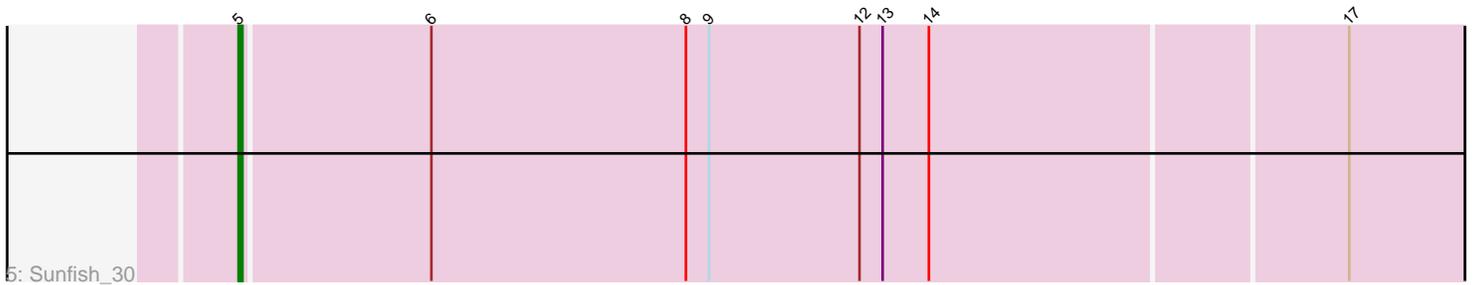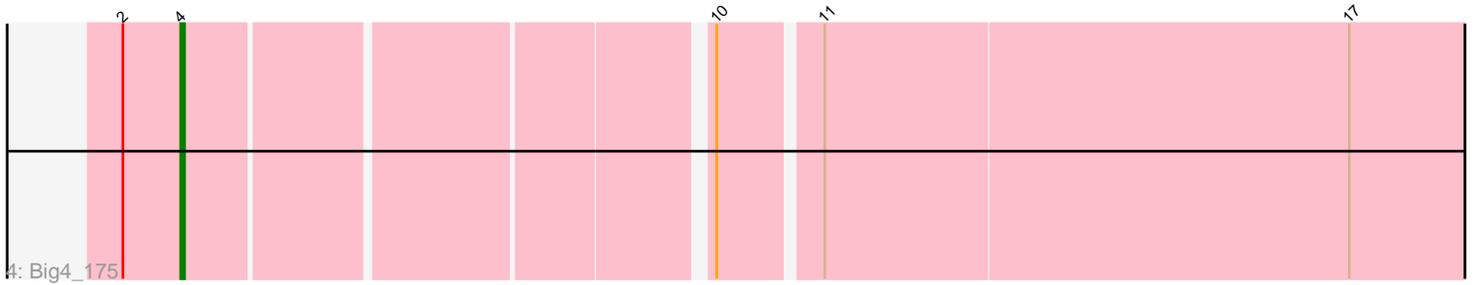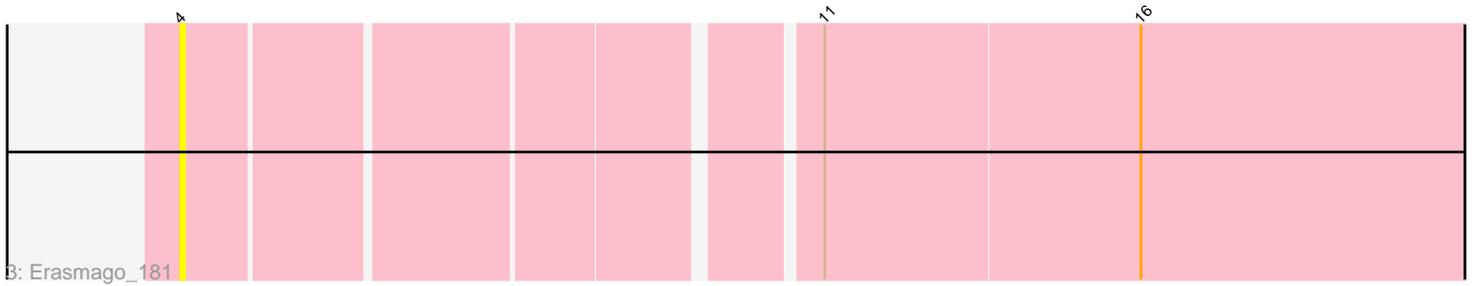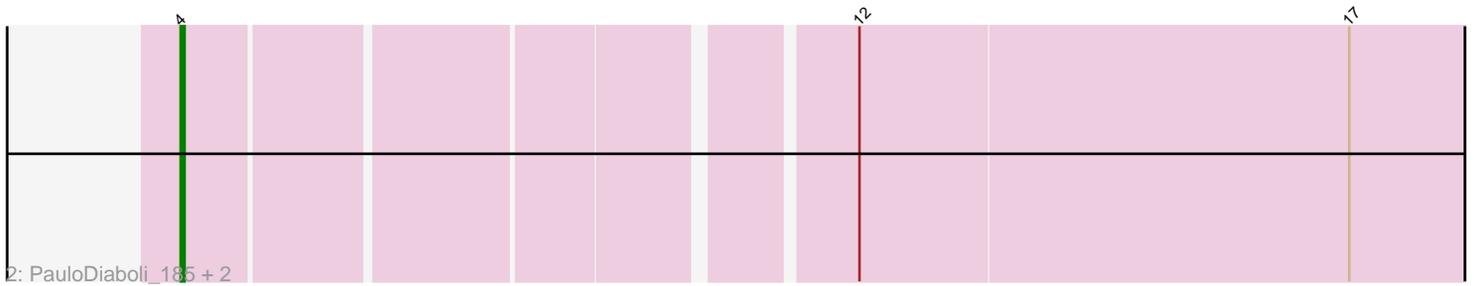# Pham 291773

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 291773 Report

This analysis was run 03/28/26 on database version 641.

Pham number 291773 has 9 members, 2 are drafts.

Phages represented in each track:
• Track 1 : Nason_2, Arcadia_2, Elsa_2
• Track 2 : PauloDiaboli_185, Dodo_187, A3Wally_186
• Track 3 : Erasmago_181
• Track 4 : Big4_175
• Track 5 : Sunfish_30

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 6 of the 7 non-draft genes in the pham.

Genes that call this "Most Annotated" start:
• A3Wally_186, Arcadia_2, Big4_175, Dodo_187, Elsa_2, Erasmago_181, Nason_2, PauloDiaboli_185,

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

Genes that do not have the "Most Annotated" start:
• Sunfish_30,

Summary by start number:

Start 4:
• Found in 8 of 9 (88.9%) of genes in pham
• Manual Annotations of this start: 6 of 7
• Called 100.0% of time when present
• Phage (with cluster) where this start called: A3Wally_186 (GD1), Arcadia_2 (AM), Big4_175 (GD2), Dodo_187 (GD1), Elsa_2 (AM), Erasmago_181 (GD2), Nason_2 (AM), PauloDiaboli_185 (GD1),

Start 5:
• Found in 1 of 9 (11.1%) of genes in pham
• Manual Annotations of this start: 1 of 7

• Called 100.0% of time when present
• Phage (with cluster) where this start called: Sunfish_30 (singleton),

Summary by clusters:

There are 4 clusters represented in this pham: GD1, GD2, AM, singleton,

Info for manual annotations of cluster AM:
•Start number 4 was manually annotated 3 times for cluster AM.

Info for manual annotations of cluster GD1:
•Start number 4 was manually annotated 2 times for cluster GD1.

Info for manual annotations of cluster GD2:
•Start number 4 was manually annotated 1 time for cluster GD2.

Gene Information:
Gene: A3Wally_186 Start: 101532, Stop: 101870, Start Num: 4
Candidate Starts for A3Wally_186:
(Start: 4 @101532 has 6 MA's), (12, 101691), (17, 101817),

Gene: Arcadia_2 Start: 525, Stop: 869, Start Num: 4
Candidate Starts for Arcadia_2:
(1, 486), (3, 522), (Start: 4 @525 has 6 MA's), (7, 591), (13, 702), (14, 714), (15, 750),

Gene: Big4_175 Start: 98404, Stop: 98742, Start Num: 4
Candidate Starts for Big4_175:
(2, 98389), (Start: 4 @98404 has 6 MA's), (10, 98530), (11, 98554), (17, 98689),

Gene: Dodo_187 Start: 101144, Stop: 101482, Start Num: 4
Candidate Starts for Dodo_187:
(Start: 4 @101144 has 6 MA's), (12, 101303), (17, 101429),

Gene: Elsa_2 Start: 525, Stop: 869, Start Num: 4
Candidate Starts for Elsa_2:
(1, 486), (3, 522), (Start: 4 @525 has 6 MA's), (7, 591), (13, 702), (14, 714), (15, 750),

Gene: Erasmago_181 Start: 98728, Stop: 99066, Start Num: 4
Candidate Starts for Erasmago_181:
(Start: 4 @98728 has 6 MA's), (11, 98878), (16, 98959),

Gene: Nason_2 Start: 525, Stop: 869, Start Num: 4
Candidate Starts for Nason_2:
(1, 486), (3, 522), (Start: 4 @525 has 6 MA's), (7, 591), (13, 702), (14, 714), (15, 750),

Gene: PauloDiaboli_185 Start: 99579, Stop: 99917, Start Num: 4
Candidate Starts for PauloDiaboli_185:
(Start: 4 @99579 has 6 MA's), (12, 99738), (17, 99864),

Gene: Sunfish_30 Start: 14435, Stop: 14770, Start Num: 5

Candidate Starts for Sunfish_30:
(Start: 5 @14435 has 1 MA's), (6, 14483), (8, 14549), (9, 14555), (12, 14594), (13, 14600), (14, 14612), (17, 14717),