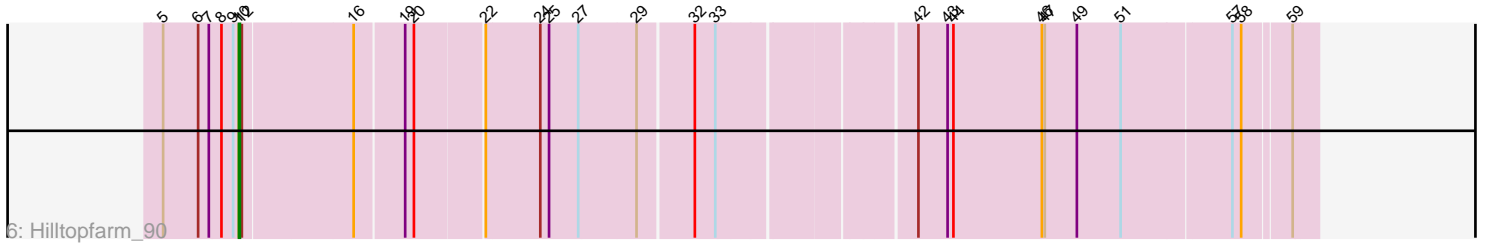
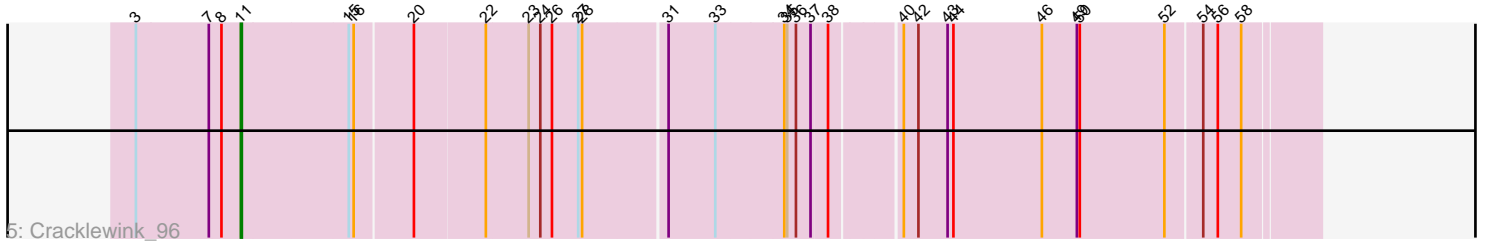
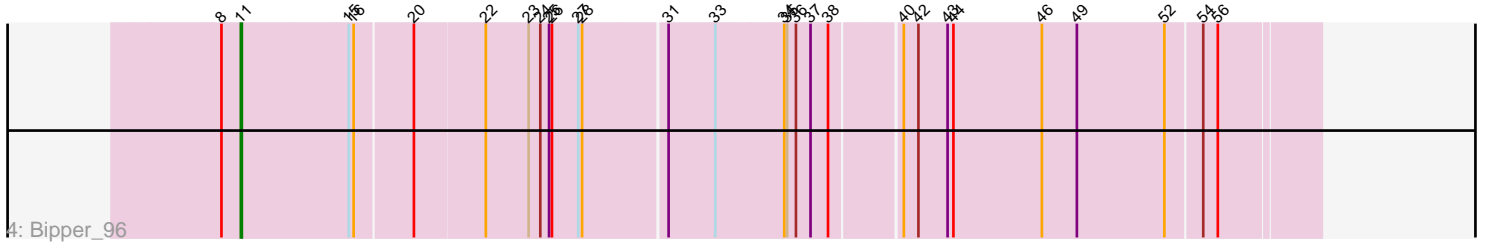
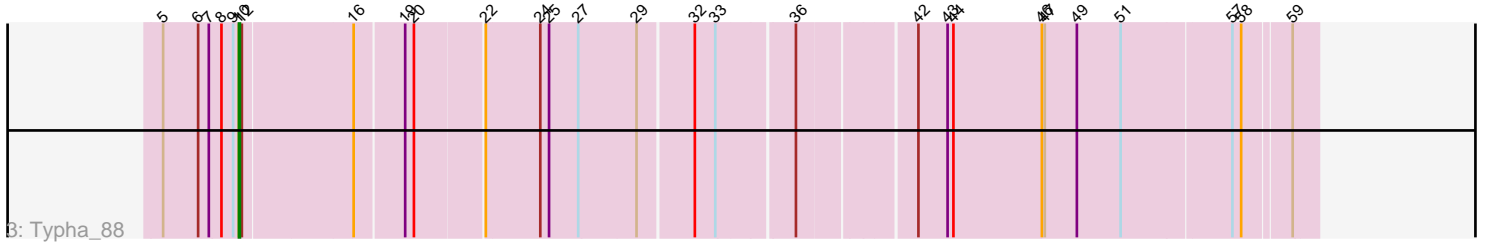
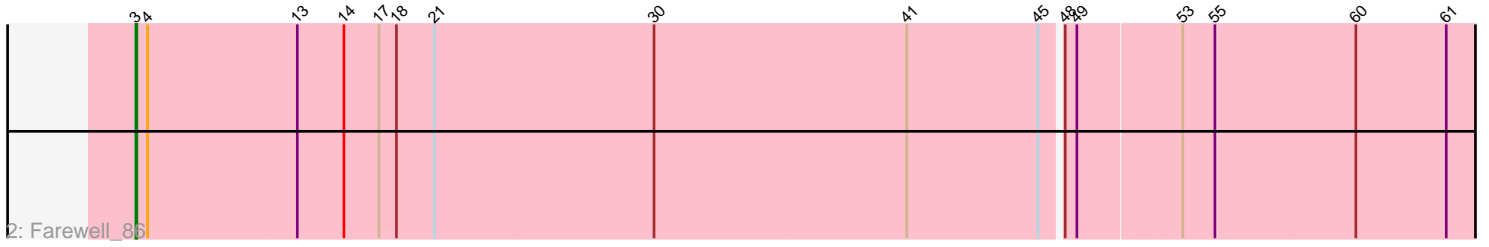
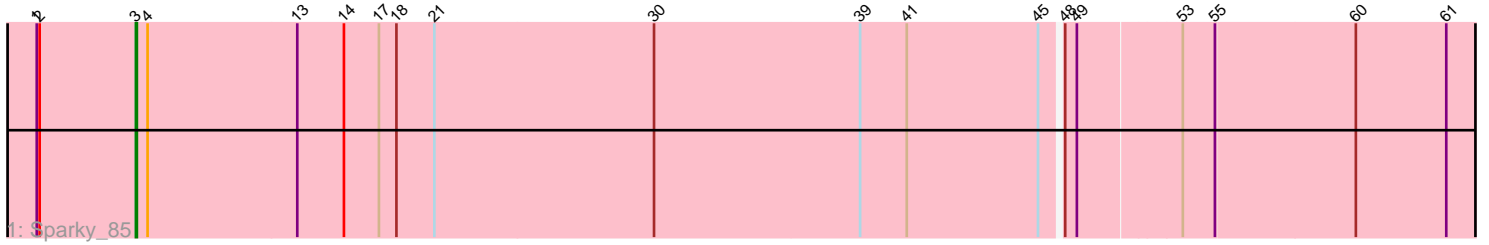


Pham 167332



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 167332 Report

This analysis was run 07/09/24 on database version 566.

Pham number 167332 has 6 members, 0 are drafts.

Phages represented in each track:

- Track 1 : Sparky_85
- Track 2 : Farewell_86
- Track 3 : Typha_88
- Track 4 : Bipper_96
- Track 5 : Cracklewink_96
- Track 6 : Hilltopfarm_90

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

The start number called the most often in the published annotations is 11, it was called in 2 of the 6 non-draft genes in the pham.

Genes that call this "Most Annotated" start:

- Bipper_96, Cracklewink_96,

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

-

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

- Farewell_86, Hilltopfarm_90, Sparky_85, Typha_88,

Summary by start number:

Start 3:

- Found in 3 of 6 (50.0%) of genes in pham
- Manual Annotations of this start: 2 of 6
- Called 66.7% of time when present
- Phage (with cluster) where this start called: Farewell_86 (AF), Sparky_85 (AF),

Start 10:

- Found in 2 of 6 (33.3%) of genes in pham
- Manual Annotations of this start: 2 of 6
- Called 100.0% of time when present
- Phage (with cluster) where this start called: Hilltopfarm_90 (Y), Typha_88 (Y),

Start 11:

- Found in 2 of 6 (33.3%) of genes in pham
- Manual Annotations of this start: 2 of 6
- Called 100.0% of time when present
- Phage (with cluster) where this start called: Bipper_96 (Y), Cracklewink_96 (Y),

Summary by clusters:

There are 2 clusters represented in this pham: Y, AF,

Info for manual annotations of cluster AF:

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

Info for manual annotations of cluster Y:

- Start number 10 was manually annotated 2 times for cluster Y.
- Start number 11 was manually annotated 2 times for cluster Y.

Gene Information:

Gene: Bipper_96 Start: 60606, Stop: 61649, Start Num: 11

Candidate Starts for Bipper_96:

(8, 60588), (Start: 11 @60606 has 2 MA's), (15, 60711), (16, 60714), (20, 60771), (22, 60840), (23, 60882), (24, 60894), (25, 60903), (26, 60906), (27, 60933), (28, 60936), (31, 61017), (33, 61065), (34, 61131), (35, 61134), (36, 61143), (37, 61158), (38, 61173), (40, 61236), (42, 61251), (43, 61281), (44, 61287), (46, 61377), (49, 61413), (52, 61503), (54, 61536), (56, 61551),

Gene: Cracklewink_96 Start: 59841, Stop: 60884, Start Num: 11

Candidate Starts for Cracklewink_96:

(Start: 3 @59739 has 2 MA's), (7, 59811), (8, 59823), (Start: 11 @59841 has 2 MA's), (15, 59946), (16, 59949), (20, 60006), (22, 60075), (23, 60117), (24, 60129), (26, 60141), (27, 60168), (28, 60171), (31, 60252), (33, 60300), (34, 60366), (35, 60369), (36, 60378), (37, 60393), (38, 60408), (40, 60471), (42, 60486), (43, 60516), (44, 60522), (46, 60612), (49, 60648), (50, 60651), (52, 60738), (54, 60771), (56, 60786), (58, 60810),

Gene: Farewell_86 Start: 55341, Stop: 56711, Start Num: 3

Candidate Starts for Farewell_86:

(Start: 3 @55341 has 2 MA's), (4, 55353), (13, 55506), (14, 55554), (17, 55590), (18, 55608), (21, 55647), (30, 55872), (41, 56130), (45, 56265), (48, 56283), (49, 56295), (53, 56397), (55, 56430), (60, 56571), (61, 56664),

Gene: Hilltopfarm_90 Start: 58232, Stop: 59260, Start Num: 10

Candidate Starts for Hilltopfarm_90:

(5, 58157), (6, 58193), (7, 58202), (8, 58214), (9, 58226), (Start: 10 @58232 has 2 MA's), (12, 58235), (16, 58340), (19, 58388), (20, 58397), (22, 58463), (24, 58517), (25, 58526), (27, 58556), (29, 58613), (32, 58667), (33, 58688), (42, 58868), (43, 58898), (44, 58904), (46, 58994), (47, 58997), (49, 59030), (51, 59075), (57, 59183), (58, 59192), (59, 59237),

Gene: Sparky_85 Start: 56935, Stop: 58305, Start Num: 3

Candidate Starts for Sparky_85:

(1, 56833), (2, 56836), (Start: 3 @56935 has 2 MA's), (4, 56947), (13, 57100), (14, 57148), (17, 57184), (18, 57202), (21, 57241), (30, 57466), (39, 57676), (41, 57724), (45, 57859), (48, 57877), (49, 57889), (53, 57991), (55, 58024), (60, 58165), (61, 58258),

Gene: Typha_88 Start: 57847, Stop: 58875, Start Num: 10

Candidate Starts for Typha_88:

(5, 57772), (6, 57808), (7, 57817), (8, 57829), (9, 57841), (Start: 10 @57847 has 2 MA's), (12, 57850), (16, 57955), (19, 58003), (20, 58012), (22, 58078), (24, 58132), (25, 58141), (27, 58171), (29, 58228), (32, 58282), (33, 58303), (36, 58375), (42, 58483), (43, 58513), (44, 58519), (46, 58609), (47, 58612), (49, 58645), (51, 58690), (57, 58798), (58, 58807), (59, 58852),