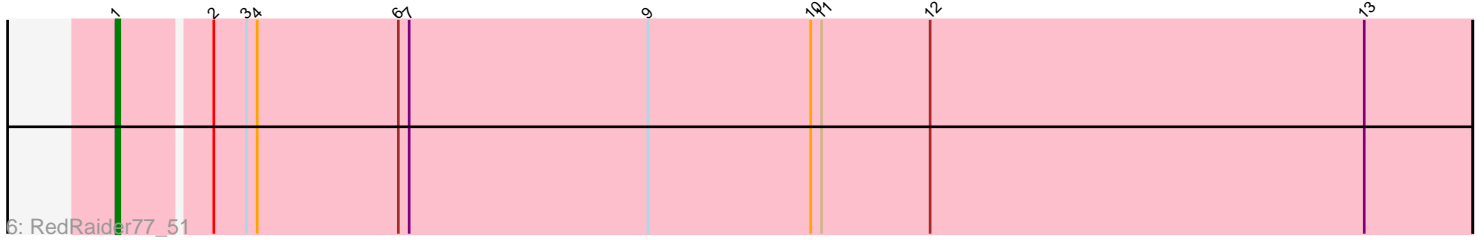
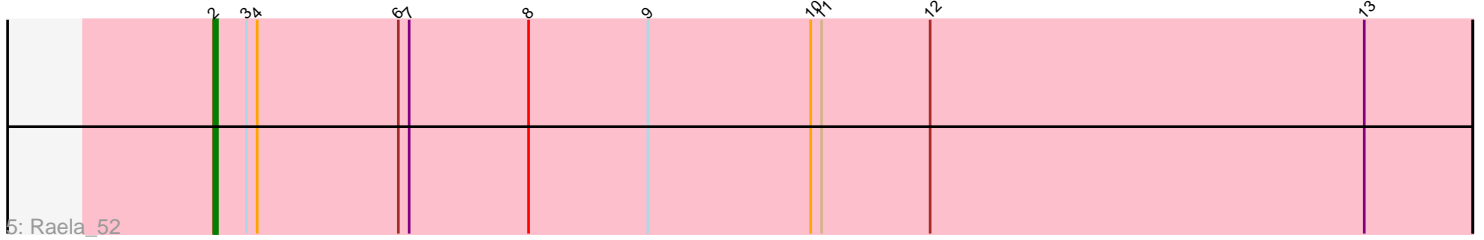
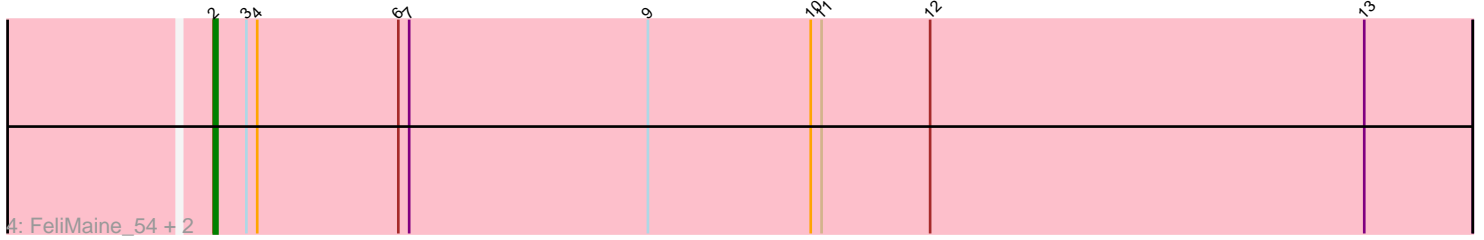
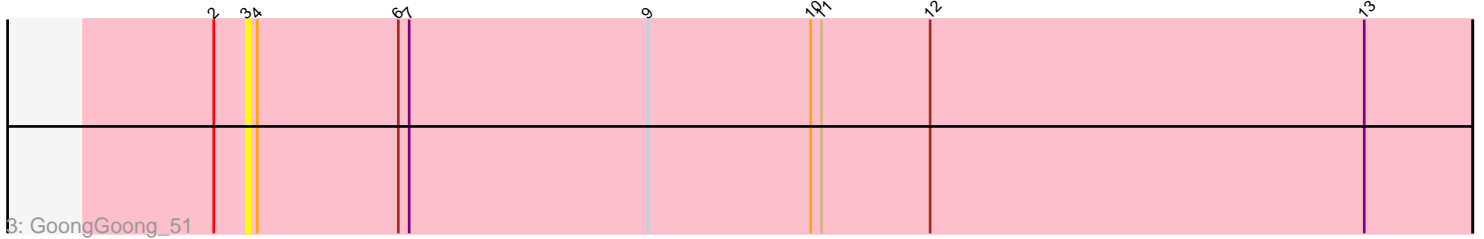
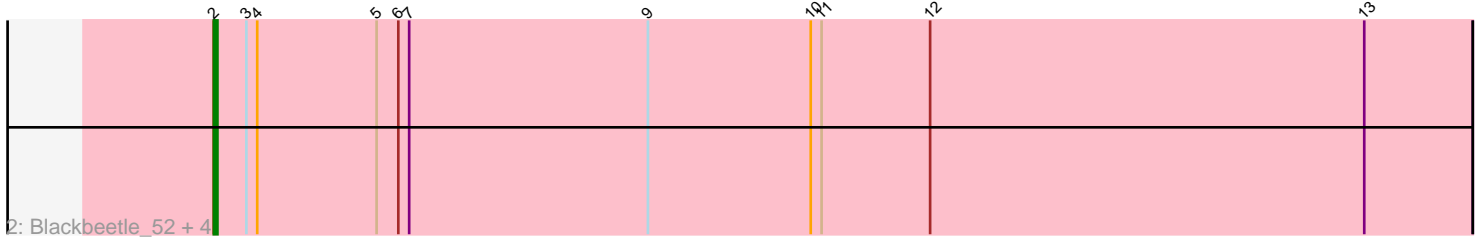
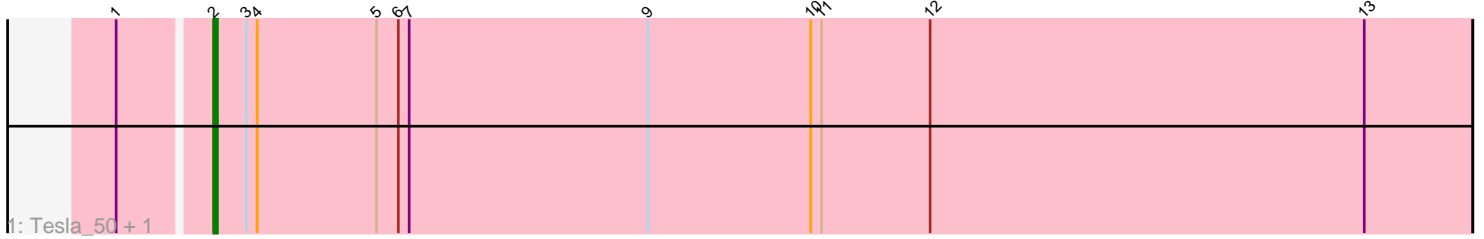


Pham 214798



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 214798 Report

This analysis was run 02/22/25 on database version 588.

Pham number 214798 has 13 members, 1 are drafts.

Phages represented in each track:

- Track 1 : Tesla_50, Marvin_49
- Track 2 : Blackbeetle_52, Pringar_52, VasuNzinga_52, Beelzebub_56, Poise_52
- Track 3 : GoongGoong_51
- Track 4 : FeliMaine_54, MosMoris_50, Gattaca_51
- Track 5 : Raela_52
- Track 6 : RedRaider77_51

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 11 of the 12 non-draft genes in the pham.

Genes that call this "Most Annotated" start:

- Beelzebub_56, Blackbeetle_52, FeliMaine_54, Gattaca_51, Marvin_49, MosMoris_50, Poise_52, Pringar_52, Raela_52, Tesla_50, VasuNzinga_52,

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

- GoongGoong_51, RedRaider77_51,

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

-

Summary by start number:

Start 1:

- Found in 3 of 13 (23.1%) of genes in pham
- Manual Annotations of this start: 1 of 12
- Called 33.3% of time when present
- Phage (with cluster) where this start called: RedRaider77_51 (S),

Start 2:

- Found in 13 of 13 (100.0%) of genes in pham
- Manual Annotations of this start: 11 of 12
- Called 84.6% of time when present

- Phage (with cluster) where this start called: Beelzebub_56 (S), Blackbeetle_52 (S), FeliMaine_54 (S), Gattaca_51 (S), Marvin_49 (S), MosMoris_50 (S), Poise_52 (S), Pringar_52 (S), Raela_52 (S), Tesla_50 (S), VasuNzinga_52 (S),

Start 3:

- Found in 13 of 13 (100.0%) of genes in pham
- No Manual Annotations of this start.
- Called 7.7% of time when present
- Phage (with cluster) where this start called: GoongGoong_51 (S),

Summary by clusters:

There is one cluster represented in this pham: S

Info for manual annotations of cluster S:

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

Gene Information:

Gene: Beelzebub_56 Start: 29928, Stop: 30275, Start Num: 2

Candidate Starts for Beelzebub_56:

(Start: 2 @29928 has 11 MA's), (3, 29937), (4, 29940), (5, 29973), (6, 29979), (7, 29982), (9, 30048), (10, 30093), (11, 30096), (12, 30126), (13, 30246),

Gene: Blackbeetle_52 Start: 28974, Stop: 29321, Start Num: 2

Candidate Starts for Blackbeetle_52:

(Start: 2 @28974 has 11 MA's), (3, 28983), (4, 28986), (5, 29019), (6, 29025), (7, 29028), (9, 29094), (10, 29139), (11, 29142), (12, 29172), (13, 29292),

Gene: FeliMaine_54 Start: 29898, Stop: 30245, Start Num: 2

Candidate Starts for FeliMaine_54:

(Start: 2 @29898 has 11 MA's), (3, 29907), (4, 29910), (6, 29949), (7, 29952), (9, 30018), (10, 30063), (11, 30066), (12, 30096), (13, 30216),

Gene: Gattaca_51 Start: 29077, Stop: 29424, Start Num: 2

Candidate Starts for Gattaca_51:

(Start: 2 @29077 has 11 MA's), (3, 29086), (4, 29089), (6, 29128), (7, 29131), (9, 29197), (10, 29242), (11, 29245), (12, 29275), (13, 29395),

Gene: GoongGoong_51 Start: 28885, Stop: 29223, Start Num: 3

Candidate Starts for GoongGoong_51:

(Start: 2 @28876 has 11 MA's), (3, 28885), (4, 28888), (6, 28927), (7, 28930), (9, 28996), (10, 29041), (11, 29044), (12, 29074), (13, 29194),

Gene: Marvin_49 Start: 29144, Stop: 29491, Start Num: 2

Candidate Starts for Marvin_49:

(Start: 1 @29120 has 1 MA's), (Start: 2 @29144 has 11 MA's), (3, 29153), (4, 29156), (5, 29189), (6, 29195), (7, 29198), (9, 29264), (10, 29309), (11, 29312), (12, 29342), (13, 29462),

Gene: MosMoris_50 Start: 29077, Stop: 29424, Start Num: 2

Candidate Starts for MosMoris_50:

(Start: 2 @29077 has 11 MA's), (3, 29086), (4, 29089), (6, 29128), (7, 29131), (9, 29197), (10, 29242), (11, 29245), (12, 29275), (13, 29395),

Gene: Poise_52 Start: 28974, Stop: 29321, Start Num: 2

Candidate Starts for Poise_52:

(Start: 2 @28974 has 11 MA's), (3, 28983), (4, 28986), (5, 29019), (6, 29025), (7, 29028), (9, 29094), (10, 29139), (11, 29142), (12, 29172), (13, 29292),

Gene: Pringar_52 Start: 29262, Stop: 29609, Start Num: 2

Candidate Starts for Pringar_52:

(Start: 2 @29262 has 11 MA's), (3, 29271), (4, 29274), (5, 29307), (6, 29313), (7, 29316), (9, 29382), (10, 29427), (11, 29430), (12, 29460), (13, 29580),

Gene: Raela_52 Start: 29529, Stop: 29876, Start Num: 2

Candidate Starts for Raela_52:

(Start: 2 @29529 has 11 MA's), (3, 29538), (4, 29541), (6, 29580), (7, 29583), (8, 29616), (9, 29649), (10, 29694), (11, 29697), (12, 29727), (13, 29847),

Gene: RedRaider77_51 Start: 28894, Stop: 29265, Start Num: 1

Candidate Starts for RedRaider77_51:

(Start: 1 @28894 has 1 MA's), (Start: 2 @28918 has 11 MA's), (3, 28927), (4, 28930), (6, 28969), (7, 28972), (9, 29038), (10, 29083), (11, 29086), (12, 29116), (13, 29236),

Gene: Tesla_50 Start: 28767, Stop: 29114, Start Num: 2

Candidate Starts for Tesla_50:

(Start: 1 @28743 has 1 MA's), (Start: 2 @28767 has 11 MA's), (3, 28776), (4, 28779), (5, 28812), (6, 28818), (7, 28821), (9, 28887), (10, 28932), (11, 28935), (12, 28965), (13, 29085),

Gene: VasuNzinga_52 Start: 28836, Stop: 29183, Start Num: 2

Candidate Starts for VasuNzinga_52:

(Start: 2 @28836 has 11 MA's), (3, 28845), (4, 28848), (5, 28881), (6, 28887), (7, 28890), (9, 28956), (10, 29001), (11, 29004), (12, 29034), (13, 29154),