



1: Neobush\_33 + 10

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 5240 Report

This analysis was run 07/09/24 on database version 566.

Pham number 5240 has 11 members, 0 are drafts.

Phages represented in each track:

- Track 1 : Neobush\_33, Maridalia\_32, Eudoria\_33, Kita\_33, Antonio\_33, Polly\_30, Manasvini\_33, Zameen\_33, Suscepit\_33, Trumpet\_33, Tayonia\_33

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

The start number called the most often in the published annotations is 1, it was called in 11 of the 11 non-draft genes in the pham.

Genes that call this "Most Annotated" start:

- Antonio\_33, Eudoria\_33, Kita\_33, Manasvini\_33, Maridalia\_32, Neobush\_33, Polly\_30, Suscepit\_33, Tayonia\_33, Trumpet\_33, Zameen\_33,

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

- 

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

- 

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

Start 1:

- Found in 11 of 11 ( 100.0% ) of genes in pham
- Manual Annotations of this start: 11 of 11
- Called 100.0% of time when present
- Phage (with cluster) where this start called: Antonio\_33 (CZ1), Eudoria\_33 (CZ1), Kita\_33 (CZ1), Manasvini\_33 (CZ1), Maridalia\_32 (CZ1), Neobush\_33 (CZ1), Polly\_30 (CZ1), Suscepit\_33 (CZ1), Tayonia\_33 (CZ1), Trumpet\_33 (CZ1), Zameen\_33 (CZ1),

### **Summary by clusters:**

There is one cluster represented in this pham: CZ1

Info for manual annotations of cluster CZ1:

- Start number 1 was manually annotated 11 times for cluster CZ1.

**Gene Information:**

Gene: Antonio\_33 Start: 28264, Stop: 28623, Start Num: 1

Candidate Starts for Antonio\_33:

(Start: 1 @28264 has 11 MA's), (2, 28447), (3, 28525), (4, 28528), (5, 28561),

Gene: Eudoria\_33 Start: 28264, Stop: 28623, Start Num: 1

Candidate Starts for Eudoria\_33:

(Start: 1 @28264 has 11 MA's), (2, 28447), (3, 28525), (4, 28528), (5, 28561),

Gene: Kita\_33 Start: 28273, Stop: 28632, Start Num: 1

Candidate Starts for Kita\_33:

(Start: 1 @28273 has 11 MA's), (2, 28456), (3, 28534), (4, 28537), (5, 28570),

Gene: Manasvini\_33 Start: 28264, Stop: 28623, Start Num: 1

Candidate Starts for Manasvini\_33:

(Start: 1 @28264 has 11 MA's), (2, 28447), (3, 28525), (4, 28528), (5, 28561),

Gene: Maridalia\_32 Start: 28264, Stop: 28623, Start Num: 1

Candidate Starts for Maridalia\_32:

(Start: 1 @28264 has 11 MA's), (2, 28447), (3, 28525), (4, 28528), (5, 28561),

Gene: Neobush\_33 Start: 28264, Stop: 28623, Start Num: 1

Candidate Starts for Neobush\_33:

(Start: 1 @28264 has 11 MA's), (2, 28447), (3, 28525), (4, 28528), (5, 28561),

Gene: Polly\_30 Start: 27707, Stop: 28066, Start Num: 1

Candidate Starts for Polly\_30:

(Start: 1 @27707 has 11 MA's), (2, 27890), (3, 27968), (4, 27971), (5, 28004),

Gene: Suscepit\_33 Start: 28265, Stop: 28624, Start Num: 1

Candidate Starts for Suscepit\_33:

(Start: 1 @28265 has 11 MA's), (2, 28448), (3, 28526), (4, 28529), (5, 28562),

Gene: Tayonia\_33 Start: 28264, Stop: 28623, Start Num: 1

Candidate Starts for Tayonia\_33:

(Start: 1 @28264 has 11 MA's), (2, 28447), (3, 28525), (4, 28528), (5, 28561),

Gene: Trumpet\_33 Start: 28265, Stop: 28624, Start Num: 1

Candidate Starts for Trumpet\_33:

(Start: 1 @28265 has 11 MA's), (2, 28448), (3, 28526), (4, 28529), (5, 28562),

Gene: Zameen\_33 Start: 28264, Stop: 28623, Start Num: 1

Candidate Starts for Zameen\_33:

(Start: 1 @28264 has 11 MA's), (2, 28447), (3, 28525), (4, 28528), (5, 28561),