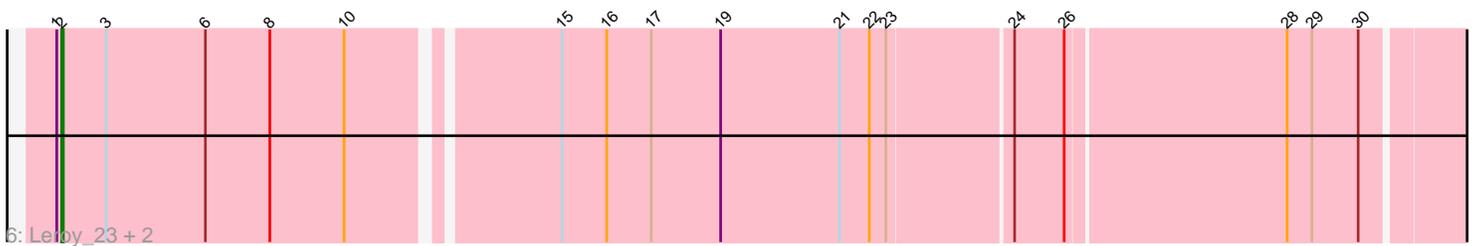
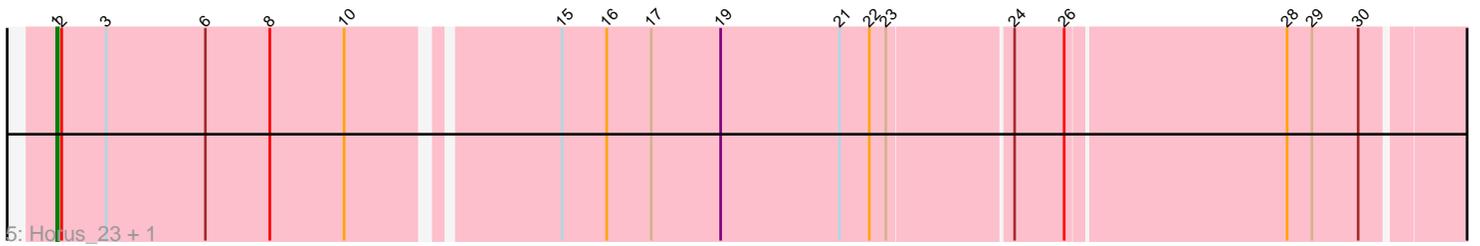
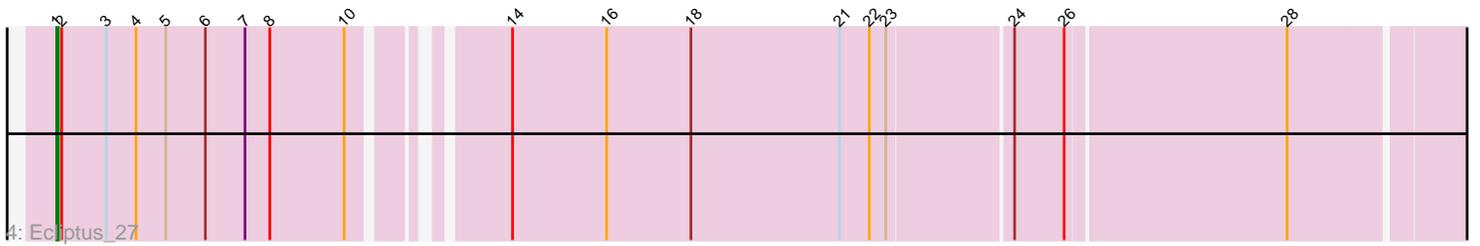
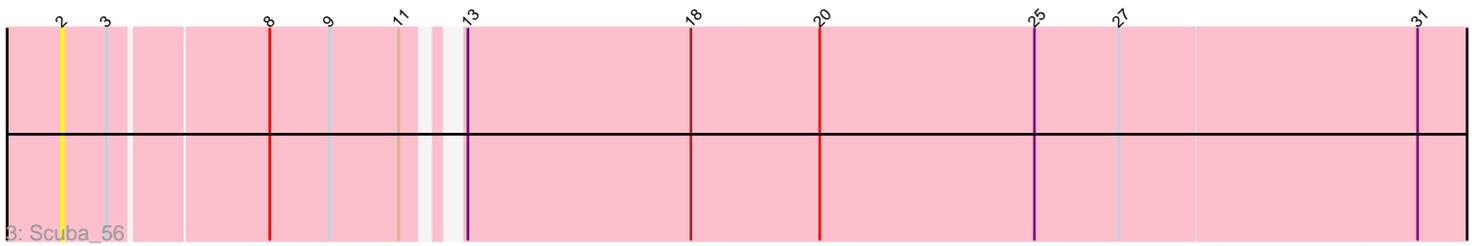
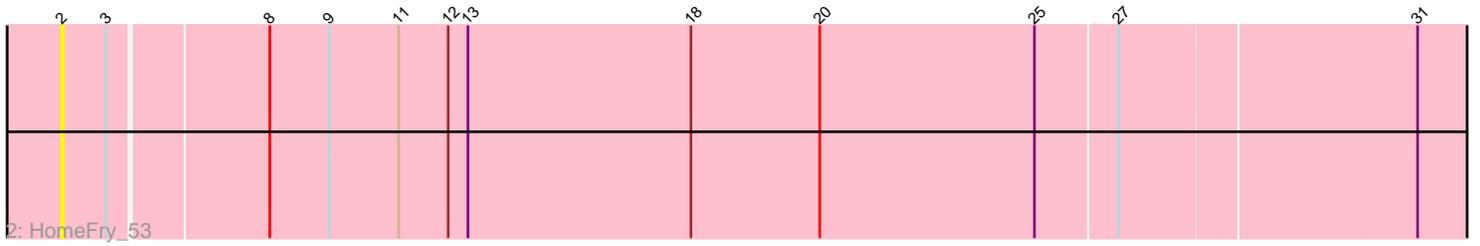
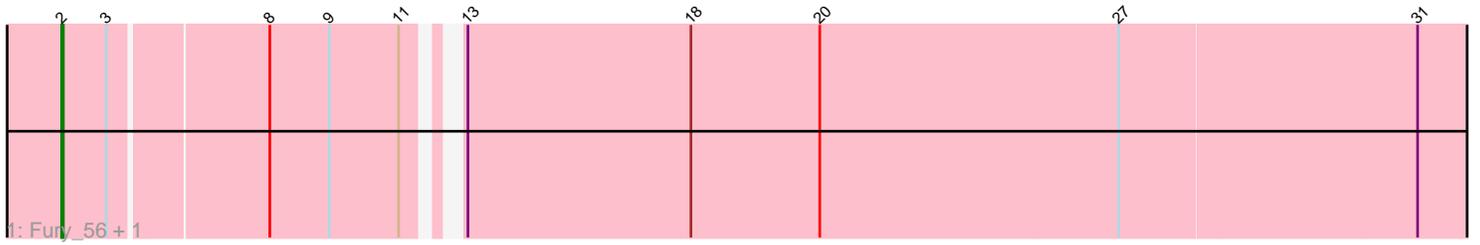


Pham 284418



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 284418 Report

This analysis was run 02/23/26 on database version 636.

Pham number 284418 has 10 members, 2 are drafts.

Phages represented in each track:

- Track 1 : Fury_56, Pleakley_56
- Track 2 : HomeFry_53
- Track 3 : Scuba_56
- Track 4 : Ecliptus_27
- Track 5 : Horus_23, Apricot_22
- Track 6 : Leroy_23, Phistory_24, Crater_21

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 5 of the 8 non-draft genes in the pham.

Genes that call this "Most Annotated" start:

- Crater_21, Fury_56, HomeFry_53, Leroy_23, Phistory_24, Pleakley_56, Scuba_56,

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

- Apricot_22, Ecliptus_27, Horus_23,

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

-

Summary by start number:

Start 1:

- Found in 6 of 10 (60.0%) of genes in pham
- Manual Annotations of this start: 3 of 8
- Called 50.0% of time when present
- Phage (with cluster) where this start called: Apricot_22 (DN3), Ecliptus_27 (DN), Horus_23 (DN1),

Start 2:

- Found in 10 of 10 (100.0%) of genes in pham
- Manual Annotations of this start: 5 of 8
- Called 70.0% of time when present

- Phage (with cluster) where this start called: Crater_21 (DN3), Fury_56 (CR5), HomeFry_53 (CR5), Leroy_23 (DN1), Phistory_24 (DN1), Pleakley_56 (CR5), Scuba_56 (CR5),

Summary by clusters:

There are 4 clusters represented in this pham: DN, DN1, DN3, CR5,

Info for manual annotations of cluster CR5:

- Start number 2 was manually annotated 2 times for cluster CR5.

Info for manual annotations of cluster DN:

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

Info for manual annotations of cluster DN1:

- Start number 1 was manually annotated 1 time for cluster DN1.
- Start number 2 was manually annotated 2 times for cluster DN1.

Info for manual annotations of cluster DN3:

- Start number 1 was manually annotated 1 time for cluster DN3.
- Start number 2 was manually annotated 1 time for cluster DN3.

Gene Information:

Gene: Apricot_22 Start: 18919, Stop: 19752, Start Num: 1

Candidate Starts for Apricot_22:

(Start: 1 @18919 has 3 MA's), (Start: 2 @18922 has 5 MA's), (3, 18949), (6, 19009), (8, 19048), (10, 19093), (15, 19207), (16, 19234), (17, 19261), (19, 19303), (21, 19375), (22, 19393), (23, 19402), (24, 19474), (26, 19504), (28, 19633), (29, 19648), (30, 19675),

Gene: Crater_21 Start: 18922, Stop: 19752, Start Num: 2

Candidate Starts for Crater_21:

(Start: 1 @18919 has 3 MA's), (Start: 2 @18922 has 5 MA's), (3, 18949), (6, 19009), (8, 19048), (10, 19093), (15, 19207), (16, 19234), (17, 19261), (19, 19303), (21, 19375), (22, 19393), (23, 19402), (24, 19474), (26, 19504), (28, 19633), (29, 19648), (30, 19675),

Gene: Ecliptus_27 Start: 21245, Stop: 22066, Start Num: 1

Candidate Starts for Ecliptus_27:

(Start: 1 @21245 has 3 MA's), (Start: 2 @21248 has 5 MA's), (3, 21275), (4, 21293), (5, 21311), (6, 21335), (7, 21359), (8, 21374), (10, 21419), (14, 21491), (16, 21548), (18, 21599), (21, 21689), (22, 21707), (23, 21716), (24, 21788), (26, 21818), (28, 21947),

Gene: Fury_56 Start: 37521, Stop: 38534, Start Num: 2

Candidate Starts for Fury_56:

(Start: 2 @37521 has 5 MA's), (3, 37548), (8, 37638), (9, 37674), (11, 37716), (13, 37734), (18, 37869), (20, 37947), (27, 38127), (31, 38304),

Gene: HomeFry_53 Start: 36153, Stop: 37187, Start Num: 2

Candidate Starts for HomeFry_53:

(Start: 2 @36153 has 5 MA's), (3, 36180), (8, 36270), (9, 36306), (11, 36348), (12, 36378), (13, 36390), (18, 36525), (20, 36603), (25, 36732), (27, 36780), (31, 36954),

Gene: Horus_23 Start: 19383, Stop: 20216, Start Num: 1

Candidate Starts for Horus_23:

(Start: 1 @19383 has 3 MA's), (Start: 2 @19386 has 5 MA's), (3, 19413), (6, 19473), (8, 19512), (10, 19557), (15, 19671), (16, 19698), (17, 19725), (19, 19767), (21, 19839), (22, 19857), (23, 19866), (24, 19938), (26, 19968), (28, 20097), (29, 20112), (30, 20139),

Gene: Leroy_23 Start: 19387, Stop: 20217, Start Num: 2

Candidate Starts for Leroy_23:

(Start: 1 @19384 has 3 MA's), (Start: 2 @19387 has 5 MA's), (3, 19414), (6, 19474), (8, 19513), (10, 19558), (15, 19672), (16, 19699), (17, 19726), (19, 19768), (21, 19840), (22, 19858), (23, 19867), (24, 19939), (26, 19969), (28, 20098), (29, 20113), (30, 20140),

Gene: Phistory_24 Start: 20152, Stop: 20982, Start Num: 2

Candidate Starts for Phistory_24:

(Start: 1 @20149 has 3 MA's), (Start: 2 @20152 has 5 MA's), (3, 20179), (6, 20239), (8, 20278), (10, 20323), (15, 20437), (16, 20464), (17, 20491), (19, 20533), (21, 20605), (22, 20623), (23, 20632), (24, 20704), (26, 20734), (28, 20863), (29, 20878), (30, 20905),

Gene: Pleakley_56 Start: 37522, Stop: 38535, Start Num: 2

Candidate Starts for Pleakley_56:

(Start: 2 @37522 has 5 MA's), (3, 37549), (8, 37639), (9, 37675), (11, 37717), (13, 37735), (18, 37870), (20, 37948), (27, 38128), (31, 38305),

Gene: Scuba_56 Start: 37596, Stop: 38609, Start Num: 2

Candidate Starts for Scuba_56:

(Start: 2 @37596 has 5 MA's), (3, 37623), (8, 37713), (9, 37749), (11, 37791), (13, 37809), (18, 37944), (20, 38022), (25, 38151), (27, 38202), (31, 38379),