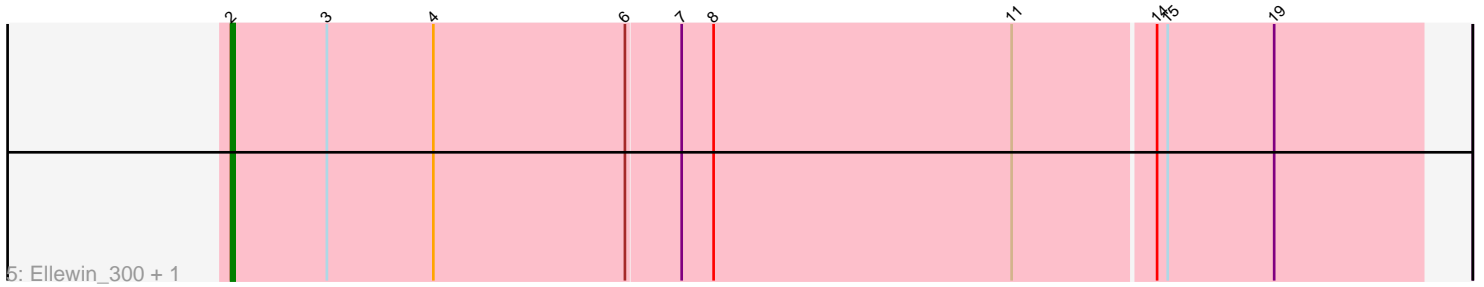
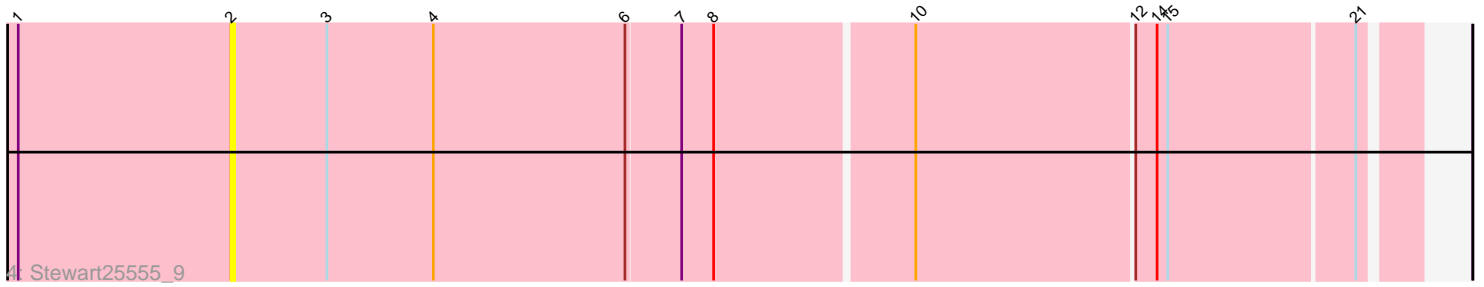
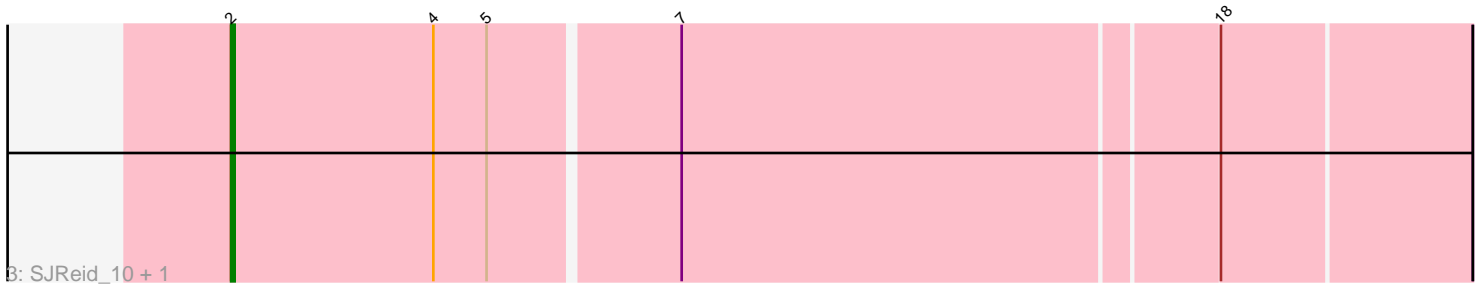
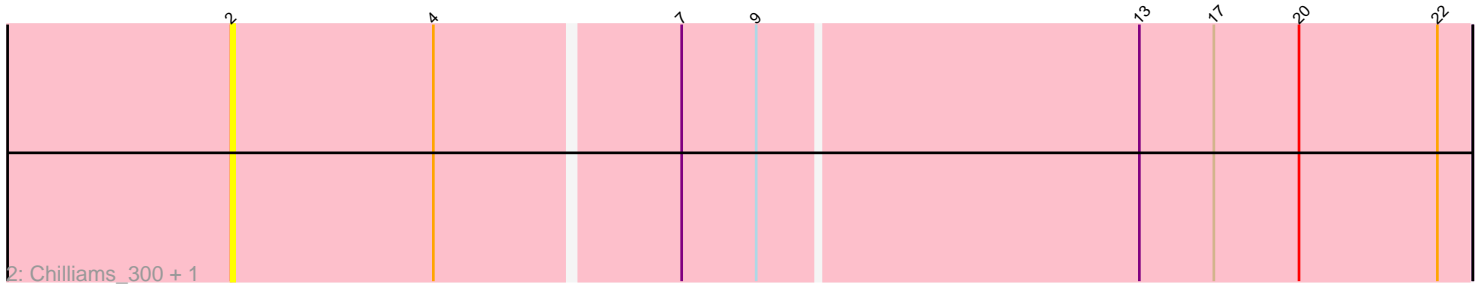
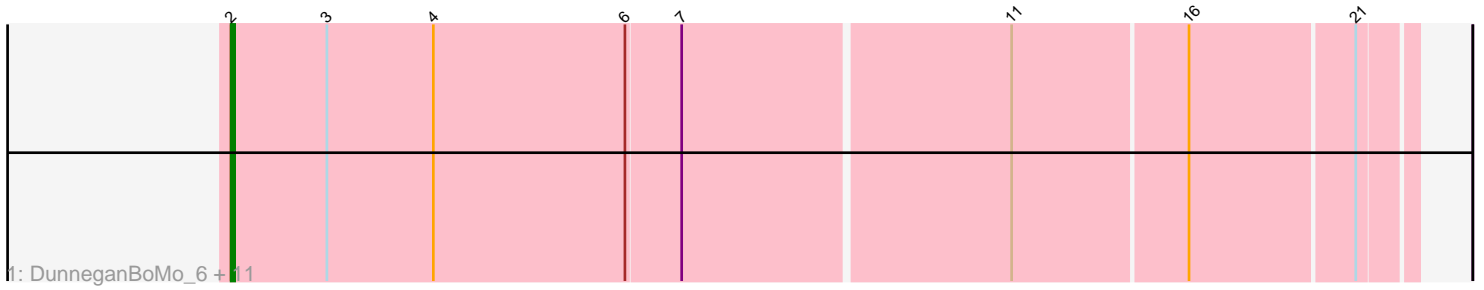


Pham 309370



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 309370 Report

This analysis was run 06/27/26 on database version 652.

Pham number 309370 has 19 members, 11 are drafts.

Phages represented in each track:

- Track 1 : DunneganBoMo\_6, WaddleDee\_297, Artu\_8, KSunshine22\_9, BooTeria\_306, Artu\_295, DunneganBoMo\_301, WaddleDee\_6, BooTeria\_7, Emmetator\_6, Emmetator\_300, KSunshine22\_301
- Track 2 : Chilliams\_300, Chilliams\_9
- Track 3 : SJReid\_10, SJReid\_319
- Track 4 : Stewart25555\_9
- Track 5 : Ellewin\_300, Ellewin\_7

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

Genes that call this "Most Annotated" start:

- Artu\_295, Artu\_8, BooTeria\_306, BooTeria\_7, Chilliams\_300, Chilliams\_9, DunneganBoMo\_301, DunneganBoMo\_6, Ellewin\_300, Ellewin\_7, Emmetator\_300, Emmetator\_6, KSunshine22\_301, KSunshine22\_9, SJReid\_10, SJReid\_319, Stewart25555\_9, WaddleDee\_297, WaddleDee\_6,

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 19 of 19 ( 100.0% ) 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: Artu\_295 (FC), Artu\_8 (FC), BooTeria\_306 (FC), BooTeria\_7 (FC), Chilliams\_300 (FC), Chilliams\_9 (FC), DunneganBoMo\_301 (FC), DunneganBoMo\_6 (FC), Ellewin\_300 (FC), Ellewin\_7

(FC), Emmetator\_300 (FC), Emmetator\_6 (FC), KSunshine22\_301 (FC), KSunshine22\_9 (FC), SJReid\_10 (FC), SJReid\_319 (FC), Stewart25555\_9 (FC), WaddleDee\_297 (FC), WaddleDee\_6 (FC),

### **Summary by clusters:**

There is one cluster represented in this pham: FC

Info for manual annotations of cluster FC:

- Start number 2 was manually annotated 8 times for cluster FC.

### **Gene Information:**

Gene: Artu\_8 Start: 3867, Stop: 4190, Start Num: 2

Candidate Starts for Artu\_8:

(Start: 2 @3867 has 8 MA's), (3, 3894), (4, 3924), (6, 3978), (7, 3993), (11, 4083), (16, 4131), (21, 4176),

Gene: Artu\_295 Start: 183021, Stop: 183344, Start Num: 2

Candidate Starts for Artu\_295:

(Start: 2 @183021 has 8 MA's), (3, 183048), (4, 183078), (6, 183132), (7, 183147), (11, 183237), (16, 183285), (21, 183330),

Gene: BooTeria\_306 Start: 182814, Stop: 183137, Start Num: 2

Candidate Starts for BooTeria\_306:

(Start: 2 @182814 has 8 MA's), (3, 182841), (4, 182871), (6, 182925), (7, 182940), (11, 183030), (16, 183078), (21, 183123),

Gene: BooTeria\_7 Start: 3905, Stop: 4228, Start Num: 2

Candidate Starts for BooTeria\_7:

(Start: 2 @3905 has 8 MA's), (3, 3932), (4, 3962), (6, 4016), (7, 4031), (11, 4121), (16, 4169), (21, 4214),

Gene: Chilliams\_300 Start: 177470, Stop: 177811, Start Num: 2

Candidate Starts for Chilliams\_300:

(Start: 2 @177470 has 8 MA's), (4, 177527), (7, 177593), (9, 177614), (13, 177719), (17, 177740), (20, 177764), (22, 177803),

Gene: Chilliams\_9 Start: 4736, Stop: 5077, Start Num: 2

Candidate Starts for Chilliams\_9:

(Start: 2 @4736 has 8 MA's), (4, 4793), (7, 4859), (9, 4880), (13, 4985), (17, 5006), (20, 5030), (22, 5069),

Gene: DunneganBoMo\_6 Start: 3916, Stop: 4239, Start Num: 2

Candidate Starts for DunneganBoMo\_6:

(Start: 2 @3916 has 8 MA's), (3, 3943), (4, 3973), (6, 4027), (7, 4042), (11, 4132), (16, 4180), (21, 4225),

Gene: DunneganBoMo\_301 Start: 183328, Stop: 183651, Start Num: 2

Candidate Starts for DunneganBoMo\_301:

(Start: 2 @183328 has 8 MA's), (3, 183355), (4, 183385), (6, 183439), (7, 183454), (11, 183544), (16, 183592), (21, 183637),

Gene: Ellewin\_300 Start: 182937, Stop: 183269, Start Num: 2

Candidate Starts for Ellewin\_300:

(Start: 2 @182937 has 8 MA's), (3, 182964), (4, 182994), (6, 183048), (7, 183063), (8, 183072), (11, 183156), (14, 183195), (15, 183198), (19, 183228),

Gene: Ellewin\_7 Start: 3823, Stop: 4155, Start Num: 2

Candidate Starts for Ellewin\_7:

(Start: 2 @3823 has 8 MA's), (3, 3850), (4, 3880), (6, 3934), (7, 3949), (8, 3958), (11, 4042), (14, 4081), (15, 4084), (19, 4114),

Gene: Emmetator\_6 Start: 4075, Stop: 4398, Start Num: 2

Candidate Starts for Emmetator\_6:

(Start: 2 @4075 has 8 MA's), (3, 4102), (4, 4132), (6, 4186), (7, 4201), (11, 4291), (16, 4339), (21, 4384),

Gene: Emmetator\_300 Start: 182375, Stop: 182698, Start Num: 2

Candidate Starts for Emmetator\_300:

(Start: 2 @182375 has 8 MA's), (3, 182402), (4, 182432), (6, 182486), (7, 182501), (11, 182591), (16, 182639), (21, 182684),

Gene: KSunshine22\_9 Start: 4506, Stop: 4829, Start Num: 2

Candidate Starts for KSunshine22\_9:

(Start: 2 @4506 has 8 MA's), (3, 4533), (4, 4563), (6, 4617), (7, 4632), (11, 4722), (16, 4770), (21, 4815),

Gene: KSunshine22\_301 Start: 181407, Stop: 181730, Start Num: 2

Candidate Starts for KSunshine22\_301:

(Start: 2 @181407 has 8 MA's), (3, 181434), (4, 181464), (6, 181518), (7, 181533), (11, 181623), (16, 181671), (21, 181716),

Gene: SJReid\_10 Start: 5211, Stop: 5549, Start Num: 2

Candidate Starts for SJReid\_10:

(Start: 2 @5211 has 8 MA's), (4, 5268), (5, 5283), (7, 5334), (18, 5481),

Gene: SJReid\_319 Start: 178050, Stop: 178388, Start Num: 2

Candidate Starts for SJReid\_319:

(Start: 2 @178050 has 8 MA's), (4, 178107), (5, 178122), (7, 178173), (18, 178320),

Gene: Stewart25555\_9 Start: 4661, Stop: 4984, Start Num: 2

Candidate Starts for Stewart25555\_9:

(1, 4601), (Start: 2 @4661 has 8 MA's), (3, 4688), (4, 4718), (6, 4772), (7, 4787), (8, 4796), (10, 4850), (12, 4910), (14, 4916), (15, 4919), (21, 4970),

Gene: WaddleDee\_297 Start: 182111, Stop: 182434, Start Num: 2

Candidate Starts for WaddleDee\_297:

(Start: 2 @182111 has 8 MA's), (3, 182138), (4, 182168), (6, 182222), (7, 182237), (11, 182327), (16, 182375), (21, 182420),

Gene: WaddleDee\_6 Start: 3916, Stop: 4239, Start Num: 2

Candidate Starts for WaddleDee\_6:

(Start: 2 @3916 has 8 MA's), (3, 3943), (4, 3973), (6, 4027), (7, 4042), (11, 4132), (16, 4180), (21, 4225),