



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 203937 Report

This analysis was run 01/18/25 on database version 583.

Pham number 203937 has 3 members, 1 are drafts.

Phages represented in each track:

Track 1 : MCubed\_55, Phanita\_54

Track 2 : Zenitsu\_55

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

The start number called the most often in the published annotations is 2, it was called in 1 of the 2 non-draft genes in the pham.

Genes that call this "Most Annotated" start:

• Zenitsu 55.

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

MCubed 55, Phanita 54,

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

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

#### Start 1:

- Found in 3 of 3 ( 100.0% ) of genes in pham
- Manual Annotations of this start: 1 of 2
- Called 66.7% of time when present
- Phage (with cluster) where this start called: MCubed\_55 (EA2), Phanita\_54 (EA2),

#### Start 2:

- Found in 3 of 3 (100.0%) of genes in pham
- Manual Annotations of this start: 1 of 2
- Called 33.3% of time when present
- Phage (with cluster) where this start called: Zenitsu\_55 (EA2),

## Summary by clusters:

There is one cluster represented in this pham: EA2

Info for manual annotations of cluster EA2:

- •Start number 1 was manually annotated 1 time for cluster EA2.
- •Start number 2 was manually annotated 1 time for cluster EA2.

## Gene Information:

Gene: MCubed\_55 Start: 36889, Stop: 36686, Start Num: 1

Candidate Starts for MCubed 55:

(Start: 1 @ 36889 has 1 MA's), (Start: 2 @ 36886 has 1 MA's), (3, 36790), (4, 36781), (5, 36754),

Gene: Phanita\_54 Start: 36801, Stop: 36598, Start Num: 1

Candidate Starts for Phanita\_54:

(Start: 1 @ 36801 has 1 MA's), (Start: 2 @ 36798 has 1 MA's), (3, 36702), (4, 36693), (5, 36666),

Gene: Zenitsu\_55 Start: 36891, Stop: 36691, Start Num: 2

Candidate Starts for Zenitsu\_55:

(Start: 1 @ 36894 has 1 MA's), (Start: 2 @ 36891 has 1 MA's), (3, 36795), (4, 36786), (5, 36759),