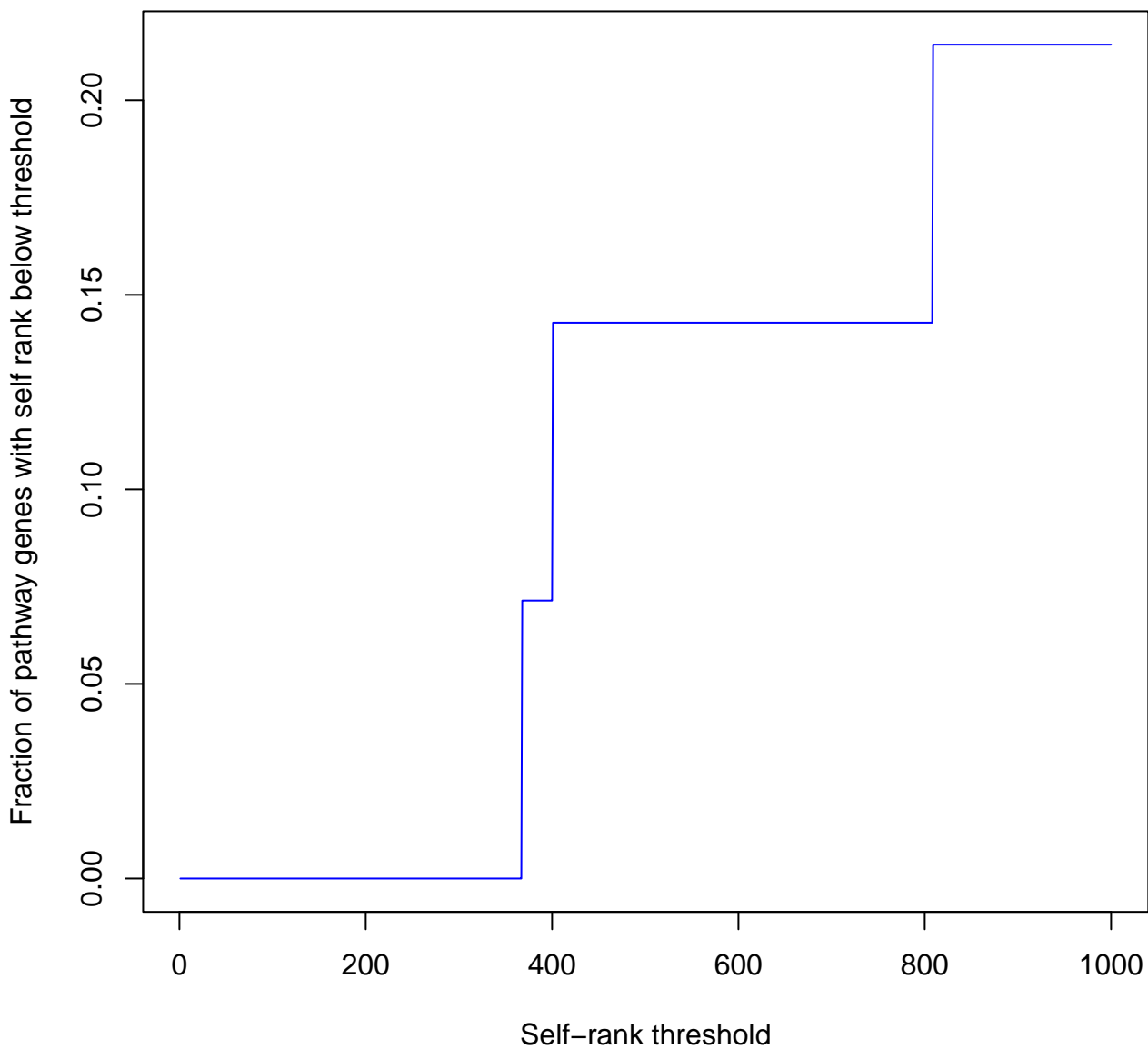
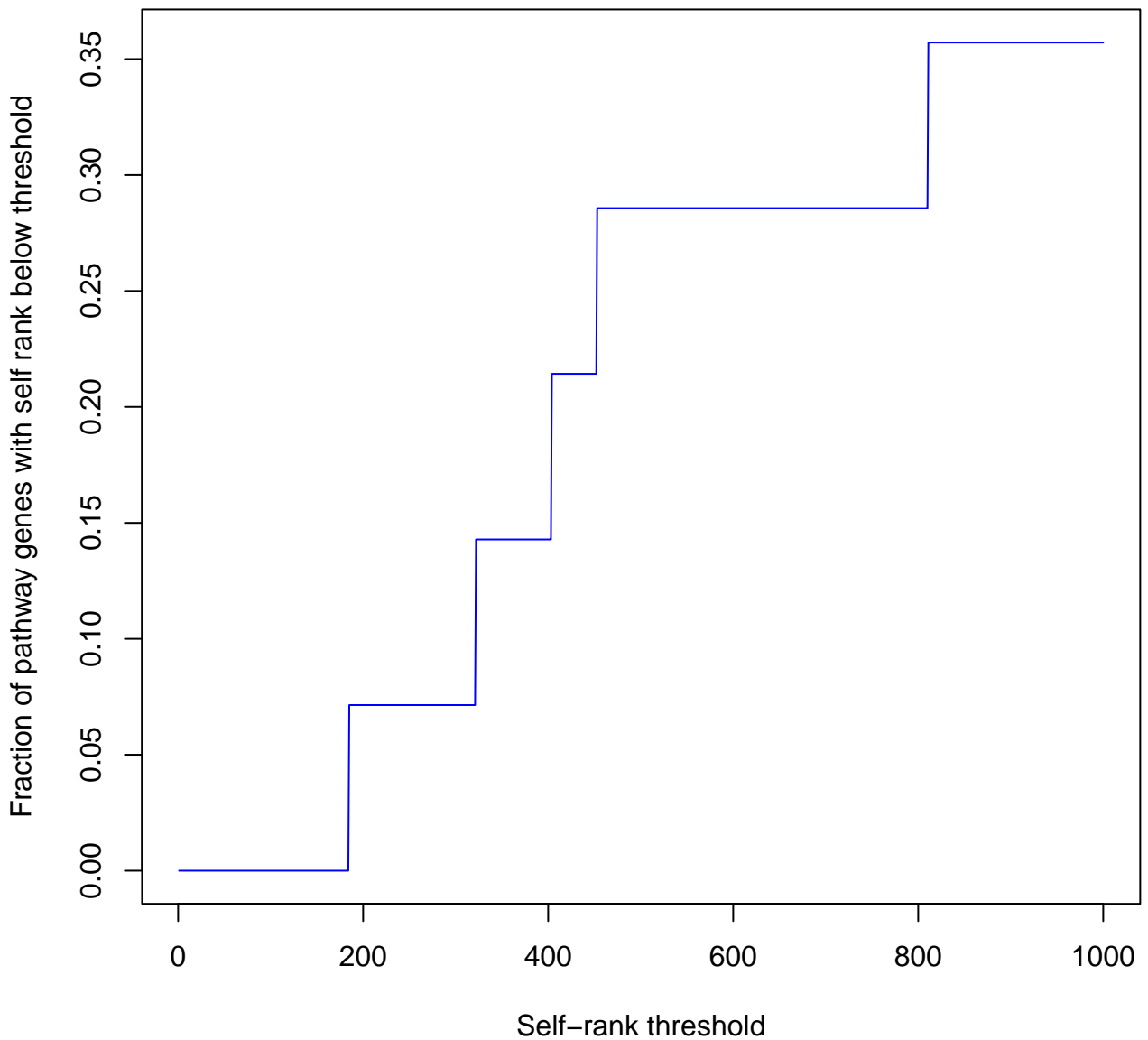


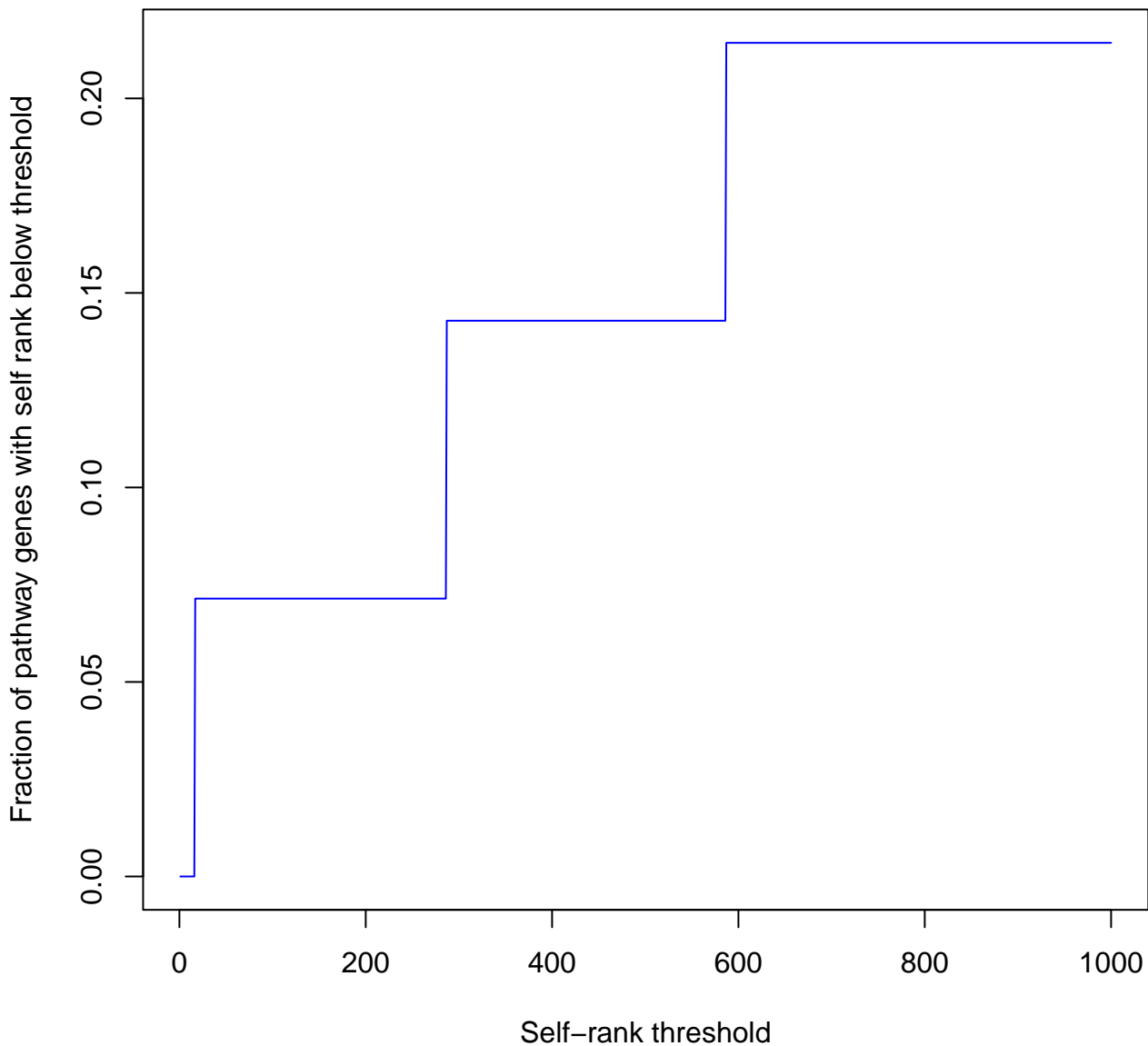
**methylation\_dataset.txt with kmeansmethylation.txt**



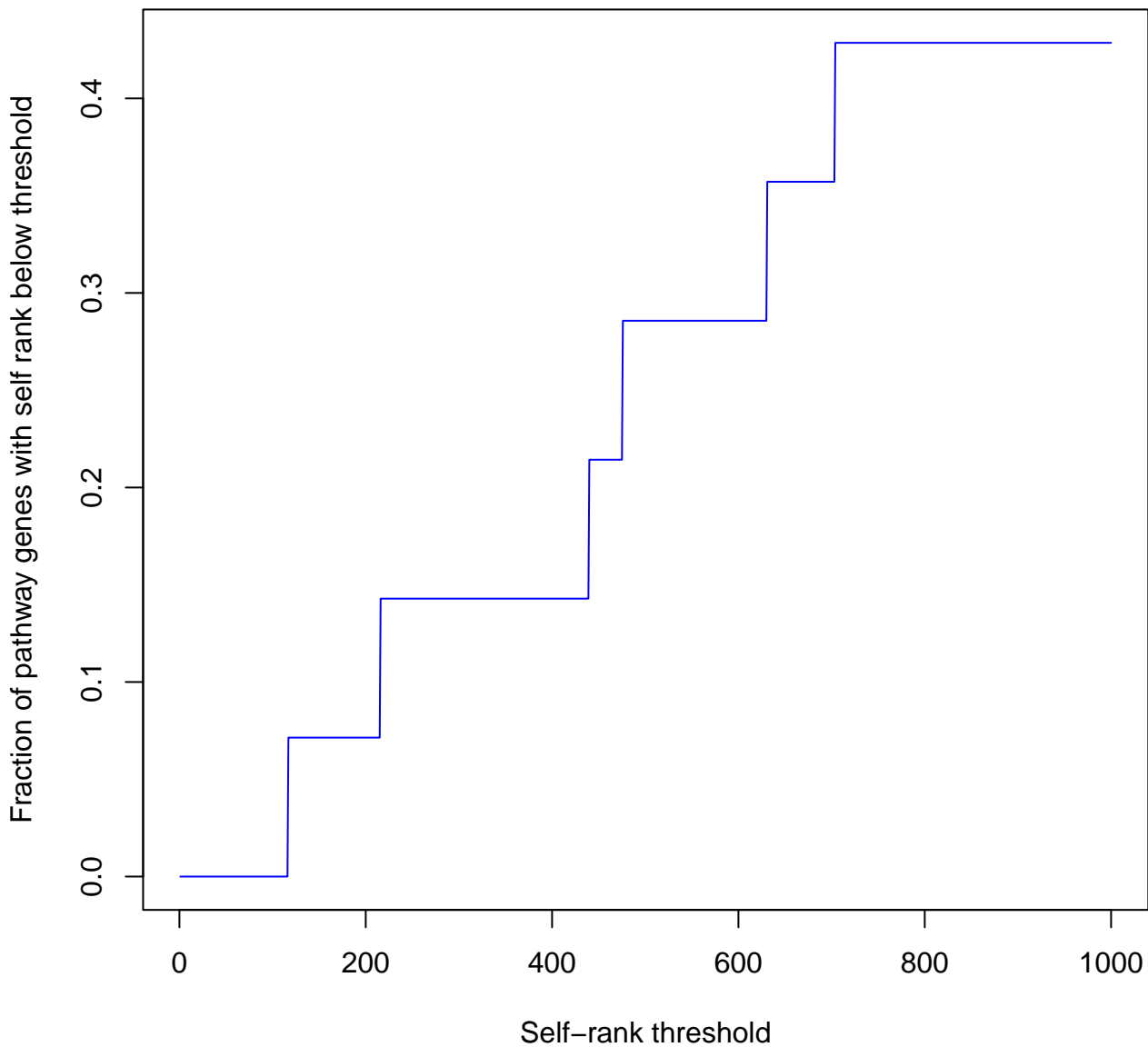
# methylation\_dataset.txt with sommethylation.txt



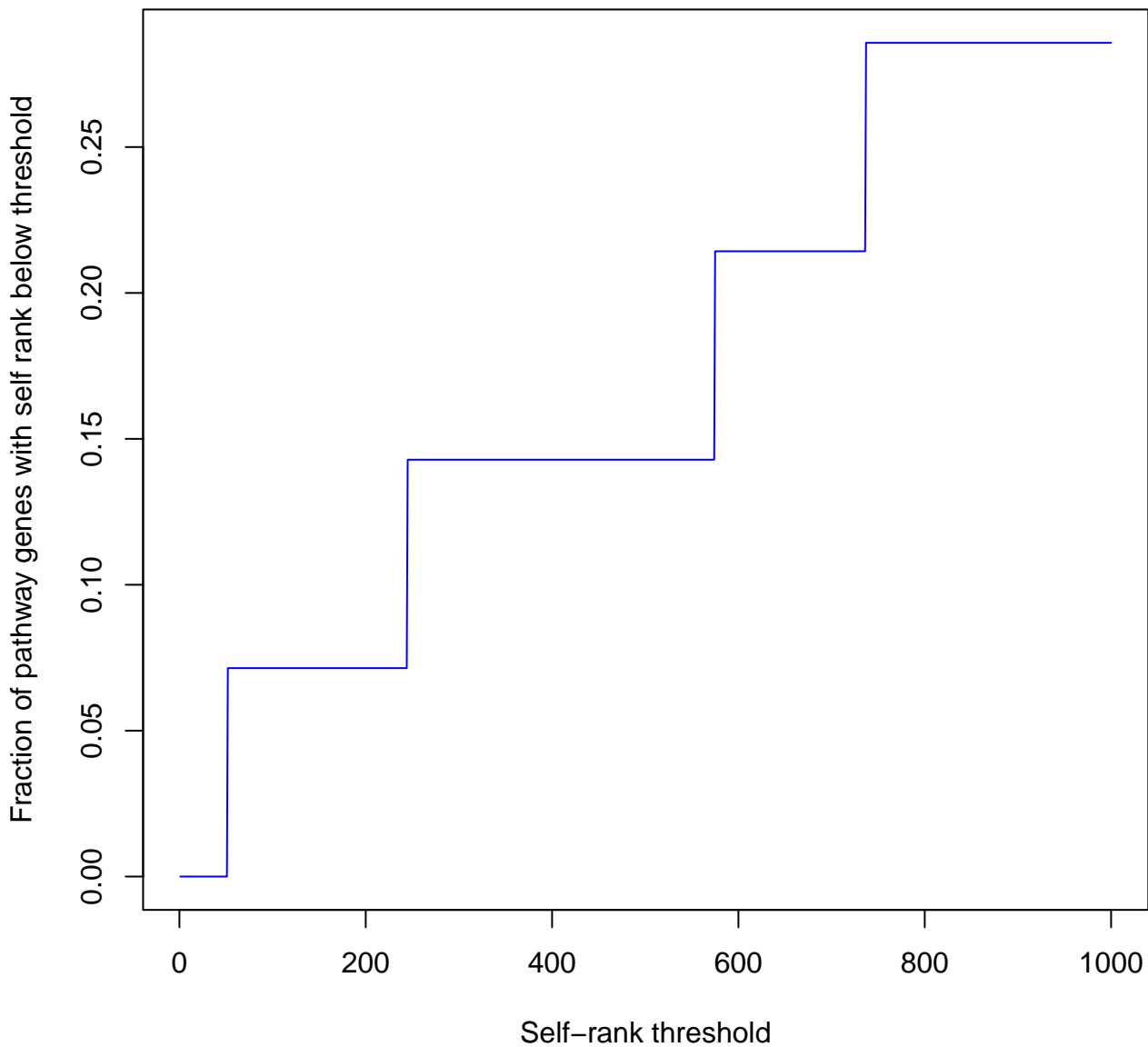
**drug\_cholesterol\_toxin.txt with kmeansdrug.txt**



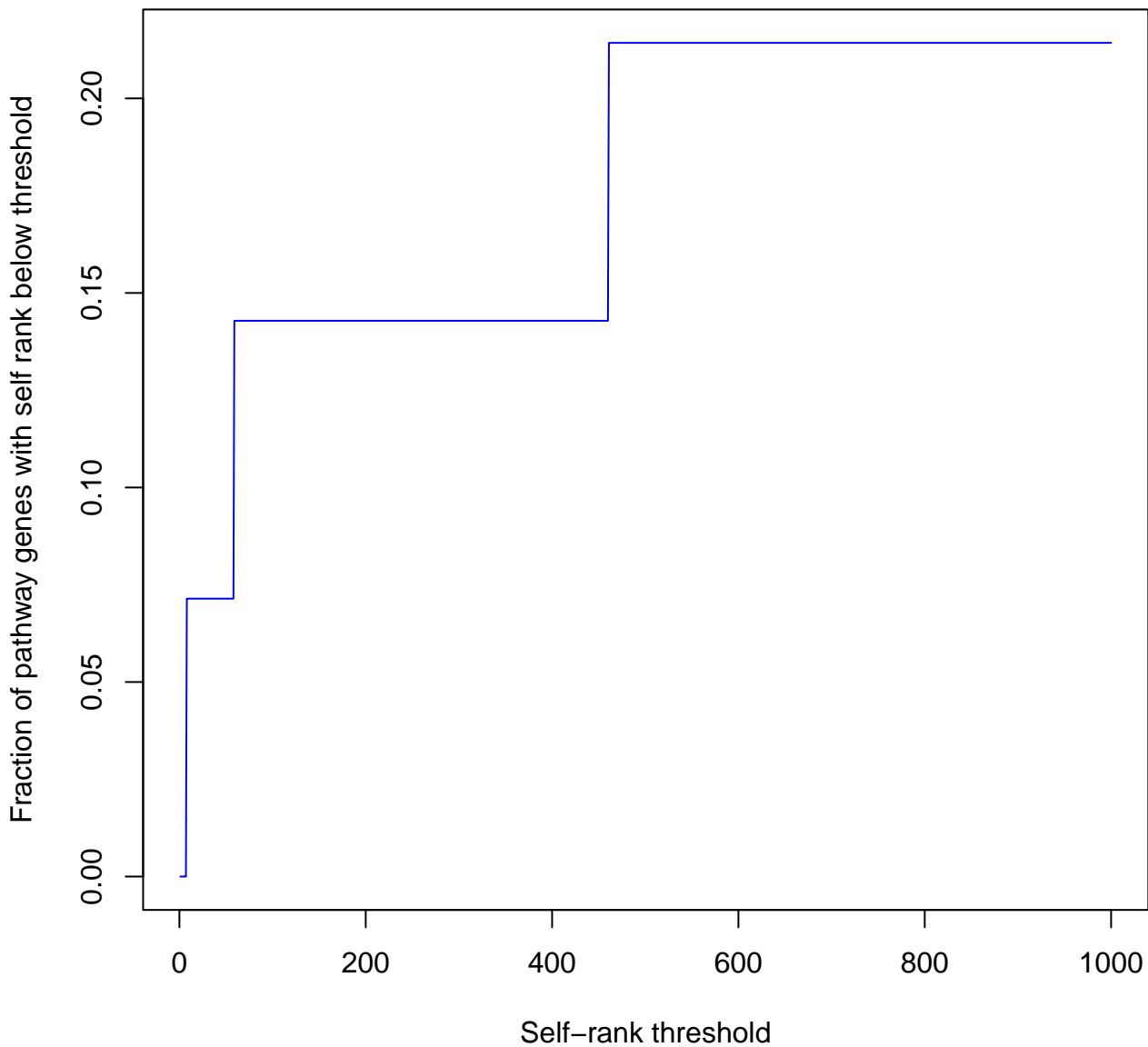
# drug\_cholesterol\_toxin.txt with somdrug.txt



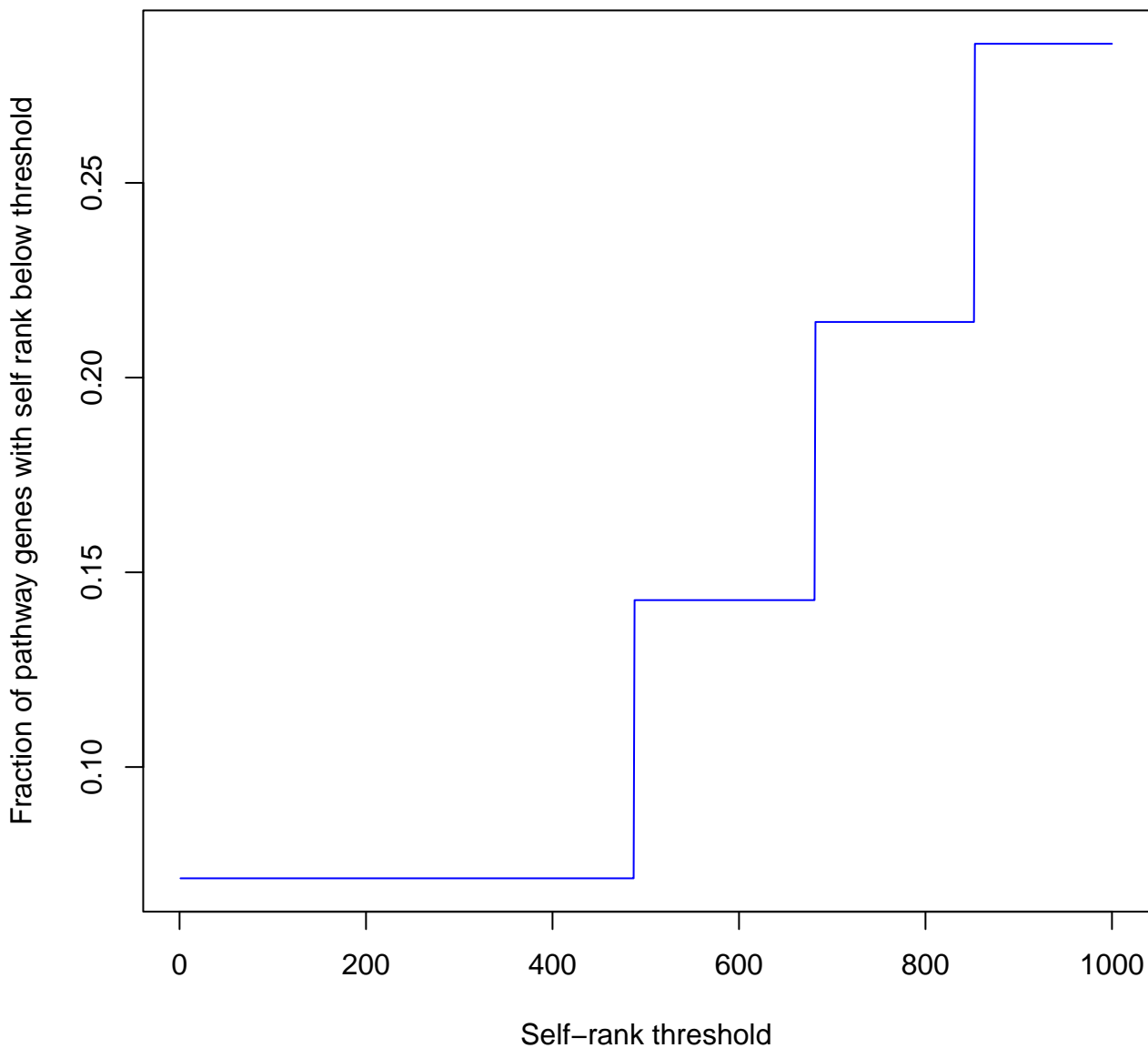
# ESX\_WT\_mutant.txt with kmeansESX.txt



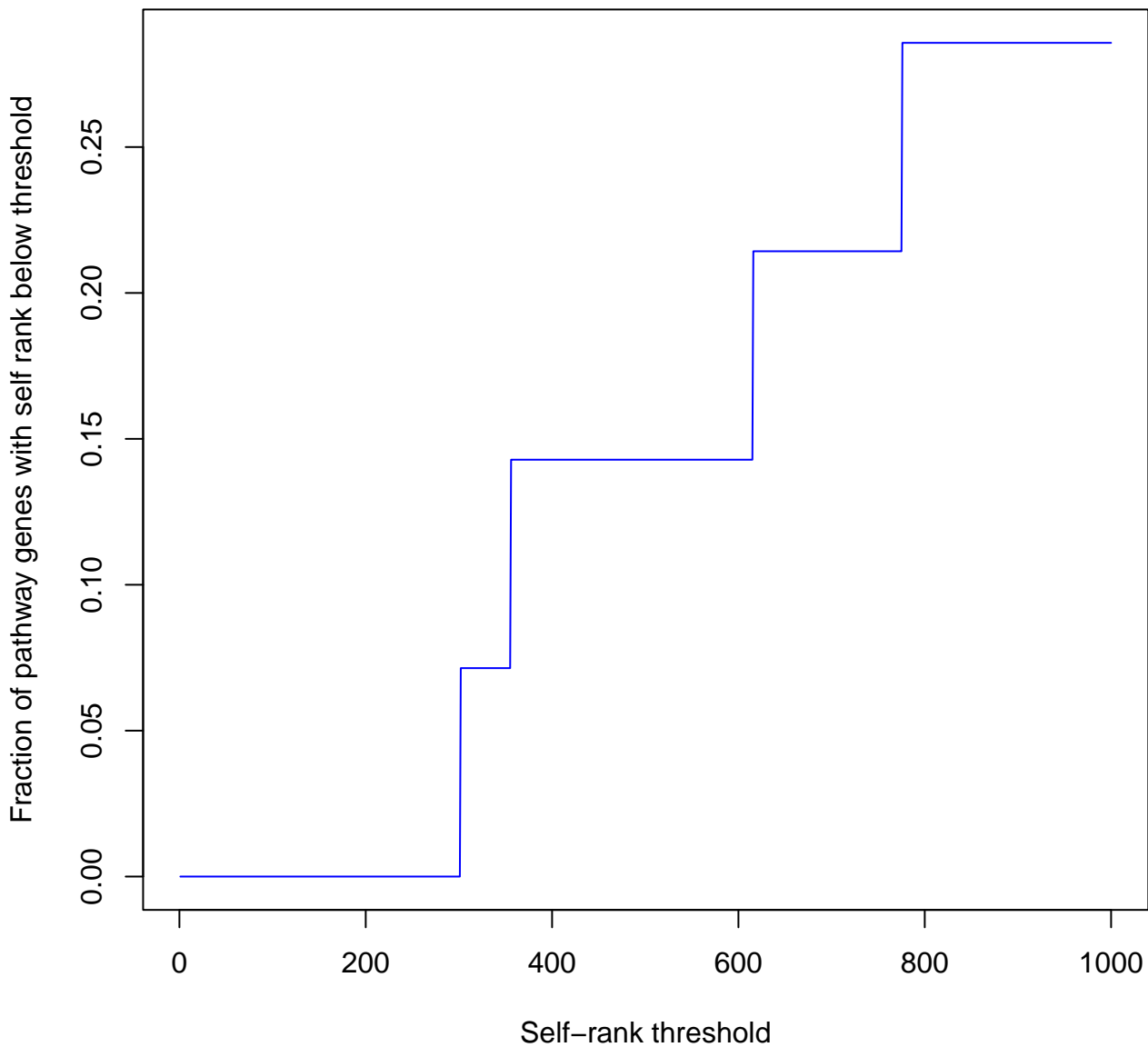
# ESX\_WT\_mutant.txt with somESX.txt



# MTBc\_all\_lineages.txt with kmeansMTBc\_lineages.txt

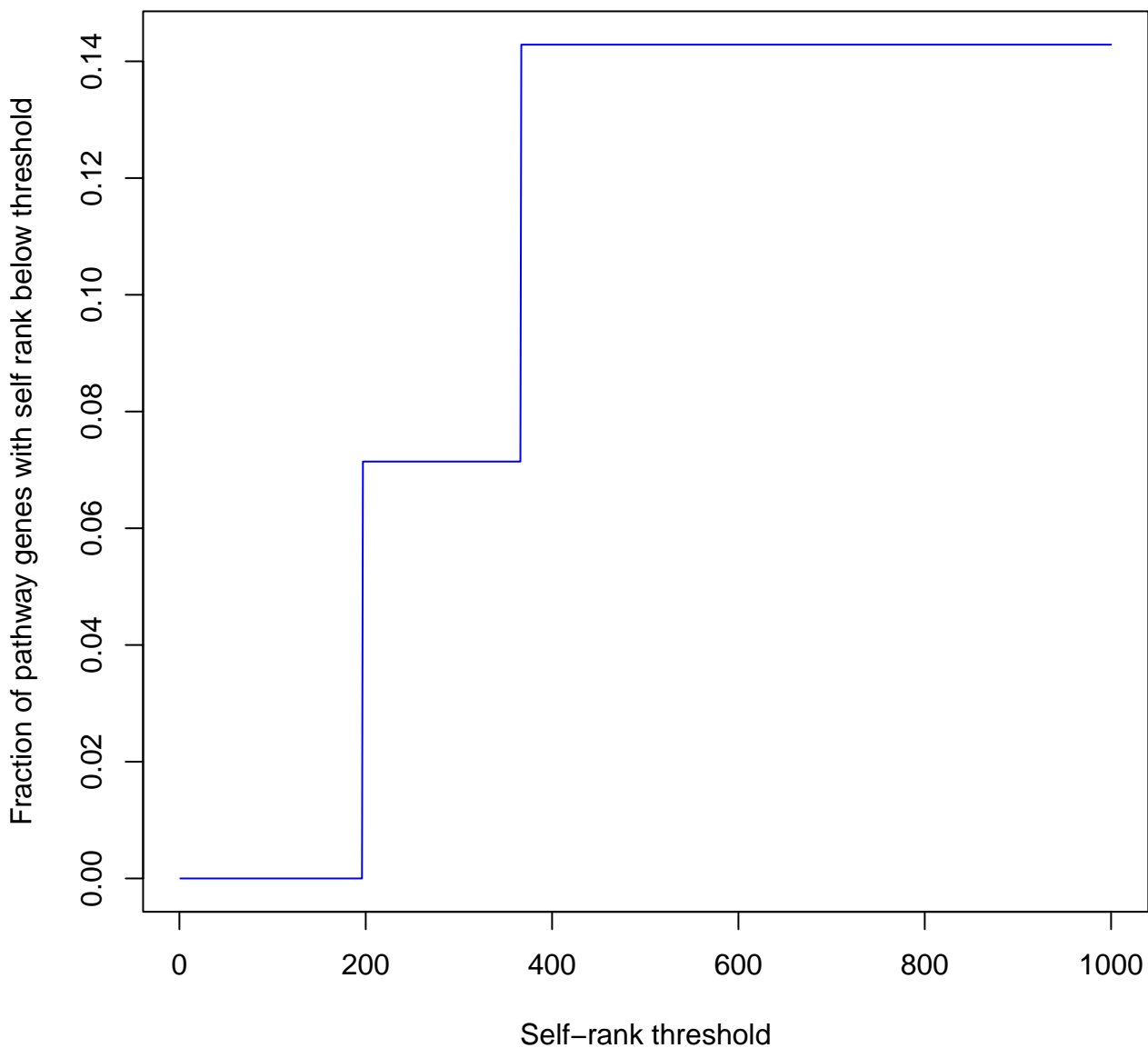


# MTBc\_all\_lineages.txt with somMTBc\_lineages.txt

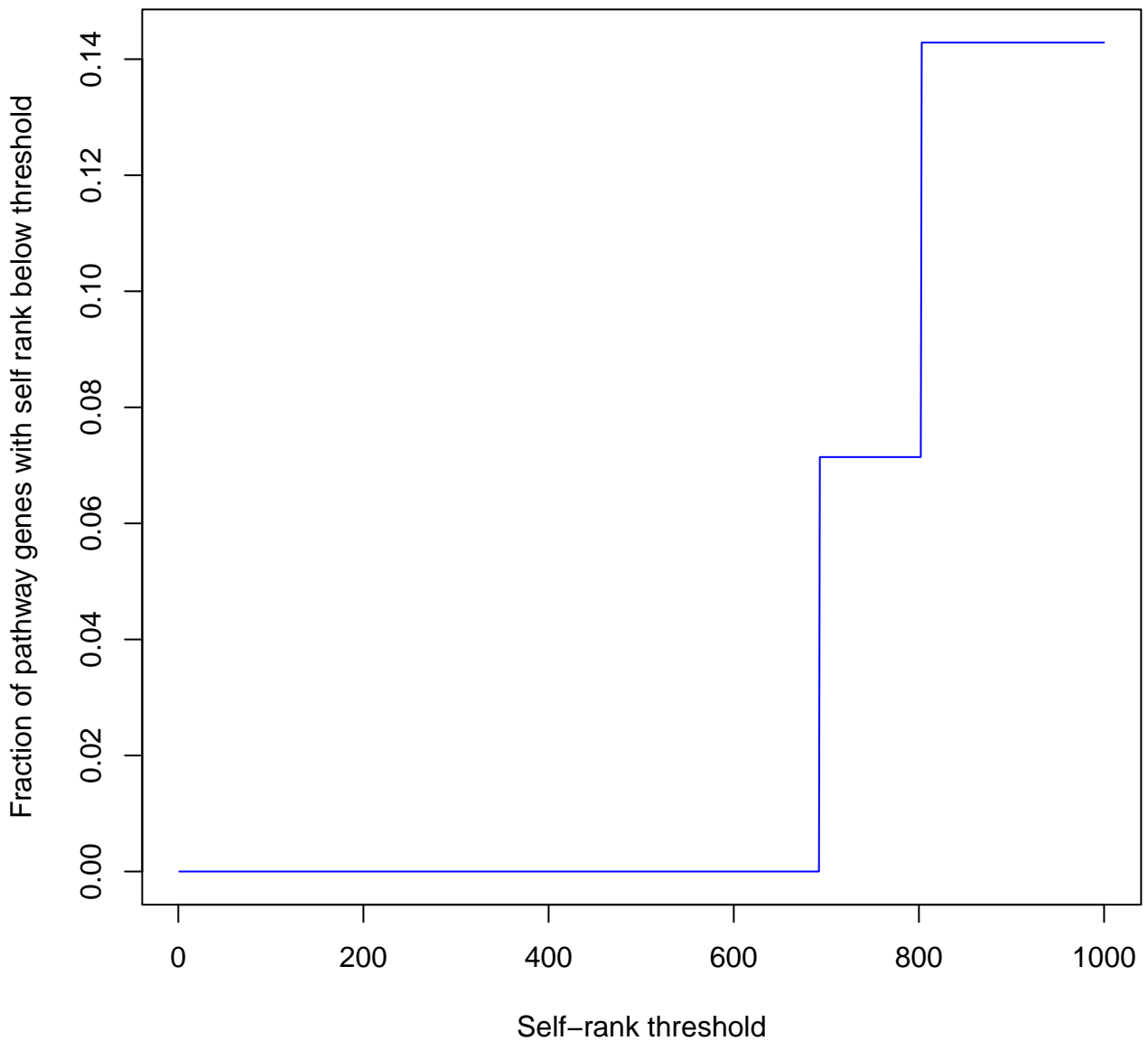




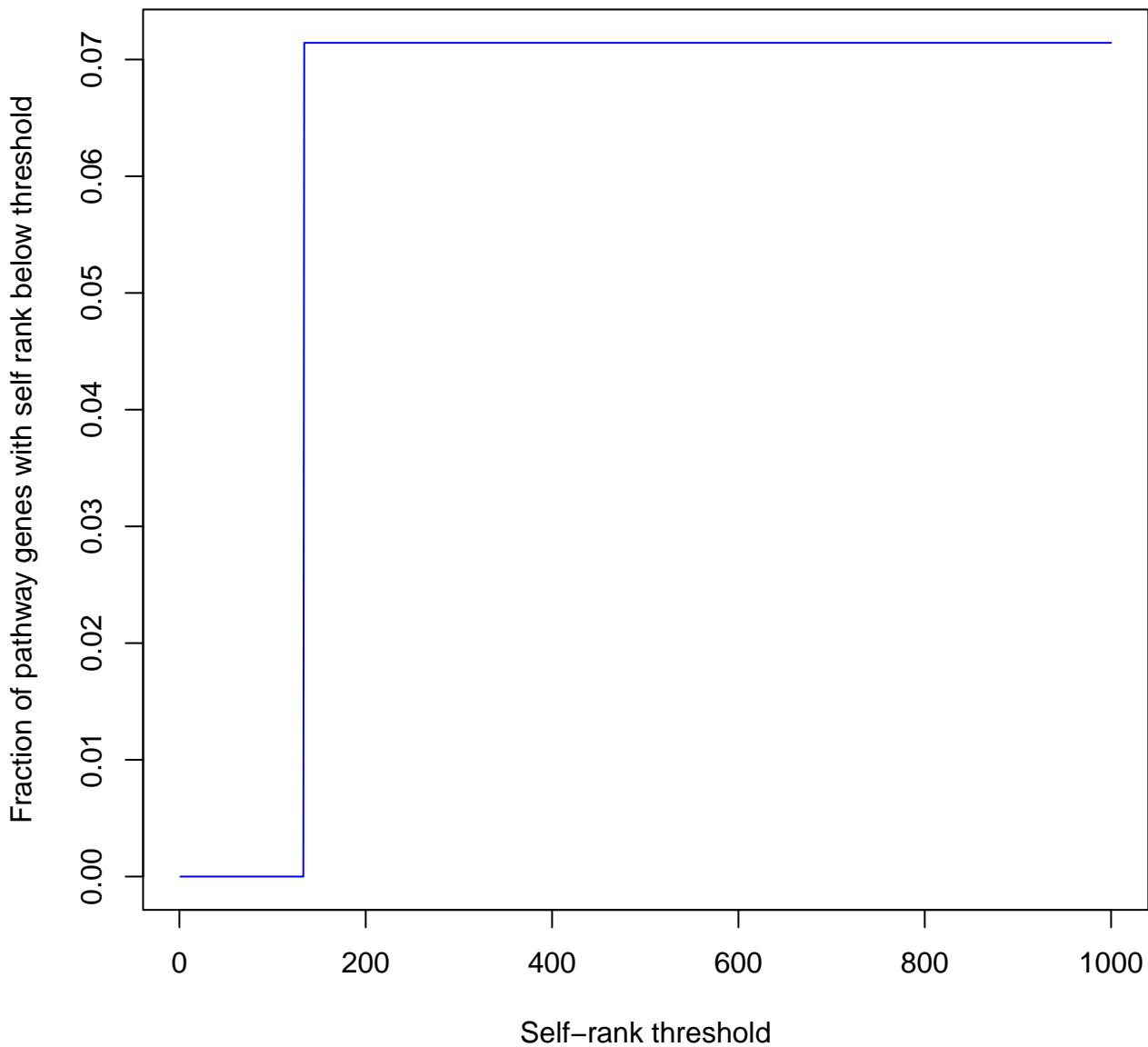
# primary\_drug.txt with kmeansprimary.txt



# primary\_drug.txt with somprimary.txt



**timecourse\_nitricacid.txt with kmeanstimecourse.txt**



**timecourse\_nitricacid.txt with somtimecourse.txt**

