Gene: VanDeWege_78 Start: 51535, Stop: 52170, Start Num: 1

Candidate Starts for VanDeWege_78:

(Start: 1 @51535 has 9 MA's), (2, 51667), (3, 51676), (4, 51952), (6, 52012), (7, 52063), (8, 52066),