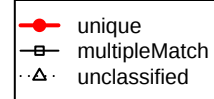
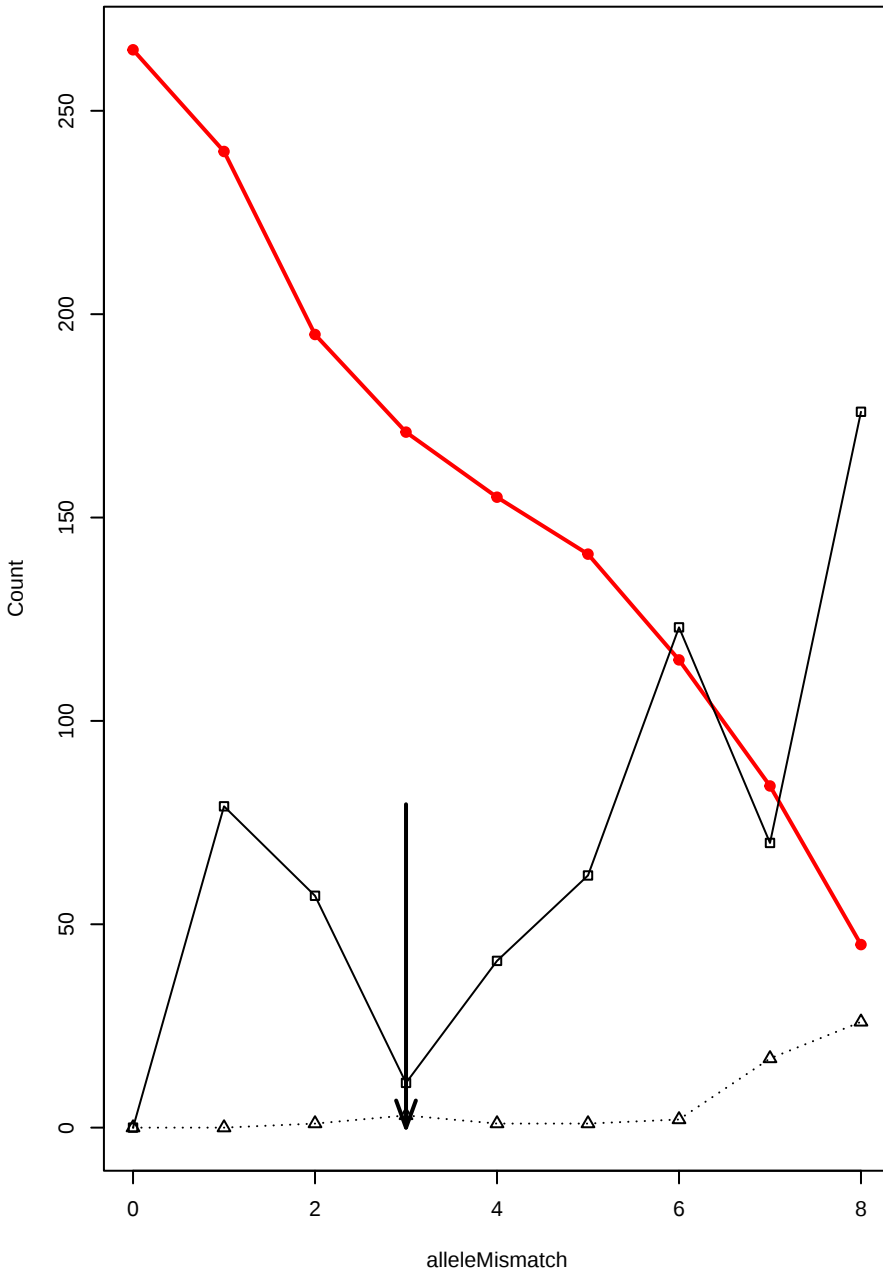


# allelematch

amUniqueProfile()

missingDataLoad=0.095

allelicDiversity=6.1



Best guess for optimum:  
alleleMismatch=3

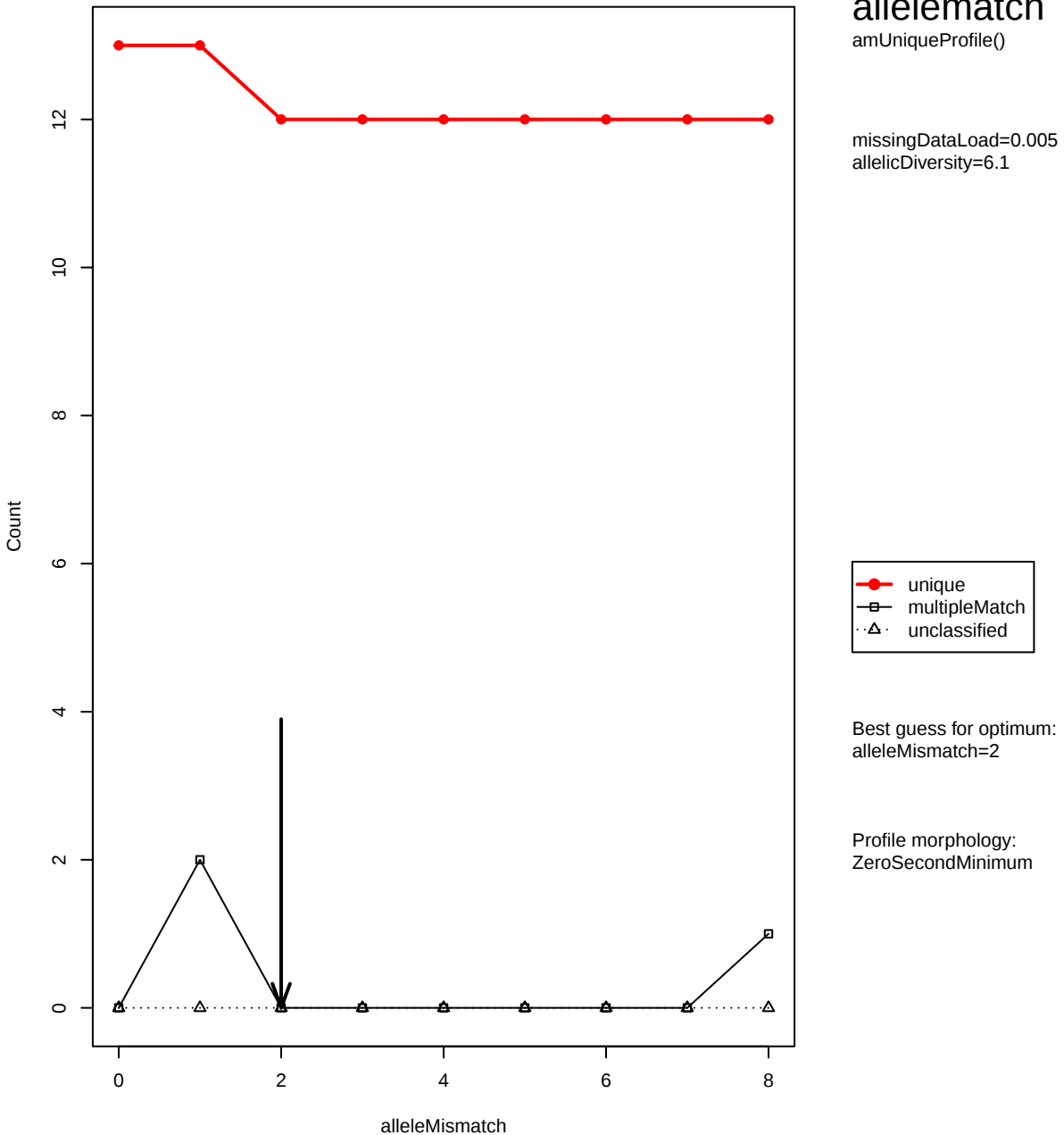
Profile morphology:  
NonZeroSecondMinimum  
Caution with optimum

# allelematch

amUniqueProfile()

missingDataLoad=0.005

allelicDiversity=6.1

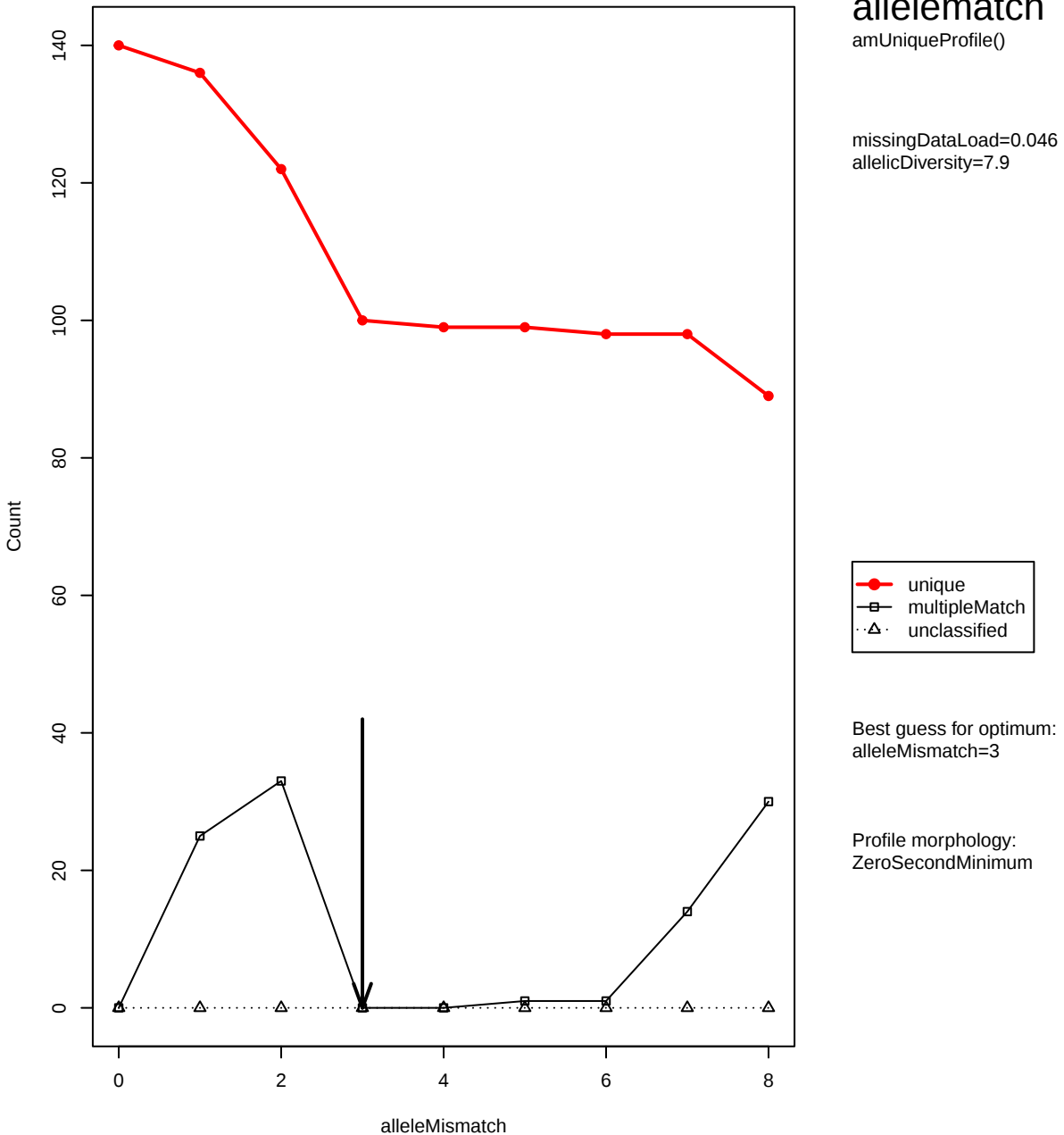


# allelematch

amUniqueProfile()

missingDataLoad=0.046

allelicDiversity=7.9

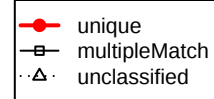
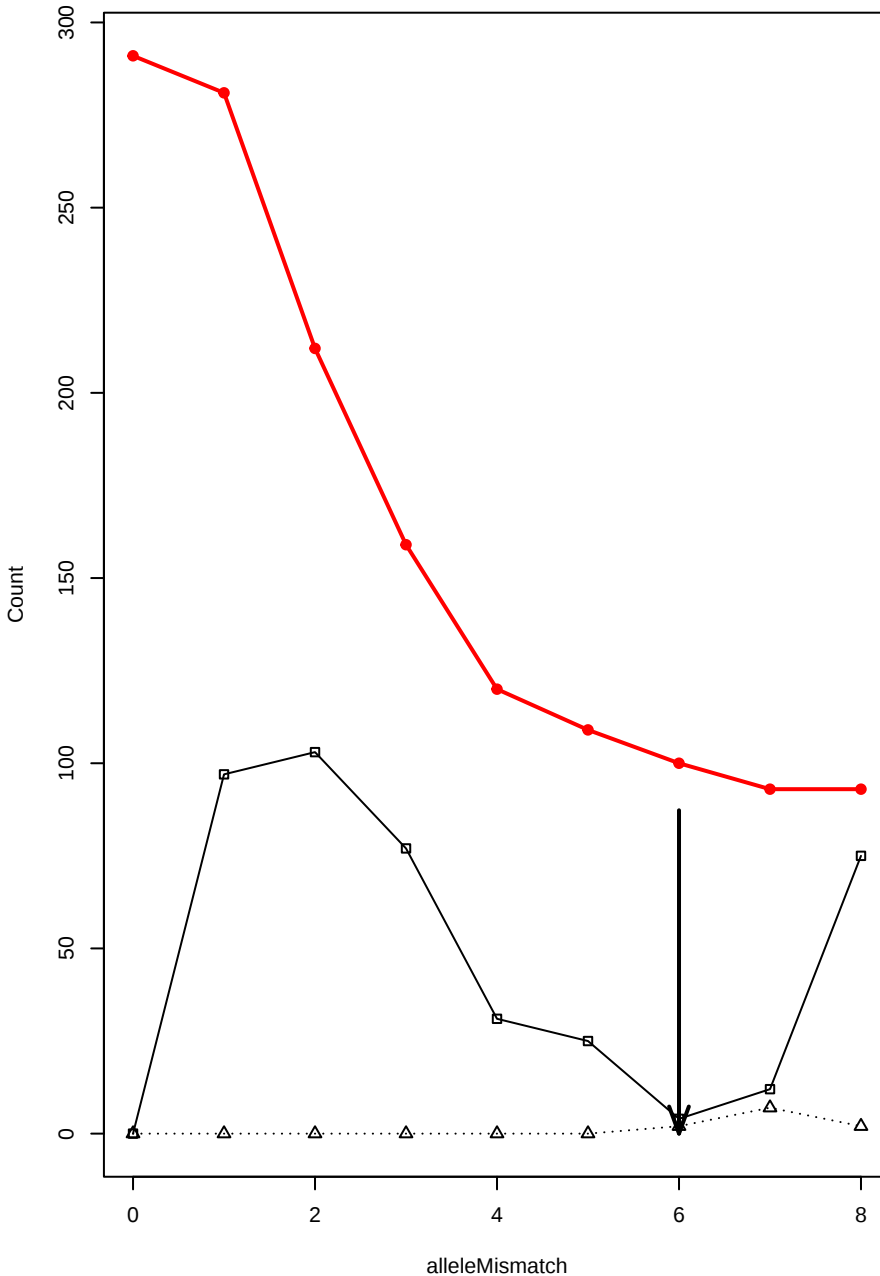


# allelematch

amUniqueProfile()

missingDataLoad=0.097

allelicDiversity=8.2



Best guess for optimum:  
alleleMismatch=6

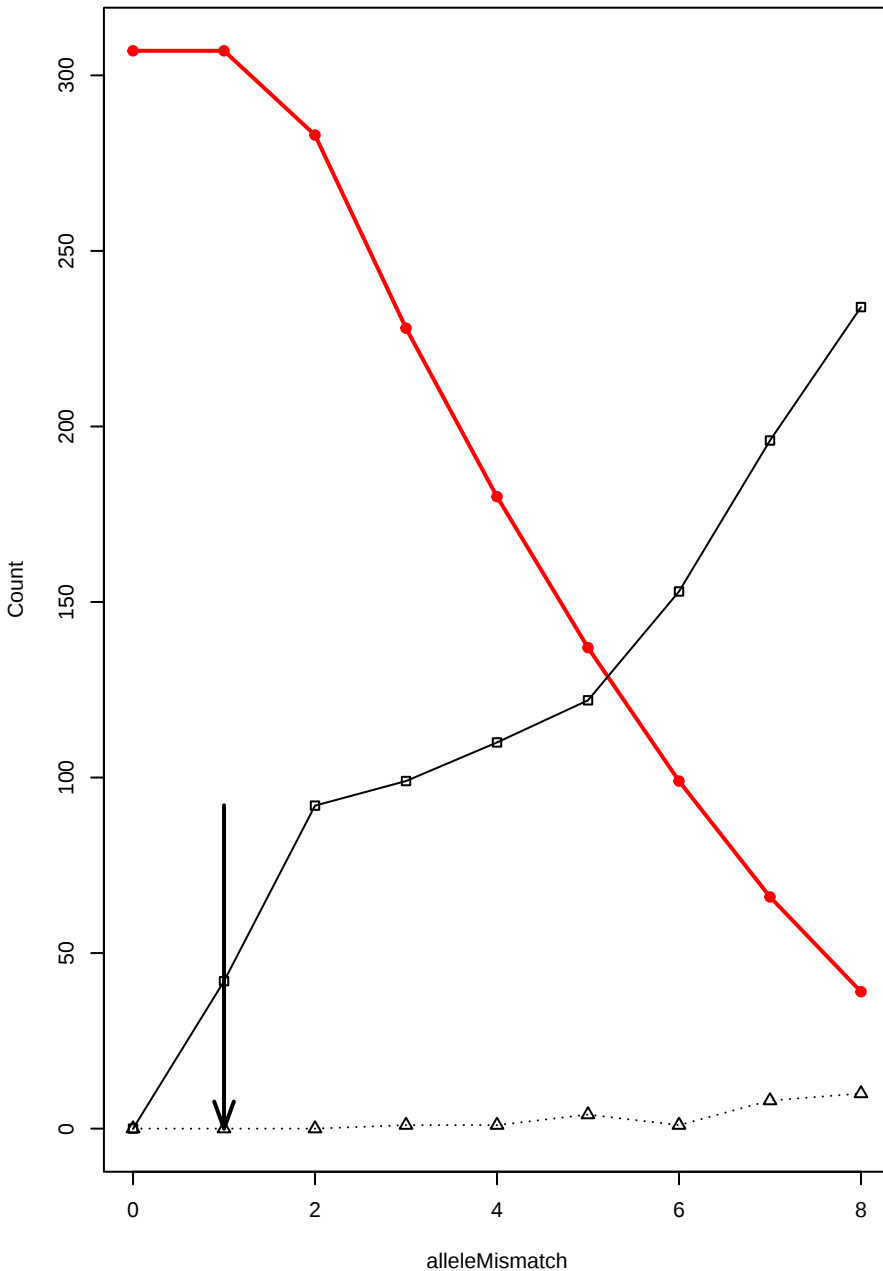
Profile morphology:  
NonZeroSecondMinimum  
Caution with optimum

# allelematch

amUniqueProfile()

missingDataLoad=0.199

allelicDiversity=4.8



Best guess for optimum:  
alleleMismatch=1

Profile morphology:  
NoSecondMinimum  
Caution with optimum