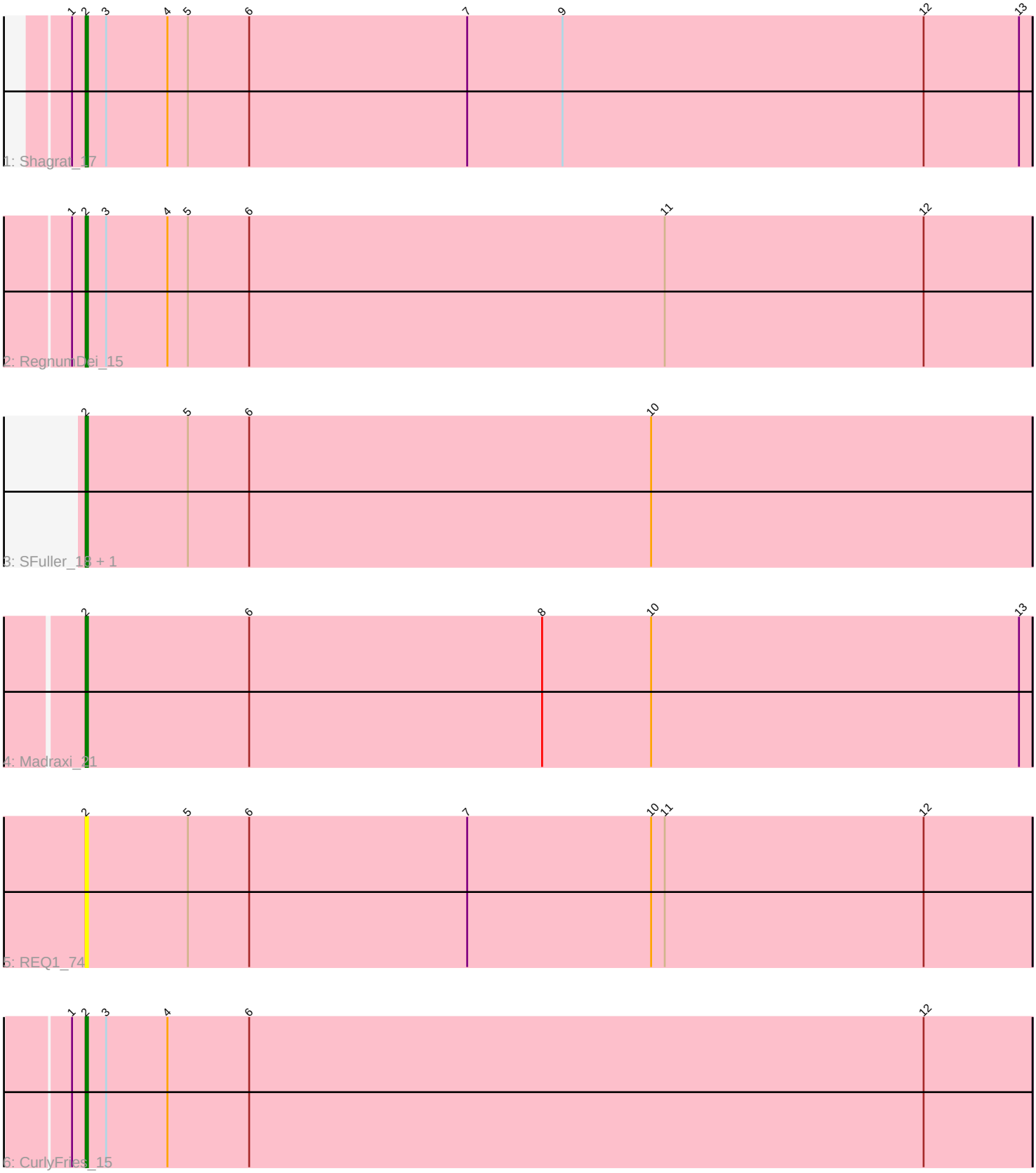
Pham 346535


1 2 3 4 5 6 7 9 12 13
1: Shagrat_17
1 2 3 4 5 6 11 12
2: RegnumDei_15
2 5 6 10
3: SFuller_13 + 1
2 6 8 10 13
4: Madraxi_21
2 5 6 7 10 11 12
5: REQ1_74
1 2 3 4 6 12
6: CurlyFries_15

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 346535 Report

This analysis was run 11/14/26 on database version 672.

Pham number 346535 has 7 members, 1 are drafts.

Phages represented in each track:
• Track 1 : Shagrat_17
• Track 2 : RegnumDei_15
• Track 3 : SFuller_18, Jflix2_16
• Track 4 : Madraxi_21
• Track 5 : REQ1_74
• Track 6 : CurlyFries_15

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 6 non-draft genes in the pham.

Genes that call this "Most Annotated" start:
• CurlyFries_15, Jflix2_16, Madraxi_21, REQ1_74, RegnumDei_15, SFuller_18, Shagrat_17,

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

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

Summary by start number:

Start 2:
• Found in 7 of 7 (100.0%) of genes in pham
• Manual Annotations of this start: 6 of 6
• Called 100.0% of time when present
• Phage (with cluster) where this start called: CurlyFries_15 (CF), Jflix2_16 (CF), Madraxi_21 (CF), REQ1_74 (CF), RegnumDei_15 (CF), SFuller_18 (CF), Shagrat_17 (CF),

Summary by clusters:

There is one cluster represented in this pham: CF

Info for manual annotations of cluster CF:
•Start number 2 was manually annotated 6 times for cluster CF.

Gene Information:

Gene: CurlyFries_15 Start: 10745, Stop: 11161, Start Num: 2
Candidate Starts for CurlyFries_15:
(1, 10739), (Start: 2 @10745 has 6 MA's), (3, 10754), (4, 10781), (6, 10817), (12, 11114),

Gene: Jflix2_16 Start: 12854, Stop: 13270, Start Num: 2
Candidate Starts for Jflix2_16:
(Start: 2 @12854 has 6 MA's), (5, 12899), (6, 12926), (10, 13103),

Gene: Madraxi_21 Start: 15288, Stop: 15704, Start Num: 2
Candidate Starts for Madraxi_21:
(Start: 2 @15288 has 6 MA's), (6, 15360), (8, 15489), (10, 15537), (13, 15699),

Gene: REQ1_74 Start: 36756, Stop: 37172, Start Num: 2
Candidate Starts for REQ1_74:
(Start: 2 @36756 has 6 MA's), (5, 36801), (6, 36828), (7, 36924), (10, 37005), (11, 37011), (12, 37125),

Gene: RegnumDei_15 Start: 11238, Stop: 11654, Start Num: 2
Candidate Starts for RegnumDei_15:
(1, 11232), (Start: 2 @11238 has 6 MA's), (3, 11247), (4, 11274), (5, 11283), (6, 11310), (11, 11493), (12, 11607),

Gene: SFuller_18 Start: 13469, Stop: 13885, Start Num: 2
Candidate Starts for SFuller_18:
(Start: 2 @13469 has 6 MA's), (5, 13514), (6, 13541), (10, 13718),

Gene: Shagrat_17 Start: 12520, Stop: 12936, Start Num: 2
Candidate Starts for Shagrat_17:
(1, 12514), (Start: 2 @12520 has 6 MA's), (3, 12529), (4, 12556), (5, 12565), (6, 12592), (7, 12688), (9, 12730), (12, 12889), (13, 12931),