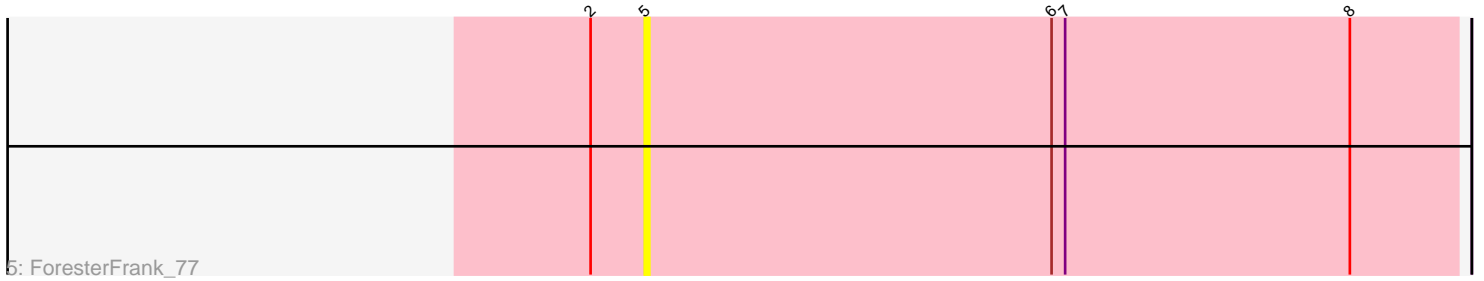
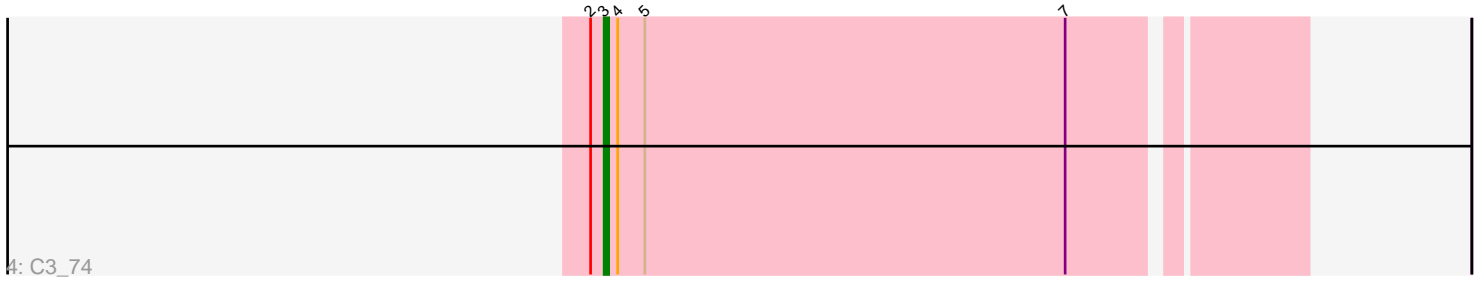
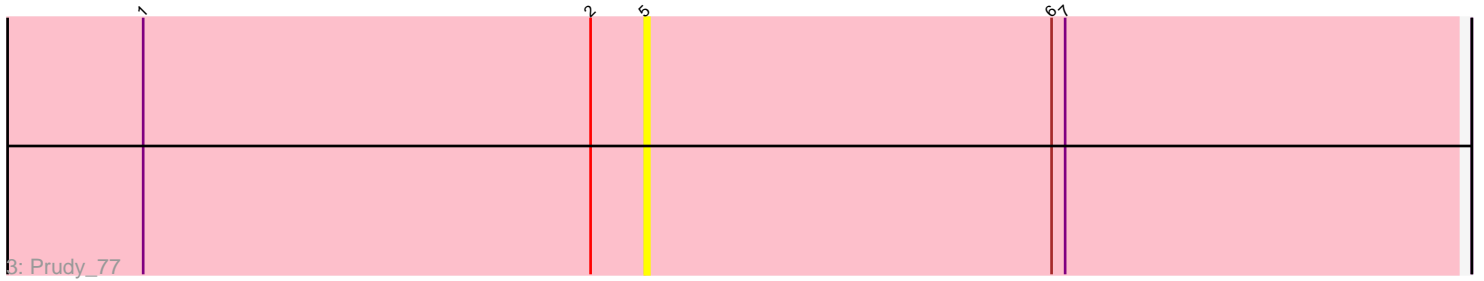
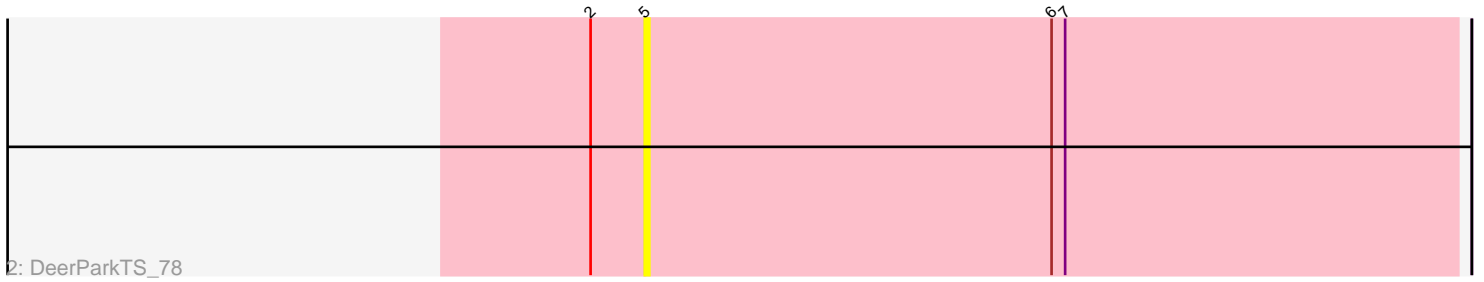
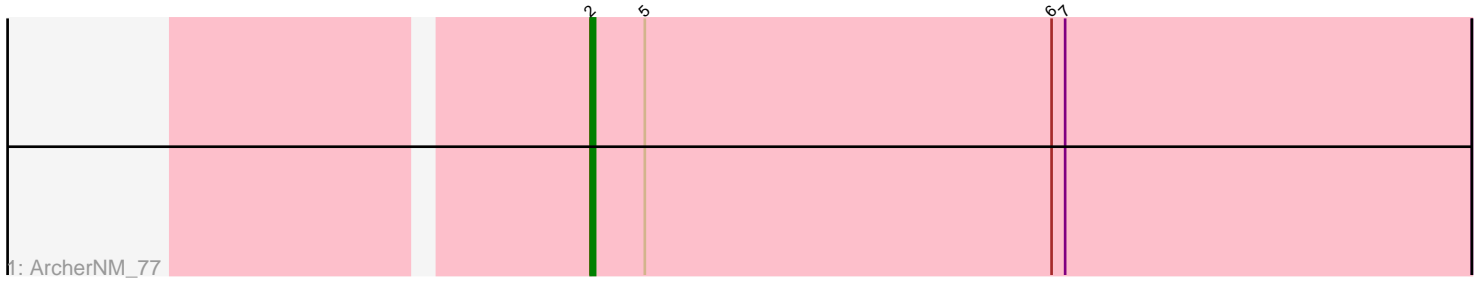


Pham 297426



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 297426 Report

This analysis was run 04/25/26 on database version 644.

Pham number 297426 has 5 members, 3 are drafts.

Phages represented in each track:

- Track 1 : ArcherNM_77
- Track 2 : DeerParkTS_78
- Track 3 : Prudy_77
- Track 4 : C3_74
- Track 5 : ForesterFrank_77

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 1 of the 2 non-draft genes in the pham.

Genes that call this "Most Annotated" start:

- ArcherNM_77,

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

- C3_74, DeerParkTS_78, ForesterFrank_77, Prudy_77,

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

-

Summary by start number:

Start 2:

- Found in 5 of 5 (100.0%) of genes in pham
- Manual Annotations of this start: 1 of 2
- Called 20.0% of time when present
- Phage (with cluster) where this start called: ArcherNM_77 (A2),

Start 3:

- Found in 1 of 5 (20.0%) of genes in pham
- Manual Annotations of this start: 1 of 2
- Called 100.0% of time when present
- Phage (with cluster) where this start called: C3_74 (A2),

Start 5:

- Found in 5 of 5 (100.0%) of genes in pham
- No Manual Annotations of this start.
- Called 60.0% of time when present
- Phage (with cluster) where this start called: DeerParkTS_78 (A2), ForesterFrank_77 (A2), Prudy_77 (A2),

Summary by clusters:

There is one cluster represented in this pham: A2

Info for manual annotations of cluster A2:

- Start number 2 was manually annotated 1 time for cluster A2.
- Start number 3 was manually annotated 1 time for cluster A2.

Gene Information:

Gene: ArcherNM_77 Start: 46958, Stop: 46764, Start Num: 2

Candidate Starts for ArcherNM_77:

(Start: 2 @46958 has 1 MA's), (5, 46946), (6, 46856), (7, 46853),

Gene: C3_74 Start: 45863, Stop: 45714, Start Num: 3

Candidate Starts for C3_74:

(Start: 2 @45866 has 1 MA's), (Start: 3 @45863 has 1 MA's), (4, 45860), (5, 45854), (7, 45761),

Gene: DeerParkTS_78 Start: 45959, Stop: 45780, Start Num: 5

Candidate Starts for DeerParkTS_78:

(Start: 2 @45971 has 1 MA's), (5, 45959), (6, 45869), (7, 45866),

Gene: ForesterFrank_77 Start: 47178, Stop: 46999, Start Num: 5

Candidate Starts for ForesterFrank_77:

(Start: 2 @47190 has 1 MA's), (5, 47178), (6, 47088), (7, 47085), (8, 47022),

Gene: Prudy_77 Start: 47444, Stop: 47265, Start Num: 5

Candidate Starts for Prudy_77:

(1, 47555), (Start: 2 @47456 has 1 MA's), (5, 47444), (6, 47354), (7, 47351),