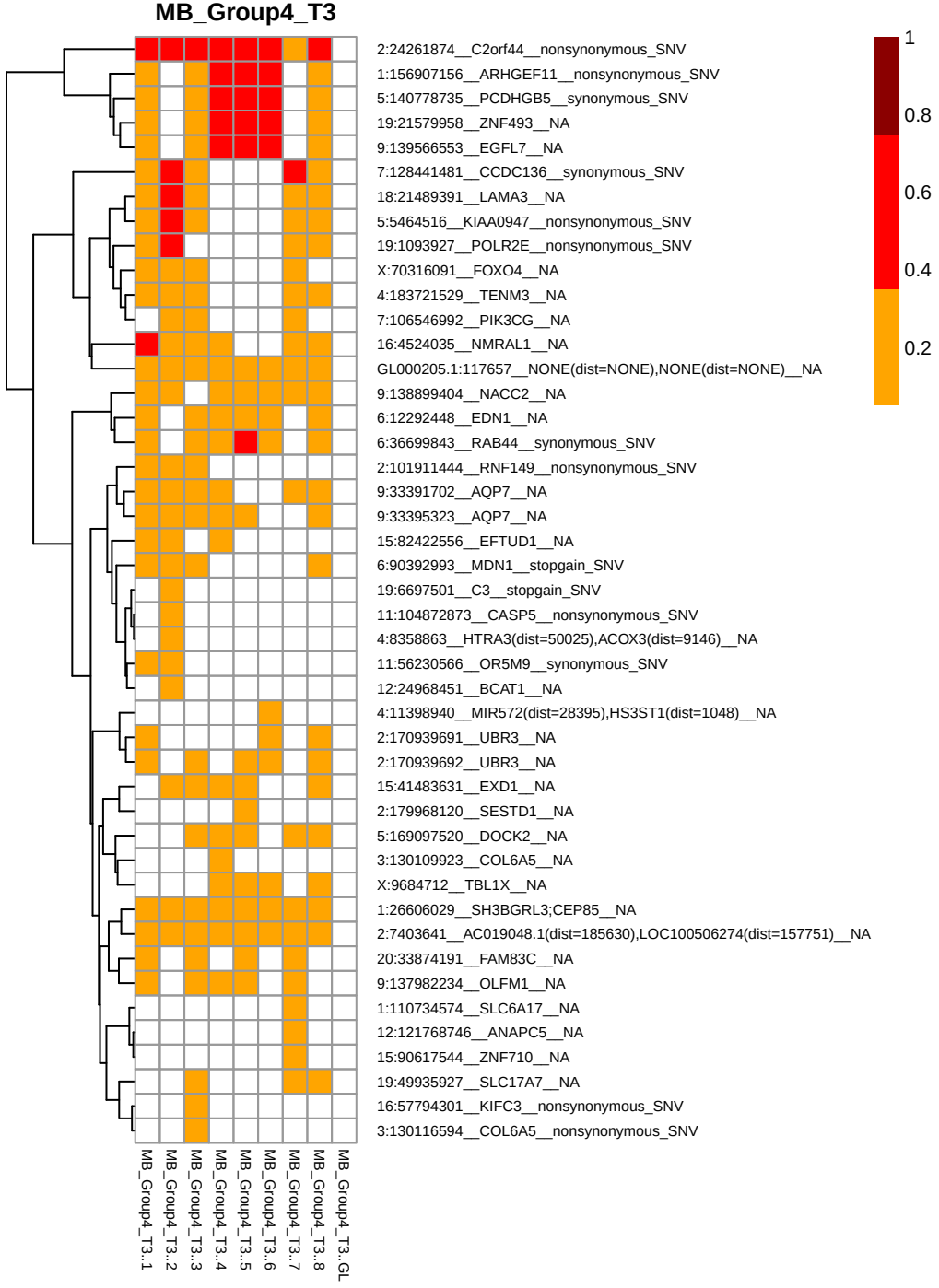
MB_Group4_T3
2:24261874__C2orf44__nonsynonymous_SNV
1:156907156__ARHGEF11__nonsynonymous_SNV
5:140778735__PCDHGB5__synonymous_SNV
19:21579958__ZNF493__NA
9:139566553__EGFL7__NA
7:128441481__CCDC136__synonymous_SNV
18:21489391__LAMA3__NA
5:5464516__KIAA0947__nonsynonymous_SNV
19:1093927__POLR2E__nonsynonymous_SNV
X:70316091__FOXO4__NA
4:183721529__TENM3__NA
7:106546992__PIK3CG__NA
16:4524035__NMRAL1__NA
GL000205.1:117657__NONE(dist=NONE),NONE(dist=NONE)__NA
9:138899404__NACC2__NA
6:12292448__EDN1__NA
6:36699843__RAB44__synonymous_SNV
2:101911444__RNF149__nonsynonymous_SNV
9:33391702__AQP7__NA
9:33395323__AQP7__NA
15:82422556__EFTUD1__NA
6:90392993__MDN1__stopgain_SNV
19:6697501__C3__stopgain_SNV
11:104872873__CASP5__nonsynonymous_SNV
4:8358863__HTRA3(dist=50025),ACOX3(dist=9146)__NA
11:56230566__OR5M9__synonymous_SNV
12:24968451__BCAT1__NA
4:11398940__MIR572(dist=28395),HS3ST1(dist=1048)__NA
2:170939691__UBR3__NA
2:170939692__UBR3__NA
15:41483631__EXD1__NA
2:179968120__SESTD1__NA
5:169097520__DOCK2__NA
3:130109923__COL6A5__NA
X:9684712__TBL1X__NA
1:26606029__SH3BGRL3;CEP85__NA
2:7403641__AC019048.1(dist=185630),LOC100506274(dist=157751)__NA
20:33874191__FAM83C__NA
9:137982234__OLFM1__NA
1:110734574__SLC6A17__NA
12:121768746__ANAPC5__NA
15:90617544__ZNF710__NA
19:49935927__SLC17A7__NA
16:57794301__KIFC3__nonsynonymous_SNV
3:130116594__COL6A5__nonsynonymous_SNV
MB_Group4_T3..1
MB_Group4_T3..2
MB_Group4_T3..3
MB_Group4_T3..4
MB_Group4_T3..5
MB_Group4_T3..6
MB_Group4_T3..7
MB_Group4_T3..8
MB_Group4_T3..GL
1
0.8
0.6
0.4
0.2