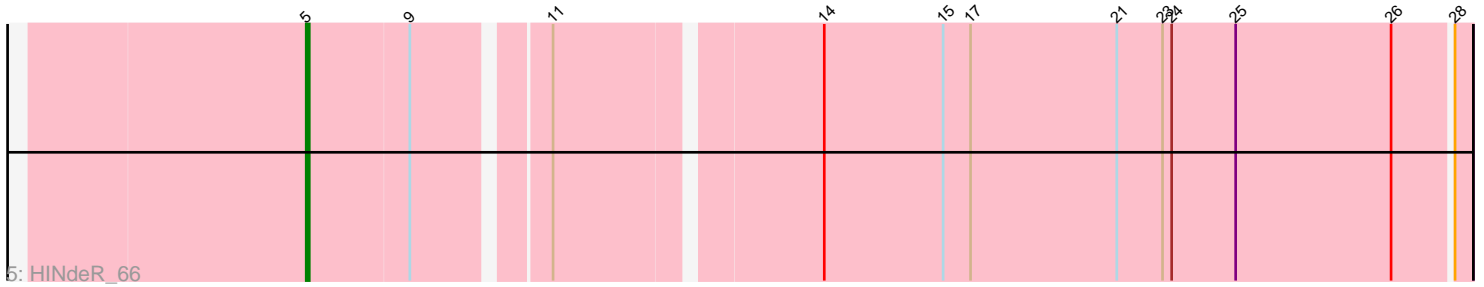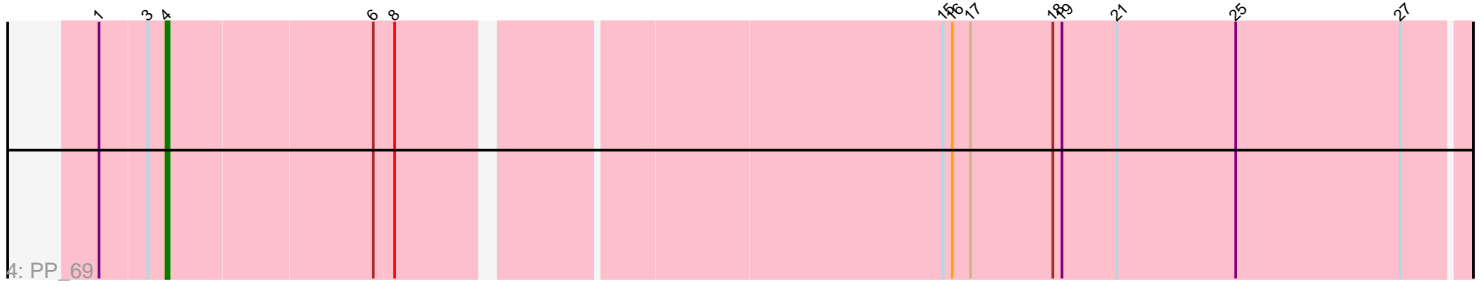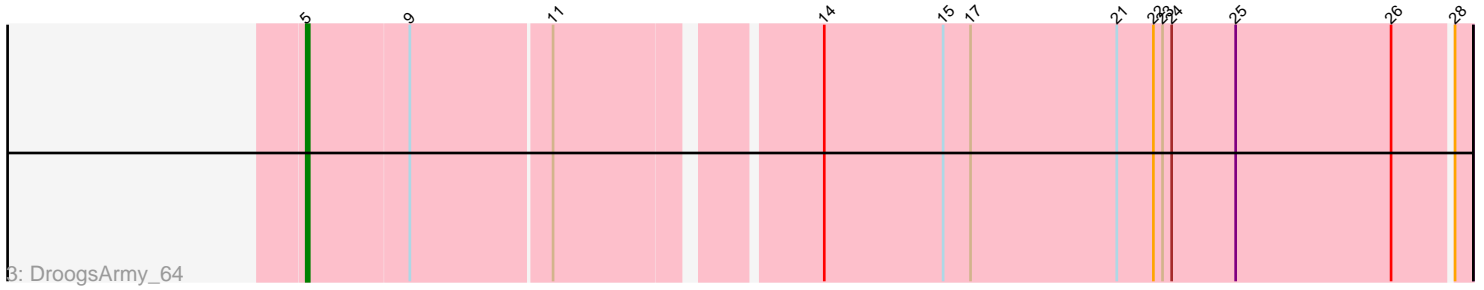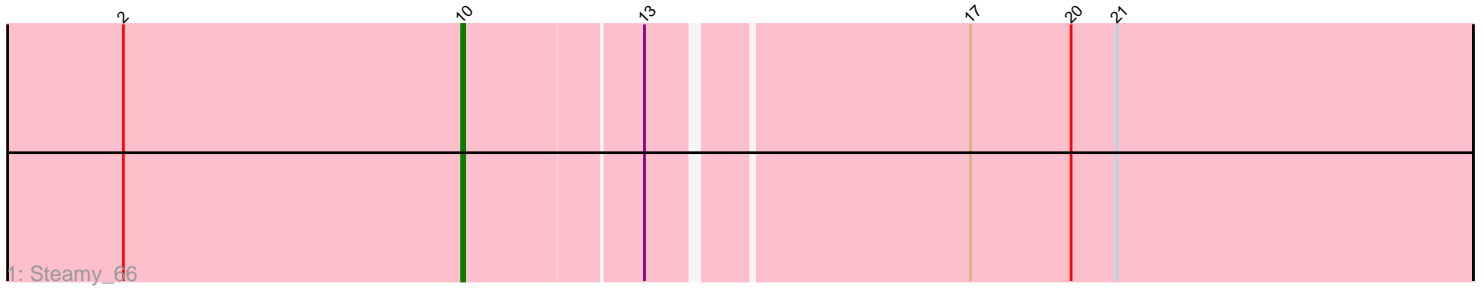# Pham 172125

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 172125 Report

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

Pham number 172125 has 5 members, 0 are drafts.

Phages represented in each track:
• Track 1 : Steamy_66
• Track 2 : Kimona_64
• Track 3 : DroogsArmy_64
• Track 4 : PP_69
• Track 5 : HINdeR_66

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

The start number called the most often in the published annotations is 5, it was called in 2 of the 5 non-draft genes in the pham.

Genes that call this "Most Annotated" start:
• DroogsArmy_64, HINdeR_66,

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

Genes that do not have the "Most Annotated" start:
• Kimona_64, PP_69, Steamy_66,

**Summary by start number:**

Start 4:
• Found in 1 of 5 ( 20.0% ) of genes in pham
• Manual Annotations of this start: 1 of 5
• Called 100.0% of time when present
• Phage (with cluster) where this start called: PP_69 (A7),

Start 5:
• Found in 2 of 5 ( 40.0% ) of genes in pham
• Manual Annotations of this start: 2 of 5
• Called 100.0% of time when present
• Phage (with cluster) where this start called: DroogsArmy_64 (A7), HINdeR_66 (A7),

Start 7:
• Found in 1 of 5 ( 20.0% ) of genes in pham
• Manual Annotations of this start: 1 of 5
• Called 100.0% of time when present
• Phage (with cluster) where this start called: Kimona_64 (A19),

Start 10:
• Found in 1 of 5 ( 20.0% ) of genes in pham
• Manual Annotations of this start: 1 of 5
• Called 100.0% of time when present
• Phage (with cluster) where this start called: Steamy_66 (A12),

**Summary by clusters:**

There are 3 clusters represented in this pham: A19, A7, A12,

Info for manual annotations of cluster A12:
•Start number 10 was manually annotated 1 time for cluster A12.

Info for manual annotations of cluster A19:
•Start number 7 was manually annotated 1 time for cluster A19.

Info for manual annotations of cluster A7:
•Start number 4 was manually annotated 1 time for cluster A7.
•Start number 5 was manually annotated 2 times for cluster A7.

*Gene Information:*

Gene: DroogsArmy_64 Start: 45309, Stop: 44944, Start Num: 5
Candidate Starts for DroogsArmy_64:
(Start: 5 @45309 has 2 MA's), (9, 45276), (11, 45231), (14, 45153), (15, 45114), (17, 45105), (21, 45057), (22, 45045), (23, 45042), (24, 45039), (25, 45018), (26, 44967), (28, 44949),

Gene: HINdeR_66 Start: 44898, Stop: 44536, Start Num: 5
Candidate Starts for HINdeR_66:
(Start: 5 @44898 has 2 MA's), (9, 44865), (11, 44826), (14, 44745), (15, 44706), (17, 44697), (21, 44649), (23, 44634), (24, 44631), (25, 44610), (26, 44559), (28, 44541),

Gene: Kimona_64 Start: 41048, Stop: 40713, Start Num: 7
Candidate Starts for Kimona_64:
(Start: 7 @41048 has 1 MA's), (12, 40988), (15, 40883), (21, 40826), (24, 40808),

Gene: PP_69 Start: 46873, Stop: 46463, Start Num: 4
Candidate Starts for PP_69:
(1, 46894), (3, 46879), (Start: 4 @46873 has 1 MA's), (6, 46807), (8, 46801), (15, 46633), (16, 46630), (17, 46624), (18, 46597), (19, 46594), (21, 46576), (25, 46537), (27, 46483),

Gene: Steamy_66 Start: 42070, Stop: 41750, Start Num: 10
Candidate Starts for Steamy_66:
(2, 42181), (Start: 10 @42070 has 1 MA's), (13, 42013), (17, 41914), (20, 41881), (21, 41866),