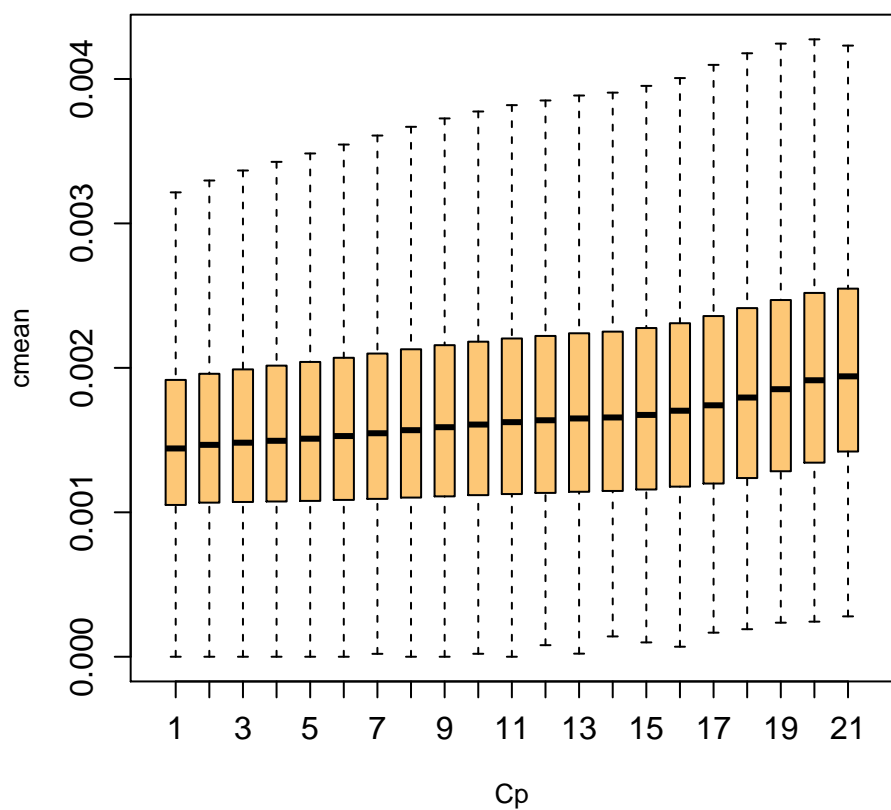
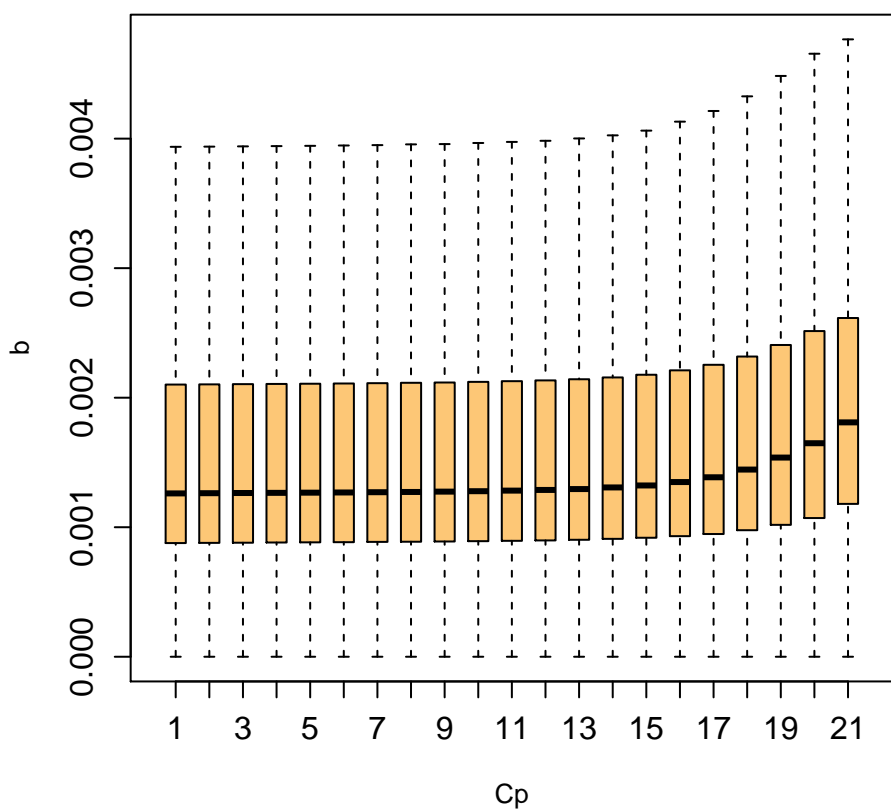


ms20b_donor_centric_PCAWG_Hg19_TTC RefSeq.16, Intron, Intergenic, sig3pcentric, -1, 8, Nucleosome
chr22:163279356del(100%)100(100)37663
22chr22:163279356del(100%)100(100)37663
chr1:15327368, 123212, 297759, 7149484, 61322, 411579, 3786434, 2850464, 2550801, 1563200, 1161818, 844579, 697945, 284190, 177321, 38822, 44852, 13205
chr1:15327368, 123212, 297759, 7149484, 61322, 411579, 3786434, 2850464, 2550801, 1563200, 1161818, 844579, 697945, 284190, 177321, 38822, 44852, 13205
chr1:15327368, 123212, 297759, 7149484, 61322, 411579, 3786434, 2850464, 2550801, 1563200, 1161818, 844579, 697945, 284190, 177321, 38822, 44852, 13205
chr1:15327368, 123212, 297759, 7149484, 61322, 411579, 3786434, 2850464, 2550801, 1563200, 1161818, 844579, 697945, 284190, 177321, 38822, 44852, 13205



Box plot showing the distribution of CSD (Y-axis, ranging from 0.0000 to 0.0025) across different values of C_p (X-axis, ranging from 1 to 21). The plot displays the median, quartiles, and range (whiskers) for each C_p value. The median CSD is approximately 0.00055 for all C_p values. The interquartile range (IQR) is approximately 0.0004 for $C_p=1$ and decreases slightly as C_p increases. The whiskers extend from approximately 0.0001 to 0.0024 for $C_p=1$ and decrease as C_p increases.