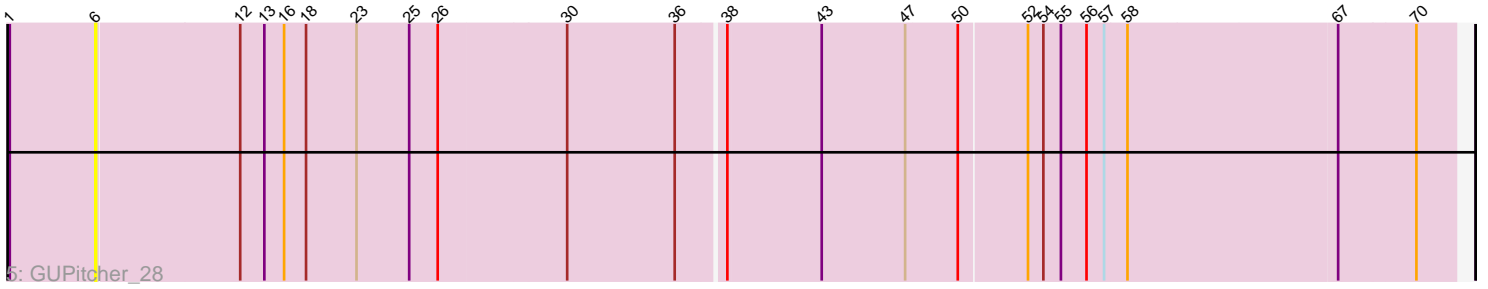
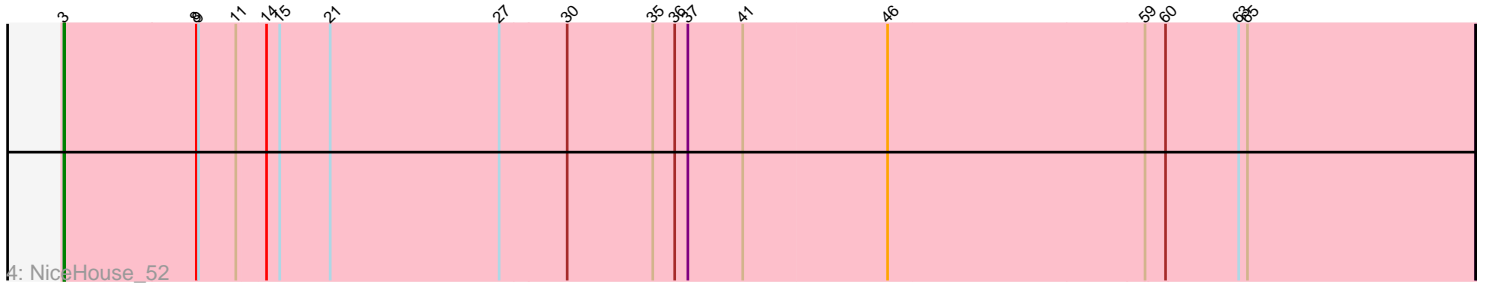
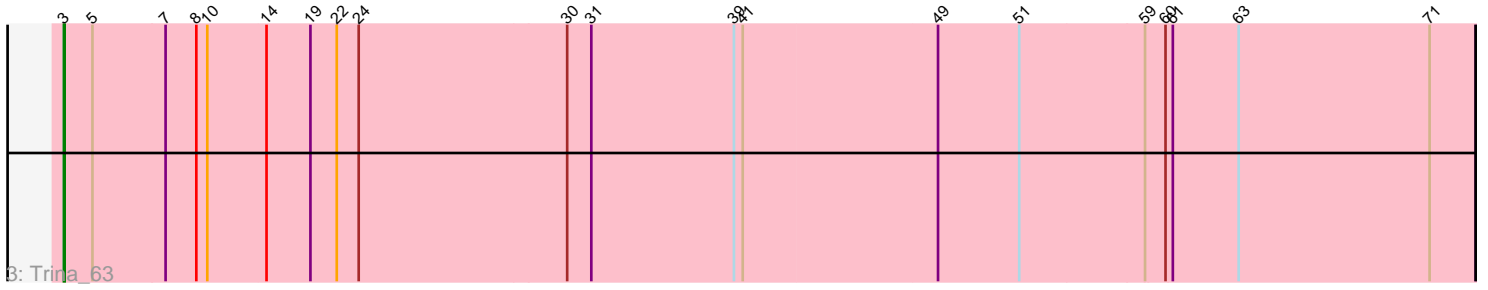
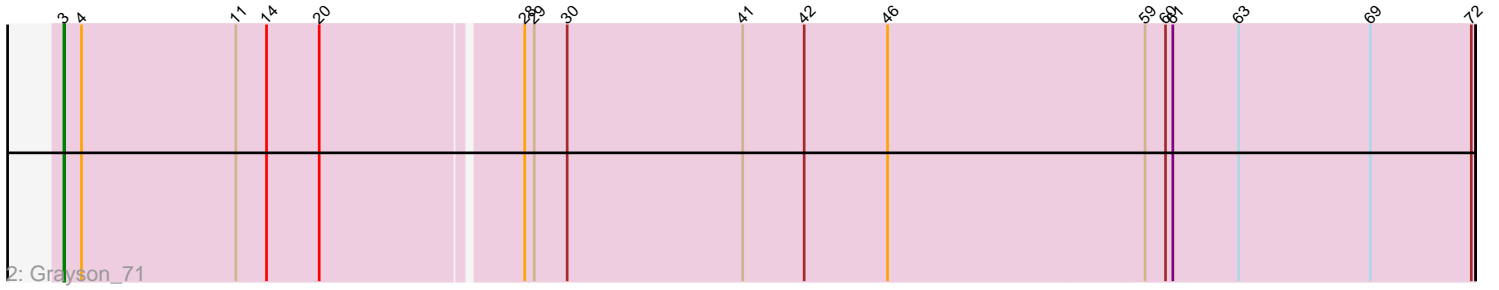
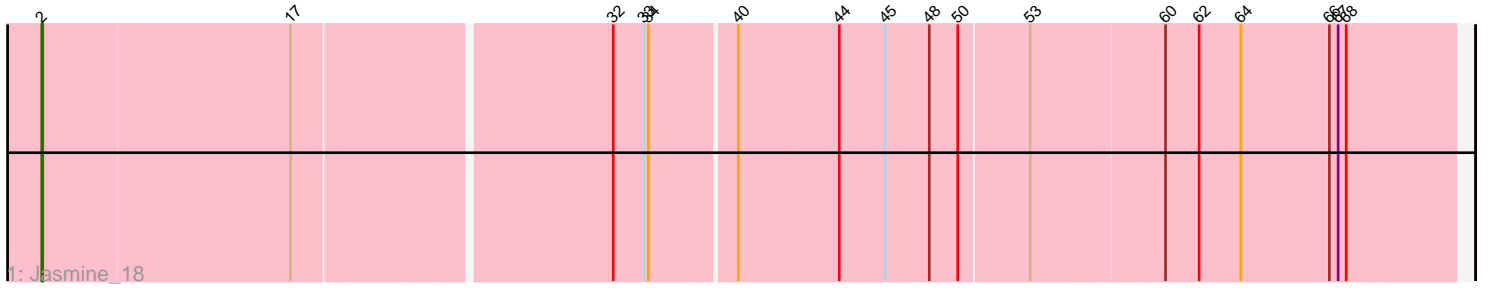


Pham 203756



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 203756 Report

This analysis was run 01/18/25 on database version 583.

Pham number 203756 has 5 members, 1 are drafts.

Phages represented in each track:

- Track 1 : Jasmine_18
- Track 2 : Grayson_71
- Track 3 : Trina_63
- Track 4 : NiceHouse_52
- Track 5 : GUPitcher_28

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

The start number called the most often in the published annotations is 3, it was called in 3 of the 4 non-draft genes in the pham.

Genes that call this "Most Annotated" start:

- Grayson_71, NiceHouse_52, Trina_63,

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

-

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

- GUPitcher_28, Jasmine_18,

Summary by start number:

Start 2:

- Found in 1 of 5 (20.0%) of genes in pham
- Manual Annotations of this start: 1 of 4
- Called 100.0% of time when present
- Phage (with cluster) where this start called: Jasmine_18 (AV),

Start 3:

- Found in 3 of 5 (60.0%) of genes in pham
- Manual Annotations of this start: 3 of 4
- Called 100.0% of time when present
- Phage (with cluster) where this start called: Grayson_71 (CB), NiceHouse_52 (CE), Trina_63 (CE),

Start 6:

- Found in 1 of 5 (20.0%) of genes in pham
- No Manual Annotations of this start.
- Called 100.0% of time when present
- Phage (with cluster) where this start called: GUPitcher_28 (FJ),

Summary by clusters:

There are 4 clusters represented in this pham: CB, FJ, CE, AV,

Info for manual annotations of cluster AV:

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

Info for manual annotations of cluster CB:

- Start number 3 was manually annotated 1 time for cluster CB.

Info for manual annotations of cluster CE:

- Start number 3 was manually annotated 2 times for cluster CE.

Gene Information:

Gene: GUPitcher_28 Start: 22041, Stop: 23840, Start Num: 6

Candidate Starts for GUPitcher_28:

(1, 21924), (6, 22041), (12, 22233), (13, 22266), (16, 22293), (18, 22323), (23, 22392), (25, 22458), (26, 22497), (30, 22668), (36, 22815), (38, 22875), (43, 23001), (47, 23115), (50, 23184), (52, 23274), (54, 23295), (55, 23319), (56, 23352), (57, 23376), (58, 23406), (67, 23685), (70, 23787),

Gene: Grayson_71 Start: 30665, Stop: 32554, Start Num: 3

Candidate Starts for Grayson_71:

(Start: 3 @30665 has 3 MA's), (4, 30689), (11, 30899), (14, 30938), (20, 31010), (28, 31268), (29, 31280), (30, 31325), (41, 31565), (42, 31646), (46, 31760), (59, 32105), (60, 32132), (61, 32141), (63, 32231), (69, 32411), (72, 32549),

Gene: Jasmine_18 Start: 11823, Stop: 13667, Start Num: 2

Candidate Starts for Jasmine_18:

(Start: 2 @11823 has 1 MA's), (17, 12159), (32, 12564), (33, 12606), (34, 12612), (40, 12720), (44, 12855), (45, 12918), (48, 12978), (50, 13017), (53, 13110), (60, 13284), (62, 13329), (64, 13383), (66, 13500), (67, 13512), (68, 13521),

Gene: NiceHouse_52 Start: 24818, Stop: 26725, Start Num: 3

Candidate Starts for NiceHouse_52:

(Start: 3 @24818 has 3 MA's), (8, 24998), (9, 25001), (11, 25052), (14, 25088), (15, 25106), (21, 25175), (27, 25406), (30, 25496), (35, 25613), (36, 25643), (37, 25661), (41, 25736), (46, 25931), (59, 26276), (60, 26303), (63, 26402), (65, 26414),

Gene: Trina_63 Start: 28559, Stop: 30469, Start Num: 3

Candidate Starts for Trina_63:

(Start: 3 @28559 has 3 MA's), (5, 28598), (7, 28697), (8, 28739), (10, 28754), (14, 28832), (19, 28892), (22, 28928), (24, 28958), (30, 29240), (31, 29273), (39, 29468), (41, 29480), (49, 29741), (51, 29852),

(59, 30020), (60, 30047), (61, 30056), (63, 30146), (71, 30407),