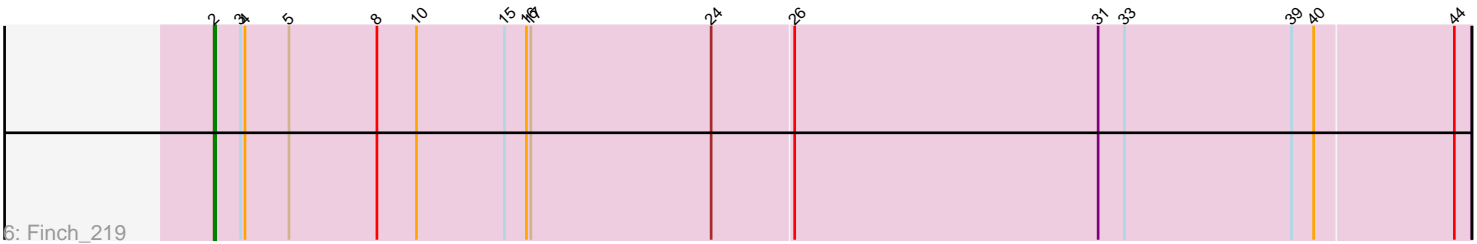
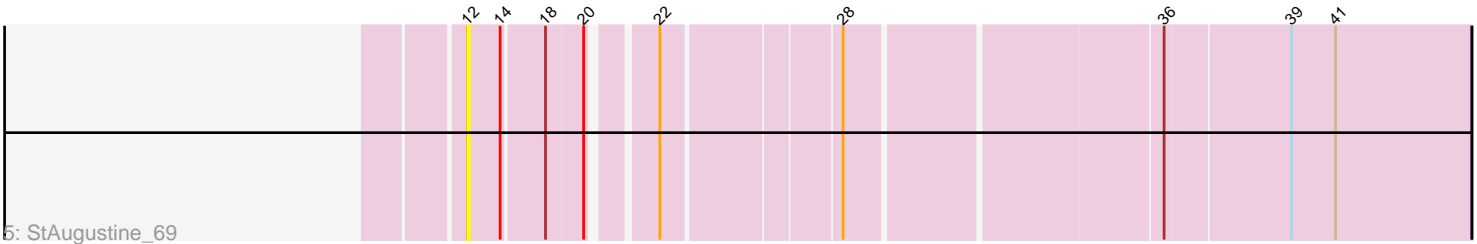
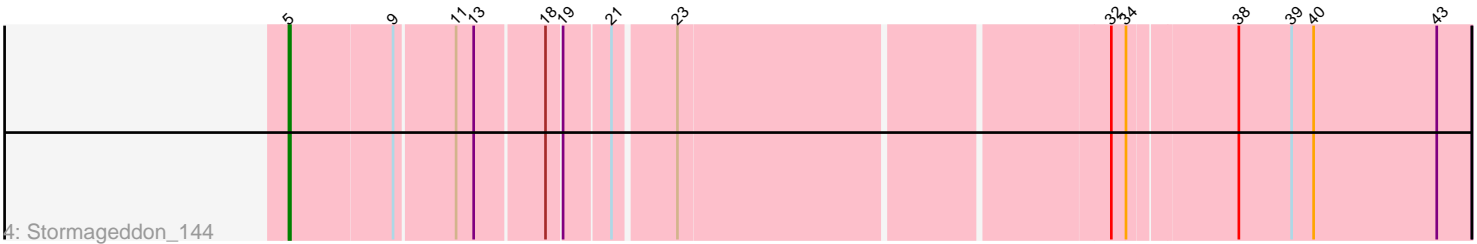
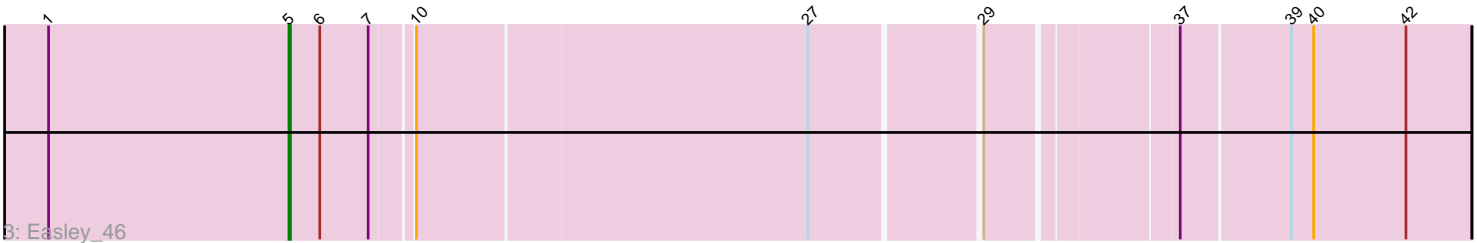
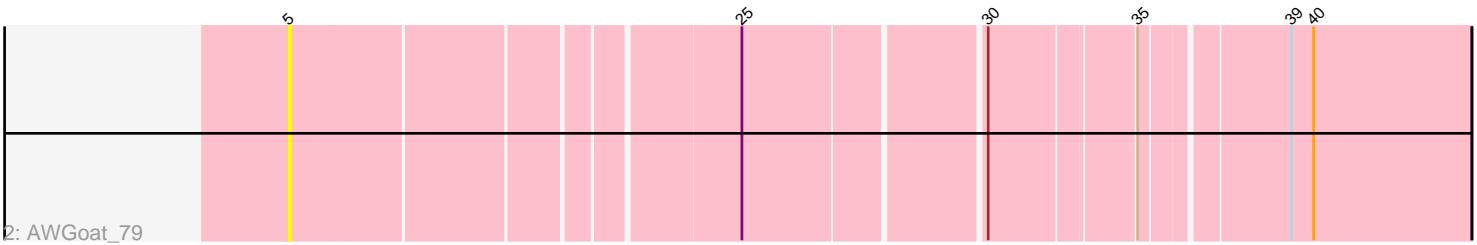
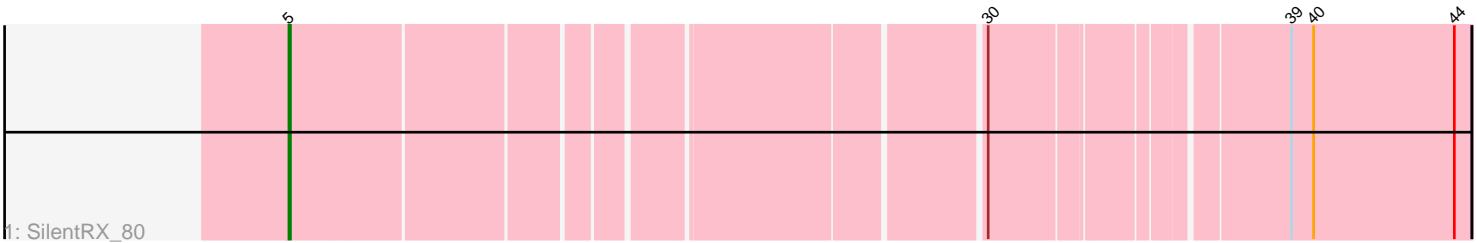


Pham 88886



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 88886 Report

This analysis was run 04/05/24 on database version 557.

Pham number 88886 has 6 members, 2 are drafts.

Phages represented in each track:

- Track 1 : SilentRX_80
- Track 2 : AWGoat_79
- Track 3 : Easley_46
- Track 4 : Stormageddon_144
- Track 5 : StAugustine_69
- Track 6 : Finch_219

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

The start number called the most often in the published annotations is 5, it was called in 3 of the 4 non-draft genes in the pham.

Genes that call this "Most Annotated" start:

- AWGoat_79, Easley_46, SilentRX_80, Stormageddon_144,

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

- Finch_219,

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

- StAugustine_69,

Summary by start number:

Start 2:

- Found in 1 of 6 (16.7%) of genes in pham
- Manual Annotations of this start: 1 of 4
- Called 100.0% of time when present
- Phage (with cluster) where this start called: Finch_219 (singleton),

Start 5:

- Found in 5 of 6 (83.3%) of genes in pham
- Manual Annotations of this start: 3 of 4
- Called 80.0% of time when present

- Phage (with cluster) where this start called: AWGoat_79 (AP4), Easley_46 (CZ4), SilentRX_80 (AP4), Stormageddon_144 (DX),

Start 12:

- Found in 1 of 6 (16.7%) of genes in pham
- No Manual Annotations of this start.
- Called 100.0% of time when present
- Phage (with cluster) where this start called: StAugustine_69 (singleton),

Summary by clusters:

There are 4 clusters represented in this pham: CZ4, singleton, DX, AP4,

Info for manual annotations of cluster AP4:

- Start number 5 was manually annotated 1 time for cluster AP4.

Info for manual annotations of cluster CZ4:

- Start number 5 was manually annotated 1 time for cluster CZ4.

Info for manual annotations of cluster DX:

- Start number 5 was manually annotated 1 time for cluster DX.

Gene Information:

Gene: AWGoat_79 Start: 54602, Stop: 53865, Start Num: 5

Candidate Starts for AWGoat_79:

(Start: 5 @54602 has 3 MA's), (25, 54320), (30, 54167), (35, 54077), (39, 53987), (40, 53972),

Gene: Easley_46 Start: 33954, Stop: 34712, Start Num: 5

Candidate Starts for Easley_46:

(1, 33792), (Start: 5 @33954 has 3 MA's), (6, 33975), (7, 34008), (10, 34032), (27, 34293), (29, 34401), (37, 34518), (39, 34590), (40, 34605), (42, 34668),

Gene: Finch_219 Start: 127269, Stop: 128120, Start Num: 2

Candidate Starts for Finch_219:

(Start: 2 @127269 has 1 MA's), (3, 127287), (4, 127290), (Start: 5 @127320 has 3 MA's), (8, 127380), (10, 127407), (15, 127467), (16, 127482), (17, 127485), (24, 127608), (26, 127662), (31, 127869), (33, 127887), (39, 128001), (40, 128016), (44, 128109),

Gene: SilentRX_80 Start: 55474, Stop: 54743, Start Num: 5

Candidate Starts for SilentRX_80:

(Start: 5 @55474 has 3 MA's), (30, 55045), (39, 54865), (40, 54850), (44, 54754),

Gene: StAugustine_69 Start: 54268, Stop: 53645, Start Num: 12

Candidate Starts for StAugustine_69:

(12, 54268), (14, 54247), (18, 54220), (20, 54196), (22, 54160), (28, 54049), (36, 53848), (39, 53767), (41, 53737),

Gene: Stormageddon_144 Start: 101018, Stop: 101776, Start Num: 5

Candidate Starts for Stormageddon_144:

(Start: 5 @101018 has 3 MA's), (9, 101087), (11, 101126), (13, 101138), (18, 101183), (19, 101192), (21, 101222), (23, 101261), (32, 101540), (34, 101549), (38, 101618), (39, 101654), (40, 101669), (43, 101753),