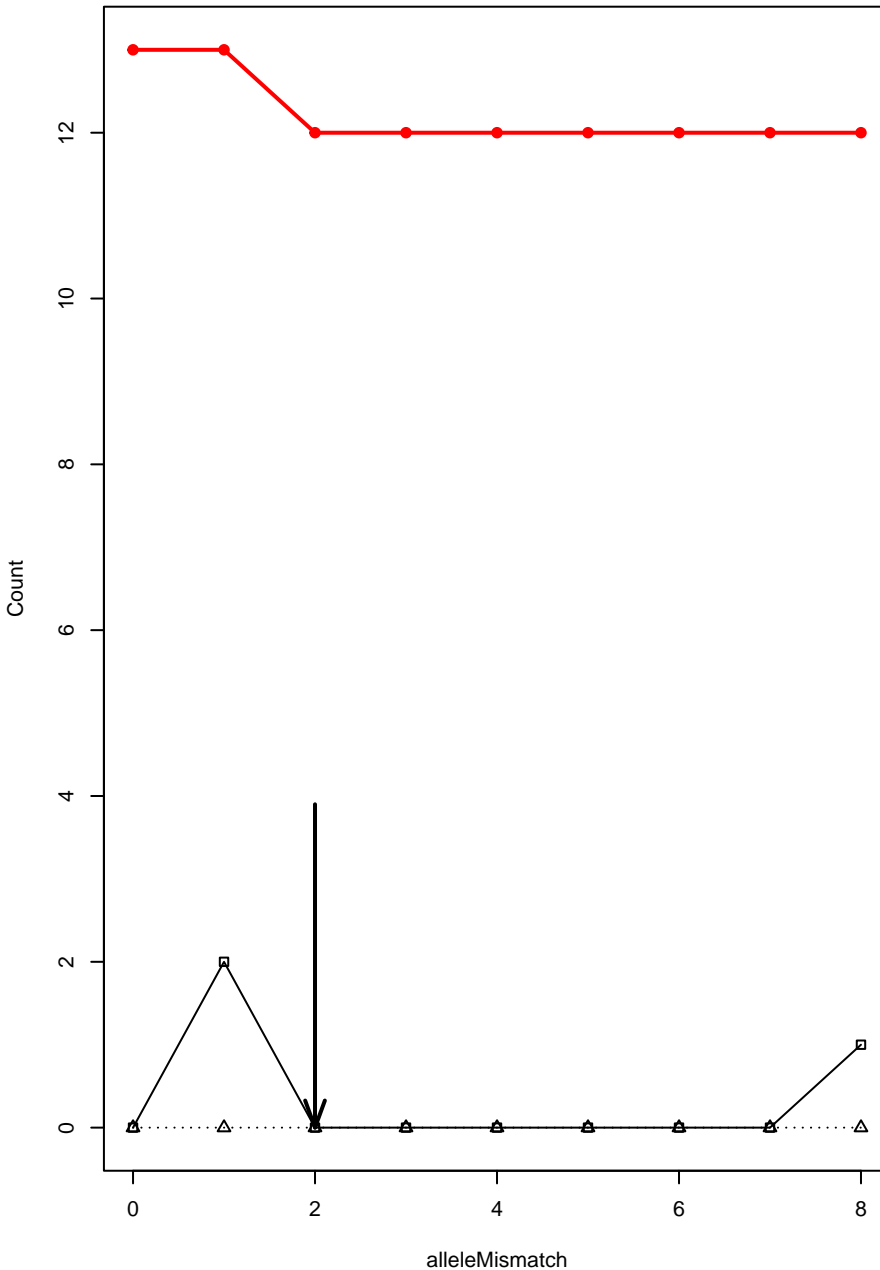


# allelematch

amUniqueProfile()

missingDataLoad=0.005

allelicDiversity=6.1



—●— unique  
—■— multipleMatch  
···△··· unclassified

Best guess for optimum:  
alleleMismatch=2

Profile morphology:  
ZeroSecondMinimum

# allelematch

amUniqueProfile()

missingDataLoad=0.046

allelicDiversity=7.9

