

| 4: RosaAsantewaa_27 | 7 |  |  |  |
|---------------------|---|--|--|--|

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 221984 Report

This analysis was run 03/28/25 on database version 593.

Pham number 221984 has 11 members, 0 are drafts.

Phages represented in each track: • Track 1 : Moozy\_29, Doxi13\_29, Annihilus\_29, PherryCruz\_28, GoblinVoyage\_29, HotFries\_27, SheRa\_28, RavenPuff\_28 • Track 2 : Scap1\_28 • Track 3 : Bilo\_28 • Track 4 : RosaAsantewaa 27

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

The start number called the most often in the published annotations is 4, it was called in 10 of the 11 non-draft genes in the pham.

Genes that call this "Most Annotated" start: • Annihilus\_29, Bilo\_28, Doxi13\_29, GoblinVoyage\_29, HotFries\_27, Moozy\_29, PherryCruz\_28, RavenPuff\_28, RosaAsantewaa\_27, SheRa\_28,

Genes that have the "Most Annotated" start but do not call it: • Scap1\_28,

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

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## Summary by start number:

Start 1:

- Found in 2 of 11 (18.2%) of genes in pham
- Manual Annotations of this start: 1 of 11
- Called 50.0% of time when present
- Phage (with cluster) where this start called: Scap1\_28 (BI2),

### Start 4:

- Found in 11 of 11 (100.0%) of genes in pham
- Manual Annotations of this start: 10 of 11
- Called 90.9% of time when present

• Phage (with cluster) where this start called: Annihilus\_29 (Bl2), Bilo\_28 (Bl2), Doxi13\_29 (Bl2), GoblinVoyage\_29 (Bl2), HotFries\_27 (Bl2), Moozy\_29 (Bl2), PherryCruz\_28 (Bl2), RavenPuff\_28 (Bl2), RosaAsantewaa\_27 (Bl2), SheRa\_28 (Bl2),

### Summary by clusters:

There is one cluster represented in this pham: BI2

Info for manual annotations of cluster BI2:Start number 1 was manually annotated 1 time for cluster BI2.Start number 4 was manually annotated 10 times for cluster BI2.

#### Gene Information:

Gene: Annihilus\_29 Start: 23803, Stop: 24042, Start Num: 4 Candidate Starts for Annihilus\_29: (2, 23773), (3, 23776), (Start: 4 @23803 has 10 MA's), (5, 23833), (6, 23839), (10, 23935), (11, 23965),

Gene: Bilo\_28 Start: 24017, Stop: 24256, Start Num: 4 Candidate Starts for Bilo\_28: (Start: 1 @23960 has 1 MA's), (3, 23990), (Start: 4 @24017 has 10 MA's), (5, 24047), (6, 24053), (10, 24149), (11, 24179),

Gene: Doxi13\_29 Start: 23766, Stop: 24005, Start Num: 4 Candidate Starts for Doxi13\_29: (2, 23736), (3, 23739), (Start: 4 @23766 has 10 MA's), (5, 23796), (6, 23802), (10, 23898), (11, 23928),

Gene: GoblinVoyage\_29 Start: 23766, Stop: 24005, Start Num: 4 Candidate Starts for GoblinVoyage\_29: (2, 23736), (3, 23739), (Start: 4 @23766 has 10 MA's), (5, 23796), (6, 23802), (10, 23898), (11, 23928),

Gene: HotFries\_27 Start: 23496, Stop: 23735, Start Num: 4 Candidate Starts for HotFries\_27: (2, 23466), (3, 23469), (Start: 4 @23496 has 10 MA's), (5, 23526), (6, 23532), (10, 23628), (11, 23658),

Gene: Moozy\_29 Start: 23790, Stop: 24029, Start Num: 4 Candidate Starts for Moozy\_29: (2, 23760), (3, 23763), (Start: 4 @23790 has 10 MA's), (5, 23820), (6, 23826), (10, 23922), (11, 23952),

Gene: PherryCruz\_28 Start: 23765, Stop: 24004, Start Num: 4 Candidate Starts for PherryCruz\_28: (2, 23735), (3, 23738), (Start: 4 @23765 has 10 MA's), (5, 23795), (6, 23801), (10, 23897), (11, 23927),

Gene: RavenPuff\_28 Start: 23765, Stop: 24004, Start Num: 4 Candidate Starts for RavenPuff\_28: (2, 23735), (3, 23738), (Start: 4 @23765 has 10 MA's), (5, 23795), (6, 23801), (10, 23897), (11, 23927),

Gene: RosaAsantewaa\_27 Start: 23272, Stop: 23511, Start Num: 4 Candidate Starts for RosaAsantewaa\_27: (3, 23245), (Start: 4 @23272 has 10 MA's), (6, 23308), (8, 23350), (9, 23353), (10, 23404), (12, 23452), Gene: Scap1\_28 Start: 23389, Stop: 23688, Start Num: 1 Candidate Starts for Scap1\_28: (Start: 1 @23389 has 1 MA's), (2, 23416), (3, 23419), (Start: 4 @23446 has 10 MA's), (5, 23476), (7, 23485), (10, 23581), (11, 23611),

Gene: SheRa\_28 Start: 23497, Stop: 23736, Start Num: 4 Candidate Starts for SheRa\_28: (2, 23467), (3, 23470), (Start: 4 @23497 has 10 MA's), (5, 23527), (6, 23533), (10, 23629), (11, 23659),