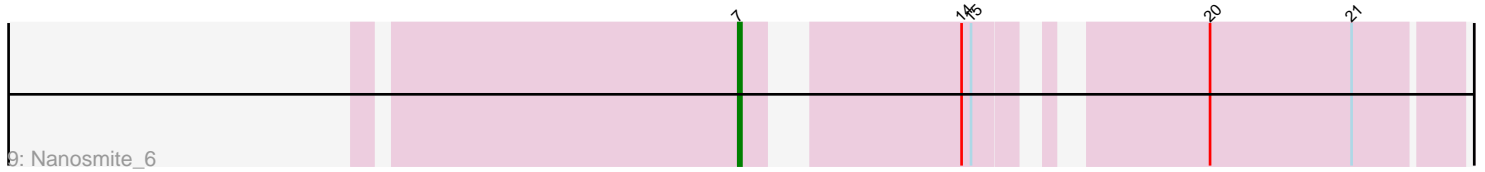
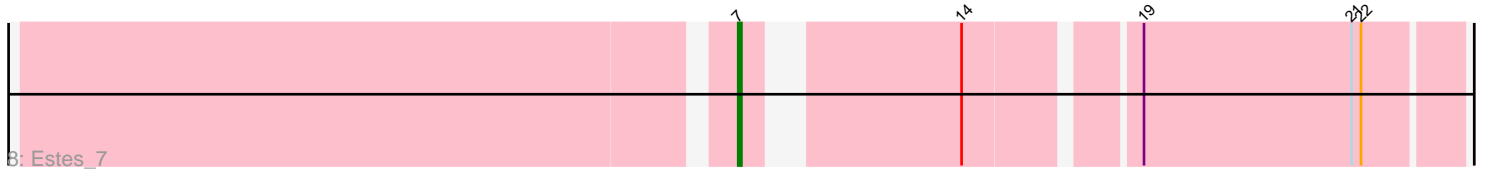
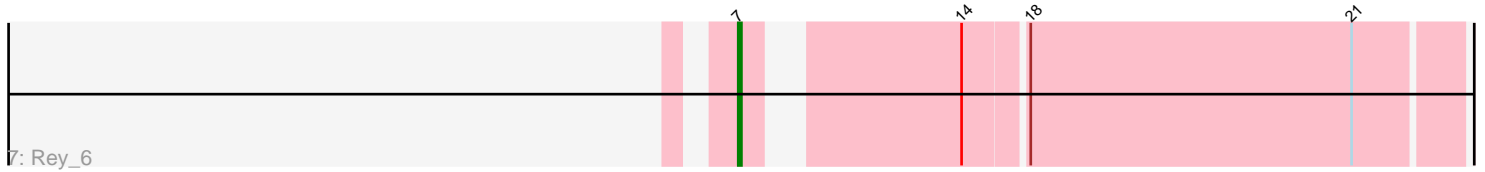
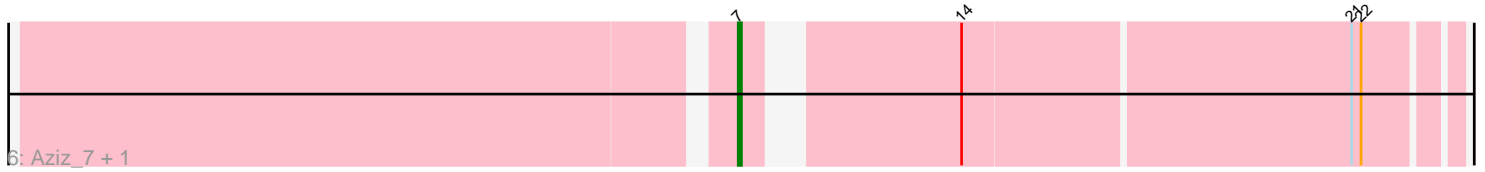
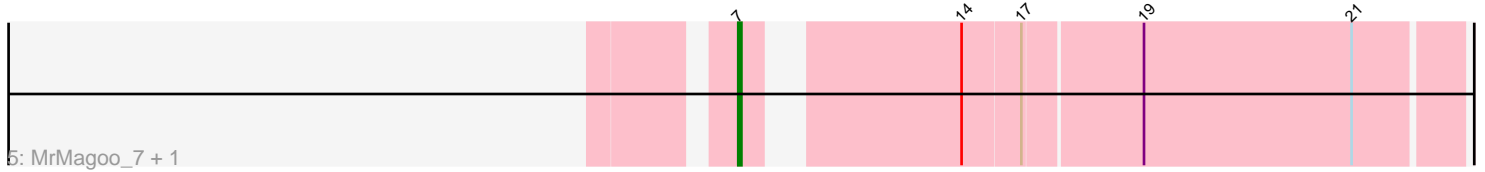
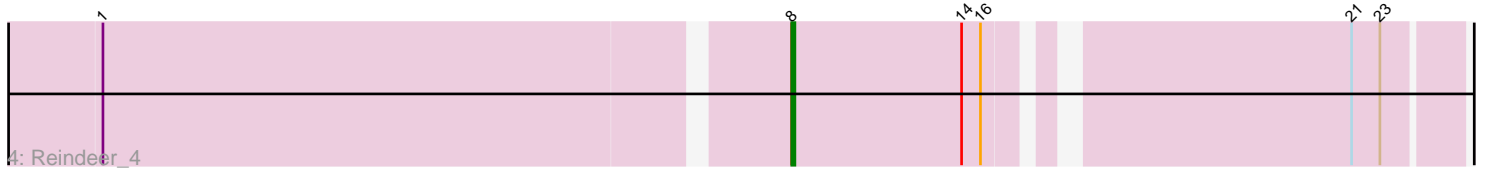
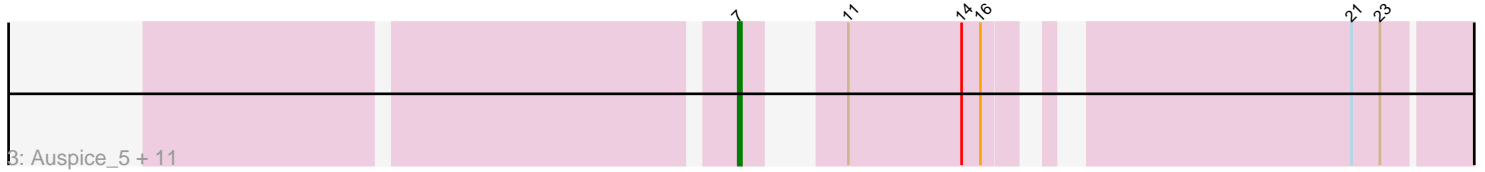
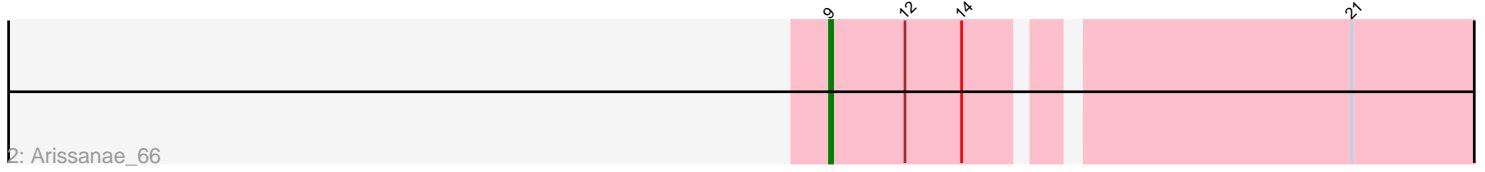
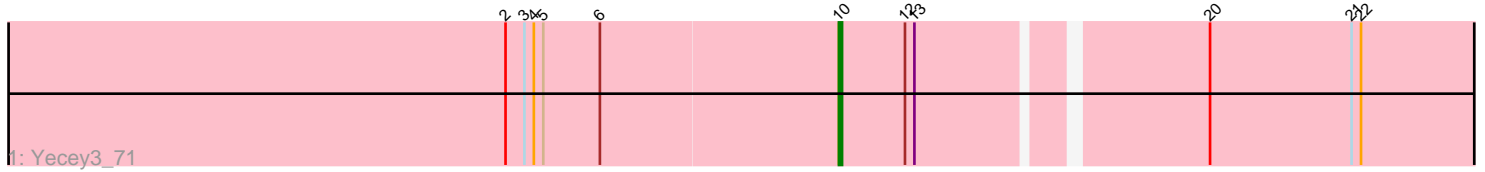


Pham 171755



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 171755 Report

This analysis was run 07/10/24 on database version 566.

Pham number 171755 has 22 members, 0 are drafts.

Phages represented in each track:

- Track 1 : Yecey3_71
- Track 2 : Arissanae_66
- Track 3 : Auspice_5, Diminimus_6, TyDawg_5, Dulcita_6, PegLeg_5, Bongo_5, LilhomieP_5, Glaske16_6, SlimJimmy_5, Bricole_5, Skinny_6, IPhane7_5
- Track 4 : Reindeer_4
- Track 5 : MrMagoo_7, GardenSalsa_7
- Track 6 : Aziz_7, GenevaB15_7
- Track 7 : Rey_6
- Track 8 : Estes_7
- Track 9 : Nanosmite_6

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

The start number called the most often in the published annotations is 7, it was called in 19 of the 22 non-draft genes in the pham.

Genes that call this "Most Annotated" start:

- Auspice_5, Aziz_7, Bongo_5, Bricole_5, Diminimus_6, Dulcita_6, Estes_7, GardenSalsa_7, GenevaB15_7, Glaske16_6, IPhane7_5, LilhomieP_5, MrMagoo_7, Nanosmite_6, PegLeg_5, Rey_6, Skinny_6, SlimJimmy_5, TyDawg_5,

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

-

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

- Arissanae_66, Reindeer_4, Yecey3_71,

Summary by start number:

Start 7:

- Found in 19 of 22 (86.4%) of genes in pham
- Manual Annotations of this start: 19 of 22
- Called 100.0% of time when present

- Phage (with cluster) where this start called: Auspice_5 (M1), Aziz_7 (M2), Bongo_5 (M1), Bricole_5 (M1), Diminimus_6 (M1), Dulcita_6 (M1), Estes_7 (M2), GardenSalsa_7 (M2), GenevaB15_7 (M2), Glaske16_6 (M1), IPHane7_5 (M1), LilhomieP_5 (M1), MrMagoo_7 (M2), Nanosmite_6 (M3), PegLeg_5 (M1), Rey_6 (M2), Skinny_6 (M1), SlimJimmy_5 (M1), TyDawg_5 (M1),

Start 8:

- Found in 1 of 22 (4.5%) of genes in pham
- Manual Annotations of this start: 1 of 22
- Called 100.0% of time when present
- Phage (with cluster) where this start called: Reindeer_4 (M1),

Start 9:

- Found in 1 of 22 (4.5%) of genes in pham
- Manual Annotations of this start: 1 of 22
- Called 100.0% of time when present
- Phage (with cluster) where this start called: Arissanae_66 (A9),

Start 10:

- Found in 1 of 22 (4.5%) of genes in pham
- Manual Annotations of this start: 1 of 22
- Called 100.0% of time when present
- Phage (with cluster) where this start called: Yecey3_71 (A9),

Summary by clusters:

There are 4 clusters represented in this pham: A9, M1, M3, M2,

Info for manual annotations of cluster A9:

- Start number 9 was manually annotated 1 time for cluster A9.
- Start number 10 was manually annotated 1 time for cluster A9.

Info for manual annotations of cluster M1:

- Start number 7 was manually annotated 12 times for cluster M1.
- Start number 8 was manually annotated 1 time for cluster M1.

Info for manual annotations of cluster M2:

- Start number 7 was manually annotated 6 times for cluster M2.

Info for manual annotations of cluster M3:

- Start number 7 was manually annotated 1 time for cluster M3.

Gene Information:

Gene: Arissanae_66 Start: 41220, Stop: 40978, Start Num: 9

Candidate Starts for Arissanae_66:

(Start: 9 @41220 has 1 MA's), (12, 41196), (14, 41178), (21, 41067),

Gene: Auspice_5 Start: 2366, Stop: 2133, Start Num: 7

Candidate Starts for Auspice_5:

(Start: 7 @2366 has 19 MA's), (11, 2348), (14, 2312), (16, 2306), (21, 2207), (23, 2198),

Gene: Aziz_7 Start: 2636, Stop: 2391, Start Num: 7

Candidate Starts for Aziz_7:

(Start: 7 @2636 has 19 MA's), (14, 2579), (21, 2459), (22, 2456),

Gene: Bongo_5 Start: 2366, Stop: 2133, Start Num: 7

Candidate Starts for Bongo_5:

(Start: 7 @2366 has 19 MA's), (11, 2348), (14, 2312), (16, 2306), (21, 2207), (23, 2198),

Gene: Bricole_5 Start: 2365, Stop: 2132, Start Num: 7

Candidate Starts for Bricole_5:

(Start: 7 @2365 has 19 MA's), (11, 2347), (14, 2311), (16, 2305), (21, 2206), (23, 2197),

Gene: Diminimus_6 Start: 2365, Stop: 2132, Start Num: 7

Candidate Starts for Diminimus_6:

(Start: 7 @2365 has 19 MA's), (11, 2347), (14, 2311), (16, 2305), (21, 2206), (23, 2197),

Gene: Dulcita_6 Start: 2365, Stop: 2132, Start Num: 7

Candidate Starts for Dulcita_6:

(Start: 7 @2365 has 19 MA's), (11, 2347), (14, 2311), (16, 2305), (21, 2206), (23, 2197),

Gene: Estes_7 Start: 2641, Stop: 2399, Start Num: 7

Candidate Starts for Estes_7:

(Start: 7 @2641 has 19 MA's), (14, 2584), (19, 2536), (21, 2470), (22, 2467),

Gene: GardenSalsa_7 Start: 2660, Stop: 2412, Start Num: 7

Candidate Starts for GardenSalsa_7:

(Start: 7 @2660 has 19 MA's), (14, 2603), (17, 2585), (19, 2549), (21, 2483),

Gene: GenevaB15_7 Start: 2636, Stop: 2391, Start Num: 7

Candidate Starts for GenevaB15_7:

(Start: 7 @2636 has 19 MA's), (14, 2579), (21, 2459), (22, 2456),

Gene: Glaske16_6 Start: 2365, Stop: 2132, Start Num: 7

Candidate Starts for Glaske16_6:

(Start: 7 @2365 has 19 MA's), (11, 2347), (14, 2311), (16, 2305), (21, 2206), (23, 2197),

Gene: IPHane7_5 Start: 2366, Stop: 2133, Start Num: 7

Candidate Starts for IPHane7_5:

(Start: 7 @2366 has 19 MA's), (11, 2348), (14, 2312), (16, 2306), (21, 2207), (23, 2198),

Gene: LilhomieP_5 Start: 2366, Stop: 2133, Start Num: 7

Candidate Starts for LilhomieP_5:

(Start: 7 @2366 has 19 MA's), (11, 2348), (14, 2312), (16, 2306), (21, 2207), (23, 2198),

Gene: MrMagoo_7 Start: 2660, Stop: 2412, Start Num: 7

Candidate Starts for MrMagoo_7:

(Start: 7 @2660 has 19 MA's), (14, 2603), (17, 2585), (19, 2549), (21, 2483),

Gene: Nanosmite_6 Start: 2681, Stop: 2448, Start Num: 7

Candidate Starts for Nanosmite_6:

(Start: 7 @2681 has 19 MA's), (14, 2624), (15, 2621), (20, 2564), (21, 2519),

Gene: PegLeg_5 Start: 2365, Stop: 2132, Start Num: 7

Candidate Starts for PegLeg_5:

(Start: 7 @2365 has 19 MA's), (11, 2347), (14, 2311), (16, 2305), (21, 2206), (23, 2197),

Gene: Reindeer_4 Start: 2200, Stop: 1967, Start Num: 8

Candidate Starts for Reindeer_4:

(1, 2410), (Start: 8 @2200 has 1 MA's), (14, 2146), (16, 2140), (21, 2038), (23, 2029),

Gene: Rey_6 Start: 2648, Stop: 2400, Start Num: 7

Candidate Starts for Rey_6:

(Start: 7 @2648 has 19 MA's), (14, 2591), (18, 2573), (21, 2471),

Gene: Skinny_6 Start: 2365, Stop: 2132, Start Num: 7

Candidate Starts for Skinny_6:

(Start: 7 @2365 has 19 MA's), (11, 2347), (14, 2311), (16, 2305), (21, 2206), (23, 2197),

Gene: SlimJimmy_5 Start: 2365, Stop: 2132, Start Num: 7

Candidate Starts for SlimJimmy_5:

(Start: 7 @2365 has 19 MA's), (11, 2347), (14, 2311), (16, 2305), (21, 2206), (23, 2197),

Gene: TyDawg_5 Start: 2366, Stop: 2133, Start Num: 7

Candidate Starts for TyDawg_5:

(Start: 7 @2366 has 19 MA's), (11, 2348), (14, 2312), (16, 2306), (21, 2207), (23, 2198),

Gene: Yecey3_71 Start: 41130, Stop: 40897, Start Num: 10

Candidate Starts for Yecey3_71:

(2, 41235), (3, 41229), (4, 41226), (5, 41223), (6, 41205), (Start: 10 @41130 has 1 MA's), (12, 41109), (13, 41106), (20, 41022), (21, 40977), (22, 40974),