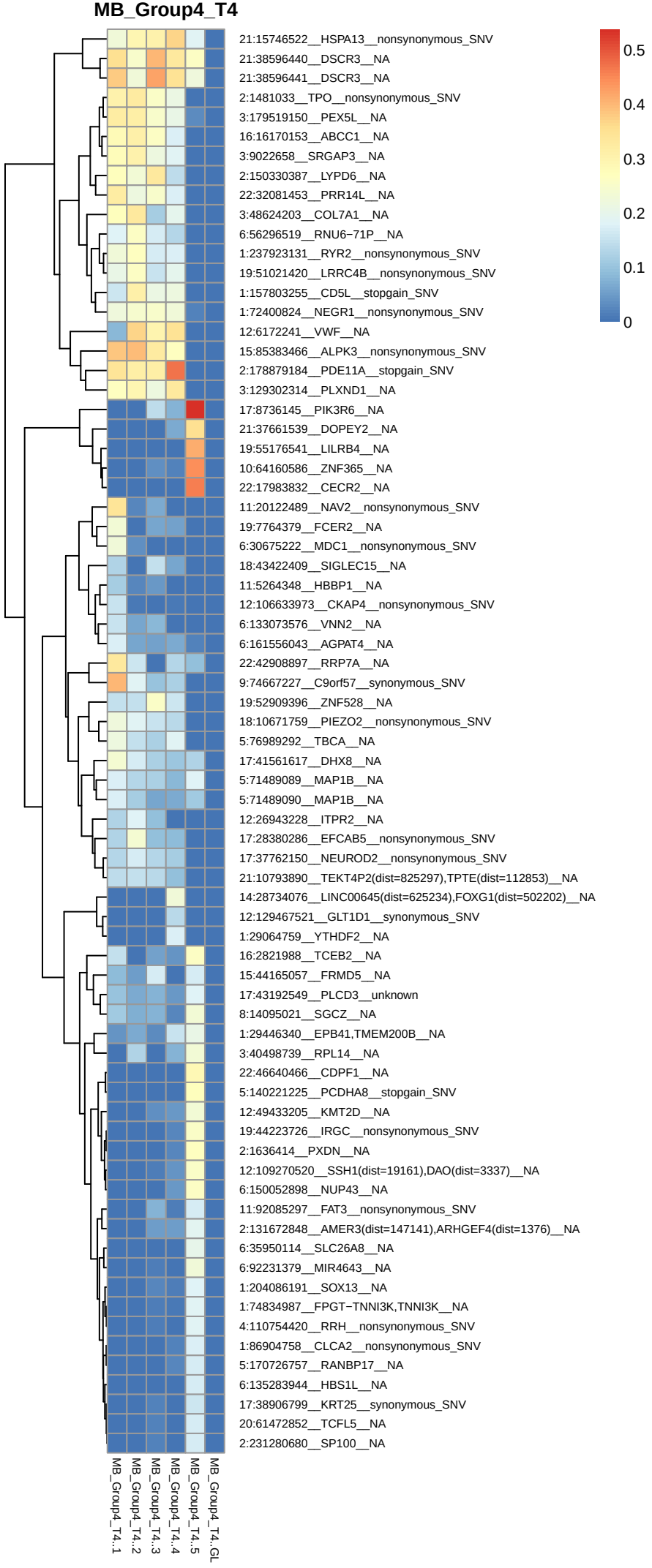
MB_Group4_T4
21:15746522__HSPA13__nonsynonymous_SNV
21:38596440__DSCR3__NA
21:38596441__DSCR3__NA
2:1481033__TPO__nonsynonymous_SNV
3:179519150__PEX5L__NA
16:16170153__ABCC1__NA
3:9022658__SRGAP3__NA
2:150330387__LYPD6__NA
22:32081453__PRR14L__NA
3:48624203__COL7A1__NA
6:56296519__RNU6−71P__NA
1:237923131__RYR2__nonsynonymous_SNV
19:51021420__LRRC4B__nonsynonymous_SNV
1:157803255__CD5L__stopgain_SNV
1:72400824__NEGR1__nonsynonymous_SNV
12:6172241__VWF__NA
15:85383466__ALPK3__nonsynonymous_SNV
2:178879184__PDE11A__stopgain_SNV
3:129302314__PLXND1__NA
17:8736145__PIK3R6__NA
21:37661539__DOPEY2__NA
19:55176541__LILRB4__NA
10:64160586__ZNF365__NA
22:17983832__CECR2__NA
11:20122489__NAV2__nonsynonymous_SNV
19:7764379__FCER2__NA
6:30675222__MDC1__nonsynonymous_SNV
18:43422409__SIGLEC15__NA
11:5264348__HBBP1__NA
12:106633973__CKAP4__nonsynonymous_SNV
6:133073576__VNN2__NA
6:161556043__AGPAT4__NA
22:42908897__RRP7A__NA
9:74667227__C9orf57__synonymous_SNV
19:52909396__ZNF528__NA
18:10671759__PIEZO2__nonsynonymous_SNV
5:76989292__TBCA__NA
17:41561617__DHX8__NA
5:71489089__MAP1B__NA
5:71489090__MAP1B__NA
12:26943228__ITPR2__NA
17:28380286__EFCAB5__nonsynonymous_SNV
17:37762150__NEUROD2__nonsynonymous_SNV
21:10793890__TEKT4P2(dist=825297),TPTE(dist=112853)__NA
14:28734076__LINC00645(dist=625234),FOXG1(dist=502202)__NA
12:129467521__GLT1D1__synonymous_SNV
1:29064759__YTHDF2__NA
16:2821988__TCEB2__NA
15:44165057__FRMD5__NA
17:43192549__PLCD3__unknown
8:14095021__SGCZ__NA
1:29446340__EPB41,TMEM200B__NA
3:40498739__RPL14__NA
22:46640466__CDPF1__NA
5:140221225__PCDHA8__stopgain_SNV
12:49433205__KMT2D__NA
19:44223726__IRGC__nonsynonymous_SNV
2:1636414__PXDN__NA
12:109270520__SSH1(dist=19161),DAO(dist=3337)__NA
6:150052898__NUP43__NA
11:92085297__FAT3__nonsynonymous_SNV
2:131672848__AMER3(dist=147141),ARHGEF4(dist=1376)__NA
6:35950114__SLC26A8__NA
6:92231379__MIR4643__NA
1:204086191__SOX13__NA
1:74834987__FPGT−TNNI3K,TNNI3K__NA
4:110754420__RRH__nonsynonymous_SNV
1:86904758__CLCA2__nonsynonymous_SNV
5:170726757__RANBP17__NA
6:135283944__HBS1L__NA
17:38906799__KRT25__synonymous_SNV
20:61472852__TCFL5__NA
2:231280680__SP100__NA
MB_Group4_T4..1
MB_Group4_T4..2
MB_Group4_T4..3
MB_Group4_T4..4
MB_Group4_T4..5
MB_Group4_T4..GL
0.5
0.4
0.3
0.2
0.1
0