Kratio Annotation Problems

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| Gene | Problem |
| Gene 2  | This gene is cut into two genes in Mycobacteriophage larva genome. |
| 3 tRNA | Gap 74  |
| Gene 4 | Gap 74 |
| Gene7 | Agrees with genemark |
| Gene 9  | Gap 133 with no match in genbank. No possible ORF. Also is the largest coding potential. There is others SD with highest score but are shorter. |
| Gene 17 | 110 gap from previous gene. |
| Gene 20 | This gene in larva is cutted in two genes. |
| Gene 28 | Agrees with glimmer |
| Gene 29 | Agrees with glimmer |
| Gene 32 | Agrees with genemark |
| Gene 35 | 123 bp gap with no idf possible ORF. The gene is the longest possible ORF. |
| Gene 38 | 117 bp gap |
| Gene 41 | It not appears on glimmer or genemark but it appears in larva genome.  |
| Gene 52 | Too short 168 bp but the other start site overlap the previous gene by 96bp. |
| Gene 54 | it does not appear in Genemark- M. smegmatis |
| Gene 56 | It is short 180 bp but The longest ORF have low SD scor, 570 best score, 364 score longest ORF |
| gene 84 | It is too short 144 bp, but the next start site have a lower SD score and it overlaps the previous gene. |
| Trna 93 | IT does not appear in DNA master but is predicted by Aragorn |
| Gene 95 | Gap 209 same as in larva |
| Gene 102 | Gap 678 same as in larva |