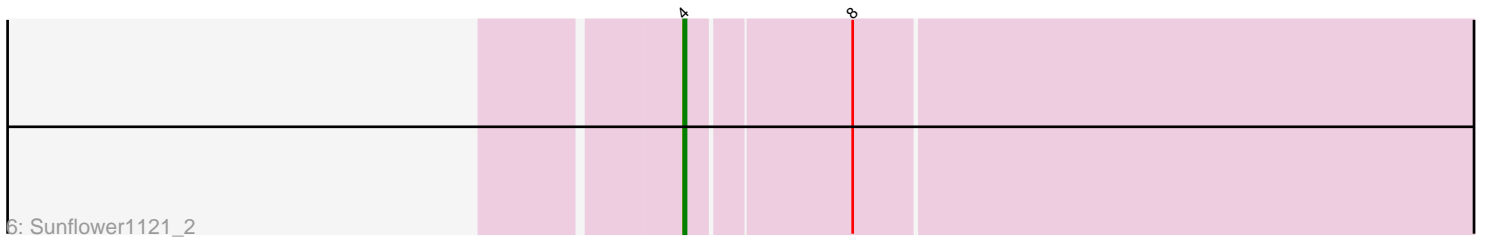
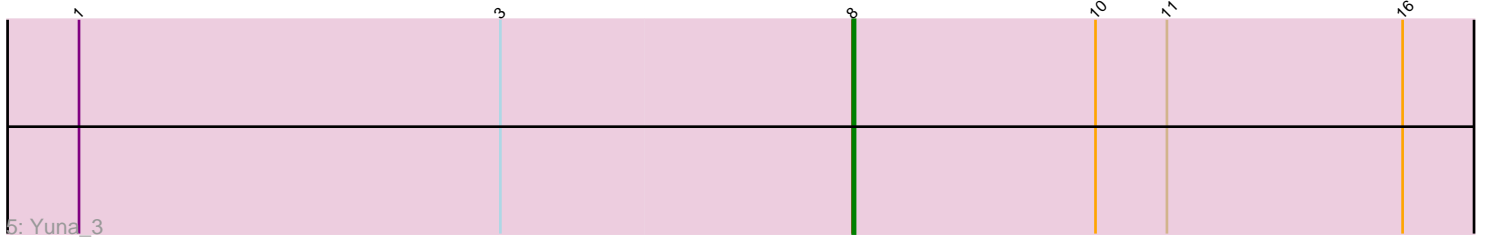
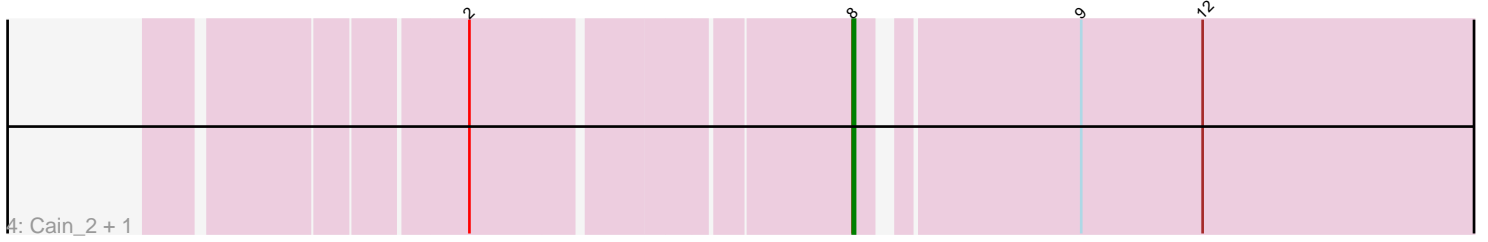
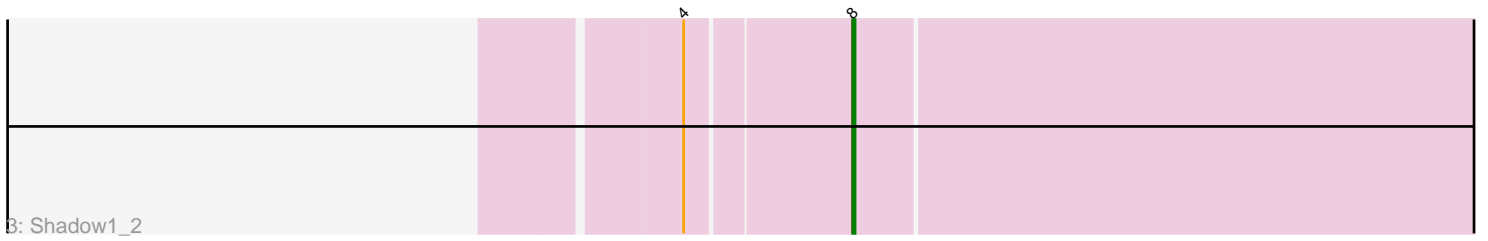
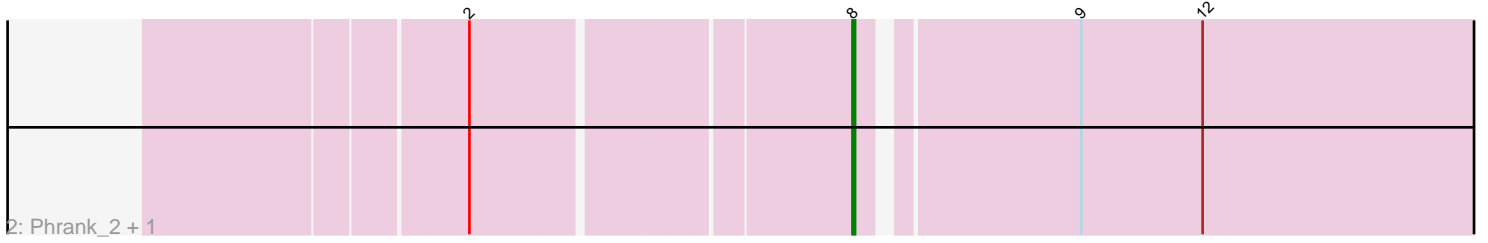
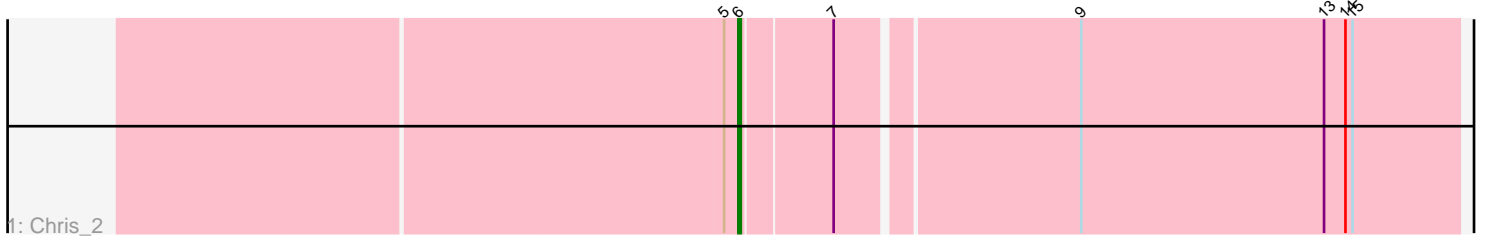


Pham 194592



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 194592 Report

This analysis was run 11/02/24 on database version 579.

Pham number 194592 has 8 members, 0 are drafts.

Phages represented in each track:

- Track 1 : Chris_2
- Track 2 : Phrank_2, Tierra_2
- Track 3 : Shadow1_2
- Track 4 : Cain_2, Bryler_2
- Track 5 : Yuna_3
- Track 6 : Sunflower1121_2

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

The start number called the most often in the published annotations is 8, it was called in 6 of the 8 non-draft genes in the pham.

Genes that call this "Most Annotated" start:

- Bryler_2, Cain_2, Phrank_2, Shadow1_2, Tierra_2, Yuna_3,

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

- Sunflower1121_2,

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

- Chris_2,

Summary by start number:

Start 4:

- Found in 2 of 8 (25.0%) of genes in pham
- Manual Annotations of this start: 1 of 8
- Called 50.0% of time when present
- Phage (with cluster) where this start called: Sunflower1121_2 (K6),

Start 6:

- Found in 1 of 8 (12.5%) of genes in pham
- Manual Annotations of this start: 1 of 8
- Called 100.0% of time when present
- Phage (with cluster) where this start called: Chris_2 (K1),

Start 8:

- Found in 7 of 8 (87.5%) of genes in pham
- Manual Annotations of this start: 6 of 8
- Called 85.7% of time when present
- Phage (with cluster) where this start called: Bryler_2 (K6), Cain_2 (K6), Phrank_2 (K6), Shadow1_2 (K6), Tierra_2 (K6), Yuna_3 (K6),

Summary by clusters:

There are 2 clusters represented in this pham: K1, K6,

Info for manual annotations of cluster K1:

- Start number 6 was manually annotated 1 time for cluster K1.

Info for manual annotations of cluster K6:

- Start number 4 was manually annotated 1 time for cluster K6.
- Start number 8 was manually annotated 6 times for cluster K6.

Gene Information:

Gene: Bryler_2 Start: 388, Stop: 648, Start Num: 8

Candidate Starts for Bryler_2:

(2, 241), (Start: 8 @388 has 6 MA's), (9, 472), (12, 523),

Gene: Cain_2 Start: 388, Stop: 648, Start Num: 8

Candidate Starts for Cain_2:

(2, 241), (Start: 8 @388 has 6 MA's), (9, 472), (12, 523),

Gene: Chris_2 Start: 375, Stop: 665, Start Num: 6

Candidate Starts for Chris_2:

(5, 369), (Start: 6 @375 has 1 MA's), (7, 411), (9, 507), (13, 609), (14, 618), (15, 621),

Gene: Phrank_2 Start: 394, Stop: 654, Start Num: 8

Candidate Starts for Phrank_2:

(2, 247), (Start: 8 @394 has 6 MA's), (9, 478), (12, 529),

Gene: Shadow1_2 Start: 396, Stop: 665, Start Num: 8

Candidate Starts for Shadow1_2:

(Start: 4 @330 has 1 MA's), (Start: 8 @396 has 6 MA's),

Gene: Sunflower1121_2 Start: 329, Stop: 664, Start Num: 4

Candidate Starts for Sunflower1121_2:

(Start: 4 @329 has 1 MA's), (Start: 8 @395 has 6 MA's),

Gene: Tierra_2 Start: 395, Stop: 655, Start Num: 8

Candidate Starts for Tierra_2:

(2, 248), (Start: 8 @395 has 6 MA's), (9, 479), (12, 530),

Gene: Yuna_3 Start: 525, Stop: 797, Start Num: 8

Candidate Starts for Yuna_3:

(1, 201), (3, 378), (Start: 8 @525 has 6 MA's), (10, 627), (11, 657), (16, 756),