

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 8768 Report

This analysis was run 04/28/24 on database version 559.

Pham number 8768 has 9 members, 1 are drafts.

Phages represented in each track:

- Track 1 : Persimmon_195, PinkiePie_193, Navo_195, NootNoot_190, Squillium_196, Braelyn_191, Paradiddles_187, Liandry_193, Bartholomune_194

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 8 of the 8 non-draft genes in the pham.

Genes that call this "Most Annotated" start:

- Bartholomune_194, Braelyn_191, Liandry_193, Navo_195, NootNoot_190, Paradiddles_187, Persimmon_195, PinkiePie_193, Squillium_196,

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 9 of 9 (100.0%) of genes in pham
- Manual Annotations of this start: 8 of 8
- Called 100.0% of time when present
- Phage (with cluster) where this start called: Bartholomune_194 (BE1), Braelyn_191 (BE1), Liandry_193 (BE1), Navo_195 (BE1), NootNoot_190 (BE1), Paradiddles_187 (BE1), Persimmon_195 (BE1), PinkiePie_193 (BE1), Squillium_196 (BE1),

Summary by clusters:

There is one cluster represented in this pham: BE1

Info for manual annotations of cluster BE1:

- Start number 1 was manually annotated 8 times for cluster BE1.

Gene Information:

Gene: Bartholomune_194 Start: 101361, Stop: 101561, Start Num: 1

Candidate Starts for Bartholomune_194:

(Start: 1 @101361 has 8 MA's), (2, 101424), (3, 101481), (4, 101511),

Gene: Braelyn_191 Start: 101011, Stop: 101211, Start Num: 1

Candidate Starts for Braelyn_191:

(Start: 1 @101011 has 8 MA's), (2, 101074), (3, 101131), (4, 101161),

Gene: Liandry_193 Start: 101680, Stop: 101880, Start Num: 1

Candidate Starts for Liandry_193:

(Start: 1 @101680 has 8 MA's), (2, 101743), (3, 101800), (4, 101830),

Gene: Navo_195 Start: 100215, Stop: 100415, Start Num: 1

Candidate Starts for Navo_195:

(Start: 1 @100215 has 8 MA's), (2, 100278), (3, 100335), (4, 100365),

Gene: NootNoot_190 Start: 100469, Stop: 100669, Start Num: 1

Candidate Starts for NootNoot_190:

(Start: 1 @100469 has 8 MA's), (2, 100532), (3, 100589), (4, 100619),

Gene: Paradiddles_187 Start: 103138, Stop: 103338, Start Num: 1

Candidate Starts for Paradiddles_187:

(Start: 1 @103138 has 8 MA's), (2, 103201), (3, 103258), (4, 103288),

Gene: Persimmon_195 Start: 100587, Stop: 100787, Start Num: 1

Candidate Starts for Persimmon_195:

(Start: 1 @100587 has 8 MA's), (2, 100650), (3, 100707), (4, 100737),

Gene: PinkiePie_193 Start: 101680, Stop: 101880, Start Num: 1

Candidate Starts for PinkiePie_193:

(Start: 1 @101680 has 8 MA's), (2, 101743), (3, 101800), (4, 101830),

Gene: Squillium_196 Start: 101682, Stop: 101882, Start Num: 1

Candidate Starts for Squillium_196:

(Start: 1 @101682 has 8 MA's), (2, 101745), (3, 101802), (4, 101832),