test set error for classification algorithm
IncRNA group = allLncRNA ::: data cols = lpa
49 / 3453 ) = (func/total lncRNA) :: biotype = lincProcTra

(49/3453) = (func/total IncRNA) ::: biotype = lincProcTrans traning set Size = 1367, trials= 10 ratio = 0.7 scale = FALSE

