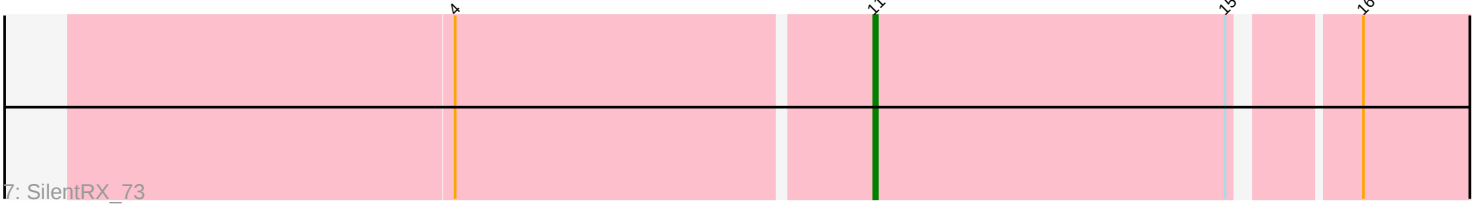
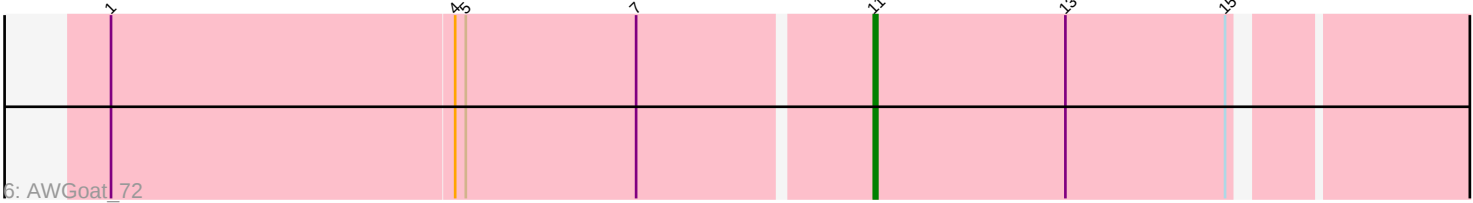
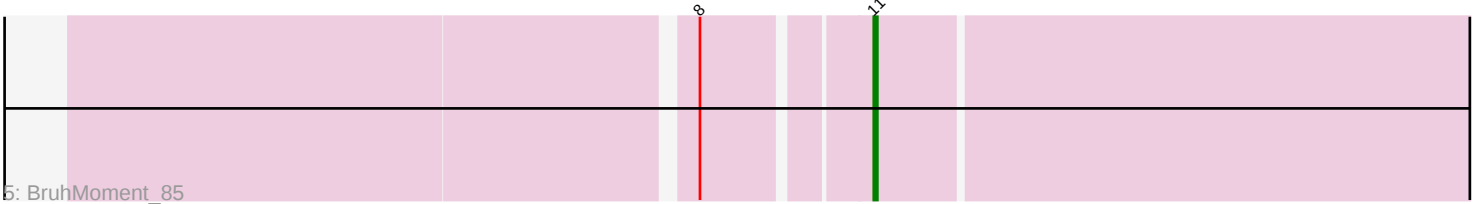
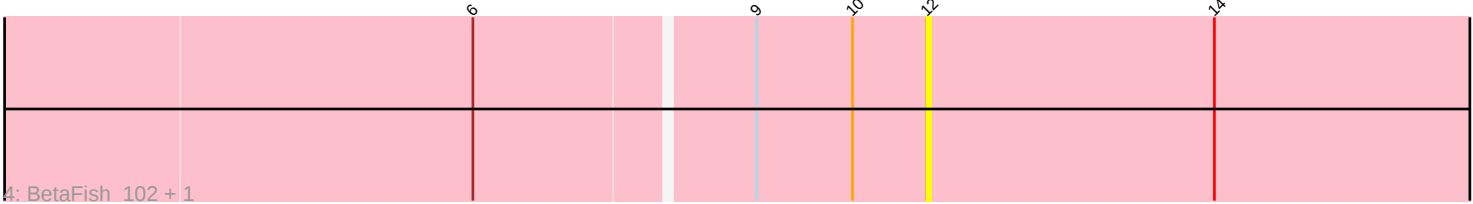
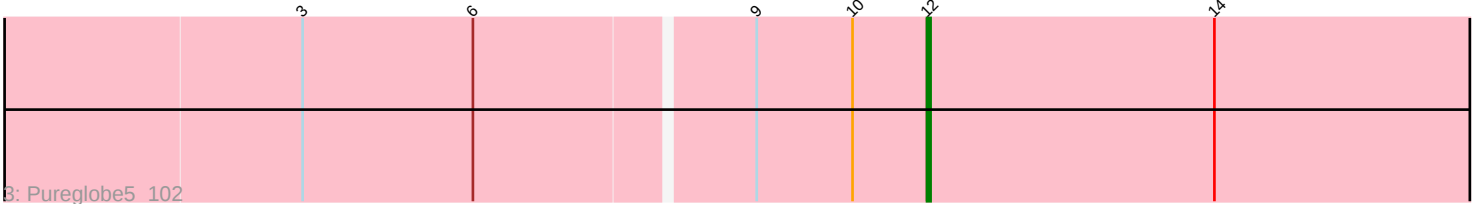
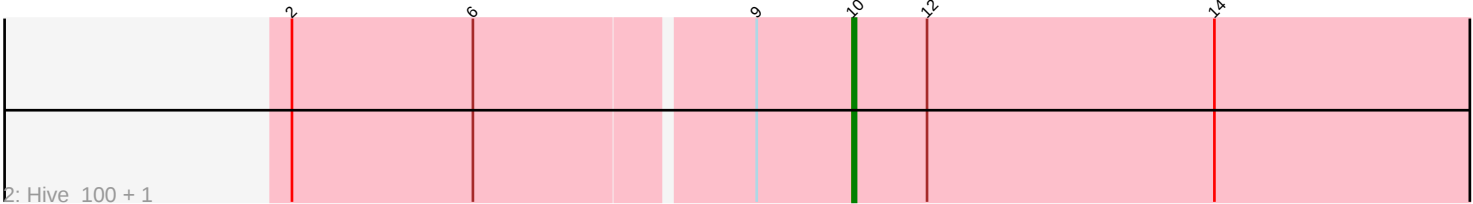
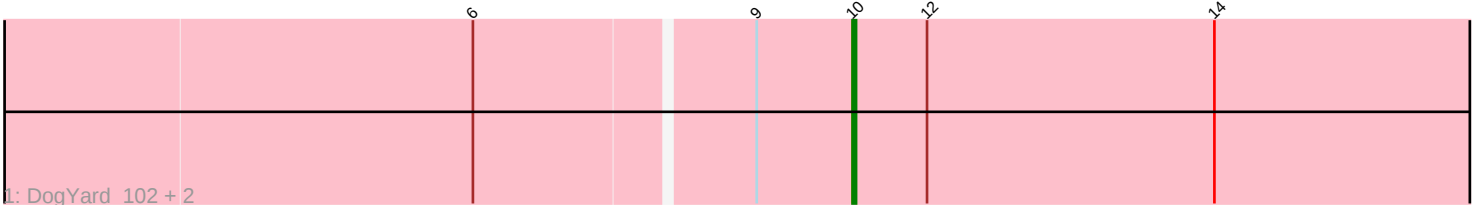


Pham 284376



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 284376 Report

This analysis was run 02/23/26 on database version 636.

Pham number 284376 has 11 members, 4 are drafts.

Phages represented in each track:

- Track 1 : DogYard_102, PhuzzTulsa_101, Odyssey395_102
- Track 2 : Hive_100, RazzB_101
- Track 3 : Pureglobe5_102
- Track 4 : BetaFish_102, Popstraw_98
- Track 5 : BruhMoment_85
- Track 6 : AWGoat_72
- Track 7 : SilentRX_73

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

The start number called the most often in the published annotations is 10, it was called in 3 of the 7 non-draft genes in the pham.

Genes that call this "Most Annotated" start:

- DogYard_102, Hive_100, Odyssey395_102, PhuzzTulsa_101, RazzB_101,

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

- BetaFish_102, Popstraw_98, Pureglobe5_102,

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

- AWGoat_72, BruhMoment_85, SilentRX_73,

Summary by start number:

Start 10:

- Found in 8 of 11 (72.7%) of genes in pham
- Manual Annotations of this start: 3 of 7
- Called 62.5% of time when present
- Phage (with cluster) where this start called: DogYard_102 (AP2), Hive_100 (AP2), Odyssey395_102 (AP2), PhuzzTulsa_101 (AP2), RazzB_101 (AP2),

Start 11:

- Found in 3 of 11 (27.3%) of genes in pham
- Manual Annotations of this start: 3 of 7

- Called 100.0% of time when present
- Phage (with cluster) where this start called: AWGoat_72 (AP4), BruhMoment_85 (AP3), SilentRX_73 (AP4),

Start 12:

- Found in 8 of 11 (72.7%) of genes in pham
- Manual Annotations of this start: 1 of 7
- Called 37.5% of time when present
- Phage (with cluster) where this start called: BetaFish_102 (AP2), Popstraw_98 (AP2), Pureglobe5_102 (AP2),

Summary by clusters:

There are 3 clusters represented in this pham: AP2, AP3, AP4,

Info for manual annotations of cluster AP2:

- Start number 10 was manually annotated 3 times for cluster AP2.
- Start number 12 was manually annotated 1 time for cluster AP2.

Info for manual annotations of cluster AP3:

- Start number 11 was manually annotated 1 time for cluster AP3.

Info for manual annotations of cluster AP4:

- Start number 11 was manually annotated 2 times for cluster AP4.

Gene Information:

Gene: AWGoat_72 Start: 50701, Stop: 50540, Start Num: 11

Candidate Starts for AWGoat_72:

(1, 50911), (4, 50815), (5, 50812), (7, 50764), (Start: 11 @50701 has 3 MA's), (13, 50647), (15, 50602),

Gene: BetaFish_102 Start: 61576, Stop: 61403, Start Num: 12

Candidate Starts for BetaFish_102:

(6, 61699), (9, 61624), (Start: 10 @61597 has 3 MA's), (Start: 12 @61576 has 1 MA's), (14, 61495),

Gene: BruhMoment_85 Start: 55806, Stop: 55639, Start Num: 11

Candidate Starts for BruhMoment_85:

(8, 55845), (Start: 11 @55806 has 3 MA's),

Gene: DogYard_102 Start: 61090, Stop: 60896, Start Num: 10

Candidate Starts for DogYard_102:

(6, 61192), (9, 61117), (Start: 10 @61090 has 3 MA's), (Start: 12 @61069 has 1 MA's), (14, 60988),

Gene: Hive_100 Start: 61072, Stop: 60878, Start Num: 10

Candidate Starts for Hive_100:

(2, 61225), (6, 61174), (9, 61099), (Start: 10 @61072 has 3 MA's), (Start: 12 @61051 has 1 MA's), (14, 60970),

Gene: Odyssey395_102 Start: 60780, Stop: 60586, Start Num: 10

Candidate Starts for Odyssey395_102:

(6, 60882), (9, 60807), (Start: 10 @60780 has 3 MA's), (Start: 12 @60759 has 1 MA's), (14, 60678),

Gene: PhuzzTulsa_101 Start: 61228, Stop: 61034, Start Num: 10

Candidate Starts for PhuzzTulsa_101:

(6, 61330), (9, 61255), (Start: 10 @61228 has 3 MA's), (Start: 12 @61207 has 1 MA's), (14, 61126),

Gene: Popstraw_98 Start: 60828, Stop: 60655, Start Num: 12

Candidate Starts for Popstraw_98:

(6, 60951), (9, 60876), (Start: 10 @60849 has 3 MA's), (Start: 12 @60828 has 1 MA's), (14, 60747),

Gene: Pureglobe5_102 Start: 61317, Stop: 61144, Start Num: 12

Candidate Starts for Pureglobe5_102:

(3, 61488), (6, 61440), (9, 61365), (Start: 10 @61338 has 3 MA's), (Start: 12 @61317 has 1 MA's), (14, 61236),

Gene: RazzB_101 Start: 60213, Stop: 60019, Start Num: 10

Candidate Starts for RazzB_101:

(2, 60366), (6, 60315), (9, 60240), (Start: 10 @60213 has 3 MA's), (Start: 12 @60192 has 1 MA's), (14, 60111),

Gene: SilentRX_73 Start: 51541, Stop: 51380, Start Num: 11

Candidate Starts for SilentRX_73:

(4, 51655), (Start: 11 @51541 has 3 MA's), (15, 51442), (16, 51412),