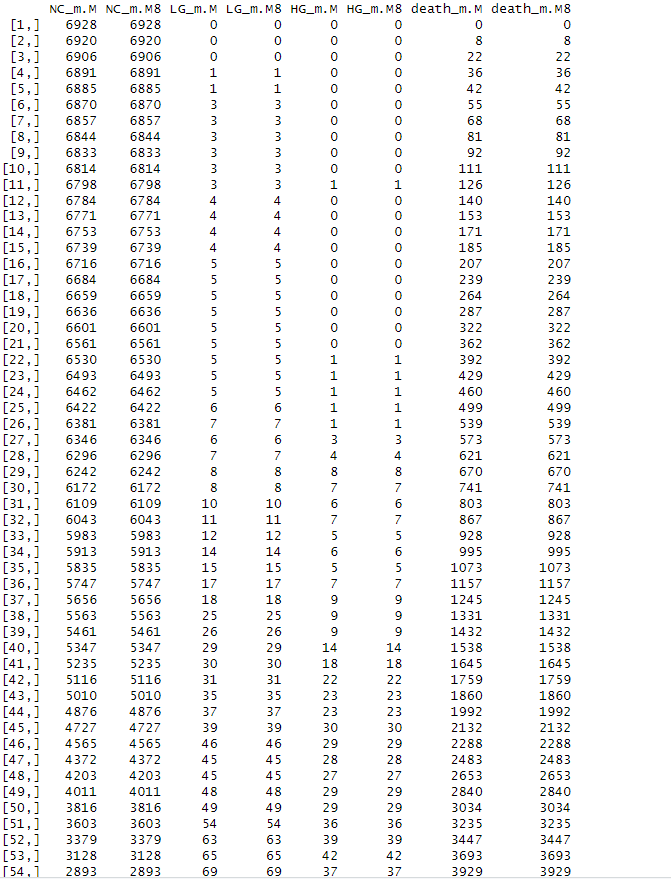
# Run the model for screening pop

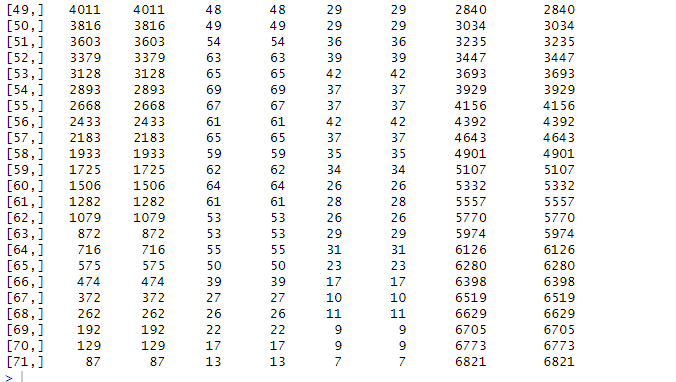
DS\_screen =1 #Set whether the screening with dipstick happens, 0 - no, 1- yes

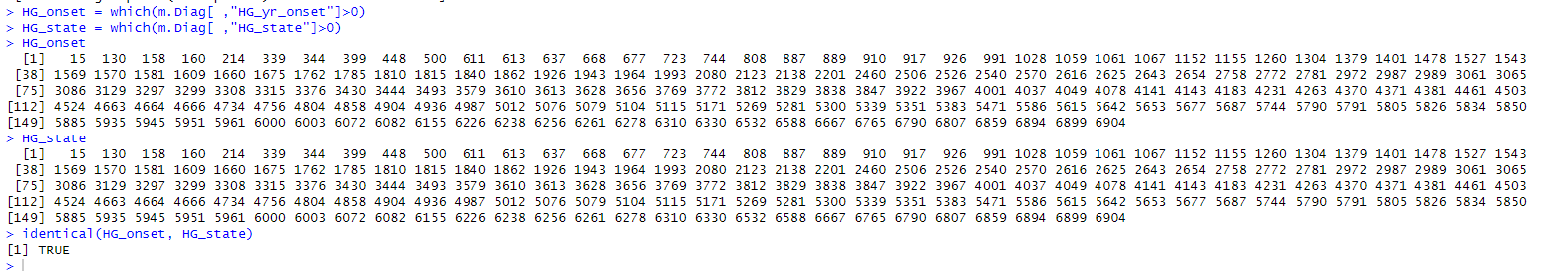
DS\_age = 50 #Set the age of the dipstick if screening happens, set to zero if no screening

DS\_round = 15 #' DS\_round - number of the screening rounds

DS\_freq =2







# check mort from BC is reflected

m.Diag =cbind(m.Diag,pop[ ,"PID"])

m.Diag\_c = m.Diag[cancer\_m.Diag, ]

p15\_a=83

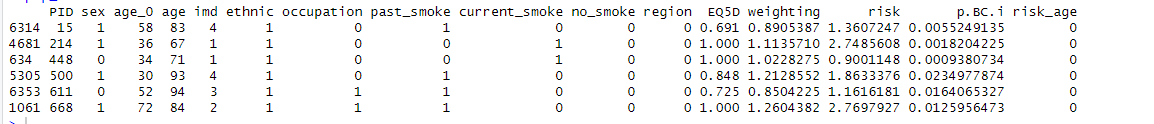
p214\_a=67

p448\_a=71

p500\_a=93

p611\_a=94

p668\_a=84



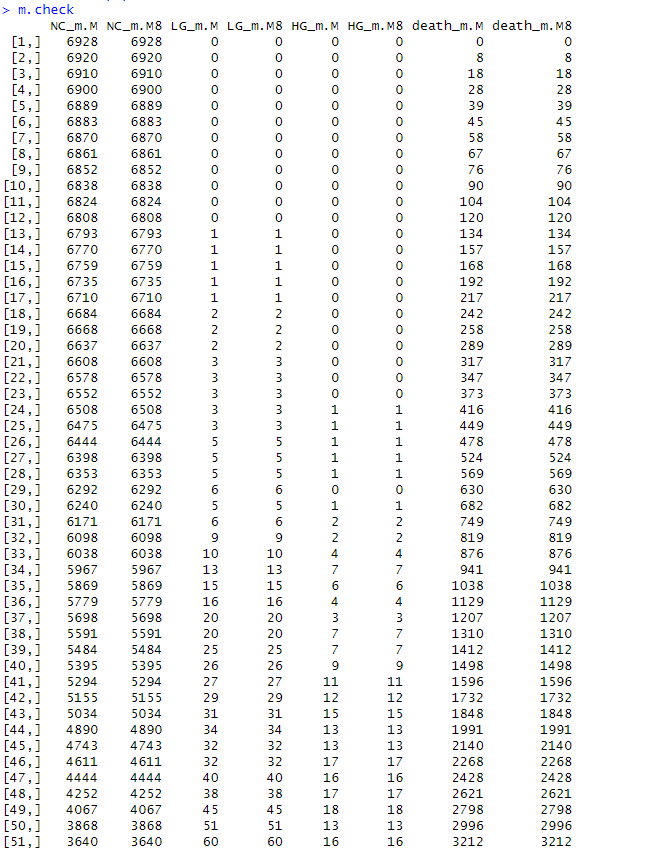
# Run the model for screening pop

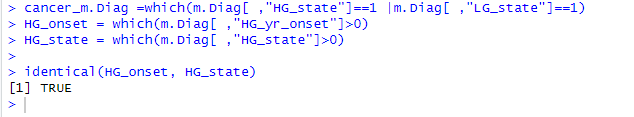
DS\_screen =1 #Set whether the screening with dipstick happens, 0 - no, 1- yes

DS\_age = 50 #Set the age of the dipstick if screening happens, set to zero if no screening

DS\_round = 15 #' DS\_round - number of the screening rounds

DS\_freq =1 #' D

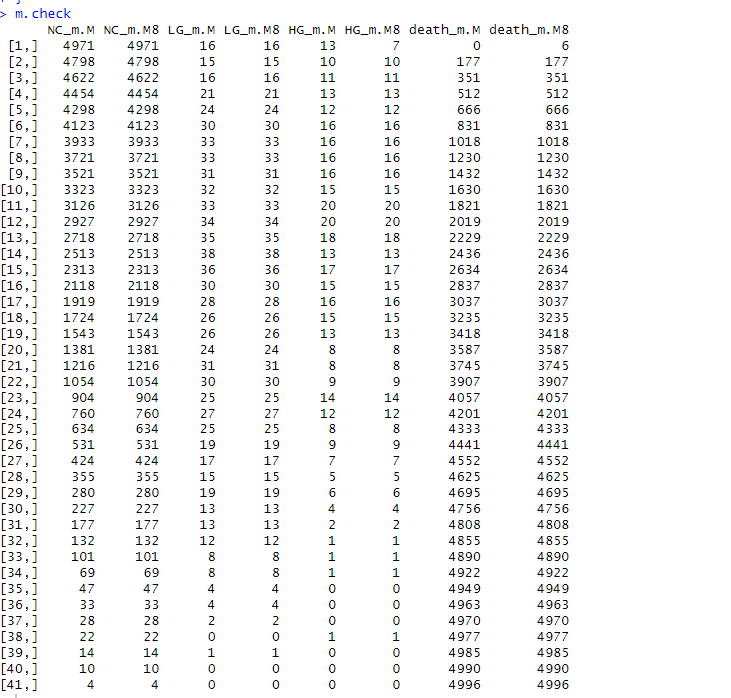




All-England pop (non cohort)

DS\_screen =0 #Set whether the screening with dipstick happens, 0 - no, 1- yes

DS\_age = 100



# Run the model for screening pop

DS\_screen =1 #Set whether the screening with dipstick happens, 0 - no, 1- yes

DS\_age = 50 #Set the age of the dipstick if screening happens, set to zero if no screening

DS\_round = 15 #' DS\_round - number of the screening rounds

DS\_freq =1 #' DS\_freq - frequency of the screening rounds (either 1 (annual) or 2 (biennial))

#iter=1

