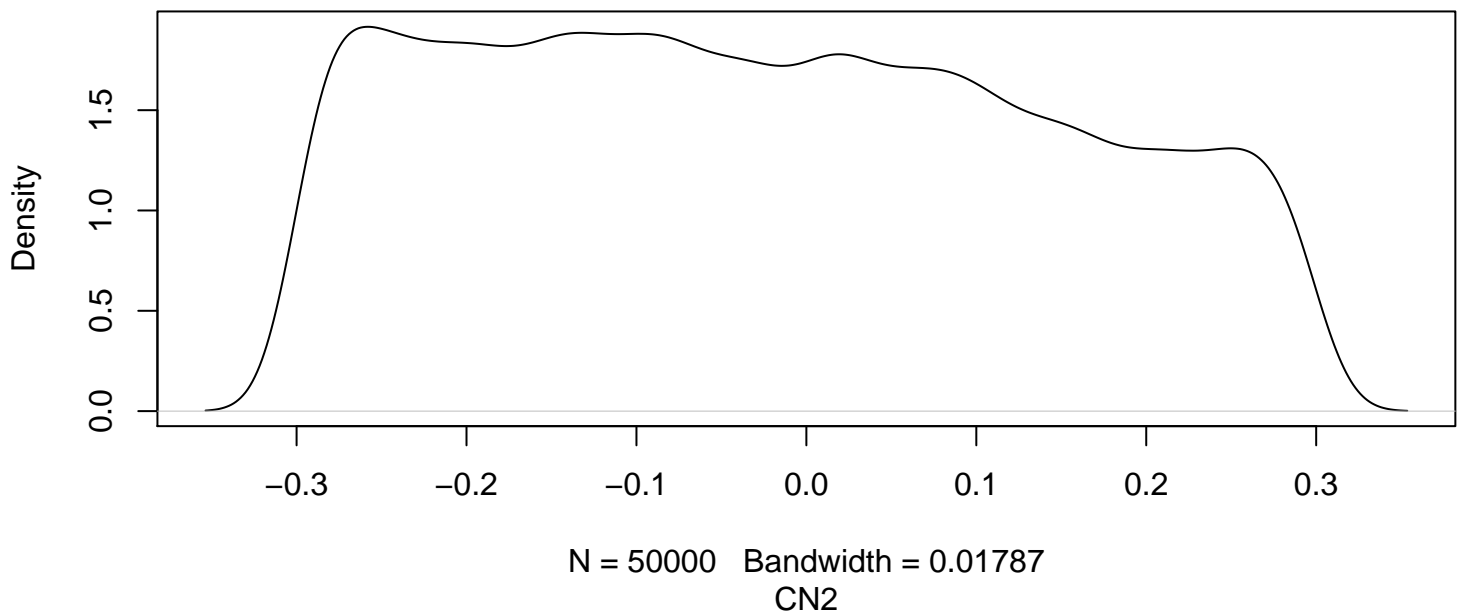
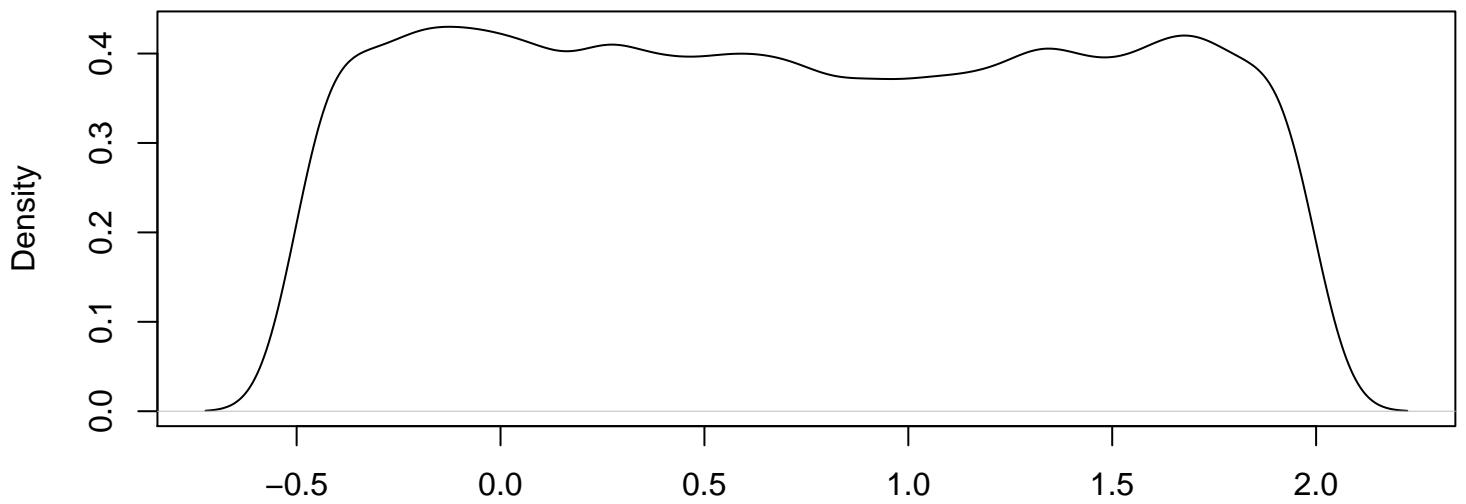


density.default(x = x, weights = wt_total)

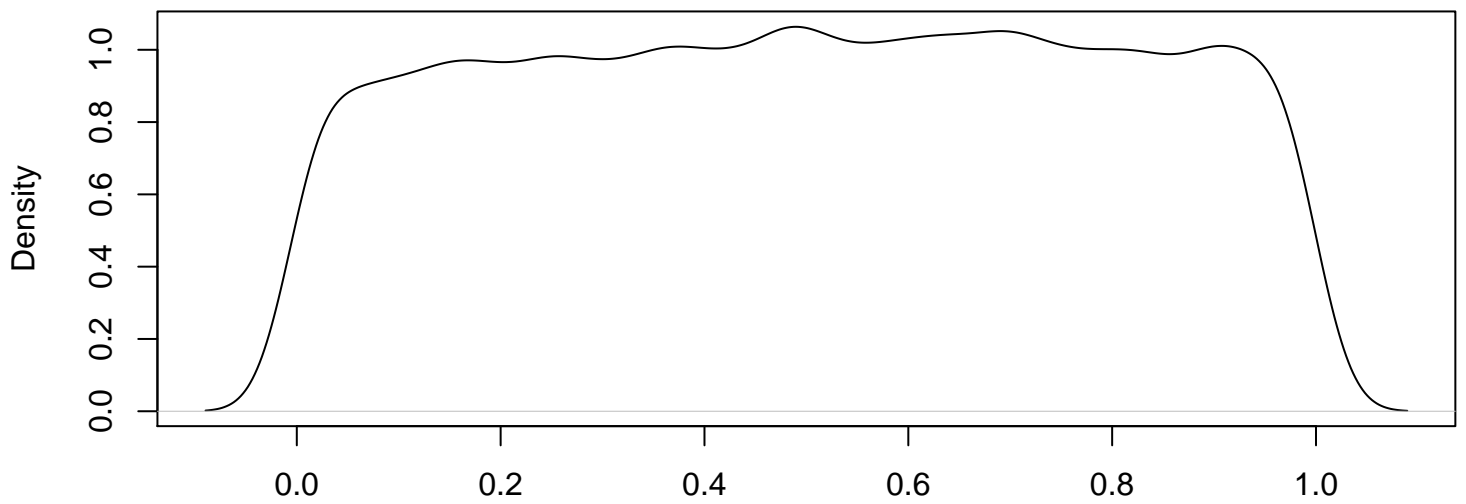


density.default(x = x, weights = wt_total)



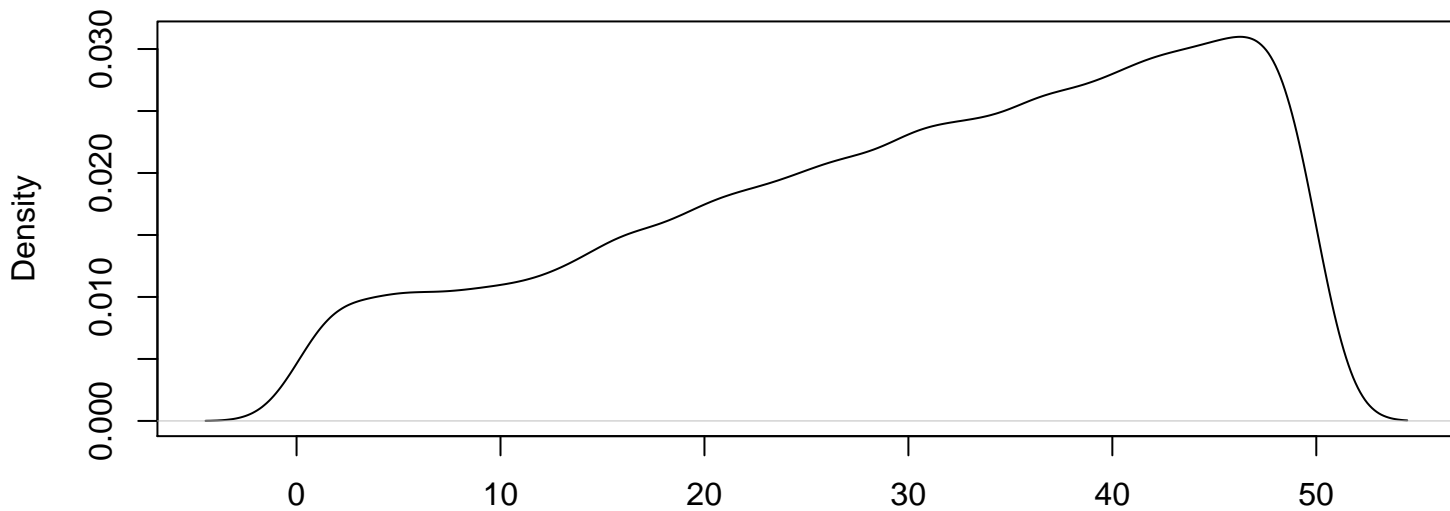
N = 50000 Bandwidth = 0.07455
GWQMN

density.default(x = x, weights = wt_total)



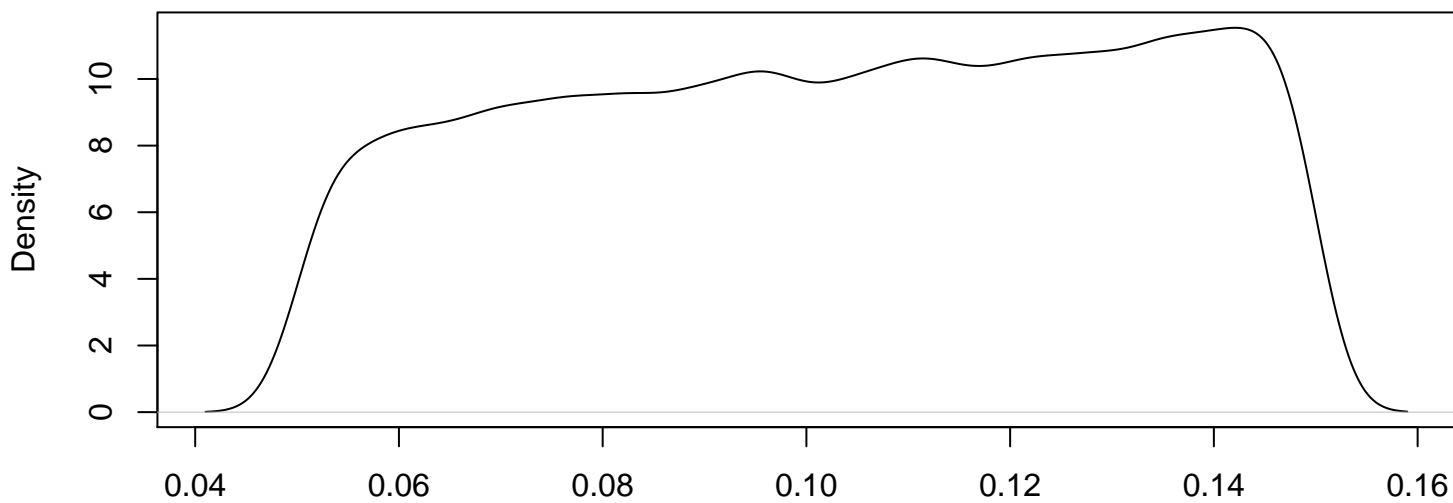
N = 50000 Bandwidth = 0.02986
ALPHA_BNK

density.default(x = x, weights = wt_total)



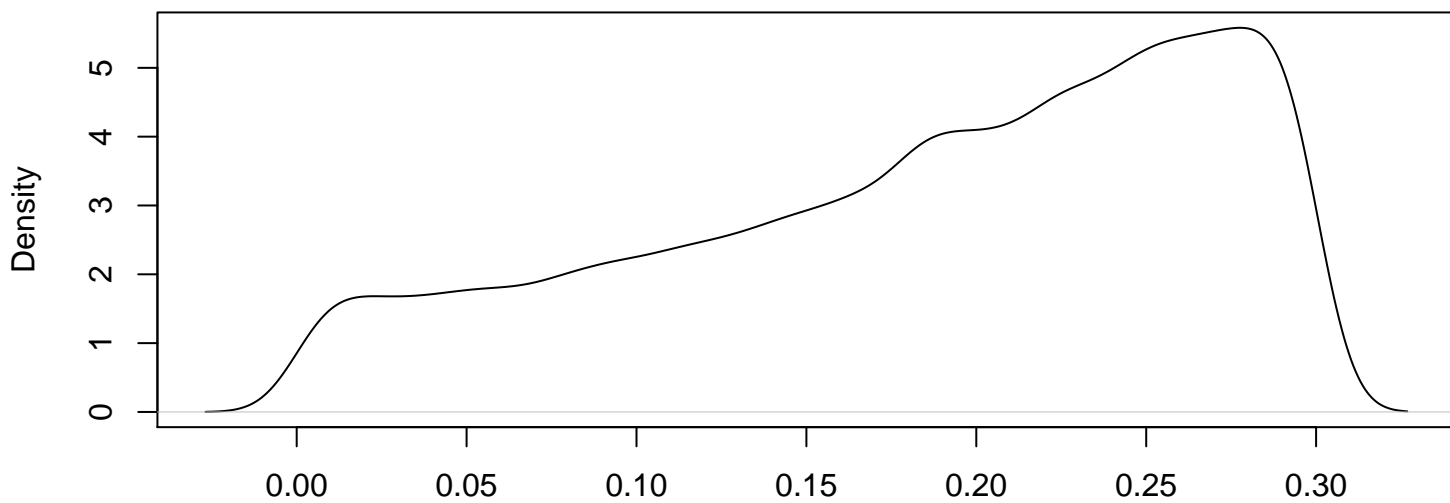
N = 50000 Bandwidth = 1.488
CH_K2

density.default(x = x, weights = wt_total)



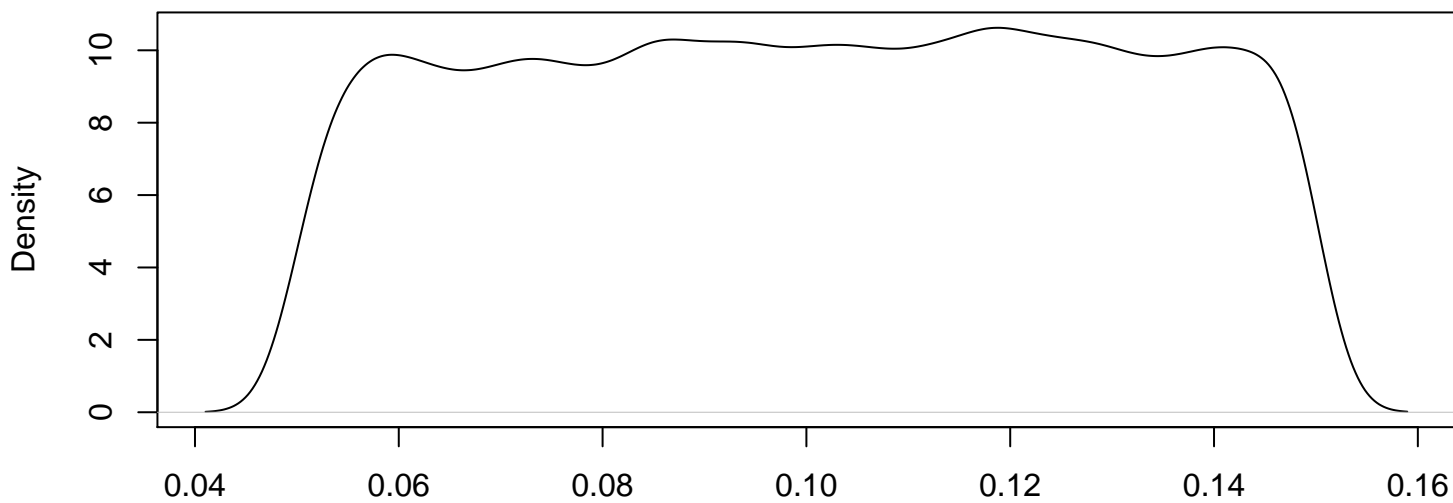
N = 50000 Bandwidth = 0.002996
CH_N2

density.default(x = x, weights = wt_total)



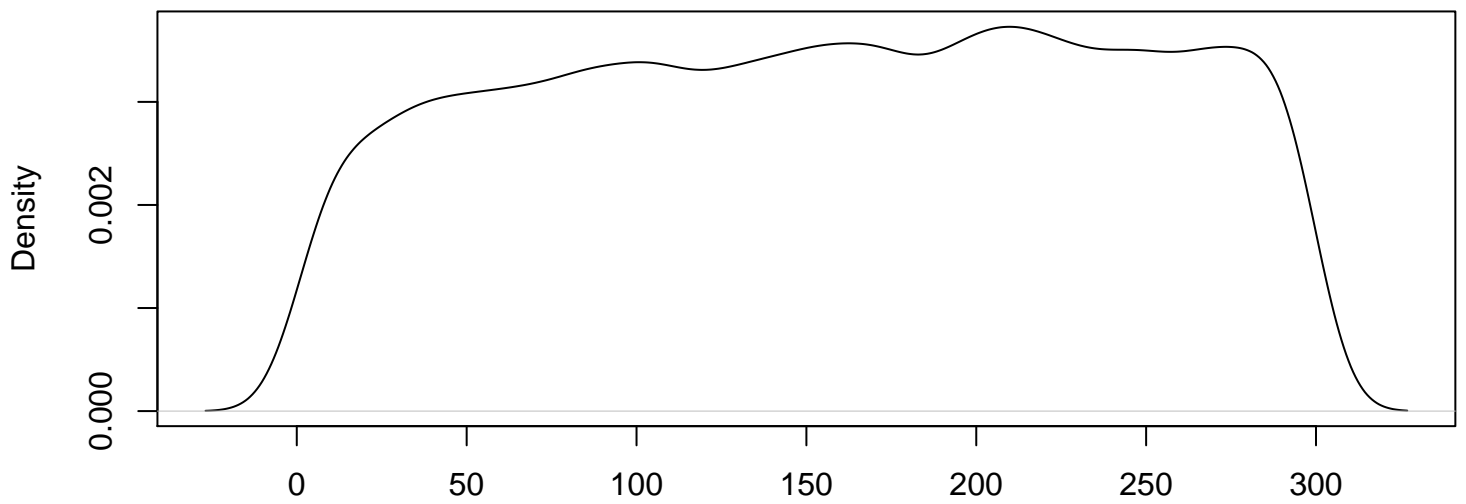
N = 50000 Bandwidth = 0.008944
TRNSRCH

density.default(x = x, weights = wt_total)



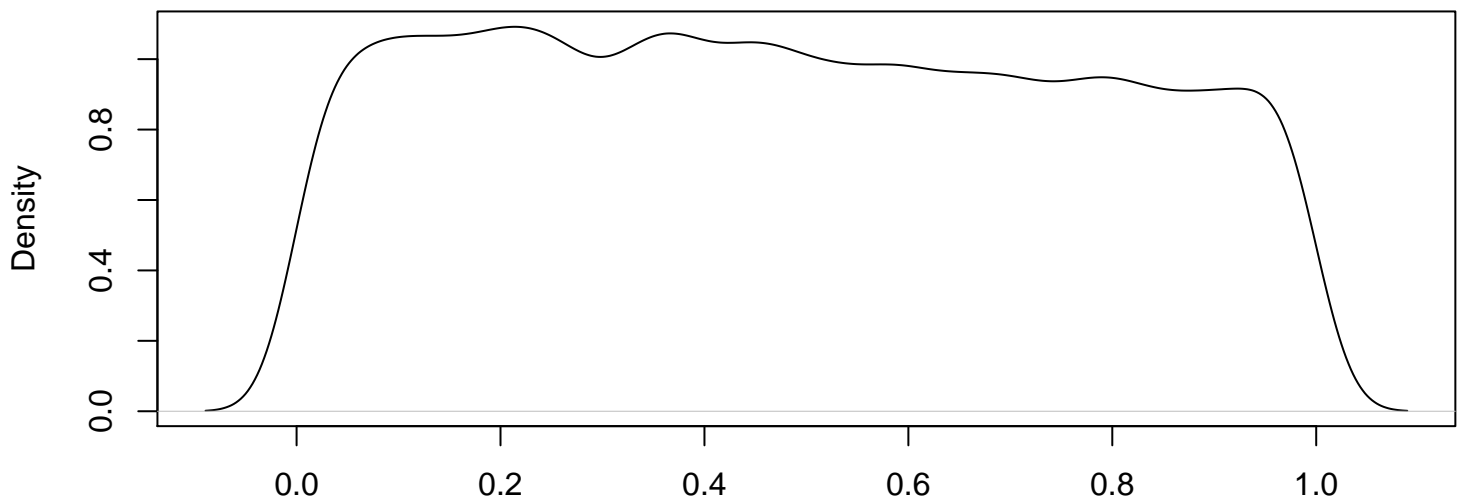
N = 50000 Bandwidth = 0.002989
CH_N1

density.default(x = x, weights = wt_total)



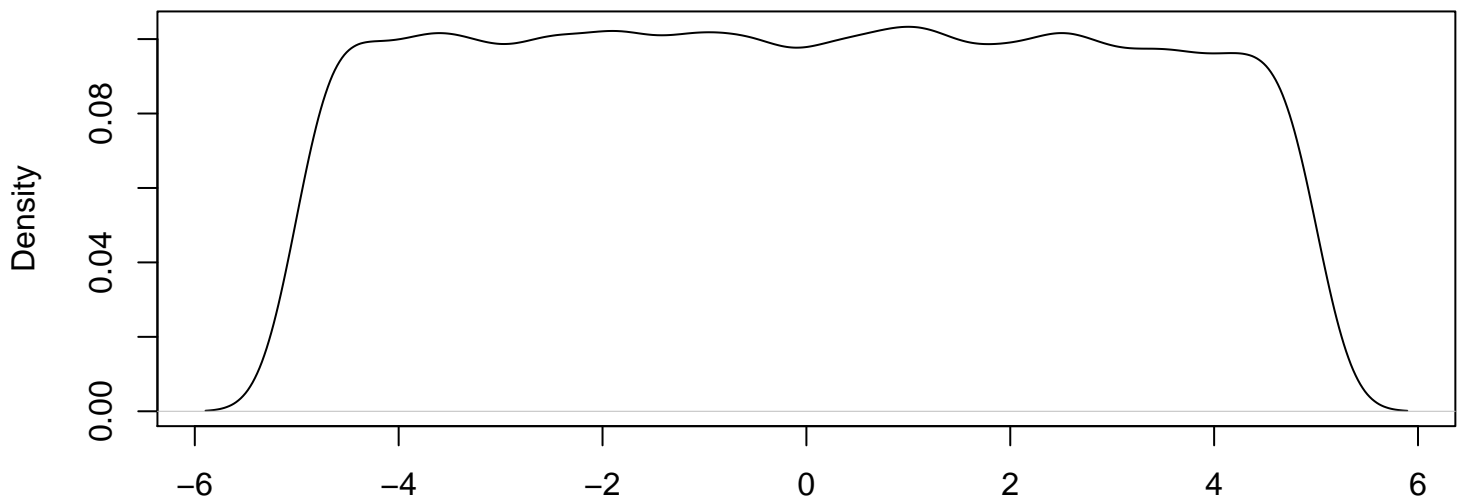
N = 50000 Bandwidth = 8.958
CH_K1

density.default(x = x, weights = wt_total)



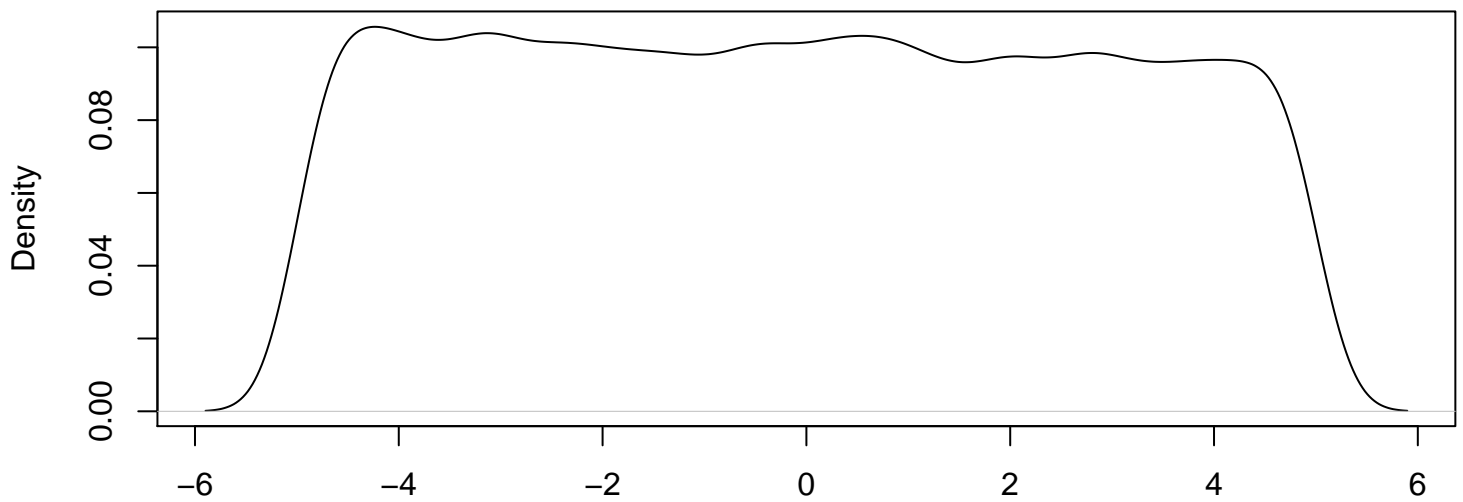
N = 50000 Bandwidth = 0.02977
RCHRG_DP

density.default(x = x, weights = wt_total)



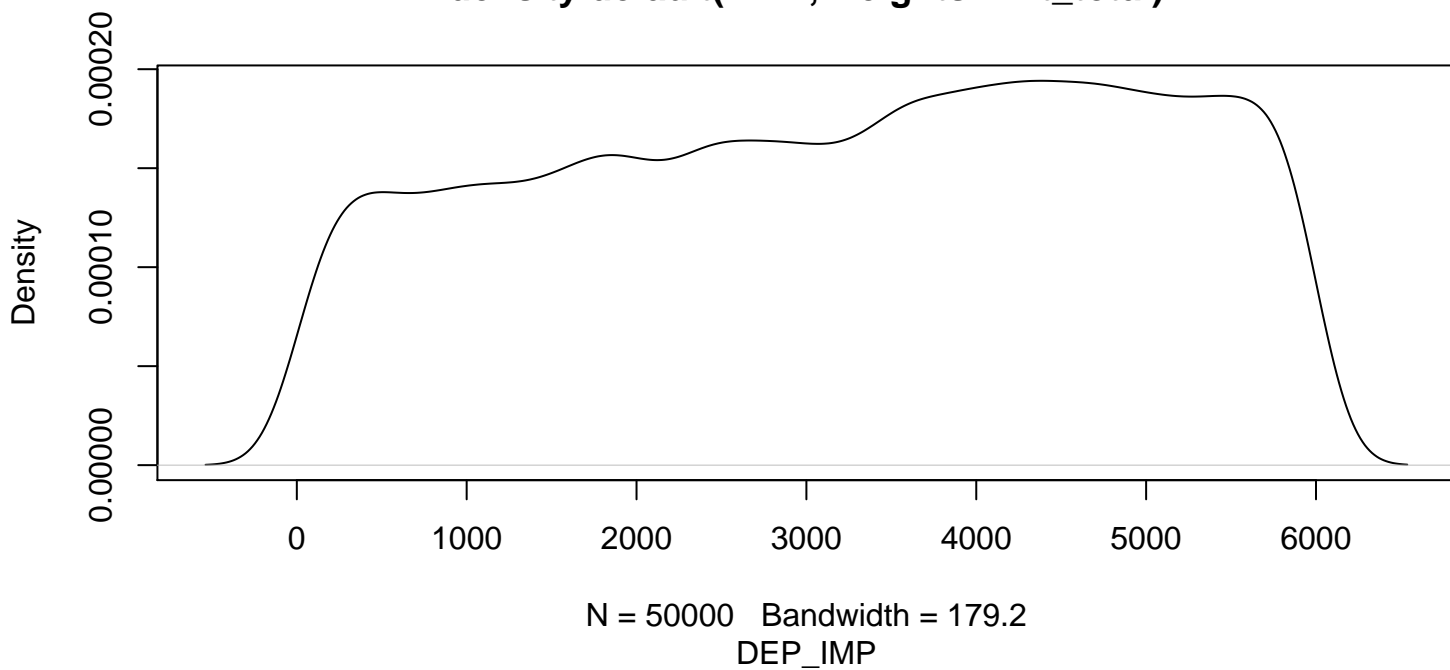
N = 50000 Bandwidth = 0.2983
SFTMP

density.default(x = x, weights = wt_total)

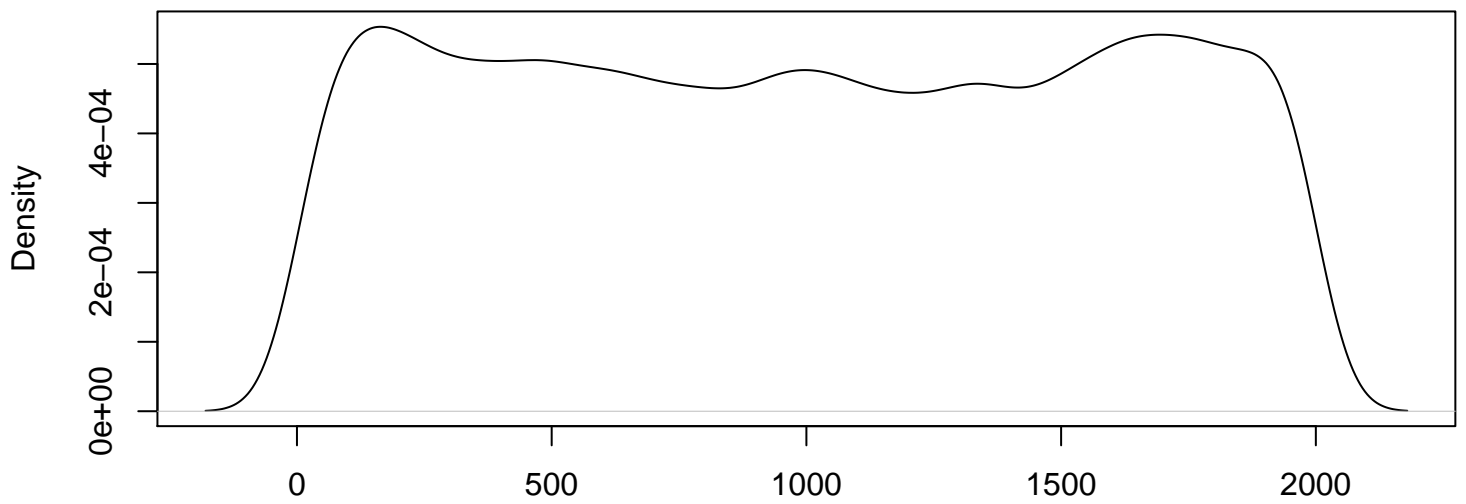


N = 50000 Bandwidth = 0.2989
SMTMP

density.default(x = x, weights = wt_total)

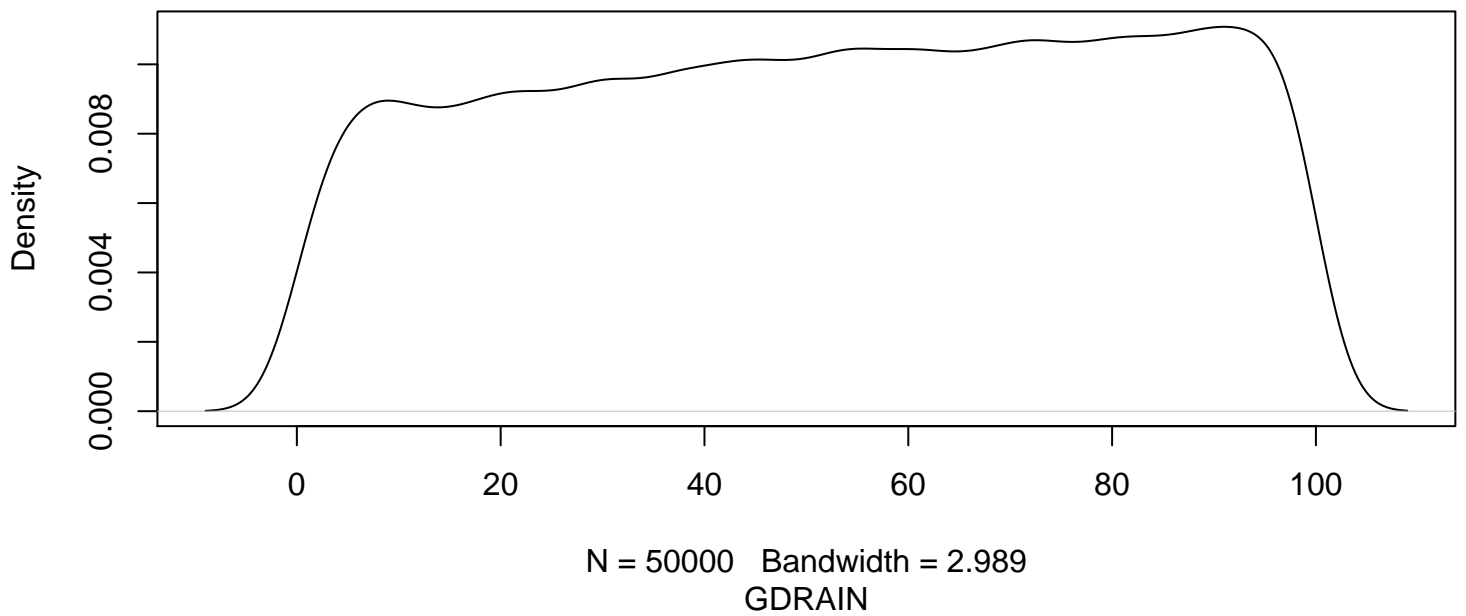


density.default(x = x, weights = wt_total)

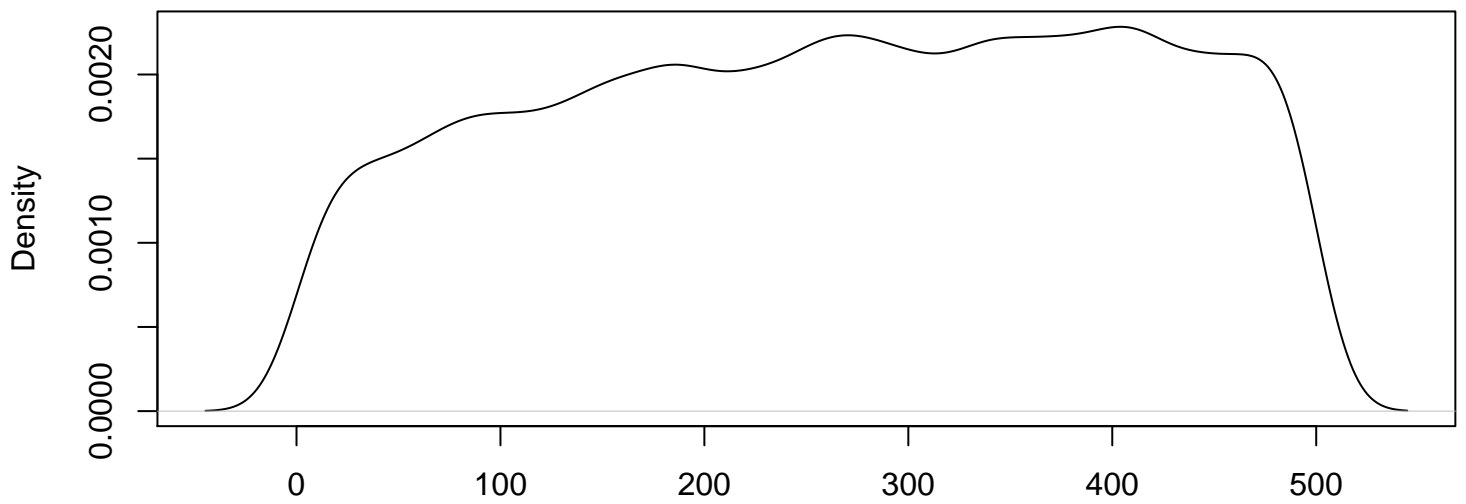


N = 50000 Bandwidth = 59.86
DDRAIN

density.default(x = x, weights = wt_total)

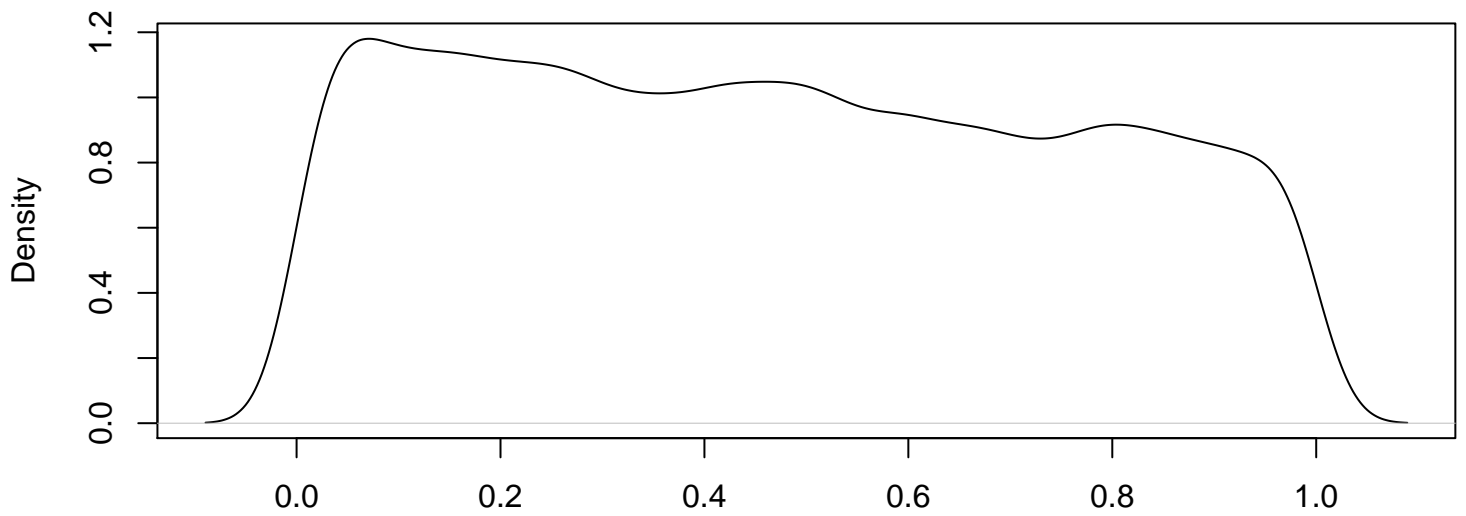


density.default(x = x, weights = wt_total)



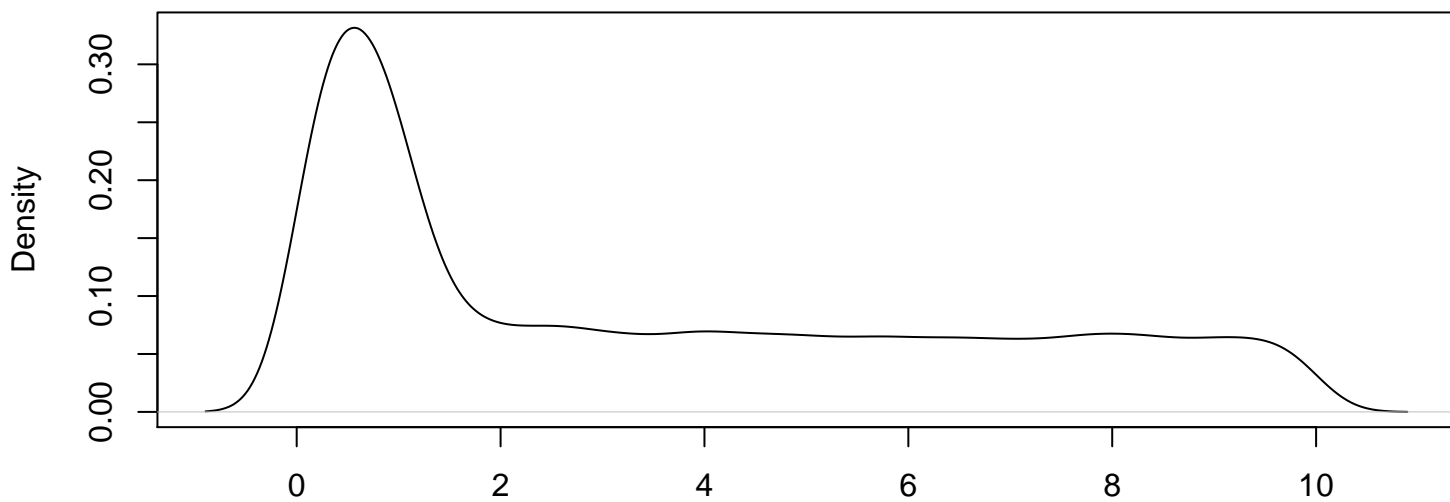
N = 50000 Bandwidth = 14.89
BACTKDQ

density.default(x = x, weights = wt_total)



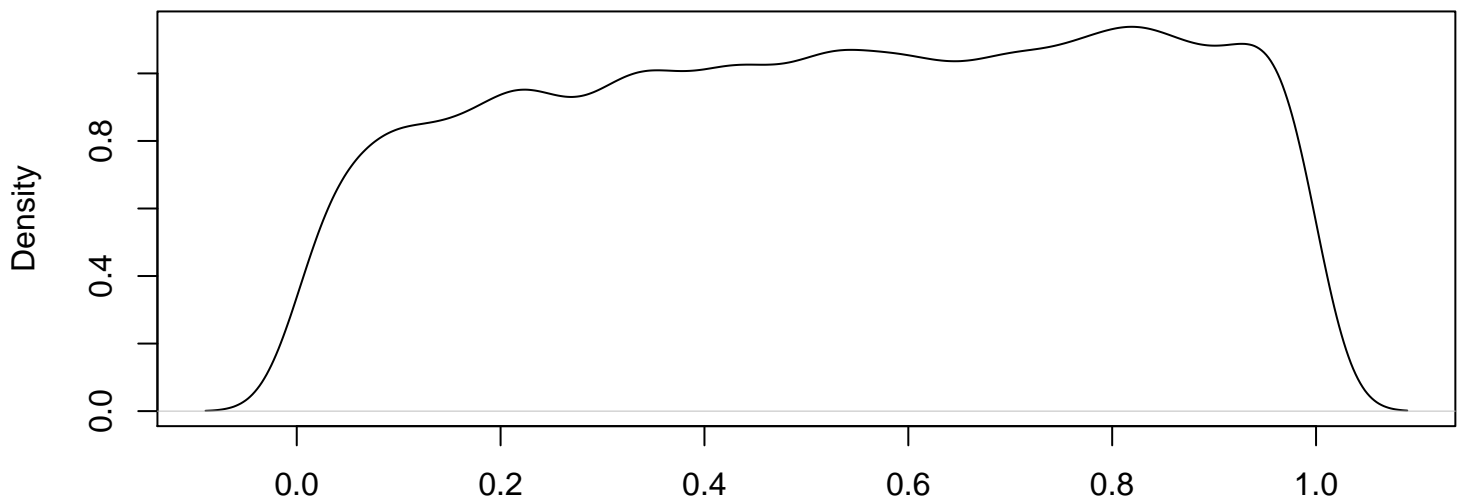
N = 50000 Bandwidth = 0.02978
BACT_SWF

density.default(x = x, weights = wt_total)



N = 50000 Bandwidth = 0.2982
THBACT

density.default(x = x, weights = wt_total)



N = 50000 Bandwidth = 0.02984
WDPRCH