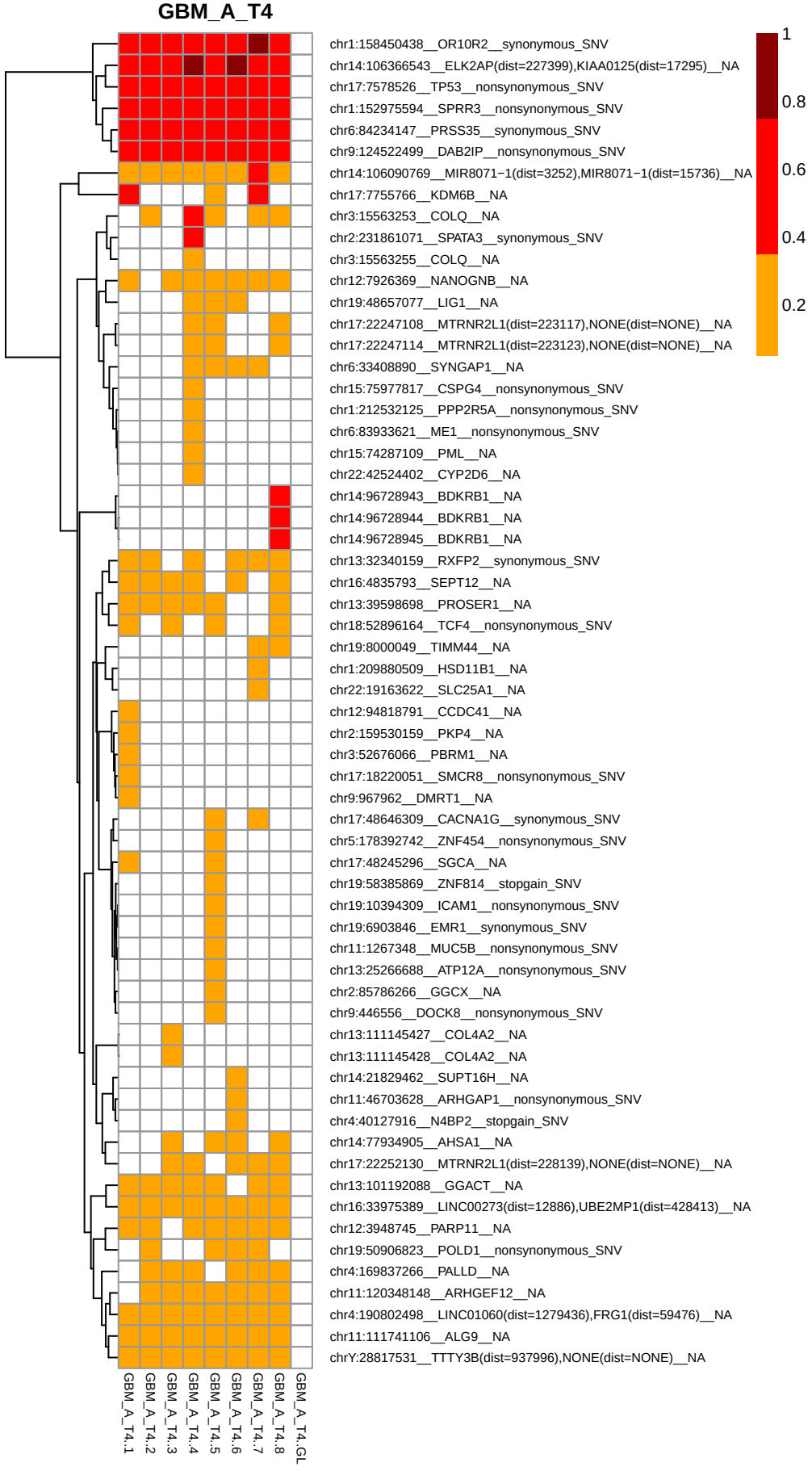
GBM_A_T4
chr1:158450438__OR10R2__synonymous_SNV
chr14:106366543__ELK2AP(dist=227399),KIAA0125(dist=17295)__NA
chr17:7578526__TP53__nonsynonymous_SNV
chr1:152975594__SPRR3__nonsynonymous_SNV
chr6:84234147__PRSS35__synonymous_SNV
chr9:124522499__DAB2IP__nonsynonymous_SNV
chr14:106090769__MIR8071−1(dist=3252),MIR8071−1(dist=15736)__NA
chr17:7755766__KDM6B__NA
chr3:15563253__COLQ__NA
chr2:231861071__SPATA3__synonymous_SNV
chr3:15563255__COLQ__NA
chr12:7926369__NANOGNB__NA
chr19:48657077__LIG1__NA
chr17:22247108__MTRNR2L1(dist=223117),NONE(dist=NONE)__NA
chr17:22247114__MTRNR2L1(dist=223123),NONE(dist=NONE)__NA
chr6:33408890__SYNGAP1__NA
chr15:75977817__CSPG4__nonsynonymous_SNV
chr1:212532125__PPP2R5A__nonsynonymous_SNV
chr6:83933621__ME1__nonsynonymous_SNV
chr15:74287109__PML__NA
chr22:42524402__CYP2D6__NA
chr14:96728943__BDKRB1__NA
chr14:96728944__BDKRB1__NA
chr14:96728945__BDKRB1__NA
chr13:32340159__RXFP2__synonymous_SNV
chr16:4835793__SEPT12__NA
chr13:39598698__PROSER1__NA
chr18:52896164__TCF4__nonsynonymous_SNV
chr19:8000049__TIMM44__NA
chr1:209880509__HSD11B1__NA
chr22:19163622__SLC25A1__NA
chr12:94818791__CCDC41__NA
chr2:159530159__PKP4__NA
chr3:52676066__PBRM1__NA
chr17:18220051__SMCR8__nonsynonymous_SNV
chr9:967962__DMRT1__NA
chr17:48646309__CACNA1G__synonymous_SNV
chr5:178392742__ZNF454__nonsynonymous_SNV
chr17:48245296__SGCA__NA
chr19:58385869__ZNF814__stopgain_SNV
chr19:10394309__ICAM1__nonsynonymous_SNV
chr19:6903846__EMR1__synonymous_SNV
chr11:1267348__MUC5B__nonsynonymous_SNV
chr13:25266688__ATP12A__nonsynonymous_SNV
chr2:85786266__GGCX__NA
chr9:446556__DOCK8__nonsynonymous_SNV
chr13:111145427__COL4A2__NA
chr13:111145428__COL4A2__NA
chr14:21829462__SUPT16H__NA
chr11:46703628__ARHGAP1__nonsynonymous_SNV
chr4:40127916__N4BP2__stopgain_SNV
chr14:77934905__AHSA1__NA
chr17:22252130__MTRNR2L1(dist=228139),NONE(dist=NONE)__NA
chr13:101192088__GGACT__NA
chr16:33975389__LINC00273(dist=12886),UBE2MP1(dist=428413)__NA
chr12:3948745__PARP11__NA
chr19:50906823__POLD1__nonsynonymous_SNV
chr4:169837266__PALLD__NA
chr11:120348148__ARHGEF12__NA
chr4:190802498__LINC01060(dist=1279436),FRG1(dist=59476)__NA
chr11:111741106__ALG9__NA
chrY:28817531__TTTY3B(dist=937996),NONE(dist=NONE)__NA
GBM_A_T4.1
GBM_A_T4.2
GBM_A_T4.3
GBM_A_T4.4
GBM_A_T4.5
GBM_A_T4.6
GBM_A_T4.7
GBM_A_T4.8
GBM_A_T4.GL
1
0.8
0.6
0.4
0.2