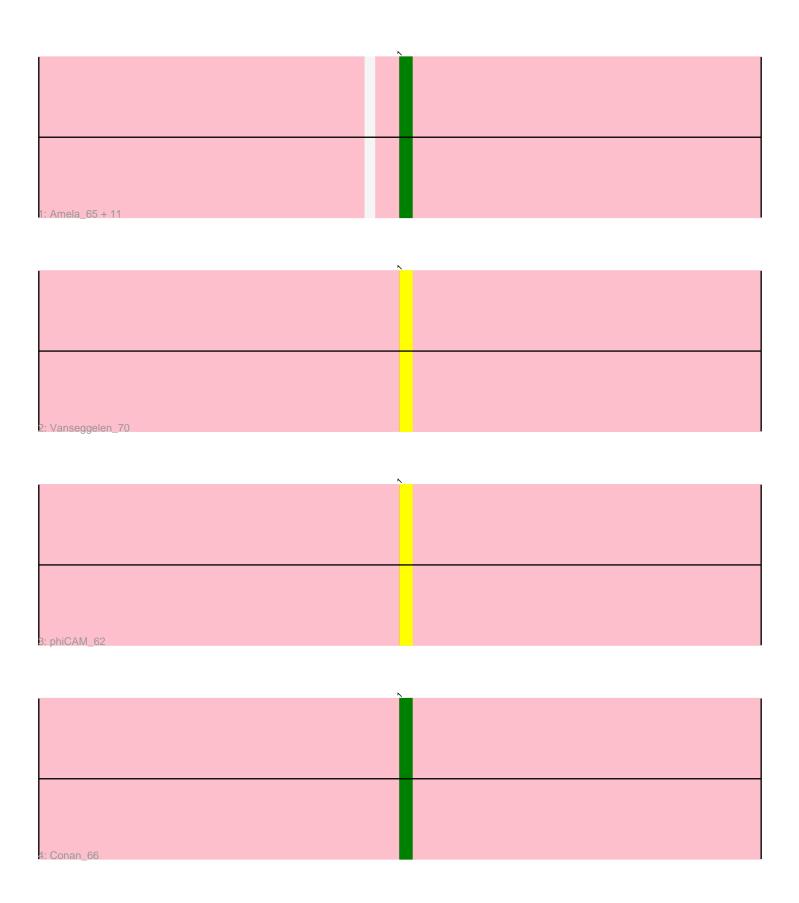
# Pham 203427



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 203427 Report

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

Pham number 203427 has 15 members, 4 are drafts.

Phages represented in each track:

• Track 1 : Amela\_65, Dexers\_63, ElGato\_66, Provolone\_66, Speedwell\_68, Celery\_68, Pavo\_66, Saftant\_63, Verse\_66, Kaine\_65, Verabelle\_68, Alsaber\_65

Track 2 : Vanseggelen\_70

Track 3 : phiCAM\_62Track 4 : Conan 66

### 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:

• Alsaber\_65, Amela\_65, Celery\_68, Conan\_66, Dexers\_63, ElGato\_66, Kaine\_65, Pavo\_66, Provolone\_66, Saftant\_63, Speedwell\_68, Vanseggelen\_70, Verabelle\_68, Verse\_66, phiCAM\_62,

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 15 of 15 (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: Alsaber\_65 (BD3), Amela\_65 (BD3), Celery\_68 (BD3), Conan\_66 (BD3), Dexers\_63 (BD3), ElGato\_66 (BD3), Kaine\_65 (BD3), Pavo\_66 (BD3), Provolone\_66 (BD3), Saftant\_63 (BD3), Speedwell\_68 (BD3), Vanseggelen\_70 (BD3), Verabelle\_68 (BD3), Verse\_66 (BD3), phiCAM\_62 (BD3),

#### Summary by clusters:

There is one cluster represented in this pham: BD3

Info for manual annotations of cluster BD3:

•Start number 1 was manually annotated 11 times for cluster BD3.

#### Gene Information:

Gene: Alsaber\_65 Start: 43432, Stop: 43151, Start Num: 1

Candidate Starts for Alsaber\_65: (Start: 1 @43432 has 11 MA's),

Gene: Amela\_65 Start: 44414, Stop: 44145, Start Num: 1

Candidate Starts for Amela\_65: (Start: 1 @44414 has 11 MA's),

Gene: Celery\_68 Start: 43343, Stop: 43065, Start Num: 1

Candidate Starts for Celery\_68: (Start: 1 @43343 has 11 MA's),

Gene: Conan\_66 Start: 43683, Stop: 43417, Start Num: 1

Candidate Starts for Conan\_66: (Start: 1 @43683 has 11 MA's),

Gene: Dexers\_63 Start: 43658, Stop: 43389, Start Num: 1

Candidate Starts for Dexers\_63: (Start: 1 @43658 has 11 MA's),

Gene: ElGato\_66 Start: 43561, Stop: 43289, Start Num: 1

Candidate Starts for ElGato\_66: (Start: 1 @43561 has 11 MA's),

Gene: Kaine 65 Start: 43456, Stop: 43175, Start Num: 1

Candidate Starts for Kaine\_65: (Start: 1 @43456 has 11 MA's),

Gene: Pavo 66 Start: 43632, Stop: 43360, Start Num: 1

Candidate Starts for Pavo\_66: (Start: 1 @43632 has 11 MA's),

Gene: Provolone 66 Start: 43773, Stop: 43501, Start Num: 1

Candidate Starts for Provolone\_66: (Start: 1 @43773 has 11 MA's),

Gene: Saftant\_63 Start: 43520, Stop: 43239, Start Num: 1

Candidate Starts for Saftant\_63: (Start: 1 @43520 has 11 MA's),

Gene: Speedwell\_68 Start: 44565, Stop: 44284, Start Num: 1

Candidate Starts for Speedwell\_68: (Start: 1 @44565 has 11 MA's),

Gene: Vanseggelen\_70 Start: 43501, Stop: 43232, Start Num: 1

Candidate Starts for Vanseggelen\_70:

(Start: 1 @43501 has 11 MA's),

Gene: Verabelle\_68 Start: 43108, Stop: 42839, Start Num: 1

Candidate Starts for Verabelle\_68: (Start: 1 @43108 has 11 MA's),

Gene: Verse\_66 Start: 44405, Stop: 44136, Start Num: 1

Candidate Starts for Verse\_66: (Start: 1 @44405 has 11 MA's),

Gene: phiCAM\_62 Start: 45181, Stop: 44900, Start Num: 1

Candidate Starts for phiCAM\_62: (Start: 1 @45181 has 11 MA's),