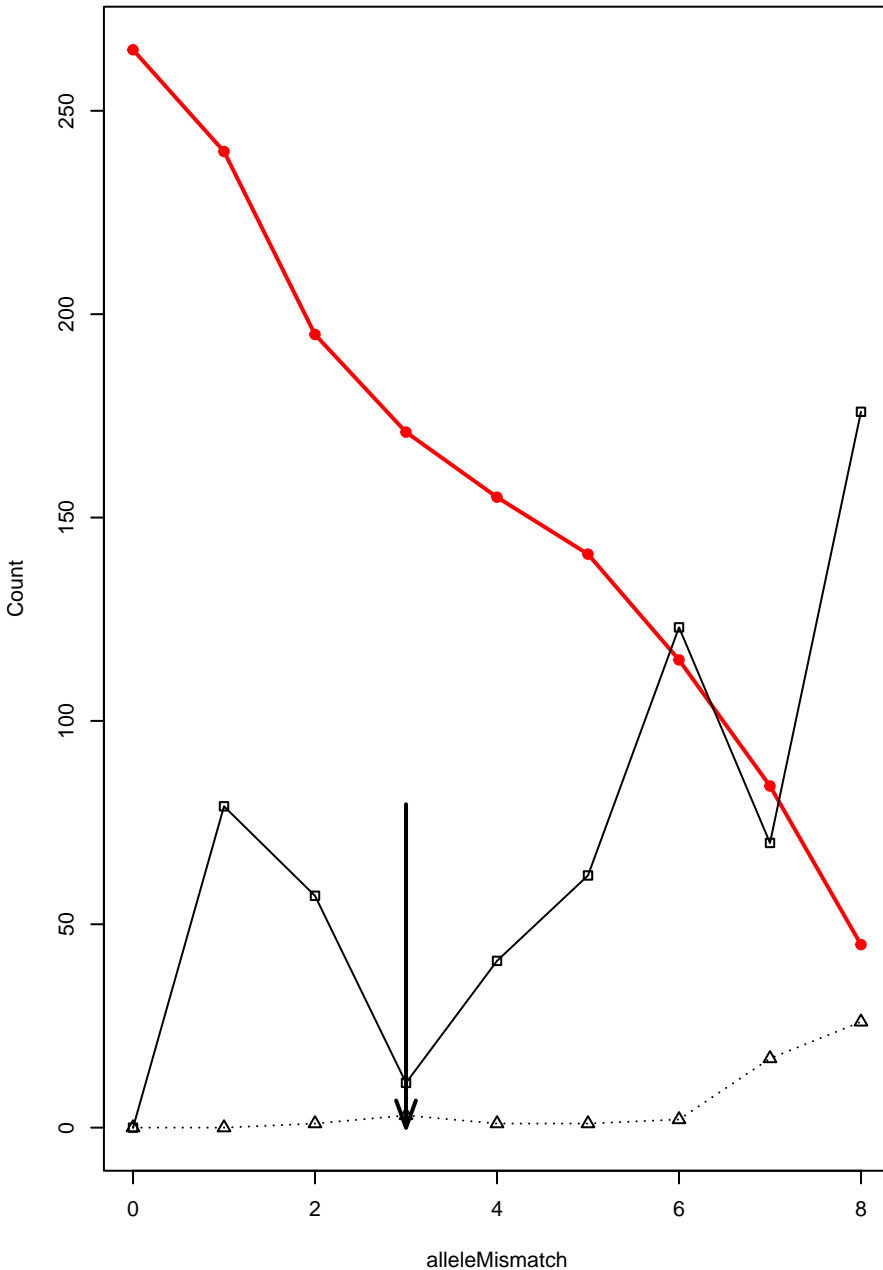


# allelematch

amUniqueProfile()

missingDataLoad=0.095

allelicDiversity=6.1



—●— unique  
—□— multipleMatch  
···△··· unclassified

Best guess for optimum:  
alleleMismatch=3

Profile morphology:  
NonZeroSecondMinimum  
Caution with optimum