Accompanying notes for *Rhodococcus* phage Measels

As one of the few *Rhodococcus* phages in the current database, the Measels annotation saw quite a number of genes without a blast hit. However, because Genemark and Glimmer indicated there was coding potential in these areas, most of these no blast hit genes were left in the genome, as it appeared further examinations with a more extensive (and updated) database could reveal similarities unseen this time around. It should be noted that most of the no blast hits genes occurred towards the end of the genome, and it may warrant extra attention. The annotation revealed 10 tRNAs mainly centralized in the middle of the genome, but in a scattered formation. Measels saw many hits to *Rhodococcus* species and *Streptomyces* species, but particularly to *Streptomyces* phage Jay2Jay. By consulting phamerator, we feel confident in our annotation for Measels with the current database available to us.