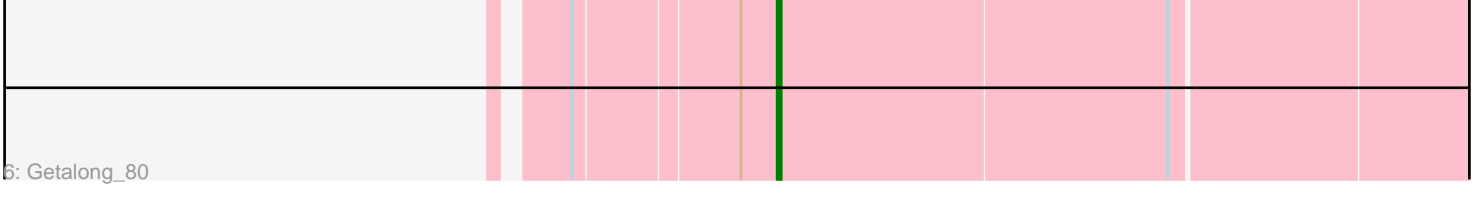
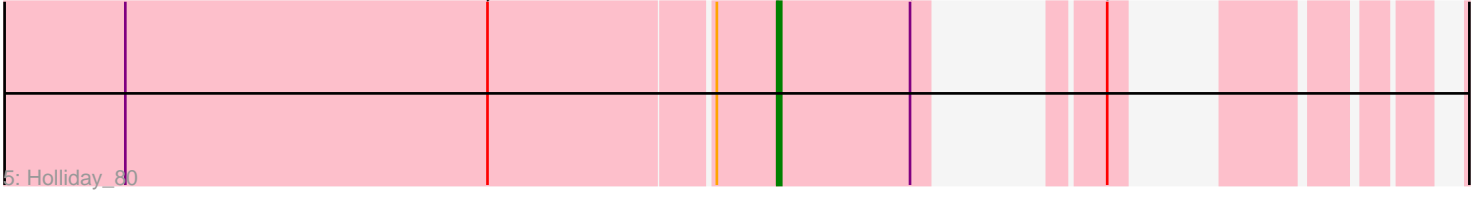
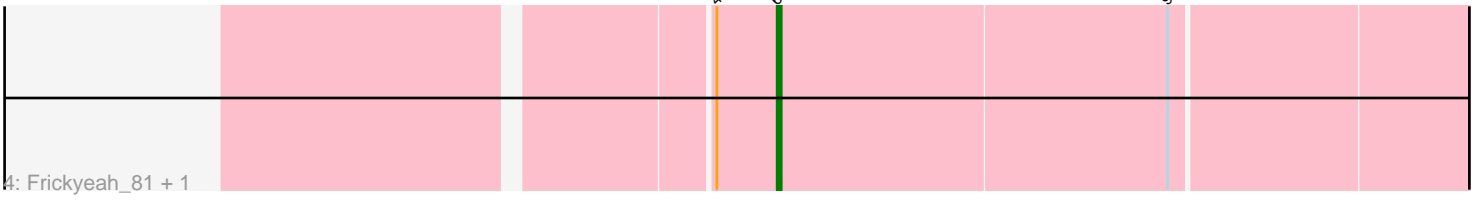
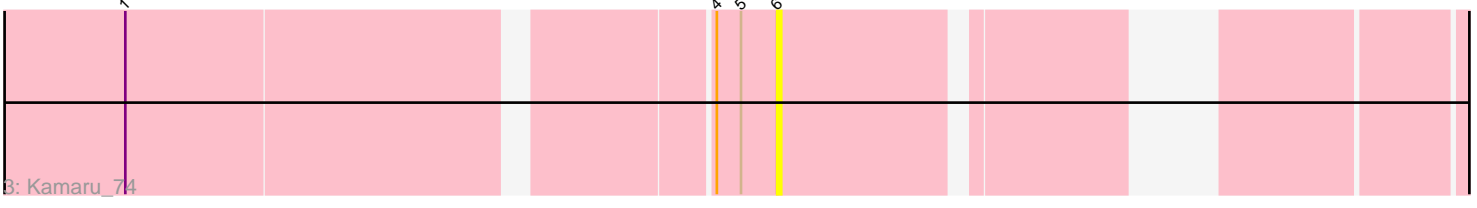
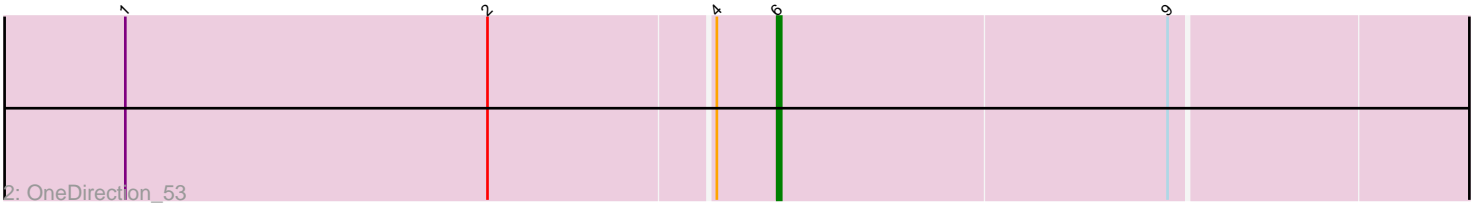
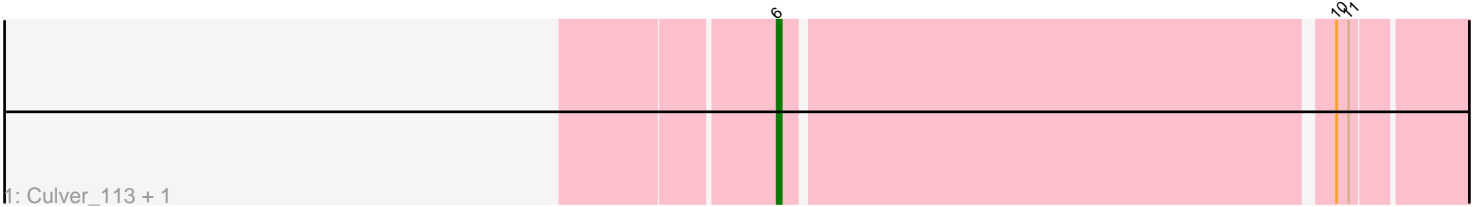


Pham 160959



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 160959 Report

This analysis was run 05/04/24 on database version 560.

Pham number 160959 has 9 members, 3 are drafts.

Phages represented in each track:

- Track 1 : Culver_113, Abscondus_112
- Track 2 : OneDirection_53
- Track 3 : Kamaru_74
- Track 4 : Frickyeh_81, BENtherdunthat_79
- Track 5 : Holliday_80
- Track 6 : Getalong_80
- Track 7 : Leroy_76

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 6 of the 6 non-draft genes in the pham.

Genes that call this "Most Annotated" start:

- Abscondus_112, BENtherdunthat_79, Culver_113, Frickyeh_81, Getalong_80, Holliday_80, Kamaru_74, Leroy_76, OneDirection_53,

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

-

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

-

Summary by start number:

Start 6:

- Found in 9 of 9 (100.0%) of genes in pham
- Manual Annotations of this start: 6 of 6
- Called 100.0% of time when present
- Phage (with cluster) where this start called: Abscondus_112 (CQ), BENtherdunthat_79 (DN1), Culver_113 (CQ1), Frickyeh_81 (DN1), Getalong_80 (DN1), Holliday_80 (DN1), Kamaru_74 (DN1), Leroy_76 (DN1), OneDirection_53 (CZ6),

Summary by clusters:

There are 4 clusters represented in this pham: CZ6, CQ1, DN1, CQ,

Info for manual annotations of cluster CQ1:

- Start number 6 was manually annotated 1 time for cluster CQ1.

Info for manual annotations of cluster CZ6:

- Start number 6 was manually annotated 1 time for cluster CZ6.

Info for manual annotations of cluster DN1:

- Start number 6 was manually annotated 4 times for cluster DN1.

Gene Information:

Gene: Abscondus_112 Start: 65388, Stop: 65648, Start Num: 6

Candidate Starts for Abscondus_112:

(Start: 6 @65388 has 6 MA's), (10, 65520), (11, 65523),

Gene: BENtherdunthat_79 Start: 44974, Stop: 45240, Start Num: 6

Candidate Starts for BENtherdunthat_79:

(4, 44959), (Start: 6 @44974 has 6 MA's), (9, 45070),

Gene: Culver_113 Start: 64860, Stop: 65120, Start Num: 6

Candidate Starts for Culver_113:

(Start: 6 @64860 has 6 MA's), (10, 64992), (11, 64995),

Gene: Frickyeah_81 Start: 45105, Stop: 45371, Start Num: 6

Candidate Starts for Frickyeah_81:

(4, 45090), (Start: 6 @45105 has 6 MA's), (9, 45201),

Gene: Getalong_80 Start: 46400, Stop: 46666, Start Num: 6

Candidate Starts for Getalong_80:

(3, 46352), (5, 46391), (Start: 6 @46400 has 6 MA's), (9, 46496),

Gene: Holliday_80 Start: 48011, Stop: 48211, Start Num: 6

Candidate Starts for Holliday_80:

(1, 47852), (2, 47942), (4, 47996), (Start: 6 @48011 has 6 MA's), (7, 48044), (8, 48062),

Gene: Kamaru_74 Start: 43594, Stop: 43830, Start Num: 6

Candidate Starts for Kamaru_74:

(1, 43444), (4, 43579), (5, 43585), (Start: 6 @43594 has 6 MA's),

Gene: Leroy_76 Start: 44184, Stop: 44435, Start Num: 6

Candidate Starts for Leroy_76:

(Start: 6 @44184 has 6 MA's),

Gene: OneDirection_53 Start: 33352, Stop: 33618, Start Num: 6

Candidate Starts for OneDirection_53:

(1, 33193), (2, 33283), (4, 33337), (Start: 6 @33352 has 6 MA's), (9, 33448),

