



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 302071 Report

This analysis was run 06/08/26 on database version 649.

Pham number 302071 has 8 members, 7 are drafts.

Phages represented in each track:

- Track 1 : Rowdy\_1, DustyMartin\_1, Shapes\_1
- Track 2 : LogiBear\_4
- Track 3 : Caraxes\_4
- Track 4 : Candra\_4
- Track 5 : Garak\_4
- Track 6 : Partridge\_2

### ***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 1 of the 1 non-draft genes in the pham.

Genes that call this "Most Annotated" start:

- Candra\_4, Caraxes\_4, DustyMartin\_1, LogiBear\_4, Partridge\_2, Rowdy\_1, Shapes\_1,

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

- Garak\_4,

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

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### **Summary by start number:**

Start 4:

- Found in 8 of 8 ( 100.0% ) of genes in pham
- Manual Annotations of this start: 1 of 1
- Called 87.5% of time when present
- Phage (with cluster) where this start called: Candra\_4 (A6), Caraxes\_4 (A2), DustyMartin\_1 (A10), LogiBear\_4 (A15), Partridge\_2 (CA), Rowdy\_1 (A10), Shapes\_1 (A10),

Start 5:

- Found in 8 of 8 ( 100.0% ) of genes in pham

- No Manual Annotations of this start.
- Called 12.5% of time when present
- Phage (with cluster) where this start called: Garak\_4 (A6),

### **Summary by clusters:**

There are 5 clusters represented in this pham: A15, CA, A2, A10, A6,

Info for manual annotations of cluster CA:

- Start number 4 was manually annotated 1 time for cluster CA.

### **Gene Information:**

Gene: Candra\_4 Start: 2340, Stop: 2191, Start Num: 4

Candidate Starts for Candra\_4:

(Start: 4 @2340 has 1 MA's), (5, 2319),

Gene: Caraxes\_4 Start: 2318, Stop: 2187, Start Num: 4

Candidate Starts for Caraxes\_4:

(Start: 4 @2318 has 1 MA's), (5, 2297),

Gene: DustyMartin\_1 Start: 635, Stop: 486, Start Num: 4

Candidate Starts for DustyMartin\_1:

(2, 674), (Start: 4 @635 has 1 MA's), (5, 614),

Gene: Garak\_4 Start: 2286, Stop: 2158, Start Num: 5

Candidate Starts for Garak\_4:

(Start: 4 @2307 has 1 MA's), (5, 2286),

Gene: LogiBear\_4 Start: 2500, Stop: 2369, Start Num: 4

Candidate Starts for LogiBear\_4:

(2, 2539), (Start: 4 @2500 has 1 MA's), (5, 2479), (6, 2395),

Gene: Partridge\_2 Start: 841, Stop: 713, Start Num: 4

Candidate Starts for Partridge\_2:

(1, 907), (2, 880), (3, 874), (Start: 4 @841 has 1 MA's), (5, 820),

Gene: Rowdy\_1 Start: 635, Stop: 486, Start Num: 4

Candidate Starts for Rowdy\_1:

(2, 674), (Start: 4 @635 has 1 MA's), (5, 614),

Gene: Shapes\_1 Start: 635, Stop: 486, Start Num: 4

Candidate Starts for Shapes\_1:

(2, 674), (Start: 4 @635 has 1 MA's), (5, 614),