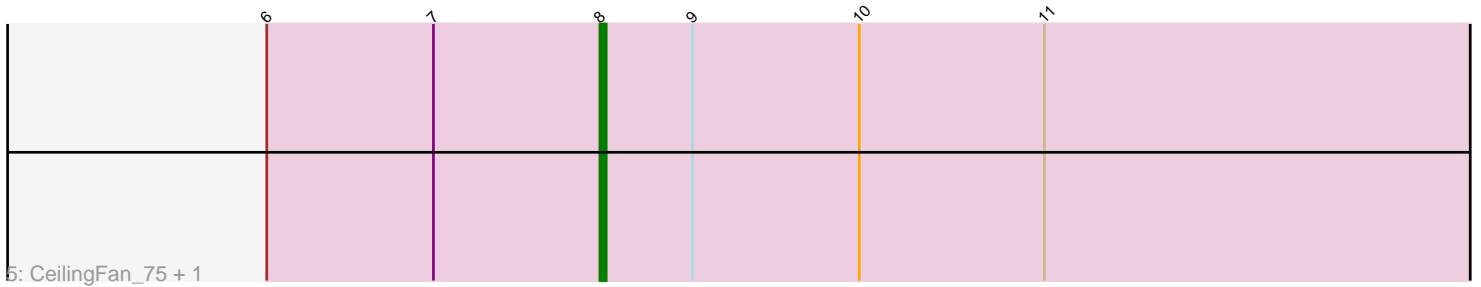
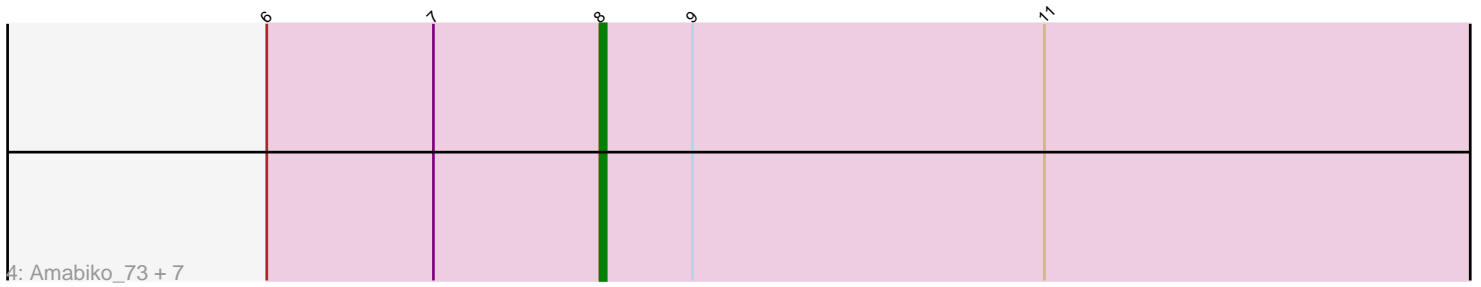
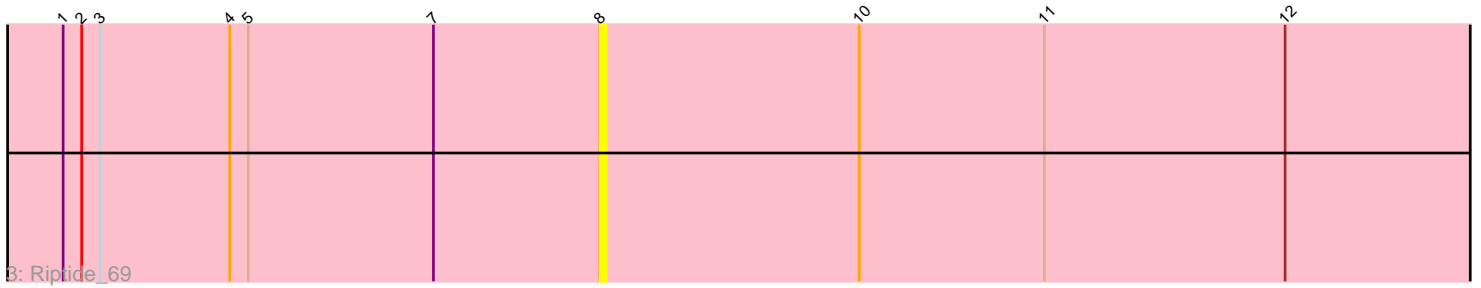
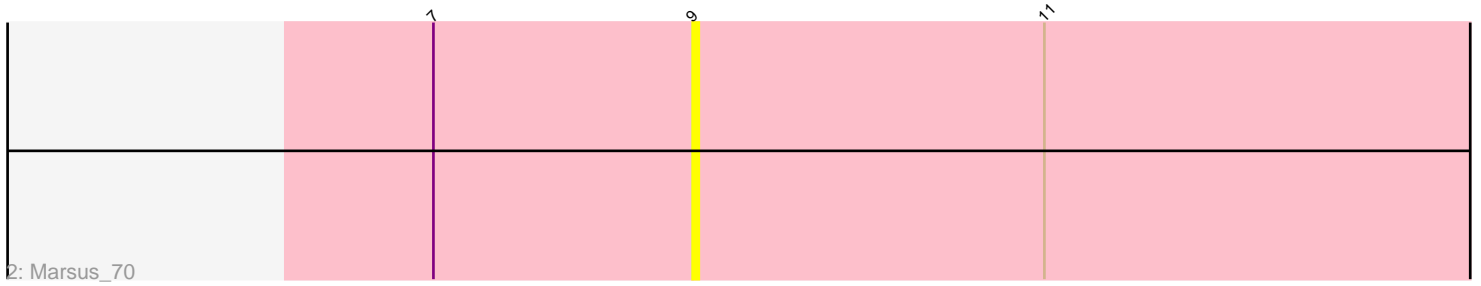
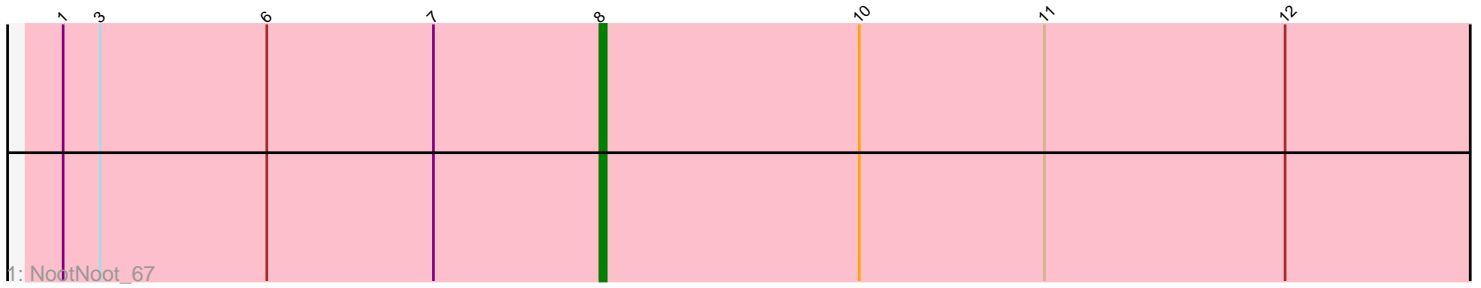


Pham 203460



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 203460 Report

This analysis was run 01/18/25 on database version 583.

Pham number 203460 has 13 members, 5 are drafts.

Phages represented in each track:

- Track 1 : NootNoot\_67
- Track 2 : Marsus\_70
- Track 3 : Riptide\_69
- Track 4 : Amabiko\_73, Gibbi\_77, Starbow\_71, Bordeaux\_71, Rikishi\_76, Battuta\_71, Birchlyn\_70, Quaran19\_73
- Track 5 : CeilingFan\_75, Karimac\_72

### ***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 8 of the 8 non-draft genes in the pham.

Genes that call this "Most Annotated" start:

- Amabiko\_73, Battuta\_71, Birchlyn\_70, Bordeaux\_71, CeilingFan\_75, Gibbi\_77, Karimac\_72, NootNoot\_67, Quaran19\_73, Rikishi\_76, Riptide\_69, Starbow\_71,

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

- 

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

- Marsus\_70,

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

Start 8:

- Found in 12 of 13 ( 92.3% ) of genes in pham
- Manual Annotations of this start: 8 of 8
- Called 100.0% of time when present
- Phage (with cluster) where this start called: Amabiko\_73 (BE2), Battuta\_71 (BE2), Birchlyn\_70 (BE2), Bordeaux\_71 (BE2), CeilingFan\_75 (BE2), Gibbi\_77 (BE2), Karimac\_72 (BE2), NootNoot\_67 (BE1), Quaran19\_73 (BE2), Rikishi\_76 (BE2), Riptide\_69 (BE1), Starbow\_71 (BE2),

Start 9:

- Found in 11 of 13 ( 84.6% ) of genes in pham
- No Manual Annotations of this start.
- Called 9.1% of time when present
- Phage (with cluster) where this start called: Marsus\_70 (BE1),

### **Summary by clusters:**

There are 2 clusters represented in this pham: BE2, BE1,

Info for manual annotations of cluster BE1:

- Start number 8 was manually annotated 1 time for cluster BE1.

Info for manual annotations of cluster BE2:

- Start number 8 was manually annotated 7 times for cluster BE2.

### **Gene Information:**

Gene: Amabiko\_73 Start: 53394, Stop: 53546, Start Num: 8

Candidate Starts for Amabiko\_73:

(6, 53340), (7, 53367), (Start: 8 @53394 has 8 MA's), (9, 53409), (11, 53466),

Gene: Battuta\_71 Start: 53368, Stop: 53520, Start Num: 8

Candidate Starts for Battuta\_71:

(6, 53314), (7, 53341), (Start: 8 @53368 has 8 MA's), (9, 53383), (11, 53440),

Gene: Birchlyn\_70 Start: 51251, Stop: 51403, Start Num: 8

Candidate Starts for Birchlyn\_70:

(6, 51197), (7, 51224), (Start: 8 @51251 has 8 MA's), (9, 51266), (11, 51323),

Gene: Bordeaux\_71 Start: 53523, Stop: 53675, Start Num: 8

Candidate Starts for Bordeaux\_71:

(6, 53469), (7, 53496), (Start: 8 @53523 has 8 MA's), (9, 53538), (11, 53595),

Gene: CeilingFan\_75 Start: 53283, Stop: 53435, Start Num: 8

Candidate Starts for CeilingFan\_75:

(6, 53229), (7, 53256), (Start: 8 @53283 has 8 MA's), (9, 53298), (10, 53325), (11, 53355),

Gene: Gibbi\_77 Start: 53506, Stop: 53658, Start Num: 8

Candidate Starts for Gibbi\_77:

(6, 53452), (7, 53479), (Start: 8 @53506 has 8 MA's), (9, 53521), (11, 53578),

Gene: Karimac\_72 Start: 53500, Stop: 53652, Start Num: 8

Candidate Starts for Karimac\_72:

(6, 53446), (7, 53473), (Start: 8 @53500 has 8 MA's), (9, 53515), (10, 53542), (11, 53572),

Gene: Marsus\_70 Start: 52661, Stop: 52798, Start Num: 9

Candidate Starts for Marsus\_70:

(7, 52619), (9, 52661), (11, 52718),

Gene: NootNoot\_67 Start: 52455, Stop: 52607, Start Num: 8

Candidate Starts for NootNoot\_67:

(1, 52368), (3, 52374), (6, 52401), (7, 52428), (Start: 8 @52455 has 8 MA's), (10, 52497), (11, 52527), (12, 52566),

Gene: Quaran19\_73 Start: 53385, Stop: 53537, Start Num: 8

Candidate Starts for Quaran19\_73:

(6, 53331), (7, 53358), (Start: 8 @53385 has 8 MA's), (9, 53400), (11, 53457),

Gene: Rikishi\_76 Start: 53300, Stop: 53452, Start Num: 8

Candidate Starts for Rikishi\_76:

(6, 53246), (7, 53273), (Start: 8 @53300 has 8 MA's), (9, 53315), (11, 53372),

Gene: Riptide\_69 Start: 52188, Stop: 52340, Start Num: 8

Candidate Starts for Riptide\_69:

(1, 52101), (2, 52104), (3, 52107), (4, 52128), (5, 52131), (7, 52161), (Start: 8 @52188 has 8 MA's), (10, 52230), (11, 52260), (12, 52299),

Gene: Starbow\_71 Start: 53370, Stop: 53522, Start Num: 8

Candidate Starts for Starbow\_71:

(6, 53316), (7, 53343), (Start: 8 @53370 has 8 MA's), (9, 53385), (11, 53442),