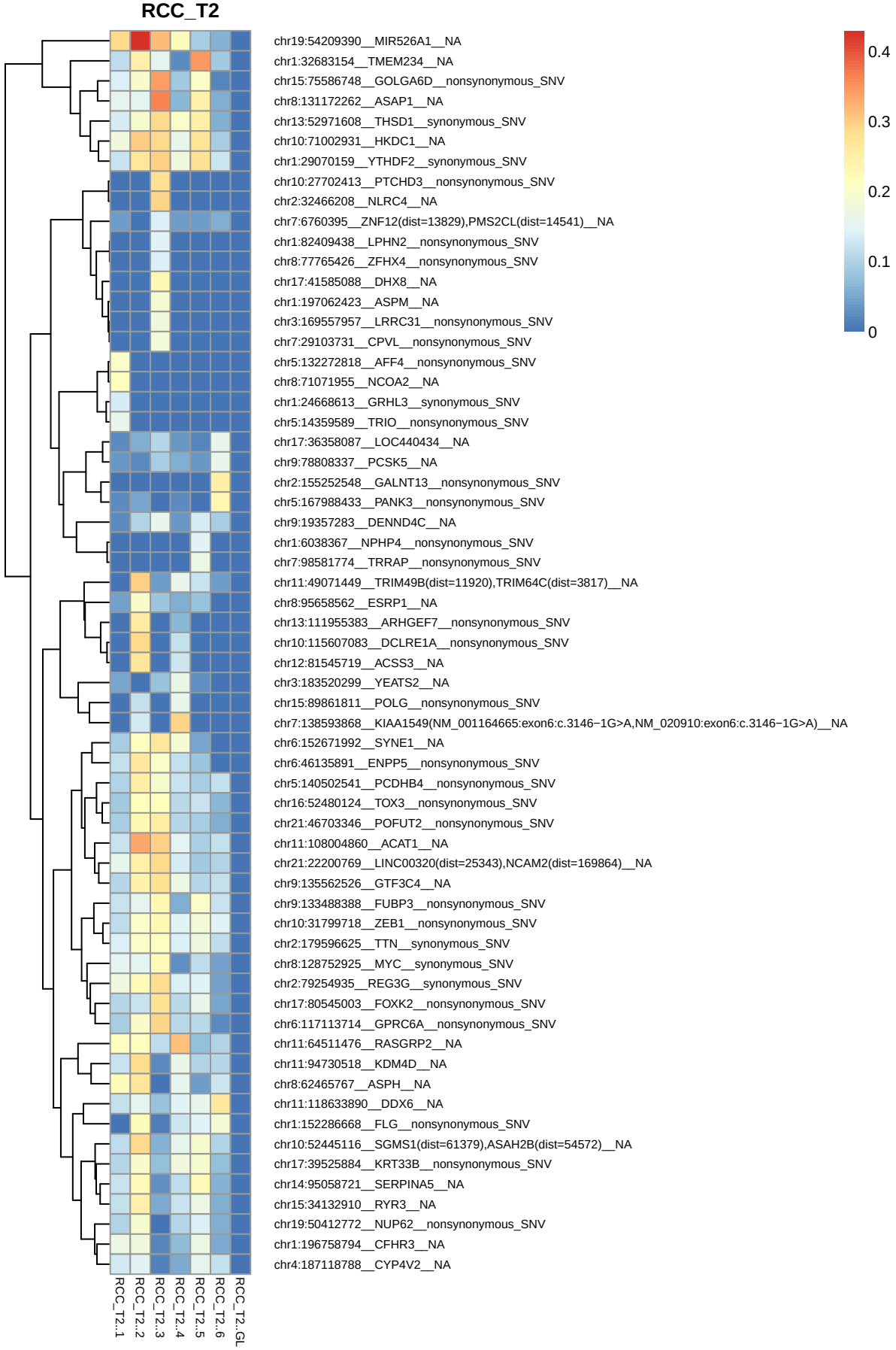
RCC_T2
chr19:54209390__MIR526A1__NA
chr1:32683154__TMEM234__NA
chr15:75586748__GOLGA6D__nonsynonymous_SNV
chr8:131172262__ASAP1__NA
chr13:52971608__THSD1__synonymous_SNV
chr10:71002931__HKDC1__NA
chr1:29070159__YTHDF2__synonymous_SNV
chr10:27702413__PTCHD3__nonsynonymous_SNV
chr2:32466208__NLRC4__NA
chr7:6760395__ZNF12(dist=13829),PMS2CL(dist=14541)__NA
chr1:82409438__LPHN2__nonsynonymous_SNV
chr8:77765426__ZFHX4__nonsynonymous_SNV
chr17:41585088__DHX8__NA
chr1:197062423__ASPM__NA
chr3:169557957__LRRC31__nonsynonymous_SNV
chr7:29103731__CPVL__nonsynonymous_SNV
chr5:132272818__AFF4__nonsynonymous_SNV
chr8:71071955__NCOA2__NA
chr1:24668613__GRHL3__synonymous_SNV
chr5:14359589__TRIO__nonsynonymous_SNV
chr17:36358087__LOC440434__NA
chr9:78808337__PCSK5__NA
chr2:155252548__GALNT13__nonsynonymous_SNV
chr5:167988433__PANK3__nonsynonymous_SNV
chr9:19357283__DENND4C__NA
chr1:6038367__NPHP4__nonsynonymous_SNV
chr7:98581774__TRRAP__nonsynonymous_SNV
chr11:49071449__TRIM49B(dist=11920),TRIM64C(dist=3817)__NA
chr8:95658562__ESRP1__NA
chr13:111955383__ARHGEF7__nonsynonymous_SNV
chr10:115607083__DCLRE1A__nonsynonymous_SNV
chr12:81545719__ACSS3__NA
chr3:183520299__YEATS2__NA
chr15:89861811__POLG__nonsynonymous_SNV
chr7:138593868__KIAA1549(NM_001164665:exon6:c.3146−1G>A,NM_020910:exon6:c.3146−1G>A)__NA
chr6:152671992__SYNE1__NA
chr6:46135891__ENPP5__nonsynonymous_SNV
chr5:140502541__PCDHB4__nonsynonymous_SNV
chr16:52480124__TOX3__nonsynonymous_SNV
chr21:46703346__POFUT2__nonsynonymous_SNV
chr11:108004860__ACAT1__NA
chr21:22200769__LINC00320(dist=25343),NCAM2(dist=169864)__NA
chr9:135562526__GTF3C4__NA
chr9:133488388__FUBP3__nonsynonymous_SNV
chr10:31799718__ZEB1__nonsynonymous_SNV
chr2:179596625__TTN__synonymous_SNV
chr8:128752925__MYC__synonymous_SNV
chr2:79254935__REG3G__synonymous_SNV
chr17:80545003__FOXK2__nonsynonymous_SNV
chr6:117113714__GPRC6A__nonsynonymous_SNV
chr11:64511476__RASGRP2__NA
chr11:94730518__KDM4D__NA
chr8:62465767__ASPH__NA
chr11:118633890__DDX6__NA
chr1:152286668__FLG__nonsynonymous_SNV
chr10:52445116__SGMS1(dist=61379),ASAH2B(dist=54572)__NA
chr17:39525884__KRT33B__nonsynonymous_SNV
chr14:95058721__SERPINA5__NA
chr15:34132910__RYR3__NA
chr19:50412772__NUP62__nonsynonymous_SNV
chr1:196758794__CFHR3__NA
chr4:187118788__CYP4V2__NA
RCC_T2..1
RCC_T2..2
RCC_T2..3
RCC_T2..4
RCC_T2..5
RCC_T2..6
RCC_T2..GL
0.4
0.3
0.2
0.1
0