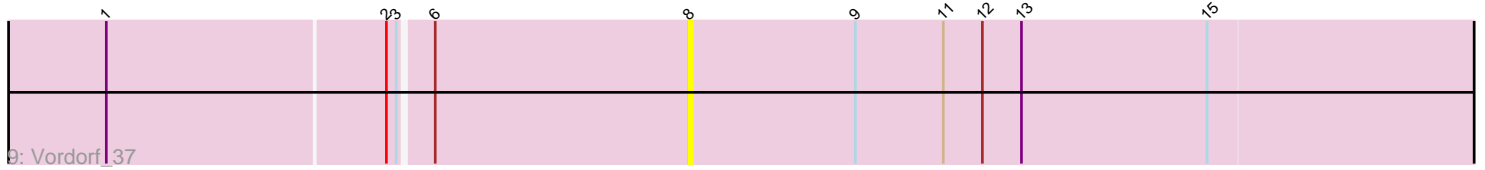
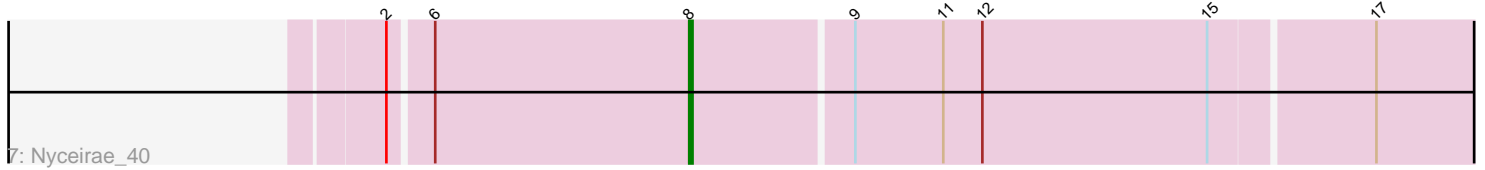
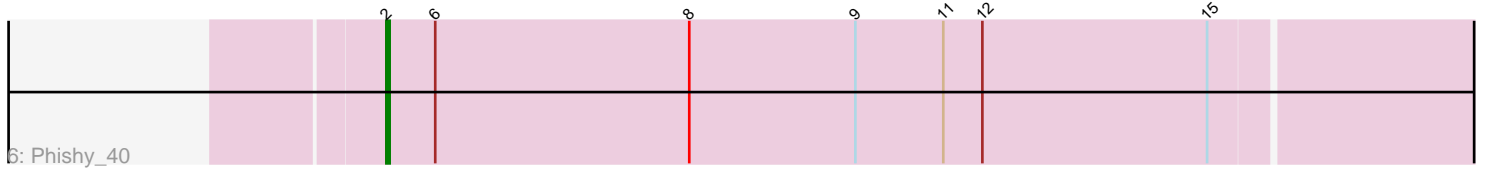
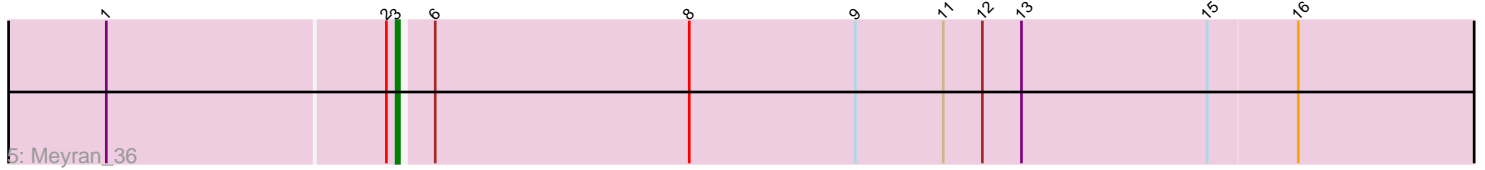
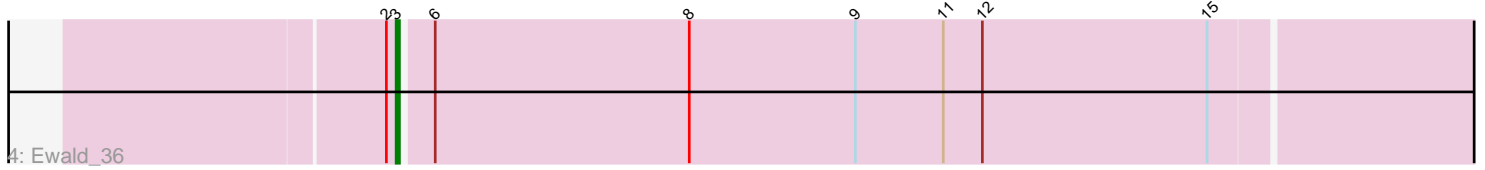
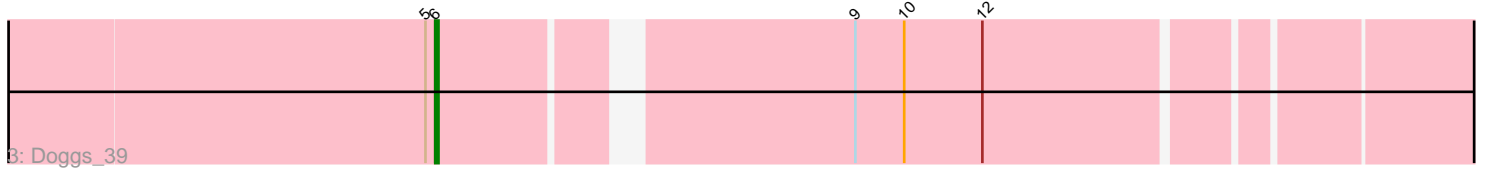
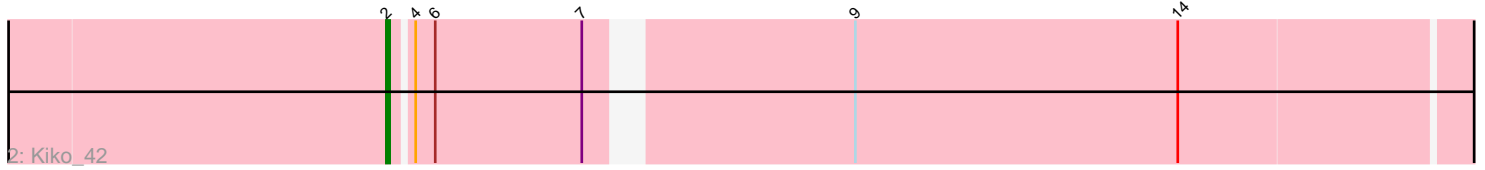
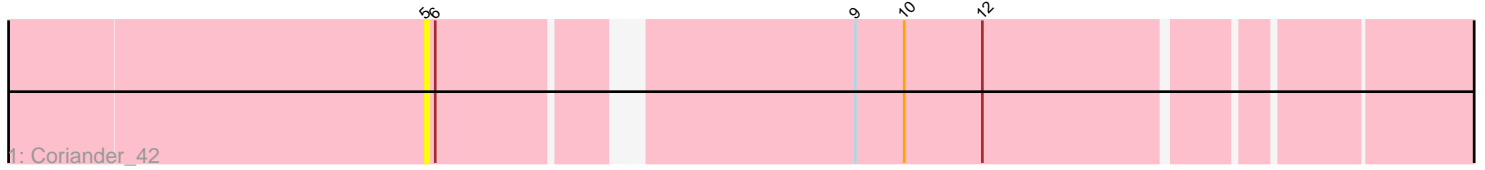


Pham 193105



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 193105 Report

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

Pham number 193105 has 9 members, 2 are drafts.

Phages represented in each track:

- Track 1 : Coriander_42
- Track 2 : Kiko_42
- Track 3 : Doggs_39
- Track 4 : Ewald_36
- Track 5 : Meyran_36
- Track 6 : Phishy_40
- Track 7 : Nyceirae_40
- Track 8 : Dogfish_36
- Track 9 : Vordorf_37

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

The start number called the most often in the published annotations is 2, it was called in 3 of the 7 non-draft genes in the pham.

Genes that call this "Most Annotated" start:

- Dogfish_36, Kiko_42, Phishy_40,

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

- Ewald_36, Meyran_36, Nyceirae_40, Vordorf_37,

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

- Coriander_42, Doggs_39,

Summary by start number:

Start 2:

- Found in 7 of 9 (77.8%) of genes in pham
- Manual Annotations of this start: 3 of 7
- Called 42.9% of time when present
- Phage (with cluster) where this start called: Dogfish_36 (DT), Kiko_42 (DB), Phishy_40 (DT),

Start 3:

- Found in 3 of 9 (33.3%) of genes in pham
- Manual Annotations of this start: 2 of 7
- Called 66.7% of time when present
- Phage (with cluster) where this start called: Ewald_36 (DT), Meyran_36 (DT),

Start 5:

- Found in 2 of 9 (22.2%) of genes in pham
- No Manual Annotations of this start.
- Called 50.0% of time when present
- Phage (with cluster) where this start called: Coriander_42 (DB),

Start 6:

- Found in 9 of 9 (100.0%) of genes in pham
- Manual Annotations of this start: 1 of 7
- Called 11.1% of time when present
- Phage (with cluster) where this start called: Doggs_39 (DB),

Start 8:

- Found in 5 of 9 (55.6%) of genes in pham
- Manual Annotations of this start: 1 of 7
- Called 40.0% of time when present
- Phage (with cluster) where this start called: Nyceirae_40 (DT), Vordorf_37 (DT),

Summary by clusters:

There are 2 clusters represented in this pham: DT, DB,

Info for manual annotations of cluster DB:

- Start number 2 was manually annotated 1 time for cluster DB.
- Start number 6 was manually annotated 1 time for cluster DB.

Info for manual annotations of cluster DT:

- Start number 2 was manually annotated 2 times for cluster DT.
- Start number 3 was manually annotated 2 times for cluster DT.
- Start number 8 was manually annotated 1 time for cluster DT.

Gene Information:

Gene: Coriander_42 Start: 32123, Stop: 32419, Start Num: 5

Candidate Starts for Coriander_42:

(5, 32123), (Start: 6 @32126 has 1 MA's), (9, 32240), (10, 32255), (12, 32279),

Gene: Dogfish_36 Start: 30380, Stop: 30739, Start Num: 2

Candidate Starts for Dogfish_36:

(Start: 2 @30380 has 3 MA's), (Start: 6 @30395 has 1 MA's), (9, 30524), (11, 30551), (12, 30563), (15, 30632), (17, 30680),

Gene: Doggs_39 Start: 32855, Stop: 33148, Start Num: 6

Candidate Starts for Doggs_39:

(5, 32852), (Start: 6 @32855 has 1 MA's), (9, 32969), (10, 32984), (12, 33008),

Gene: Ewald_36 Start: 30203, Stop: 30556, Start Num: 3

Candidate Starts for Ewald_36:

(Start: 2 @30200 has 3 MA's), (Start: 3 @30203 has 2 MA's), (Start: 6 @30212 has 1 MA's), (Start: 8 @30290 has 1 MA's), (9, 30341), (11, 30368), (12, 30380), (15, 30449),

Gene: Kiko_42 Start: 32442, Stop: 32807, Start Num: 2

Candidate Starts for Kiko_42:

(Start: 2 @32442 has 3 MA's), (4, 32448), (Start: 6 @32454 has 1 MA's), (7, 32499), (9, 32571), (14, 32670),

Gene: Meyran_36 Start: 31276, Stop: 31632, Start Num: 3

Candidate Starts for Meyran_36:

(1, 31189), (Start: 2 @31273 has 3 MA's), (Start: 3 @31276 has 2 MA's), (Start: 6 @31285 has 1 MA's), (Start: 8 @31363 has 1 MA's), (9, 31414), (11, 31441), (12, 31453), (13, 31465), (15, 31522), (16, 31549),

Gene: Nyceirae_40 Start: 31180, Stop: 31443, Start Num: 8

Candidate Starts for Nyceirae_40:

(Start: 2 @31090 has 3 MA's), (Start: 6 @31102 has 1 MA's), (Start: 8 @31180 has 1 MA's), (9, 31228), (11, 31255), (12, 31267), (15, 31336), (17, 31384),

Gene: Phishy_40 Start: 32258, Stop: 32617, Start Num: 2

Candidate Starts for Phishy_40:

(Start: 2 @32258 has 3 MA's), (Start: 6 @32273 has 1 MA's), (Start: 8 @32351 has 1 MA's), (9, 32402), (11, 32429), (12, 32441), (15, 32510),

Gene: Vordorf_37 Start: 30638, Stop: 30907, Start Num: 8

Candidate Starts for Vordorf_37:

(1, 30464), (Start: 2 @30548 has 3 MA's), (Start: 3 @30551 has 2 MA's), (Start: 6 @30560 has 1 MA's), (Start: 8 @30638 has 1 MA's), (9, 30689), (11, 30716), (12, 30728), (13, 30740), (15, 30797),