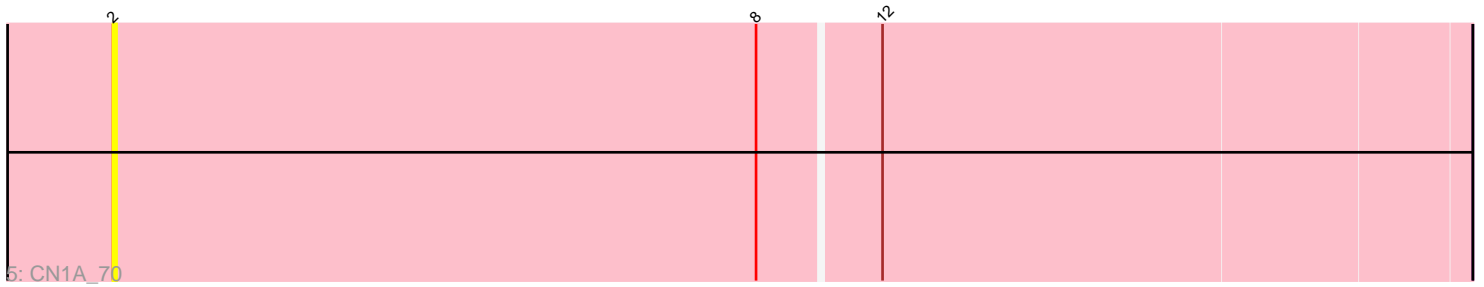
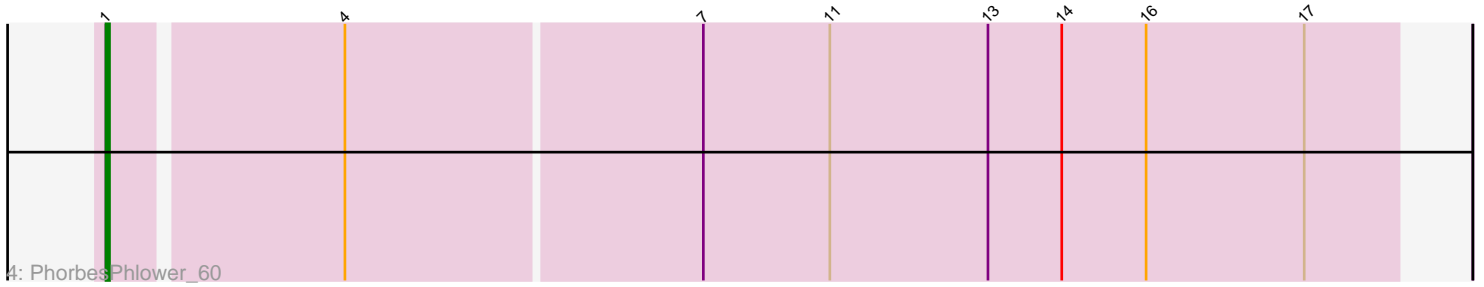
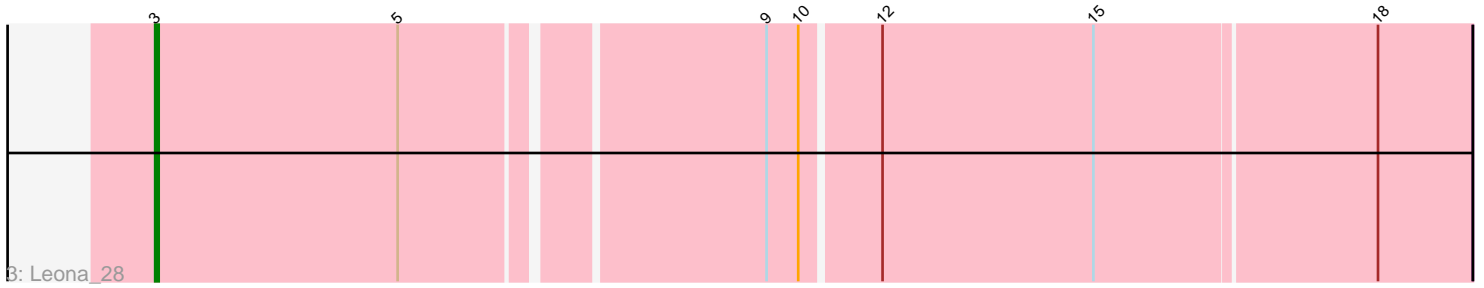
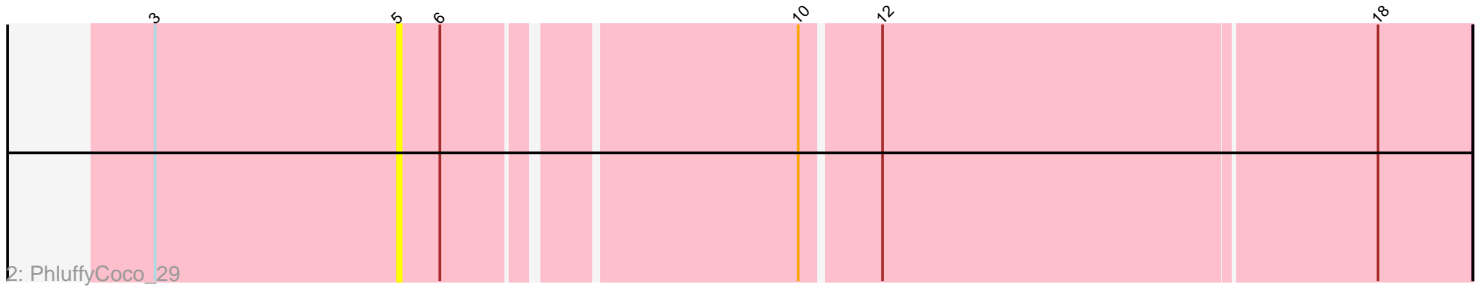
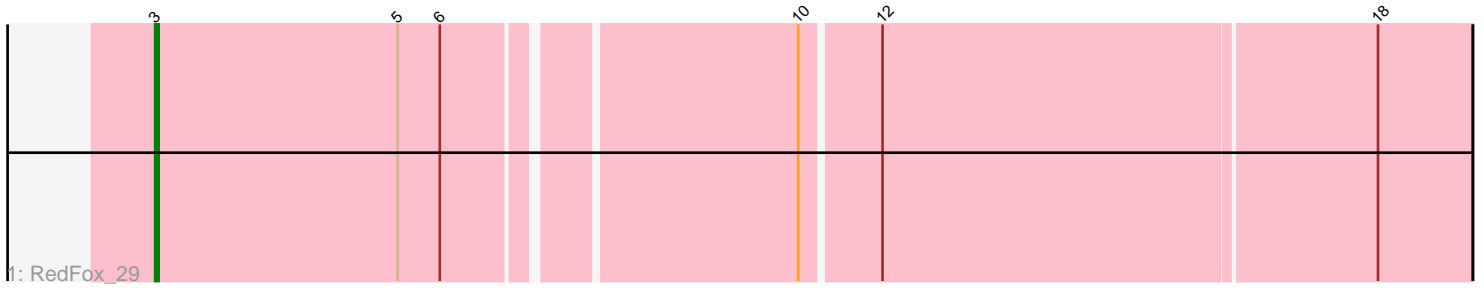


Pham 164273



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 164273 Report

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

Pham number 164273 has 5 members, 2 are drafts.

Phages represented in each track:

- Track 1 : RedFox_29
- Track 2 : PhluffyCoco_29
- Track 3 : Leona_28
- Track 4 : PhorbesPhlower_60
- Track 5 : CN1A_70

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

The start number called the most often in the published annotations is 3, it was called in 2 of the 3 non-draft genes in the pham.

Genes that call this "Most Annotated" start:

- Leona_28, RedFox_29,

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

- PhluffyCoco_29,

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

- CN1A_70, PhorbesPhlower_60,

Summary by start number:

Start 1:

- Found in 1 of 5 (20.0%) of genes in pham
- Manual Annotations of this start: 1 of 3
- Called 100.0% of time when present
- Phage (with cluster) where this start called: PhorbesPhlower_60 (DH),

Start 2:

- Found in 1 of 5 (20.0%) of genes in pham
- No Manual Annotations of this start.
- Called 100.0% of time when present
- Phage (with cluster) where this start called: CN1A_70 (singleton),

Start 3:

- Found in 3 of 5 (60.0%) of genes in pham
- Manual Annotations of this start: 2 of 3
- Called 66.7% of time when present
- Phage (with cluster) where this start called: Leona_28 (AS3), RedFox_29 (AS3),

Start 5:

- Found in 3 of 5 (60.0%) of genes in pham
- No Manual Annotations of this start.
- Called 33.3% of time when present
- Phage (with cluster) where this start called: PhluffyCoco_29 (AS3),

Summary by clusters:

There are 3 clusters represented in this pham: AS3, singleton, DH,

Info for manual annotations of cluster AS3:

- Start number 3 was manually annotated 2 times for cluster AS3.

Info for manual annotations of cluster DH:

- Start number 1 was manually annotated 1 time for cluster DH.

Gene Information:

Gene: CN1A_70 Start: 53137, Stop: 52757, Start Num: 2

Candidate Starts for CN1A_70:

(2, 53137), (8, 52954), (12, 52921),

Gene: Leona_28 Start: 20593, Stop: 20234, Start Num: 3

Candidate Starts for Leona_28:

(Start: 3 @20593 has 2 MA's), (5, 20524), (9, 20428), (10, 20419), (12, 20398), (15, 20338), (18, 20260),

Gene: PhluffyCoco_29 Start: 20437, Stop: 20147, Start Num: 5

Candidate Starts for PhluffyCoco_29:

(Start: 3 @20506 has 2 MA's), (5, 20437), (6, 20425), (10, 20332), (12, 20311), (18, 20173),

Gene: PhorbesPhlower_60 Start: 36317, Stop: 36676, Start Num: 1

Candidate Starts for PhorbesPhlower_60:

(Start: 1 @36317 has 1 MA's), (4, 36380), (7, 36479), (11, 36515), (13, 36560), (14, 36581), (16, 36605), (17, 36650),

Gene: RedFox_29 Start: 20519, Stop: 20160, Start Num: 3

Candidate Starts for RedFox_29:

(Start: 3 @20519 has 2 MA's), (5, 20450), (6, 20438), (10, 20345), (12, 20324), (18, 20186),