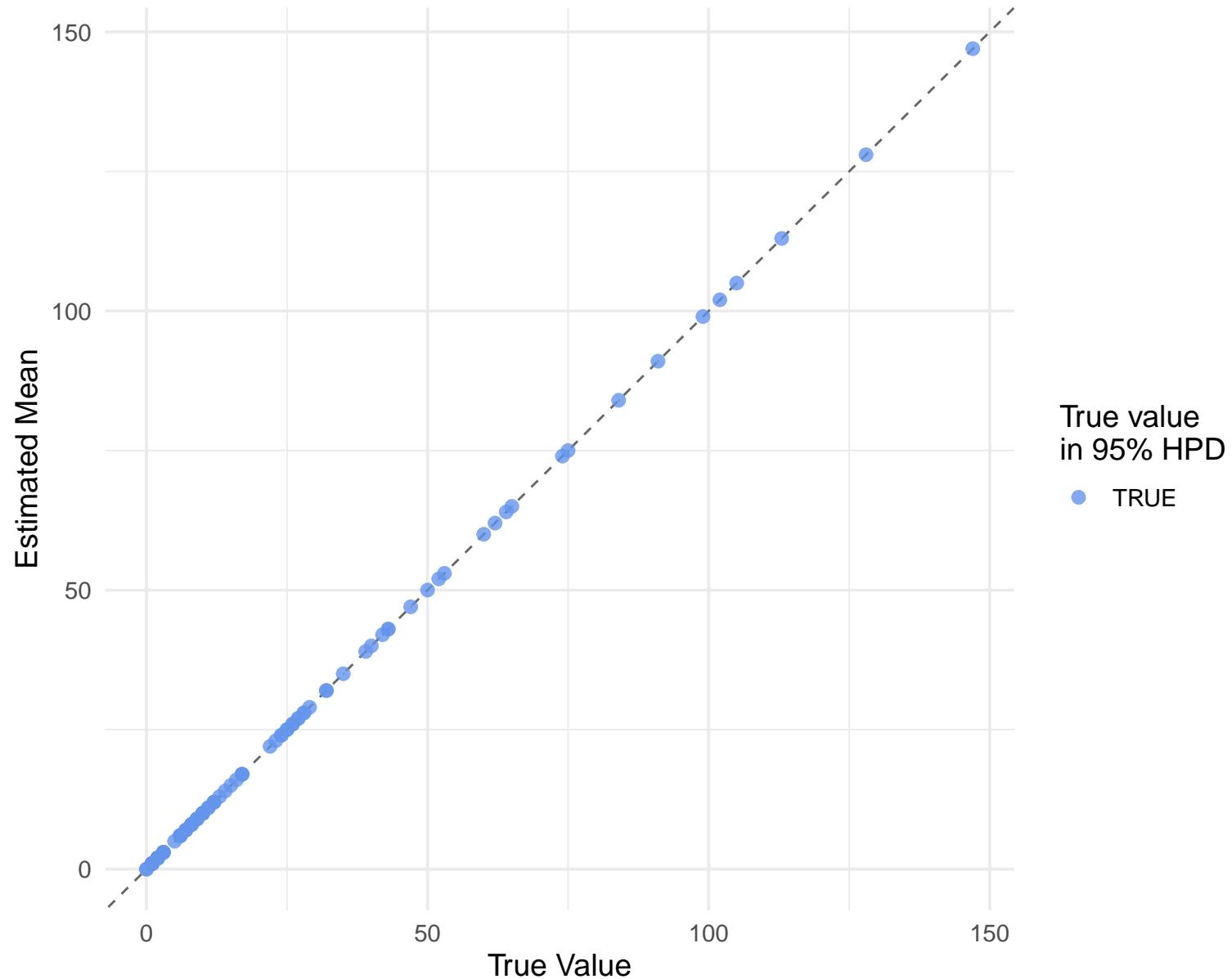


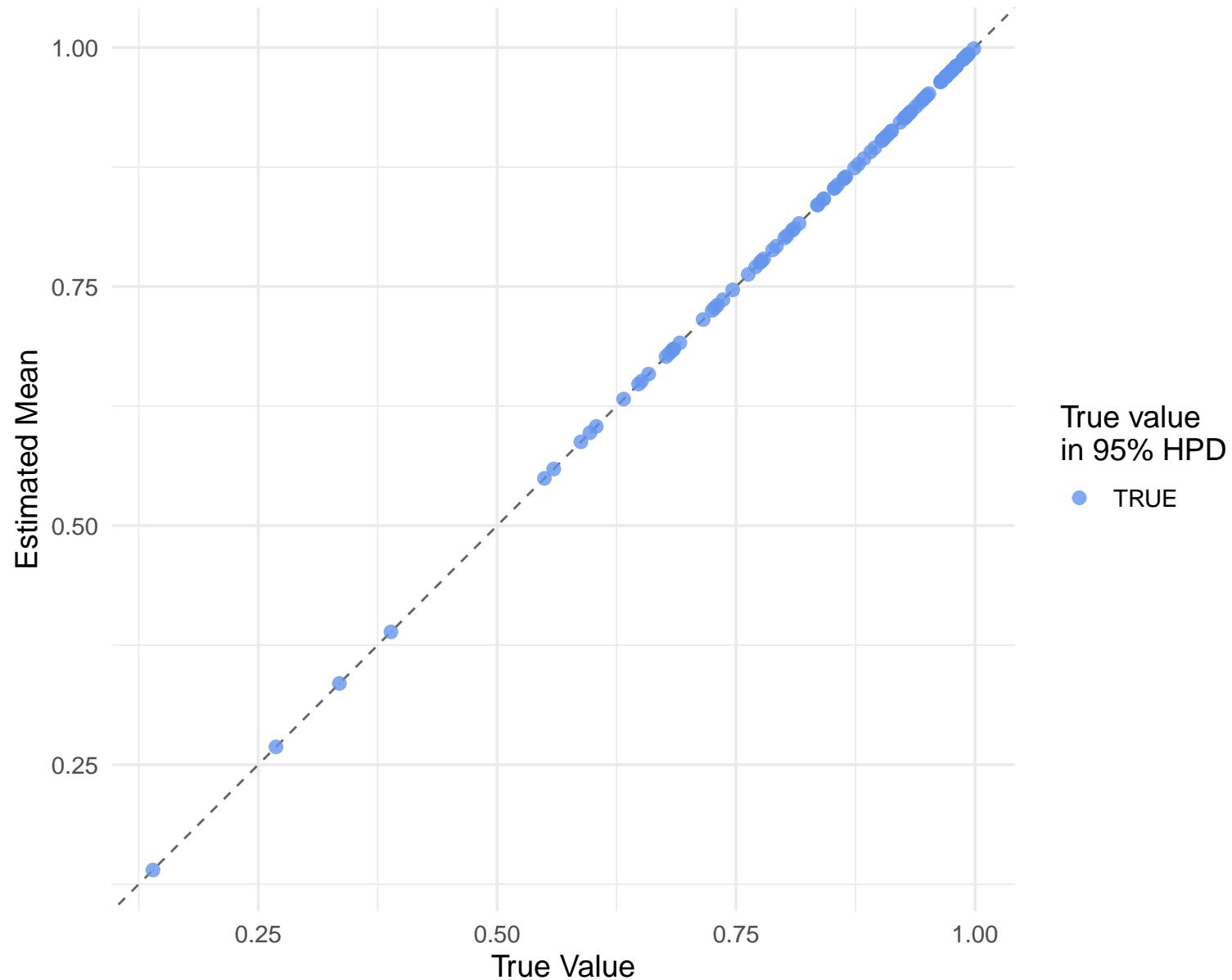
# nSampledAncestors

Coverage = 100.0%, Pearson's r = 1.000, N = 99



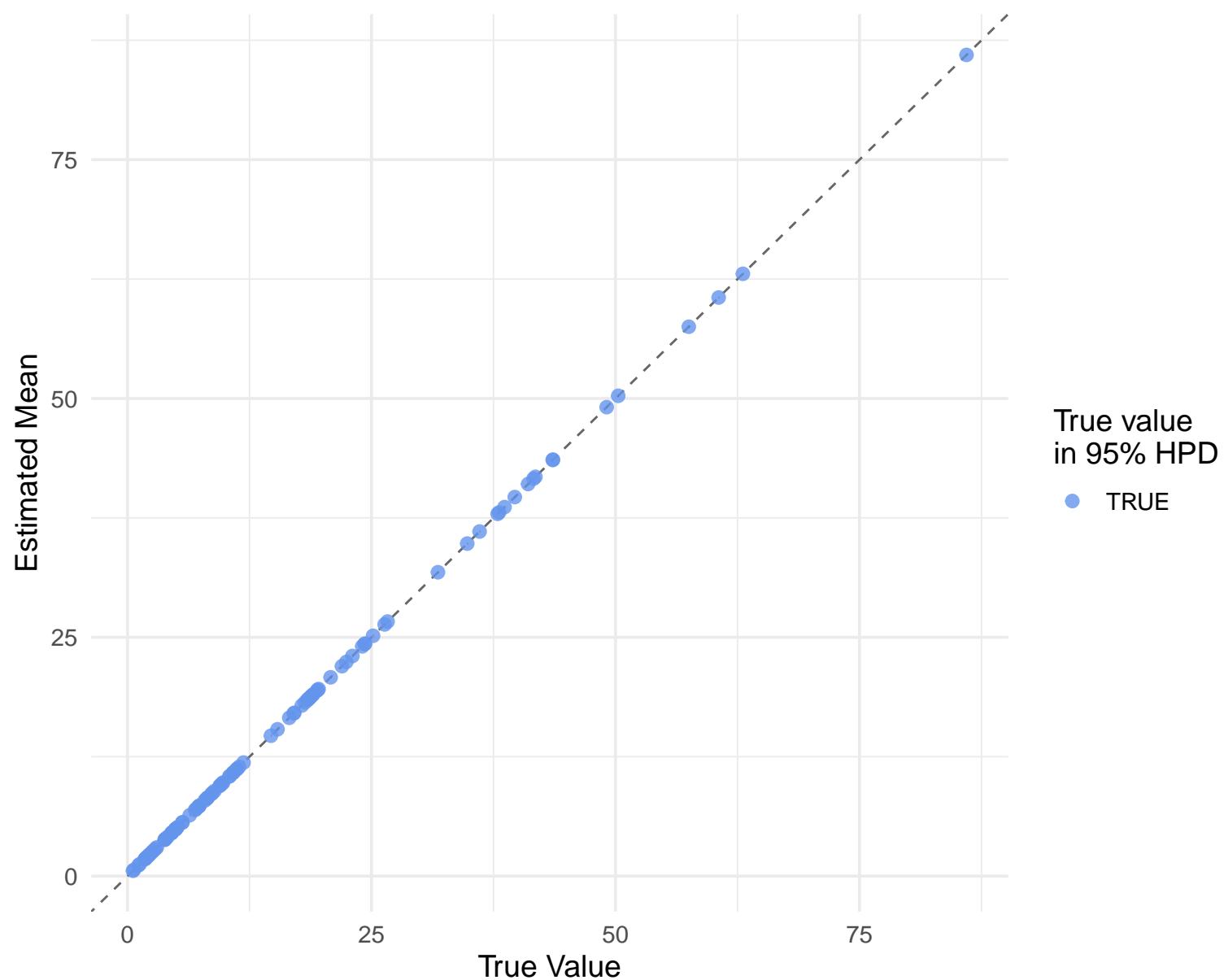
# tree.height

Coverage = 100.0%, Pearson's r = 1.000, N = 99



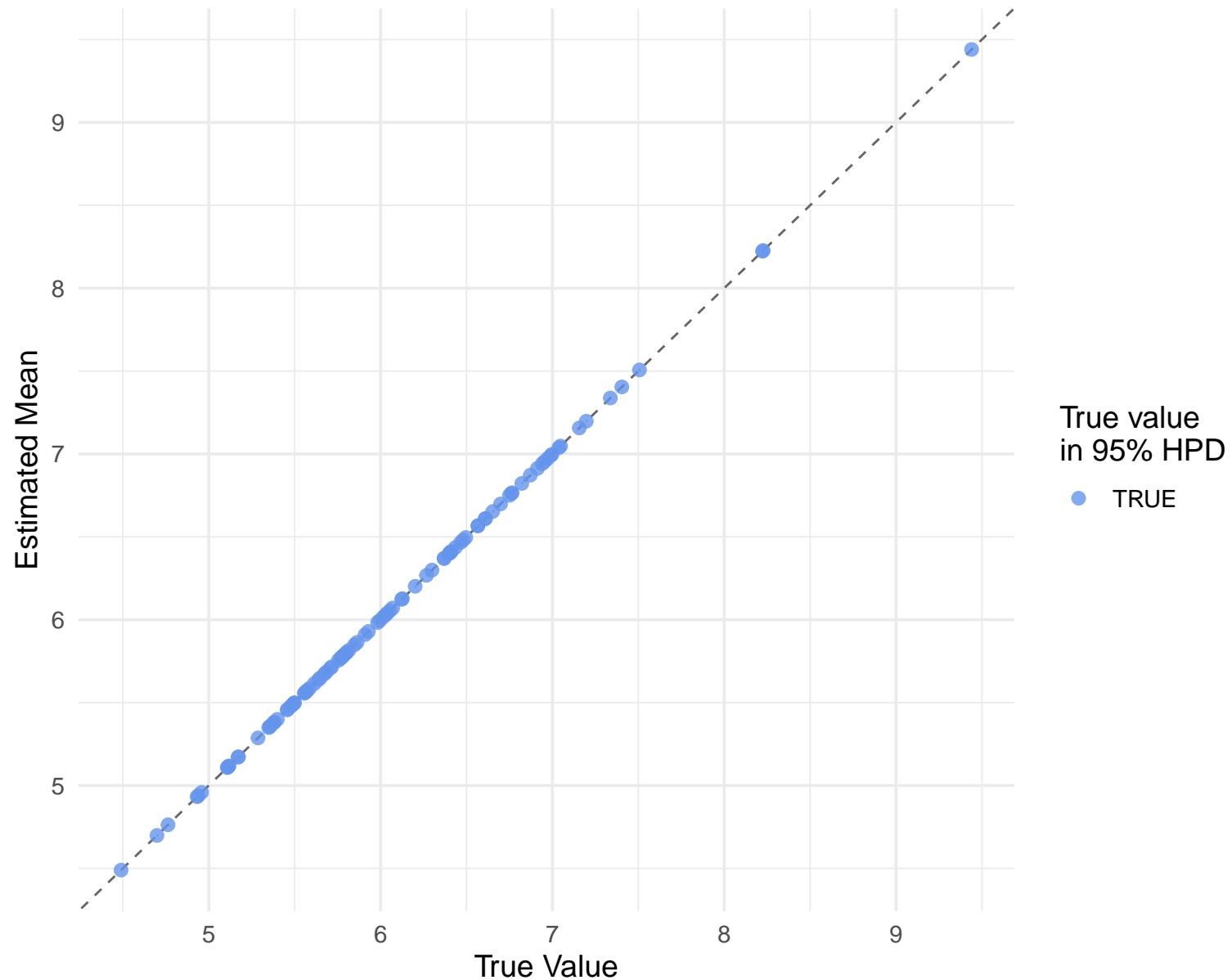
# tree.treeLength

Coverage = 100.0%, Pearson's r = 1.000, N = 99



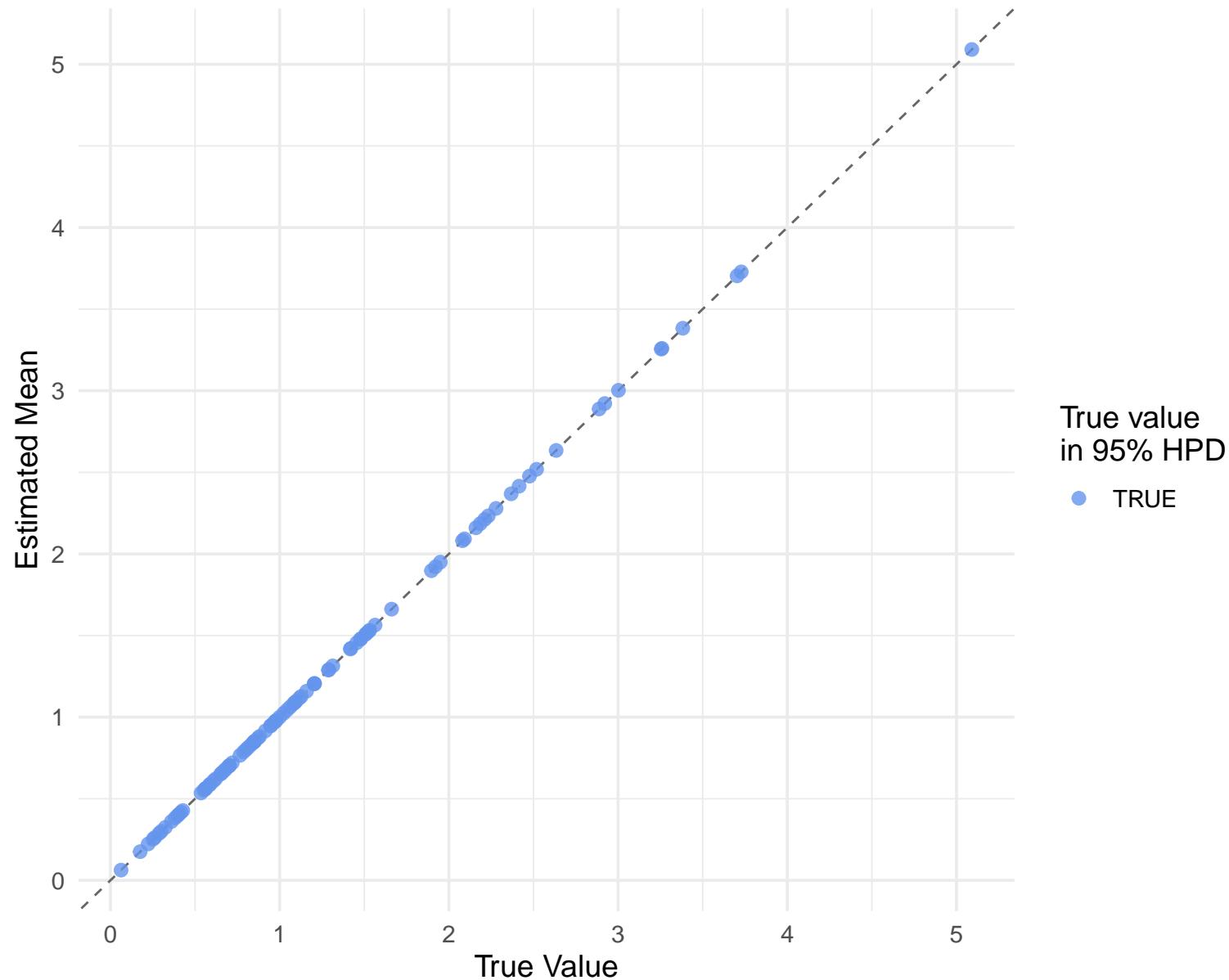
# lambda

Coverage = 100.0%, Pearson's r = 1.000, N = 99



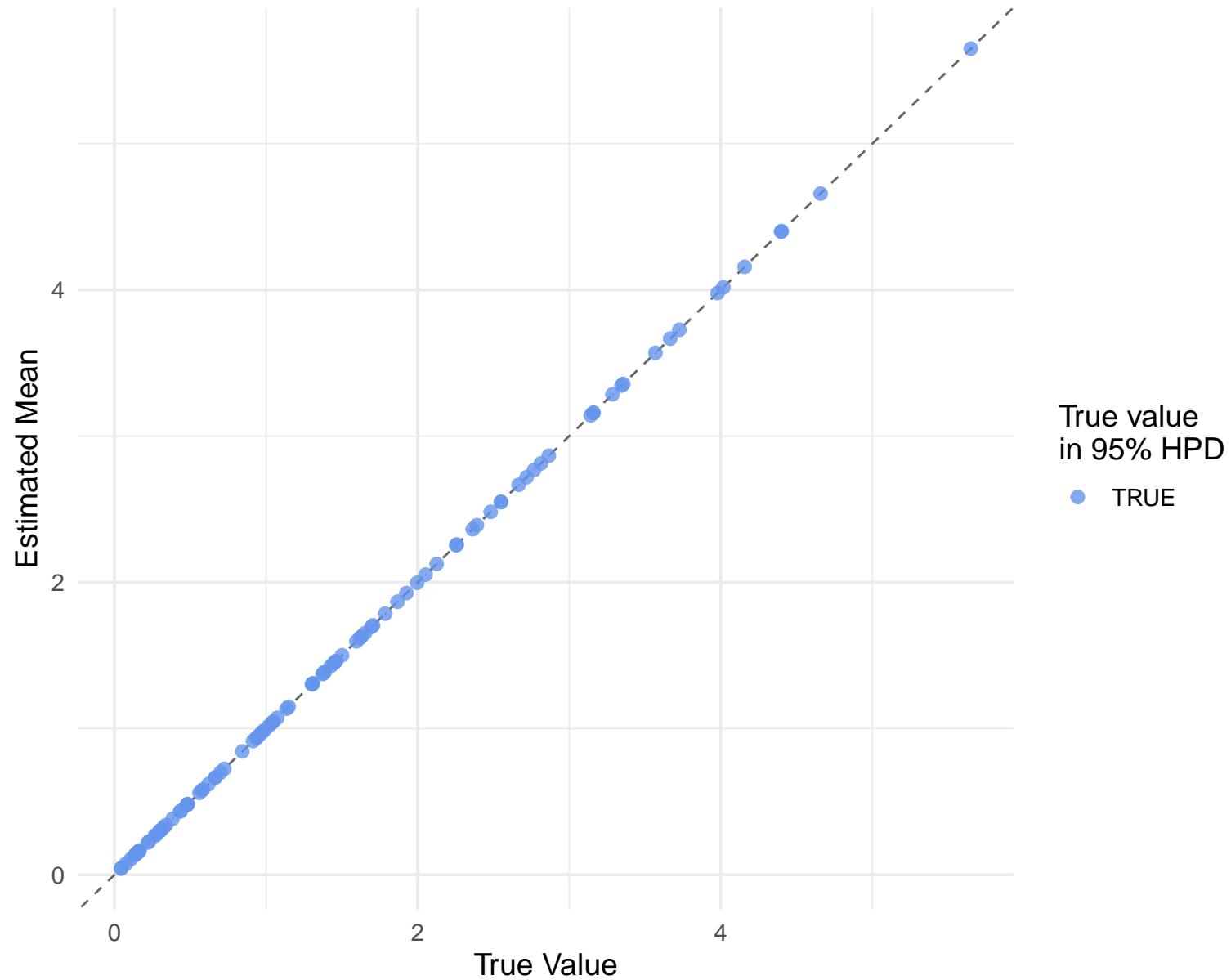
**mu**

Coverage = 100.0%, Pearson's r = 1.000, N = 99



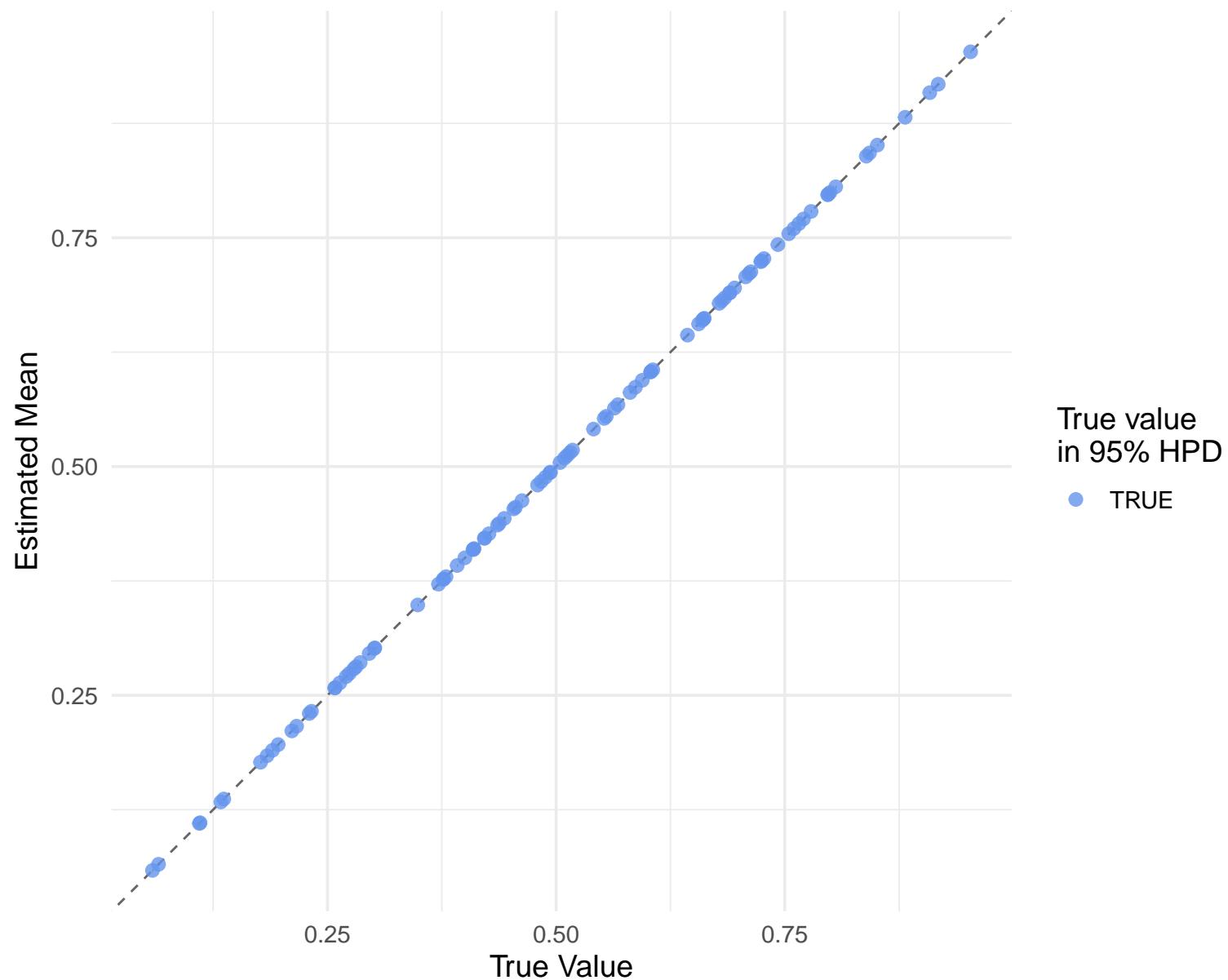
**psi**

Coverage = 100.0%, Pearson's r = 1.000, N = 99



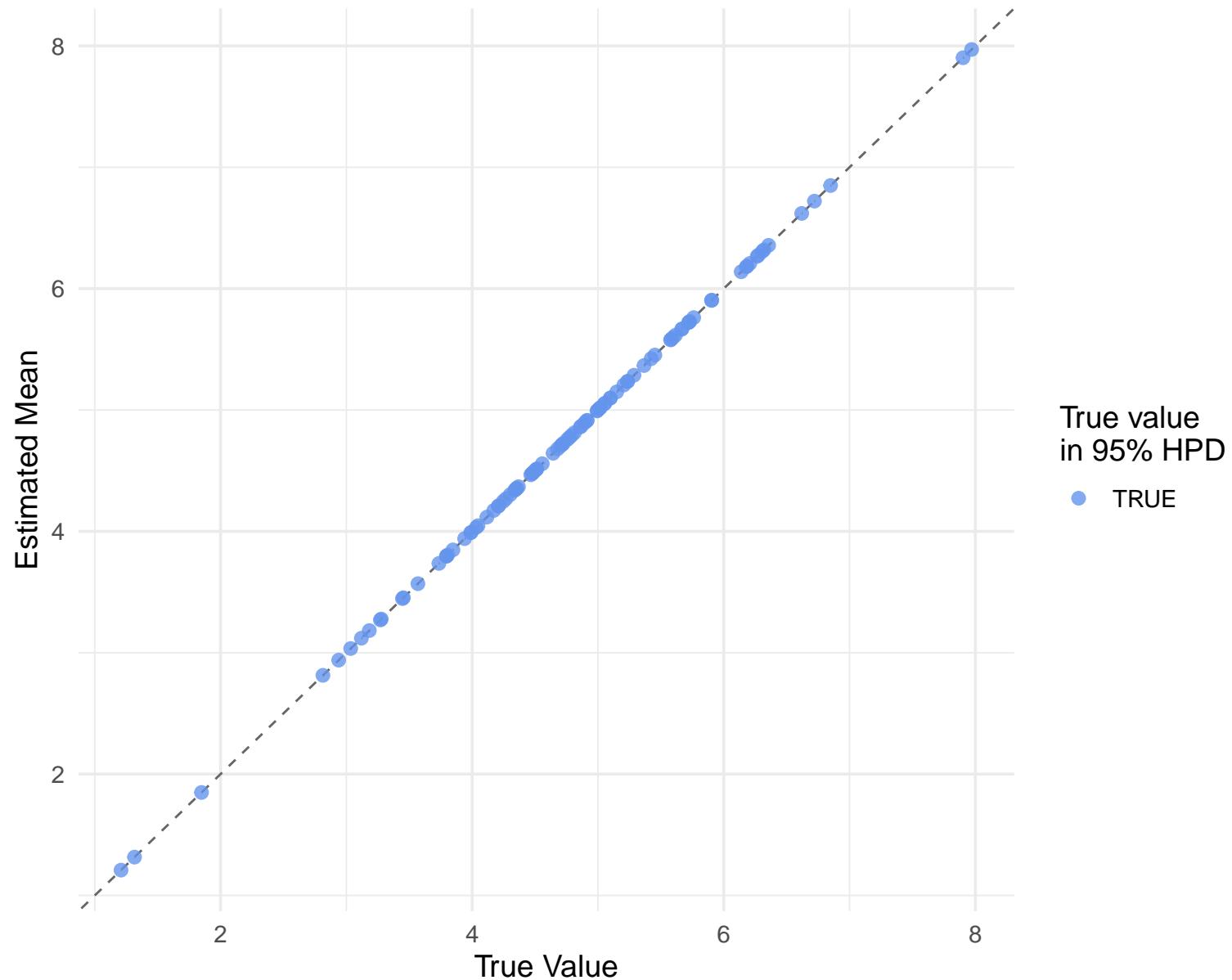
# rhoSampling

Coverage = 100.0%, Pearson's r = 1.000, N = 99



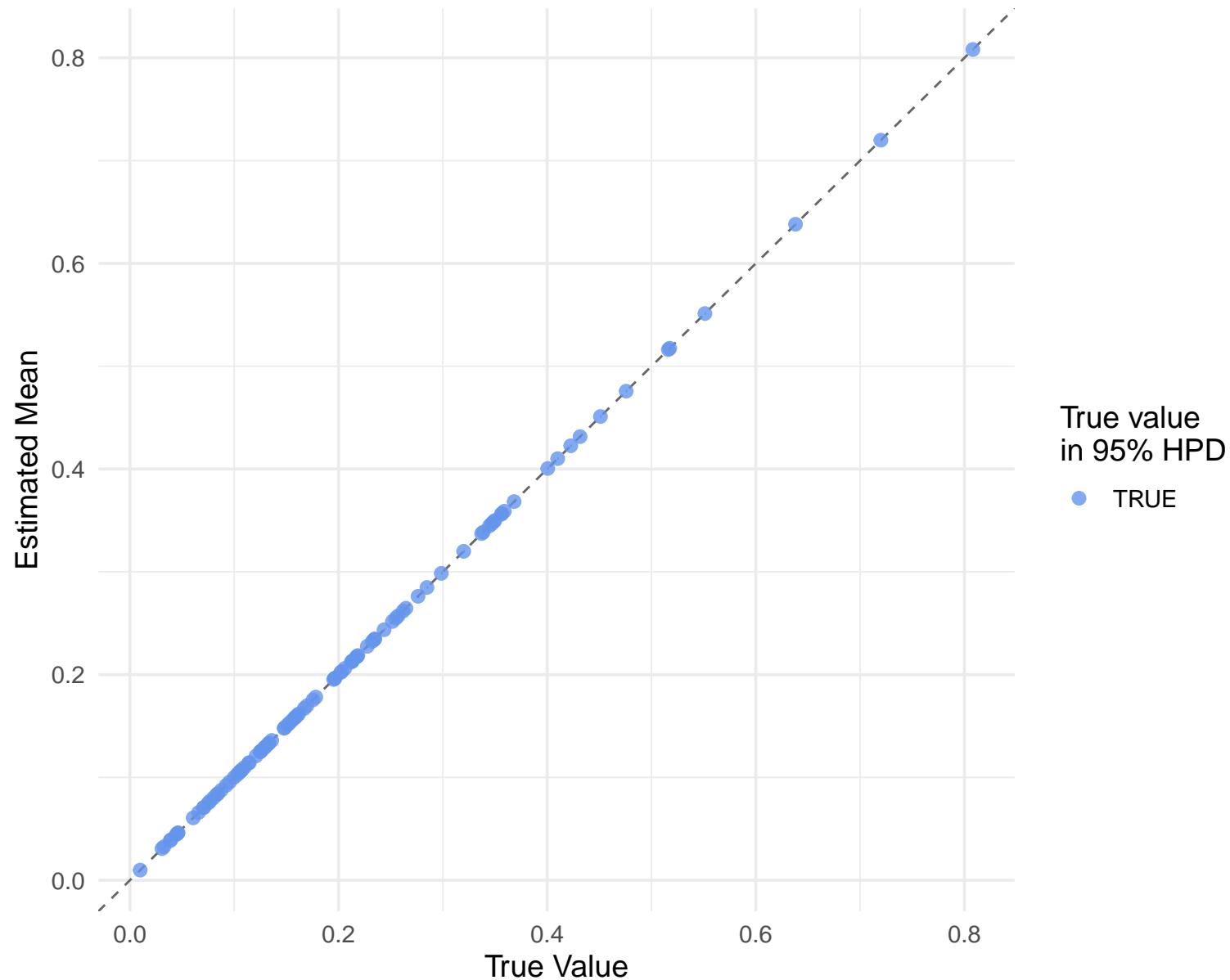
# netDiv

Coverage = 100.0%, Pearson's r = 1.000, N = 99



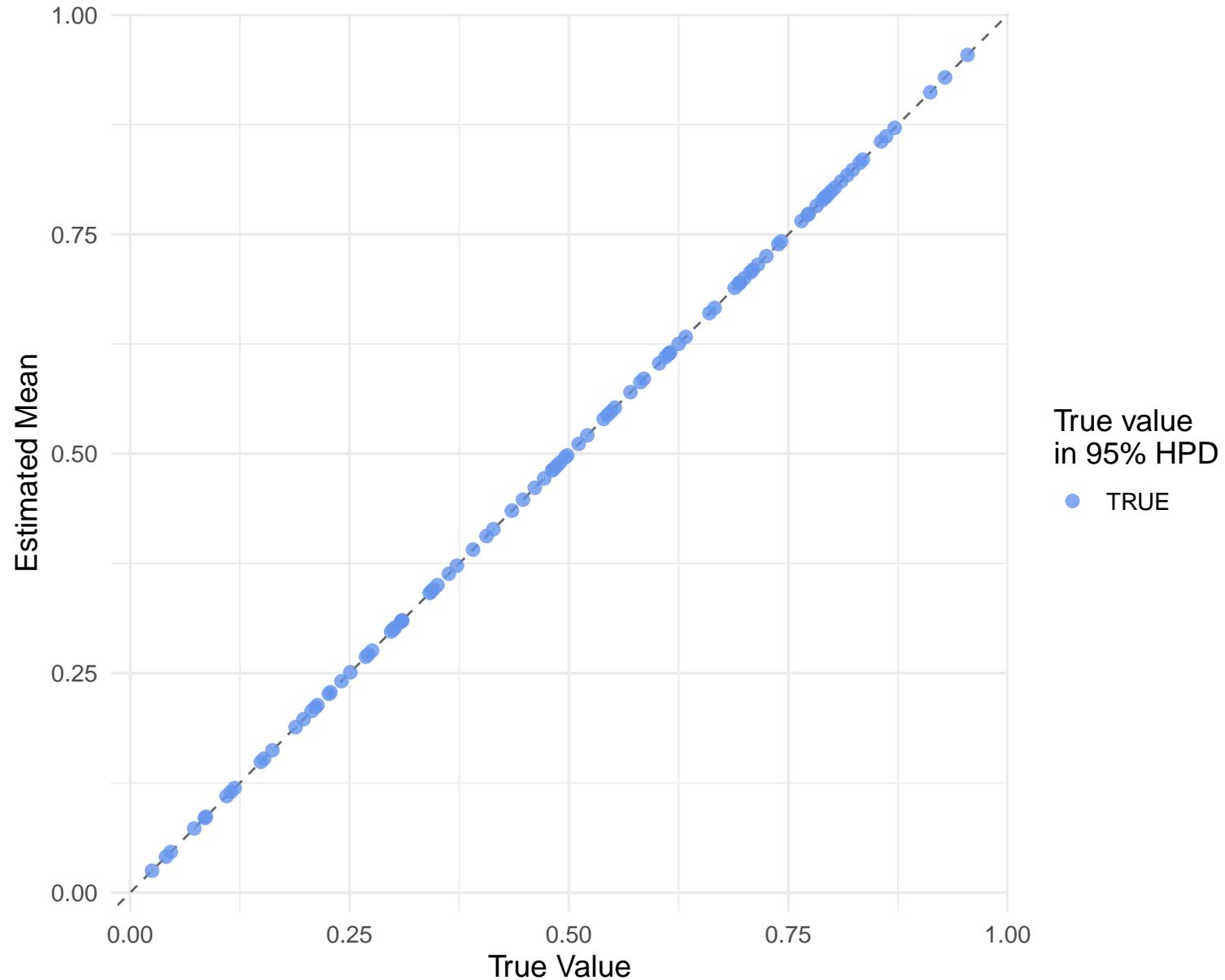
# turnover

Coverage = 100.0%, Pearson's r = 1.000, N = 99



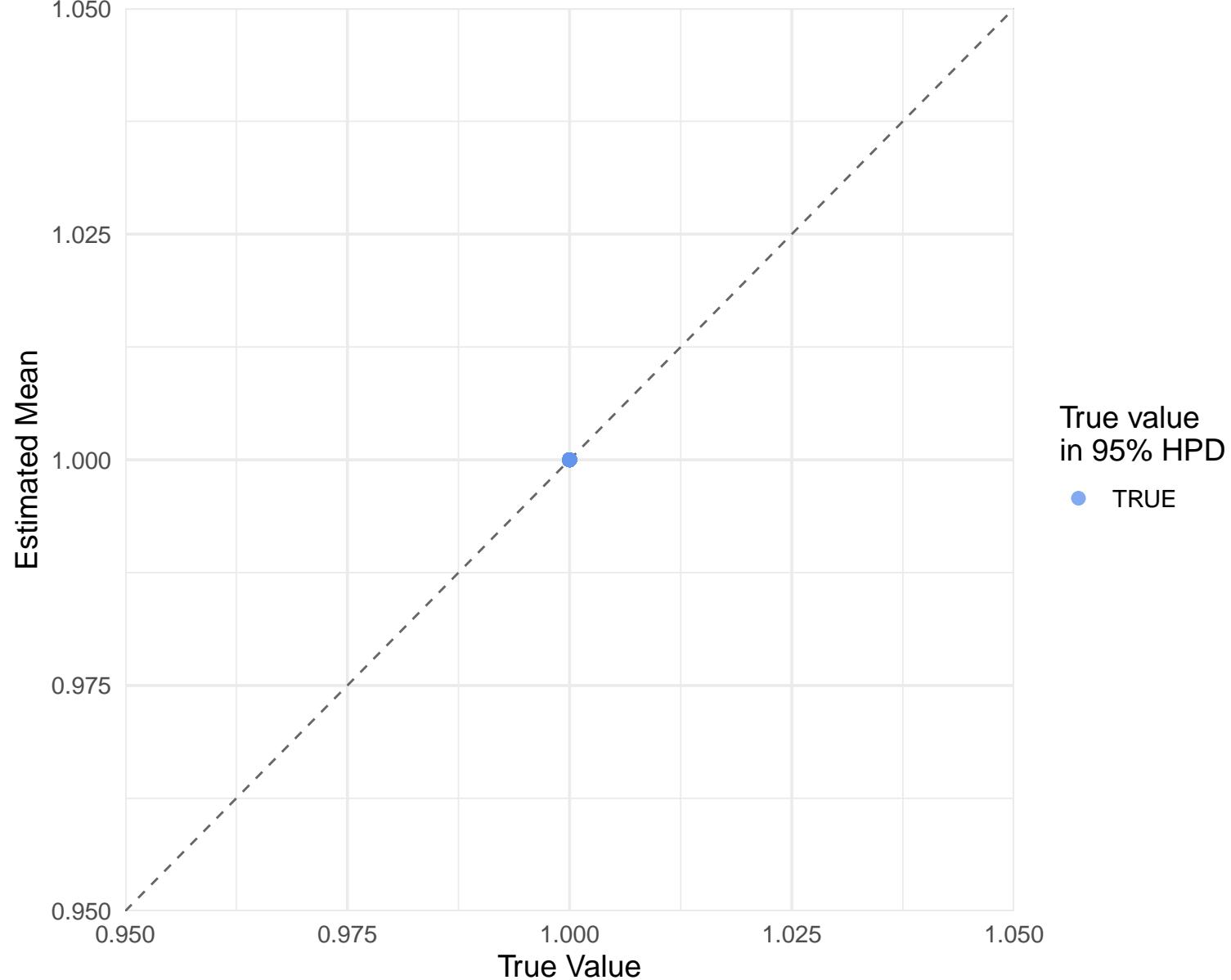
# samplingProportion

Coverage = 100.0%, Pearson's r = 1.000, N = 99



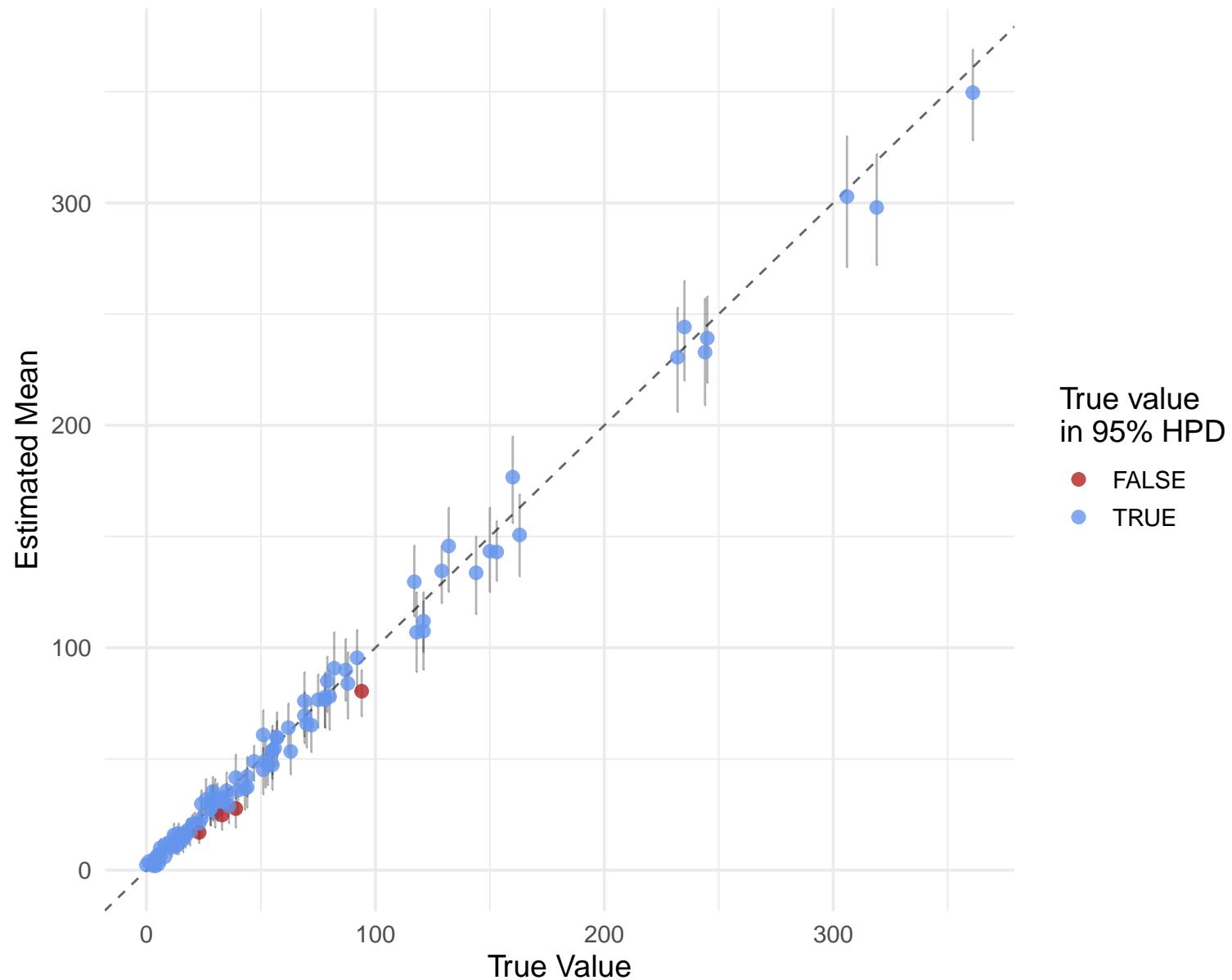
# origin

Coverage = 100.0%, Pearson's r = NA, N = 99



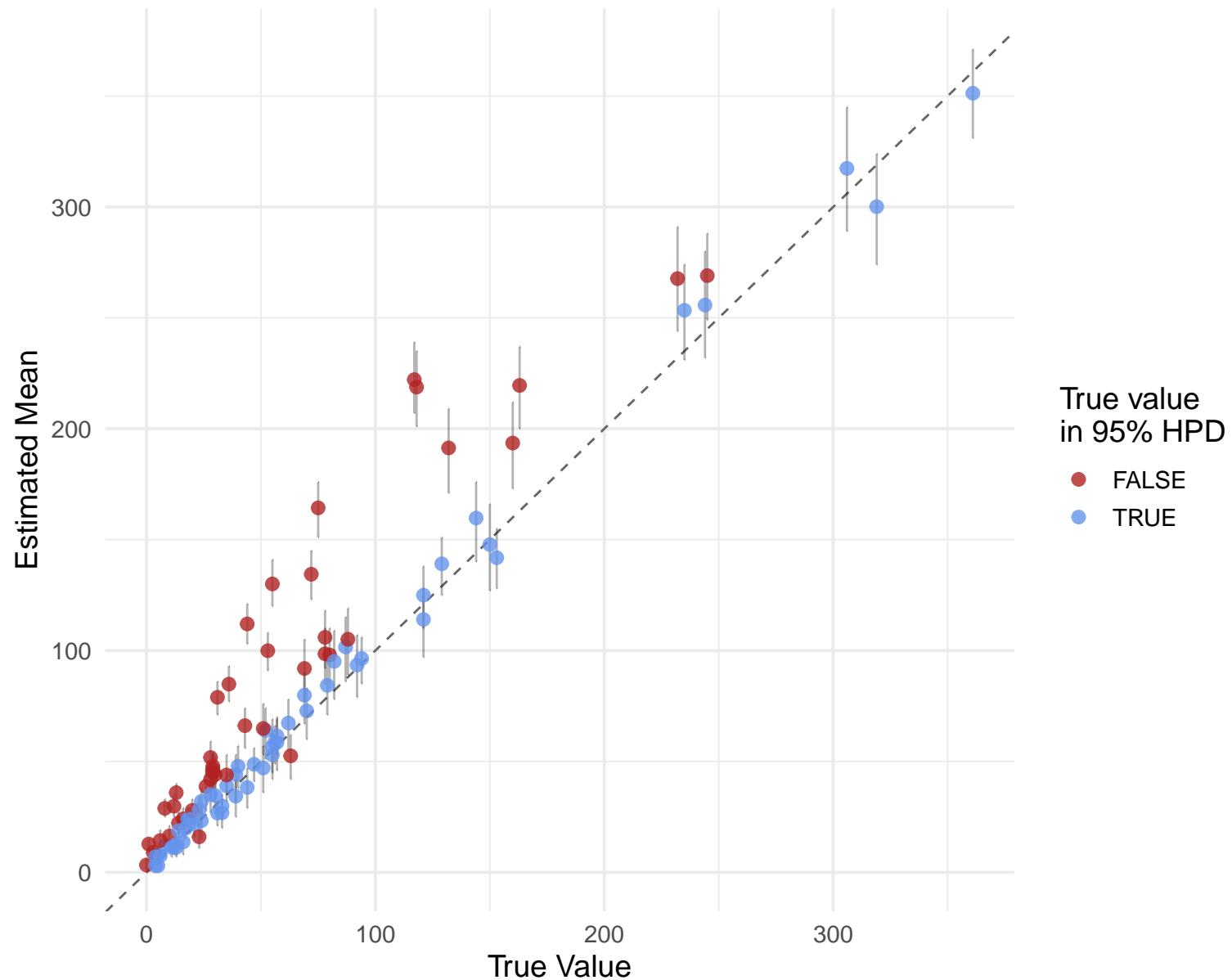
# estNrStubs.nstubs

Coverage = 96.0%, Pearson's r = 0.997, N = 99



# nrStubsSampledFromPrior\_spikePrior.nstubs

Coverage = 57.6%, Pearson's r = 0.957, N = 99



# nrStubsSampledFromPrior\_noSpikePrior.nstubs

Coverage = 97.0%, Pearson's r = 0.995, N = 99

