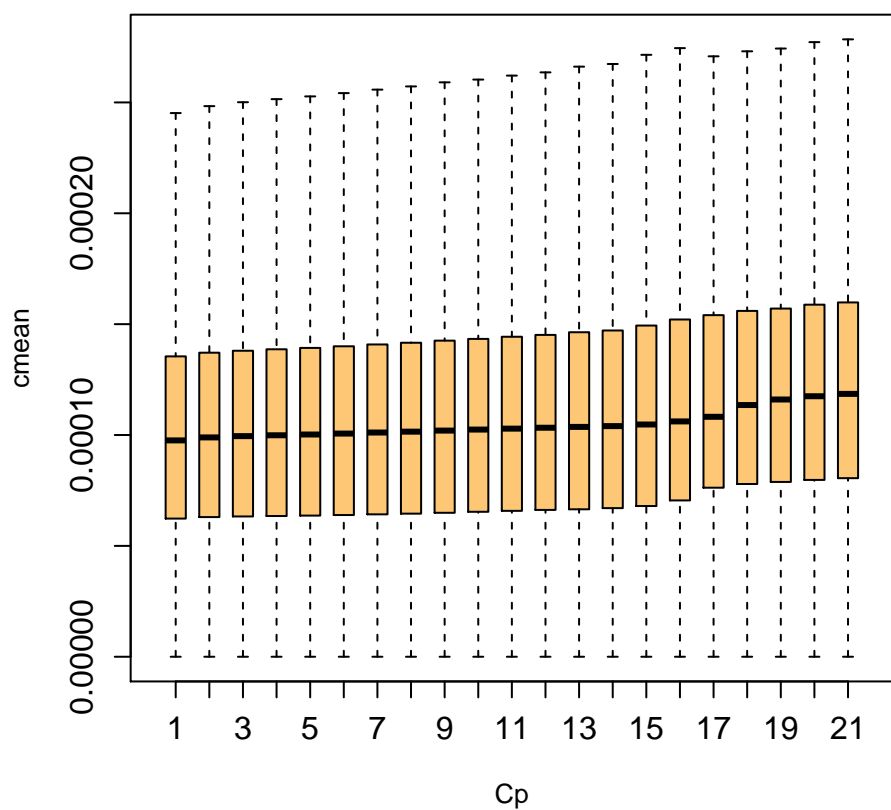
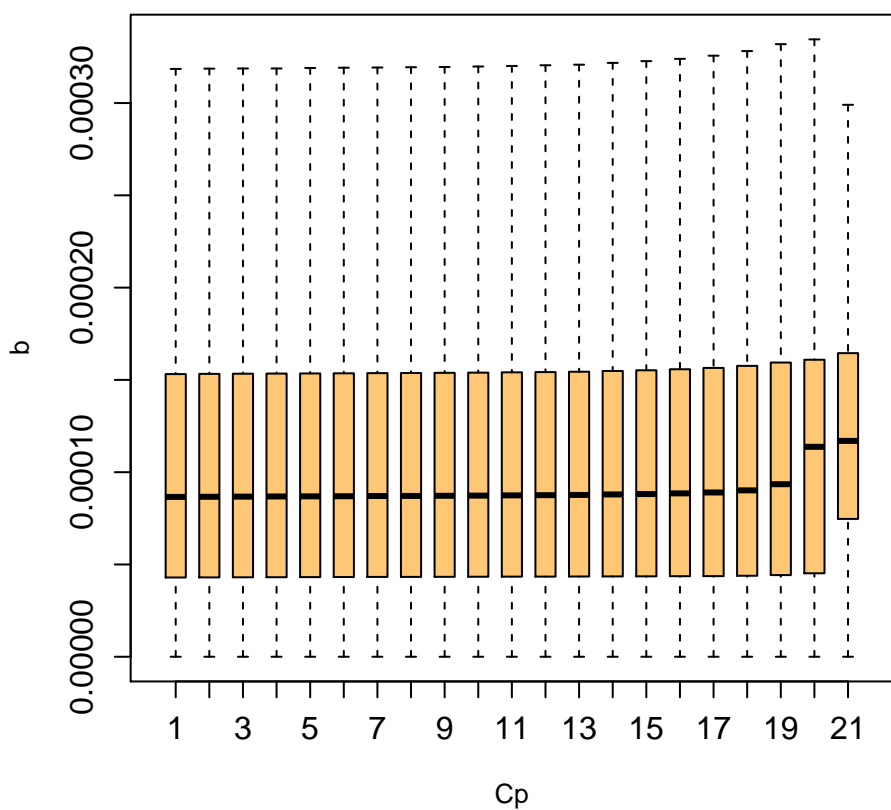


min2M, donor_centric_CAIWG, Hg19, TloC, RefSeqMMR1, exon_intron_intergenic_intronic_cass, sig1percivmis, 1000revs_HmNetman
cnaaa, 16c43d719vad100/jout100240719
23chr, chr1,chr2,chr3,chr4,chr5,chr6,chr7,chr8,chr9,chr10,chr11,chr12,chr13,chr14,chr15,chr16,chr17,chr18,chr19,chr20,chr21,chr22,chr23,chr24,chr25,chr26,chr27,chr28,chr29,chr30,chr31,chr32,chr33,chr34,chr35,chr36,chr37,chr38,chr39,chr40,chr41,chr42,chr43,chr44,chr45,chr46,chr47,chr48,chr49,chr50,chr51,chr52,chr53,chr54,chr55,chr56,chr57,chr58,chr59,chr60,chr61,chr62,chr63,chr64,chr65,chr66,chr67,chr68,chr69,chr70,chr71,chr72,chr73,chr74,chr75,chr76,chr77,chr78,chr79,chr80,chr81,chr82,chr83,chr84,chr85,chr86,chr87,chr88,chr89,chr90,chr91,chr92,chr93,chr94,chr95,chr96,chr97,chr98,chr99,chr100,chr101,chr102,chr103,chr104,chr105,chr106,chr107,chr108,chr109,chr110,chr111,chr112,chr113,chr114,chr115,chr116,chr117,chr118,chr119,chr120,chr121,chr122,chr123,chr124,chr125,chr126,chr127,chr128,chr129,chr130,chr131,chr132,chr133,chr134,chr135,chr136,chr137,chr138,chr139,chr140,chr141,chr142,chr143,chr144,chr145,chr146,chr147,chr148,chr149,chr150,chr151,chr152,chr153,chr154,chr155,chr156,chr157,chr158,chr159,chr160,chr161,chr162,chr163,chr164,chr165,chr166,chr167,chr168,chr169,chr170,chr171,chr172,chr173,chr174,chr175,chr176,chr177,chr178,chr179,chr180,chr181,chr182,chr183,chr184,chr185,chr186,chr187,chr188,chr189,chr190,chr191,chr192,chr193,chr194,chr195,chr196,chr197,chr198,chr199,chr200,chr201,chr202,chr203,chr204,chr205,chr206,chr207,chr208,chr209,chr210,chr211,chr212,chr213,chr214,chr215,chr216,chr217,chr218,chr219,chr220,chr221,chr222,chr223,chr224,chr225,chr226,chr227,chr228,chr229,chr230,chr231,chr232,chr233,chr234,chr235,chr236,chr237,chr238,chr239,chr240,chr241,chr242,chr243,chr244,chr245,chr246,chr247,chr248,chr249,chr250,chr251,chr252,chr253,chr254,chr255,chr256,chr257,chr258,chr259,chr260,chr261,chr262,chr263,chr264,chr265,chr266,chr267,chr268,chr269,chr270,chr271,chr272,chr273,chr274,chr275,chr276,chr277,chr278,chr279,chr280,chr281,chr282,chr283,chr284,chr285,chr286,chr287,chr288,chr289,chr290,chr291,chr292,chr293,chr294,chr295,chr296,chr297,chr298,chr299,chr300,chr301,chr302,chr303,chr304,chr305,chr306,chr307,chr308,chr309,chr310,chr311,chr312,chr313,chr314,chr315,chr316,chr317,chr318,chr319,chr320,chr321,chr322,chr323,chr324,chr325,chr326,chr327,chr328,chr329,chr330,chr331,chr332,chr333,chr334,chr335,chr336,chr337,chr338,chr339,chr340,chr341,chr342,chr343,chr344,chr345,chr346,chr347,chr348,chr349,chr350,chr351,chr352,chr353,chr354,chr355,chr356,chr357,chr358,chr359,chr360,chr361,chr362,chr363,chr364,chr365,chr366,chr367,chr368,chr369,chr370,chr371,chr372,chr373,chr374,chr375,chr376,chr377,chr378,chr379,chr380,chr381,chr382,chr383,chr384,chr385,chr386,chr387,chr388,chr389,chr390,chr391,chr392,chr393,chr394,chr395,chr396,chr397,chr398,chr399,chr400,chr401,chr402,chr403,chr404,chr405,chr406,chr407,chr408,chr409,chr410,chr411,chr412,chr413,chr414,chr415,chr416,chr417,chr418,chr419,chr420,chr421,chr422,chr423,chr424,chr425,chr426,chr427,chr428,chr429,chr430,chr431,chr432,chr433,chr434,chr435,chr436,chr437,chr438,chr439,chr440,chr441,chr442,chr443,chr444,chr445,chr446,chr447,chr448,chr449,chr450,chr451,chr452,chr453,chr454,chr455,chr456,chr457,chr458,chr459,chr460,chr461,chr462,chr463,chr464,chr465,chr466,chr467,chr468,chr469,chr470,chr471,chr472,chr473,chr474,chr475,chr476,chr477,chr478,chr479,chr480,chr481,chr482,chr483,chr484,chr485,chr486,chr487,chr488,chr489,chr490,chr491,chr492,chr493,chr494,chr495,chr496,chr497,chr498,chr499,chr500,chr501,chr502,chr503,chr504,chr505,chr506,chr507,chr508,chr509,chr510,chr511,chr512,chr513,chr514,chr515,chr516,chr517,chr518,chr519,chr520,chr521,chr522,chr523,chr524,chr525,chr526,chr527,chr528,chr529,chr530,chr531,chr532,chr533,chr534,chr535,chr536,chr537,chr538,chr539,chr540,chr541,chr542,chr543,chr544,chr545,chr546,chr547,chr548,chr549,chr550,chr551,chr552,chr553,chr554,chr555,chr556,chr557,chr558,chr559,chr560,chr561,chr562,chr563,chr564,chr565,chr566,chr567,chr568,chr569,chr570,chr571,chr572,chr573,chr574,chr575,chr576,chr577,chr578,chr579,chr580,chr581,chr582,chr583,chr584,chr585,chr586,chr587,chr588,chr589,chr590,chr591,chr592,chr593,chr594,chr595,chr596,chr597,chr598,chr599,chr600,chr601,chr602,chr603,chr604,chr605,chr606,chr607,chr608,chr609,chr610,chr611,chr612,chr613,chr614,chr615,chr616,chr617,chr618,chr619,chr620,chr621,chr622,chr623,chr624,chr625,chr626,chr627,chr628,chr629,chr630,chr631,chr632,chr633,chr634,chr635,chr636,chr637,chr638,chr639,chr640,chr641,chr642,chr643,chr644,chr645,chr646,chr647,chr648,chr649,chr650,chr651,chr652,chr653,chr654,chr655,chr656,chr657,chr658,chr659,chr660,chr661,chr662,chr663,chr664,chr665,chr666,chr667,chr668,chr669,chr670,chr671,chr672,chr673,chr674,chr675,chr676,chr677,chr678,chr679,chr680,chr681,chr682,chr683,chr684,chr685,chr686,chr687,chr688,chr689,chr690,chr691,chr692,chr693,chr694,chr695,chr696,chr697,chr698,chr699,chr700,chr701,chr702,chr703,chr704,chr705,chr706,chr707,chr708,chr709,chr710,chr711,chr712,chr713,chr714,chr715,chr716,chr717,chr718,chr719,chr720,chr721,chr722,chr723,chr724,chr725,chr726,chr727,chr728,chr729,chr730,chr731,chr732,chr733,chr734,chr735,chr736,chr737,chr738,chr739,chr740,chr741,chr742,chr743,chr744,chr745,chr746,chr747,chr748,chr749,chr750,chr751,chr752,chr753,chr754,chr755,chr756,chr757,chr758,chr759,chr760,chr761,chr762,chr763,chr764,chr765,chr766,chr767,chr768,chr769,chr770,chr771,chr772,chr773,chr774,chr775,chr776,chr777,chr778,chr779,chr780,chr781,chr782,chr783,chr784,chr785,chr786,chr787,chr788,chr789,chr790,chr791,chr792,chr793,chr794,chr795,chr796,chr797,chr798,chr799,chr800,chr801,chr802,chr803,chr804,chr805,chr806,chr807,chr808,chr809,chr810,chr811,chr812,chr813,chr814,chr815,chr816,chr817,chr818,chr819,chr820,chr821,chr822,chr8



A box plot showing the distribution of the coefficient of determination (csd) for various values of the parameter C_p . The x-axis is labeled C_p and has tick marks at 1, 3, 5, 7, 9, 11, 13, 15, 17, 19, and 21. The y-axis is labeled csd and ranges from 0.00000 to 0.00015 with major ticks every 0.00005. Each box plot represents the distribution of csd for a specific C_p value. The boxes are orange with black outlines, and the median is indicated by a thick black horizontal line. Dashed vertical lines extend from the boxes to the top and bottom of the plot area, representing the range of the data. The median csd is consistently around 0.00005 for all C_p values, and the spread of the data is relatively uniform across the range of C_p values.