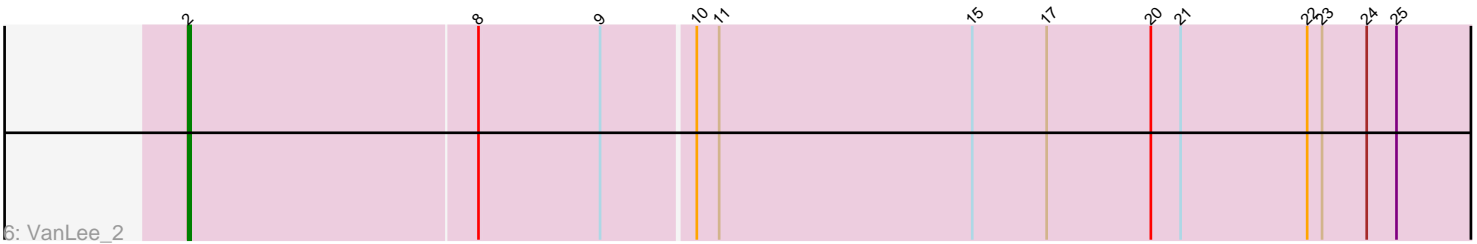
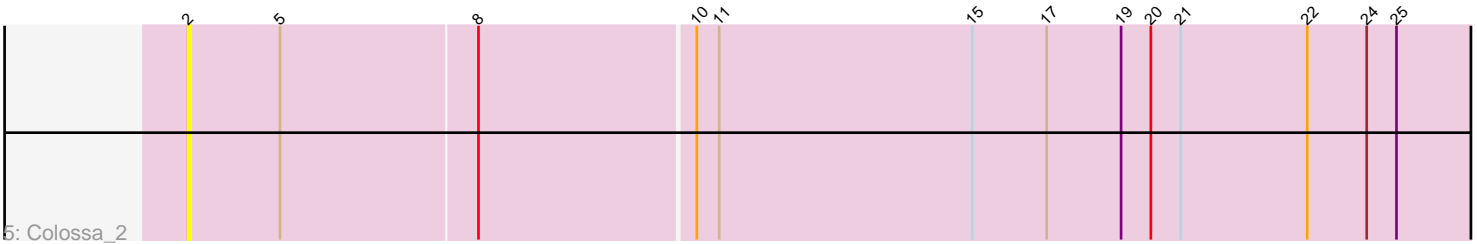
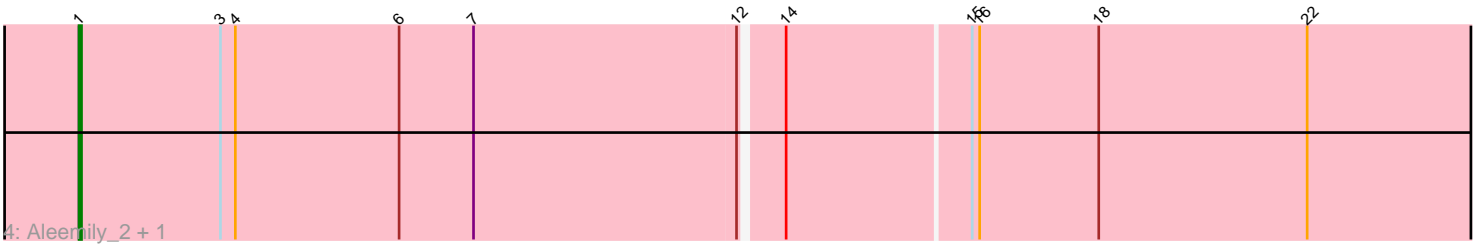
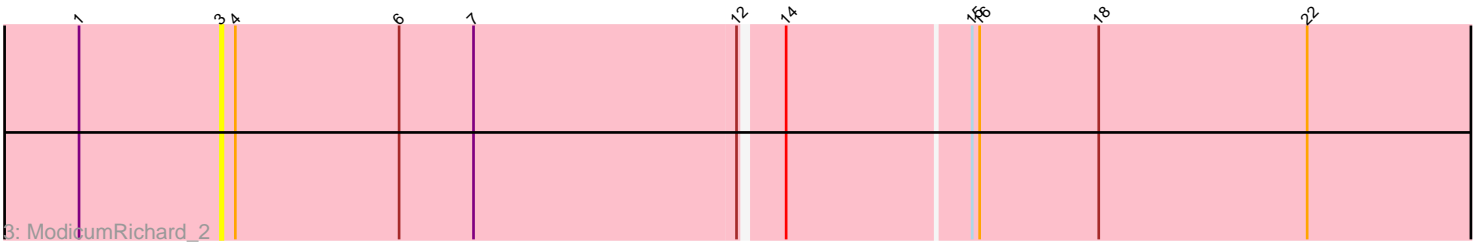
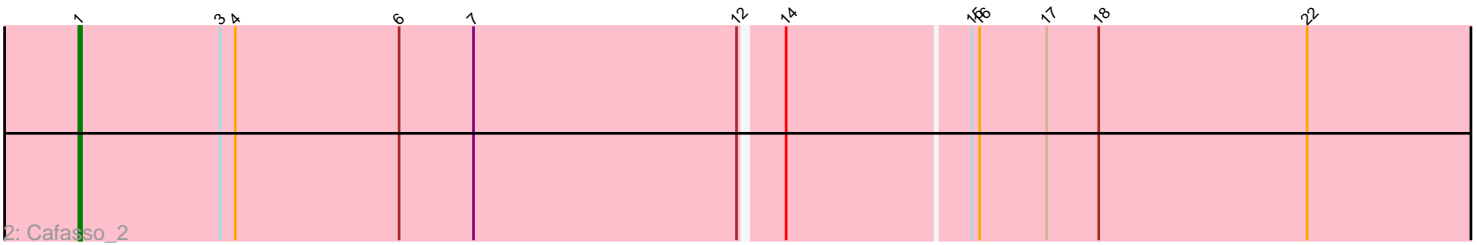
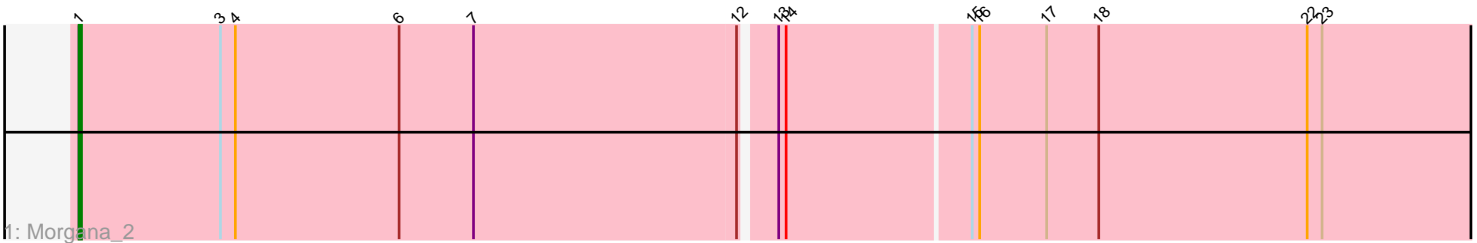


Pham 281253



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 281253 Report

This analysis was run 02/07/26 on database version 634.

Pham number 281253 has 7 members, 2 are drafts.

Phages represented in each track:

- Track 1 : Morgana_2
- Track 2 : Cafasso_2
- Track 3 : ModicumRichard_2
- Track 4 : Aleemily_2, ObLaDi_2
- Track 5 : Colossa_2
- Track 6 : VanLee_2

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 4 of the 5 non-draft genes in the pham.

Genes that call this "Most Annotated" start:

- Aleemily_2, Cafasso_2, Morgana_2, ObLaDi_2,

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

- ModicumRichard_2,

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

- Colossa_2, VanLee_2,

Summary by start number:

Start 1:

- Found in 5 of 7 (71.4%) of genes in pham
- Manual Annotations of this start: 4 of 5
- Called 80.0% of time when present
- Phage (with cluster) where this start called: Aleemily_2 (DZ), Cafasso_2 (DZ), Morgana_2 (DZ), ObLaDi_2 (DZ),

Start 2:

- Found in 2 of 7 (28.6%) of genes in pham
- Manual Annotations of this start: 1 of 5
- Called 100.0% of time when present

- Phage (with cluster) where this start called: Colossa_2 (KA), VanLee_2 (KA),

Start 3:

- Found in 5 of 7 (71.4%) of genes in pham
- No Manual Annotations of this start.
- Called 20.0% of time when present
- Phage (with cluster) where this start called: ModicumRichard_2 (DZ),

Summary by clusters:

There are 2 clusters represented in this pham: KA, DZ,

Info for manual annotations of cluster DZ:

- Start number 1 was manually annotated 4 times for cluster DZ.

Info for manual annotations of cluster KA:

- Start number 2 was manually annotated 1 time for cluster KA.

Gene Information:

Gene: Aleemily_2 Start: 374, Stop: 967, Start Num: 1

Candidate Starts for Aleemily_2:

(Start: 1 @374 has 4 MA's), (3, 431), (4, 437), (6, 503), (7, 533), (12, 638), (14, 653), (15, 725), (16, 728), (18, 776), (22, 860),

Gene: Cafasso_2 Start: 374, Stop: 967, Start Num: 1

Candidate Starts for Cafasso_2:

(Start: 1 @374 has 4 MA's), (3, 431), (4, 437), (6, 503), (7, 533), (12, 638), (14, 653), (15, 725), (16, 728), (17, 755), (18, 776), (22, 860),

Gene: Colossa_2 Start: 370, Stop: 900, Start Num: 2

Candidate Starts for Colossa_2:

(Start: 2 @370 has 1 MA's), (5, 406), (8, 484), (10, 568), (11, 577), (15, 679), (17, 709), (19, 739), (20, 751), (21, 763), (22, 814), (24, 838), (25, 850),

Gene: ModicumRichard_2 Start: 431, Stop: 967, Start Num: 3

Candidate Starts for ModicumRichard_2:

(Start: 1 @374 has 4 MA's), (3, 431), (4, 437), (6, 503), (7, 533), (12, 638), (14, 653), (15, 725), (16, 728), (18, 776), (22, 860),

Gene: Morgana_2 Start: 379, Stop: 972, Start Num: 1

Candidate Starts for Morgana_2:

(Start: 1 @379 has 4 MA's), (3, 436), (4, 442), (6, 508), (7, 538), (12, 643), (13, 655), (14, 658), (15, 730), (16, 733), (17, 760), (18, 781), (22, 865), (23, 871),

Gene: ObLaDi_2 Start: 374, Stop: 967, Start Num: 1

Candidate Starts for ObLaDi_2:

(Start: 1 @374 has 4 MA's), (3, 431), (4, 437), (6, 503), (7, 533), (12, 638), (14, 653), (15, 725), (16, 728), (18, 776), (22, 860),

Gene: VanLee_2 Start: 366, Stop: 896, Start Num: 2

Candidate Starts for VanLee_2:

(Start: 2 @366 has 1 MA's), (8, 480), (9, 528), (10, 564), (11, 573), (15, 675), (17, 705), (20, 747), (21, 759), (22, 810), (23, 816), (24, 834), (25, 846),