

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 88091 Report

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

Pham number 88091 has 7 members, 0 are drafts.

Phages represented in each track:

Track 1: PrincessTrina_63, Tophat_63, Conboy_62, Chipper1996_630,

EdgarPoe_62

Track 2 : KBurrousTX_63Track 3 : DrYang 61

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

The start number called the most often in the published annotations is 7, it was called in 5 of the 7 non-draft genes in the pham.

Genes that call this "Most Annotated" start:

• Chipper1996_630, Conboy_62, EdgarPoe_62, PrincessTrina_63, Tophat_63,

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

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

DrYang_61, KBurrousTX_63,

Summary by start number:

Start 7:

- Found in 5 of 7 (71.4%) of genes in pham
- Manual Annotations of this start: 5 of 7
- Called 100.0% of time when present
- Phage (with cluster) where this start called: Chipper1996_630 (AR), Conboy_62 (AR), EdgarPoe_62 (AR), PrincessTrina_63 (AR), Tophat_63 (AR),

Start 8:

- Found in 1 of 7 (14.3%) of genes in pham
- Manual Annotations of this start: 1 of 7
- Called 100.0% of time when present
- Phage (with cluster) where this start called: KBurrousTX 63 (AR).

Start 9:

- Found in 1 of 7 (14.3%) of genes in pham
- Manual Annotations of this start: 1 of 7
- Called 100.0% of time when present
- Phage (with cluster) where this start called: DrYang_61 (AR),

Summary by clusters:

There is one cluster represented in this pham: AR

Info for manual annotations of cluster AR:

- •Start number 7 was manually annotated 5 times for cluster AR.
- •Start number 8 was manually annotated 1 time for cluster AR.
- •Start number 9 was manually annotated 1 time for cluster AR.

Gene Information:

Gene: Chipper1996_630 Start: 44753, Stop: 44941, Start Num: 7

Candidate Starts for Chipper1996 630:

(Start: 7 @ 44753 has 5 MA's), (11, 44795), (15, 44879), (16, 44888), (17, 44891), (19, 44900), (20, 44903), (21, 44909), (23, 44918),

Gene: Conboy_62 Start: 44617, Stop: 44805, Start Num: 7

Candidate Starts for Conboy_62:

(Start: 7 @44617 has 5 MA's), (11, 44659), (15, 44743), (16, 44752), (17, 44755), (19, 44764), (20, 44767), (21, 44773), (23, 44782),

Gene: DrYang_61 Start: 46179, Stop: 46376, Start Num: 9

Candidate Starts for DrYang 61:

(1, 46050), (2, 46074), (3, 46098), (5, 46137), (6, 46164), (Start: 9 @ 46179 has 1 MA's), (11, 46227), (12, 46233), (13, 46263), (14, 46278), (18, 46332), (22, 46347), (24, 46359),

Gene: EdgarPoe 62 Start: 44617, Stop: 44805, Start Num: 7

Candidate Starts for EdgarPoe_62:

(Start: 7 @44617 has 5 MA's), (11, 44659), (15, 44743), (16, 44752), (17, 44755), (19, 44764), (20, 44767), (21, 44773), (23, 44782),

Gene: KBurrousTX 63 Start: 46148, Stop: 46339, Start Num: 8

Candidate Starts for KBurrousTX 63:

(4, 46106), (Start: 8 @46148 has 1 MA's), (10, 46154), (11, 46193), (12, 46199), (16, 46286), (17, 46289), (25, 46328),

Gene: PrincessTrina_63 Start: 44821, Stop: 45009, Start Num: 7

Candidate Starts for PrincessTrina_63:

(Start: 7 @44821 has 5 MA's), (11, 44863), (15, 44947), (16, 44956), (17, 44959), (19, 44968), (20, 44971), (21, 44977), (23, 44986),

Gene: Tophat 63 Start: 44754, Stop: 44942, Start Num: 7

Candidate Starts for Tophat 63:

(Start: 7 @ 44754 has 5 MA's), (11, 44796), (15, 44880), (16, 44889), (17, 44892), (19, 44901), (20, 44904), (21, 44910), (23, 44919),