

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 197095 Report

This analysis was run 12/09/24 on database version 580.

Pham number 197095 has 9 members, 3 are drafts.

Phages represented in each track:

Track 1 : Adaia\_20

Track 2: Coriander 41, TaronosaurasRx 43, Doggs 38, Gusanita 41

Track 3 : Dmitri\_41

Track 4 : Opie\_44Track 5 : Phayonce\_34

Track 6 : CyranoPS\_19

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

The start number called the most often in the published annotations is 5, it was called in 4 of the 6 non-draft genes in the pham.

Genes that call this "Most Annotated" start:

• Coriander\_41, Dmitri\_41, Doggs\_38, Gusanita\_41, Opie\_44, TaronosaurasRx\_43,

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

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

Adaia\_20, CyranoPS\_19, Phayonce\_34,

## Summary by start number:

### Start 2:

- Found in 1 of 9 (11.1%) of genes in pham
- Manual Annotations of this start: 1 of 6
- Called 100.0% of time when present
- Phage (with cluster) where this start called: Adaia\_20 (AX),

#### Start 3:

- Found in 1 of 9 (11.1%) of genes in pham
- No Manual Annotations of this start.
- Called 100.0% of time when present
- Phage (with cluster) where this start called: CyranoPS 19 (singleton).

### Start 4:

- Found in 1 of 9 (11.1%) of genes in pham
- Manual Annotations of this start: 1 of 6
- Called 100.0% of time when present
- Phage (with cluster) where this start called: Phayonce\_34 (P5),

## Start 5:

- Found in 6 of 9 (66.7%) of genes in pham
- Manual Annotations of this start: 4 of 6
- Called 100.0% of time when present
- Phage (with cluster) where this start called: Coriander\_41 (DB), Dmitri\_41 (DB), Doggs\_38 (DB), Gusanita\_41 (FF), Opie\_44 (DB), TaronosaurasRx\_43 (DB),

## Summary by clusters:

There are 5 clusters represented in this pham: AX, singleton, DB, FF, P5,

Info for manual annotations of cluster AX:

•Start number 2 was manually annotated 1 time for cluster AX.

Info for manual annotations of cluster DB:

•Start number 5 was manually annotated 3 times for cluster DB.

Info for manual annotations of cluster FF:

•Start number 5 was manually annotated 1 time for cluster FF.

Info for manual annotations of cluster P5:

•Start number 4 was manually annotated 1 time for cluster P5.

### Gene Information:

Gene: Adaia\_20 Start: 13289, Stop: 13507, Start Num: 2

Candidate Starts for Adaia\_20:

(Start: 2 @13289 has 1 MA's), (8, 13343), (9, 13361),

Gene: Coriander 41 Start: 31926, Stop: 32126, Start Num: 5

Candidate Starts for Coriander\_41: (Start: 5 @31926 has 4 MA's),

Gene: CyranoPS 19 Start: 12813, Stop: 13016, Start Num: 3

Candidate Starts for CyranoPS\_19:

(3, 12813), (6, 12831), (7, 12843), (10, 12906), (12, 12933), (13, 12939), (14, 12948),

Gene: Dmitri\_41 Start: 33516, Stop: 33710, Start Num: 5

Candidate Starts for Dmitri 41:

(Start: 5 @ 33516 has 4 MA's), (15, 33669),

Gene: Doggs\_38 Start: 32655, Stop: 32855, Start Num: 5

Candidate Starts for Doggs\_38: (Start: 5 @32655 has 4 MA's),

Gene: Gusanita\_41 Start: 30644, Stop: 30853, Start Num: 5

Candidate Starts for Gusanita\_41: (Start: 5 @30644 has 4 MA's),

Gene: Opie\_44 Start: 33313, Stop: 33507, Start Num: 5

Candidate Starts for Opie\_44:

(Start: 5 @33313 has 4 MA's), (8, 33340), (16, 33484),

Gene: Phayonce\_34 Start: 29517, Stop: 29735, Start Num: 4

Candidate Starts for Phayonce\_34:

(1, 29487), (Start: 4 @ 29517 has 1 MA's), (11, 29625),

Gene: TaronosaurasRx\_43 Start: 31715, Stop: 31909, Start Num: 5

Candidate Starts for TaronosaurasRx\_43:

(Start: 5 @31715 has 4 MA's),