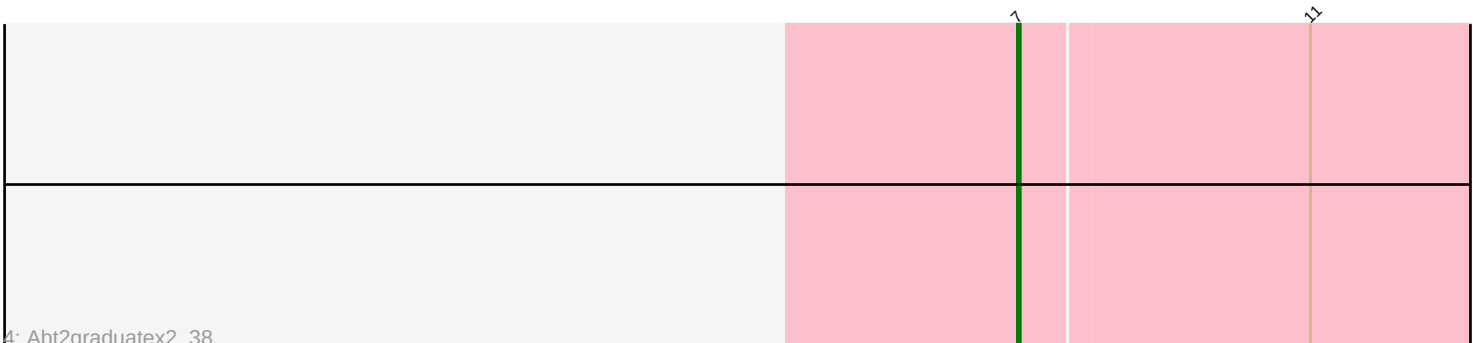
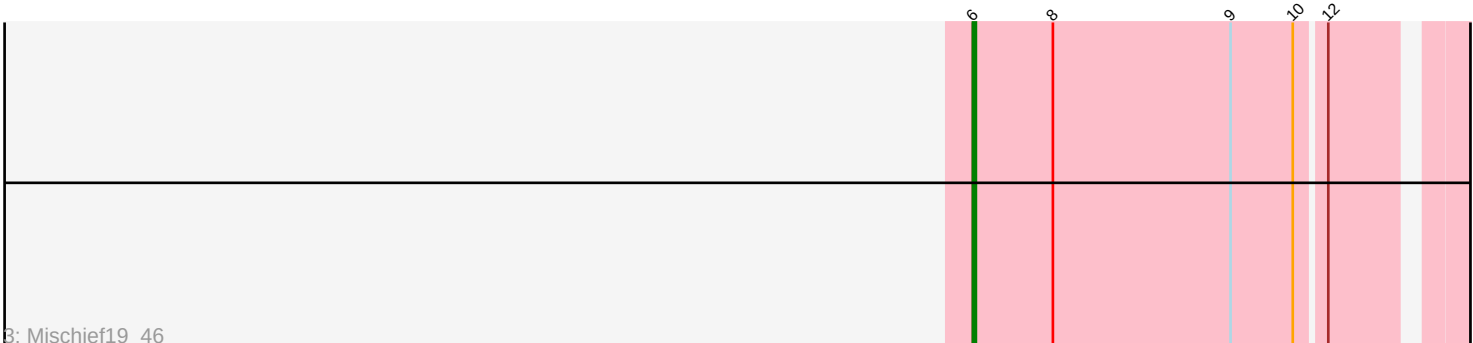
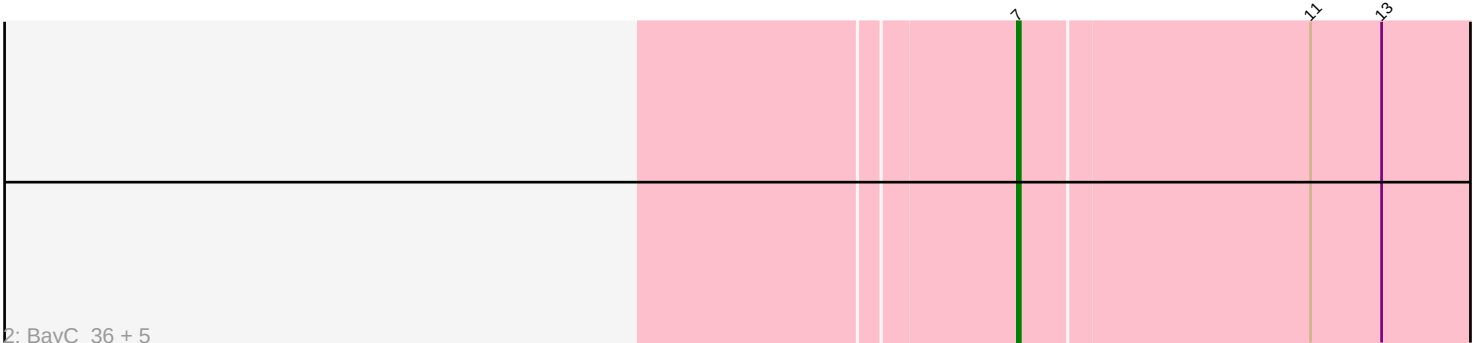




Note: Tracks are now grouped by subcluster and scaled. Switching in subcluster is indicated by changes in track color. Track scale is now set by default to display the region 30 bp upstream of start 1 to 30 bp downstream of the last possible start. If this default region is judged to be packed too tightly 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 309852 Report

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

Pham number 309852 has 9 members, 0 are drafts.

Phages represented in each track:

- Track 1 : RainyPolka_38
- Track 2 : BayC_36, Salete_36, Asis_36, BabyGotBac_37, Maih_35, TP1604_36
- Track 3 : Mischief19_46
- Track 4 : Abt2graduatex2_38

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

The start number called the most often in the published annotations is 7, it was called in 8 of the 9 non-draft genes in the pham.

Genes that call this "Most Annotated" start:

- Abt2graduatex2_38, Asis_36, BabyGotBac_37, BayC_36, Maih_35, RainyPolka_38, Salete_36, TP1604_36,

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

-

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

- Mischief19_46,

Summary by start number:

Start 6:

- Found in 1 of 9 (11.1%) 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: Mischief19_46 (BG),

Start 7:

- Found in 8 of 9 (88.9%) 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: Abt2graduatex2_38 (BG), Asis_36 (BG), BabyGotBac_37 (BG), BayC_36 (BG), Maih_35 (BG), RainyPolka_38 (BG),

Salete_36 (BG), TP1604_36 (BG),

Summary by clusters:

There is one cluster represented in this pham: BG

Info for manual annotations of cluster BG:

- Start number 6 was manually annotated 1 time for cluster BG.
- Start number 7 was manually annotated 8 times for cluster BG.

Gene Information:

Gene: Abt2graduatex2_38 Start: 30914, Stop: 30690, Start Num: 7

Candidate Starts for Abt2graduatex2_38:

(Start: 7 @30914 has 8 MA's), (11, 30818),

Gene: Asis_36 Start: 30112, Stop: 29876, Start Num: 7

Candidate Starts for Asis_36:

(Start: 7 @30112 has 8 MA's), (11, 30016), (13, 29992),

Gene: BabyGotBac_37 Start: 30112, Stop: 29876, Start Num: 7

Candidate Starts for BabyGotBac_37:

(Start: 7 @30112 has 8 MA's), (11, 30016), (13, 29992),

Gene: BayC_36 Start: 30112, Stop: 29876, Start Num: 7

Candidate Starts for BayC_36:

(Start: 7 @30112 has 8 MA's), (11, 30016), (13, 29992),

Gene: Maih_35 Start: 30111, Stop: 29875, Start Num: 7

Candidate Starts for Maih_35:

(Start: 7 @30111 has 8 MA's), (11, 30015), (13, 29991),

Gene: Mischief19_46 Start: 41935, Stop: 41714, Start Num: 6

Candidate Starts for Mischief19_46:

(Start: 6 @41935 has 1 MA's), (8, 41908), (9, 41848), (10, 41827), (12, 41818),

Gene: RainyPolka_38 Start: 30700, Stop: 30473, Start Num: 7

Candidate Starts for RainyPolka_38:

(1, 31036), (2, 31027), (3, 31018), (4, 30916), (5, 30883), (Start: 7 @30700 has 8 MA's), (11, 30604),

Gene: Salete_36 Start: 30112, Stop: 29876, Start Num: 7

Candidate Starts for Salete_36:

(Start: 7 @30112 has 8 MA's), (11, 30016), (13, 29992),

Gene: TP1604_36 Start: 30112, Stop: 29876, Start Num: 7

Candidate Starts for TP1604_36:

(Start: 7 @30112 has 8 MA's), (11, 30016), (13, 29992),