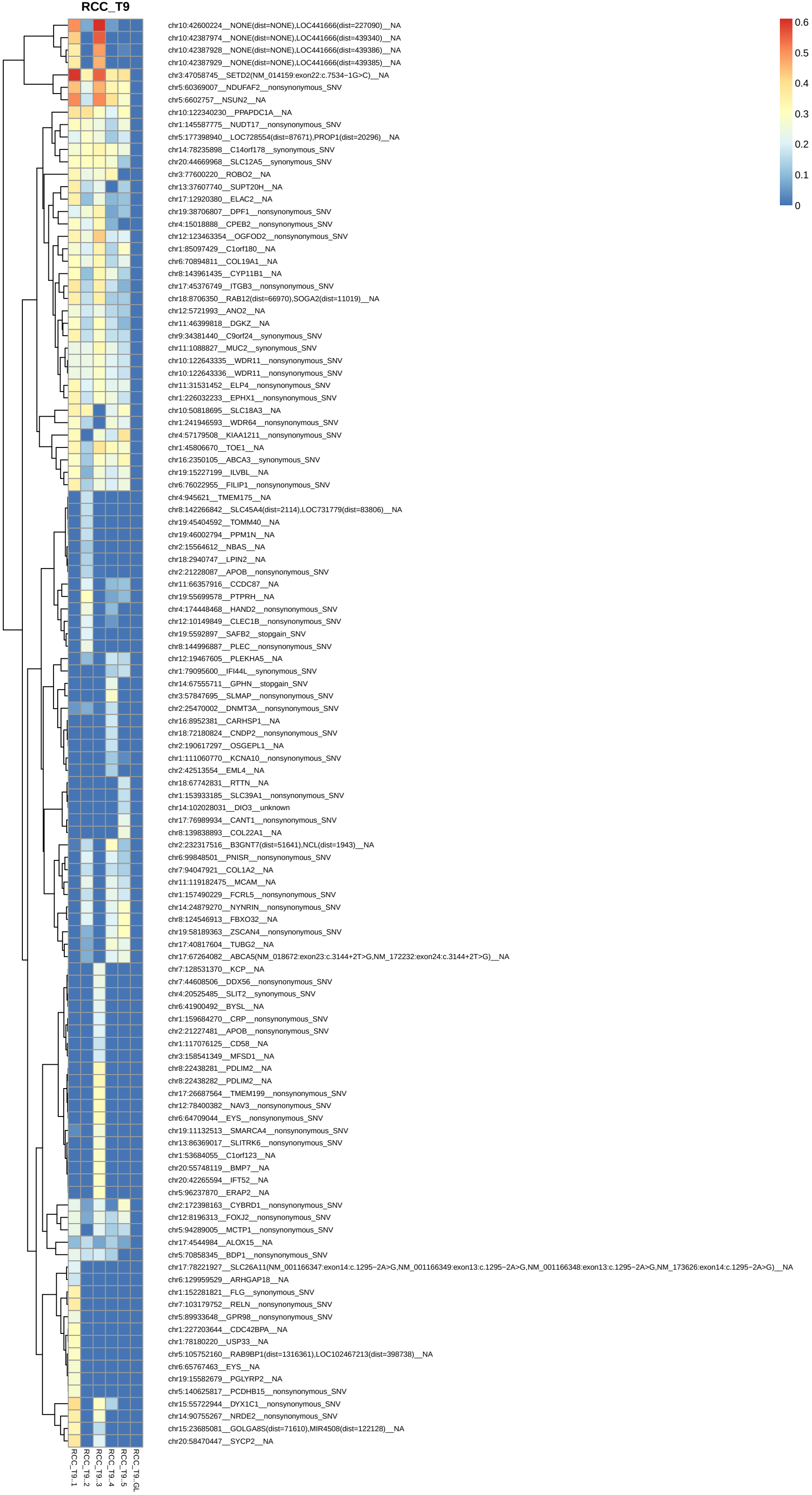
RCC_T9
chr10:42600224__NONE(dist=NONE),LOC441666(dist=227090)__NA
chr10:42387974__NONE(dist=NONE),LOC441666(dist=439340)__NA
chr10:42387928__NONE(dist=NONE),LOC441666(dist=439386)__NA
chr10:42387929__NONE(dist=NONE),LOC441666(dist=439385)__NA
chr3:47058745__SETD2(NM_014159:exon22:c.7534−1G>C)__NA
chr5:60369007__NDUFAF2__nonsynonymous_SNV
chr5:6602757__NSUN2__NA
chr10:122340230__PPAPDC1A__NA
chr1:145587775__NUDT17__nonsynonymous_SNV
chr5:177398940__LOC728554(dist=87671),PROP1(dist=20296)__NA
chr14:78235898__C14orf178__synonymous_SNV
chr20:44669968__SLC12A5__synonymous_SNV
chr3:77600220__ROBO2__NA
chr13:37607740__SUPT20H__NA
chr17:12920380__ELAC2__NA
chr19:38706807__DPF1__nonsynonymous_SNV
chr4:15018888__CPEB2__nonsynonymous_SNV
chr12:123463354__OGFOD2__nonsynonymous_SNV
chr1:85097429__C1orf180__NA
chr6:70894811__COL19A1__NA
chr8:143961435__CYP11B1__NA
chr17:45376749__ITGB3__nonsynonymous_SNV
chr18:8706350__RAB12(dist=66970),SOGA2(dist=11019)__NA
chr12:5721993__ANO2__NA
chr11:46399818__DGKZ__NA
chr9:34381440__C9orf24__synonymous_SNV
chr11:1088827__MUC2__synonymous_SNV
chr10:122643335__WDR11__nonsynonymous_SNV
chr10:122643336__WDR11__nonsynonymous_SNV
chr11:31531452__ELP4__nonsynonymous_SNV
chr1:226032233__EPHX1__nonsynonymous_SNV
chr10:50818695__SLC18A3__NA
chr1:241946593__WDR64__nonsynonymous_SNV
chr4:57179508__KIAA1211__nonsynonymous_SNV
chr1:45806670__TOE1__NA
chr16:2350105__ABCA3__synonymous_SNV
chr19:15227199__ILVBL__NA
chr6:76022955__FILIP1__nonsynonymous_SNV
chr4:945621__TMEM175__NA
chr8:142266842__SLC45A4(dist=2114),LOC731779(dist=83806)__NA
chr19:45404592__TOMM40__NA
chr19:46002794__PPM1N__NA
chr2:15564612__NBAS__NA
chr18:2940747__LPIN2__NA
chr2:21228087__APOB__nonsynonymous_SNV
chr11:66357916__CCDC87__NA
chr19:55699578__PTPRH__NA
chr4:174448468__HAND2__nonsynonymous_SNV
chr12:10149849__CLEC1B__nonsynonymous_SNV
chr19:5592897__SAFB2__stopgain_SNV
chr8:144996887__PLEC__nonsynonymous_SNV
chr12:19467605__PLEKHA5__NA
chr1:79095600__IFI44L__synonymous_SNV
chr14:67555711__GPHN__stopgain_SNV
chr3:57847695__SLMAP__nonsynonymous_SNV
chr2:25470002__DNMT3A__nonsynonymous_SNV
chr16:8952381__CARHSP1__NA
chr18:72180824__CNDP2__nonsynonymous_SNV
chr2:190617297__OSGEPL1__NA
chr1:111060770__KCNA10__nonsynonymous_SNV
chr2:42513554__EML4__NA
chr18:67742831__RTTN__NA
chr1:153933185__SLC39A1__nonsynonymous_SNV
chr14:102028031__DIO3__unknown
chr17:76989934__CANT1__nonsynonymous_SNV
chr8:139838893__COL22A1__NA
chr2:232317516__B3GNT7(dist=51641),NCL(dist=1943)__NA
chr6:99848501__PNISR__nonsynonymous_SNV
chr7:94047921__COL1A2__NA
chr11:119182475__MCAM__NA
chr1:157490229__FCRL5__nonsynonymous_SNV
chr14:24879270__NYNRIN__nonsynonymous_SNV
chr8:124546913__FBXO32__NA
chr19:58189363__ZSCAN4__nonsynonymous_SNV
chr17:40817604__TUBG2__NA
chr17:67264082__ABCA5(NM_018672:exon23:c.3144+2T>G,NM_172232:exon24:c.3144+2T>G)__NA
chr7:128531370__KCP__NA
chr7:44608506__DDX56__nonsynonymous_SNV
chr4:20525485__SLIT2__synonymous_SNV
chr6:41900492__BYSL__NA
chr1:159684270__CRP__nonsynonymous_SNV
chr2:21227481__APOB__nonsynonymous_SNV
chr1:117076125__CD58__NA
chr3:158541349__MFSD1__NA
chr8:22438281__PDLIM2__NA
chr8:22438282__PDLIM2__NA
chr17:26687564__TMEM199__nonsynonymous_SNV
chr12:78400382__NAV3__nonsynonymous_SNV
chr6:64709044__EYS__nonsynonymous_SNV
chr19:11132513__SMARCA4__nonsynonymous_SNV
chr13:86369017__SLITRK6__nonsynonymous_SNV
chr1:53684055__C1orf123__NA
chr20:55748119__BMP7__NA
chr20:42265594__IFT52__NA
chr5:96237870__ERAP2__NA
chr2:172398163__CYBRD1__nonsynonymous_SNV
chr12:8196313__FOXJ2__nonsynonymous_SNV
chr5:94289005__MCTP1__nonsynonymous_SNV
chr17:4544984__ALOX15__NA
chr5:70858345__BDP1__nonsynonymous_SNV
chr17:78221927__SLC26A11(NM_001166347:exon14:c.1295−2A>G,NM_001166349:exon13:c.1295−2A>G,NM_001166348:exon13:c.1295−2A>G,NM_173626:exon14:c.1295−2A>G)__NA
chr6:129959529__ARHGAP18__NA
chr1:152281821__FLG__synonymous_SNV
chr7:103179752__RELN__nonsynonymous_SNV
chr5:89933648__GPR98__nonsynonymous_SNV
chr1:227203644__CDC42BPA__NA
chr1:78180220__USP33__NA
chr5:105752160__RAB9BP1(dist=1316361),LOC102467213(dist=398738)__NA
chr6:65767463__EYS__NA
chr19:15582679__PGLYRP2__NA
chr5:140625817__PCDHB15__nonsynonymous_SNV
chr15:55722944__DYX1C1__nonsynonymous_SNV
chr14:90755267__NRDE2__nonsynonymous_SNV
chr15:23685081__GOLGA8S(dist=71610),MIR4508(dist=122128)__NA
chr20:58470447__SYCP2__NA
RCC_T9..1
RCC_T9..2
RCC_T9..3
RCC_T9..4
RCC_T9..5
RCC_T9..GL
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0.5
0.4
0.3
0.2
0.1
0