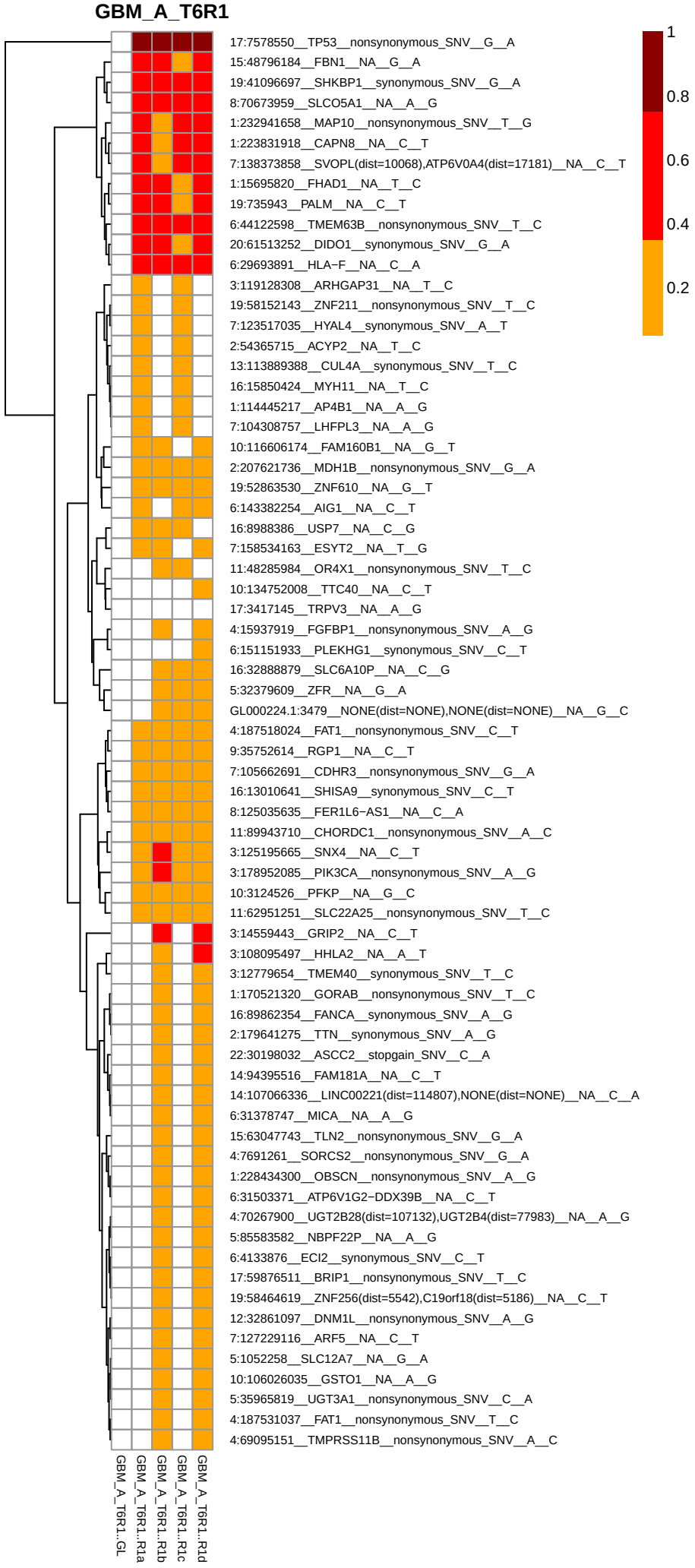
GBM_A_T6R1
17:7578550__TP53__nonsynonymous_SNV__G__A
15:48796184__FBN1__NA__G__A
19:41096697__SHKBP1__synonymous_SNV__G__A
8:70673959__SLCO5A1__NA__A__G
1:232941658__MAP10__nonsynonymous_SNV__T__G
1:223831918__CAPN8__NA__C__T
7:138373858__SVOPL(dist=10068),ATP6V0A4(dist=17181)__NA__C__T
1:15695820__FHAD1__NA__T__C
19:735943__PALM__NA__C__T
6:44122598__TMEM63B__nonsynonymous_SNV__T__C
20:61513252__DIDO1__synonymous_SNV__G__A
6:29693891__HLA−F__NA__C__A
3:119128308__ARHGAP31__NA__T__C
19:58152143__ZNF211__nonsynonymous_SNV__T__C
7:123517035__HYAL4__synonymous_SNV__A__T
2:54365715__ACYP2__NA__T__C
13:113889388__CUL4A__synonymous_SNV__T__C
16:15850424__MYH11__NA__T__C
1:114445217__AP4B1__NA__A__G
7:104308757__LHFPL3__NA__A__G
10:116606174__FAM160B1__NA__G__T
2:207621736__MDH1B__nonsynonymous_SNV__G__A
19:52863530__ZNF610__NA__G__T
6:143382254__AIG1__NA__C__T
16:8988386__USP7__NA__C__G
7:158534163__ESYT2__NA__T__G
11:48285984__OR4X1__nonsynonymous_SNV__T__C
10:134752008__TTC40__NA__C__T
17:3417145__TRPV3__NA__A__G
4:15937919__FGFBP1__nonsynonymous_SNV__A__G
6:151151933__PLEKHG1__synonymous_SNV__C__T
16:32888879__SLC6A10P__NA__C__G
5:32379609__ZFR__NA__G__A
GL000224.1:3479__NONE(dist=NONE),NONE(dist=NONE)__NA__G__C
4:187518024__FAT1__nonsynonymous_SNV__C__T
9:35752614__RGP1__NA__C__T
7:105662691__CDHR3__nonsynonymous_SNV__G__A
16:13010641__SHISA9__synonymous_SNV__C__T
8:125035635__FER1L6−AS1__NA__C__A
11:89943710__CHORDC1__nonsynonymous_SNV__A__C
3:125195665__SNX4__NA__C__T
3:178952085__PIK3CA__nonsynonymous_SNV__A__G
10:3124526__PFKP__NA__G__C
11:62951251__SLC22A25__nonsynonymous_SNV__T__C
3:14559443__GRIP2__NA__C__T
3:108095497__HHLA2__NA__A__T
3:12779654__TMEM40__synonymous_SNV__T__C
1:170521320__GORAB__nonsynonymous_SNV__T__C
16:89862354__FANCA__synonymous_SNV__A__G
2:179641275__TTN__synonymous_SNV__A__G
22:30198032__ASCC2__stopgain_SNV__C__A
14:94395516__FAM181A__NA__C__T
14:107066336__LINC00221(dist=114807),NONE(dist=NONE)__NA__C__A
6:31378747__MICA__NA__A__G
15:63047743__TLN2__nonsynonymous_SNV__G__A
4:7691261__SORCS2__nonsynonymous_SNV__G__A
1:228434300__OBSCN__nonsynonymous_SNV__A__G
6:31503371__ATP6V1G2−DDX39B__NA__C__T
4:70267900__UGT2B28(dist=107132),UGT2B4(dist=77983)__NA__A__G
5:85583582__NBPF22P__NA__A__G
6:4133876__ECI2__synonymous_SNV__C__T
17:59876511__BRIP1__nonsynonymous_SNV__T__C
19:58464619__ZNF256(dist=5542),C19orf18(dist=5186)__NA__C__T
12:32861097__DNM1L__nonsynonymous_SNV__A__G
7:127229116__ARF5__NA__C__T
5:1052258__SLC12A7__NA__G__A
10:106026035__GSTO1__NA__A__G
5:35965819__UGT3A1__nonsynonymous_SNV__C__A
4:187531037__FAT1__nonsynonymous_SNV__T__C
4:69095151__TMPRSS11B__nonsynonymous_SNV__A__C
GBM_A_T6R1..GL
GBM_A_T6R1..R1a
GBM_A_T6R1..R1b
GBM_A_T6R1..R1c
GBM_A_T6R1..R1d
1
0.8
0.6
0.4
0.2