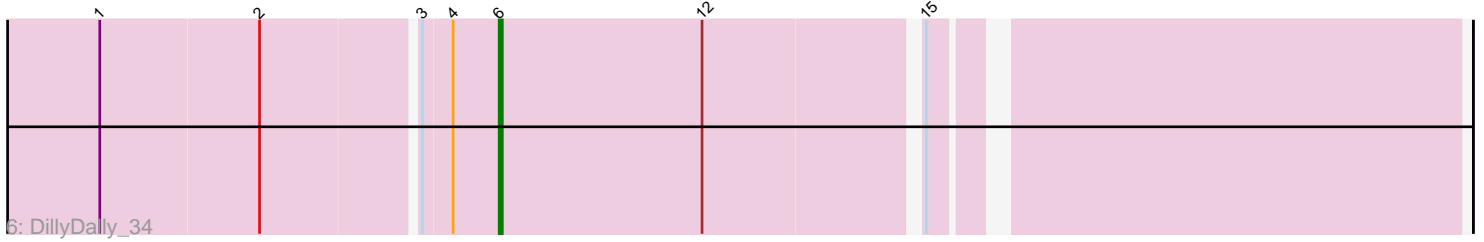
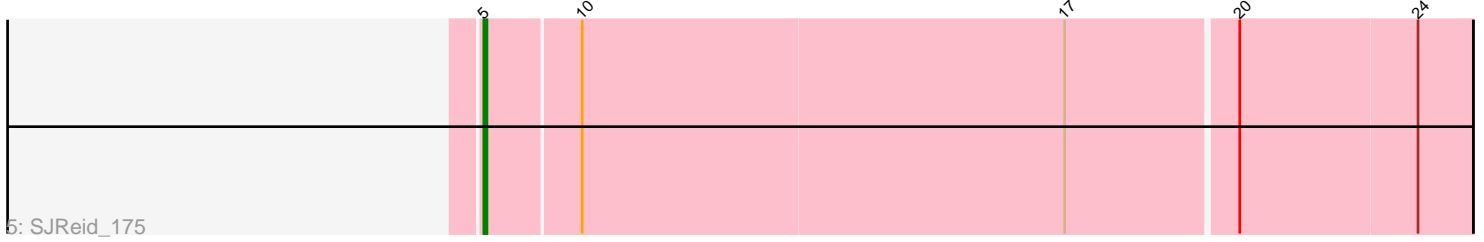
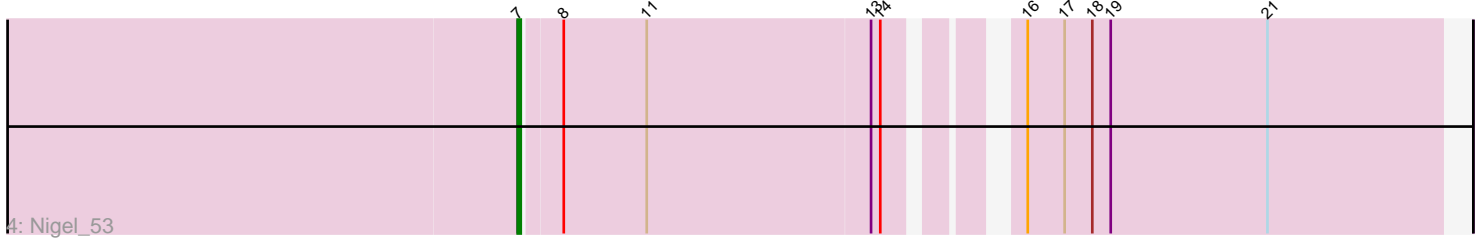
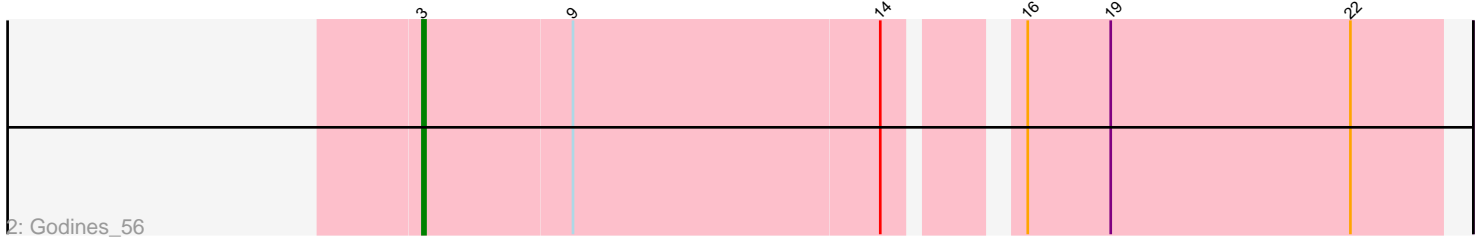


Pham 305592



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 305592 Report

This analysis was run 06/08/26 on database version 649.

Pham number 305592 has 6 members, 0 are drafts.

Phages represented in each track:

- Track 1 : Laurie\_55
- Track 2 : Godines\_56
- Track 3 : Vic9\_00055
- Track 4 : Nigel\_53
- Track 5 : SJReid\_175
- Track 6 : DillyDally\_34

### ***Summary of Final Annotations (See graph section above for start numbers):***

The start number called the most often in the published annotations is 3, it was called in 3 of the 6 non-draft genes in the pham.

Genes that call this "Most Annotated" start:

- Godines\_56, Laurie\_55, Vic9\_00055,

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

- DillyDally\_34,

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

- Nigel\_53, SJReid\_175,

### **Summary by start number:**

Start 3:

- Found in 4 of 6 ( 66.7% ) of genes in pham
- Manual Annotations of this start: 3 of 6
- Called 75.0% of time when present
- Phage (with cluster) where this start called: Godines\_56 (B2), Laurie\_55 (B2), Vic9\_00055 (B2),

Start 5:

- Found in 1 of 6 ( 16.7% ) of genes in pham
- Manual Annotations of this start: 1 of 6
- Called 100.0% of time when present

- Phage (with cluster) where this start called: SJReid\_175 (FC),

Start 6:

- Found in 1 of 6 ( 16.7% ) of genes in pham
- Manual Annotations of this start: 1 of 6
- Called 100.0% of time when present
- Phage (with cluster) where this start called: DillyDally\_34 (singleton),

Start 7:

- Found in 1 of 6 ( 16.7% ) of genes in pham
- Manual Annotations of this start: 1 of 6
- Called 100.0% of time when present
- Phage (with cluster) where this start called: Nigel\_53 (B4),

### **Summary by clusters:**

There are 4 clusters represented in this pham: B4, singleton, FC, B2,

Info for manual annotations of cluster B2:

- Start number 3 was manually annotated 3 times for cluster B2.

Info for manual annotations of cluster B4:

- Start number 7 was manually annotated 1 time for cluster B4.

Info for manual annotations of cluster FC:

- Start number 5 was manually annotated 1 time for cluster FC.

### **Gene Information:**

Gene: DillyDally\_34 Start: 25564, Stop: 25857, Start Num: 6

Candidate Starts for DillyDally\_34:

(1, 25441), (2, 25492), (Start: 3 @25540 has 3 MA's), (4, 25549), (Start: 6 @25564 has 1 MA's), (12, 25630), (15, 25696),

Gene: Godines\_56 Start: 50854, Stop: 50540, Start Num: 3

Candidate Starts for Godines\_56:

(Start: 3 @50854 has 3 MA's), (9, 50806), (14, 50707), (16, 50674), (19, 50647), (22, 50569),

Gene: Laurie\_55 Start: 50282, Stop: 49968, Start Num: 3

Candidate Starts for Laurie\_55:

(Start: 3 @50282 has 3 MA's), (9, 50234), (14, 50135), (16, 50102), (22, 49997), (23, 49988),

Gene: Nigel\_53 Start: 51747, Stop: 51469, Start Num: 7

Candidate Starts for Nigel\_53:

(Start: 7 @51747 has 1 MA's), (8, 51735), (11, 51708), (13, 51636), (14, 51633), (16, 51603), (17, 51591), (18, 51582), (19, 51576), (21, 51525),

Gene: SJReid\_175 Start: 105039, Stop: 105353, Start Num: 5

Candidate Starts for SJReid\_175:

(Start: 5 @105039 has 1 MA's), (10, 105069), (17, 105225), (20, 105279), (24, 105336),

Gene: Vic9\_00055 Start: 51306, Stop: 50992, Start Num: 3

Candidate Starts for Vic9\_00055:

(Start: 3 @51306 has 3 MA's), (9, 51258), (14, 51159), (16, 51126), (19, 51099), (22, 51021), (23, 51012),