

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 9637 Report

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

Pham number 9637 has 7 members, 3 are drafts.

Phages represented in each track:

• Track 1 : Finemlucis 127

• Track 2 : Claus\_126

Track 3: MkaliMitinis3 129

Track 4 : Baoshan\_124

Track 5 : ZhongYanYuan\_124

Track 6 : Bazzle\_124Track 7 : Tourach 129

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

The start number called the most often in the published annotations is 9, it was called in 4 of the 4 non-draft genes in the pham.

Genes that call this "Most Annotated" start:

 Baoshan\_124, Bazzle\_124, Claus\_126, Finemlucis\_127, MkaliMitinis3\_129, Tourach\_129, ZhongYanYuan\_124,

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 9:

- Found in 7 of 7 (100.0%) of genes in pham
- Manual Annotations of this start: 4 of 4
- Called 100.0% of time when present
- Phage (with cluster) where this start called: Baoshan\_124 (L2), Bazzle\_124 (L2), Claus\_126 (L2), Finemlucis\_127 (L2), MkaliMitinis3\_129 (L2), Tourach\_129 (L2), ZhongYanYuan\_124 (L2),

## Summary by clusters:

There is one cluster represented in this pham: L2

Info for manual annotations of cluster L2:

•Start number 9 was manually annotated 4 times for cluster L2.

## Gene Information:

Gene: Baoshan 124 Start: 67542, Stop: 67381, Start Num: 9

Candidate Starts for Baoshan 124:

(1, 67770), (4, 67629), (Start: 9 @67542 has 4 MA's), (11, 67482), (12, 67440), (13, 67431), (14, 67419),

Gene: Bazzle 124 Start: 67958, Stop: 67797, Start Num: 9

Candidate Starts for Bazzle 124:

(6, 67982), (8, 67964), (Start: 9 @67958 has 4 MA's), (11, 67898), (13, 67847), (14, 67835),

Gene: Claus\_126 Start: 67618, Stop: 67457, Start Num: 9

Candidate Starts for Claus 126:

(2, 67768), (4, 67705), (Start: 9 @67618 has 4 MA's), (10, 67579), (11, 67558), (13, 67507), (14, 67495),

Gene: Finemlucis\_127 Start: 68809, Stop: 68648, Start Num: 9

Candidate Starts for Finemlucis 127:

(3, 68941), (4, 68896), (8, 68815), (Start: 9 @68809 has 4 MA's), (10, 68770), (11, 68749),

Gene: MkaliMitinis3\_129 Start: 67746, Stop: 67585, Start Num: 9

Candidate Starts for MkaliMitinis3\_129:

(3, 67878), (4, 67833), (6, 67770), (7, 67764), (Start: 9 @67746 has 4 MA's), (11, 67686),

Gene: Tourach 129 Start: 69251, Stop: 69090, Start Num: 9

Candidate Starts for Tourach 129:

(3, 69383), (5, 69326), (6, 69275), (7, 69269), (8, 69257), (Start: 9 @69251 has 4 MA's), (11, 69191),

Gene: ZhongYanYuan\_124 Start: 67241, Stop: 67080, Start Num: 9

Candidate Starts for ZhongYanYuan\_124:

(4, 67328), (Start: 9 @67241 has 4 MA's), (11, 67181), (12, 67139), (13, 67130), (14, 67118),