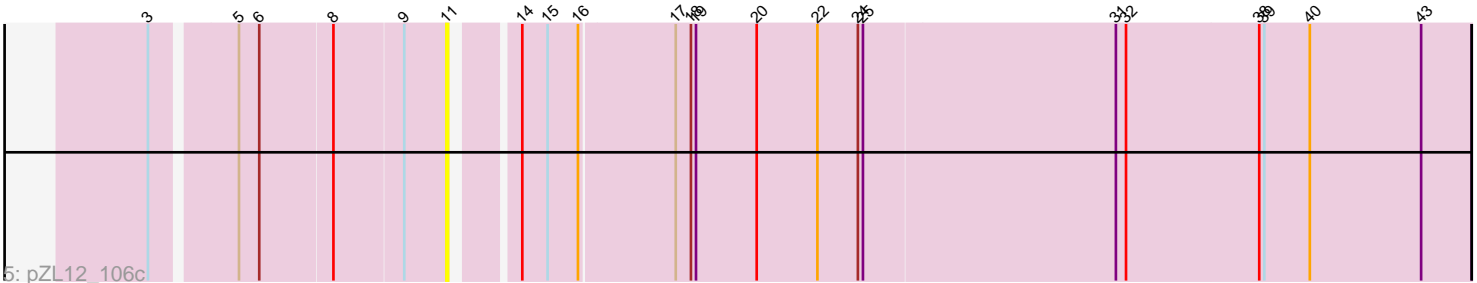
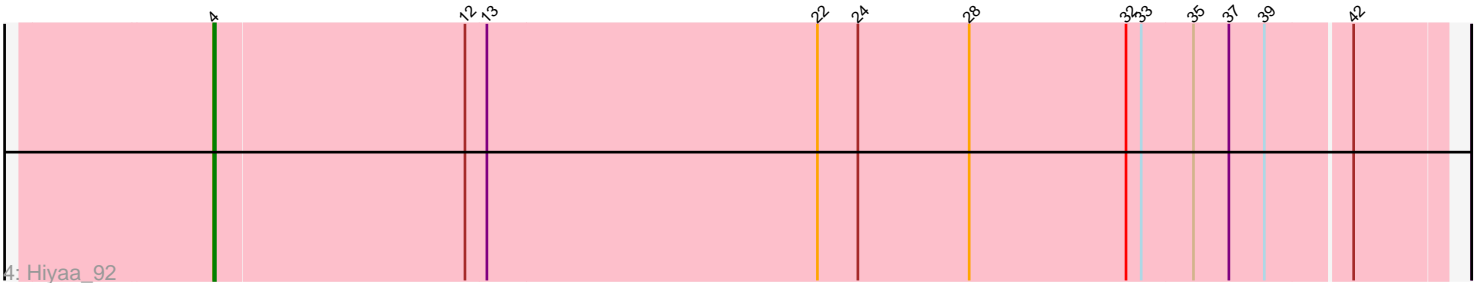
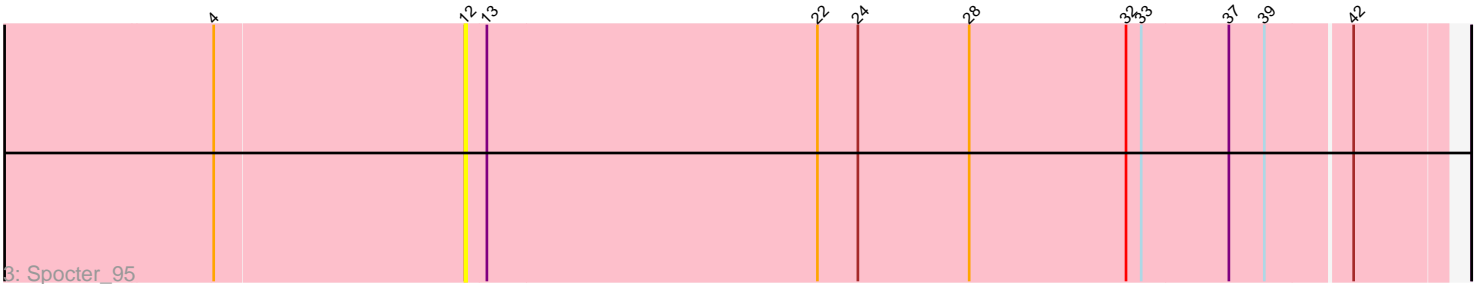
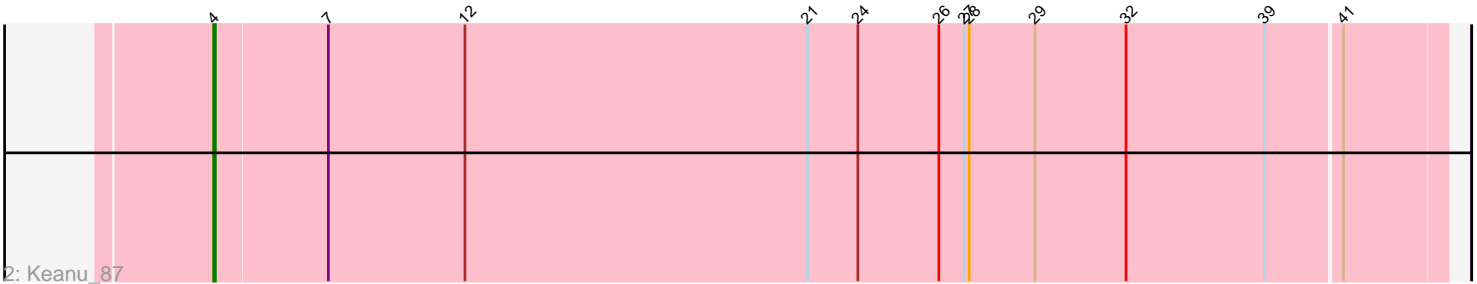
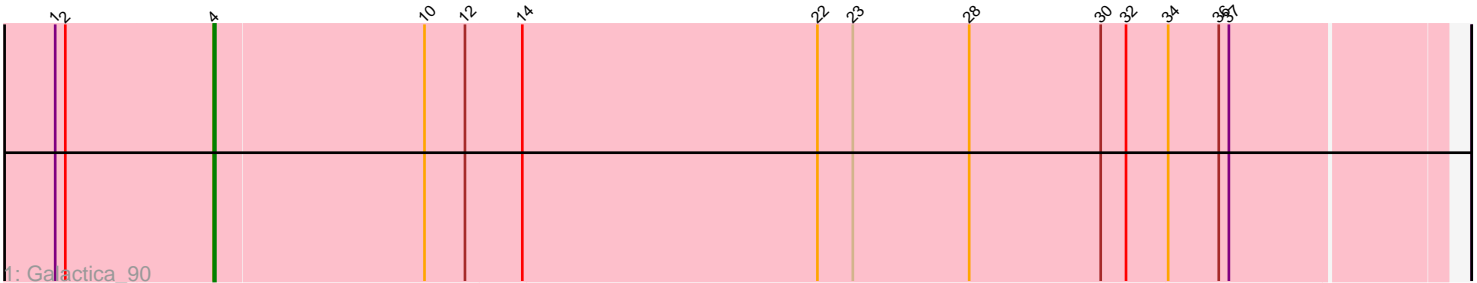


Pham 146330



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 146330 Report

This analysis was run 04/28/24 on database version 559.

Pham number 146330 has 5 members, 2 are drafts.

Phages represented in each track:

- Track 1 : Galactica_90
- Track 2 : Keanu_87
- Track 3 : Spocter_95
- Track 4 : Hiyaa_92
- Track 5 : pZL12_106c

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

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

Genes that call this "Most Annotated" start:

- Galactica_90, Hiyaa_92, Keanu_87,

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

- Spocter_95,

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

- pZL12_106c,

Summary by start number:

Start 4:

- Found in 4 of 5 (80.0%) of genes in pham
- Manual Annotations of this start: 3 of 3
- Called 75.0% of time when present
- Phage (with cluster) where this start called: Galactica_90 (BQ), Hiyaa_92 (BQ), Keanu_87 (BQ),

Start 11:

- 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: pZL12_106c (singleton),

Start 12:

- Found in 4 of 5 (80.0%) of genes in pham
- No Manual Annotations of this start.
- Called 25.0% of time when present
- Phage (with cluster) where this start called: Spocter_95 (BQ),

Summary by clusters:

There are 2 clusters represented in this pham: singleton, BQ,

Info for manual annotations of cluster BQ:

- Start number 4 was manually annotated 3 times for cluster BQ.

Gene Information:

Gene: Galactica_90 Start: 62183, Stop: 62902, Start Num: 4

Candidate Starts for Galactica_90:

(1, 62090), (2, 62096), (Start: 4 @62183 has 3 MA's), (10, 62306), (12, 62330), (14, 62363), (22, 62537), (23, 62558), (28, 62627), (30, 62705), (32, 62720), (34, 62744), (36, 62774), (37, 62780),

Gene: Hiyaa_92 Start: 62989, Stop: 63708, Start Num: 4

Candidate Starts for Hiyaa_92:

(Start: 4 @62989 has 3 MA's), (12, 63136), (13, 63148), (22, 63343), (24, 63367), (28, 63433), (32, 63526), (33, 63535), (35, 63565), (37, 63586), (39, 63607), (42, 63655),

Gene: Keanu_87 Start: 62136, Stop: 62855, Start Num: 4

Candidate Starts for Keanu_87:

(Start: 4 @62136 has 3 MA's), (7, 62202), (12, 62283), (21, 62484), (24, 62514), (26, 62562), (27, 62577), (28, 62580), (29, 62619), (32, 62673), (39, 62754), (41, 62796),

Gene: Spocter_95 Start: 63497, Stop: 64069, Start Num: 12

Candidate Starts for Spocter_95:

(Start: 4 @63350 has 3 MA's), (12, 63497), (13, 63509), (22, 63704), (24, 63728), (28, 63794), (32, 63887), (33, 63896), (37, 63947), (39, 63968), (42, 64016),

Gene: pZL12_106c Start: 86095, Stop: 85511, Start Num: 11

Candidate Starts for pZL12_106c:

(3, 86257), (5, 86212), (6, 86200), (8, 86158), (9, 86119), (11, 86095), (14, 86065), (15, 86050), (16, 86032), (17, 85978), (18, 85969), (19, 85966), (20, 85930), (22, 85894), (24, 85870), (25, 85867), (31, 85720), (32, 85714), (38, 85636), (39, 85633), (40, 85606), (43, 85540),