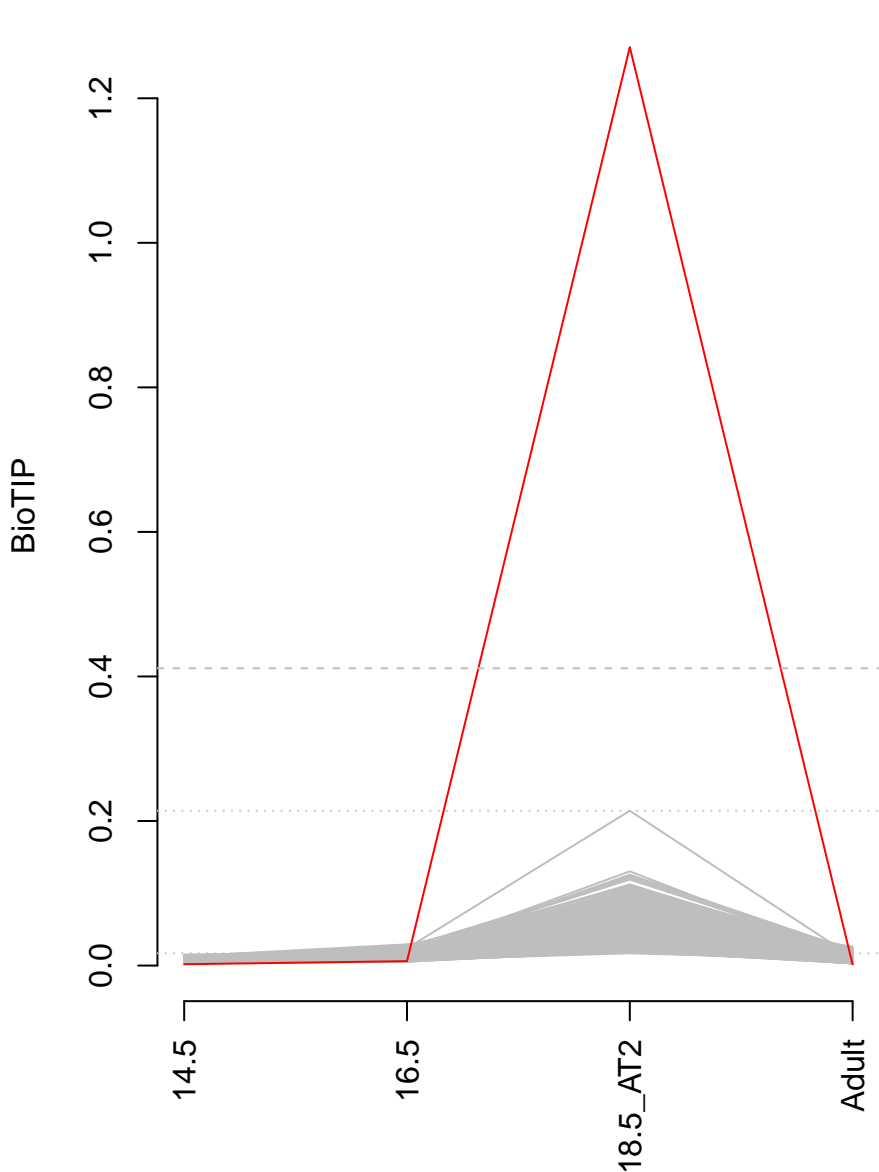


CTS.E18 (267 transcripts)



ensity.default(x = simuBioTIP_s["18.5_AT2",], ylim = c(

