mtcars

attach(mtcars)

base<-glm(am~1,data=mtcars, family=binomial())

base

fit1<-glm(am~mpg+disp+hp+wt, family=binomial())

fit1

summary(fit1)

anova(fit1)

fit2<-glm(am~mpg+hp+wt, family=binomial())

fit2

summary(fit2)

fit3<-glm(am~hp+wt, family=binomial())

fit3

summary(fit3)

library(Rcmdr)

layout(matrix(c(1,2,3,4),2,2))

plot(fit3)

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