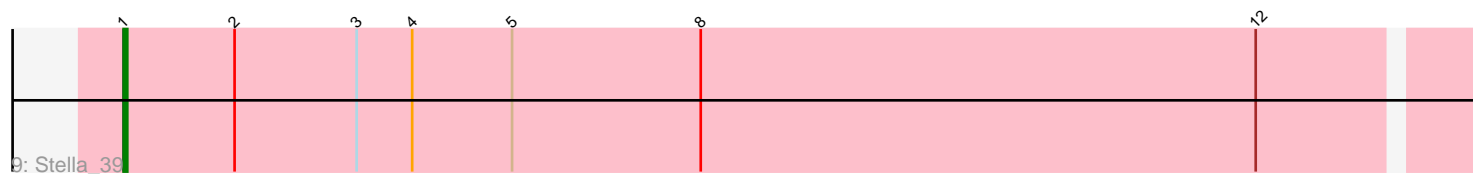
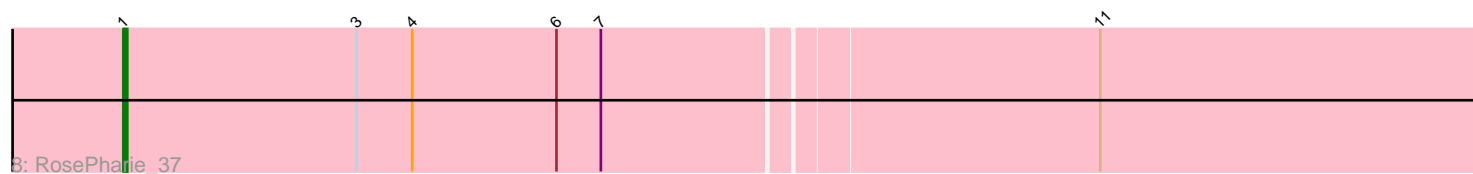
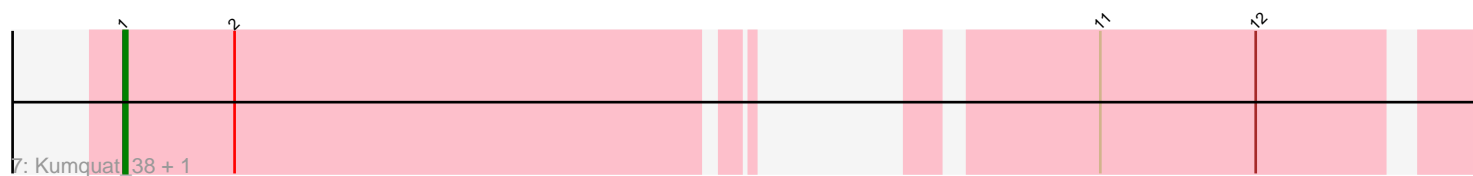
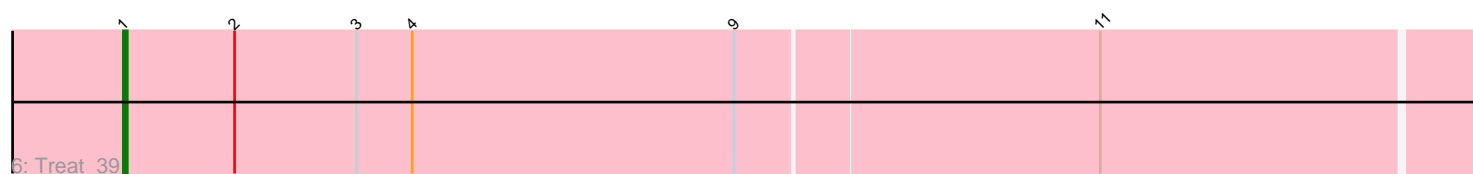
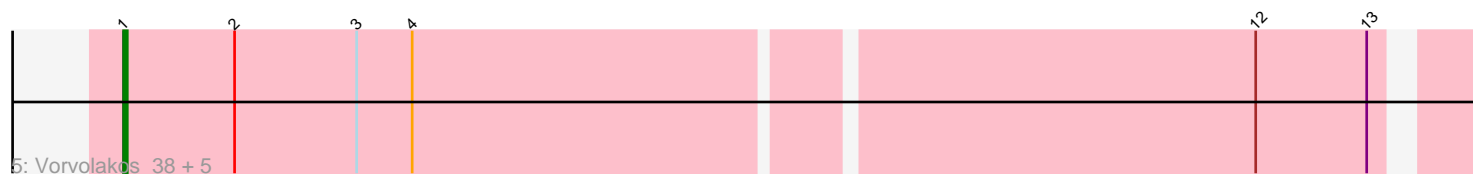
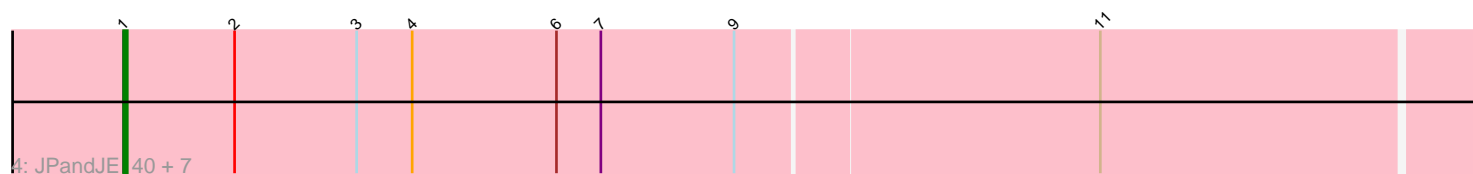
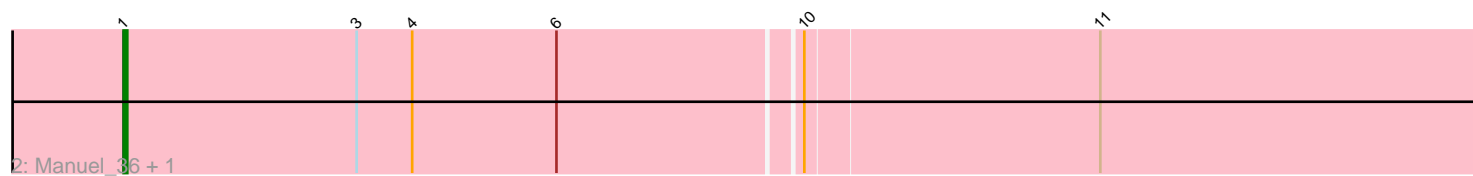


# Pham 44988



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 44988 Report

This analysis was run 04/28/24 on database version 559.

Pham number 44988 has 23 members, 2 are drafts.

Phages represented in each track:

- Track 1 : Dennebes\_39
- Track 2 : Manuel\_36, WRightOn\_40
- Track 3 : Rideau\_40
- Track 4 : JPandJE\_40, HaugeAnator\_39, Percastrophe\_39, ZooBear\_39, Romero\_39, Olicious\_39, Immanuel3\_39, ToriToki\_39
- Track 5 : Vorvolakos\_38, Fabian\_35, FlowerPower\_38, Gremlin23\_38, RetrieverFever\_38, Geostin\_33
- Track 6 : Treat\_39
- Track 7 : Kumquat\_38, Zeigle\_38
- Track 8 : RosePharie\_37
- Track 9 : Stella\_39

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

The start number called the most often in the published annotations is 1, it was called in 21 of the 21 non-draft genes in the pham.

Genes that call this "Most Annotated" start:

- Dennebes\_39, Fabian\_35, FlowerPower\_38, Geostin\_33, Gremlin23\_38, HaugeAnator\_39, Immanuel3\_39, JPandJE\_40, Kumquat\_38, Manuel\_36, Olicious\_39, Percastrophe\_39, RetrieverFever\_38, Romero\_39, RosePharie\_37, Stella\_39, ToriToki\_39, Treat\_39, Vorvolakos\_38, WRightOn\_40, Zeigle\_38, ZooBear\_39,

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

- Rideau\_40,

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

- 

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

Start 1:

- Found in 23 of 23 ( 100.0% ) of genes in pham

- Manual Annotations of this start: 21 of 21
- Called 95.7% of time when present
- Phage (with cluster) where this start called: Dennebes\_39 (BF), Fabian\_35 (BF), FlowerPower\_38 (BF), Geostin\_33 (BF), Gremlin23\_38 (BF), HaugeAnator\_39 (BF), Immanuel3\_39 (BF), JPandJE\_40 (BF), Kumquat\_38 (BF), Manuel\_36 (BF), Olicious\_39 (BF), Percastrophe\_39 (BF), RetrieverFever\_38 (BF), Romero\_39 (BF), RosePharie\_37 (BF), Stella\_39 (BF), ToriToki\_39 (BF), Treat\_39 (BF), Vorvolakos\_38 (BF), WRightOn\_40 (BF), Zeigle\_38 (BF), ZooBear\_39 (BF),

Start 4:

- Found in 21 of 23 ( 91.3% ) of genes in pham
- No Manual Annotations of this start.
- Called 4.8% of time when present
- Phage (with cluster) where this start called: Rideau\_40 (BF),

### Summary by clusters:

There is one cluster represented in this pham: BF

Info for manual annotations of cluster BF:

- Start number 1 was manually annotated 21 times for cluster BF.

### Gene Information:

Gene: Dennebes\_39 Start: 20180, Stop: 20563, Start Num: 1

Candidate Starts for Dennebes\_39:

(Start: 1 @20180 has 21 MA's), (2, 20210), (3, 20243), (4, 20258), (7, 20309), (12, 20486),

Gene: Fabian\_35 Start: 20039, Stop: 20413, Start Num: 1

Candidate Starts for Fabian\_35:

(Start: 1 @20039 has 21 MA's), (2, 20069), (3, 20102), (4, 20117), (12, 20336), (13, 20366),

Gene: FlowerPower\_38 Start: 20039, Stop: 20413, Start Num: 1

Candidate Starts for FlowerPower\_38:

(Start: 1 @20039 has 21 MA's), (2, 20069), (3, 20102), (4, 20117), (12, 20336), (13, 20366),

Gene: Geostin\_33 Start: 20039, Stop: 20413, Start Num: 1

Candidate Starts for Geostin\_33:

(Start: 1 @20039 has 21 MA's), (2, 20069), (3, 20102), (4, 20117), (12, 20336), (13, 20366),

Gene: Gremlin23\_38 Start: 20039, Stop: 20413, Start Num: 1

Candidate Starts for Gremlin23\_38:

(Start: 1 @20039 has 21 MA's), (2, 20069), (3, 20102), (4, 20117), (12, 20336), (13, 20366),

Gene: HaugeAnator\_39 Start: 19966, Stop: 20352, Start Num: 1

Candidate Starts for HaugeAnator\_39:

(Start: 1 @19966 has 21 MA's), (2, 19996), (3, 20029), (4, 20044), (6, 20083), (7, 20095), (9, 20131), (11, 20227),

Gene: Immanuel3\_39 Start: 19970, Stop: 20356, Start Num: 1

Candidate Starts for Immanuel3\_39:

(Start: 1 @19970 has 21 MA's), (2, 20000), (3, 20033), (4, 20048), (6, 20087), (7, 20099), (9, 20135), (11, 20231),

Gene: JPandJE\_40 Start: 20437, Stop: 20823, Start Num: 1

Candidate Starts for JPandJE\_40:

(Start: 1 @20437 has 21 MA's), (2, 20467), (3, 20500), (4, 20515), (6, 20554), (7, 20566), (9, 20602), (11, 20698),

Gene: Kumquat\_38 Start: 20010, Stop: 20339, Start Num: 1

Candidate Starts for Kumquat\_38:

(Start: 1 @20010 has 21 MA's), (2, 20040), (11, 20220), (12, 20262),

Gene: Manuel\_36 Start: 19984, Stop: 20370, Start Num: 1

Candidate Starts for Manuel\_36:

(Start: 1 @19984 has 21 MA's), (3, 20047), (4, 20062), (6, 20101), (10, 20164), (11, 20242),

Gene: Olicious\_39 Start: 19966, Stop: 20352, Start Num: 1

Candidate Starts for Olicious\_39:

(Start: 1 @19966 has 21 MA's), (2, 19996), (3, 20029), (4, 20044), (6, 20083), (7, 20095), (9, 20131), (11, 20227),

Gene: Percastrophe\_39 Start: 19900, Stop: 20286, Start Num: 1

Candidate Starts for Percastrophe\_39:

(Start: 1 @19900 has 21 MA's), (2, 19930), (3, 19963), (4, 19978), (6, 20017), (7, 20029), (9, 20065), (11, 20161),

Gene: RetrieverFever\_38 Start: 20039, Stop: 20413, Start Num: 1

Candidate Starts for RetrieverFever\_38:

(Start: 1 @20039 has 21 MA's), (2, 20069), (3, 20102), (4, 20117), (12, 20336), (13, 20366),

Gene: Rideau\_40 Start: 20258, Stop: 20563, Start Num: 4

Candidate Starts for Rideau\_40:

(Start: 1 @20180 has 21 MA's), (2, 20210), (3, 20243), (4, 20258), (7, 20309), (12, 20486),

Gene: Romero\_39 Start: 19959, Stop: 20345, Start Num: 1

Candidate Starts for Romero\_39:

(Start: 1 @19959 has 21 MA's), (2, 19989), (3, 20022), (4, 20037), (6, 20076), (7, 20088), (9, 20124), (11, 20220),

Gene: RosePharie\_37 Start: 20000, Stop: 20386, Start Num: 1

Candidate Starts for RosePharie\_37:

(Start: 1 @20000 has 21 MA's), (3, 20063), (4, 20078), (6, 20117), (7, 20129), (11, 20258),

Gene: Stella\_39 Start: 19724, Stop: 20110, Start Num: 1

Candidate Starts for Stella\_39:

(Start: 1 @19724 has 21 MA's), (2, 19754), (3, 19787), (4, 19802), (5, 19829), (8, 19880), (12, 20030),

Gene: ToriToki\_39 Start: 19959, Stop: 20345, Start Num: 1

Candidate Starts for ToriToki\_39:

(Start: 1 @19959 has 21 MA's), (2, 19989), (3, 20022), (4, 20037), (6, 20076), (7, 20088), (9, 20124), (11, 20220),

Gene: Treat\_39 Start: 19903, Stop: 20289, Start Num: 1

Candidate Starts for Treat\_39:

(Start: 1 @19903 has 21 MA's), (2, 19933), (3, 19966), (4, 19981), (9, 20068), (11, 20164),

Gene: Vorvolakos\_38 Start: 20038, Stop: 20412, Start Num: 1

Candidate Starts for Vorvolakos\_38:

(Start: 1 @20038 has 21 MA's), (2, 20068), (3, 20101), (4, 20116), (12, 20335), (13, 20365),

Gene: WRightOn\_40 Start: 20062, Stop: 20448, Start Num: 1

Candidate Starts for WRightOn\_40:

(Start: 1 @20062 has 21 MA's), (3, 20125), (4, 20140), (6, 20179), (10, 20242), (11, 20320),

Gene: Zeigle\_38 Start: 20010, Stop: 20339, Start Num: 1

Candidate Starts for Zeigle\_38:

(Start: 1 @20010 has 21 MA's), (2, 20040), (11, 20220), (12, 20262),

Gene: ZooBear\_39 Start: 19966, Stop: 20352, Start Num: 1

Candidate Starts for ZooBear\_39:

(Start: 1 @19966 has 21 MA's), (2, 19996), (3, 20029), (4, 20044), (6, 20083), (7, 20095), (9, 20131),  
(11, 20227),