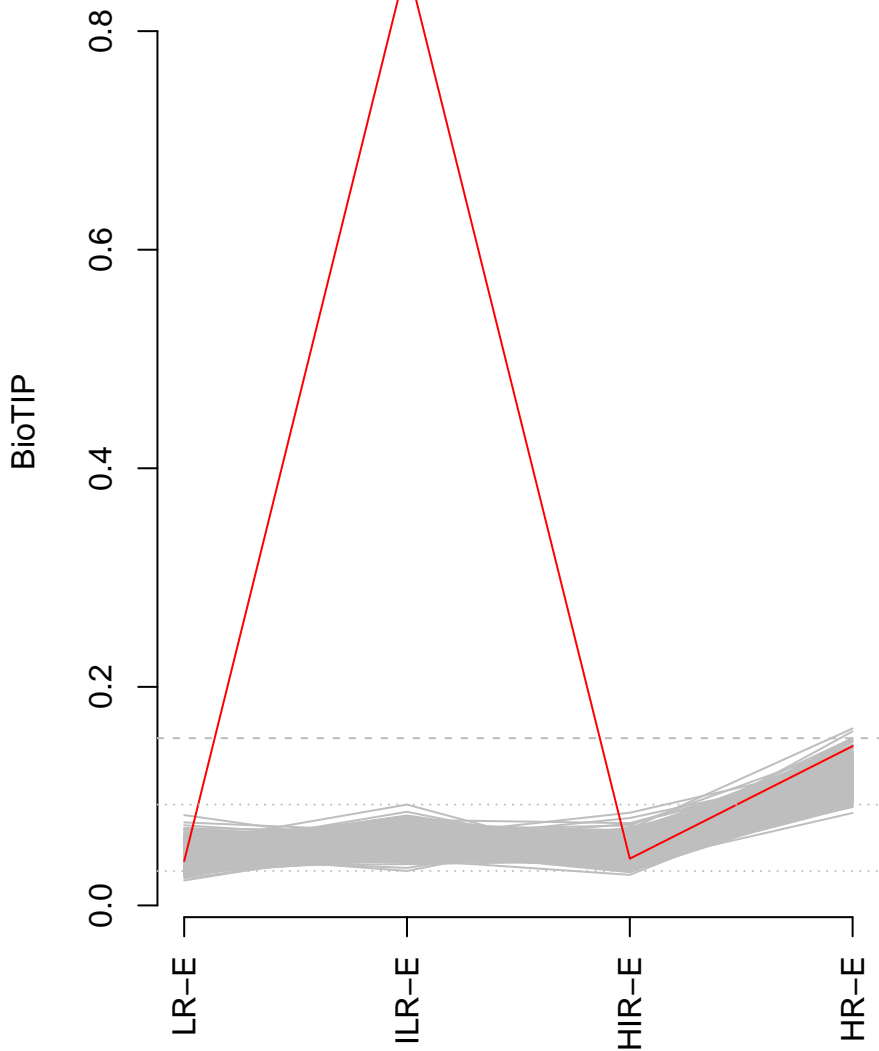


CTS.E (119 transcripts)



density.default(x = simuBioTIP\_s["ILR-E", ])

