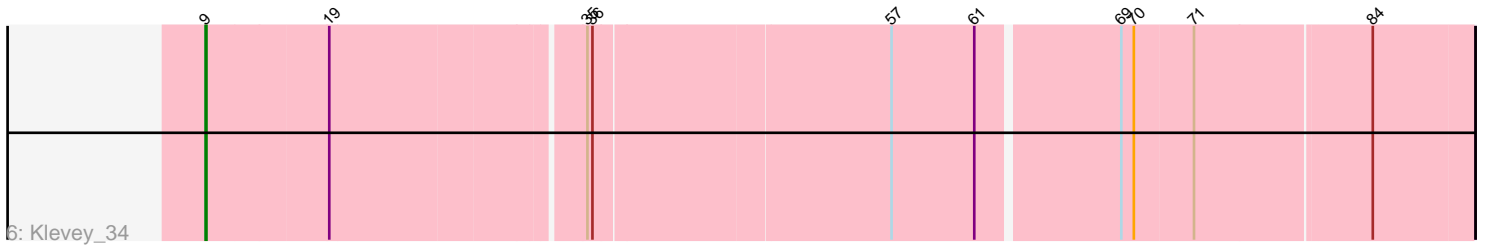
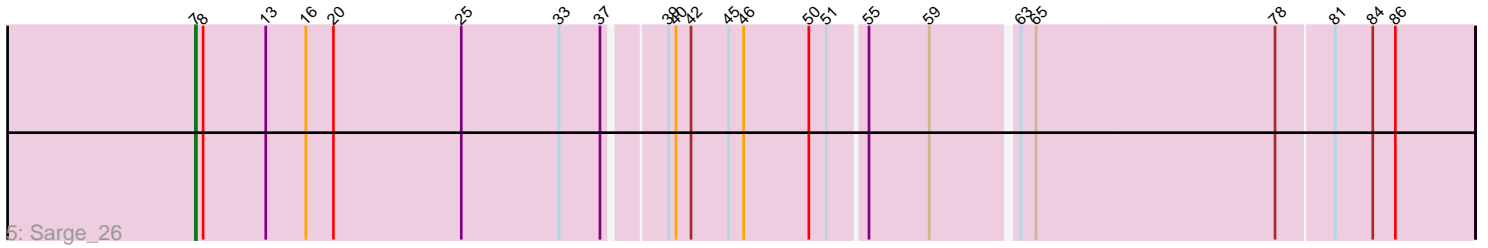
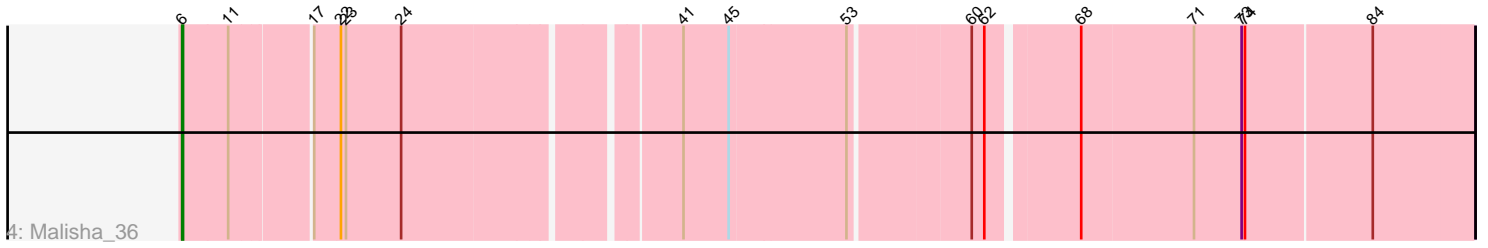
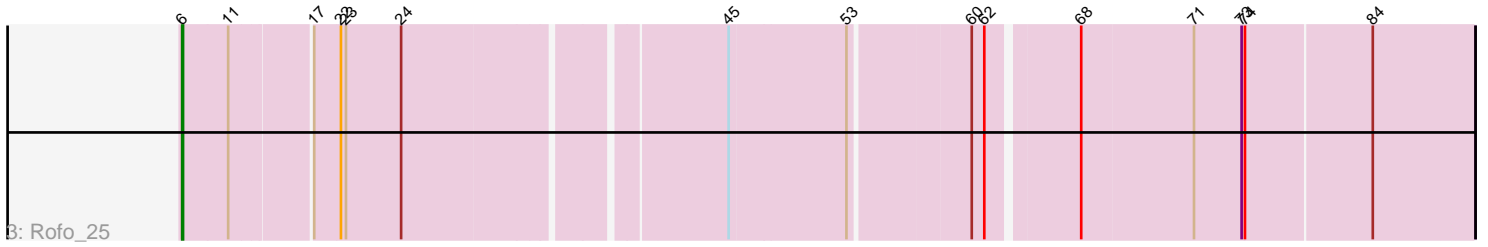
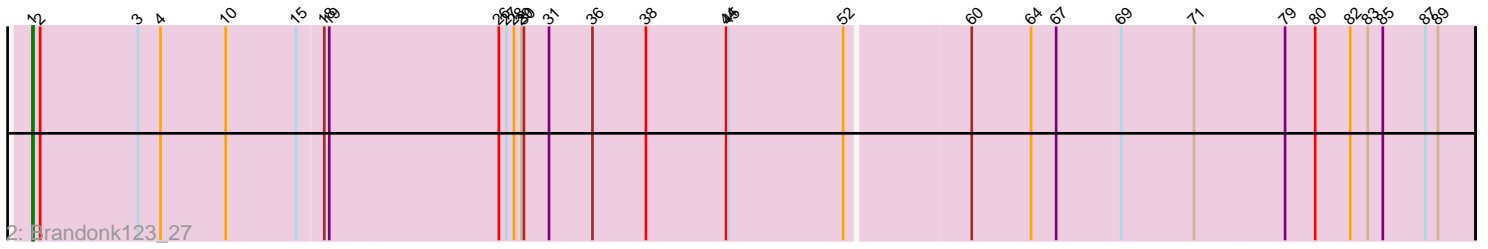
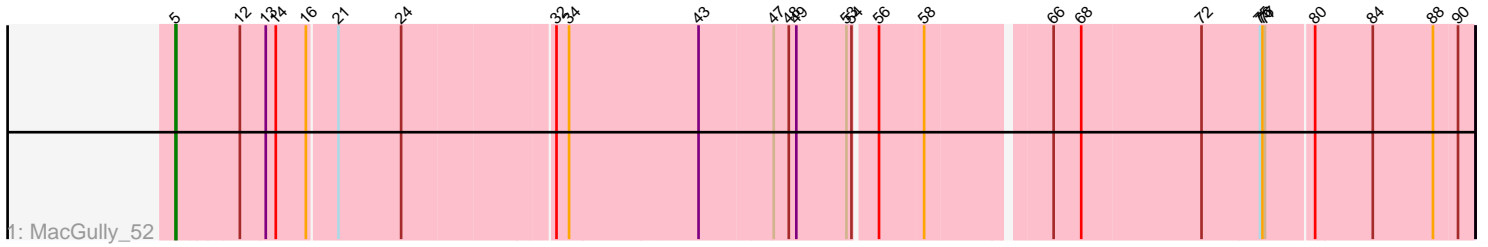


Pham 8675



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 8675 Report

This analysis was run 07/09/24 on database version 566.

Pham number 8675 has 6 members, 0 are drafts.

Phages represented in each track:

- Track 1 : MacGully_52
- Track 2 : Brandonk123_27
- Track 3 : Rofo_25
- Track 4 : Malisha_36
- Track 5 : Sarge_26
- Track 6 : Klevey_34

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

The start number called the most often in the published annotations is 6, it was called in 2 of the 6 non-draft genes in the pham.

Genes that call this "Most Annotated" start:

- Malisha_36, Rofo_25,

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

-

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

- Brandonk123_27, Klevey_34, MacGully_52, Sarge_26,

Summary by start number:

Start 1:

- Found in 1 of 6 (16.7%) of genes in pham
- Manual Annotations of this start: 1 of 6
- Called 100.0% of time when present
- Phage (with cluster) where this start called: Brandonk123_27 (DE1),

Start 5:

- Found in 1 of 6 (16.7%) of genes in pham
- Manual Annotations of this start: 1 of 6
- Called 100.0% of time when present
- Phage (with cluster) where this start called: MacGully_52 (CR),

Start 6:

- Found in 2 of 6 (33.3%) of genes in pham
- Manual Annotations of this start: 2 of 6
- Called 100.0% of time when present
- Phage (with cluster) where this start called: Malisha_36 (DN), Rofo_25 (DE1),

Start 7:

- Found in 1 of 6 (16.7%) of genes in pham
- Manual Annotations of this start: 1 of 6
- Called 100.0% of time when present
- Phage (with cluster) where this start called: Sarge_26 (FB),

Start 9:

- Found in 1 of 6 (16.7%) of genes in pham
- Manual Annotations of this start: 1 of 6
- Called 100.0% of time when present
- Phage (with cluster) where this start called: Klevey_34 (FH),

Summary by clusters:

There are 5 clusters represented in this pham: FH, DN, FB, CR, DE1,

Info for manual annotations of cluster CR:

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

Info for manual annotations of cluster DE1:

- Start number 1 was manually annotated 1 time for cluster DE1.
- Start number 6 was manually annotated 1 time for cluster DE1.

Info for manual annotations of cluster DN:

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

Info for manual annotations of cluster FB:

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

Info for manual annotations of cluster FH:

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

Gene Information:

Gene: Brandonk123_27 Start: 19364, Stop: 21058, Start Num: 1

Candidate Starts for Brandonk123_27:

(Start: 1 @19364 has 1 MA's), (2, 19373), (3, 19490), (4, 19517), (10, 19595), (15, 19679), (18, 19709), (19, 19715), (26, 19916), (27, 19925), (28, 19934), (29, 19943), (30, 19946), (31, 19976), (36, 20027), (38, 20090), (44, 20186), (45, 20189), (52, 20324), (60, 20465), (64, 20534), (67, 20564), (69, 20639), (71, 20726), (79, 20834), (80, 20870), (82, 20909), (83, 20930), (85, 20948), (87, 20999), (89, 21014),

Gene: Klevey_34 Start: 25869, Stop: 27302, Start Num: 9

Candidate Starts for Klevey_34:

(Start: 9 @25869 has 1 MA's), (19, 26010), (35, 26292), (36, 26298), (57, 26640), (61, 26739), (69, 26901), (70, 26916), (71, 26985), (84, 27186),

Gene: MacGully_52 Start: 34011, Stop: 35462, Start Num: 5

Candidate Starts for MacGully_52:

(Start: 5 @34011 has 1 MA's), (12, 34083), (13, 34110), (14, 34122), (16, 34158), (21, 34188), (24, 34263), (32, 34431), (34, 34446), (43, 34596), (47, 34680), (48, 34698), (49, 34707), (53, 34767), (54, 34773), (56, 34797), (58, 34848), (66, 34980), (68, 35013), (72, 35154), (75, 35223), (76, 35226), (77, 35229), (80, 35280), (84, 35346), (88, 35418), (90, 35445),

Gene: Malisha_36 Start: 27449, Stop: 28879, Start Num: 6

Candidate Starts for Malisha_36:

(Start: 6 @27449 has 2 MA's), (11, 27503), (17, 27590), (22, 27620), (23, 27626), (24, 27692), (41, 27986), (45, 28040), (53, 28175), (60, 28310), (62, 28325), (68, 28424), (71, 28556), (73, 28613), (74, 28616), (84, 28763),

Gene: Rofo_25 Start: 19691, Stop: 21121, Start Num: 6

Candidate Starts for Rofo_25:

(Start: 6 @19691 has 2 MA's), (11, 19745), (17, 19832), (22, 19862), (23, 19868), (24, 19934), (45, 20282), (53, 20417), (60, 20552), (62, 20567), (68, 20666), (71, 20798), (73, 20855), (74, 20858), (84, 21005),

Gene: Sarge_26 Start: 19160, Stop: 20605, Start Num: 7

Candidate Starts for Sarge_26:

(Start: 7 @19160 has 1 MA's), (8, 19169), (13, 19238), (16, 19286), (20, 19319), (25, 19472), (33, 19583), (37, 19631), (39, 19691), (40, 19700), (42, 19718), (45, 19763), (46, 19778), (50, 19853), (51, 19874), (55, 19916), (59, 19985), (63, 20081), (65, 20099), (78, 20381), (81, 20441), (84, 20486), (86, 20513),