

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 87787 Report

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

Pham number 87787 has 10 members, 2 are drafts.

Phages represented in each track:

• Track 1 : Dori 60, Mask 60

• Track 2: Mao1 62

Track 3 : Sparky\_75, Farewell\_77

• Track 4 : Tortellini\_49

• Track 5 : Xavia 49

Track 6 : Phayonce\_43

Track 7 : ThuĺaThula\_49

Track 8 : TurkishDelight\_66

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

Genes that call this "Most Annotated" start:

• Dori\_60, Mao1\_62, Mask\_60, Phayonce\_43, ThulaThula\_49, Xavia\_49,

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:

Farewell\_77, Sparky\_75, Tortellini\_49, TurkishDelight\_66,

# Summary by start number:

#### Start 10:

- Found in 2 of 10 (20.0%) of genes in pham
- Manual Annotations of this start: 1 of 8
- Called 100.0% of time when present
- Phage (with cluster) where this start called: Farewell\_77 (AF), Sparky\_75 (AF),

### Start 12:

- Found in 6 of 10 (60.0%) of genes in pham
- Manual Annotations of this start: 5 of 8

- Called 100.0% of time when present
- Phage (with cluster) where this start called: Dori\_60 (AD), Mao1\_62 (AD), Mask\_60 (AD), Phayonce\_43 (P5), ThulaThula\_49 (P5), Xavia\_49 (P3),

#### Start 14:

- Found in 1 of 10 (10.0%) of genes in pham
- Manual Annotations of this start: 1 of 8
- Called 100.0% of time when present
- Phage (with cluster) where this start called: Tortellini\_49 (P2),

### Start 17:

- Found in 1 of 10 (10.0%) of genes in pham
- Manual Annotations of this start: 1 of 8
- Called 100.0% of time when present
- Phage (with cluster) where this start called: TurkishDelight\_66 (singleton),

### Summary by clusters:

There are 6 clusters represented in this pham: P2, P3, singleton, AD, P5, AF,

Info for manual annotations of cluster AD:

•Start number 12 was manually annotated 2 times for cluster AD.

Info for manual annotations of cluster AF:

•Start number 10 was manually annotated 1 time for cluster AF.

Info for manual annotations of cluster P2:

•Start number 14 was manually annotated 1 time for cluster P2.

Info for manual annotations of cluster P3:

•Start number 12 was manually annotated 1 time for cluster P3.

Info for manual annotations of cluster P5:

•Start number 12 was manually annotated 2 times for cluster P5.

### Gene Information:

Gene: Dori\_60 Start: 48301, Stop: 48642, Start Num: 12

Candidate Starts for Dori 60:

(5, 48187), (8, 48223), (9, 48226), (Start: 12 @48301 has 5 MA's), (13, 48316), (18, 48454), (19, 48469), (20, 48475), (21, 48487), (22, 48493),

Gene: Farewell\_77 Start: 50660, Stop: 51028, Start Num: 10

Candidate Starts for Farewell\_77:

(8, 50594), (Start: 10 @50660 has 1 MA's), (11, 50675), (25, 50936), (29, 51017),

Gene: Mao1 62 Start: 46743, Stop: 47087, Start Num: 12

Candidate Starts for Mao1 62:

(1, 46494), (2, 46521), (4, 46614), (Start: 12 @46743 has 5 MA's), (13, 46758), (18, 46896), (20, 46047), (24, 46020), (24, 47042)

46917), (21, 46929), (24, 47013),

Gene: Mask 60 Start: 49044, Stop: 49385, Start Num: 12

Candidate Starts for Mask\_60:

(5, 48930), (8, 48966), (9, 48969), (Start: 12 @49044 has 5 MA's), (13, 49059), (18, 49197), (19, 49212), (20, 49218), (21, 49230), (22, 49236),

Gene: Phayonce 43 Start: 33062, Stop: 33409, Start Num: 12

Candidate Starts for Phayonce\_43:

(Start: 12 @33062 has 5 MA's), (13, 33083), (18, 33221), (20, 33242), (21, 33254),

Gene: Sparky\_75 Start: 52254, Stop: 52622, Start Num: 10

Candidate Starts for Sparky 75:

(8, 52188), (Start: 10 @52254 has 1 MA's), (11, 52269), (25, 52530), (29, 52611),

Gene: ThulaThula\_49 Start: 35441, Stop: 35788, Start Num: 12

Candidate Starts for ThulaThula\_49:

(6, 35333), (Start: 12 @35441 has 5 MA's), (13, 35462), (18, 35600), (20, 35621), (21, 35633),

Gene: Tortellini\_49 Start: 37757, Stop: 38077, Start Num: 14

Candidate Starts for Tortellini 49:

(Start: 14 @37757 has 1 MA's), (15, 37766), (16, 37769), (18, 37889), (19, 37904), (20, 37910), (21, 37922),

Gene: TurkishDelight\_66 Start: 54550, Stop: 54927, Start Num: 17

Candidate Starts for TurkishDelight\_66:

(3, 54289), (7, 54325), (Start: 17 @54550 has 1 MA's), (23, 54673), (26, 54724), (27, 54775), (28, 54787),

Gene: Xavia 49 Start: 37571, Stop: 37912, Start Num: 12

Candidate Starts for Xavia\_49:

(3, 37436), (5, 37457), (8, 37493), (9, 37496), (Start: 12 @37571 has 5 MA's), (13, 37586), (18, 37724), (19, 37739), (20, 37745), (21, 37757),