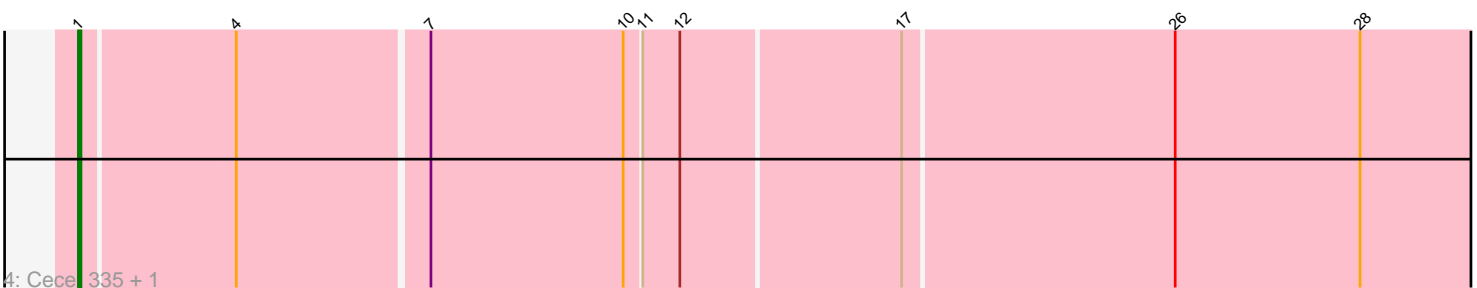
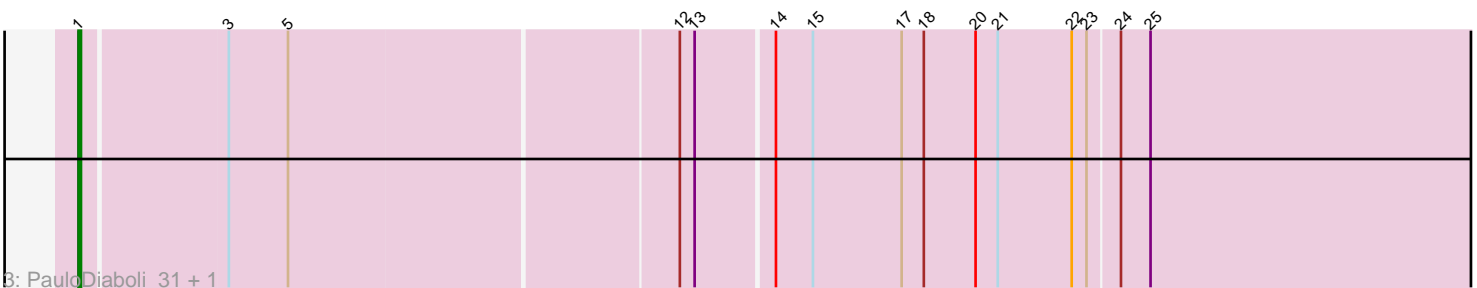
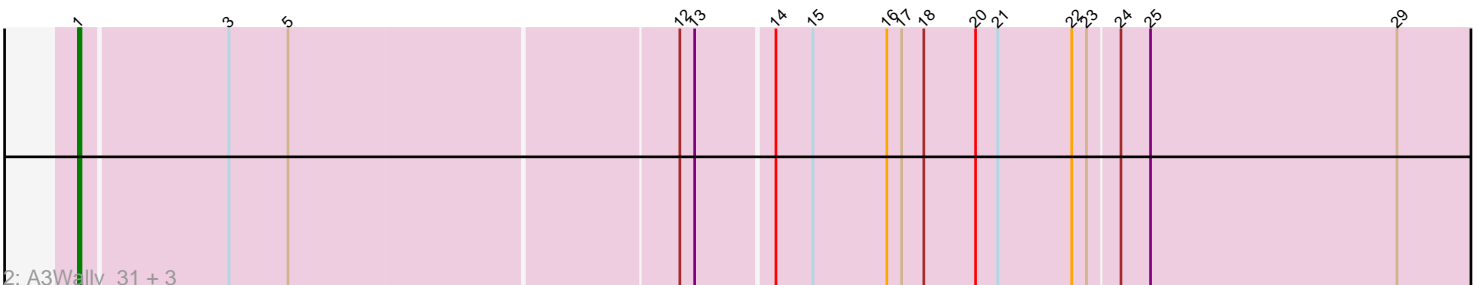
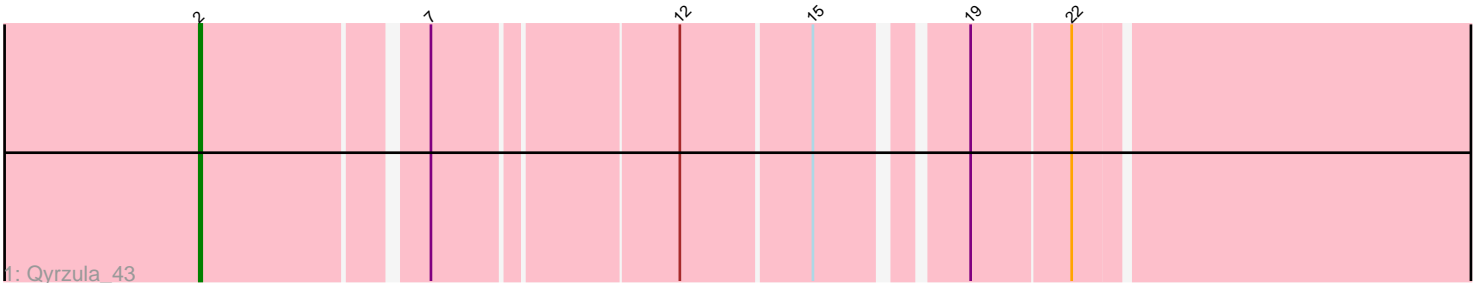


Pham 281091



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 281091 Report

This analysis was run 02/07/26 on database version 634.

Pham number 281091 has 11 members, 2 are drafts.

Phages represented in each track:

- Track 1 : Qyrzula_43
- Track 2 : A3Wally_31, Dodo_382, A3Wally_384, Dodo_32
- Track 3 : PauloDiaboli_31, PauloDiaboli_386
- Track 4 : Cece_335, Cece_33
- Track 5 : Pumpernickel_341, Pumpernickel_40

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

The start number called the most often in the published annotations is 1, it was called in 8 of the 9 non-draft genes in the pham.

Genes that call this "Most Annotated" start:

- A3Wally_31, A3Wally_384, Cece_33, Cece_335, Dodo_32, Dodo_382, PauloDiaboli_31, PauloDiaboli_386, Pumpernickel_341, Pumpernickel_40,

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

-

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

- Qyrzula_43,

Summary by start number:

Start 1:

- Found in 10 of 11 (90.9%) of genes in pham
- Manual Annotations of this start: 8 of 9
- Called 100.0% of time when present
- Phage (with cluster) where this start called: A3Wally_31 (GD1), A3Wally_384 (GD1), Cece_33 (GD3), Cece_335 (GD3), Dodo_32 (GD1), Dodo_382 (GD1), PauloDiaboli_31 (GD1), PauloDiaboli_386 (GD1), Pumpernickel_341 (GD4), Pumpernickel_40 (GD4),

Start 2:

- Found in 1 of 11 (9.1%) of genes in pham

- Manual Annotations of this start: 1 of 9
- Called 100.0% of time when present
- Phage (with cluster) where this start called: Qyrzula_43 (B2),

Summary by clusters:

There are 4 clusters represented in this pham: GD1, GD3, GD4, B2,

Info for manual annotations of cluster B2:

- Start number 2 was manually annotated 1 time for cluster B2.

Info for manual annotations of cluster GD1:

- Start number 1 was manually annotated 4 times for cluster GD1.

Info for manual annotations of cluster GD3:

- Start number 1 was manually annotated 2 times for cluster GD3.

Info for manual annotations of cluster GD4:

- Start number 1 was manually annotated 2 times for cluster GD4.

Gene Information:

Gene: A3Wally_31 Start: 10518, Stop: 11093, Start Num: 1

Candidate Starts for A3Wally_31:

(Start: 1 @10518 has 8 MA's), (3, 10575), (5, 10599), (12, 10752), (13, 10758), (14, 10788), (15, 10803), (16, 10833), (17, 10839), (18, 10848), (20, 10869), (21, 10878), (22, 10908), (23, 10914), (24, 10926), (25, 10938), (29, 11037),

Gene: A3Wally_384 Start: 189739, Stop: 190314, Start Num: 1

Candidate Starts for A3Wally_384:

(Start: 1 @189739 has 8 MA's), (3, 189796), (5, 189820), (12, 189973), (13, 189979), (14, 190009), (15, 190024), (16, 190054), (17, 190060), (18, 190069), (20, 190090), (21, 190099), (22, 190129), (23, 190135), (24, 190147), (25, 190159), (29, 190258),

Gene: Cece_335 Start: 181006, Stop: 181578, Start Num: 1

Candidate Starts for Cece_335:

(Start: 1 @181006 has 8 MA's), (4, 181066), (7, 181141), (10, 181219), (11, 181225), (12, 181240), (17, 181327), (26, 181435), (28, 181510),

Gene: Cece_33 Start: 12572, Stop: 13144, Start Num: 1

Candidate Starts for Cece_33:

(Start: 1 @12572 has 8 MA's), (4, 12632), (7, 12707), (10, 12785), (11, 12791), (12, 12806), (17, 12893), (26, 13001), (28, 13076),

Gene: Dodo_382 Start: 188565, Stop: 189140, Start Num: 1

Candidate Starts for Dodo_382:

(Start: 1 @188565 has 8 MA's), (3, 188622), (5, 188646), (12, 188799), (13, 188805), (14, 188835), (15, 188850), (16, 188880), (17, 188886), (18, 188895), (20, 188916), (21, 188925), (22, 188955), (23, 188961), (24, 188973), (25, 188985), (29, 189084),

Gene: Dodo_32 Start: 10365, Stop: 10940, Start Num: 1

Candidate Starts for Dodo_32:

(Start: 1 @10365 has 8 MA's), (3, 10422), (5, 10446), (12, 10599), (13, 10605), (14, 10635), (15, 10650), (16, 10680), (17, 10686), (18, 10695), (20, 10716), (21, 10725), (22, 10755), (23, 10761), (24, 10773), (25, 10785), (29, 10884),

Gene: PauloDiaboli_31 Start: 10358, Stop: 10933, Start Num: 1

Candidate Starts for PauloDiaboli_31:

(Start: 1 @10358 has 8 MA's), (3, 10415), (5, 10439), (12, 10592), (13, 10598), (14, 10628), (15, 10643), (17, 10679), (18, 10688), (20, 10709), (21, 10718), (22, 10748), (23, 10754), (24, 10766), (25, 10778),

Gene: PauloDiaboli_386 Start: 186987, Stop: 187562, Start Num: 1

Candidate Starts for PauloDiaboli_386:

(Start: 1 @186987 has 8 MA's), (3, 187044), (5, 187068), (12, 187221), (13, 187227), (14, 187257), (15, 187272), (17, 187308), (18, 187317), (20, 187338), (21, 187347), (22, 187377), (23, 187383), (24, 187395), (25, 187407),

Gene: Pumpernickel_341 Start: 180990, Stop: 181568, Start Num: 1

Candidate Starts for Pumpernickel_341:

(Start: 1 @180990 has 8 MA's), (6, 181086), (8, 181182), (9, 181194), (12, 181227), (27, 181443),

Gene: Pumpernickel_40 Start: 14858, Stop: 15436, Start Num: 1

Candidate Starts for Pumpernickel_40:

(Start: 1 @14858 has 8 MA's), (6, 14954), (8, 15050), (9, 15062), (12, 15095), (27, 15311),

Gene: Qyrzula_43 Start: 40579, Stop: 40082, Start Num: 2

Candidate Starts for Qyrzula_43:

(Start: 2 @40579 has 1 MA's), (7, 40495), (12, 40402), (15, 40351), (19, 40300), (22, 40261),