

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 10338 Report

This analysis was run 04/05/24 on database version 557.

Pham number 10338 has 5 members, 0 are drafts.

Phages represented in each track:

Track 1 : BiggityBass_66

Track 2 : Sisko_65Track 3 : AnClar 63

Track 4: LittleMunchkin_68

Track 5 : Yago84_64

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

The start number called the most often in the published annotations is 12, it was called in 4 of the 5 non-draft genes in the pham.

Genes that call this "Most Annotated" start:

AnClar_63, LittleMunchkin_68, Sisko_65, Yago84_64,

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

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

BiggityBass_66,

Summary by start number:

Start 12:

- Found in 4 of 5 (80.0%) of genes in pham
- Manual Annotations of this start: 4 of 5
- Called 100.0% of time when present
- Phage (with cluster) where this start called: AnClar_63 (DR), LittleMunchkin_68 (DR), Sisko_65 (DR), Yago84_64 (DR),

Start 19:

- Found in 1 of 5 (20.0%) of genes in pham
- Manual Annotations of this start: 1 of 5
- Called 100.0% of time when present
- Phage (with cluster) where this start called: BiggityBass 66 (DR).

Summary by clusters:

There is one cluster represented in this pham: DR

Info for manual annotations of cluster DR:

- •Start number 12 was manually annotated 4 times for cluster DR.
- •Start number 19 was manually annotated 1 time for cluster DR.

Gene Information:

Gene: AnClar_63 Start: 53907, Stop: 54446, Start Num: 12

Candidate Starts for AnClar_63:

(1, 53394), (2, 53397), (3, 53412), (4, 53421), (5, 53460), (8, 53604), (9, 53736), (10, 53823), (11, 53835), (Start: 12 @53907 has 4 MA's), (13, 53964), (15, 54012), (16, 54033), (18, 54057), (21, 54129), (22, 54150), (23, 54153), (26, 54321), (27, 54336), (28, 54345), (29, 54381),

Gene: BiggityBass_66 Start: 54527, Stop: 54943, Start Num: 19

Candidate Starts for BiggityBass_66:

(6, 53939), (7, 54038), (14, 54431), (17, 54479), (Start: 19 @54527 has 1 MA's), (20, 54572), (24, 54740), (30, 54911), (31, 54920),

Gene: LittleMunchkin_68 Start: 55523, Stop: 56035, Start Num: 12

Candidate Starts for LittleMunchkin 68:

(8, 55220), (9, 55352), (11, 55451), (Start: 12 @55523 has 4 MA's), (13, 55565), (25, 55850), (26, 55910), (28, 55934), (29, 55970),

Gene: Sisko_65 Start: 52972, Stop: 53511, Start Num: 12

Candidate Starts for Sisko 65:

(11, 52900), (Start: 12 @52972 has 4 MA's), (16, 53098), (18, 53122), (21, 53194), (22, 53215), (23, 53218), (26, 53386), (27, 53401), (28, 53410), (29, 53446),

Gene: Yago84 64 Start: 52364, Stop: 52903, Start Num: 12

Candidate Starts for Yago84_64:

(10, 52280), (11, 52292), (Start: 12 @52364 has 4 MA's), (16, 52490), (18, 52514), (21, 52586), (22, 52607), (23, 52610), (26, 52778), (27, 52793), (28, 52802), (29, 52838),