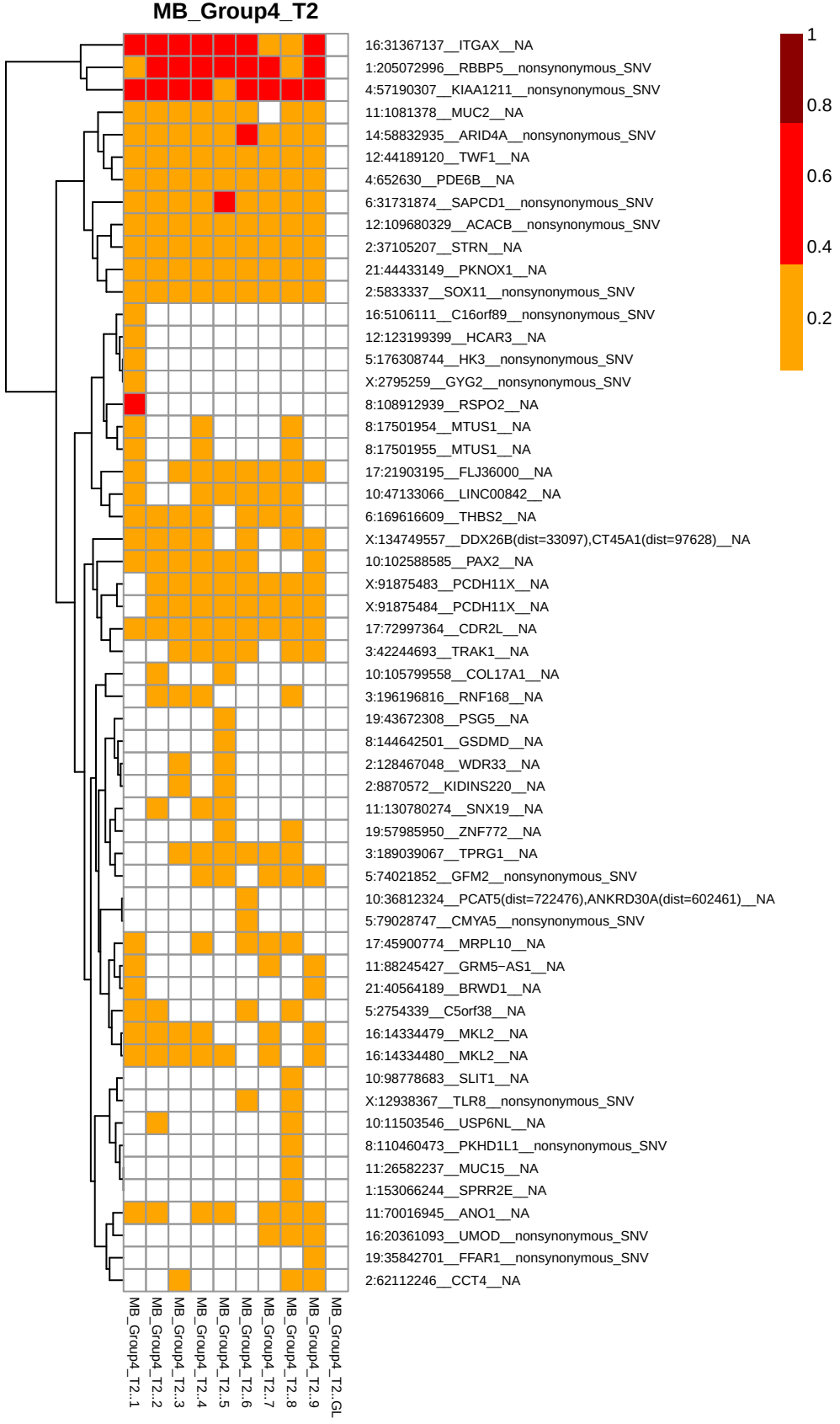
MB_Group4_T2
16:31367137__ITGAX__NA
1:205072996__RBBP5__nonsynonymous_SNV
4:57190307__KIAA1211__nonsynonymous_SNV
11:1081378__MUC2__NA
14:58832935__ARID4A__nonsynonymous_SNV
12:44189120__TWF1__NA
4:652630__PDE6B__NA
6:31731874__SAPCD1__nonsynonymous_SNV
12:109680329__ACACB__nonsynonymous_SNV
2:37105207__STRN__NA
21:44433149__PKNOX1__NA
2:5833337__SOX11__nonsynonymous_SNV
16:5106111__C16orf89__nonsynonymous_SNV
12:123199399__HCAR3__NA
5:176308744__HK3__nonsynonymous_SNV
X:2795259__GYG2__nonsynonymous_SNV
8:108912939__RSPO2__NA
8:17501954__MTUS1__NA
8:17501955__MTUS1__NA
17:21903195__FLJ36000__NA
10:47133066__LINC00842__NA
6:169616609__THBS2__NA
X:134749557__DDX26B(dist=33097),CT45A1(dist=97628)__NA
10:102588585__PAX2__NA
X:91875483__PCDH11X__NA
X:91875484__PCDH11X__NA
17:72997364__CDR2L__NA
3:42244693__TRAK1__NA
10:105799558__COL17A1__NA
3:196196816__RNF168__NA
19:43672308__PSG5__NA
8:144642501__GSDMD__NA
2:128467048__WDR33__NA
2:8870572__KIDINS220__NA
11:130780274__SNX19__NA
19:57985950__ZNF772__NA
3:189039067__TPRG1__NA
5:74021852__GFM2__nonsynonymous_SNV
10:36812324__PCAT5(dist=722476),ANKRD30A(dist=602461)__NA
5:79028747__CMYA5__nonsynonymous_SNV
17:45900774__MRPL10__NA
11:88245427__GRM5−AS1__NA
21:40564189__BRWD1__NA
5:2754339__C5orf38__NA
16:14334479__MKL2__NA
16:14334480__MKL2__NA
10:98778683__SLIT1__NA
X:12938367__TLR8__nonsynonymous_SNV
10:11503546__USP6NL__NA
8:110460473__PKHD1L1__nonsynonymous_SNV
11:26582237__MUC15__NA
1:153066244__SPRR2E__NA
11:70016945__ANO1__NA
16:20361093__UMOD__nonsynonymous_SNV
19:35842701__FFAR1__nonsynonymous_SNV
2:62112246__CCT4__NA
MB_Group4_T2..1
MB_Group4_T2..2
MB_Group4_T2..3
MB_Group4_T2..4
MB_Group4_T2..5
MB_Group4_T2..6
MB_Group4_T2..7
MB_Group4_T2..8
MB_Group4_T2..9
MB_Group4_T2..GL
1
0.8
0.6
0.4
0.2