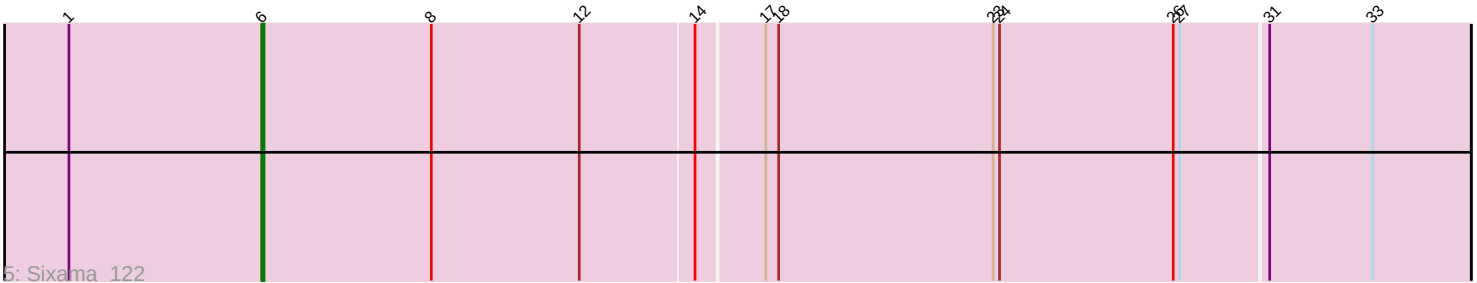
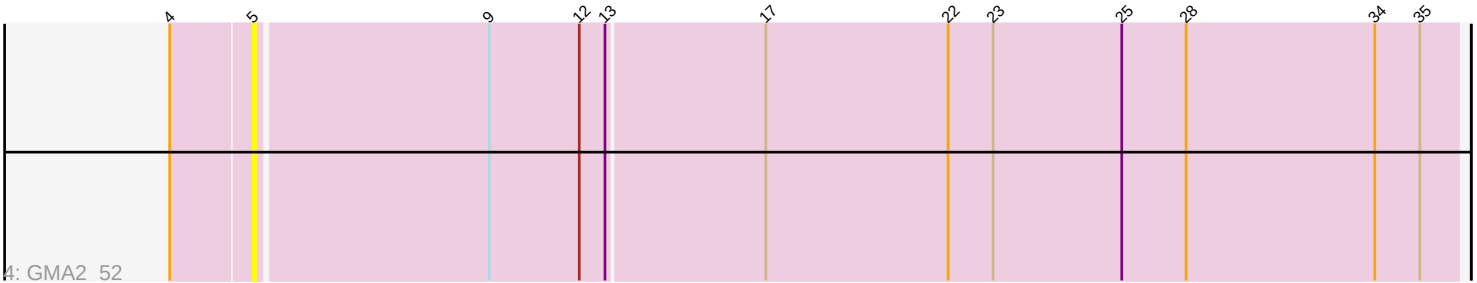
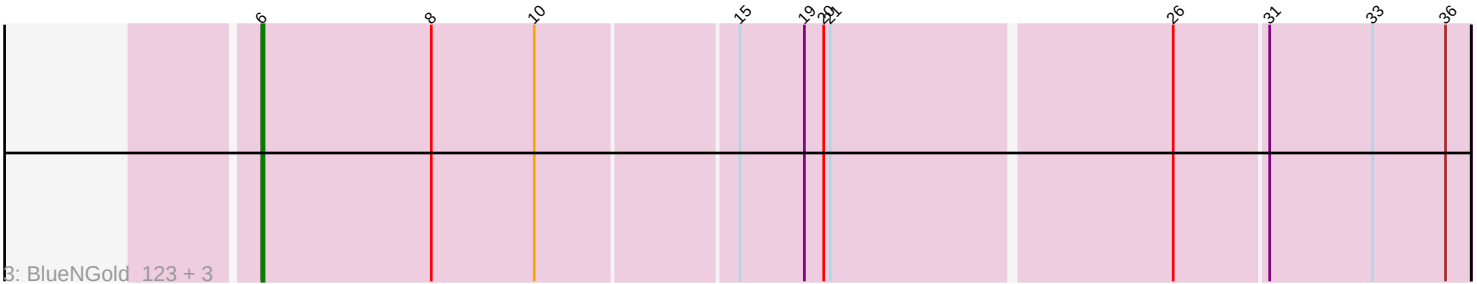
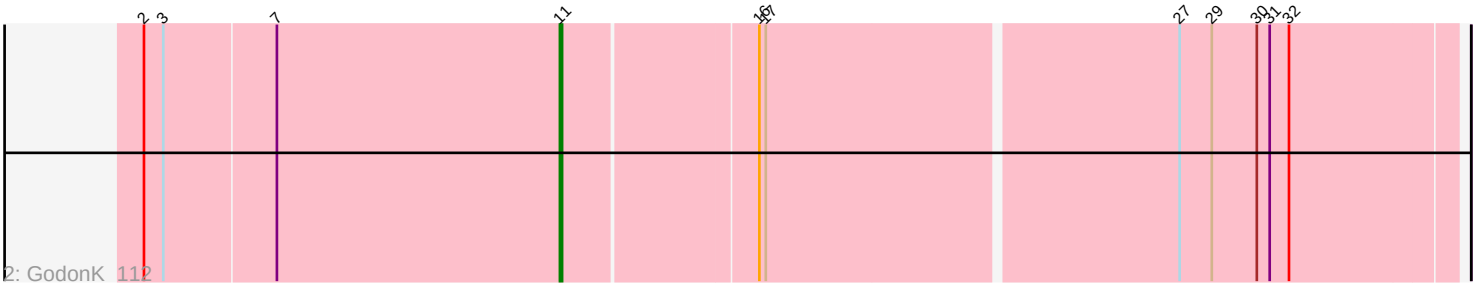
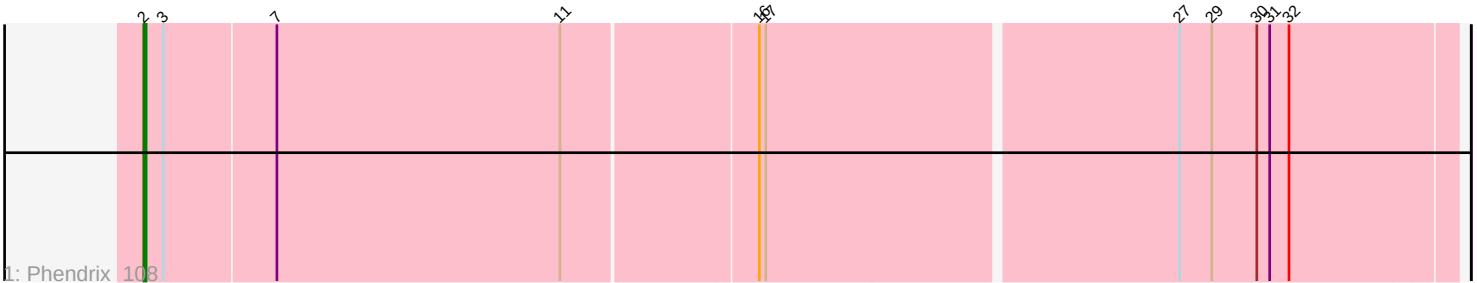


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Note: Tracks are now grouped by subcluster and scaled. Switching in subcluster is indicated by changes in track color. Track scale is now set by default to display the region 30 bp upstream of start 1 to 30 bp downstream of the last possible start. 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 318280 Report

This analysis was run 09/05/26 on database version 662.

Pham number 318280 has 8 members, 1 are drafts.

Phages represented in each track:

- Track 1 : Phendrix_108
- Track 2 : GodonK_112
- Track 3 : BlueNGold_123, Forza_124, Mareelih_122, Boopy_124
- Track 4 : GMA2_52
- Track 5 : Sixama_122

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

The start number called the most often in the published annotations is 6, it was called in 5 of the 7 non-draft genes in the pham.

Genes that call this "Most Annotated" start:

- BlueNGold_123, Boopy_124, Forza_124, Mareelih_122, Sixama_122,

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

-

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

- GMA2_52, GodonK_112, Phendrix_108,

Summary by start number:

Start 2:

- Found in 2 of 8 (25.0%) of genes in pham
- Manual Annotations of this start: 1 of 7
- Called 50.0% of time when present
- Phage (with cluster) where this start called: Phendrix_108 (DK),

Start 5:

- Found in 1 of 8 (12.5%) of genes in pham
- No Manual Annotations of this start.
- Called 100.0% of time when present
- Phage (with cluster) where this start called: GMA2_52 (DS),

Start 6:

- Found in 5 of 8 (62.5%) of genes in pham
- Manual Annotations of this start: 5 of 7
- Called 100.0% of time when present
- Phage (with cluster) where this start called: BlueNGold_123 (DS), Boopy_124 (DS), Forza_124 (DS), Mareelih_122 (DS), Sixama_122 (DS),

Start 11:

- Found in 2 of 8 (25.0%) of genes in pham
- Manual Annotations of this start: 1 of 7
- Called 50.0% of time when present
- Phage (with cluster) where this start called: GodonK_112 (DK),

Summary by clusters:

There are 2 clusters represented in this pham: DS, DK,

Info for manual annotations of cluster DK:

- Start number 2 was manually annotated 1 time for cluster DK.
- Start number 11 was manually annotated 1 time for cluster DK.

Info for manual annotations of cluster DS:

- Start number 6 was manually annotated 5 times for cluster DS.

Gene Information:

Gene: BlueNGold_123 Start: 79020, Stop: 78475, Start Num: 6

Candidate Starts for BlueNGold_123:

(Start: 6 @79020 has 5 MA's), (8, 78942), (10, 78894), (15, 78804), (19, 78774), (20, 78765), (21, 78762), (26, 78609), (31, 78567), (33, 78519), (36, 78486),

Gene: Boopy_124 Start: 79032, Stop: 78487, Start Num: 6

Candidate Starts for Boopy_124:

(Start: 6 @79032 has 5 MA's), (8, 78954), (10, 78906), (15, 78816), (19, 78786), (20, 78777), (21, 78774), (26, 78621), (31, 78579), (33, 78531), (36, 78498),

Gene: Forza_124 Start: 78948, Stop: 78403, Start Num: 6

Candidate Starts for Forza_124:

(Start: 6 @78948 has 5 MA's), (8, 78870), (10, 78822), (15, 78732), (19, 78702), (20, 78693), (21, 78690), (26, 78537), (31, 78495), (33, 78447), (36, 78414),

Gene: GMA2_52 Start: 53375, Stop: 52827, Start Num: 5

Candidate Starts for GMA2_52:

(4, 53411), (5, 53375), (9, 53270), (12, 53228), (13, 53216), (17, 53144), (22, 53060), (23, 53039), (25, 52979), (28, 52949), (34, 52865), (35, 52844),

Gene: GodonK_112 Start: 69704, Stop: 69303, Start Num: 11

Candidate Starts for GodonK_112:

(Start: 2 @69896 has 1 MA's), (3, 69887), (7, 69836), (Start: 11 @69704 has 1 MA's), (16, 69617), (17, 69614), (27, 69428), (29, 69413), (30, 69392), (31, 69386), (32, 69377),

Gene: Mareelih_122 Start: 78450, Stop: 77905, Start Num: 6

Candidate Starts for Mareelih_122:

(Start: 6 @78450 has 5 MA's), (8, 78372), (10, 78324), (15, 78234), (19, 78204), (20, 78195), (21, 78192), (26, 78039), (31, 77997), (33, 77949), (36, 77916),

Gene: Phendrix_108 Start: 70145, Stop: 69552, Start Num: 2

Candidate Starts for Phendrix_108:

(Start: 2 @70145 has 1 MA's), (3, 70136), (7, 70085), (Start: 11 @69953 has 1 MA's), (16, 69866), (17, 69863), (27, 69677), (29, 69662), (30, 69641), (31, 69635), (32, 69626),

Gene: Sixama_122 Start: 78458, Stop: 77907, Start Num: 6

Candidate Starts for Sixama_122:

(1, 78548), (Start: 6 @78458 has 5 MA's), (8, 78380), (12, 78311), (14, 78260), (17, 78230), (18, 78224), (23, 78125), (24, 78122), (26, 78041), (27, 78038), (31, 77999), (33, 77951),