Lehigh decided to delete many original *mycobacterium* phage Dusk gene products. These originally were gps 26,27,62,106,120,122,128. These deletions are consistent with other E cluster mycobacteriophage. We would like the annotation QC team to look closely at the tRNAs (gp 103, 104) bp 62931-62472 and 62477-62548 respectively. There are gaps present but this also seems to be consistent with the E cluster phages. Lehigh feels confident of the rest of this genome annotation.