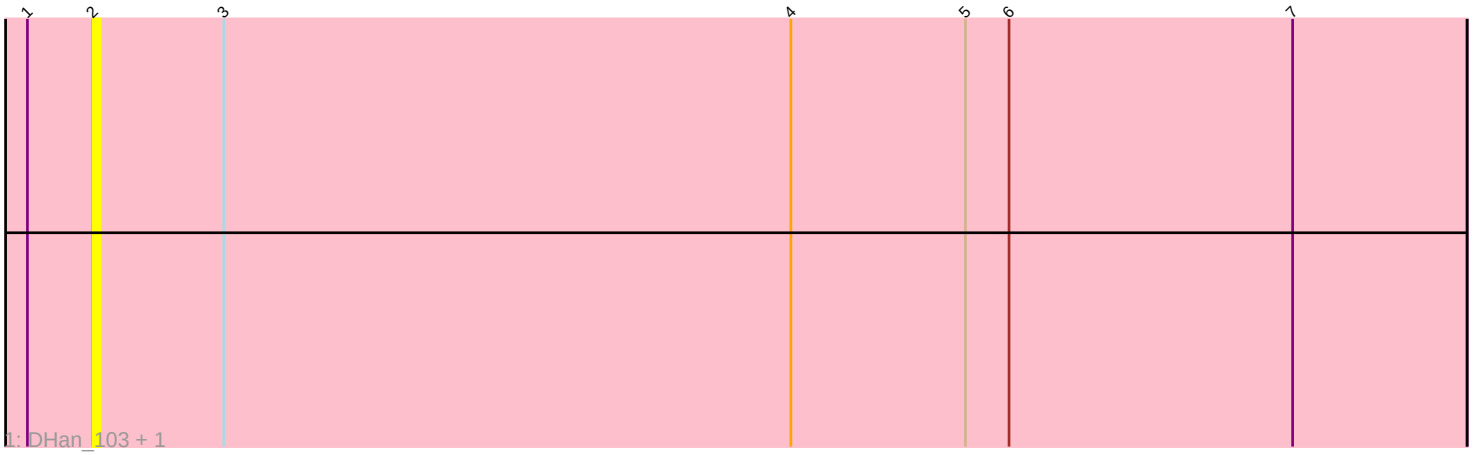
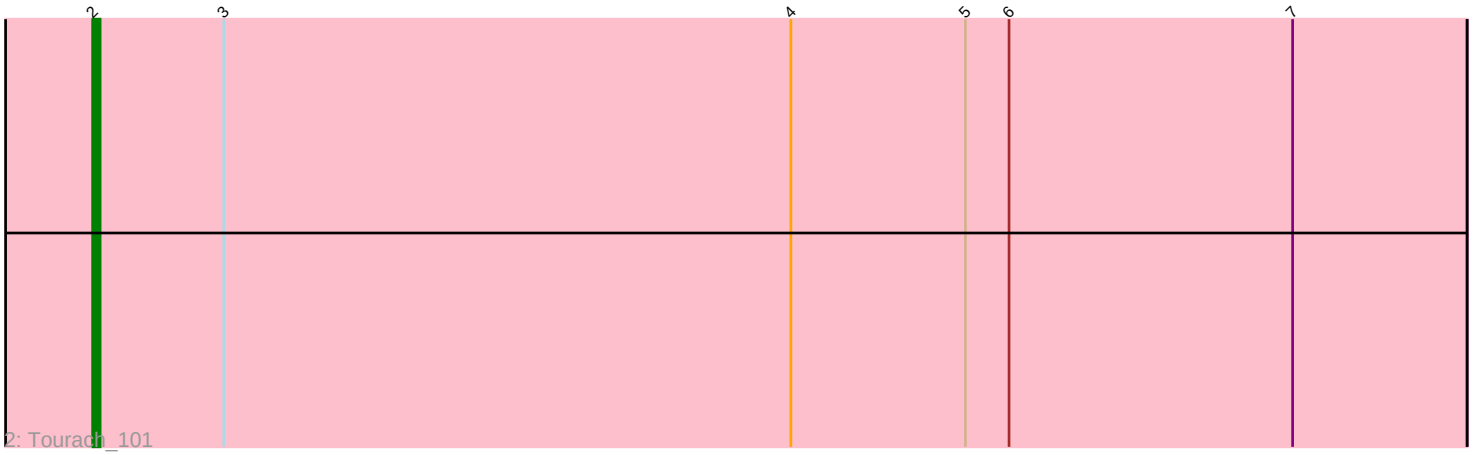
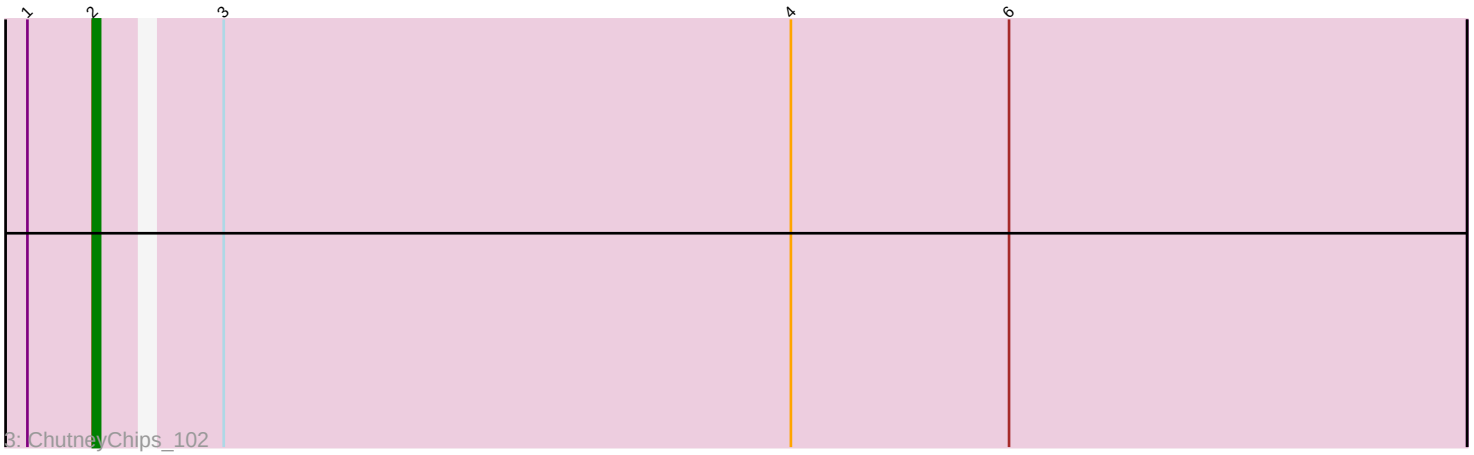
Pham 346679


1
2
3
4
5
6
7
1: DHan_103 + 1


2
3
4
5
6
7
2: Touradh_101


1
2
3
4
6
3: ChutneyChips_102

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 346679 Report

This analysis was run 11/14/26 on database version 672.

Pham number 346679 has 4 members, 2 are drafts.

Phages represented in each track:
• Track 1 : DHan_103, ZhongYanYuan_97
• Track 2 : Tourach_101
• Track 3 : ChutneyChips_102

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

Genes that call this "Most Annotated" start:
• ChutneyChips_102, DHan_103, Tourach_101, ZhongYanYuan_97,

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 4 of 4 (100.0%) of genes in pham
• Manual Annotations of this start: 2 of 2
• Called 100.0% of time when present
• Phage (with cluster) where this start called: ChutneyChips_102 (L4), DHan_103 (L2), Tourach_101 (L2), ZhongYanYuan_97 (L2),

Summary by clusters:

There are 2 clusters represented in this pham: L4, L2,

Info for manual annotations of cluster L2:
•Start number 2 was manually annotated 1 time for cluster L2.

Info for manual annotations of cluster L4:
•Start number 2 was manually annotated 1 time for cluster L4.

Gene Information:

Gene: ChutneyChips_102 Start: 61567, Stop: 61752, Start Num: 2
Candidate Starts for ChutneyChips_102:
(1, 61558), (Start: 2 @61567 has 2 MA's), (3, 61582), (4, 61660), (6, 61690),

Gene: DHan_103 Start: 61955, Stop: 62143, Start Num: 2
Candidate Starts for DHan_103:
(1, 61946), (Start: 2 @61955 has 2 MA's), (3, 61973), (4, 62051), (5, 62075), (6, 62081), (7, 62120),

Gene: Tourach_101 Start: 62411, Stop: 62599, Start Num: 2
Candidate Starts for Tourach_101:
(Start: 2 @62411 has 2 MA's), (3, 62429), (4, 62507), (5, 62531), (6, 62537), (7, 62576),

Gene: ZhongYanYuan_97 Start: 61475, Stop: 61663, Start Num: 2
Candidate Starts for ZhongYanYuan_97:
(1, 61466), (Start: 2 @61475 has 2 MA's), (3, 61493), (4, 61571), (5, 61595), (6, 61601), (7, 61640),