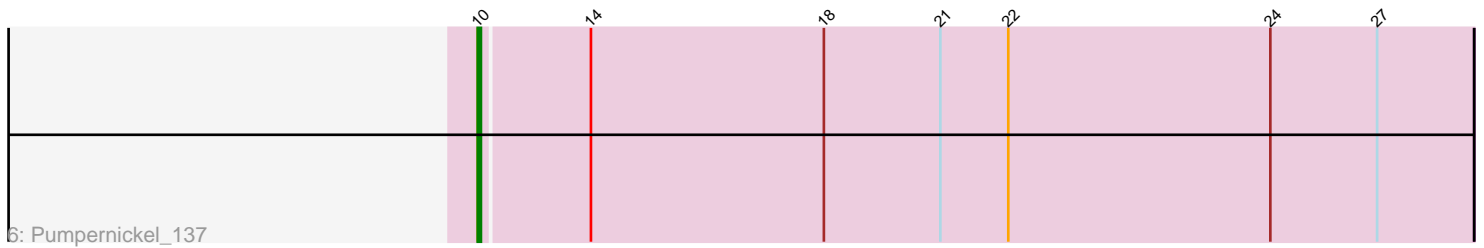
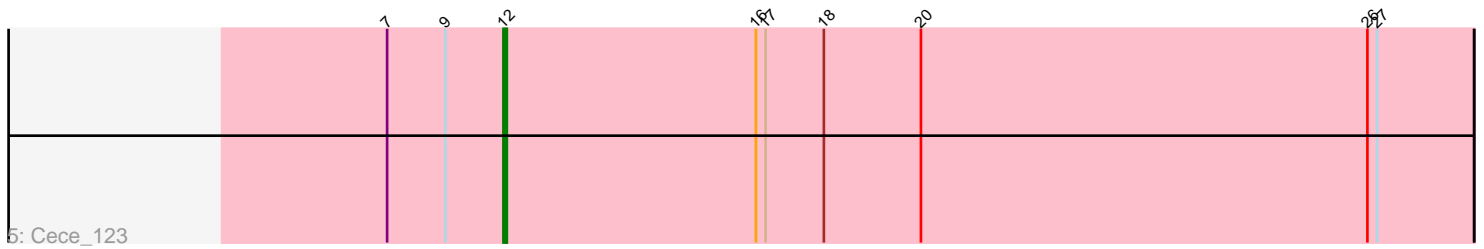
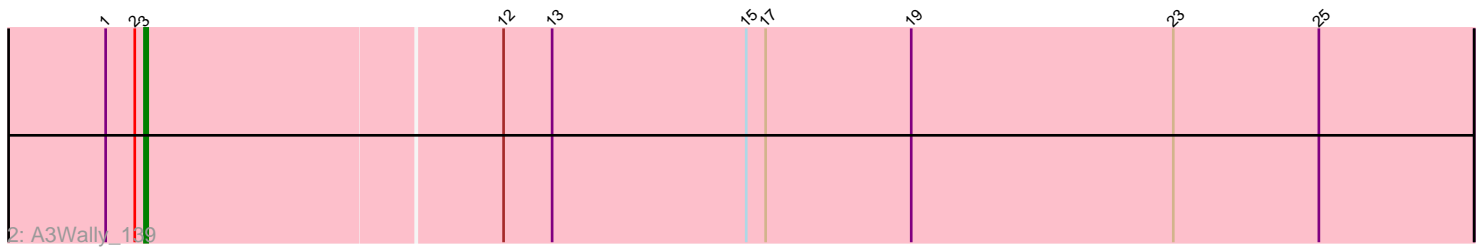
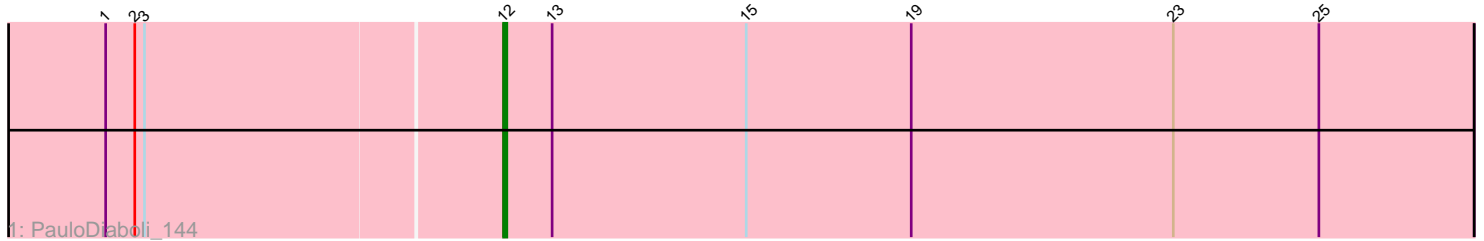


Pham 7460



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 7460 Report

This analysis was run 04/05/24 on database version 557.

Pham number 7460 has 6 members, 0 are drafts.

Phages represented in each track:

- Track 1 : PauloDiaboli_144
- Track 2 : A3Wally_139
- Track 3 : Zooman_123
- Track 4 : Big4_133
- Track 5 : Cece_123
- Track 6 : Pumpernickel_137

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

The start number called the most often in the published annotations is 12, it was called in 3 of the 6 non-draft genes in the pham.

Genes that call this "Most Annotated" start:

- Cece_123, PauloDiaboli_144, Zooman_123,

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

- A3Wally_139, Big4_133,

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

- Pumpernickel_137,

Summary by start number:

Start 3:

- Found in 2 of 6 (33.3%) of genes in pham
- Manual Annotations of this start: 1 of 6
- Called 50.0% of time when present
- Phage (with cluster) where this start called: A3Wally_139 (GD1),

Start 10:

- Found in 1 of 6 (16.7%) of genes in pham
- Manual Annotations of this start: 1 of 6
- Called 100.0% of time when present
- Phage (with cluster) where this start called: Pumpernickel_137 (GD4),

Start 11:

- Found in 2 of 6 (33.3%) of genes in pham
- Manual Annotations of this start: 1 of 6
- Called 50.0% of time when present
- Phage (with cluster) where this start called: Big4_133 (GD2),

Start 12:

- Found in 5 of 6 (83.3%) of genes in pham
- Manual Annotations of this start: 3 of 6
- Called 60.0% of time when present
- Phage (with cluster) where this start called: Cece_123 (GD3), PauloDiaboli_144 (GD1), Zooman_123 (GD2),

Summary by clusters:

There are 4 clusters represented in this pham: GD1, GD2, GD3, GD4,

Info for manual annotations of cluster GD1:

- Start number 3 was manually annotated 1 time for cluster GD1.
- Start number 12 was manually annotated 1 time for cluster GD1.

Info for manual annotations of cluster GD2:

- Start number 11 was manually annotated 1 time for cluster GD2.
- Start number 12 was manually annotated 1 time for cluster GD2.

Info for manual annotations of cluster GD3:

- Start number 12 was manually annotated 1 time for cluster GD3.

Info for manual annotations of cluster GD4:

- Start number 10 was manually annotated 1 time for cluster GD4.

Gene Information:

Gene: A3Wally_139 Start: 88444, Stop: 88851, Start Num: 3

Candidate Starts for A3Wally_139:

(1, 88432), (2, 88441), (Start: 3 @88444 has 1 MA's), (Start: 12 @88552 has 3 MA's), (13, 88567), (15, 88627), (17, 88633), (19, 88678), (23, 88759), (25, 88804),

Gene: Big4_133 Start: 87333, Stop: 87638, Start Num: 11

Candidate Starts for Big4_133:

(6, 87291), (8, 87321), (Start: 11 @87333 has 1 MA's), (Start: 12 @87339 has 3 MA's), (13, 87354), (19, 87465), (23, 87546),

Gene: Cece_123 Start: 89988, Stop: 90287, Start Num: 12

Candidate Starts for Cece_123:

(7, 89952), (9, 89970), (Start: 12 @89988 has 3 MA's), (16, 90066), (17, 90069), (18, 90087), (20, 90117), (26, 90255), (27, 90258),

Gene: PauloDiaboli_144 Start: 87894, Stop: 88193, Start Num: 12

Candidate Starts for PauloDiaboli_144:

(1, 87774), (2, 87783), (Start: 3 @87786 has 1 MA's), (Start: 12 @87894 has 3 MA's), (13, 87909), (15, 87969), (19, 88020), (23, 88101), (25, 88146),

Gene: Pumpernickel_137 Start: 89076, Stop: 89381, Start Num: 10

Candidate Starts for Pumpernickel_137:

(Start: 10 @89076 has 1 MA's), (14, 89109), (18, 89181), (21, 89217), (22, 89238), (24, 89319), (27, 89352),

Gene: Zooman_123 Start: 84687, Stop: 84986, Start Num: 12

Candidate Starts for Zooman_123:

(4, 84591), (5, 84597), (6, 84639), (8, 84669), (Start: 11 @84681 has 1 MA's), (Start: 12 @84687 has 3 MA's), (13, 84702), (19, 84813), (23, 84894), (26, 84954),