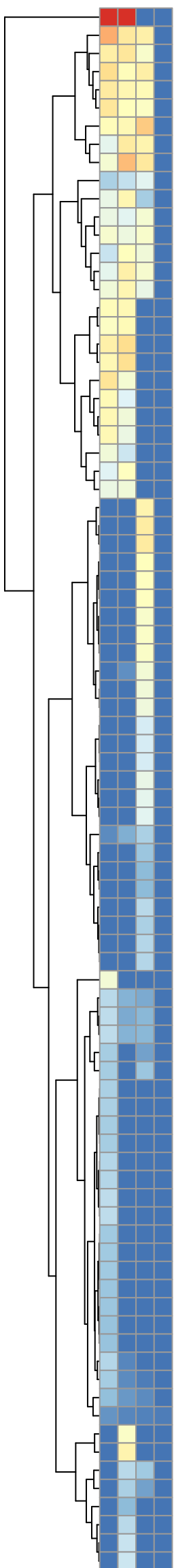
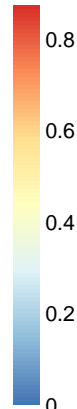


MB_SHH_T3



chr12:102060676__MYBPC1__NA
 chr8:48845580__PRKDC__unknown
 chr4:25677949__SLC34A2__nonsynonymous_SNV
 chr19:3589918__GIPC3__NA
 chr12:53040621__KRT2__nonsynonymous_SNV
 chr19:56702375__ZSCAN5B__NA
 chr5:126792963__MEGF10__stopgain_SNV
 chr12:123103008__KNTC1__NA
 chr15:101561213__LRRK1__NA
 chr9:140777280__CACNA1B__nonsynonymous_SNV
 chr7:100608268__ACHE(dist=114676),MUC12(dist=4636)__NA
 chr1:204413251__PIK3C2B__nonsynonymous_SNV
 chr17:3832036__ATP2A3__stopgain_SNV
 chrX:69366453__IGBP1__NA
 chr1:44587857__KLF17__NA
 chr9:35618245__CD72__nonsynonymous_SNV
 chr1:168262527__TBX19__NA
 chr14:71215605__MAP3K9__stopgain_SNV
 chr11:9074689__SCUBE2__synonymous_SNV
 chrX:36156085__CHDC2__synonymous_SNV
 chr5:38485877__LIFR__NA
 chr9:40702867__SPATA31A3__nonsynonymous_SNV
 chr12:21699280__GYS2__nonsynonymous_SNV
 chr3:166409304__BCHE(dist=854051),ZBBX(dist=548773)__NA
 chr22:22469383__TOP3B(dist=132143),VPREB1(dist=129817)__NA
 chr5:140744503__PCDHGA5__synonymous_SNV
 chr8:121328211__COL14A1__nonsynonymous_SNV
 chr14:54875422__CDKN3__NA
 chr12:4737717__AKAP3__synonymous_SNV
 chr6:30698483__FLOT1__synonymous_SNV
 chr15:63988312__HERC1__NA
 chr13:29004334__FLT1__NA
 chrX:119004629__RNF113A__nonsynonymous_SNV
 chr12:53202504__KRT4__nonsynonymous_SNV
 chr12:110570497__IFT81__NA
 chr5:85586898__NBPF22P__NA
 chr1:148012902__NBPF8__NA
 chr11:73675860__DNAJB13__NA
 chr7:20201492__MACC1__NA
 chr16:46428062__NONE(dist=NONE),ANKRD26P1(dist=75187)__NA
 chr1:21582557__ECE1__synonymous_SNV
 chr16:4254526__SRL__NA
 chr6:30671788__MDC1__nonsynonymous_SNV
 chr20:61511331__DIDO1__nonsynonymous_SNV
 chr3:183667641__ABCC5__nonsynonymous_SNV
 chr11:108216738__ATM__NA
 chr1:16258592__SPEN__nonsynonymous_SNV
 chr19:6420497__KHSRP__NA
 chr6:33411424__SYNGAP1__nonsynonymous_SNV
 chr16:84766768__USP10__NA
 chr22:42605851__TCF20__nonsynonymous_SNV
 chr1:155727076__GON4L__synonymous_SNV
 chr19:45409046__APOE__NA
 chr14:101514191__MIR381HG__NA
 chr4:88536955__DSPP__synonymous_SNV
 chr1:145296776__NBPF25P__NA
 chr6:170627695__FAM120B__nonsynonymous_SNV
 chr19:36365779__APLP1__NA
 chr19:8000073__TIMM44__NA
 chr1:210405167__SERTAD4-AS1__NA
 chr2:101869802__CNOT11__nonsynonymous_SNV
 chr22:30891985__SEC14L4__NA
 chr5:154179317__LARP1__nonsynonymous_SNV
 chr10:69770355__HERC4__NA
 chr22:24709505__SPECC1L-ADORA2A__NA
 chr3:138329870__FAIM__nonsynonymous_SNV
 chr9:135772999__TSC1(NM_000368:exon22:c.2626-2A>T,NM_001162427:exon21:c.2473-2A>T,NM_001162426:exon22:c.2623-2A>T)__NA
 chr1:15783667__CELA2A__nonsynonymous_SNV
 chr15:71185336__LRRRC49__NA
 chr16:30078544__ALDOA__NA
 chr22:45927276__FBLN1__NA
 chr17:17701053__RAI1__synonymous_SNV
 chr14:42367988__LRFN5__NA
 chr3:58139421__FLNB__NA
 chr17:3489091__TRPV1__nonsynonymous_SNV
 chr6:167571239__GPR31__synonymous_SNV
 chr2:162136825__AC009299.3(dist=31264),PSMD14(dist=27961)__NA
 chr9:139996380__MAN1B1__NA
 chr17:74288421__QRICH2__nonsynonymous_SNV
 chr17:74288422__QRICH2__nonsynonymous_SNV
 chr1:117122282__IGSF3__nonsynonymous_SNV
 chr12:133202328__POLE__nonsynonymous_SNV
 chr10:79589101__DLG5__NA
 chr1:155899515__KIAA0907__nonsynonymous_SNV
 chr15:71633455__THSD4__NA
 chr16:84488676__ATP2C2__NA



MB_SHH_T3.GL
 MB_SHH_T3.3
 MB_SHH_T3.2
 MB_SHH_T3.1