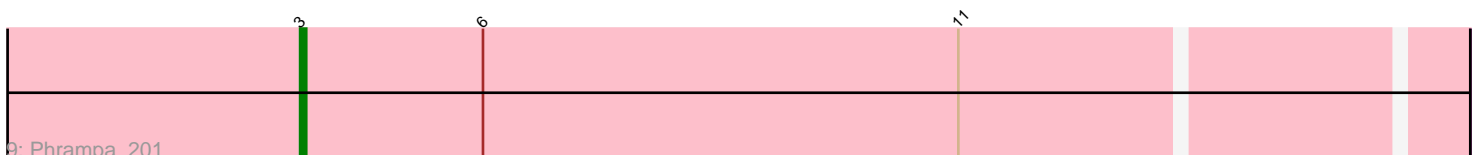
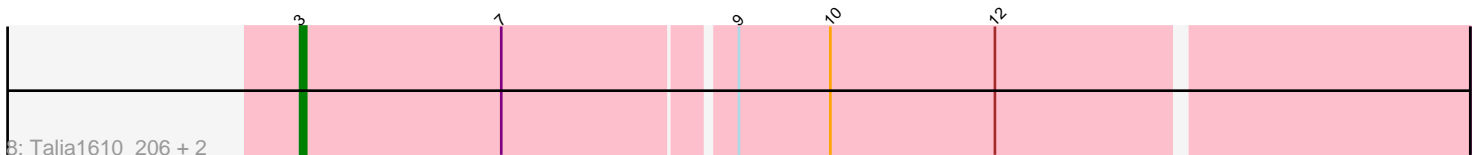
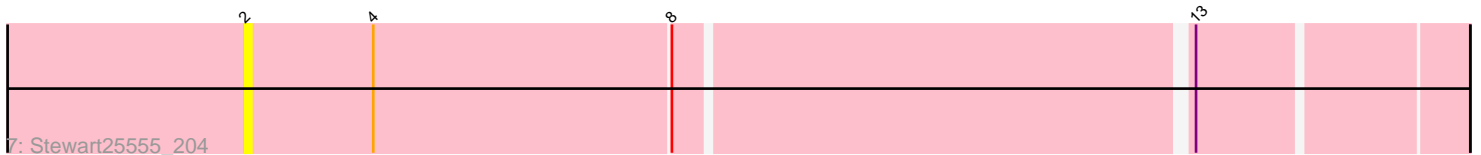
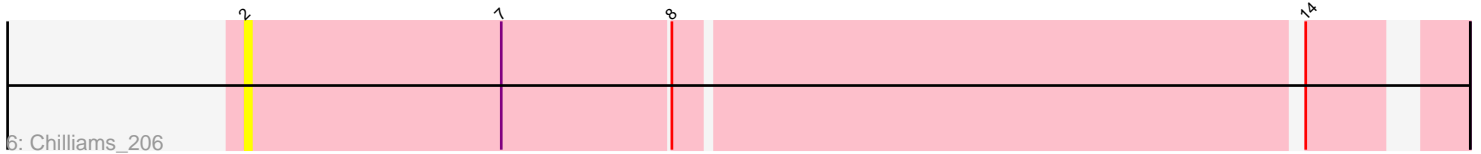
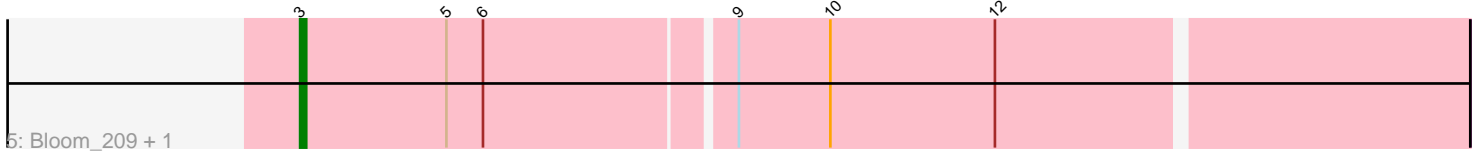
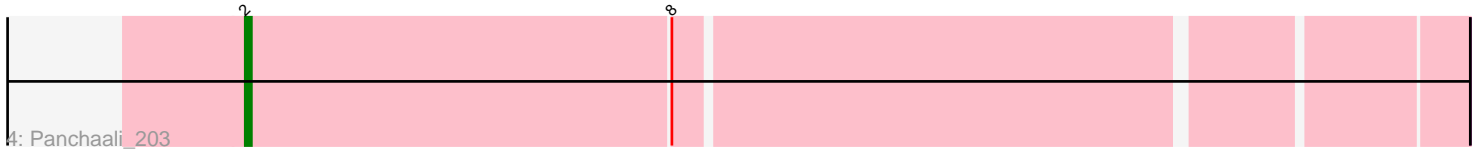
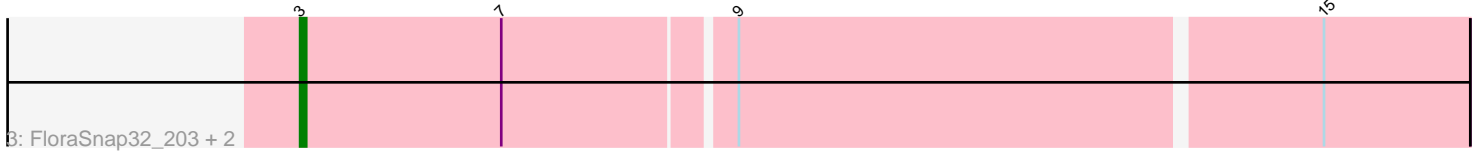
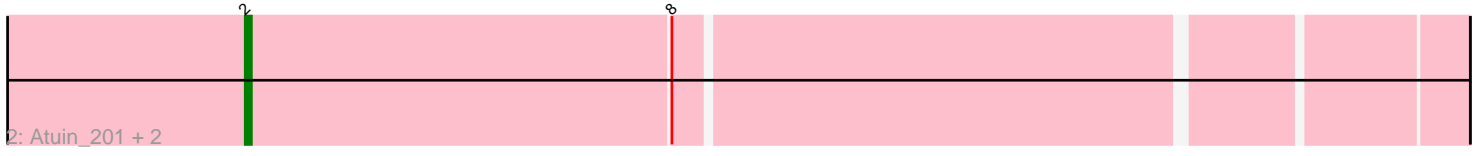
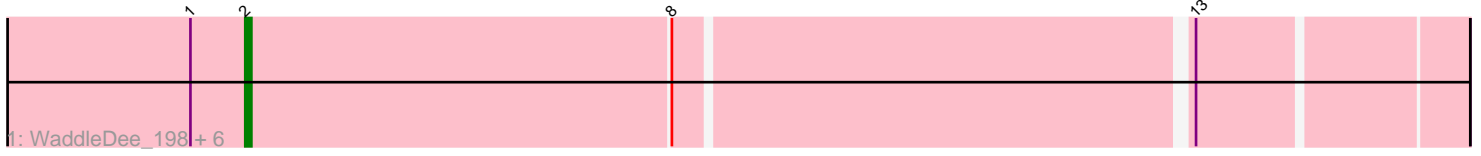


Pham 309232



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 309232 Report

This analysis was run 06/27/26 on database version 652.

Pham number 309232 has 23 members, 11 are drafts.

Phages represented in each track:

- Track 1 : WaddleDee_198, DunneganBoMo_203, Emmetator_206, Ellewin_201, KSunshine22_206, BooTeria_211, Artu_204
- Track 2 : Atuin_201, ReginaGlobina_213, LeoJr_210
- Track 3 : FloraSnap32_203, Patbob_202, GoldenEssence_193
- Track 4 : Panchaali_203
- Track 5 : Bloom_209, Racecar_206
- Track 6 : Chilliams_206
- Track 7 : Stewart25555_204
- Track 8 : Talia1610_206, Mimi_204, FrostedClock_208
- Track 9 : Phrampa_201
- Track 10 : SJReid_213

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

The start number called the most often in the published annotations is 2, it was called in 6 of the 12 non-draft genes in the pham.

Genes that call this "Most Annotated" start:

- Artu_204, Atuin_201, BooTeria_211, Chilliams_206, DunneganBoMo_203, Ellewin_201, Emmetator_206, KSunshine22_206, LeoJr_210, Panchaali_203, ReginaGlobina_213, SJReid_213, Stewart25555_204, WaddleDee_198,

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

-

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

- Bloom_209, FloraSnap32_203, FrostedClock_208, GoldenEssence_193, Mimi_204, Patbob_202, Phrampa_201, Racecar_206, Talia1610_206,

Summary by start number:

Start 2:

- Found in 14 of 23 (60.9%) of genes in pham
- Manual Annotations of this start: 6 of 12

- Called 100.0% of time when present
- Phage (with cluster) where this start called: Artu_204 (FC), Atuin_201 (FC), BooTeria_211 (FC), Chilliams_206 (FC), DunneganBoMo_203 (FC), Ellewin_201 (FC), Emmetator_206 (FC), KSunshine22_206 (FC), LeoJr_210 (FC), Panchaali_203 (FC), ReginaGlobina_213 (FC), SJReid_213 (FC), Stewart25555_204 (FC), WaddleDee_198 (FC),

Start 3:

- Found in 9 of 23 (39.1%) of genes in pham
- Manual Annotations of this start: 6 of 12
- Called 100.0% of time when present
- Phage (with cluster) where this start called: Bloom_209 (FC), FloraSnap32_203 (FC), FrostedClock_208 (FC), GoldenEssence_193 (FC), Mimi_204 (FC), Patbob_202 (FC), Phrampa_201 (FC), Racecar_206 (FC), Talia1610_206 (FC),

Summary by clusters:

There is one cluster represented in this pham: FC

Info for manual annotations of cluster FC:

- Start number 2 was manually annotated 6 times for cluster FC.
- Start number 3 was manually annotated 6 times for cluster FC.

Gene Information:

Gene: Artu_204 Start: 148044, Stop: 148235, Start Num: 2

Candidate Starts for Artu_204:

(1, 148035), (Start: 2 @148044 has 6 MA's), (8, 148113), (13, 148194),

Gene: Atuin_201 Start: 141205, Stop: 141396, Start Num: 2

Candidate Starts for Atuin_201:

(Start: 2 @141205 has 6 MA's), (8, 141274),

Gene: Bloom_209 Start: 142256, Stop: 142441, Start Num: 3

Candidate Starts for Bloom_209:

(Start: 3 @142256 has 6 MA's), (5, 142280), (6, 142286), (9, 142325), (10, 142340), (12, 142367),

Gene: BooTeria_211 Start: 147554, Stop: 147745, Start Num: 2

Candidate Starts for BooTeria_211:

(1, 147545), (Start: 2 @147554 has 6 MA's), (8, 147623), (13, 147704),

Gene: Chilliams_206 Start: 139068, Stop: 139256, Start Num: 2

Candidate Starts for Chilliams_206:

(Start: 2 @139068 has 6 MA's), (7, 139110), (8, 139137), (14, 139236),

Gene: DunneganBoMo_203 Start: 147018, Stop: 147209, Start Num: 2

Candidate Starts for DunneganBoMo_203:

(1, 147009), (Start: 2 @147018 has 6 MA's), (8, 147087), (13, 147168),

Gene: Ellewin_201 Start: 146285, Stop: 146476, Start Num: 2

Candidate Starts for Ellewin_201:

Gene: Stewart25555_204 Start: 144933, Stop: 145124, Start Num: 2
Candidate Starts for Stewart25555_204:
(Start: 2 @144933 has 6 MA's), (4, 144954), (8, 145002), (13, 145083),

Gene: Talia1610_206 Start: 142041, Stop: 142226, Start Num: 3
Candidate Starts for Talia1610_206:
(Start: 3 @142041 has 6 MA's), (7, 142074), (9, 142110), (10, 142125), (12, 142152),

Gene: WaddleDee_198 Start: 145479, Stop: 145670, Start Num: 2
Candidate Starts for WaddleDee_198:
(1, 145470), (Start: 2 @145479 has 6 MA's), (8, 145548), (13, 145629),