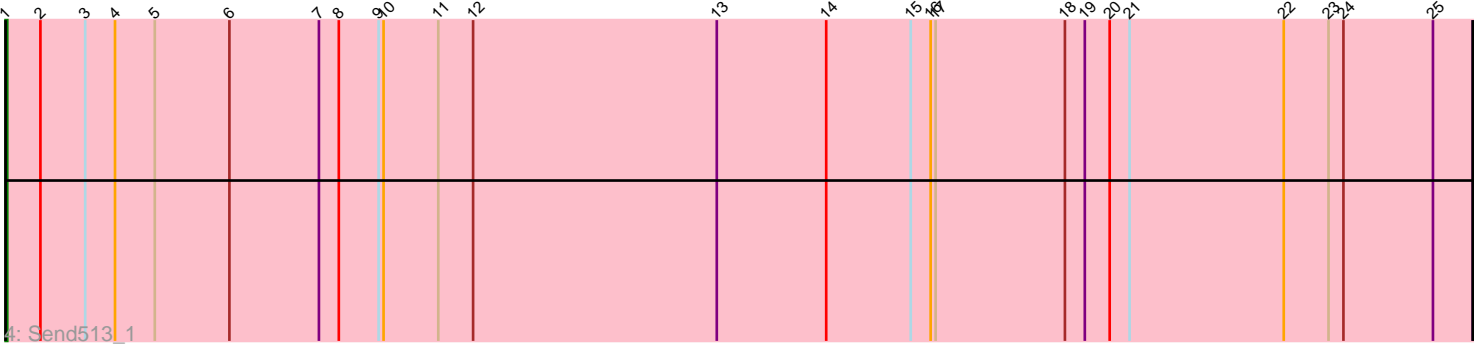
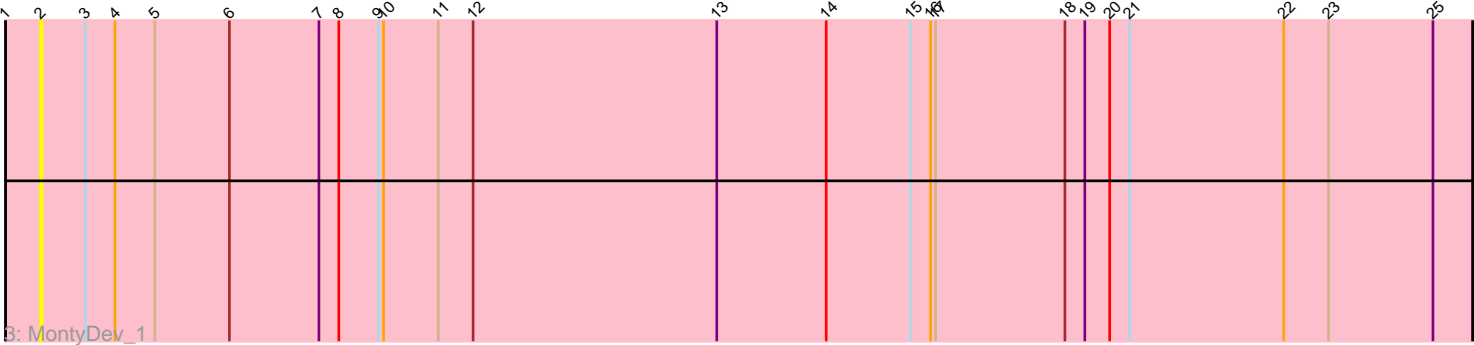
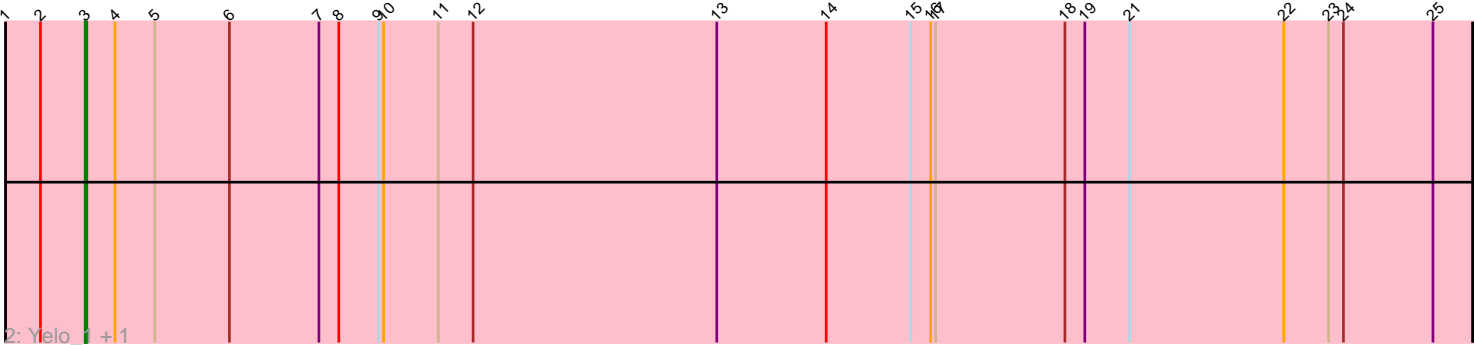
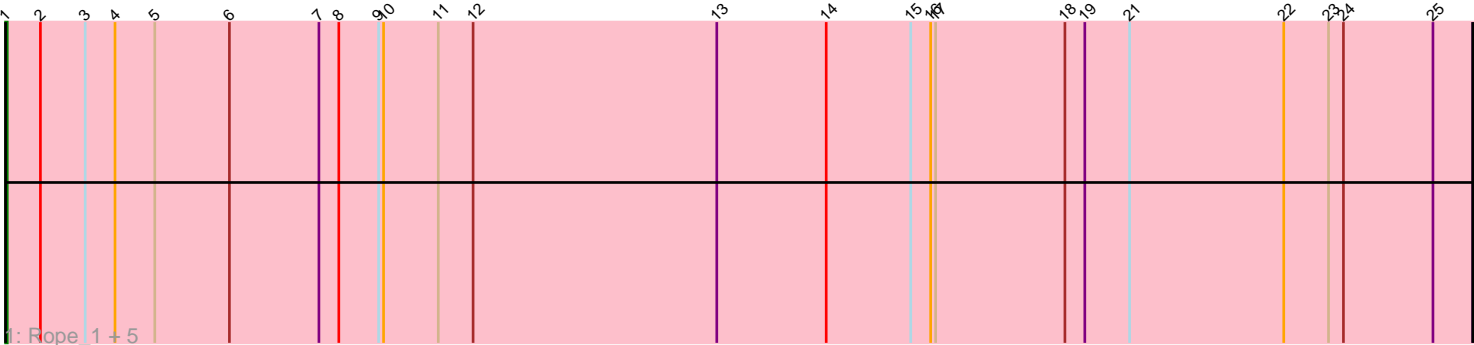


Pham 87670



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 87670 Report

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

Pham number 87670 has 10 members, 2 are drafts.

Phages represented in each track:

- Track 1 : Rope_1, Candle_1, Riparian_1, Nilo_1, Papyrus_1, Weiss13_1
- Track 2 : Yelo_1, Zenon_1
- Track 3 : MontyDev_1
- Track 4 : Send513_1

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

The start number called the most often in the published annotations is 1, it was called in 7 of the 8 non-draft genes in the pham.

Genes that call this "Most Annotated" start:

- Candle_1, Nilo_1, Papyrus_1, Riparian_1, Rope_1, Send513_1, Weiss13_1,

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

- MontyDev_1, Yelo_1, Zenon_1,

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

-

Summary by start number:

Start 1:

- Found in 10 of 10 (100.0%) of genes in pham
- Manual Annotations of this start: 7 of 8
- Called 70.0% of time when present
- Phage (with cluster) where this start called: Candle_1 (R), Nilo_1 (R), Papyrus_1 (R), Riparian_1 (R), Rope_1 (R), Send513_1 (R), Weiss13_1 (R),

Start 2:

- Found in 10 of 10 (100.0%) of genes in pham
- No Manual Annotations of this start.
- Called 10.0% of time when present
- Phage (with cluster) where this start called: MontyDev_1 (R),

Start 3:

- Found in 10 of 10 (100.0%) of genes in pham
- Manual Annotations of this start: 1 of 8
- Called 20.0% of time when present
- Phage (with cluster) where this start called: Yelo_1 (R), Zenon_1 (R),

Summary by clusters:

There is one cluster represented in this pham: R

Info for manual annotations of cluster R:

- Start number 1 was manually annotated 7 times for cluster R.
- Start number 3 was manually annotated 1 time for cluster R.

Gene Information:

Gene: Candle_1 Start: 1, Stop: 885, Start Num: 1

Candidate Starts for Candle_1:

(Start: 1 @1 has 7 MA's), (2, 22), (Start: 3 @49 has 1 MA's), (4, 67), (5, 91), (6, 136), (7, 190), (8, 202), (9, 226), (10, 229), (11, 262), (12, 283), (13, 430), (14, 496), (15, 547), (16, 559), (17, 562), (18, 640), (19, 652), (21, 679), (22, 772), (23, 799), (24, 808), (25, 862),

Gene: MontyDev_1 Start: 22, Stop: 885, Start Num: 2

Candidate Starts for MontyDev_1:

(Start: 1 @1 has 7 MA's), (2, 22), (Start: 3 @49 has 1 MA's), (4, 67), (5, 91), (6, 136), (7, 190), (8, 202), (9, 226), (10, 229), (11, 262), (12, 283), (13, 430), (14, 496), (15, 547), (16, 559), (17, 562), (18, 640), (19, 652), (20, 667), (21, 679), (22, 772), (23, 799), (25, 862),

Gene: Nilo_1 Start: 1, Stop: 885, Start Num: 1

Candidate Starts for Nilo_1:

(Start: 1 @1 has 7 MA's), (2, 22), (Start: 3 @49 has 1 MA's), (4, 67), (5, 91), (6, 136), (7, 190), (8, 202), (9, 226), (10, 229), (11, 262), (12, 283), (13, 430), (14, 496), (15, 547), (16, 559), (17, 562), (18, 640), (19, 652), (21, 679), (22, 772), (23, 799), (24, 808), (25, 862),

Gene: Papyrus_1 Start: 1, Stop: 885, Start Num: 1

Candidate Starts for Papyrus_1:

(Start: 1 @1 has 7 MA's), (2, 22), (Start: 3 @49 has 1 MA's), (4, 67), (5, 91), (6, 136), (7, 190), (8, 202), (9, 226), (10, 229), (11, 262), (12, 283), (13, 430), (14, 496), (15, 547), (16, 559), (17, 562), (18, 640), (19, 652), (21, 679), (22, 772), (23, 799), (24, 808), (25, 862),

Gene: Riparian_1 Start: 1, Stop: 885, Start Num: 1

Candidate Starts for Riparian_1:

(Start: 1 @1 has 7 MA's), (2, 22), (Start: 3 @49 has 1 MA's), (4, 67), (5, 91), (6, 136), (7, 190), (8, 202), (9, 226), (10, 229), (11, 262), (12, 283), (13, 430), (14, 496), (15, 547), (16, 559), (17, 562), (18, 640), (19, 652), (21, 679), (22, 772), (23, 799), (24, 808), (25, 862),

Gene: Rope_1 Start: 1, Stop: 885, Start Num: 1

Candidate Starts for Rope_1:

(Start: 1 @1 has 7 MA's), (2, 22), (Start: 3 @49 has 1 MA's), (4, 67), (5, 91), (6, 136), (7, 190), (8, 202), (9, 226), (10, 229), (11, 262), (12, 283), (13, 430), (14, 496), (15, 547), (16, 559), (17, 562), (18, 640), (19, 652), (21, 679), (22, 772), (23, 799), (24, 808), (25, 862),

Gene: Send513_1 Start: 1, Stop: 885, Start Num: 1

Candidate Starts for Send513_1:

(Start: 1 @1 has 7 MA's), (2, 22), (Start: 3 @49 has 1 MA's), (4, 67), (5, 91), (6, 136), (7, 190), (8, 202), (9, 226), (10, 229), (11, 262), (12, 283), (13, 430), (14, 496), (15, 547), (16, 559), (17, 562), (18, 640), (19, 652), (20, 667), (21, 679), (22, 772), (23, 799), (24, 808), (25, 862),

Gene: Weiss13_1 Start: 1, Stop: 885, Start Num: 1

Candidate Starts for Weiss13_1:

(Start: 1 @1 has 7 MA's), (2, 22), (Start: 3 @49 has 1 MA's), (4, 67), (5, 91), (6, 136), (7, 190), (8, 202), (9, 226), (10, 229), (11, 262), (12, 283), (13, 430), (14, 496), (15, 547), (16, 559), (17, 562), (18, 640), (19, 652), (21, 679), (22, 772), (23, 799), (24, 808), (25, 862),

Gene: Yelo_1 Start: 49, Stop: 885, Start Num: 3

Candidate Starts for Yelo_1:

(Start: 1 @1 has 7 MA's), (2, 22), (Start: 3 @49 has 1 MA's), (4, 67), (5, 91), (6, 136), (7, 190), (8, 202), (9, 226), (10, 229), (11, 262), (12, 283), (13, 430), (14, 496), (15, 547), (16, 559), (17, 562), (18, 640), (19, 652), (21, 679), (22, 772), (23, 799), (24, 808), (25, 862),

Gene: Zenon_1 Start: 49, Stop: 885, Start Num: 3

Candidate Starts for Zenon_1:

(Start: 1 @1 has 7 MA's), (2, 22), (Start: 3 @49 has 1 MA's), (4, 67), (5, 91), (6, 136), (7, 190), (8, 202), (9, 226), (10, 229), (11, 262), (12, 283), (13, 430), (14, 496), (15, 547), (16, 559), (17, 562), (18, 640), (19, 652), (21, 679), (22, 772), (23, 799), (24, 808), (25, 862),