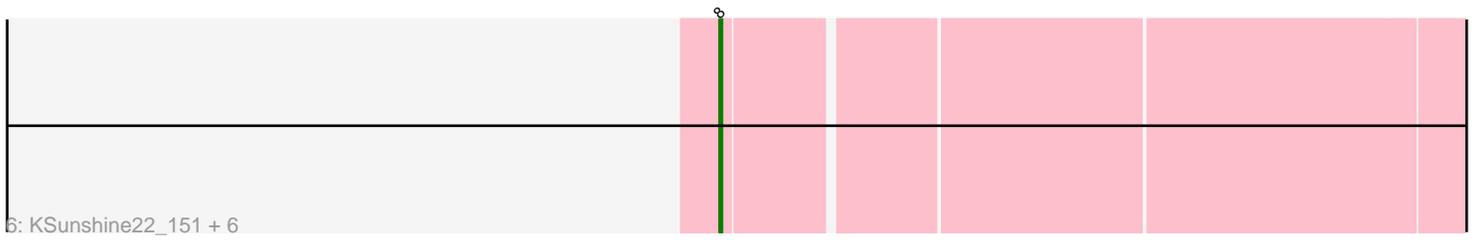
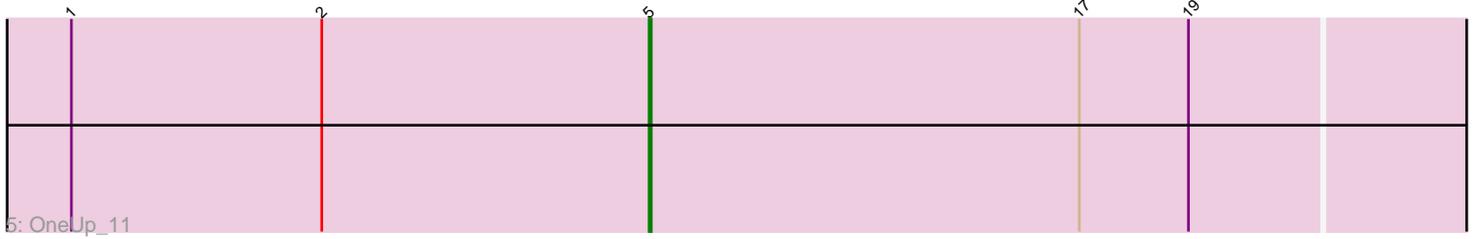
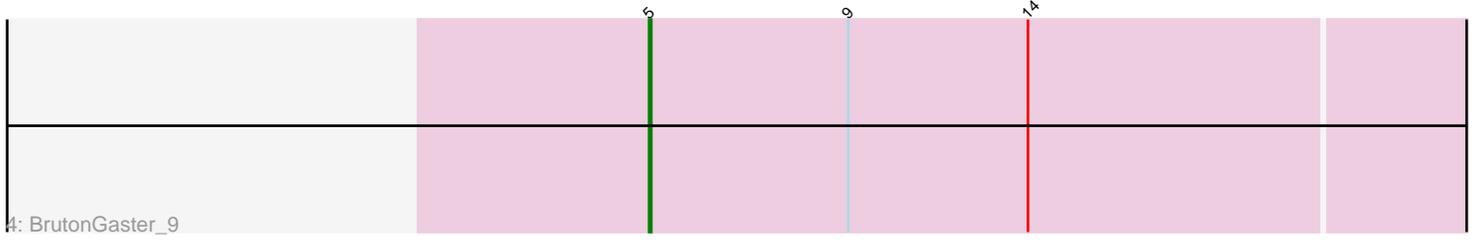
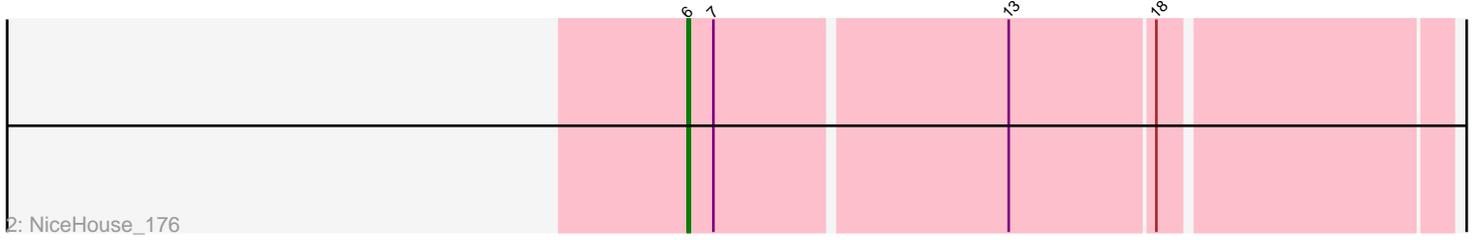
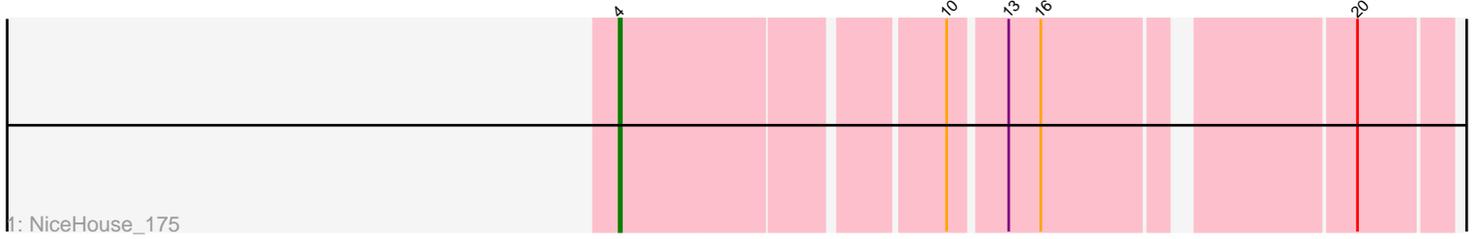


Pham 289923



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 289923 Report

This analysis was run 03/28/26 on database version 641.

Pham number 289923 has 12 members, 4 are drafts.

Phages represented in each track:

- Track 1 : NiceHouse\_175
- Track 2 : NiceHouse\_176
- Track 3 : NiceHouse\_177
- Track 4 : BrutonGaster\_9
- Track 5 : OneUp\_11
- Track 6 : KSunshine22\_151, WaddleDee\_141, BooTeria\_154, DunneganBoMo\_145, Panchaali\_147, Artu\_147, Emmetator\_148

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

The start number called the most often in the published annotations is 8, it was called in 3 of the 8 non-draft genes in the pham.

Genes that call this "Most Annotated" start:

- Artu\_147, BooTeria\_154, DunneganBoMo\_145, Emmetator\_148, KSunshine22\_151, Panchaali\_147, WaddleDee\_141,

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

- 

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

- BrutonGaster\_9, NiceHouse\_175, NiceHouse\_176, NiceHouse\_177, OneUp\_11,

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

Start 3:

- Found in 1 of 12 ( 8.3% ) of genes in pham
- Manual Annotations of this start: 1 of 8
- Called 100.0% of time when present
- Phage (with cluster) where this start called: NiceHouse\_177 (CE),

Start 4:

- Found in 1 of 12 ( 8.3% ) of genes in pham
- Manual Annotations of this start: 1 of 8

- Called 100.0% of time when present
- Phage (with cluster) where this start called: NiceHouse\_175 (CE),

Start 5:

- Found in 2 of 12 ( 16.7% ) of genes in pham
- Manual Annotations of this start: 2 of 8
- Called 100.0% of time when present
- Phage (with cluster) where this start called: BrutonGaster\_9 (CQ2), OneUp\_11 (CQ2),

Start 6:

- Found in 1 of 12 ( 8.3% ) of genes in pham
- Manual Annotations of this start: 1 of 8
- Called 100.0% of time when present
- Phage (with cluster) where this start called: NiceHouse\_176 (CE),

Start 8:

- Found in 7 of 12 ( 58.3% ) of genes in pham
- Manual Annotations of this start: 3 of 8
- Called 100.0% of time when present
- Phage (with cluster) where this start called: Artu\_147 (FC), BooTeria\_154 (FC), DunneganBoMo\_145 (FC), Emmetator\_148 (FC), KSunshine22\_151 (FC), Panchaali\_147 (FC), WaddleDee\_141 (FC),

### **Summary by clusters:**

There are 3 clusters represented in this pham: CQ2, FC, CE,

Info for manual annotations of cluster CE:

- Start number 3 was manually annotated 1 time for cluster CE.
- Start number 4 was manually annotated 1 time for cluster CE.
- Start number 6 was manually annotated 1 time for cluster CE.

Info for manual annotations of cluster CQ2:

- Start number 5 was manually annotated 2 times for cluster CQ2.

Info for manual annotations of cluster FC:

- Start number 8 was manually annotated 3 times for cluster FC.

### **Gene Information:**

Gene: Artu\_147 Start: 101698, Stop: 102027, Start Num: 8

Candidate Starts for Artu\_147:

(Start: 8 @101698 has 3 MA's),

Gene: BooTeria\_154 Start: 101992, Stop: 102321, Start Num: 8

Candidate Starts for BooTeria\_154:

(Start: 8 @101992 has 3 MA's),

Gene: BrutonGaster\_9 Start: 2765, Stop: 3145, Start Num: 5

Candidate Starts for BrutonGaster\_9:

(Start: 5 @2765 has 2 MA's), (9, 2858), (14, 2942),

Gene: DunneganBoMo\_145 Start: 101261, Stop: 101590, Start Num: 8

Candidate Starts for DunneganBoMo\_145:

(Start: 8 @101261 has 3 MA's),

Gene: Emmetator\_148 Start: 101571, Stop: 101903, Start Num: 8

Candidate Starts for Emmetator\_148:

(Start: 8 @101571 has 3 MA's),

Gene: KSunshine22\_151 Start: 102661, Stop: 102990, Start Num: 8

Candidate Starts for KSunshine22\_151:

(Start: 8 @102661 has 3 MA's),

Gene: NiceHouse\_175 Start: 99827, Stop: 100174, Start Num: 4

Candidate Starts for NiceHouse\_175:

(Start: 4 @99827 has 1 MA's), (10, 99965), (13, 99989), (16, 100004), (20, 100133),

Gene: NiceHouse\_176 Start: 100180, Stop: 100518, Start Num: 6

Candidate Starts for NiceHouse\_176:

(Start: 6 @100180 has 1 MA's), (7, 100192), (13, 100324), (18, 100390),

Gene: NiceHouse\_177 Start: 100519, Stop: 100866, Start Num: 3

Candidate Starts for NiceHouse\_177:

(Start: 3 @100519 has 1 MA's), (11, 100657), (12, 100660), (15, 100687), (21, 100846),

Gene: OneUp\_11 Start: 3220, Stop: 3600, Start Num: 5

Candidate Starts for OneUp\_11:

(1, 2950), (2, 3067), (Start: 5 @3220 has 2 MA's), (17, 3421), (19, 3472),

Gene: Panchaali\_147 Start: 101776, Stop: 102108, Start Num: 8

Candidate Starts for Panchaali\_147:

(Start: 8 @101776 has 3 MA's),

Gene: WaddleDee\_141 Start: 100447, Stop: 100776, Start Num: 8

Candidate Starts for WaddleDee\_141:

(Start: 8 @100447 has 3 MA's),