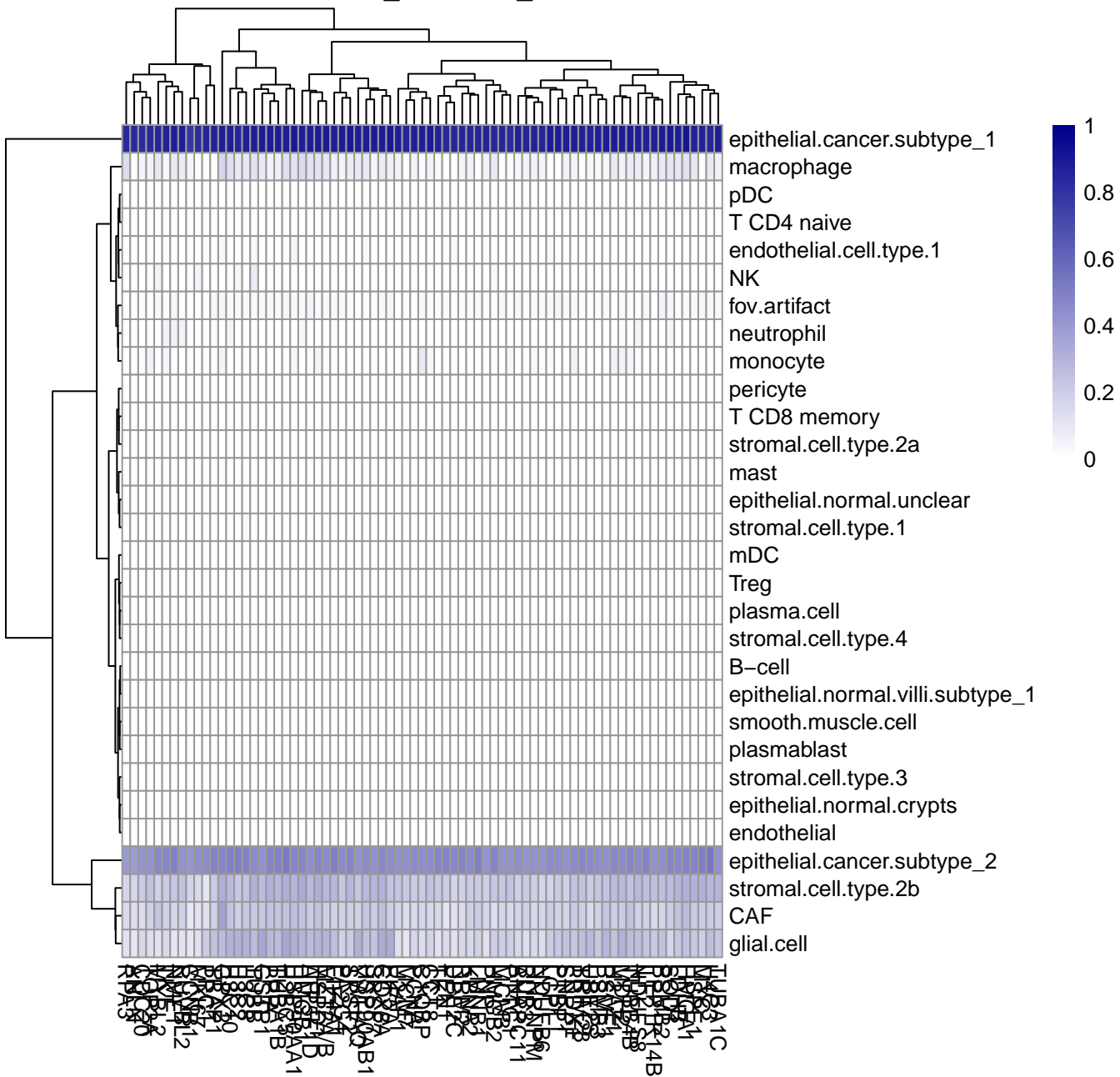
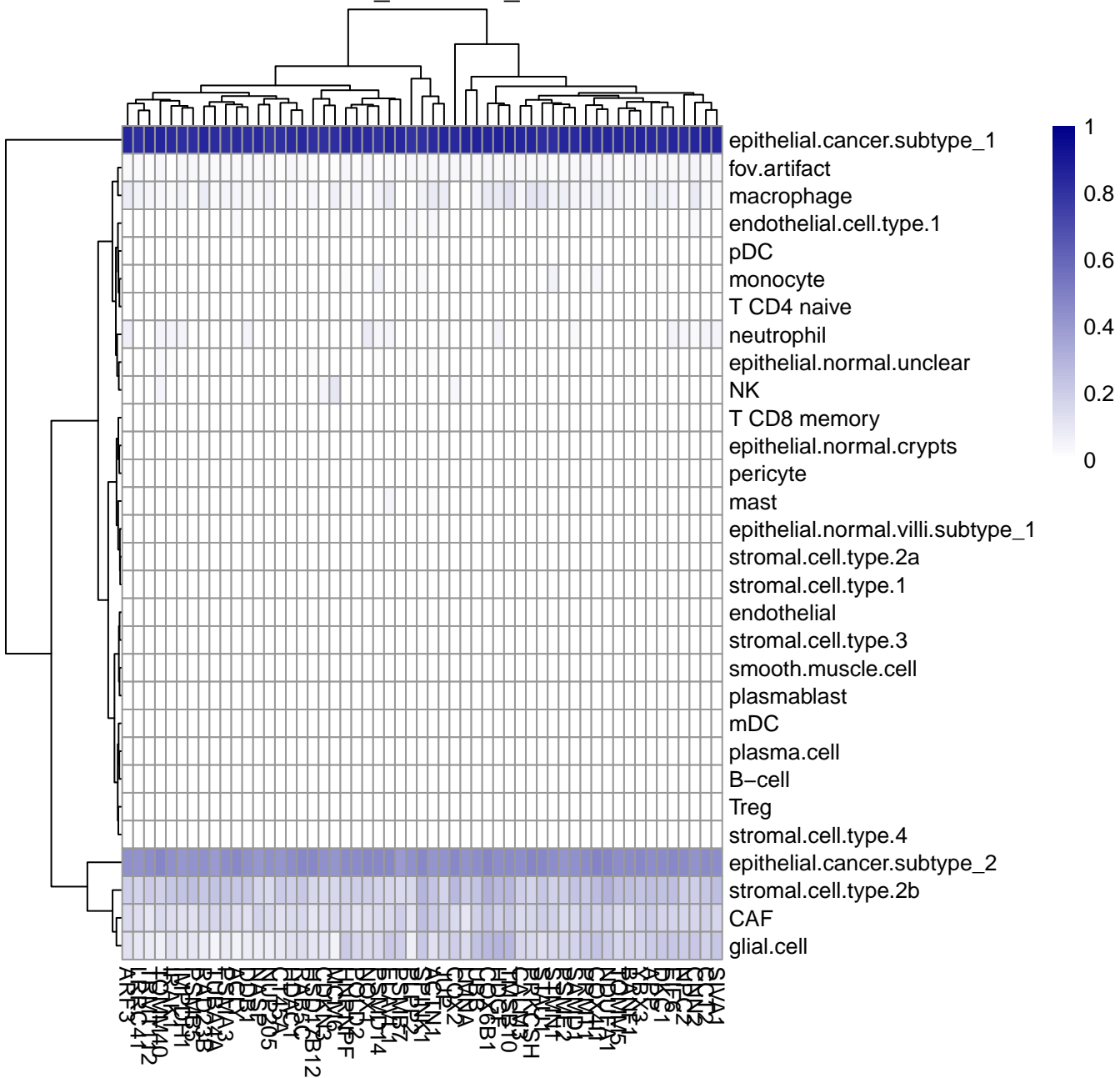


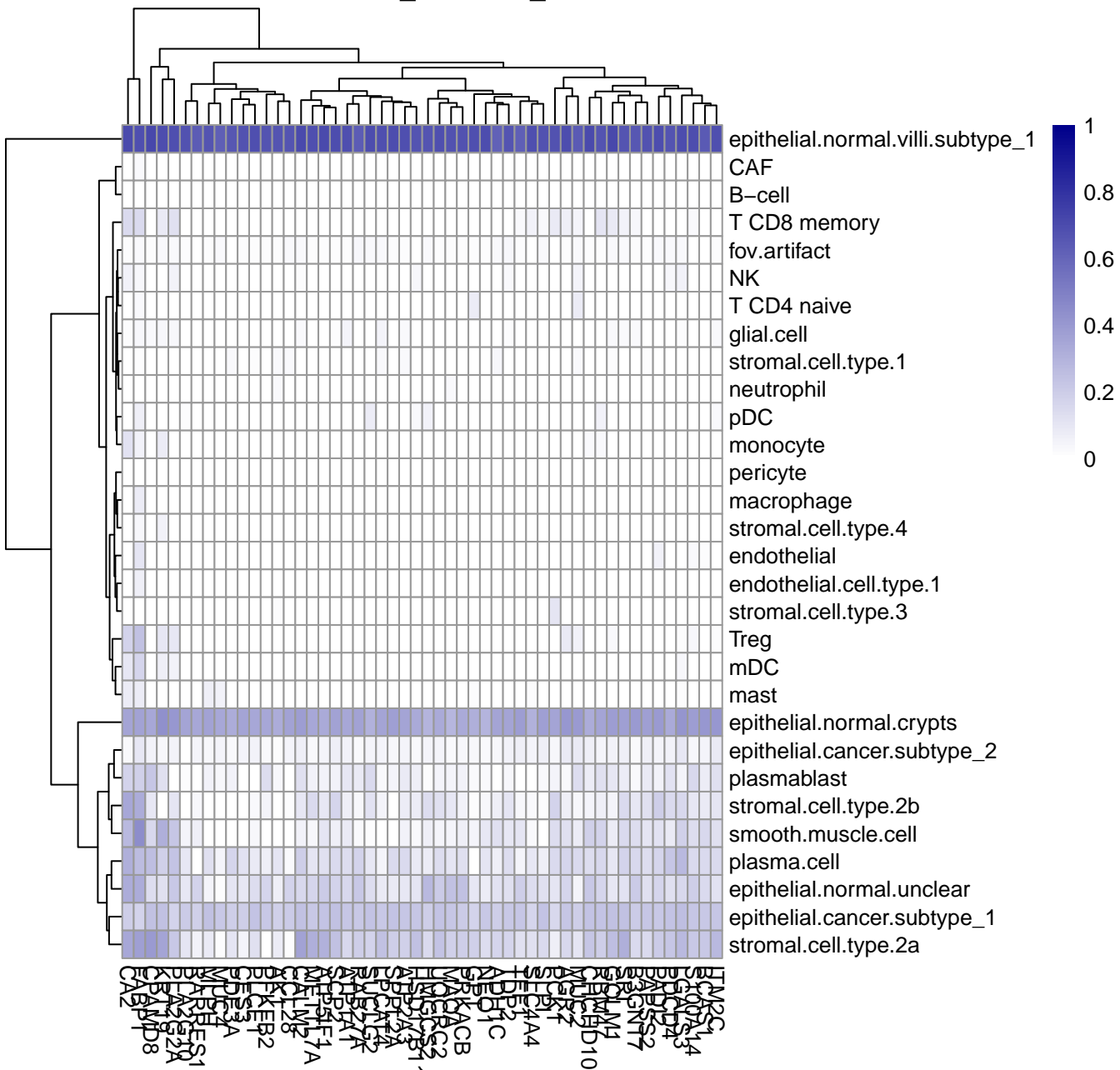
HMGA1_TUBB4B_75



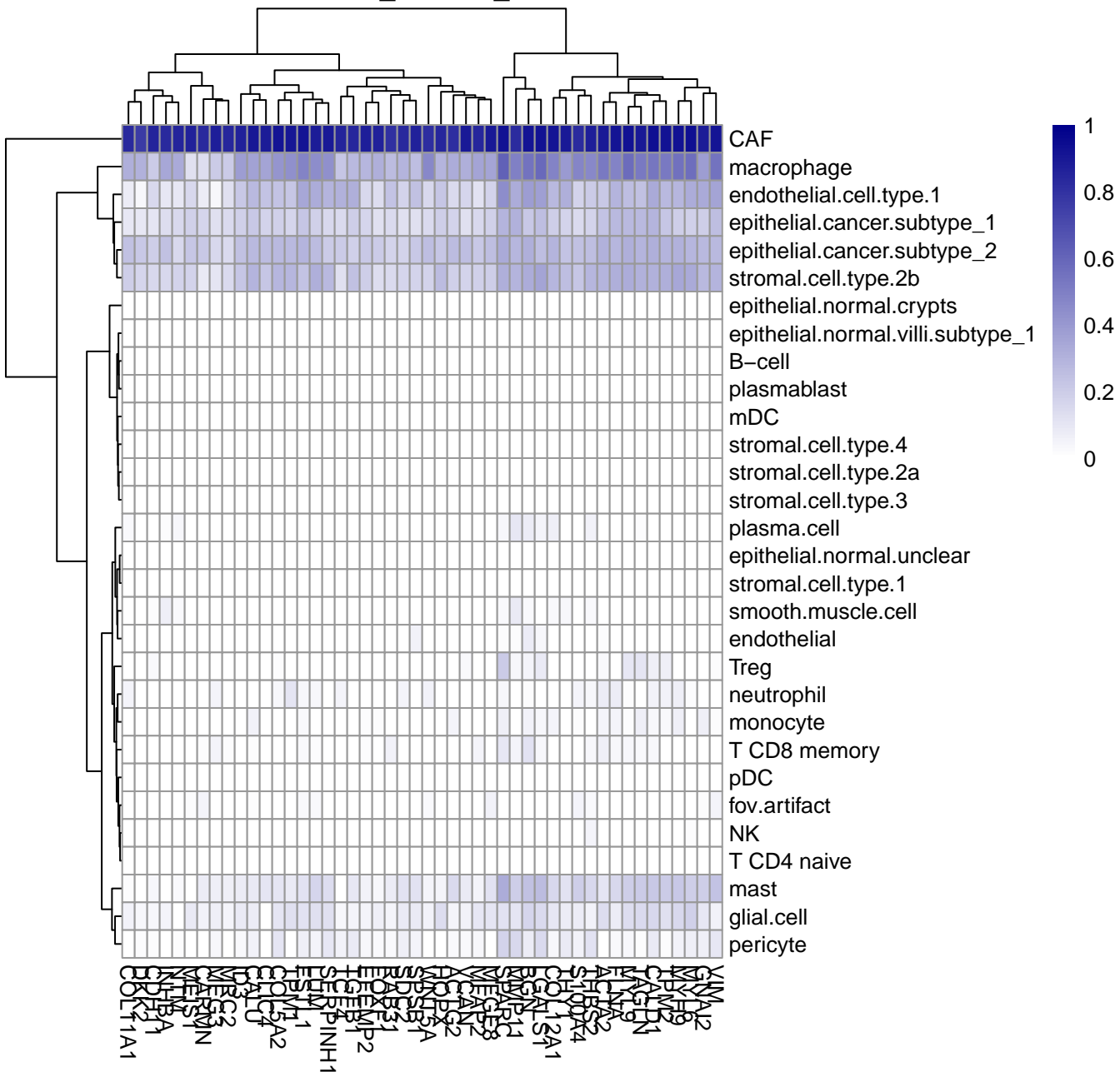
HDGF TMSB10 55



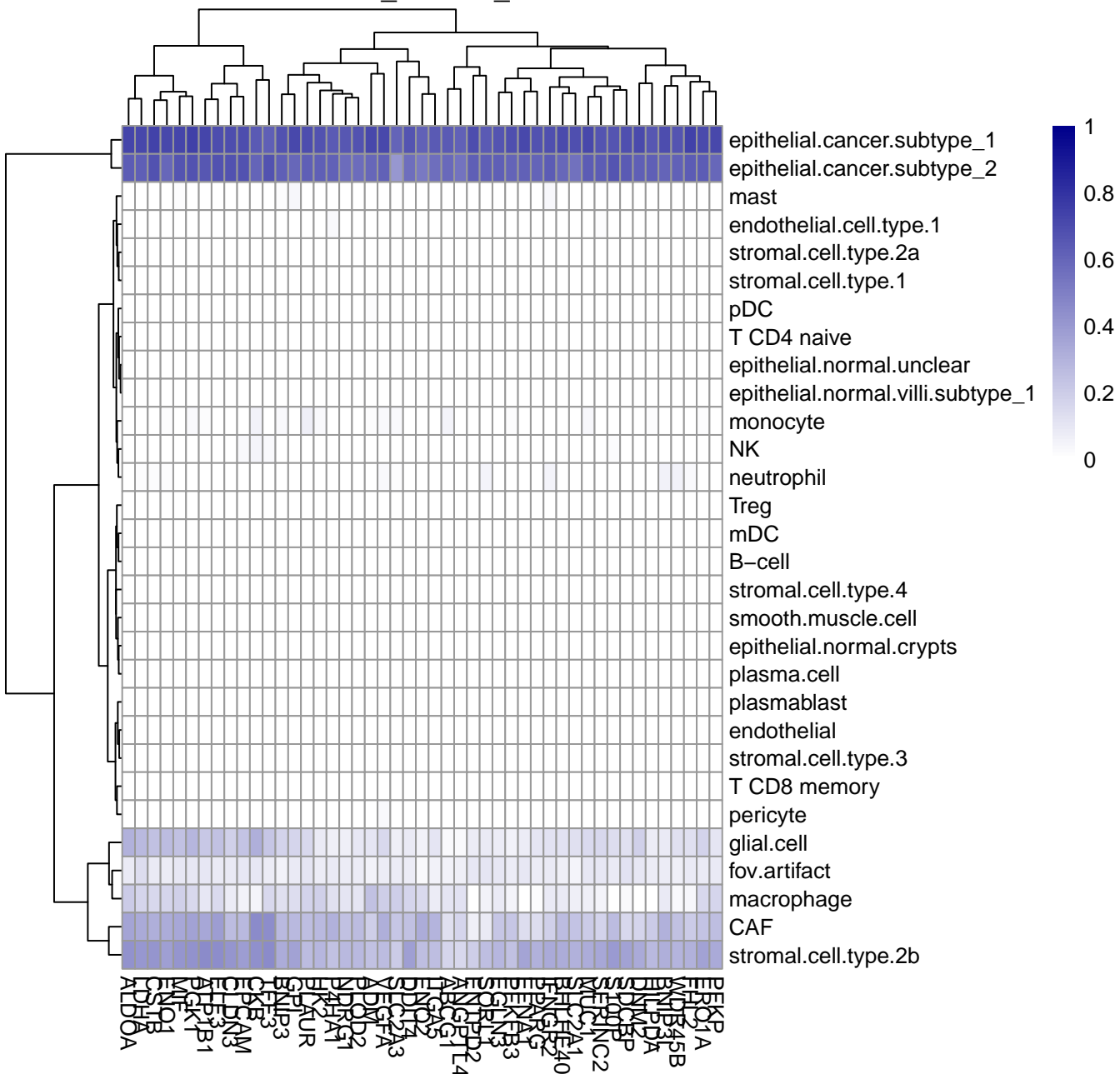
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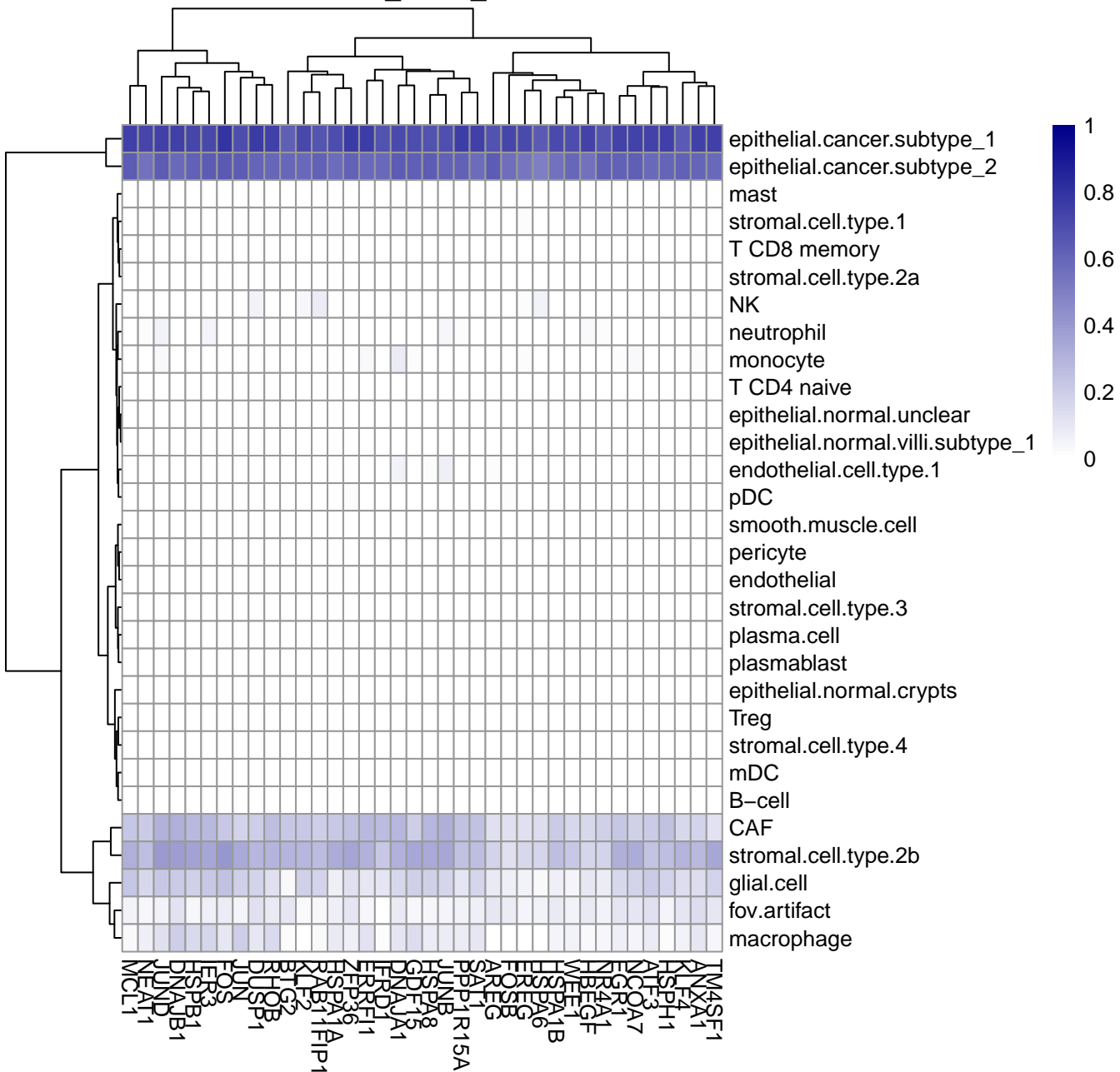
MYL9_SPARC_48



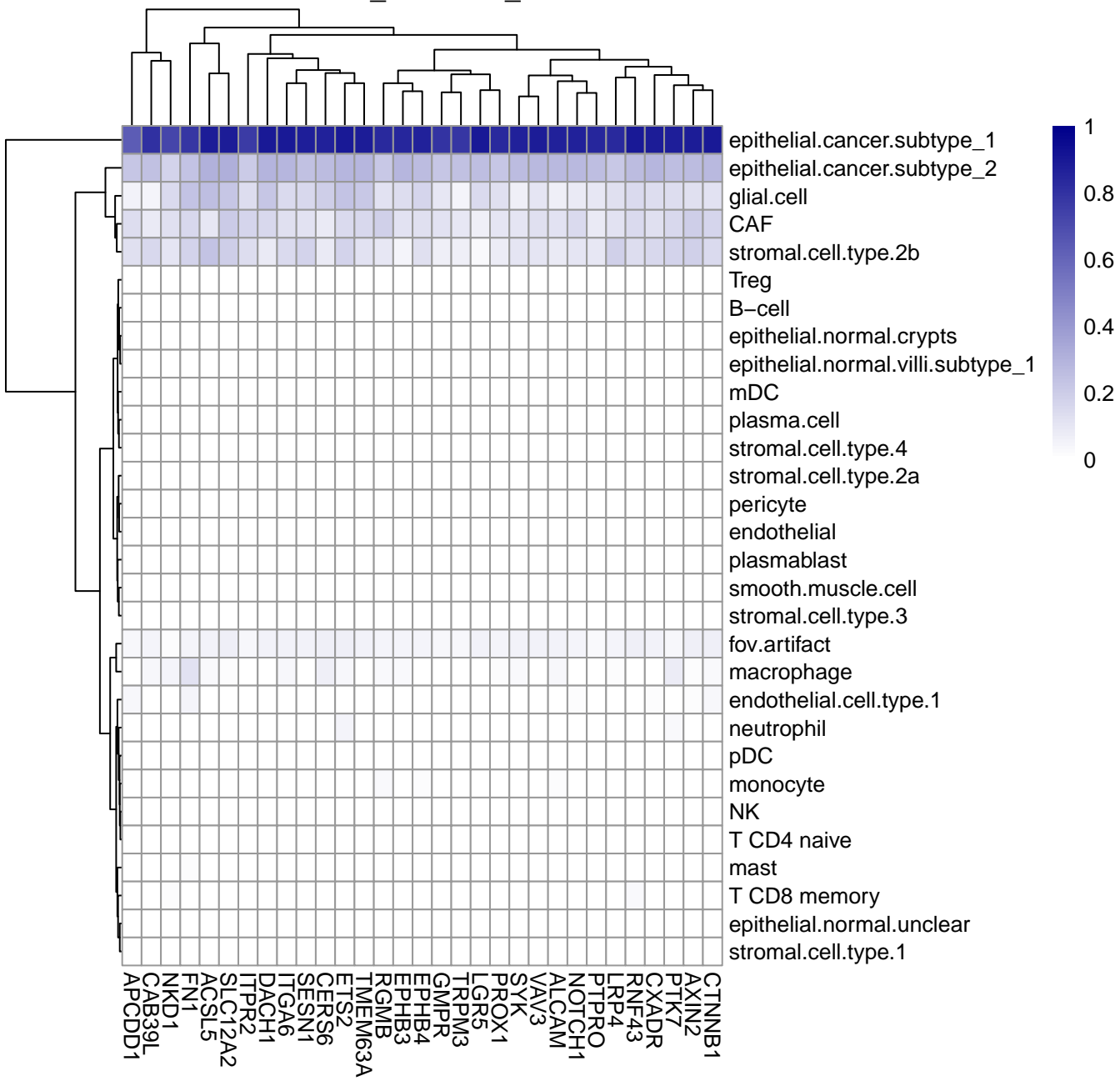
PGK1 VEGFA 47



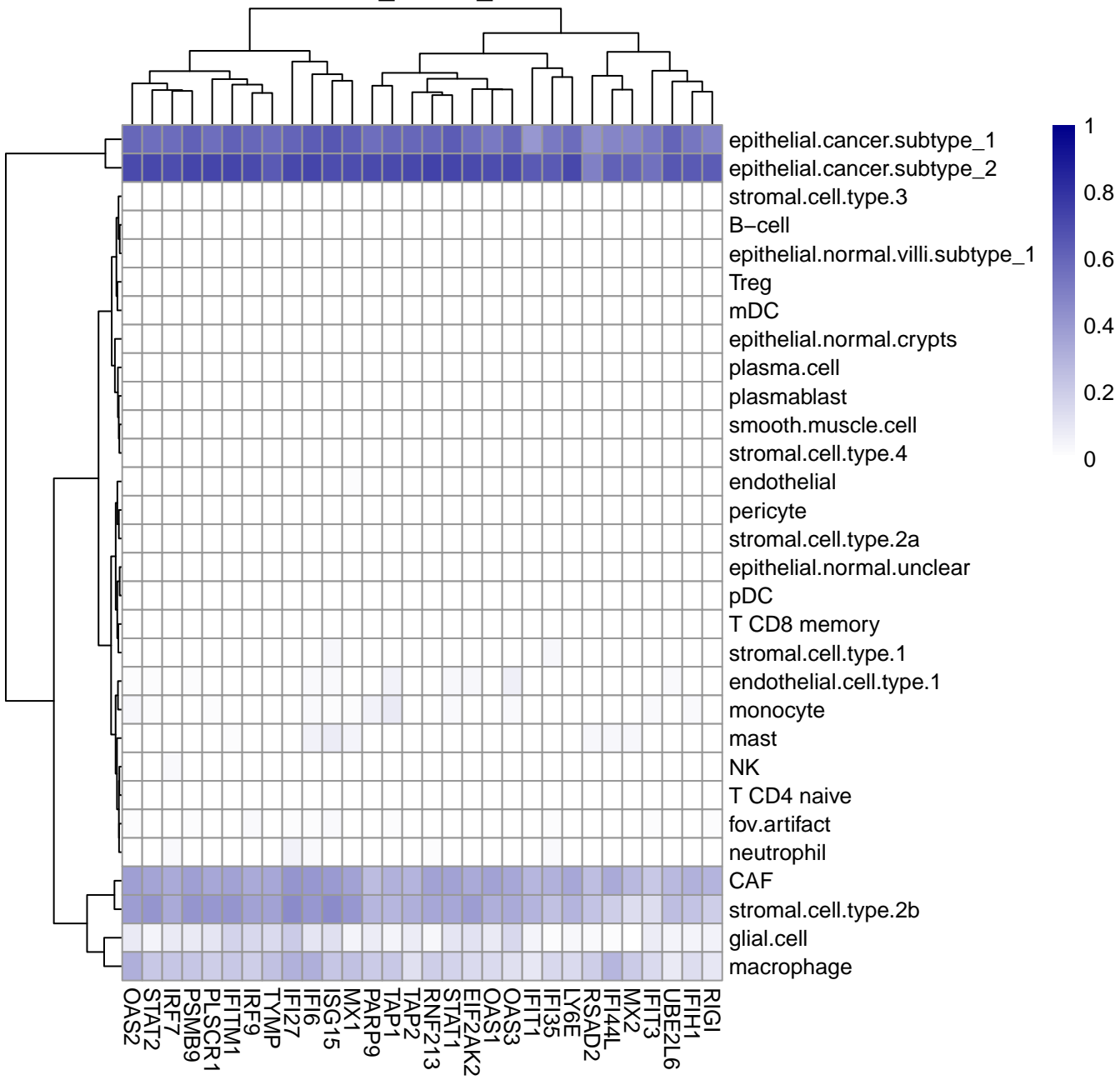
FOS_ATF3_38



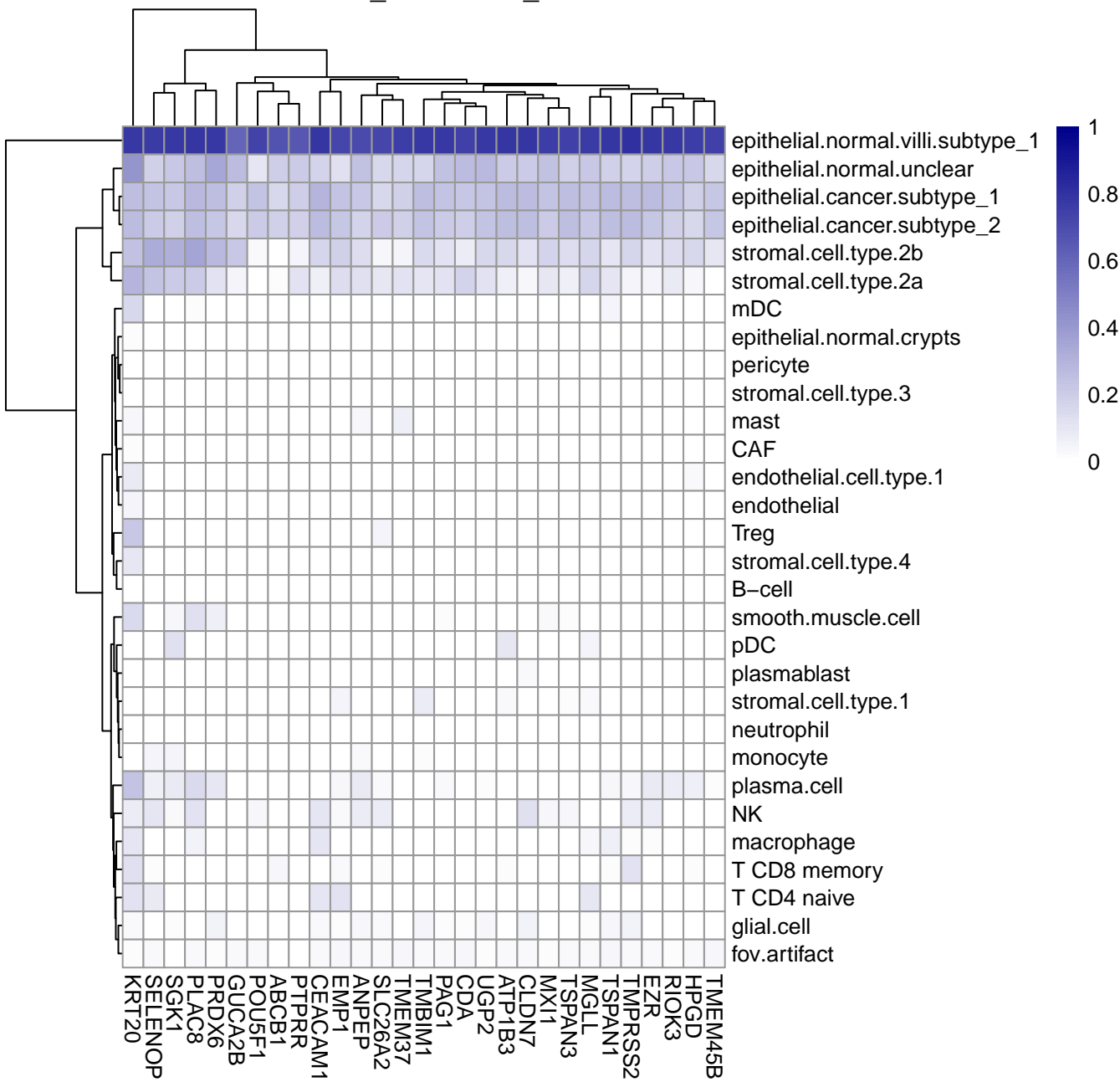
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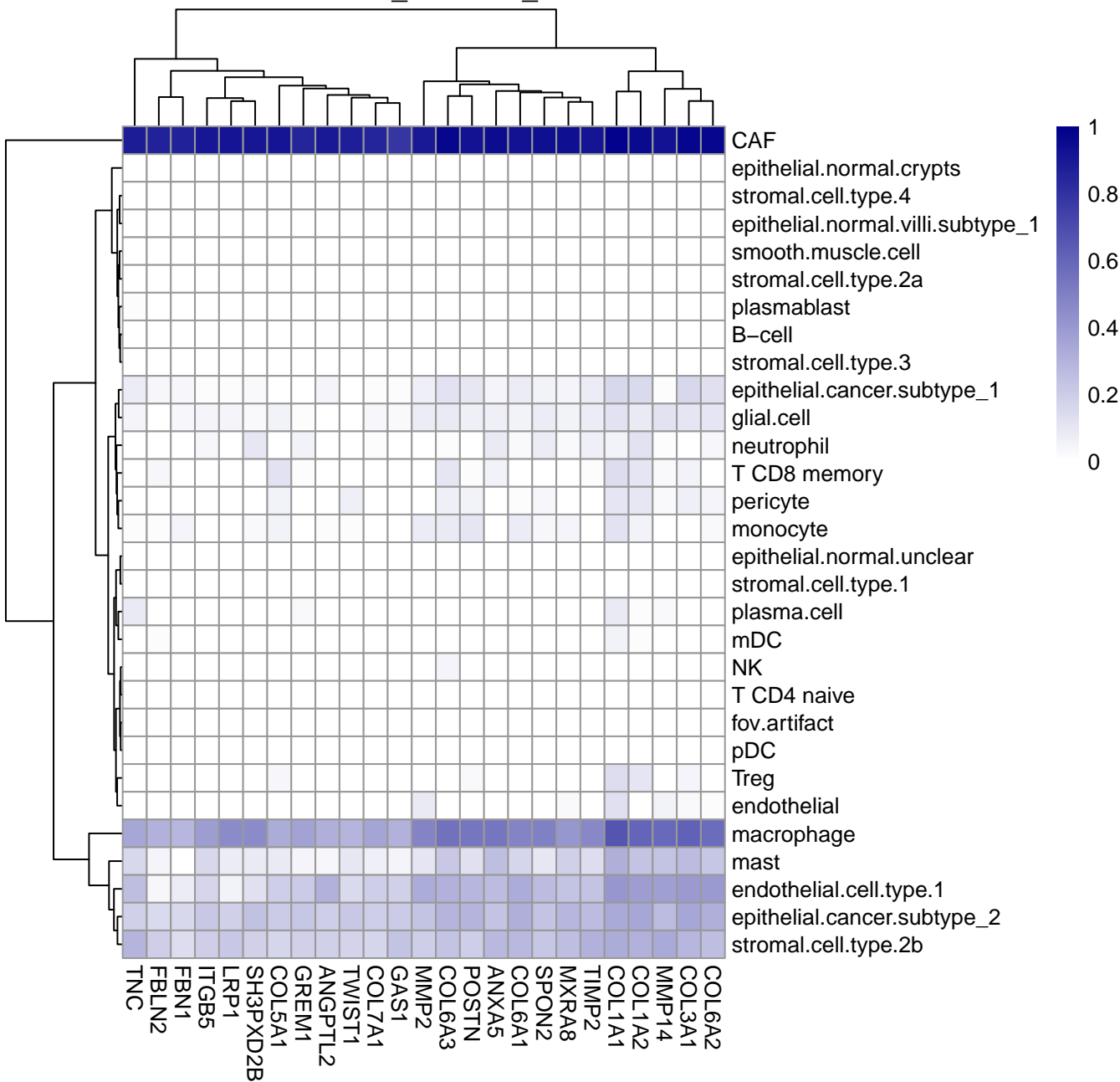
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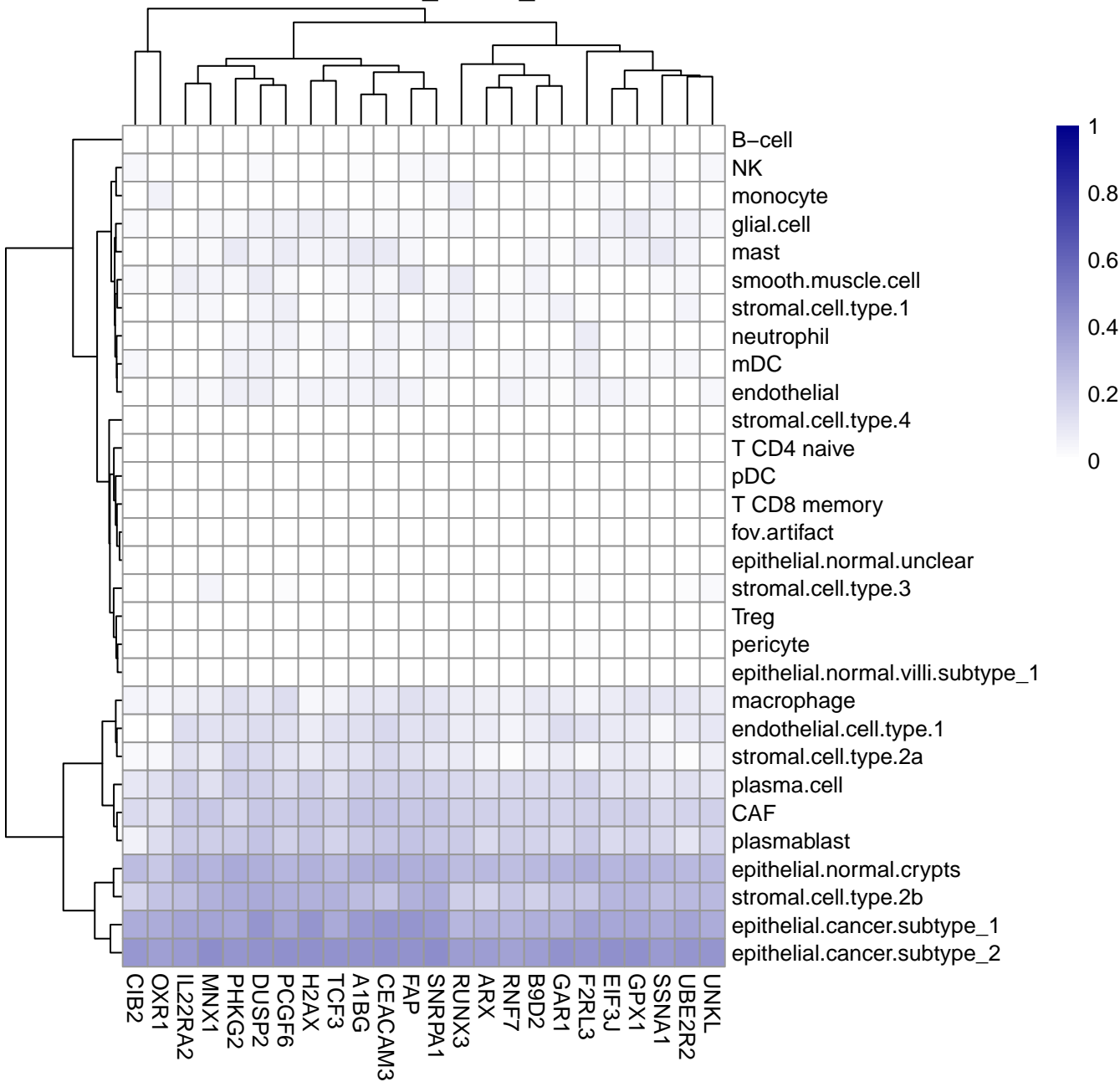
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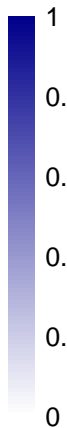


COL1A1_COL3A1_25

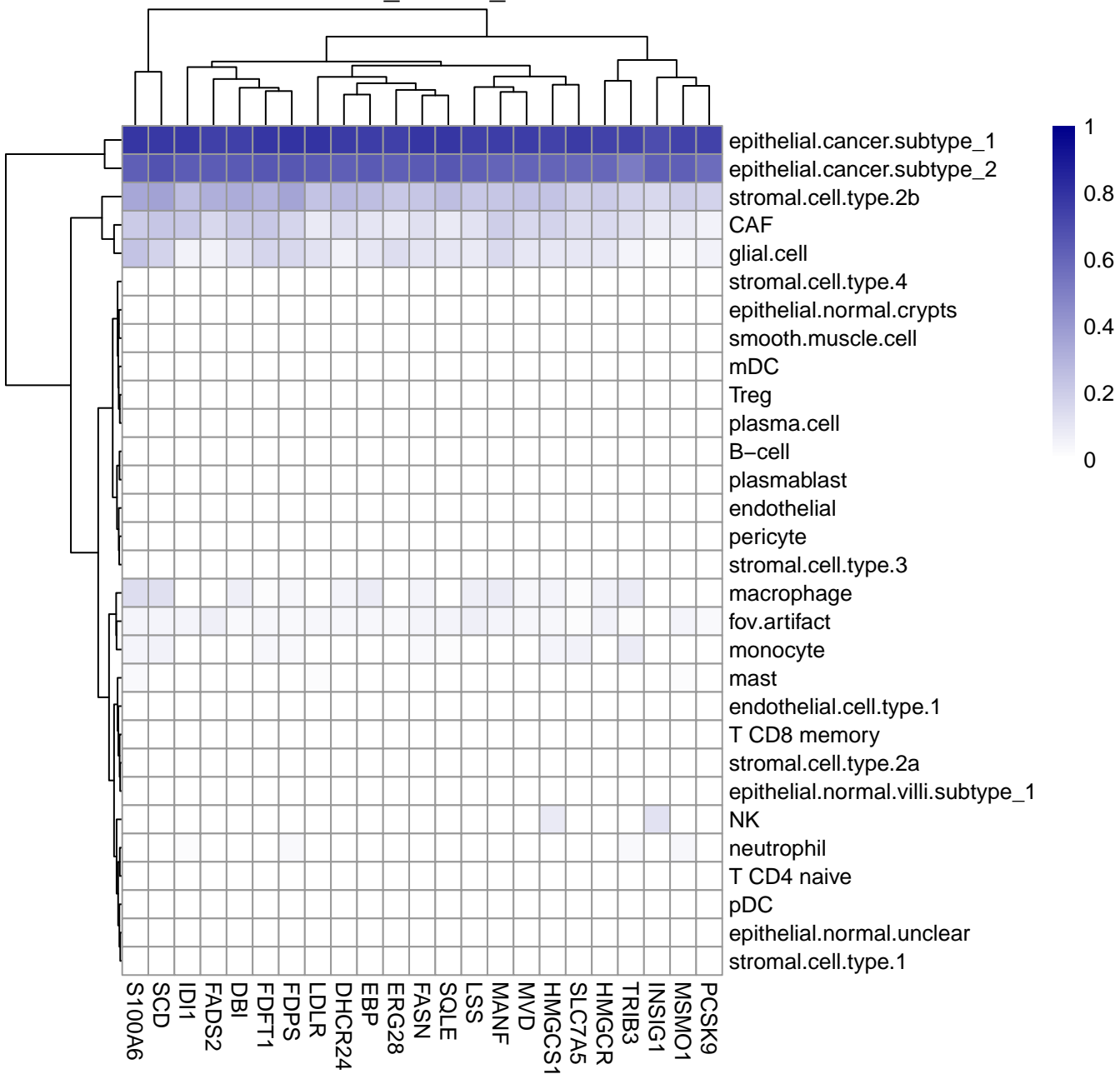


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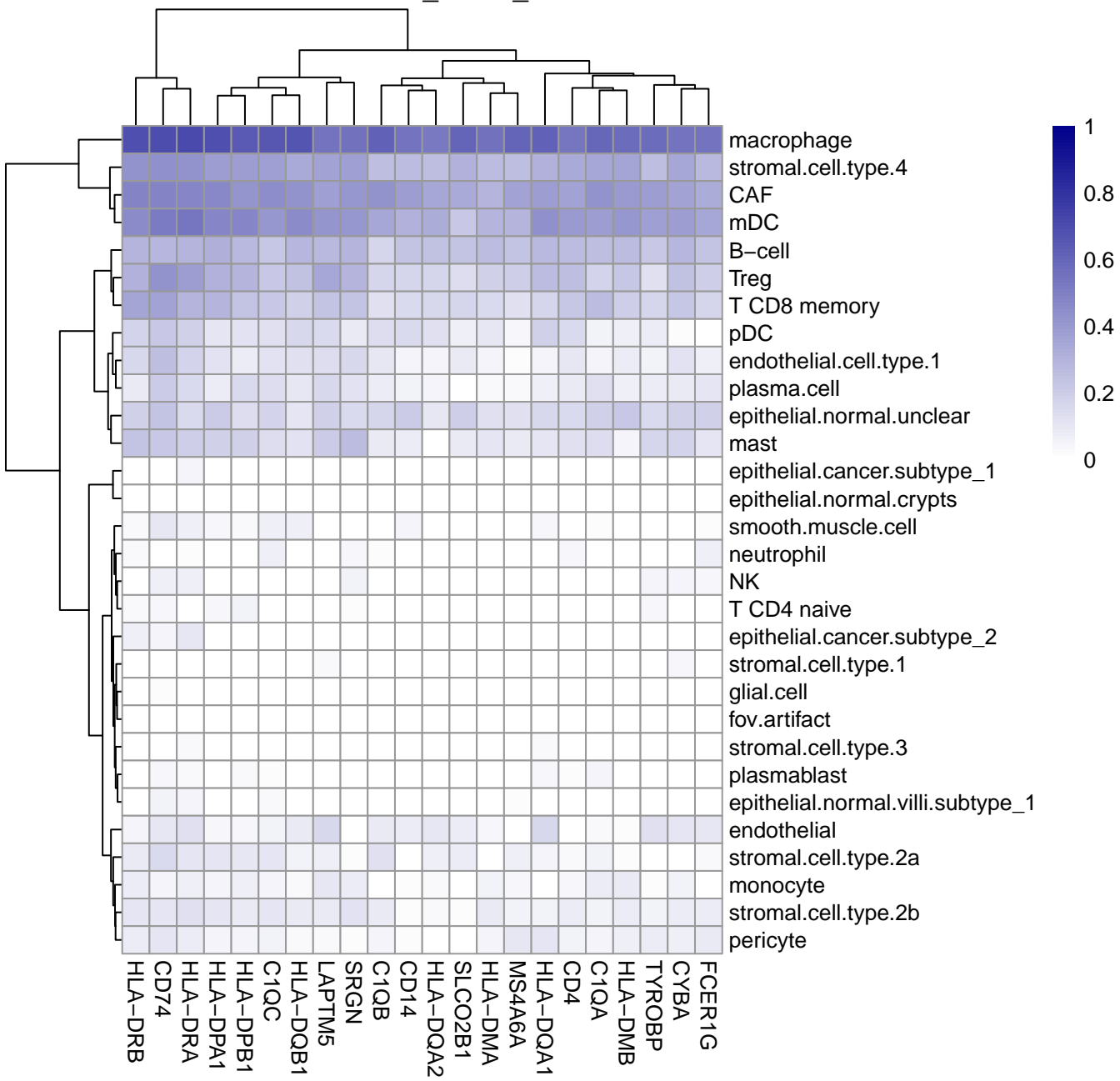




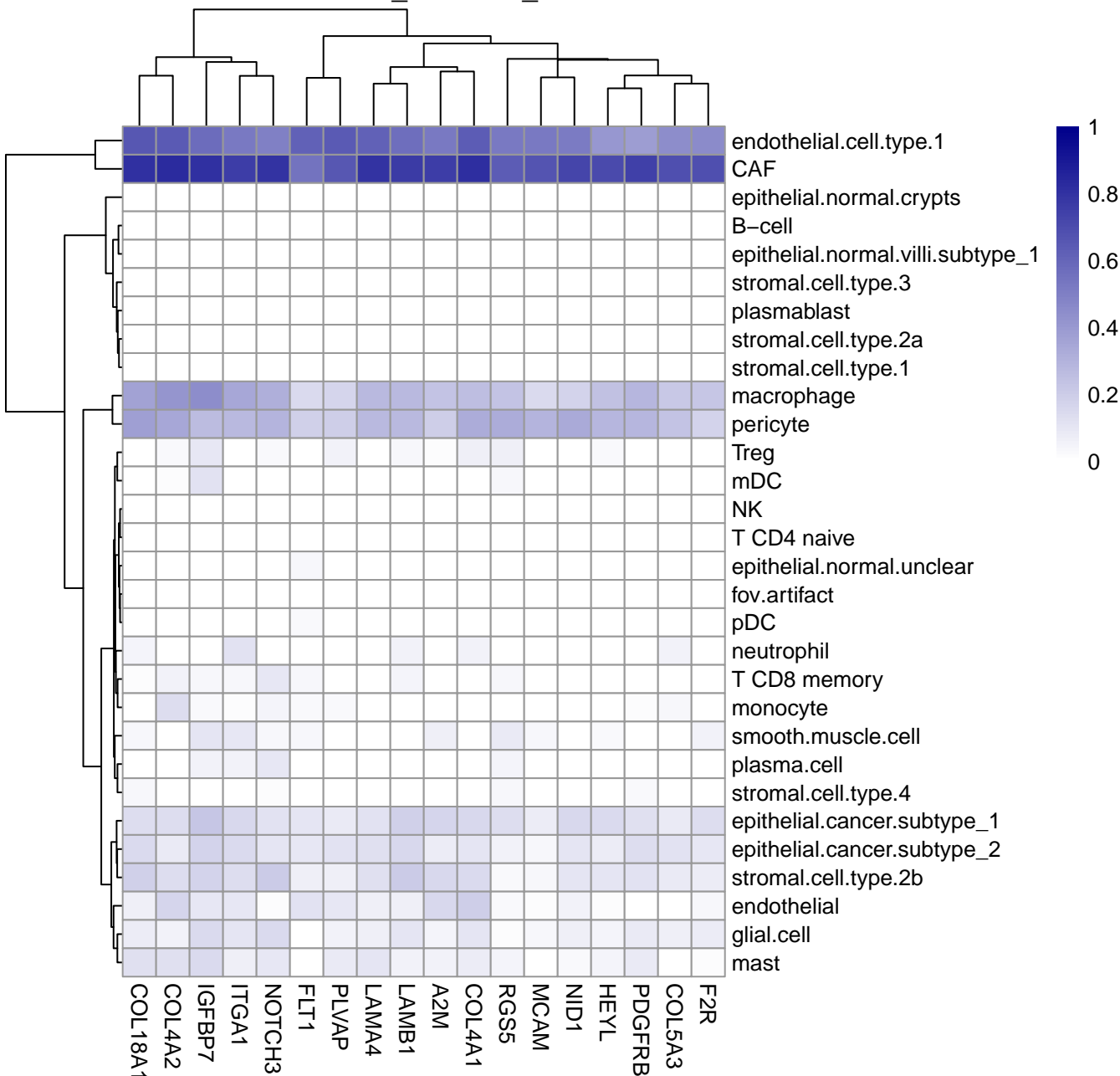
FDPS FDFT1 23



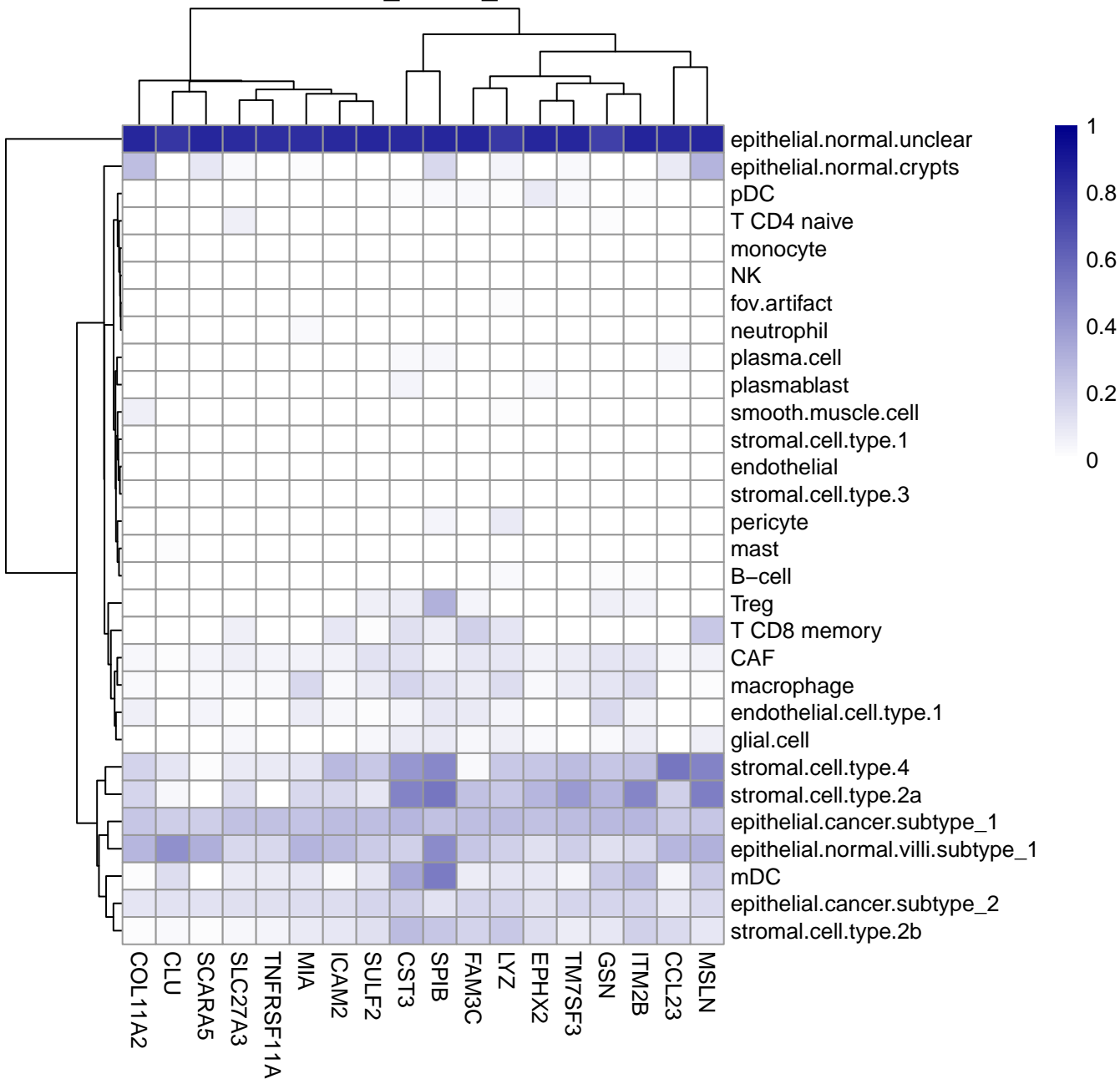
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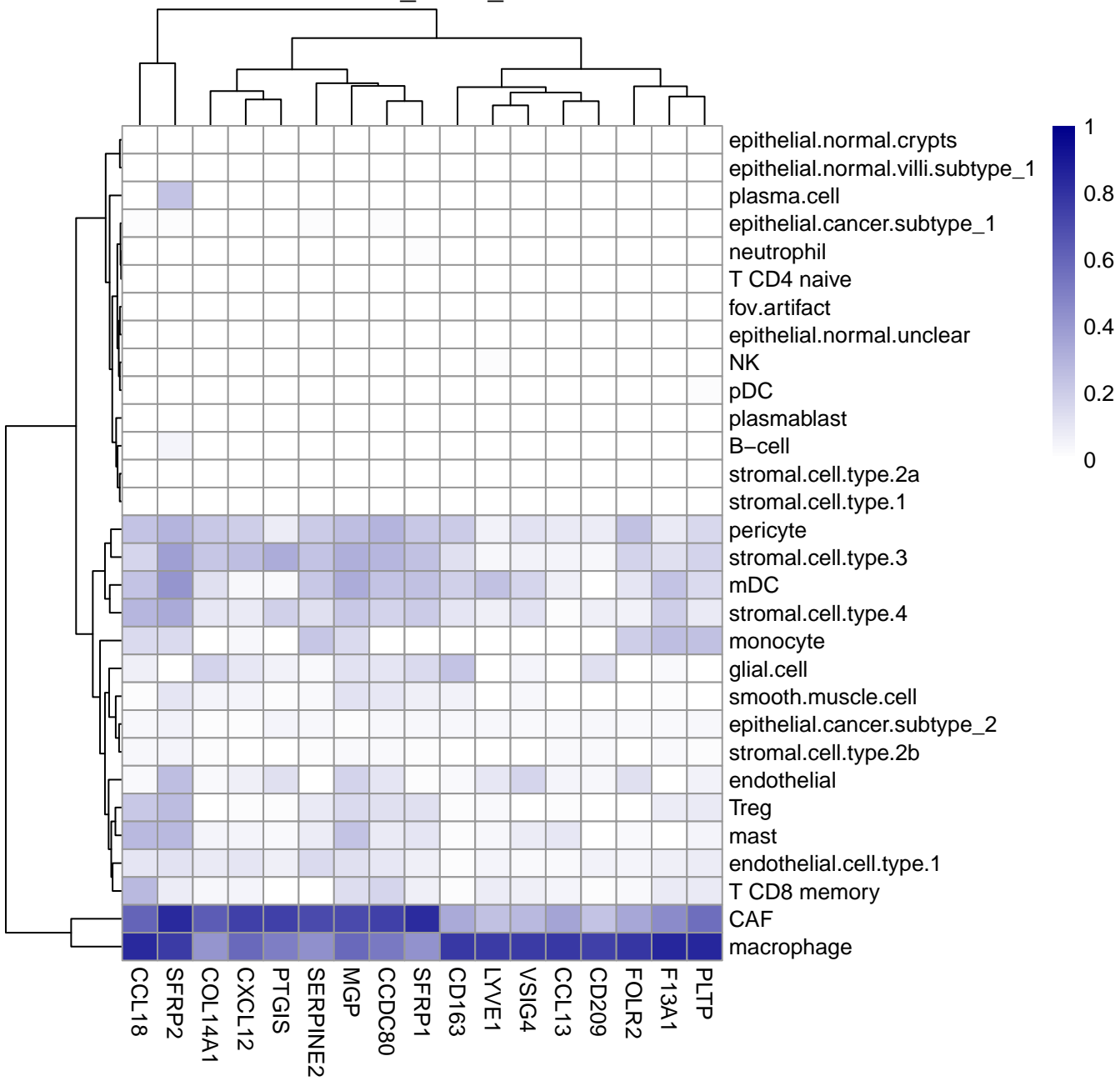
COL4A2_COL4A1_18



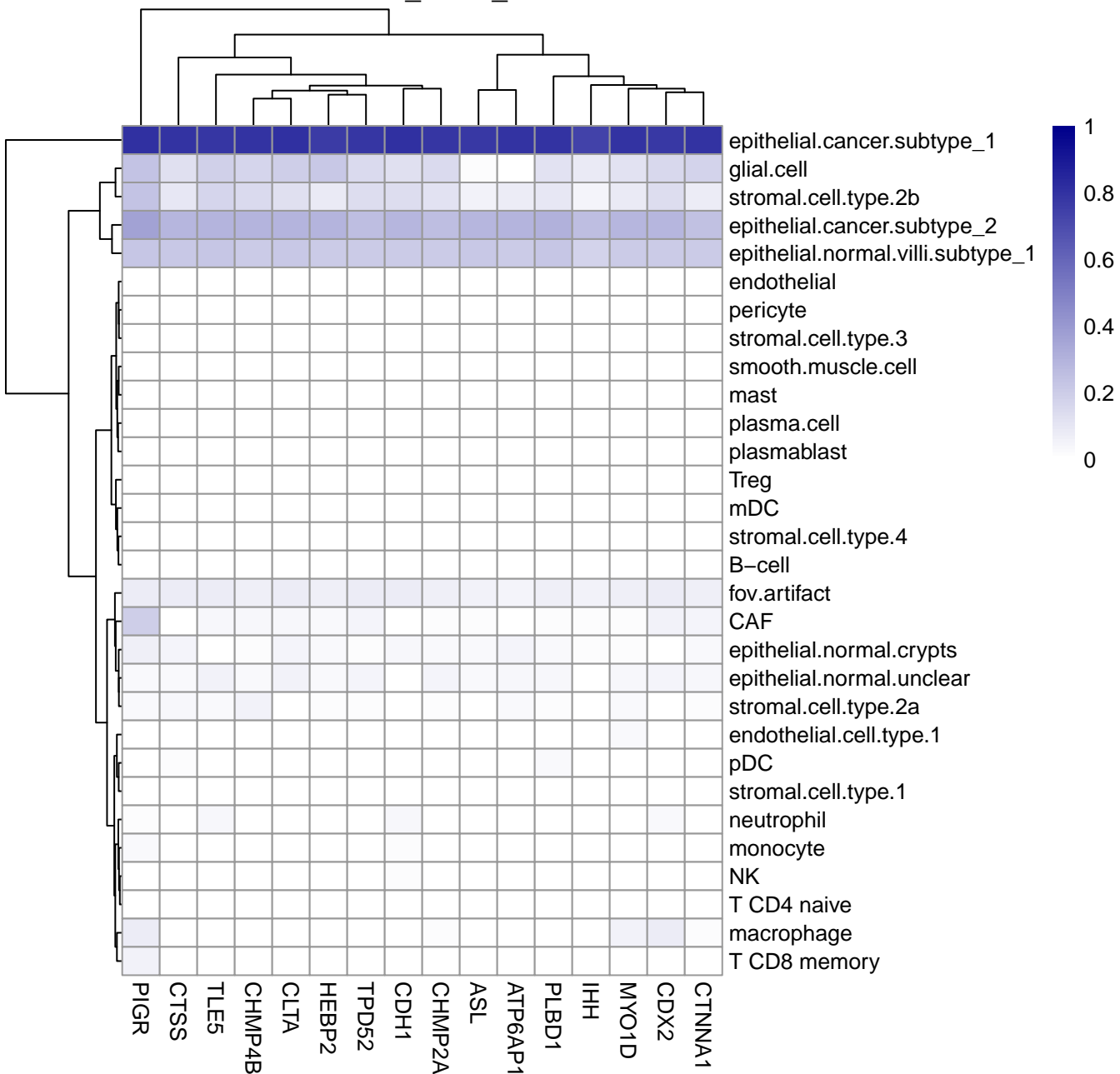
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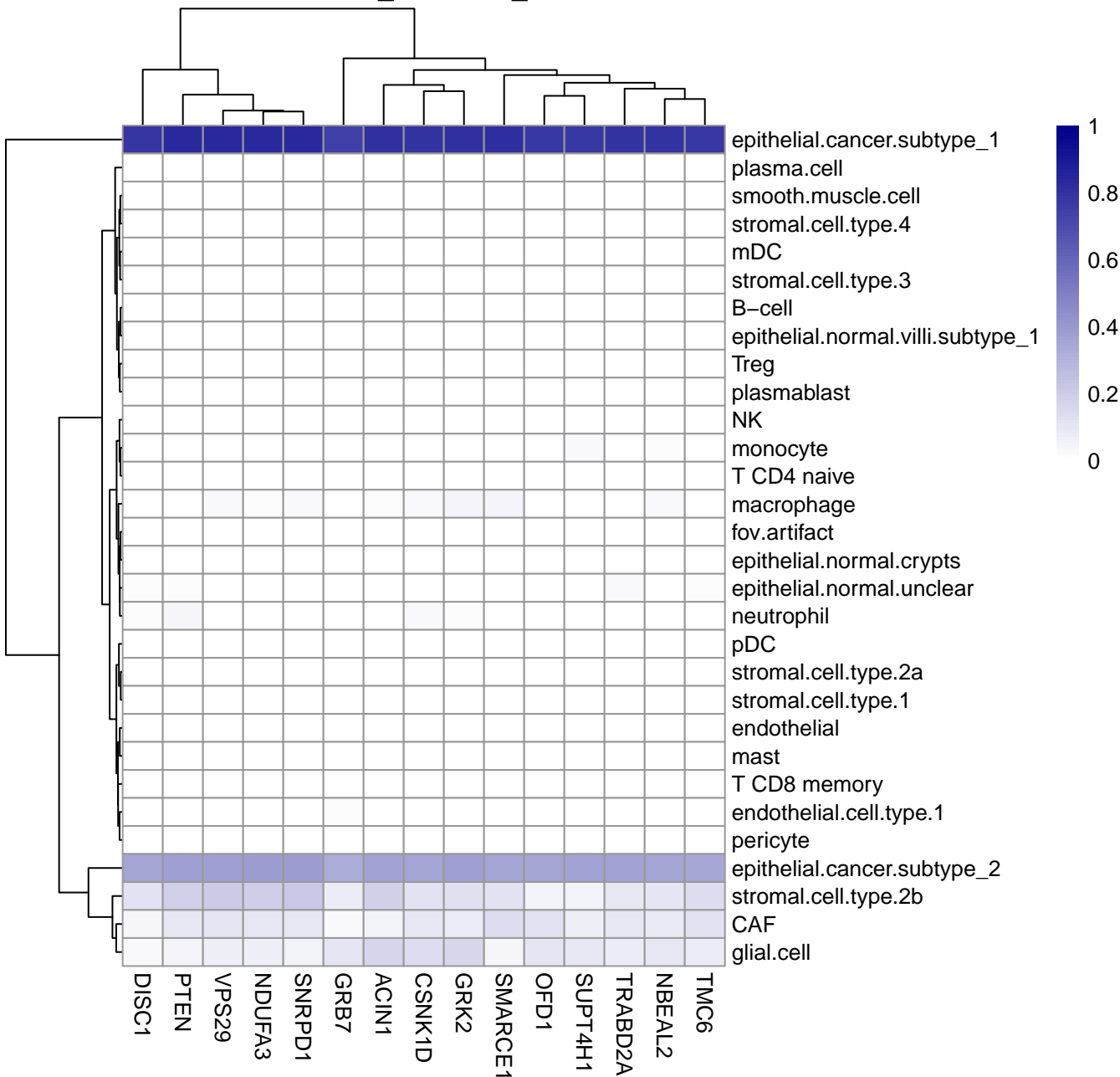
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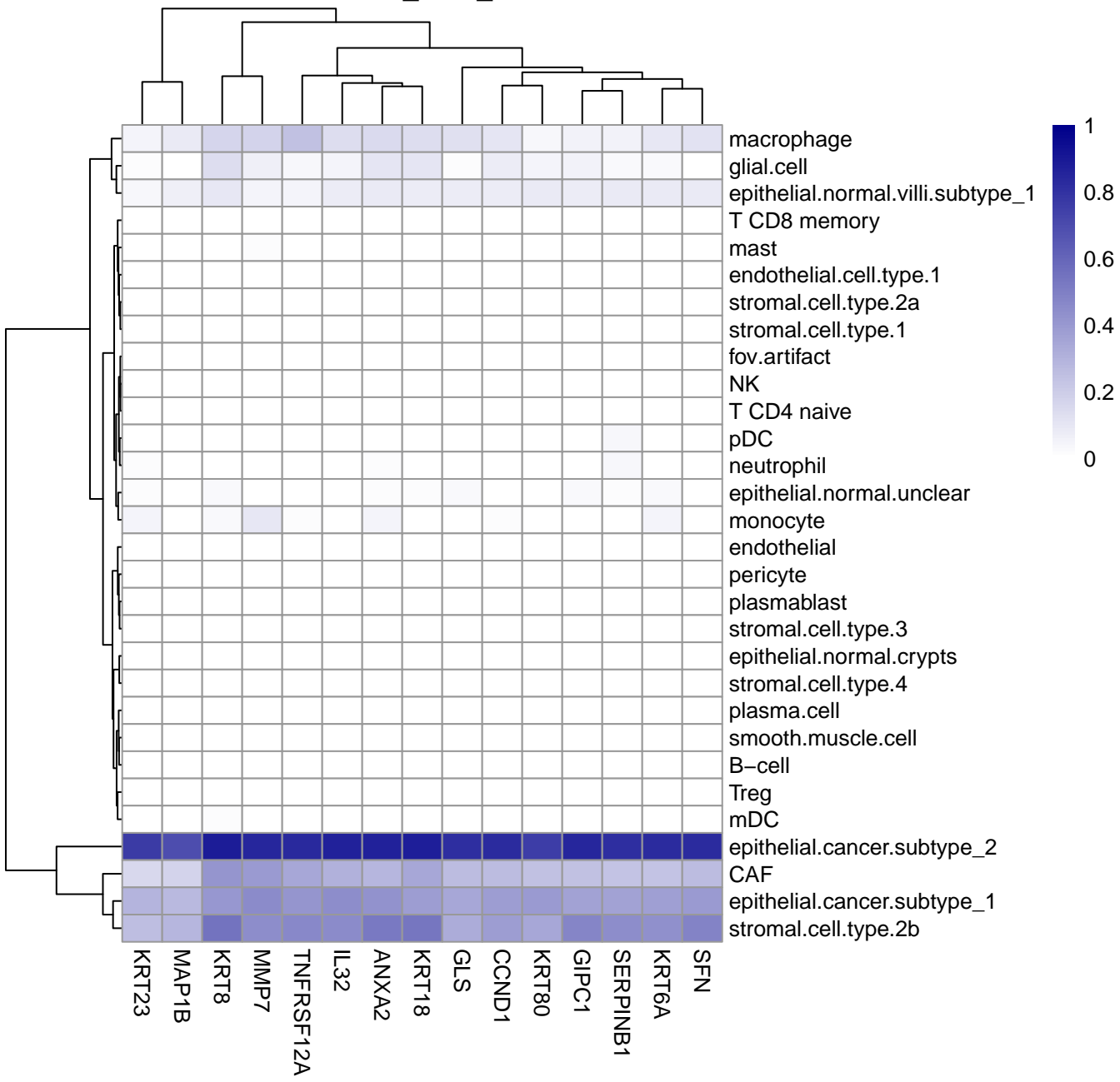
MYO1D_PIGR_16



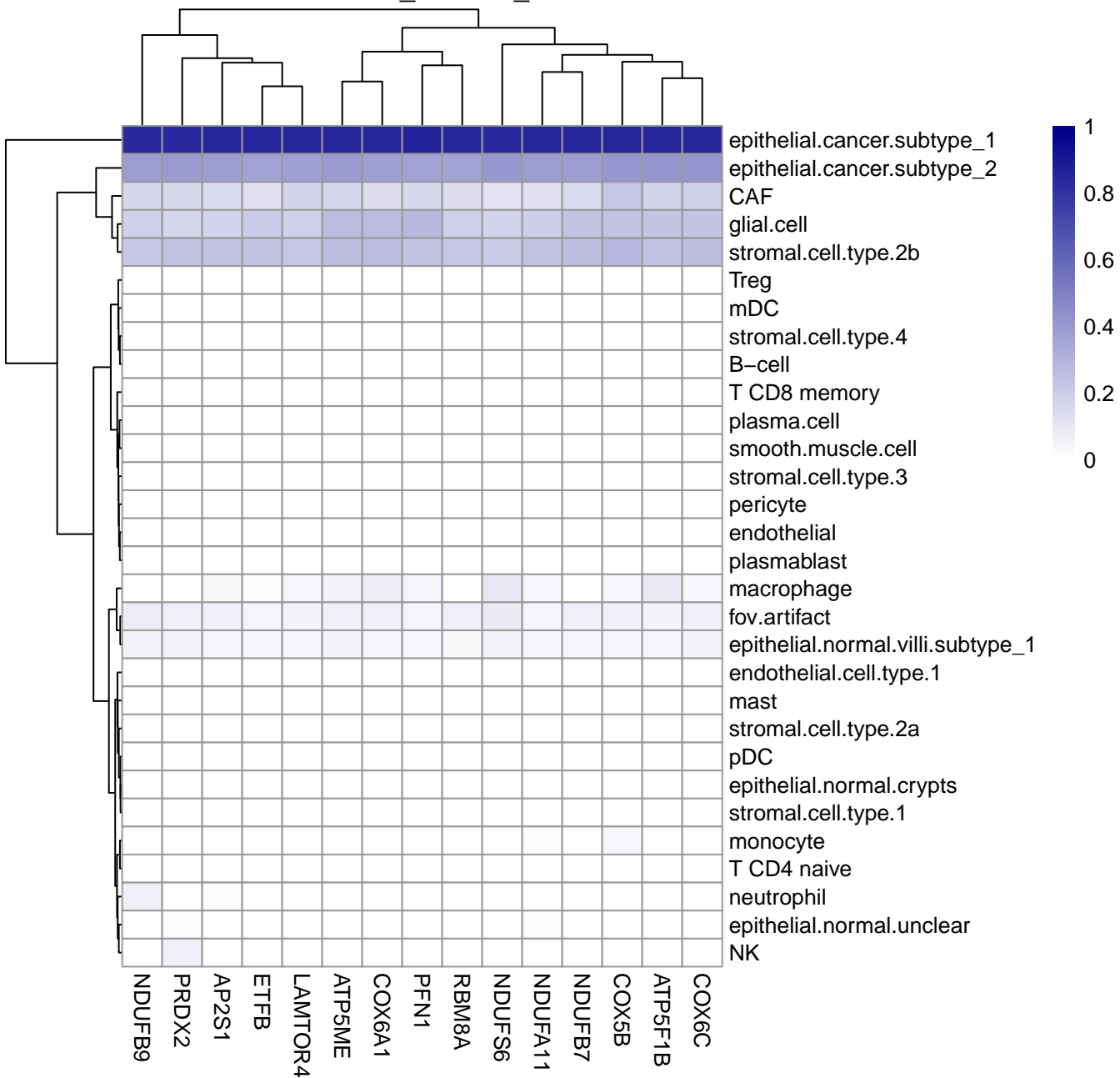
VPS29_SNRPD1_15



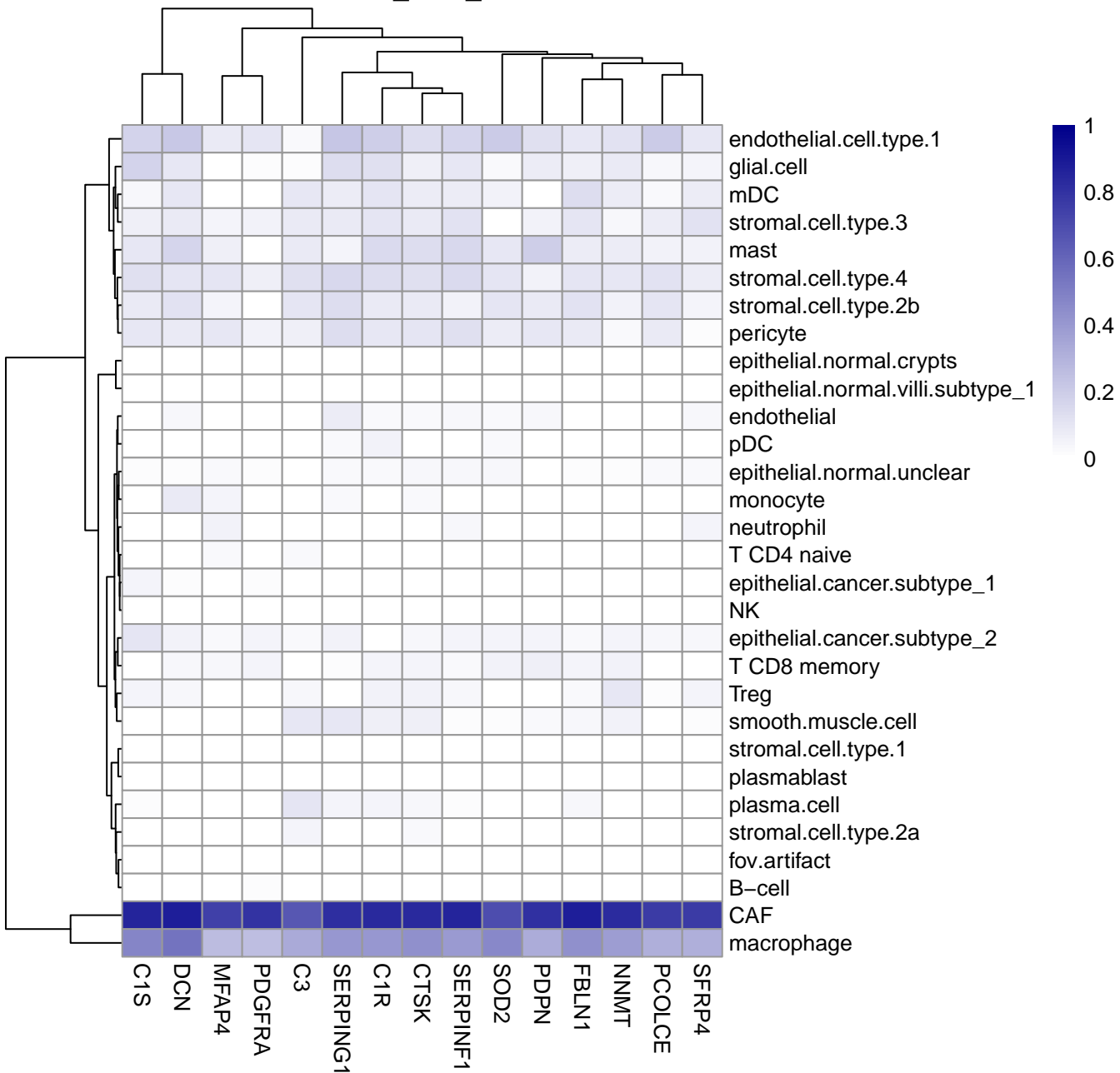
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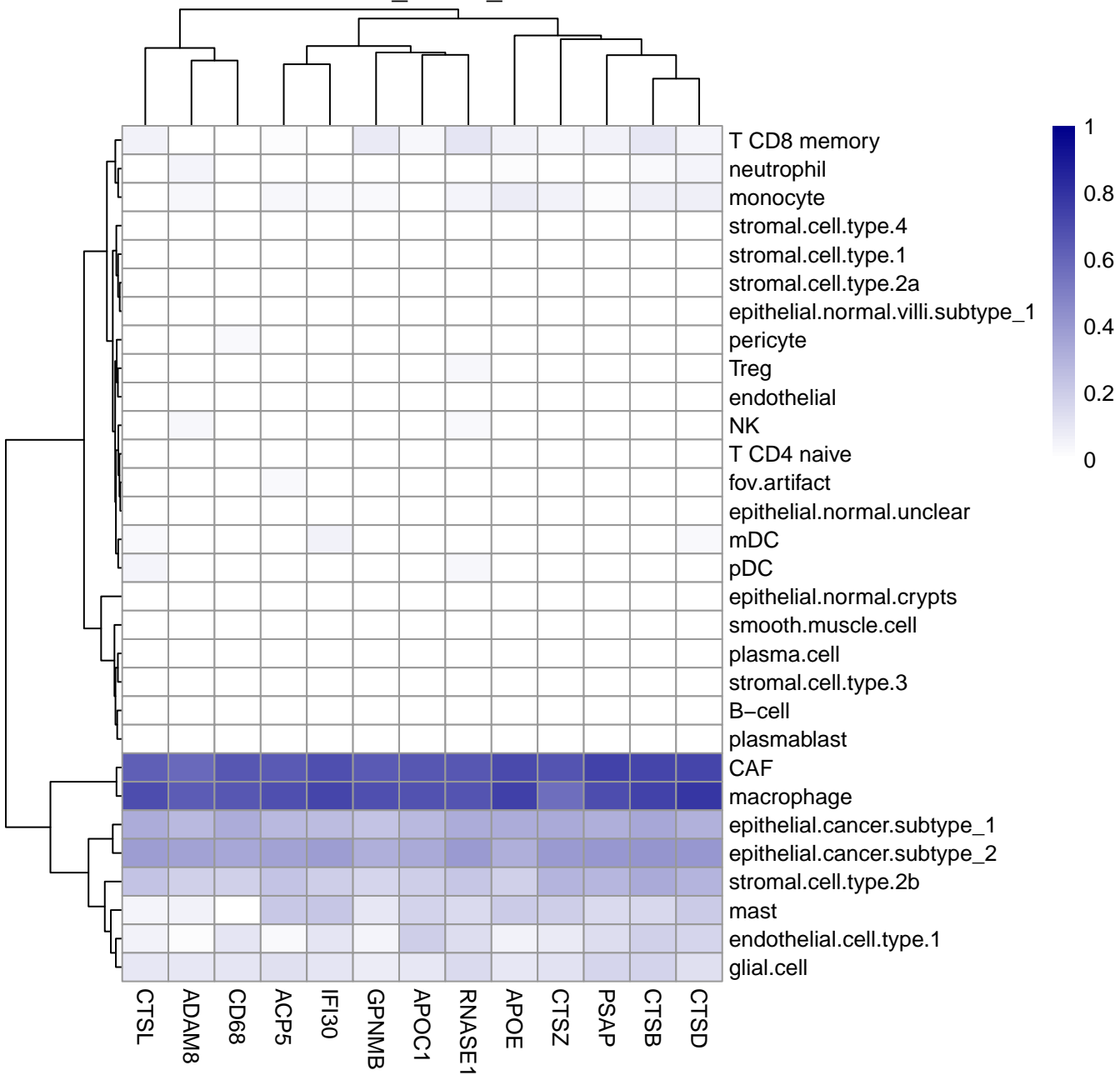
NDUFB7 COX6C 15



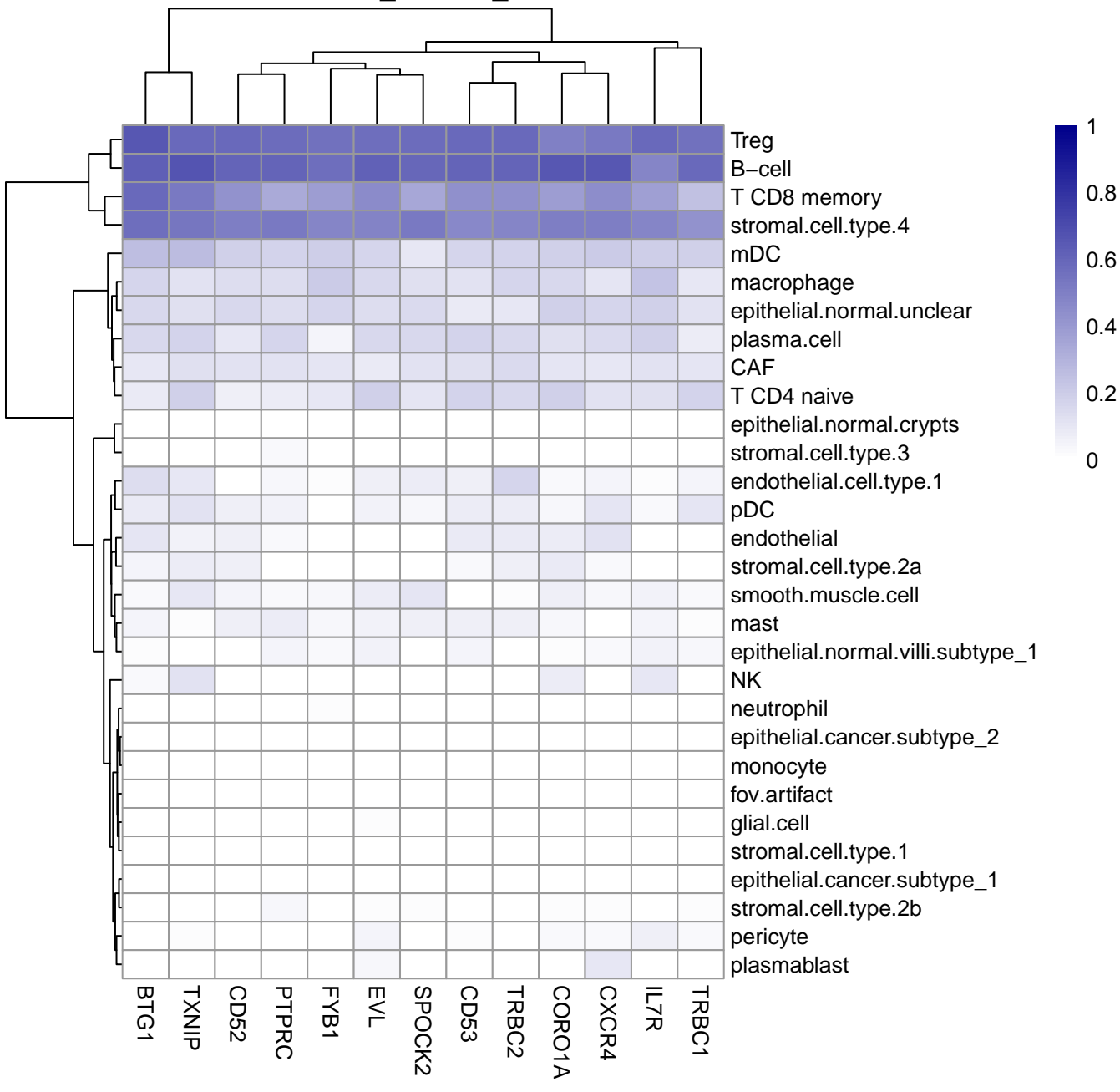
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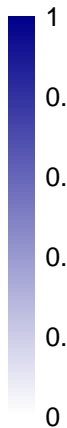


APOE_CTSD_13

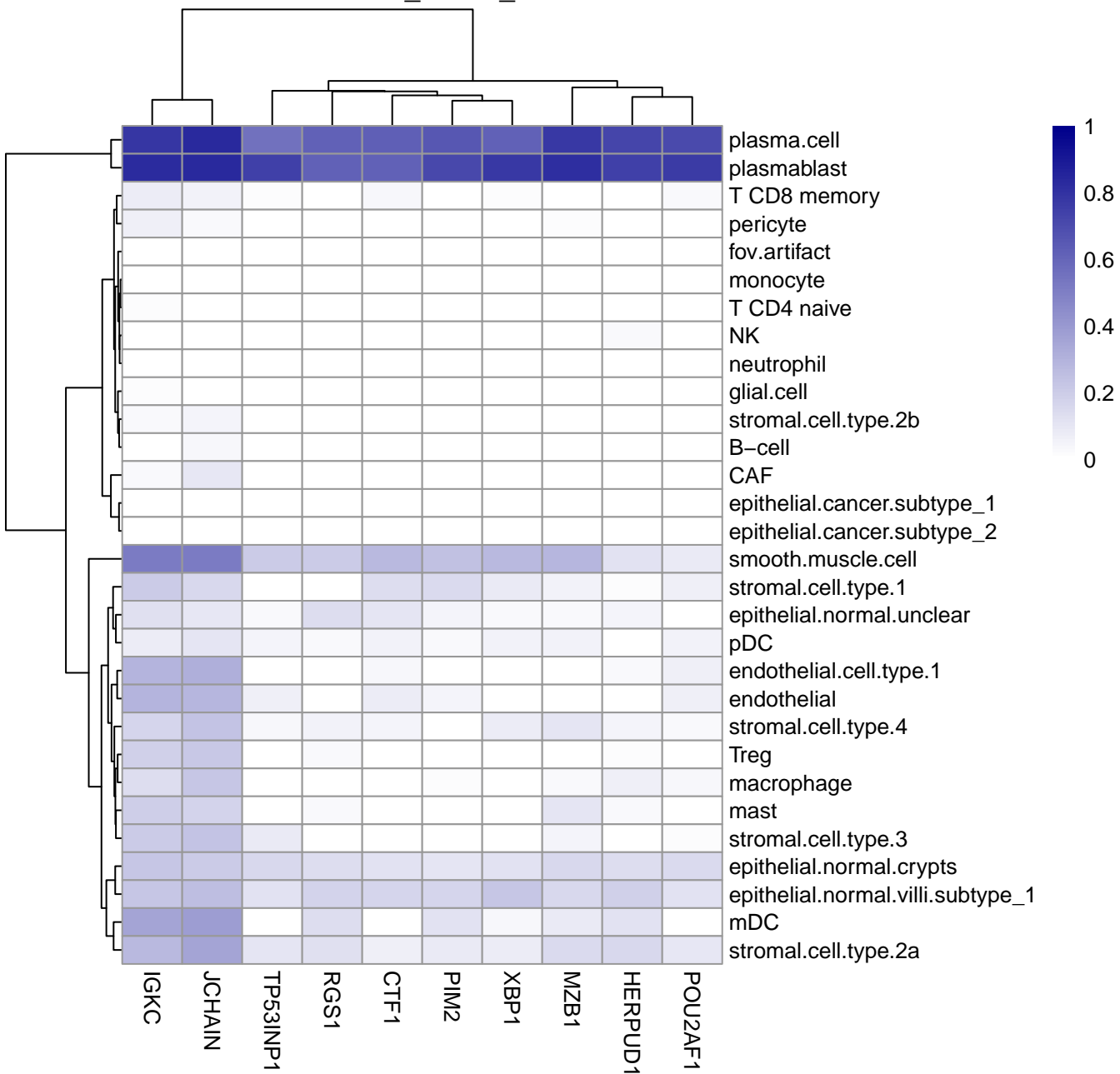


CD53_TRBC2_13

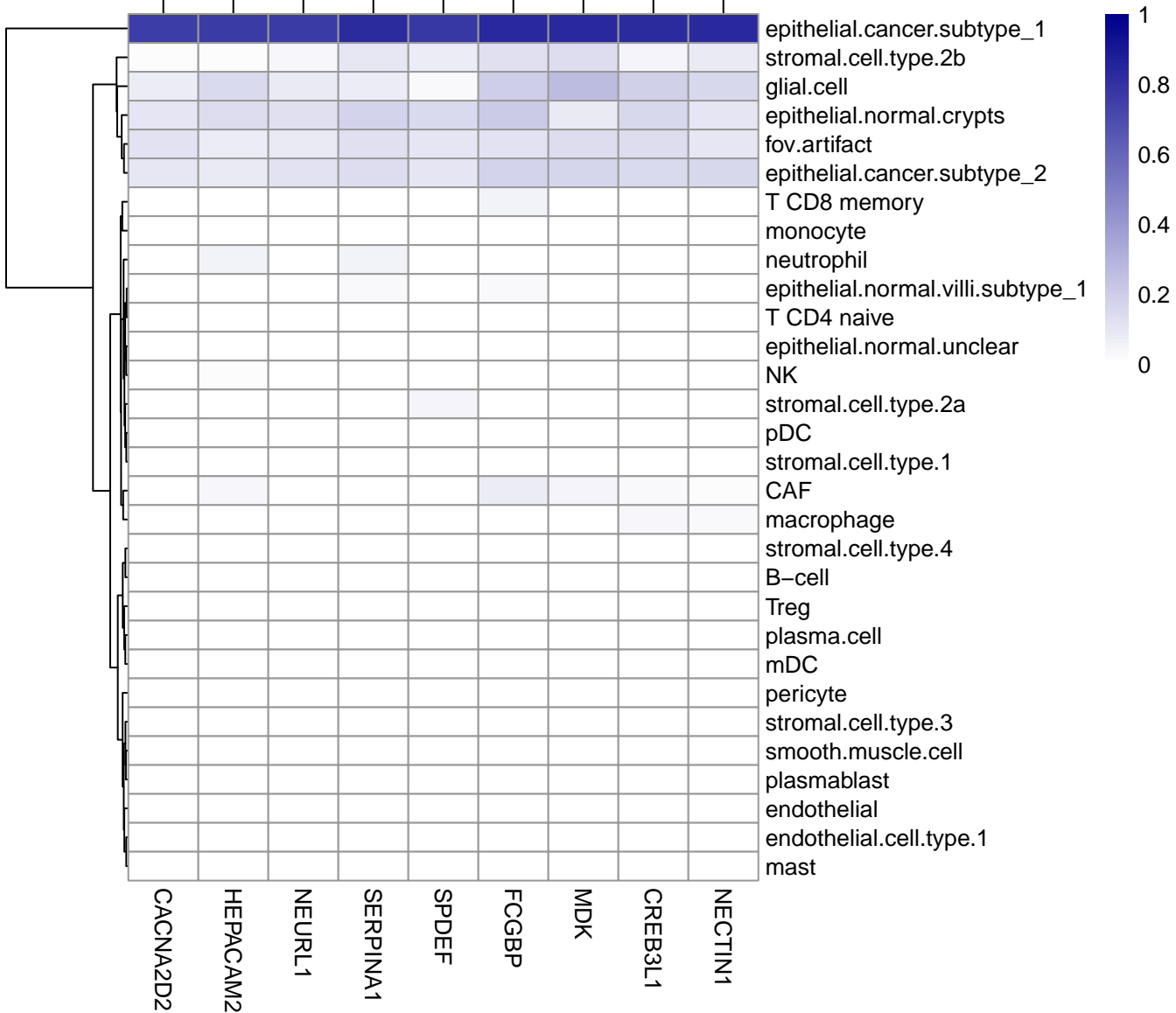


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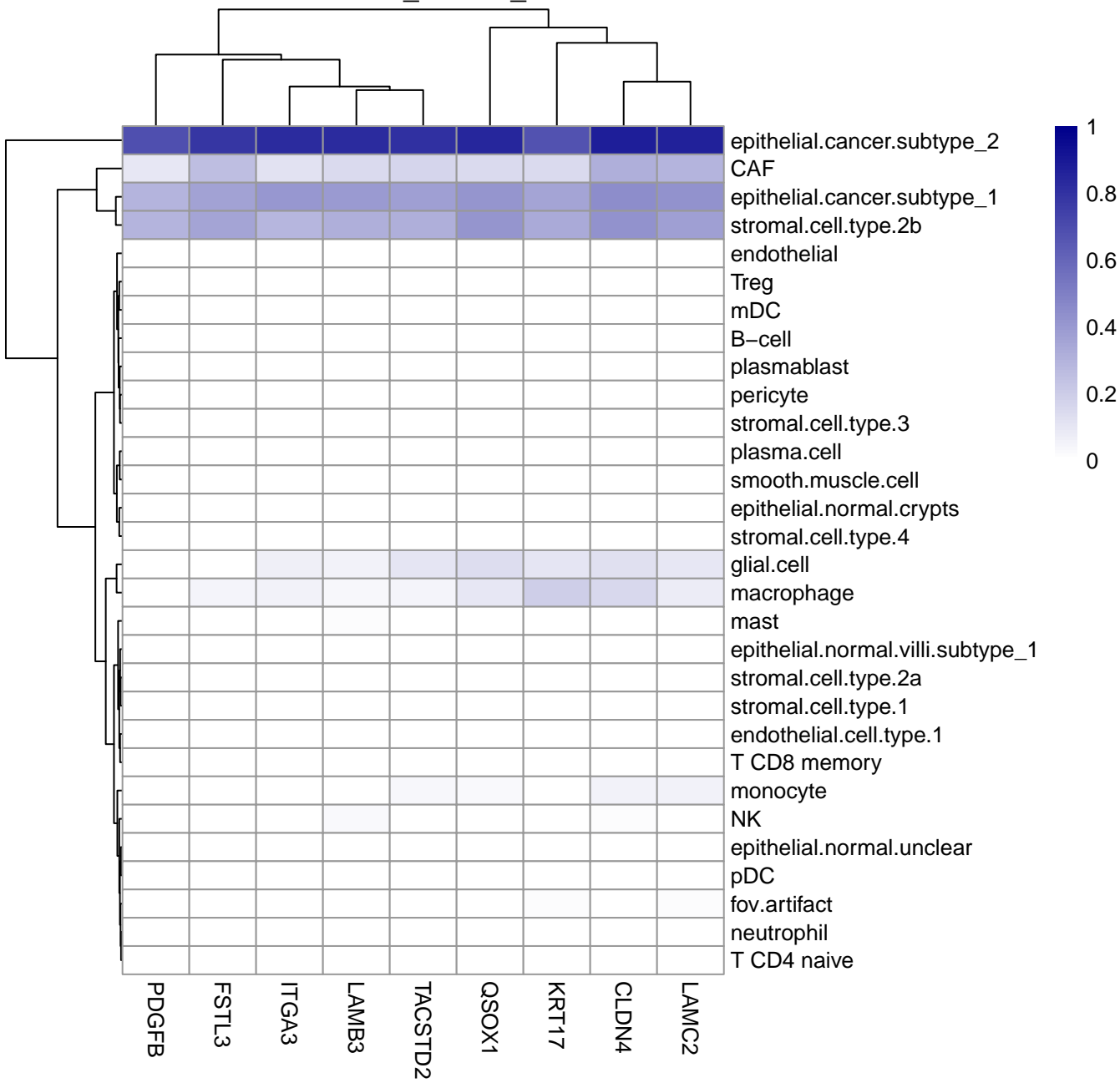
JCHAIN MZB1 10



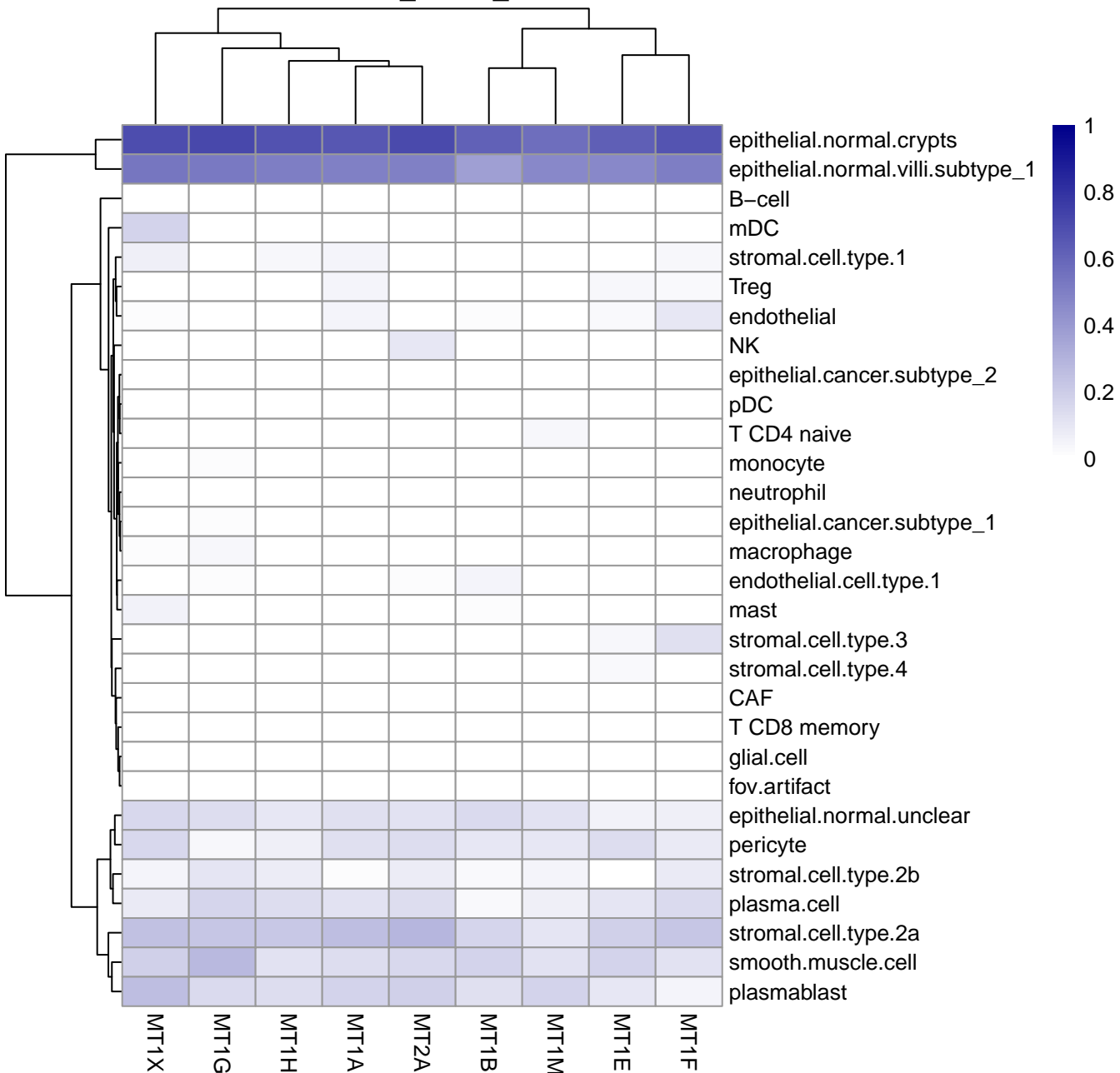
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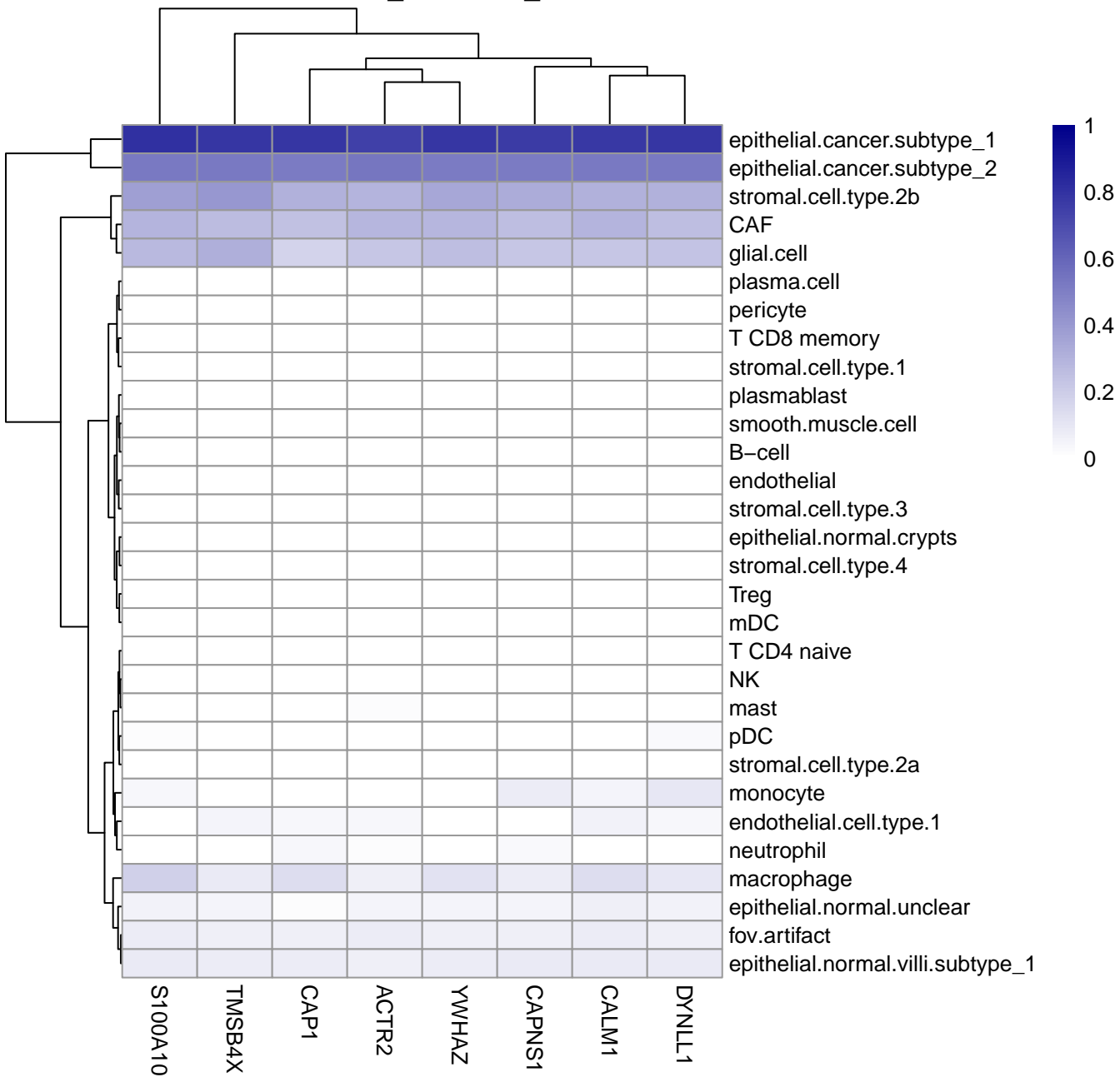
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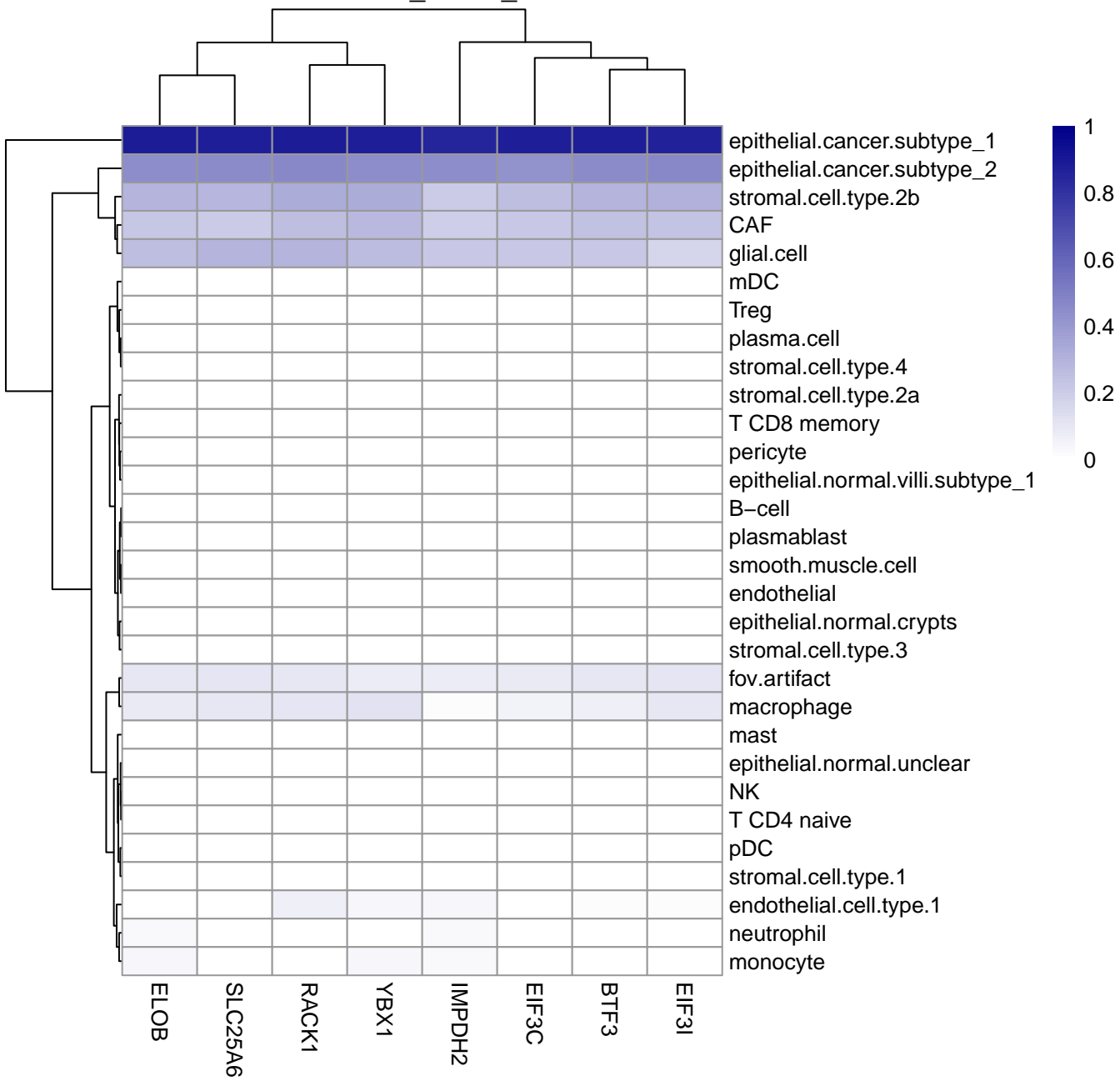
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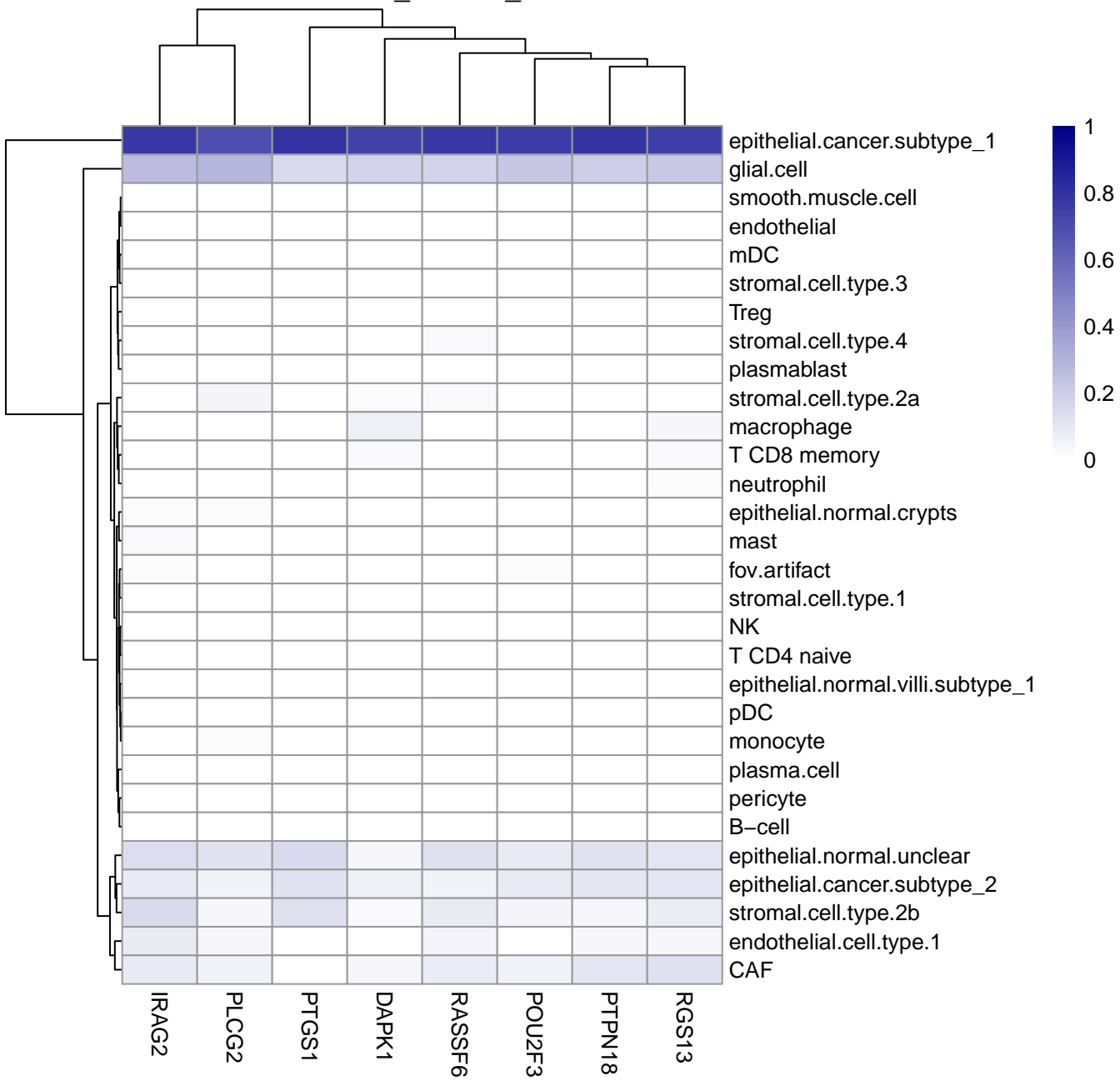
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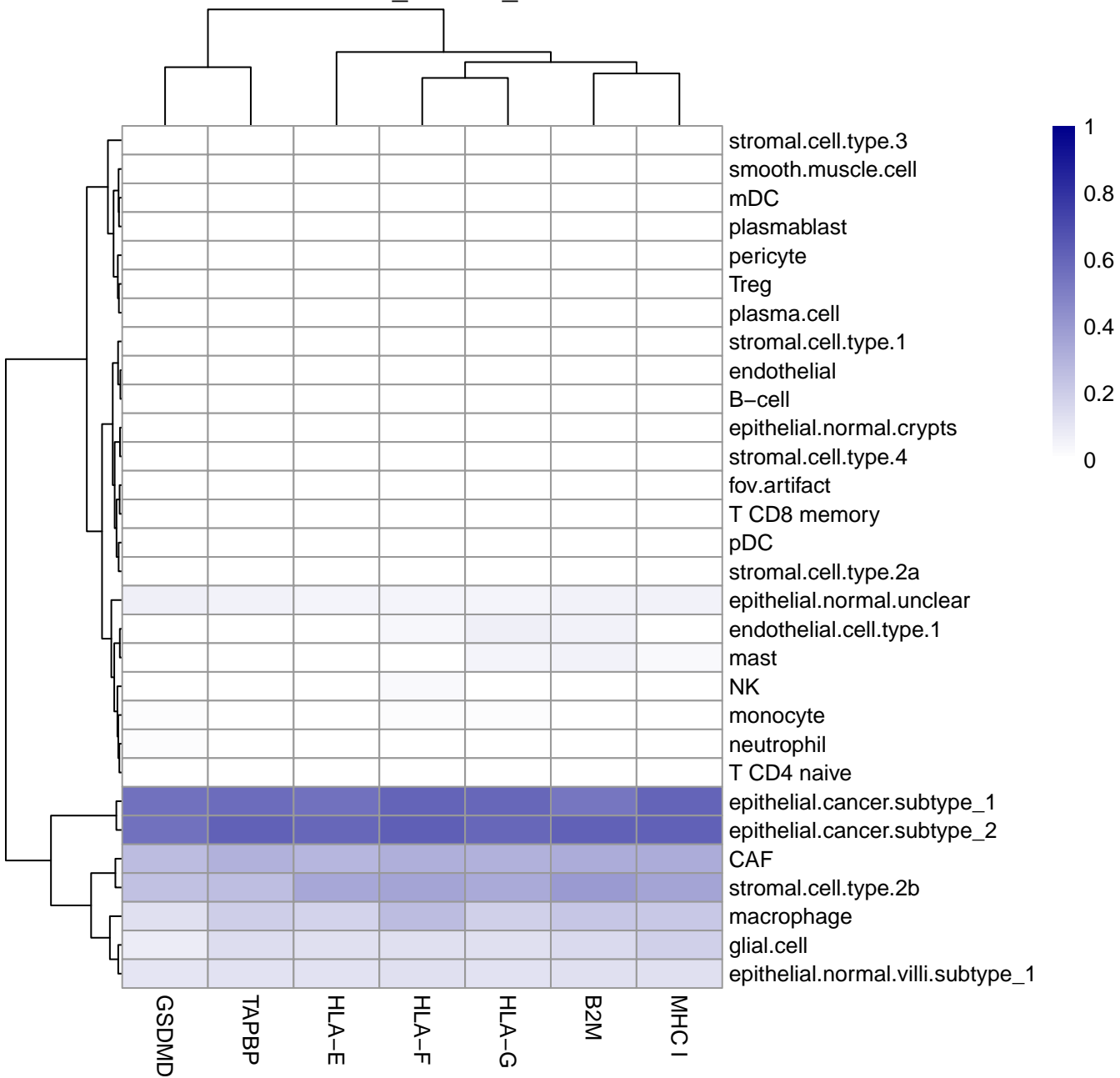
RACK1_YBX1_8



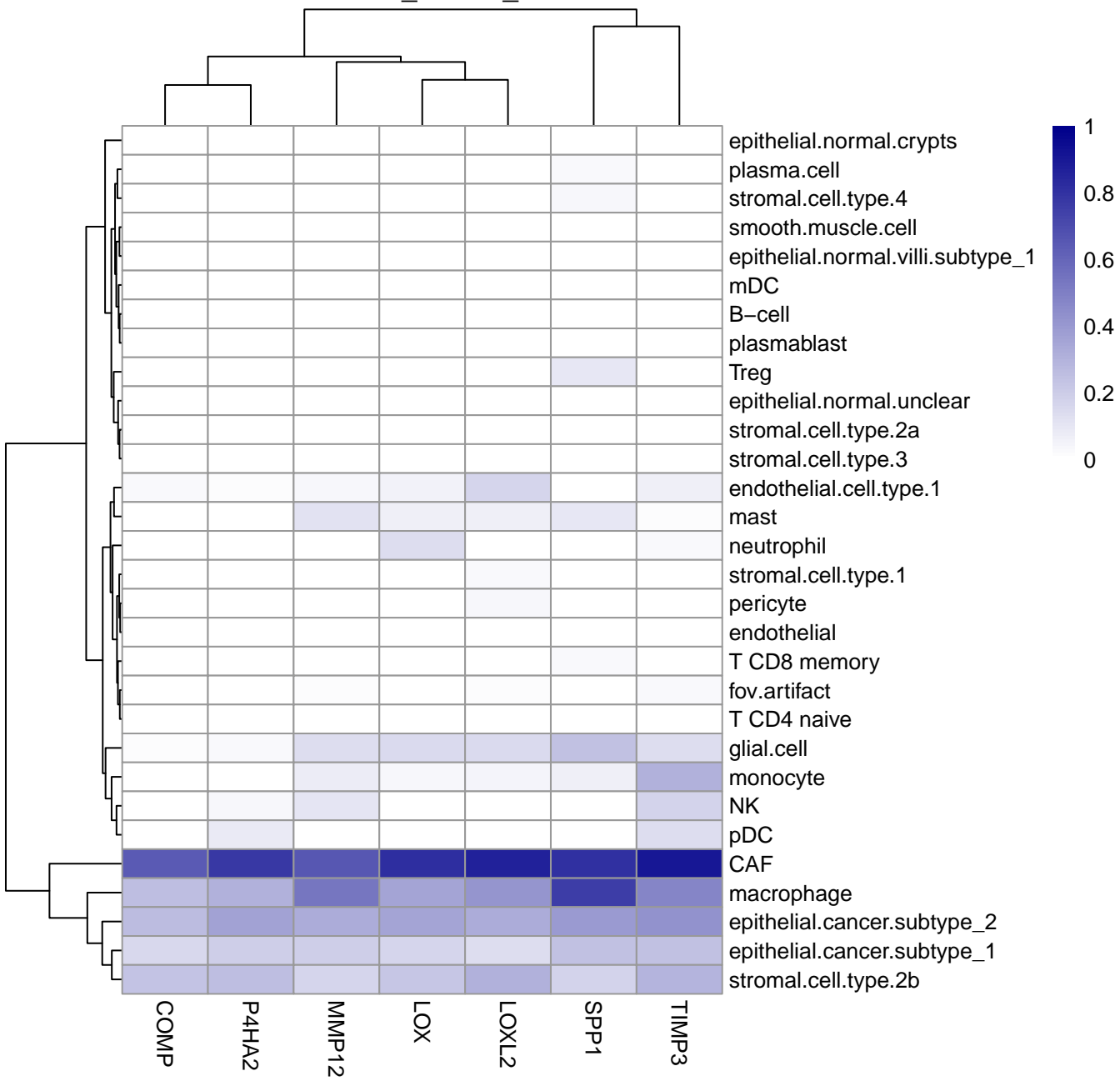
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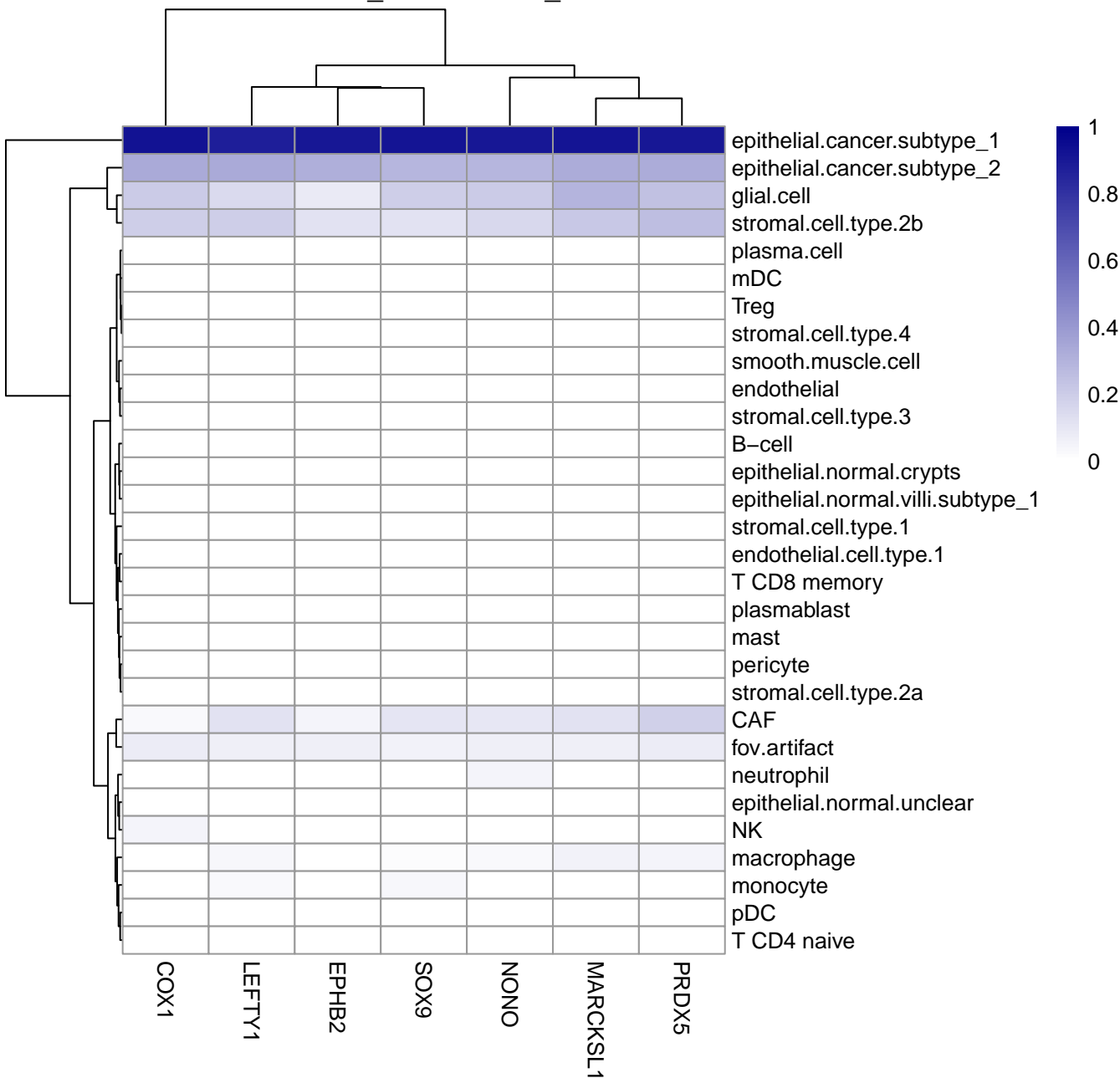
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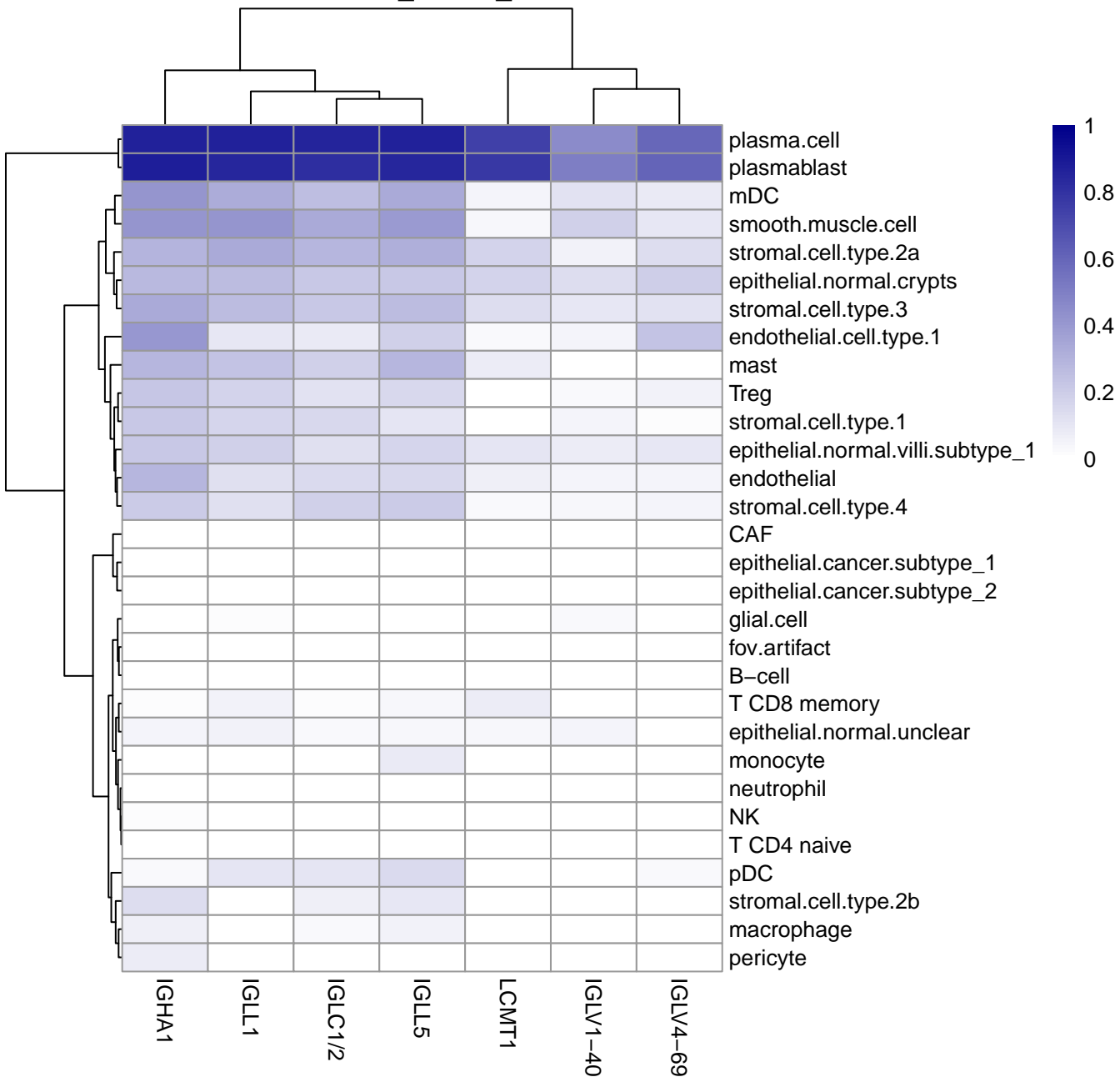
LOXL2_TIMP3_7



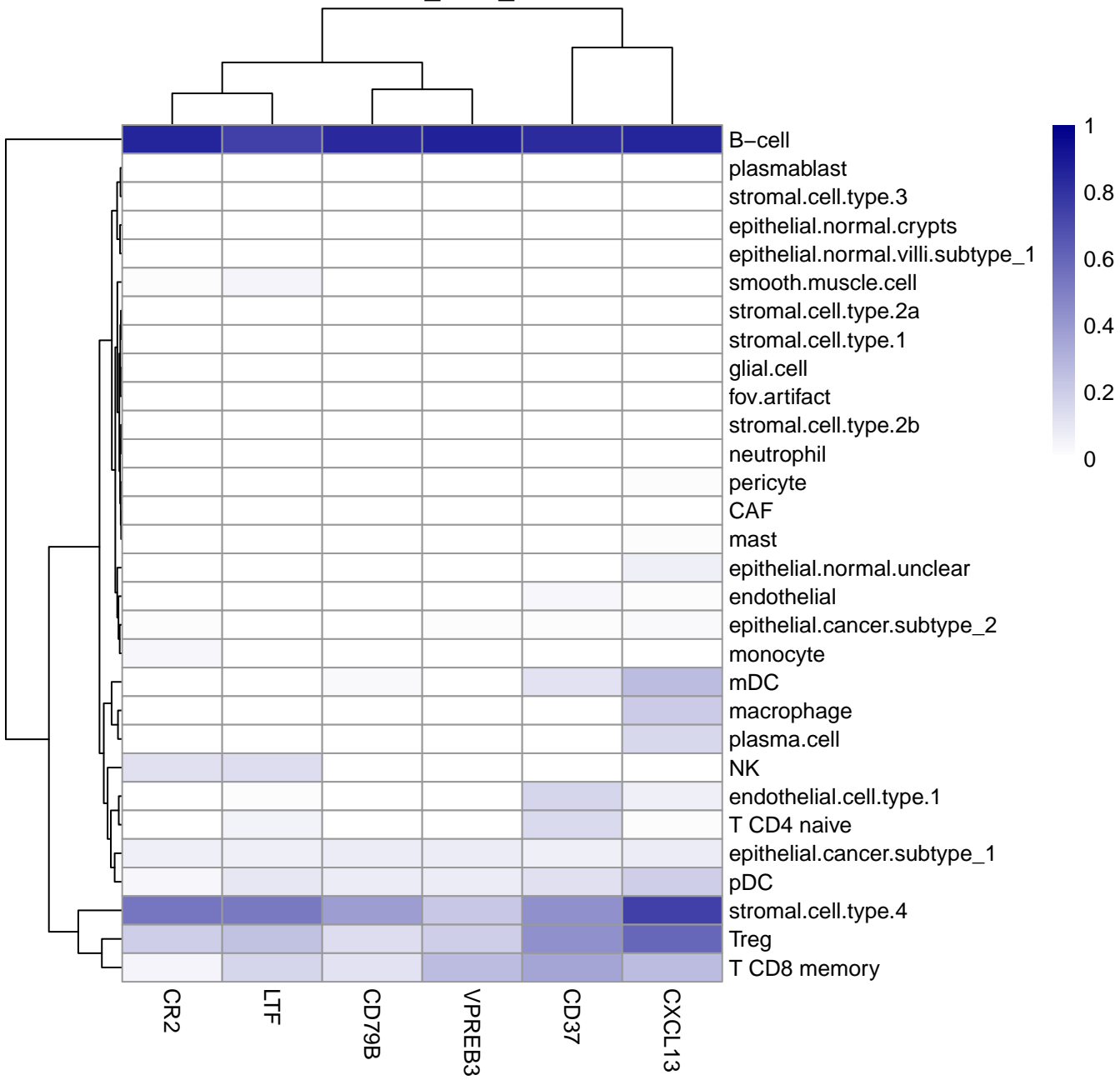
PRDX5 MARCKSL1 7



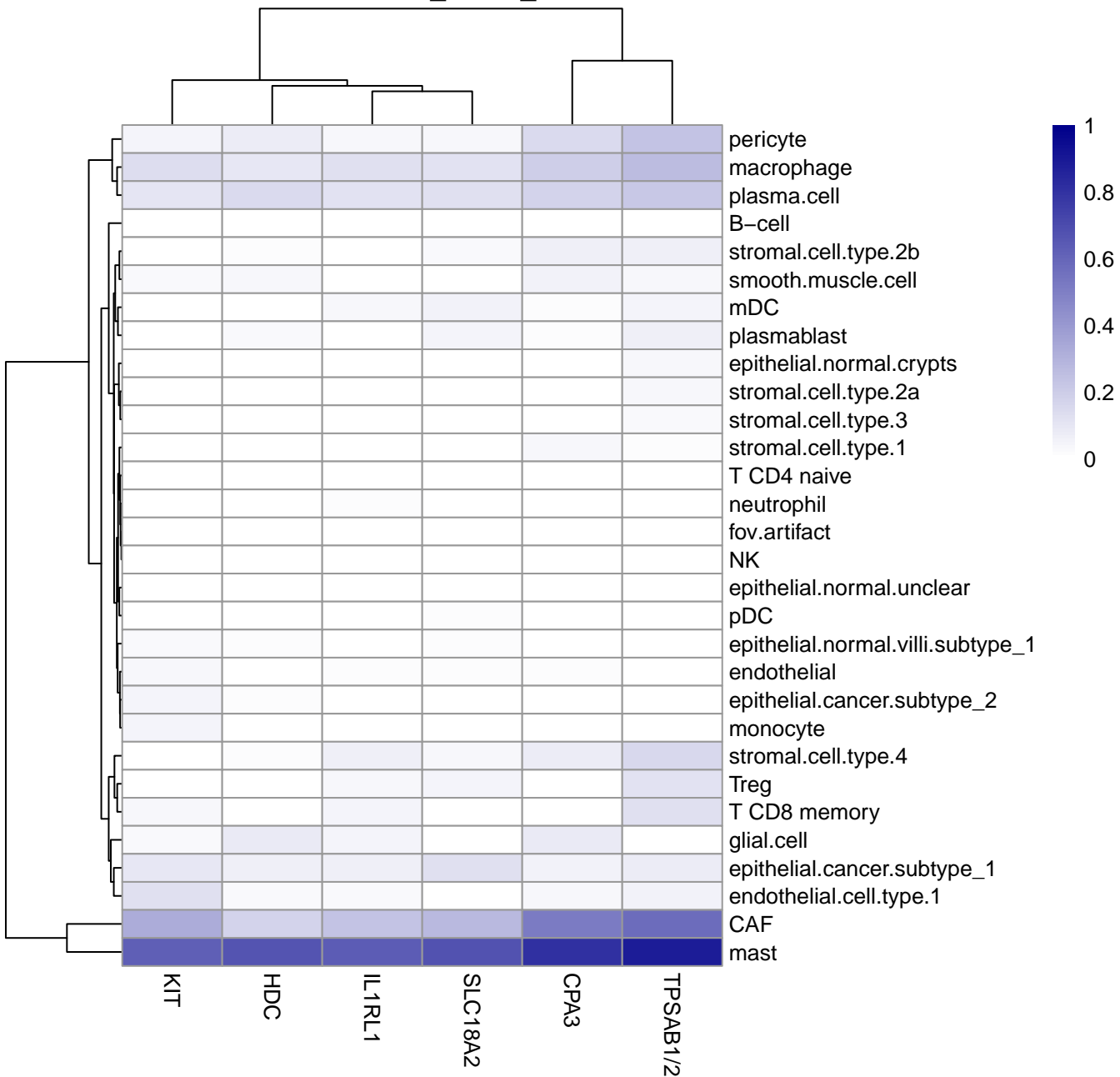
IGLL1_IGLL5_7



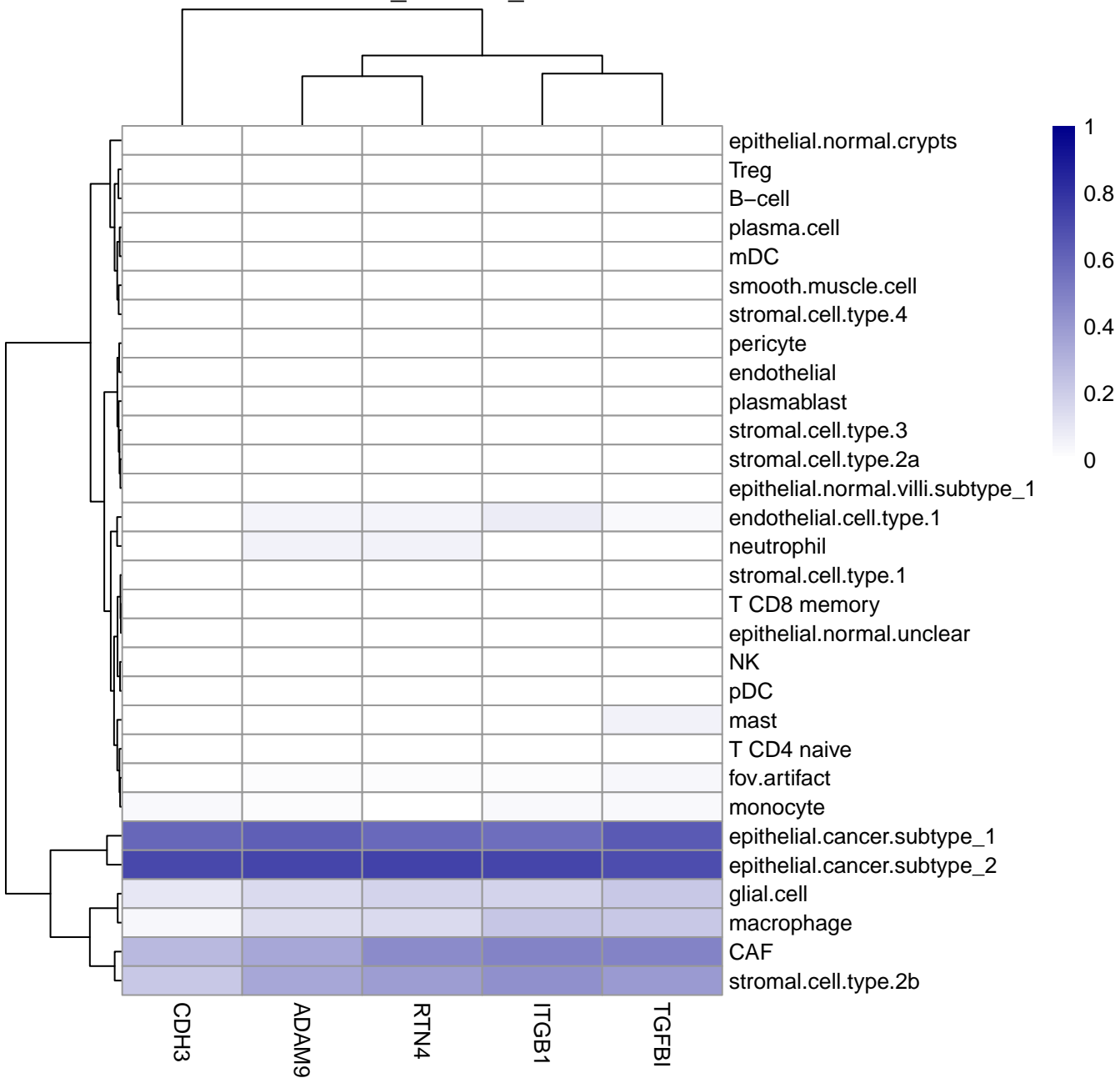
CXCL13_CR2_6



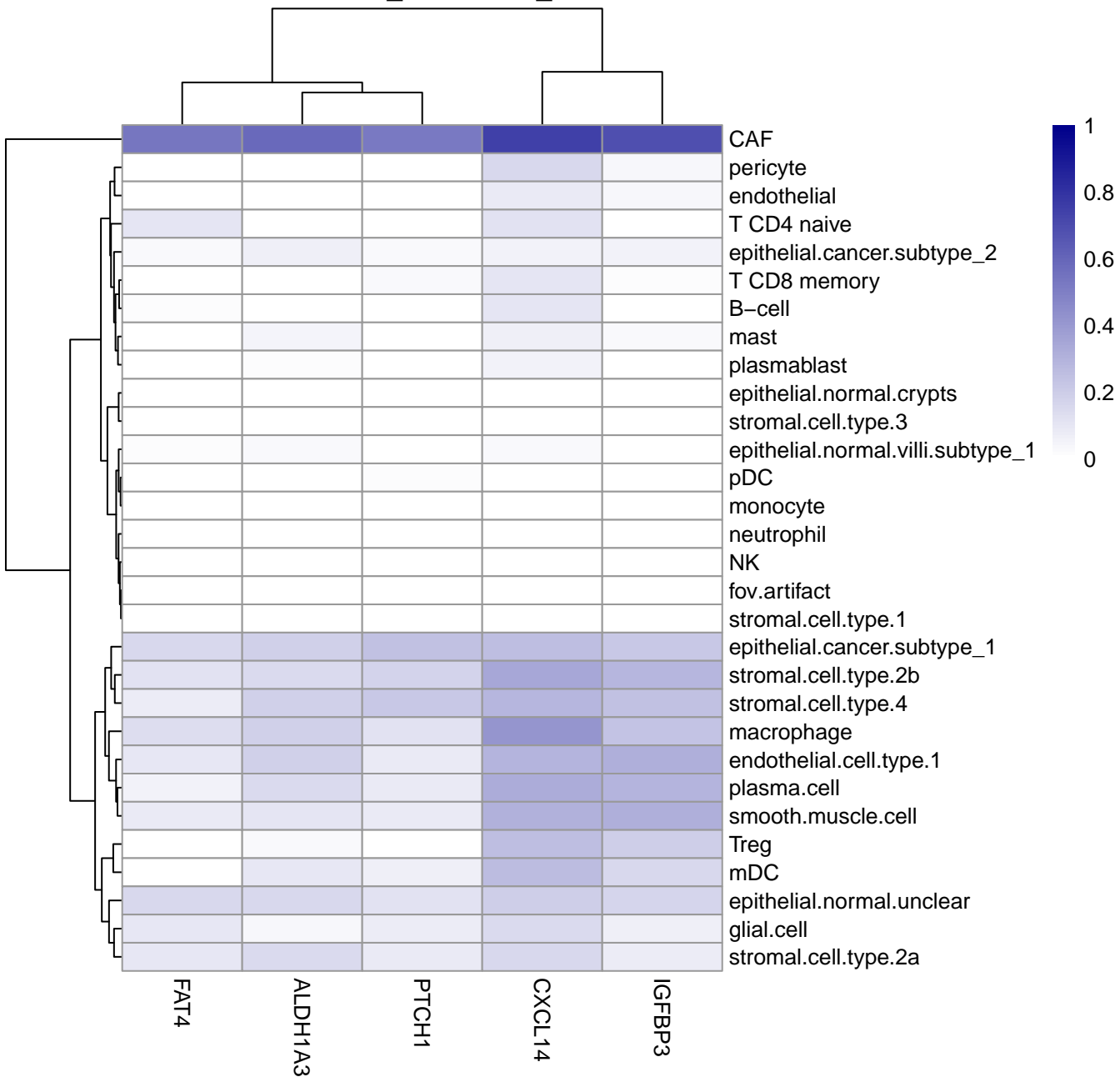
TPSAB1/2_CPA3_6



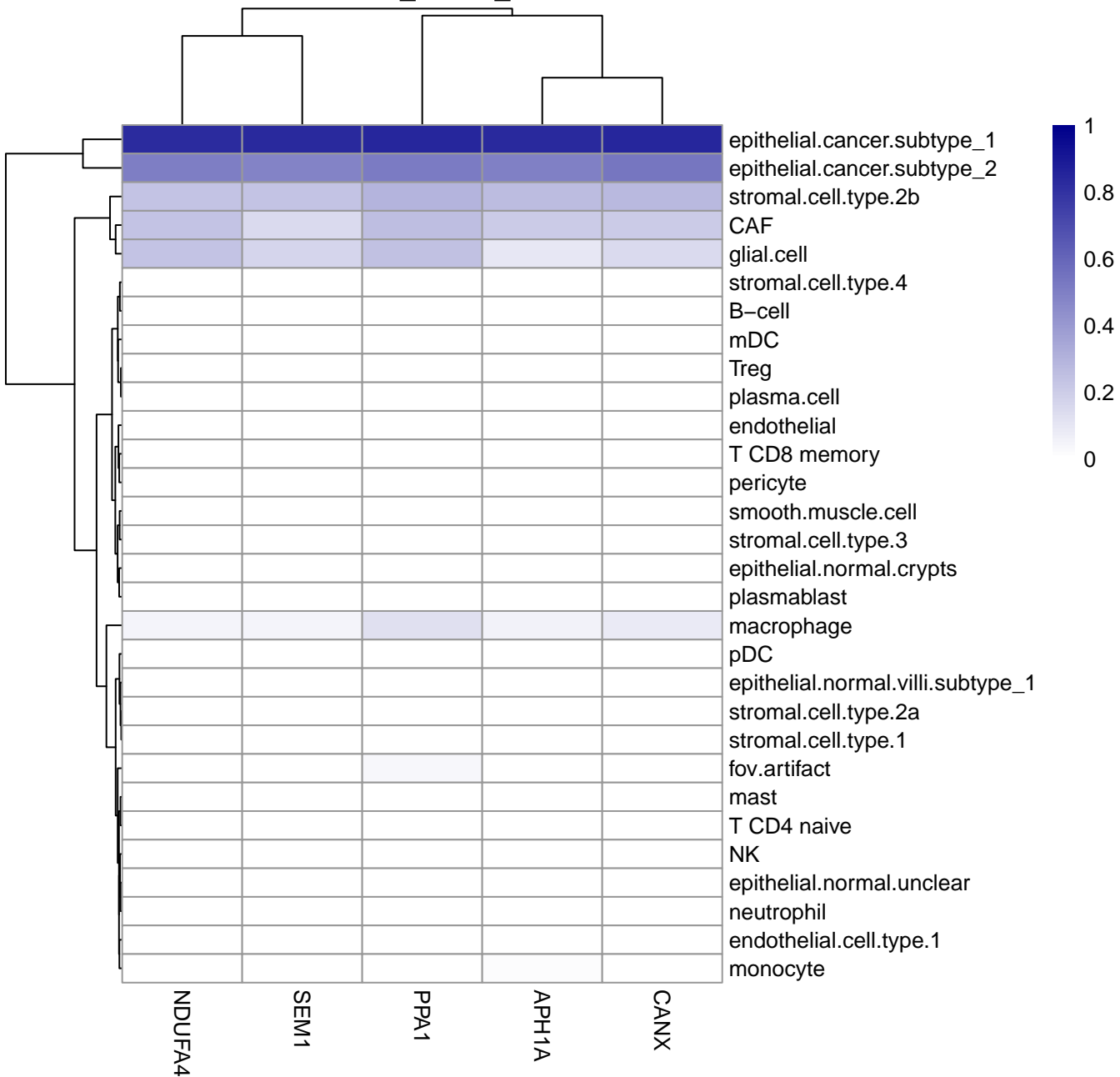
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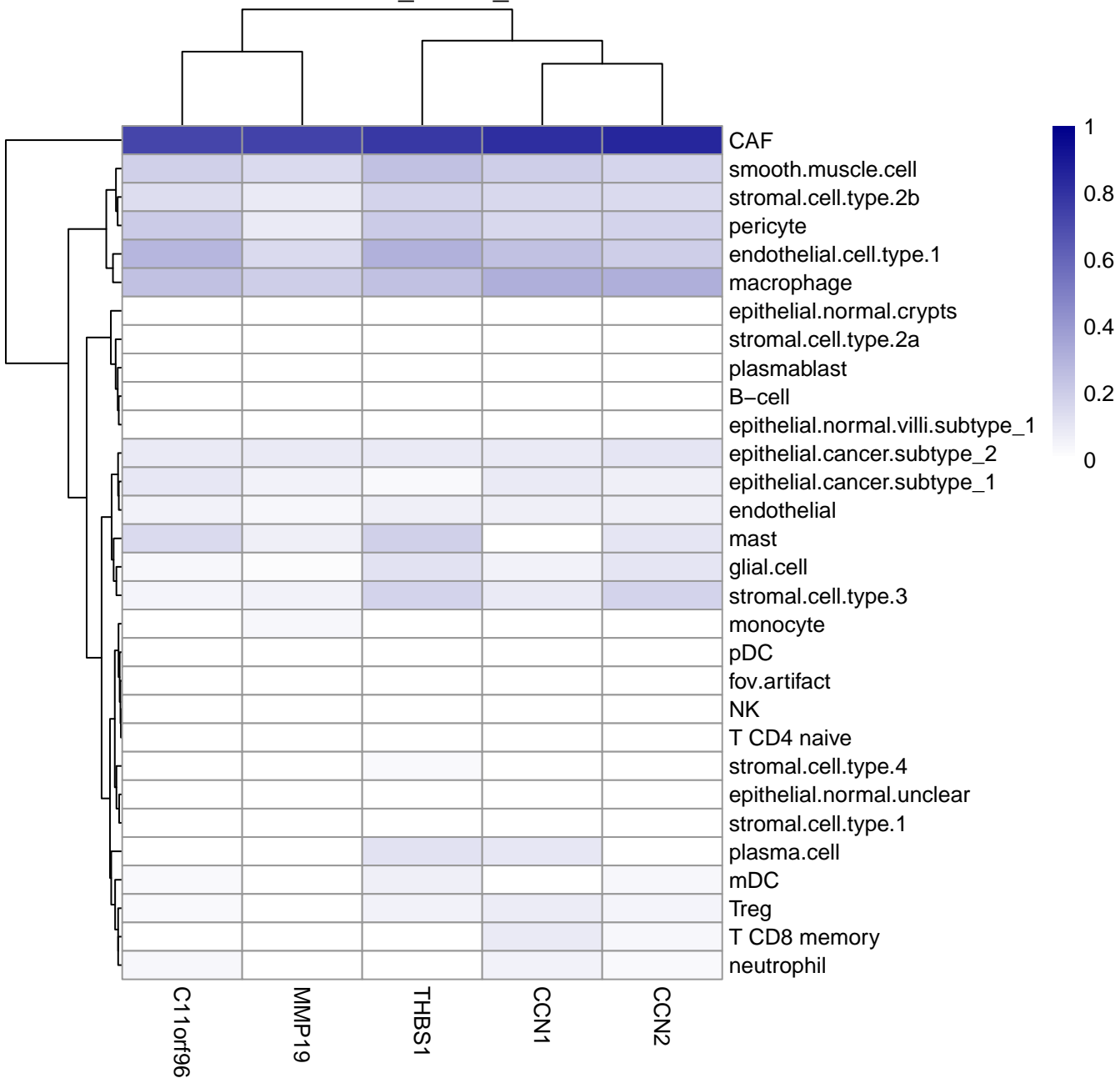
CXCL14_ALDH1A3_5



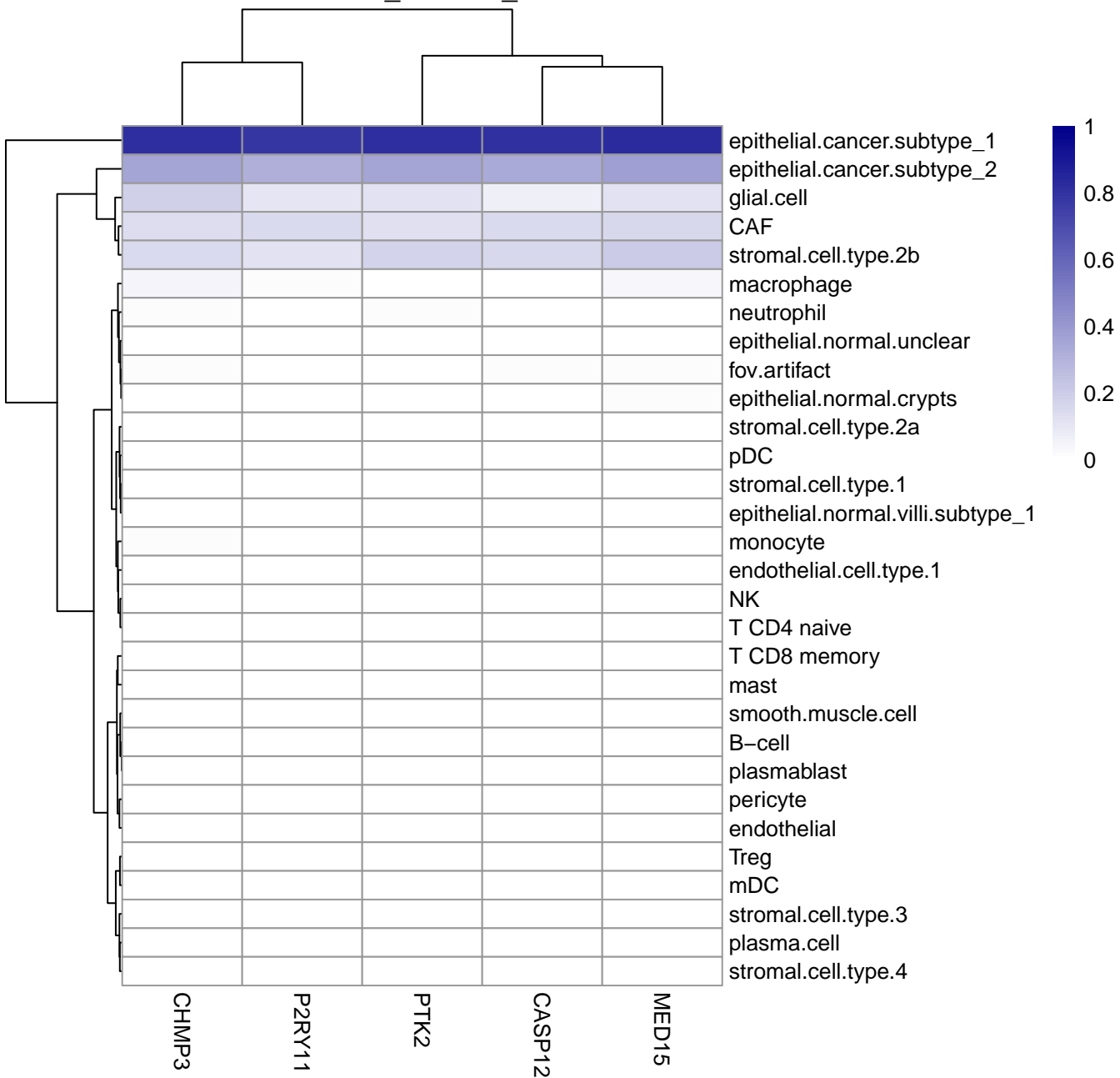
CANX_SEM1_5



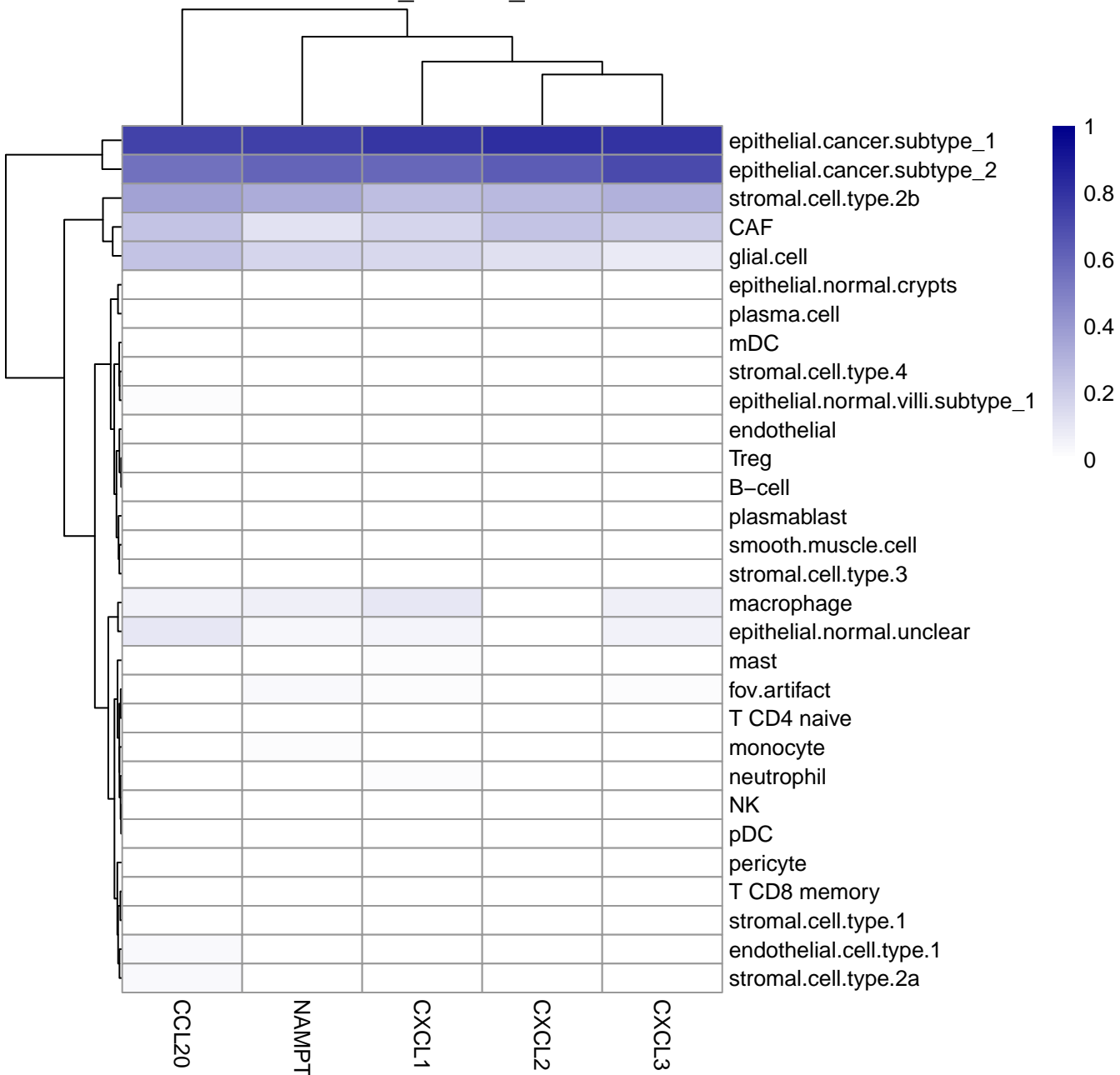
CCN1_CCN2_5



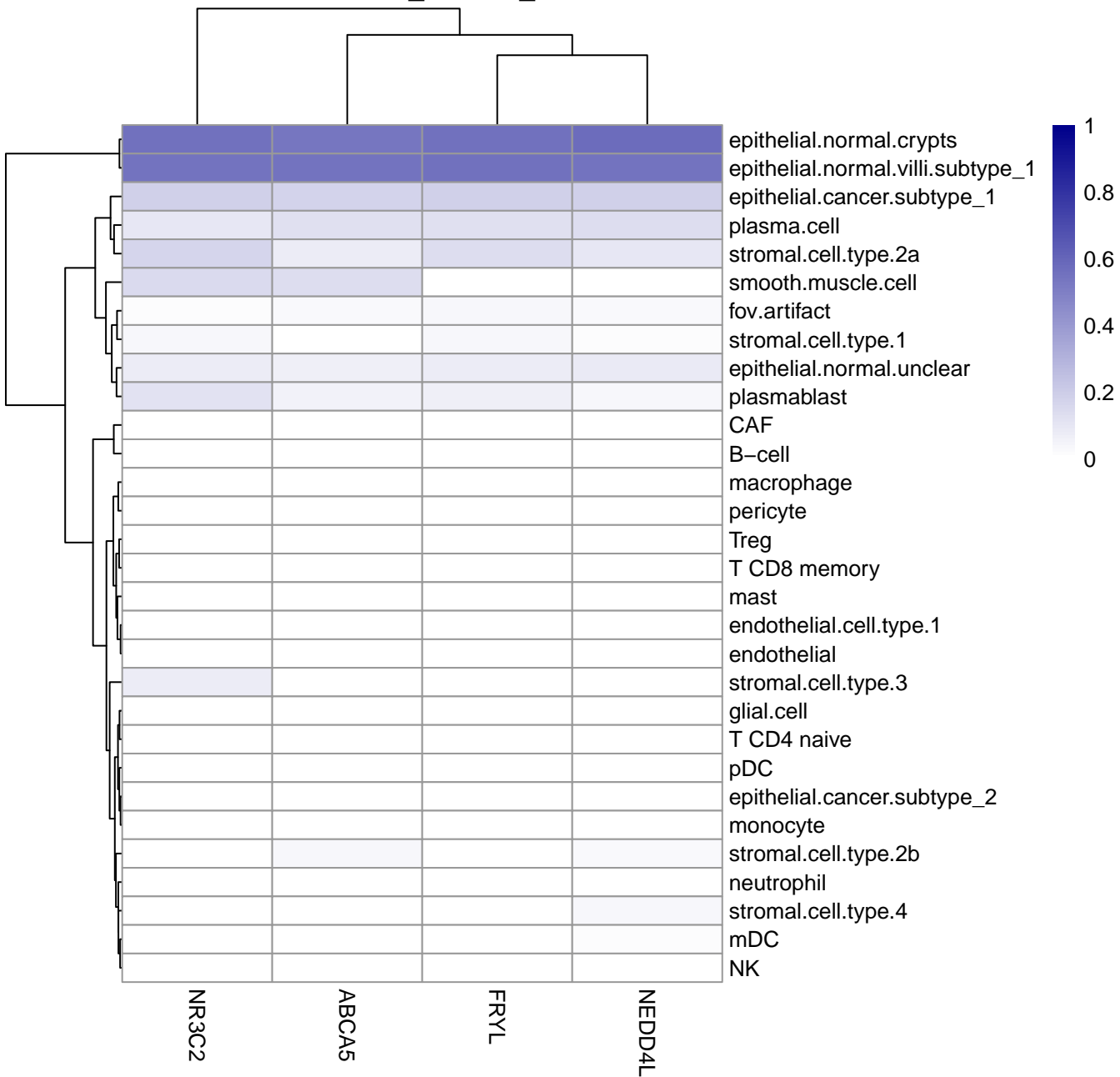
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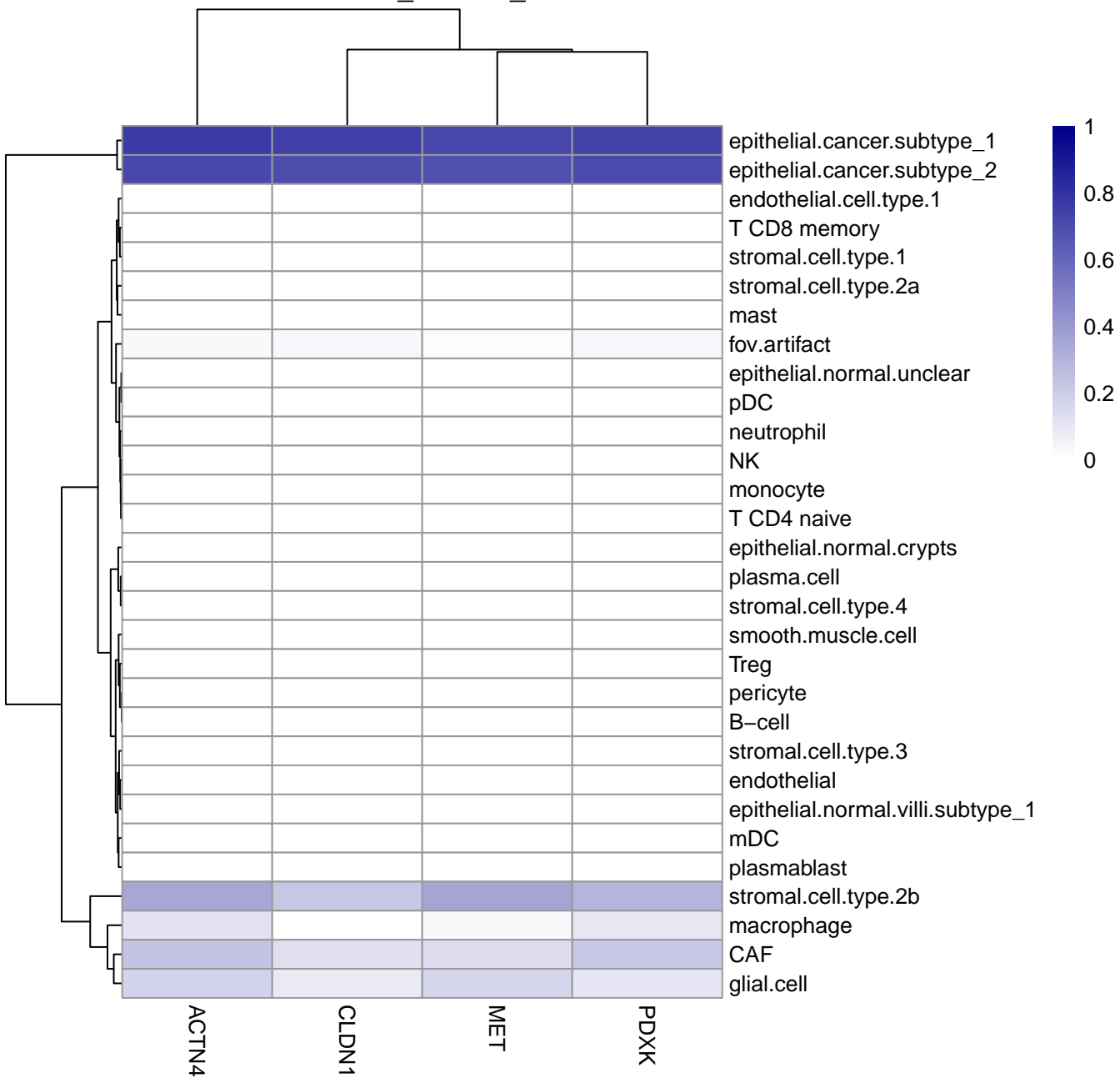
CXCL3_CXCL2_5



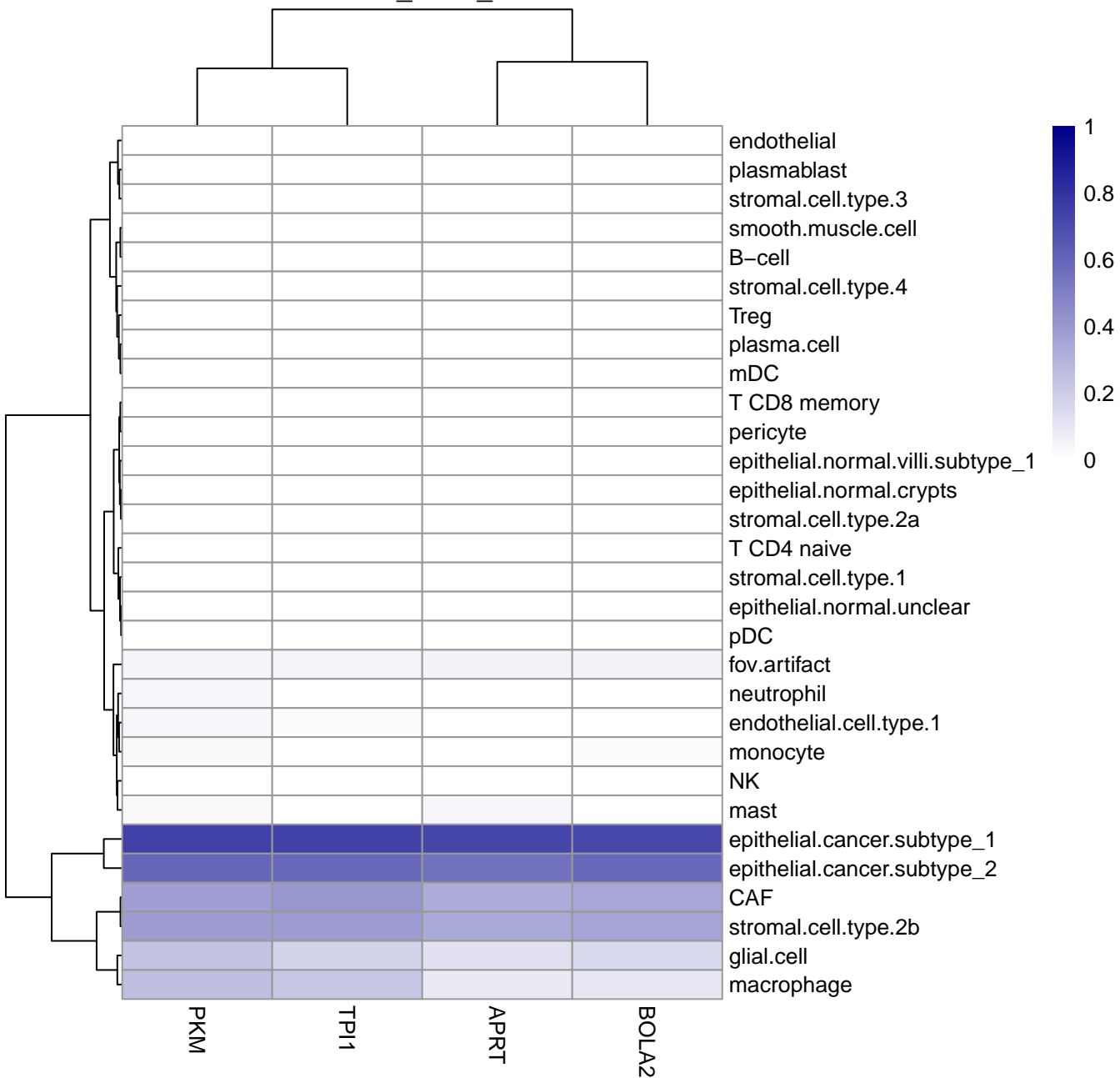
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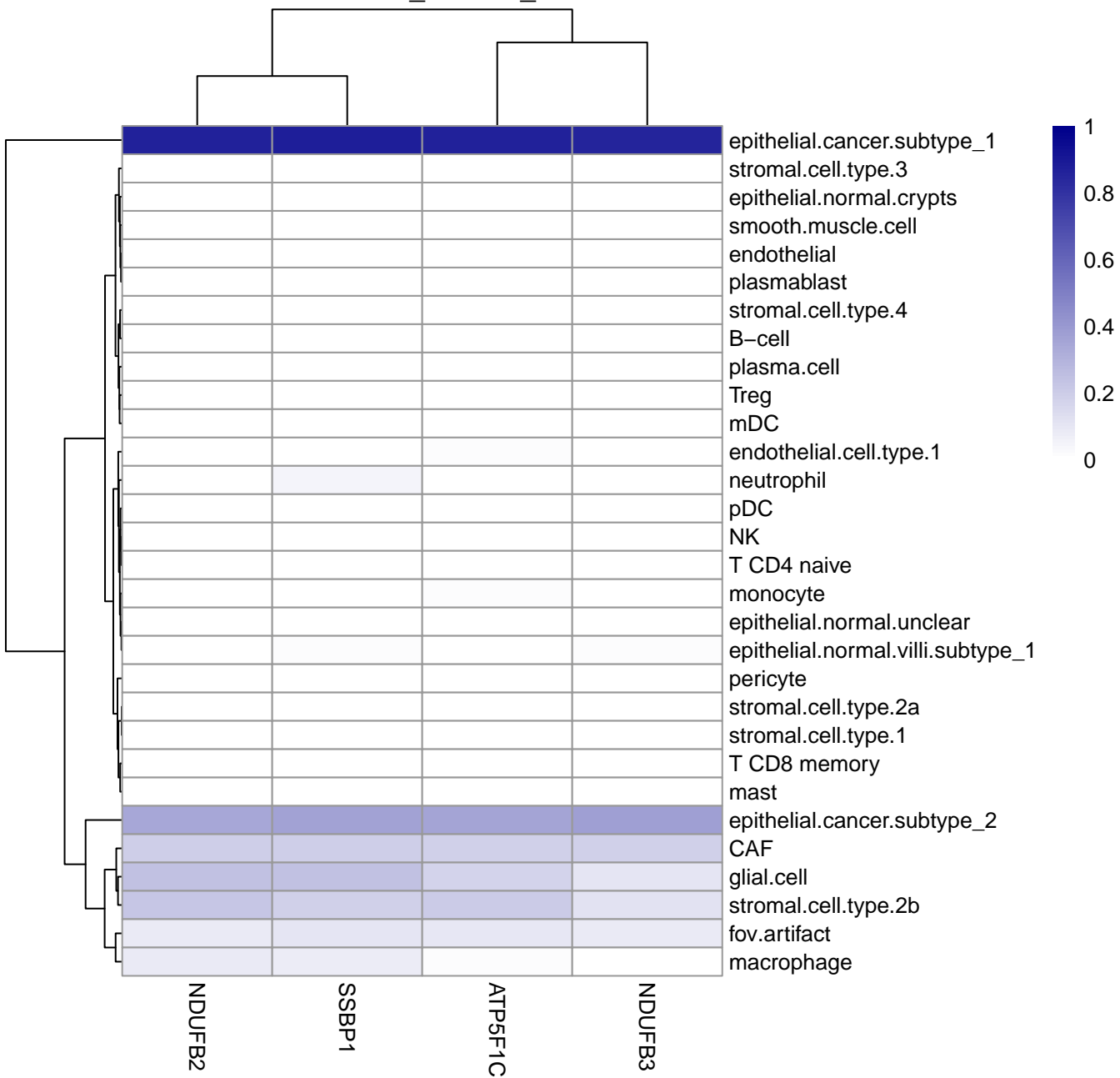
ACTN4_CLDN1_4



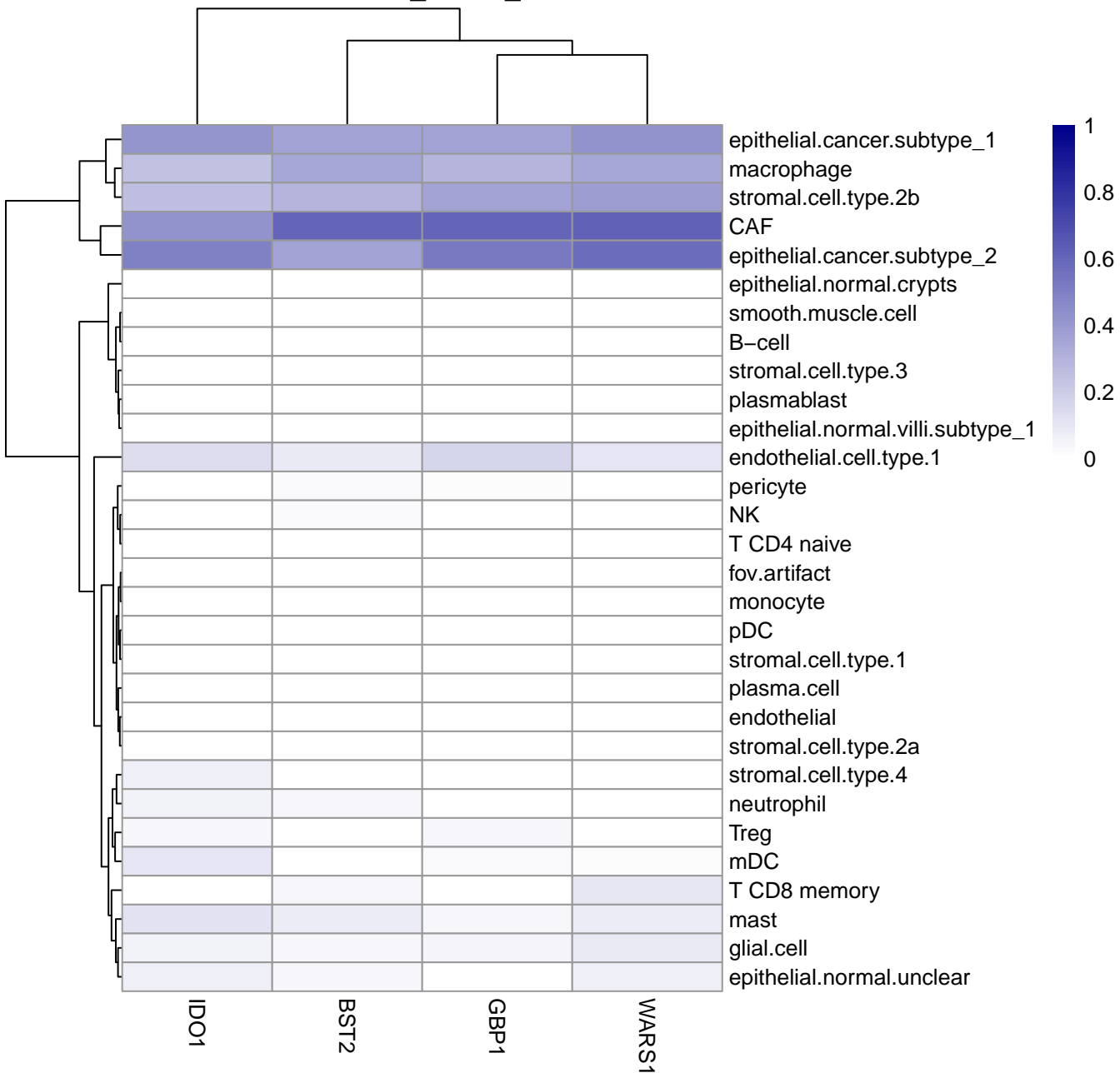
TPI1_PKM_4



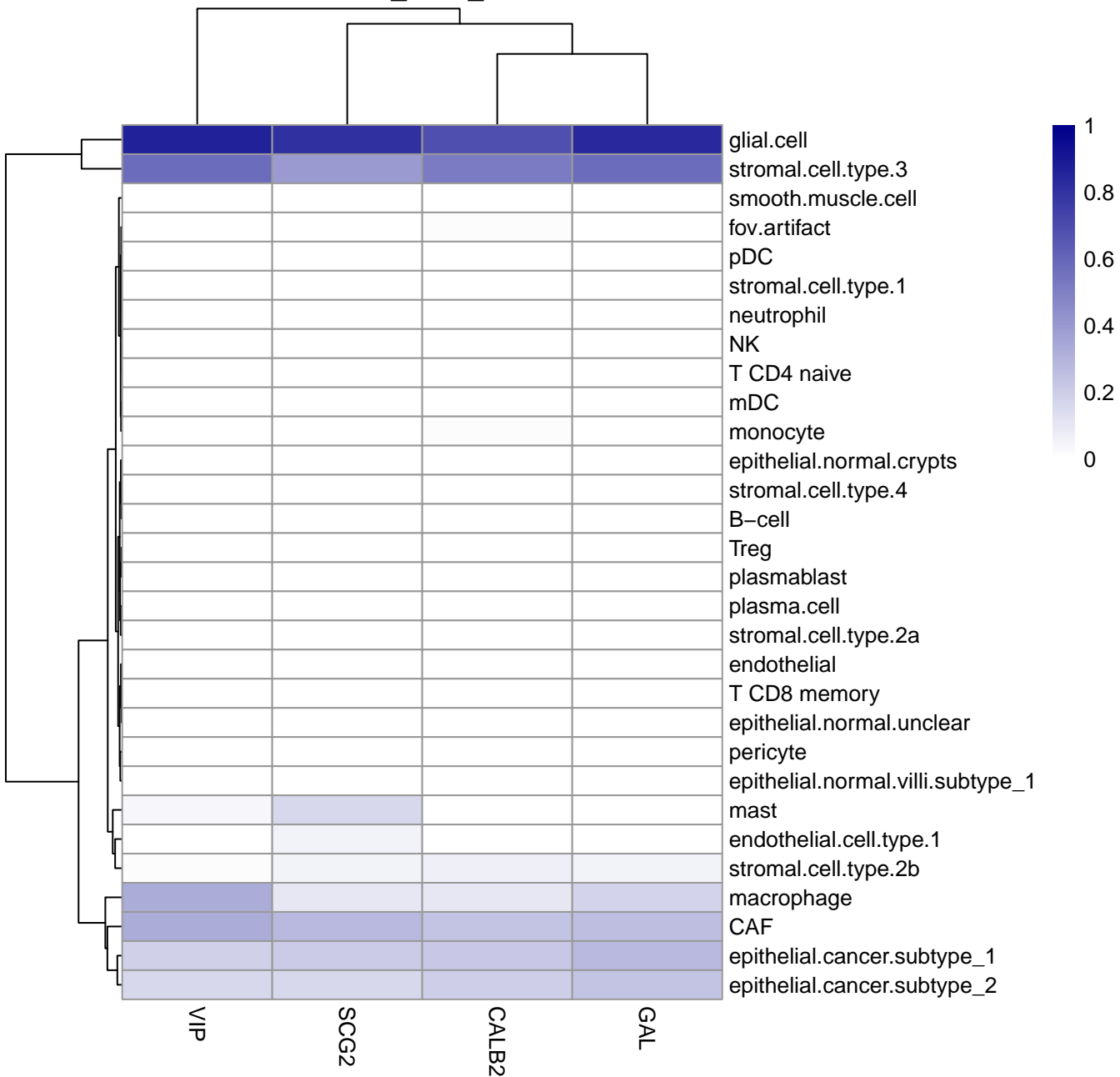
NDUFB3 SSBP1 4



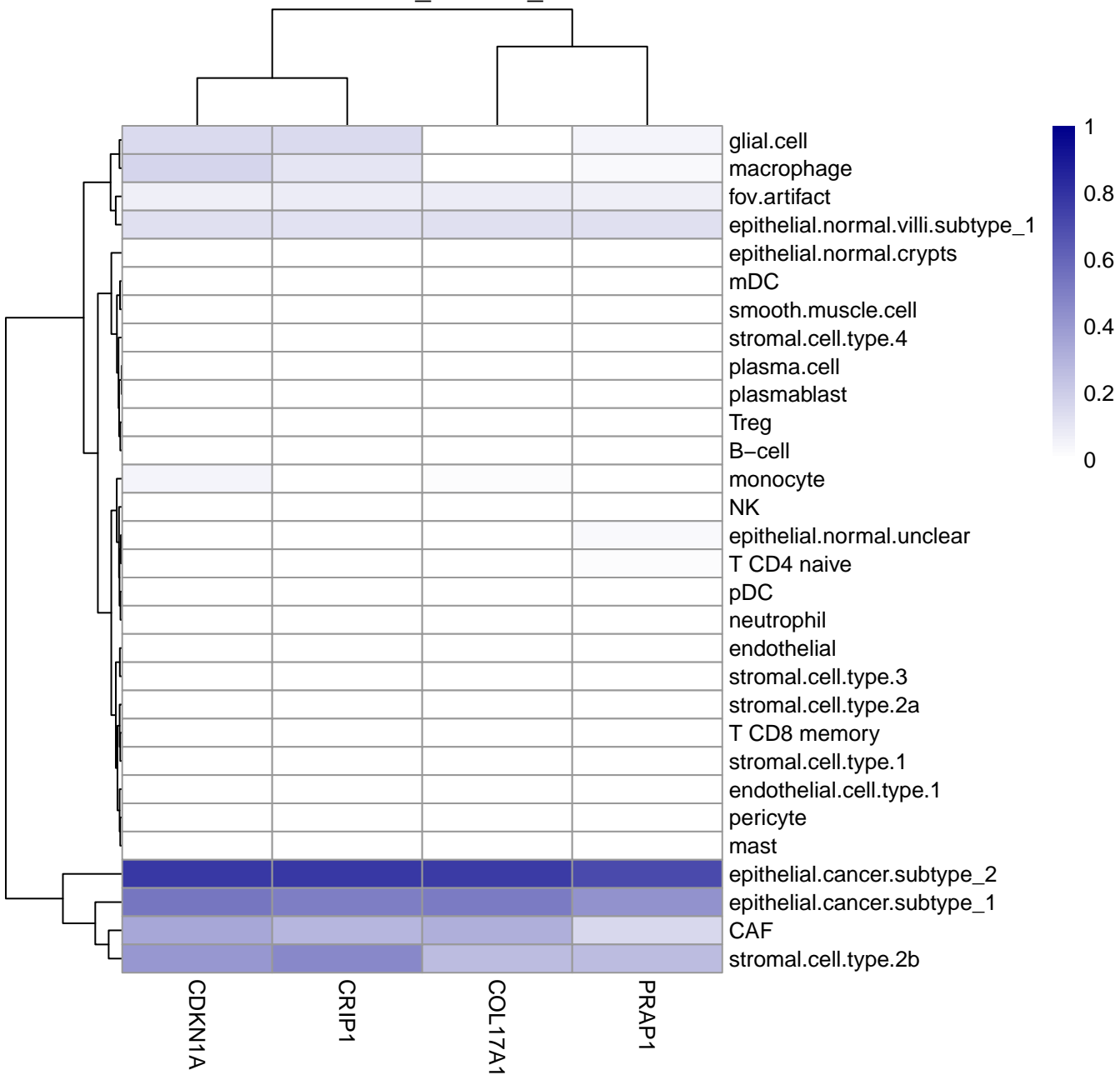
WARSI1_GBP1_4



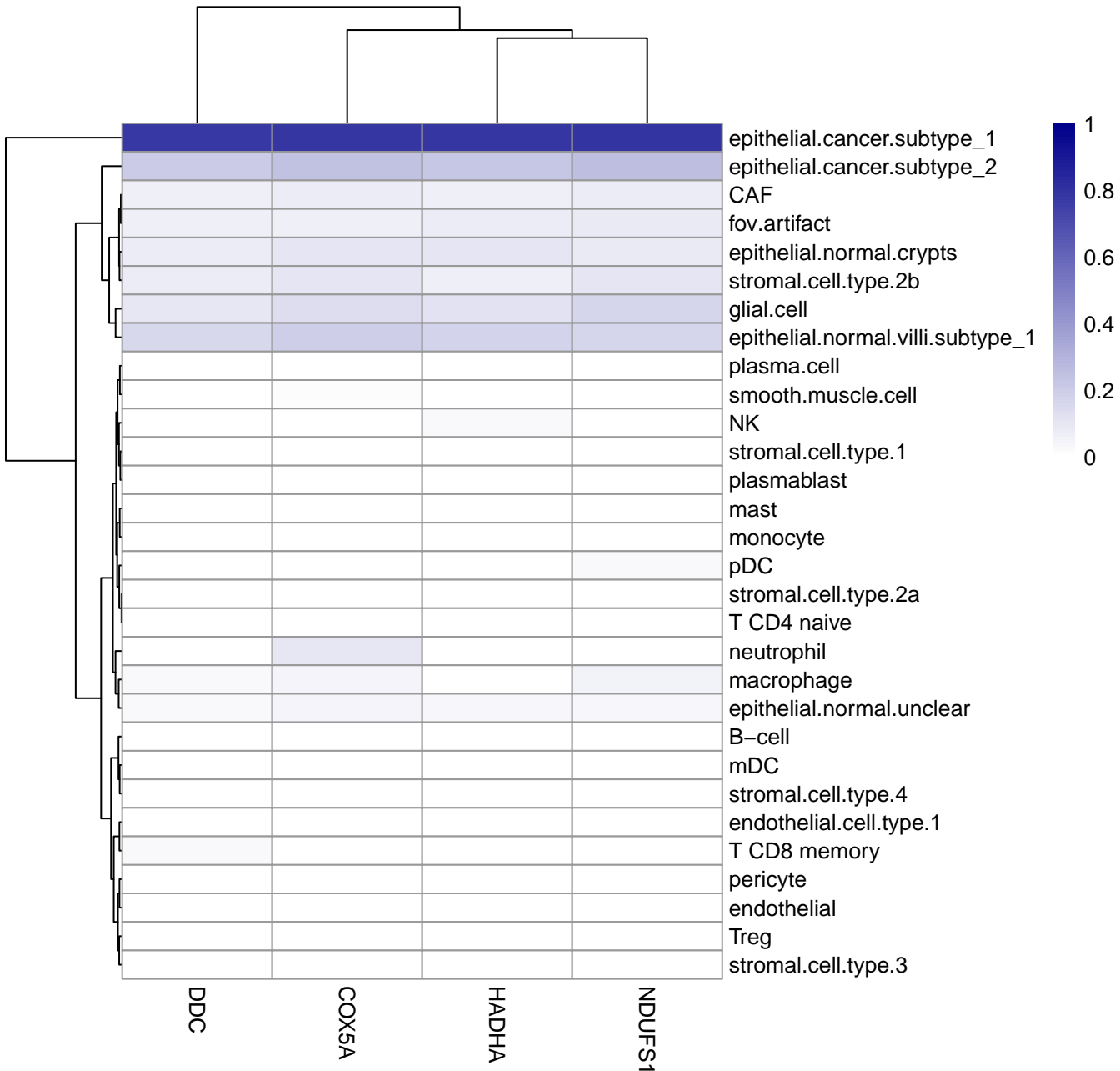
VIP_GAL_4



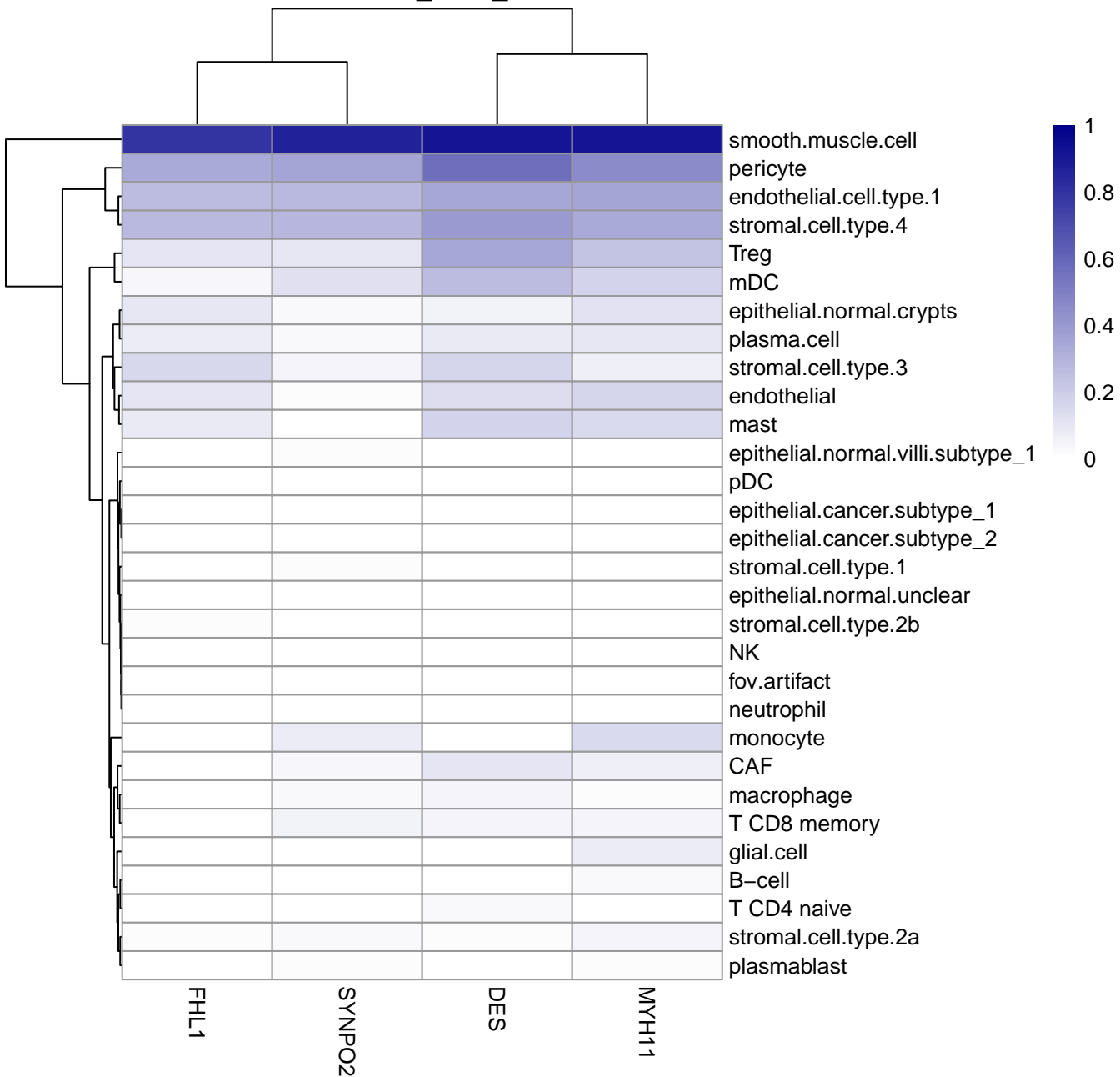
COL17A1_PRAP1_4



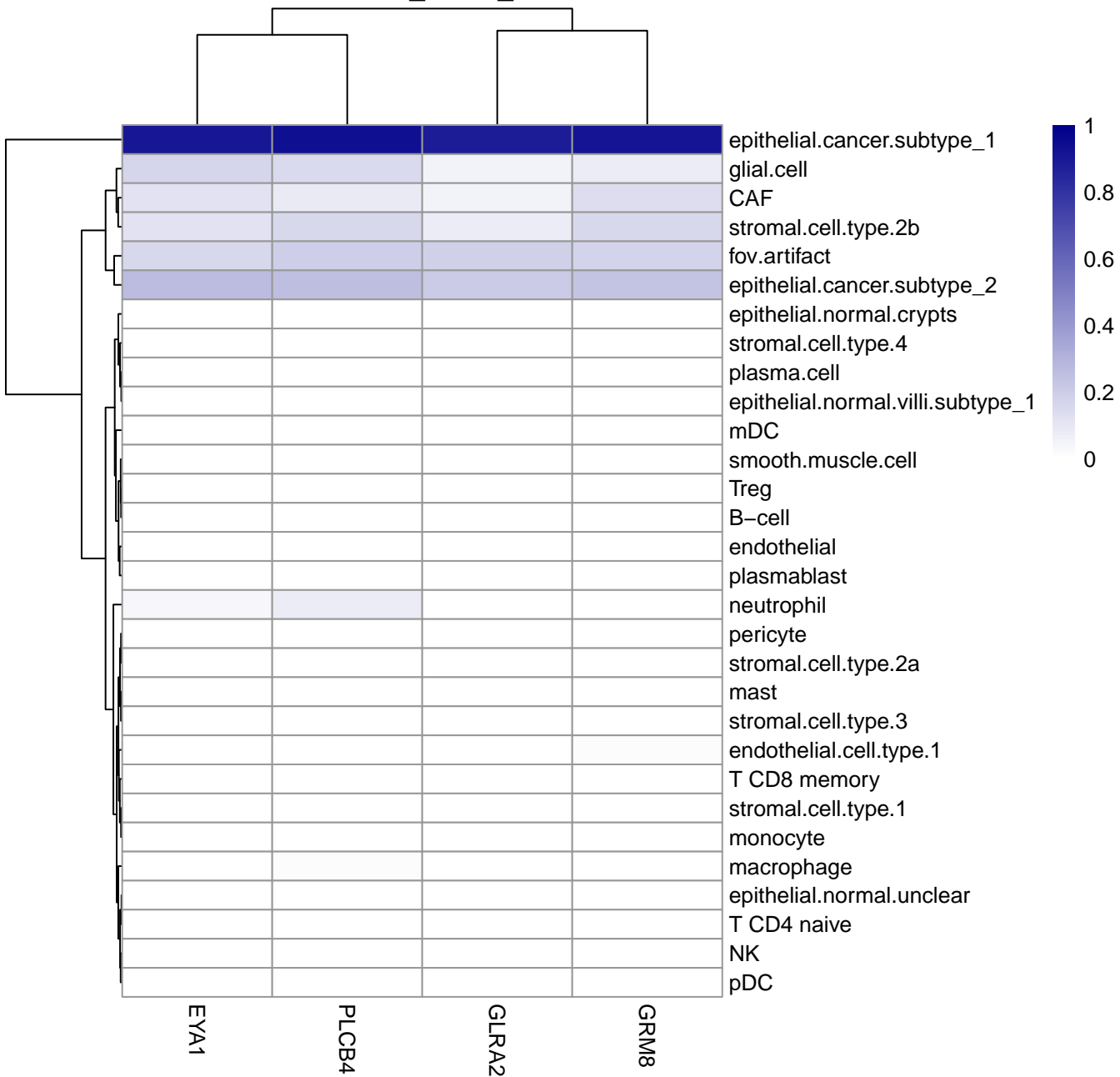
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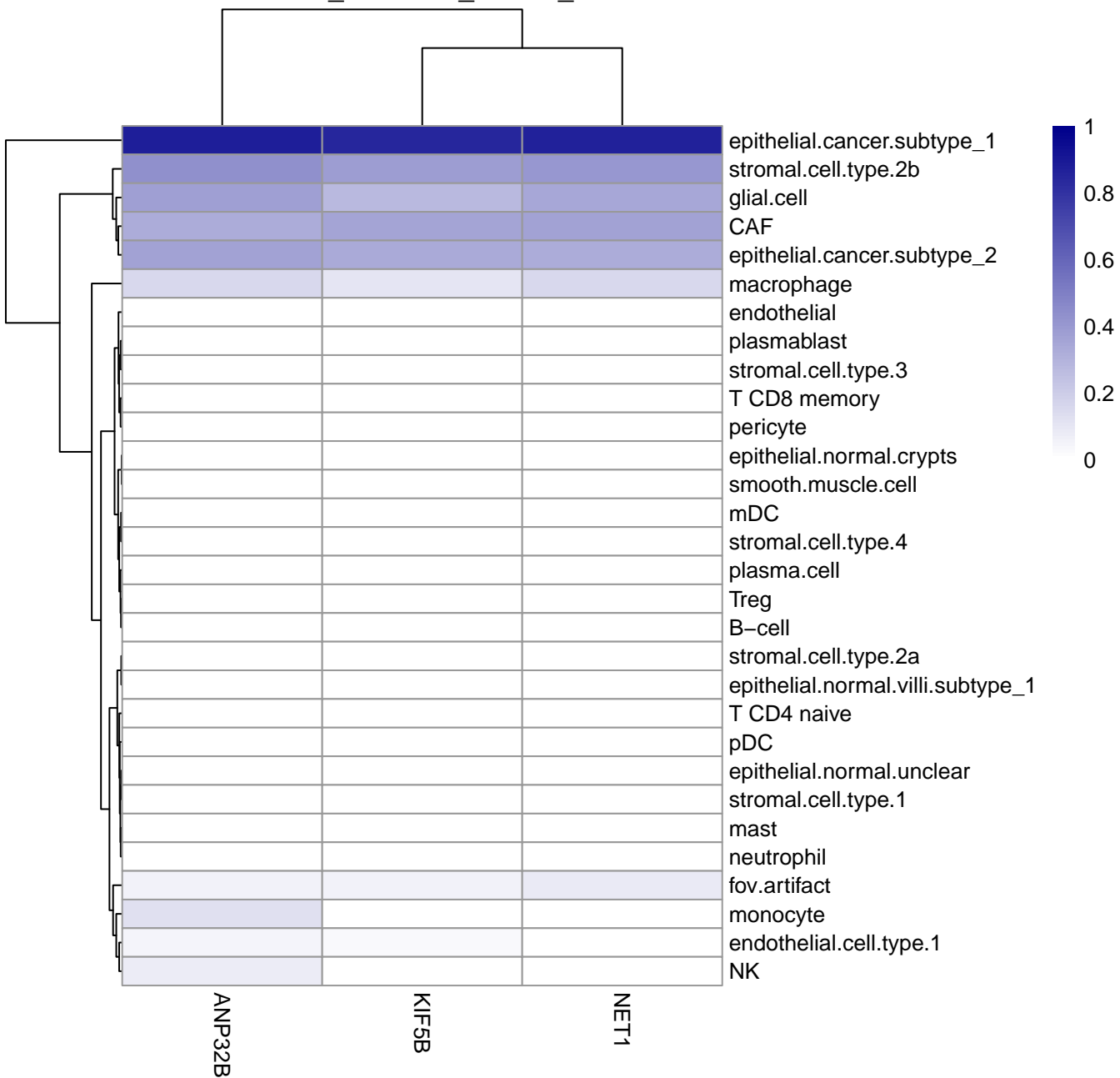
MYH11_DES_4



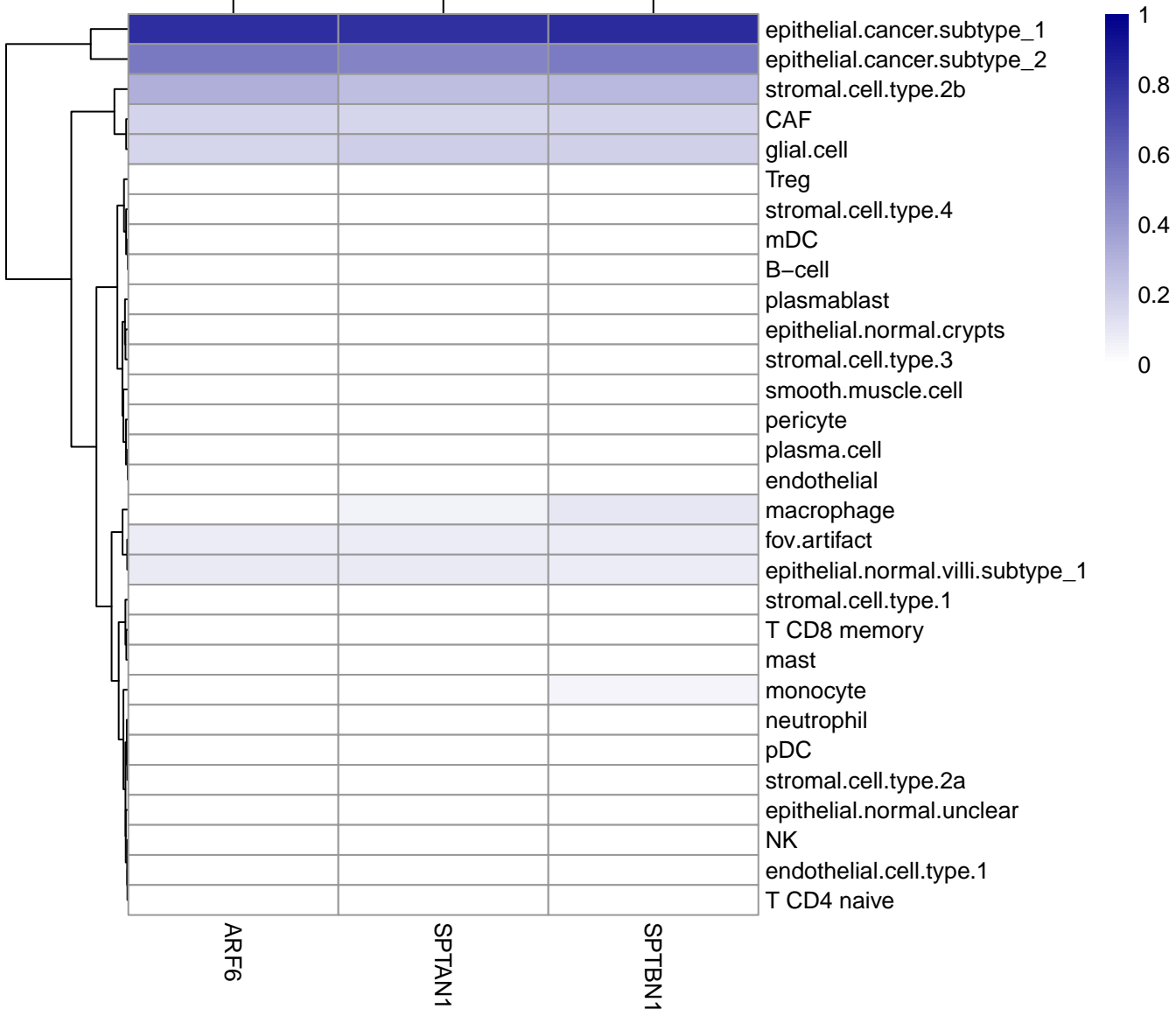
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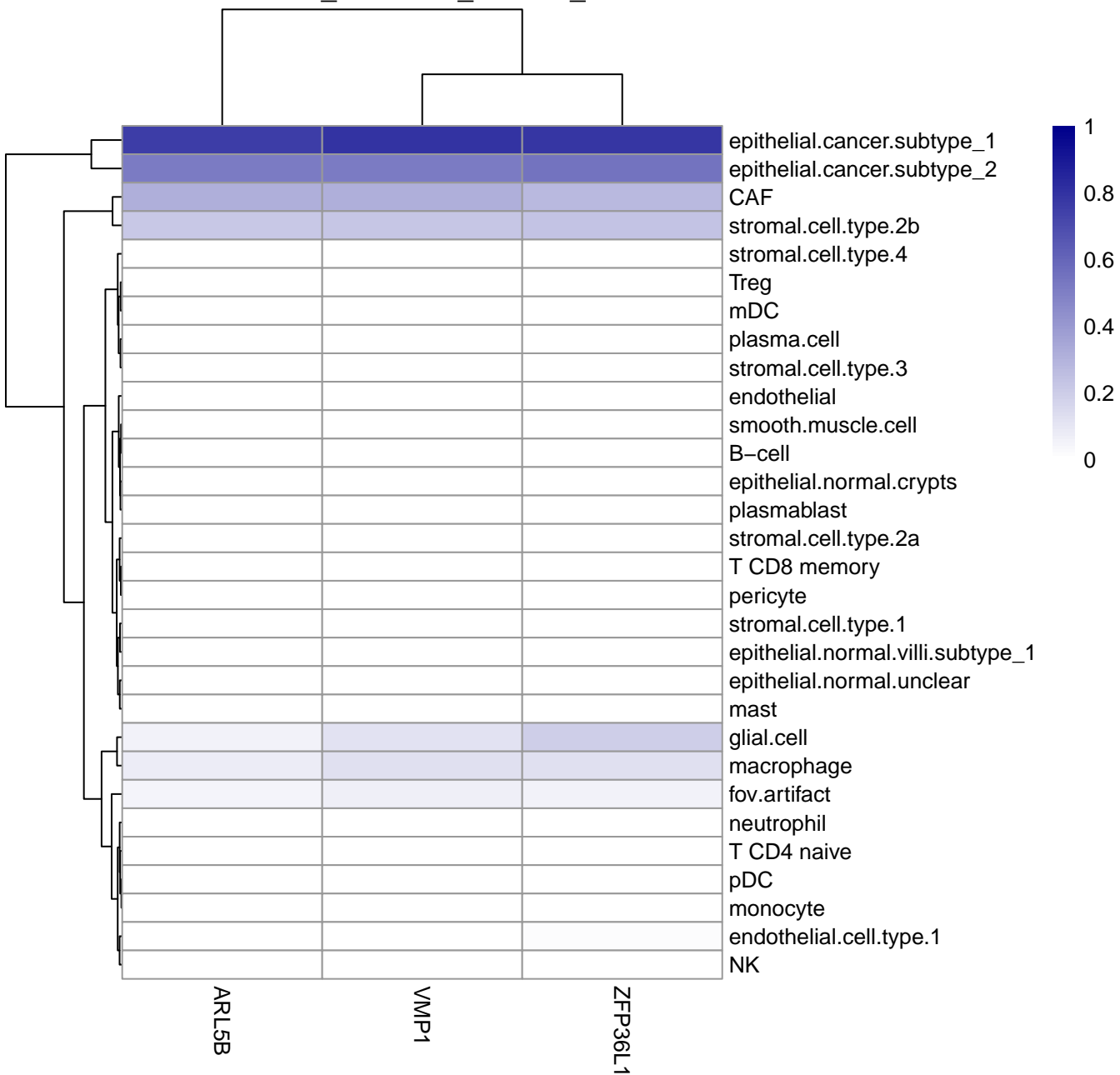
NET1_ANP32B_KIF5B_3



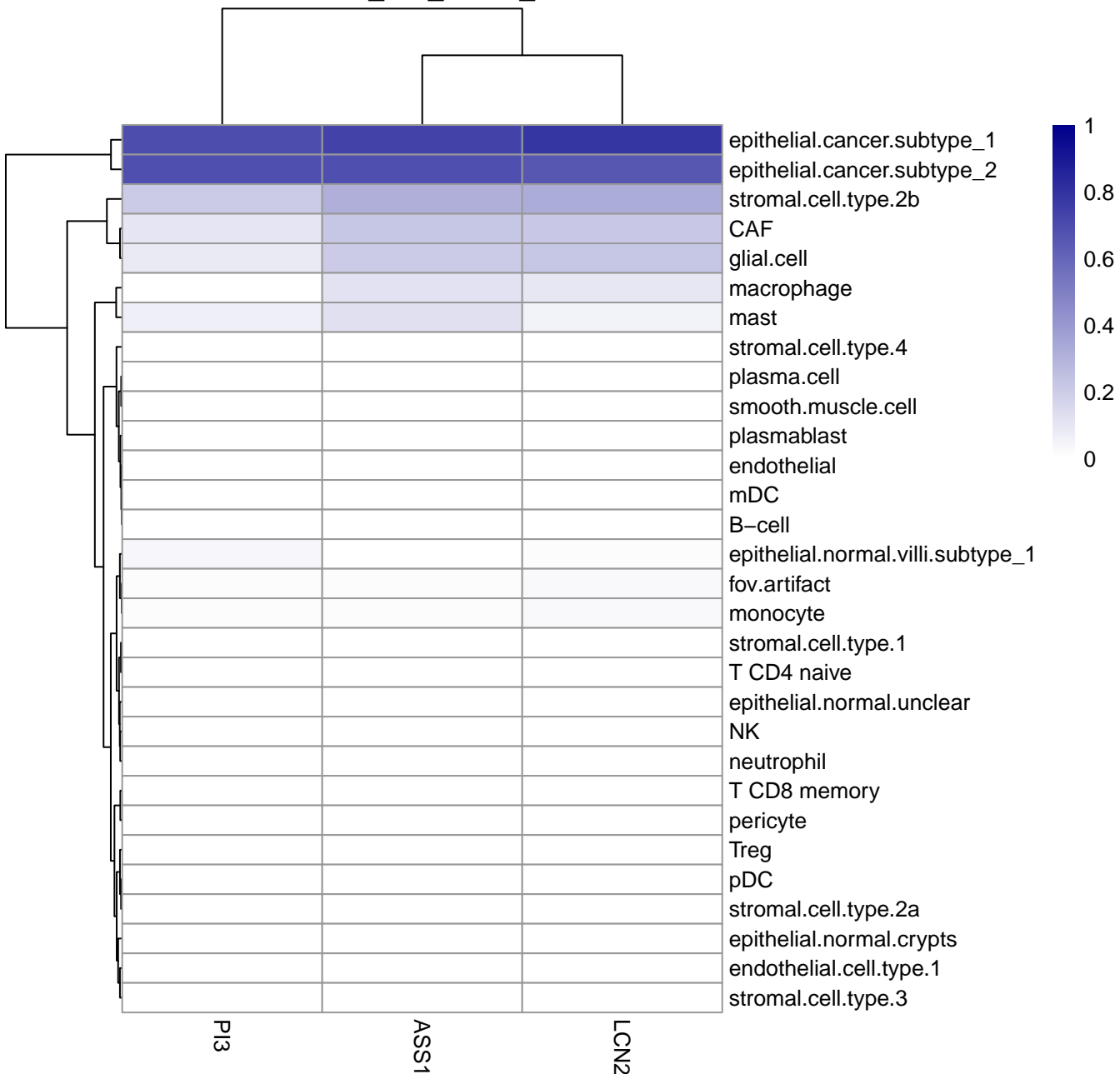
SPTBN1_SPTAN1_ARF6_3



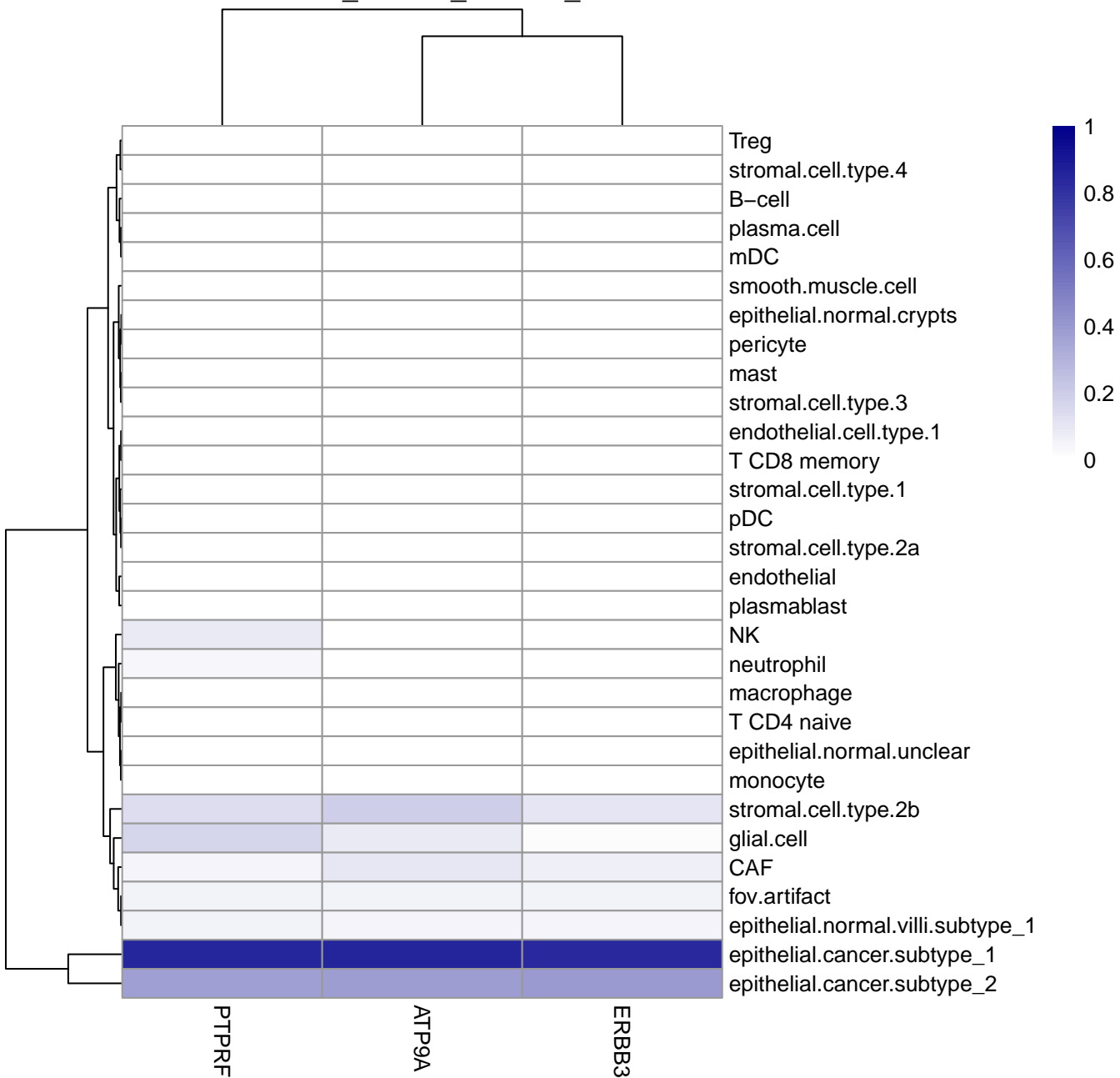
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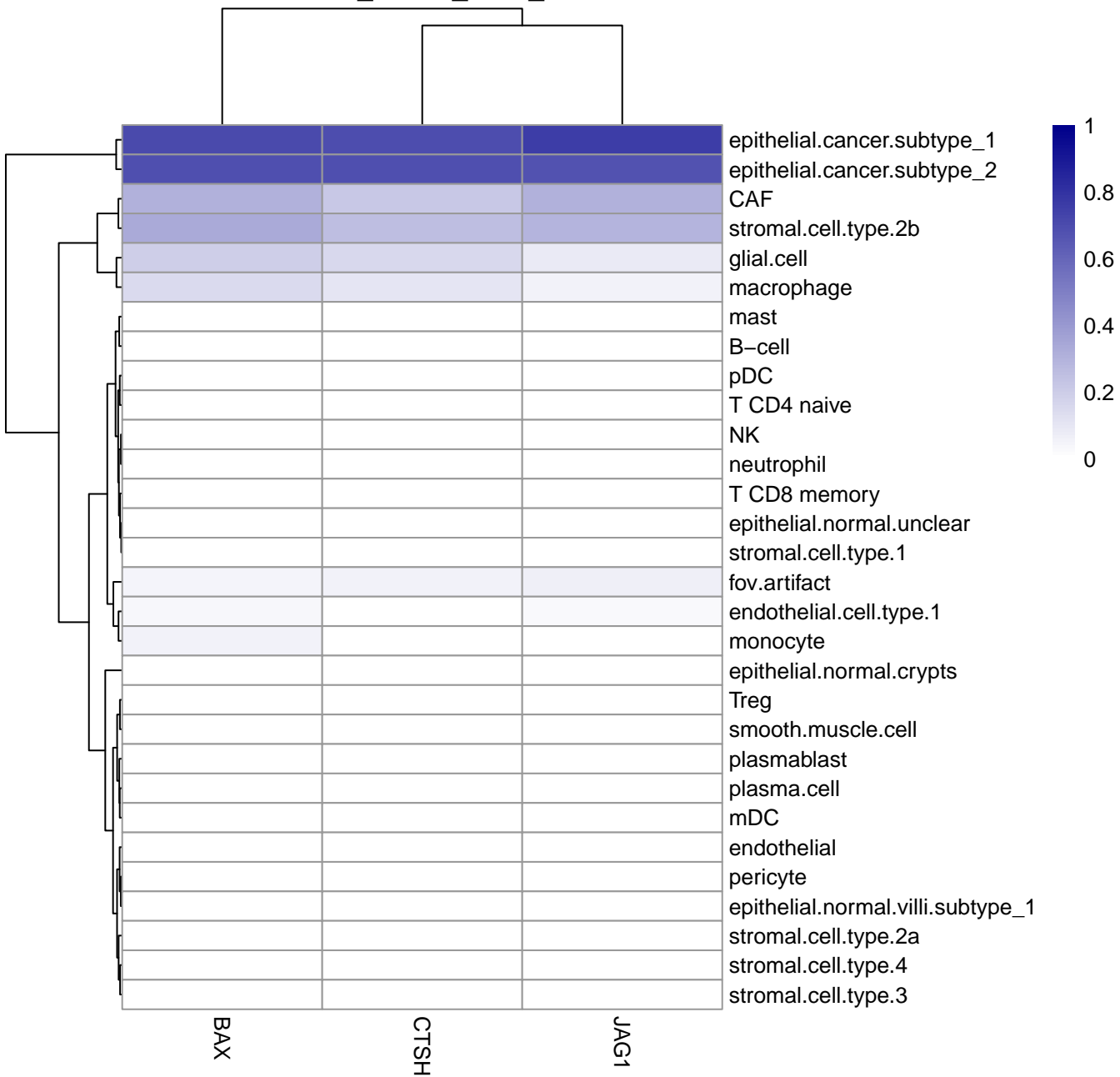
ASS1_PI3_LCN2_3



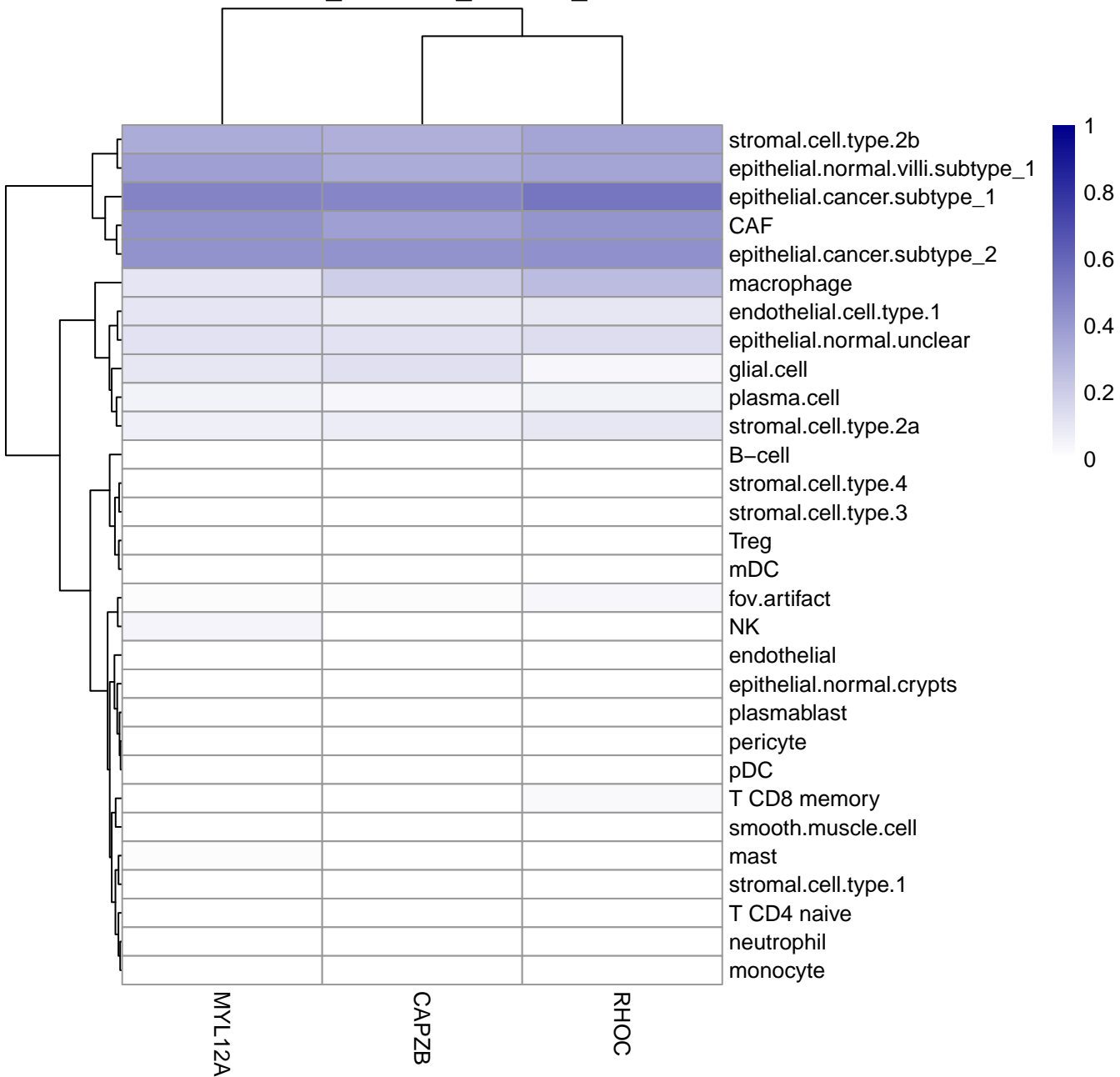
PTPRF_ATP9A_ERBB3_3



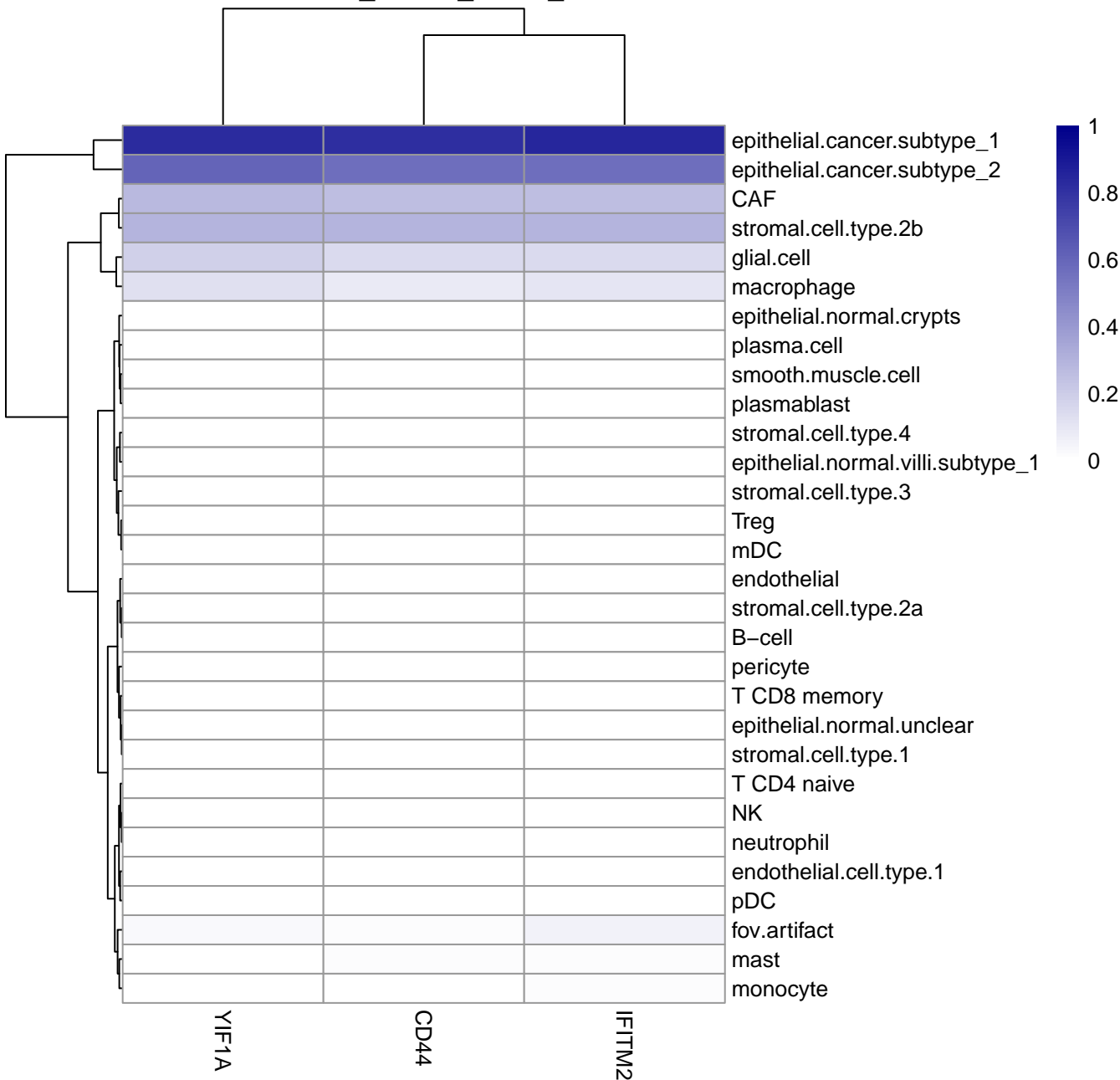
JAG1_CTSH_BAX_3



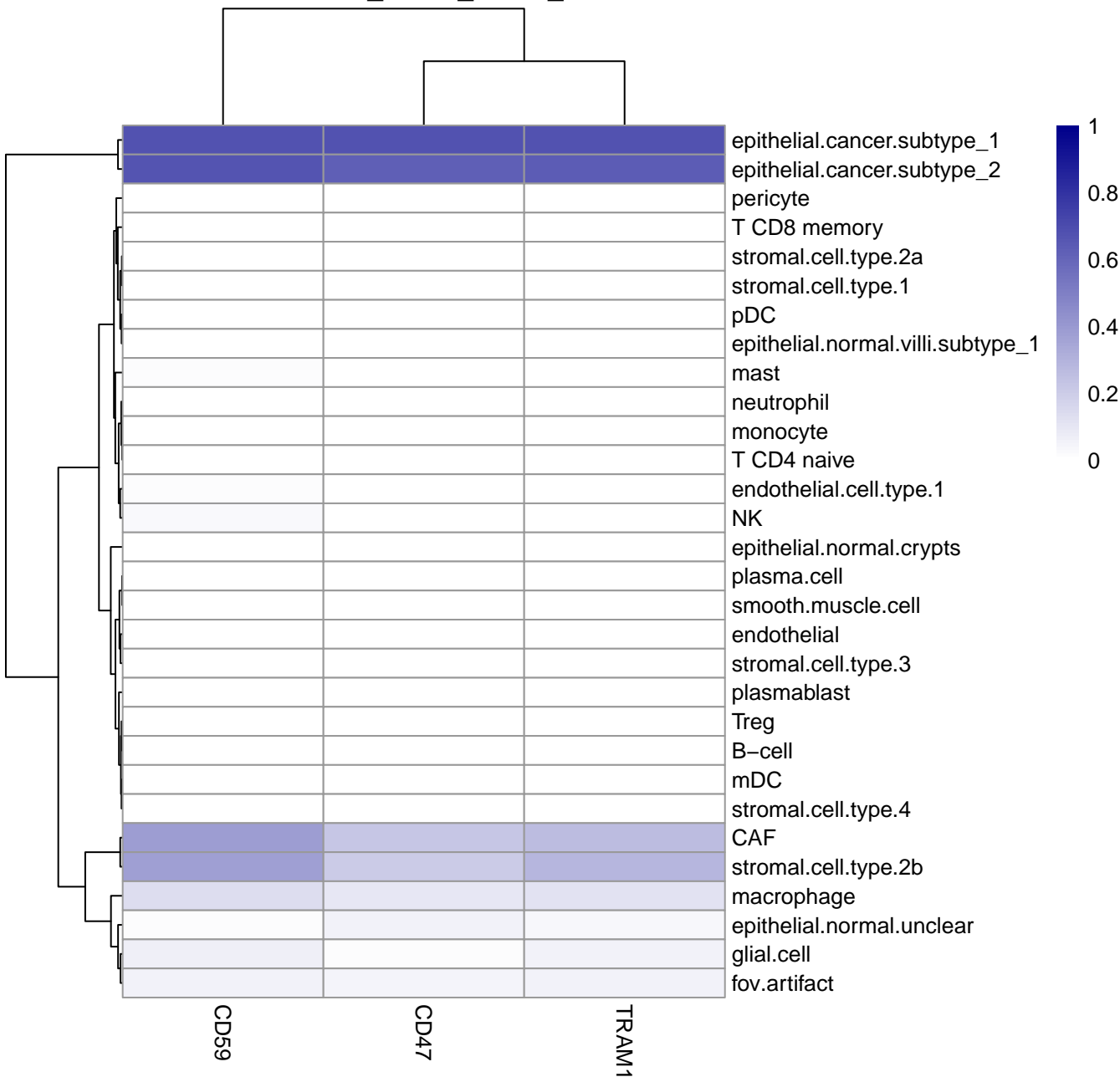
RHOC_MYL12A_CAPZB_3



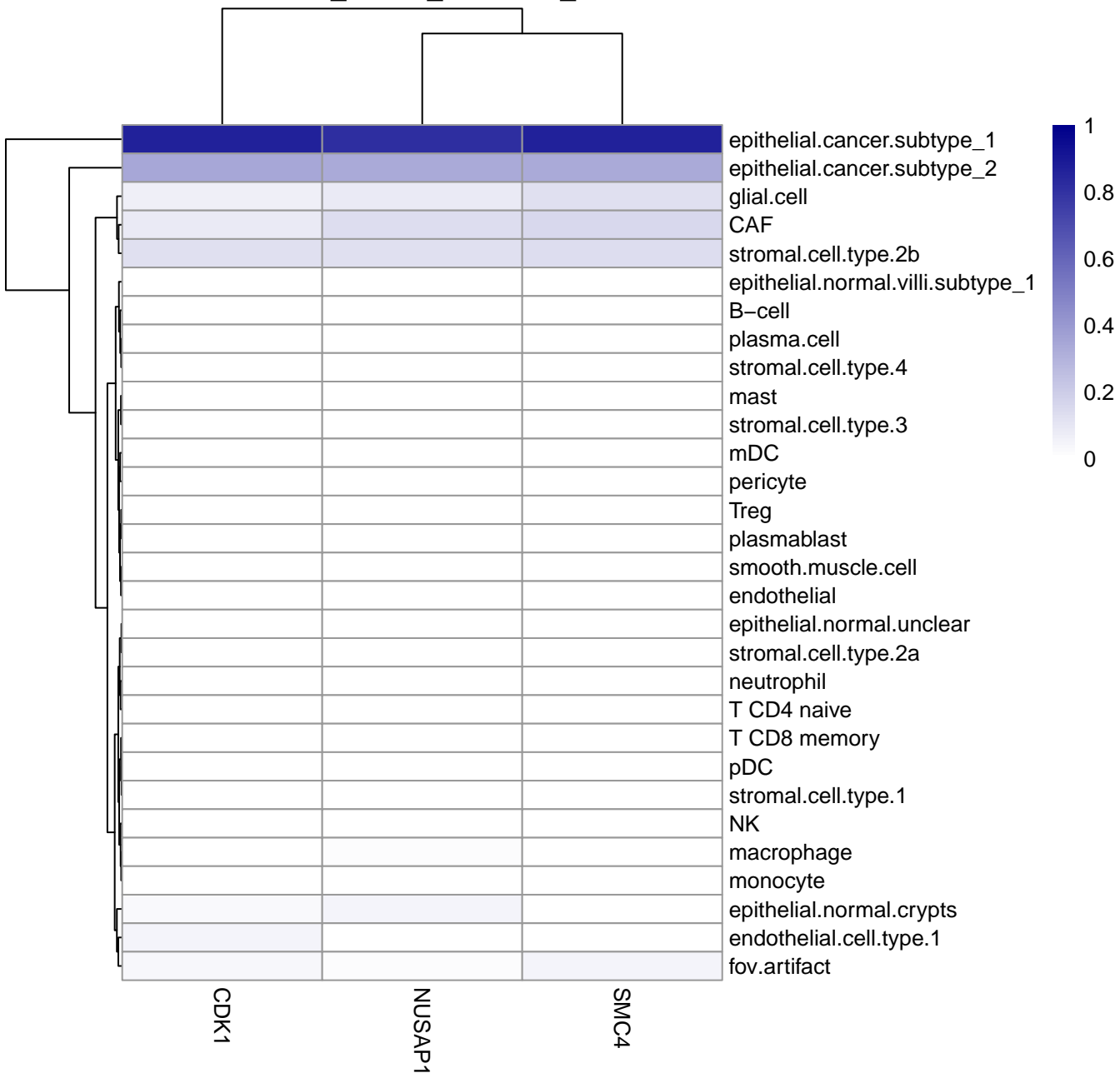
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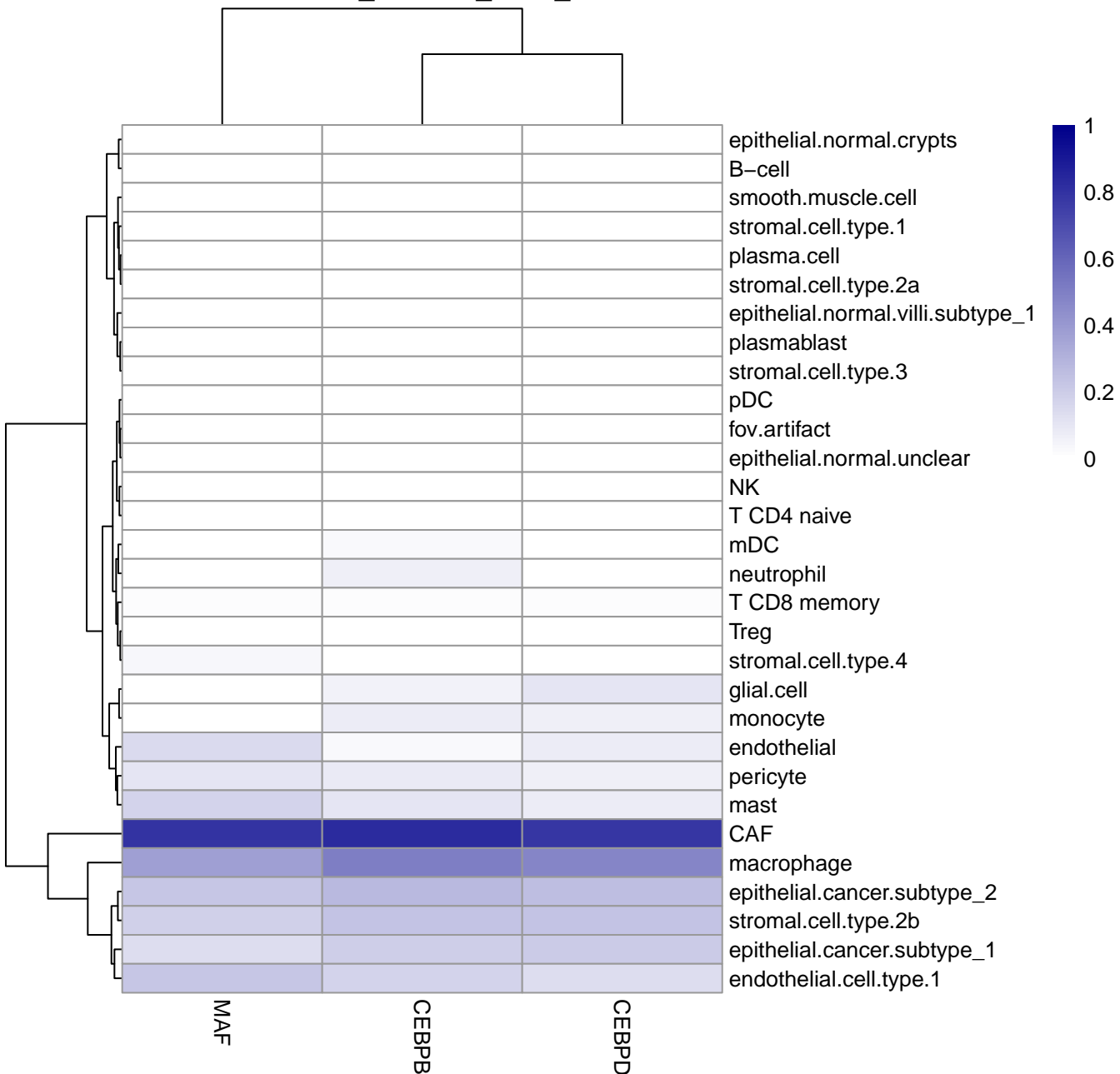
TRAM1_CD59_CD47_3



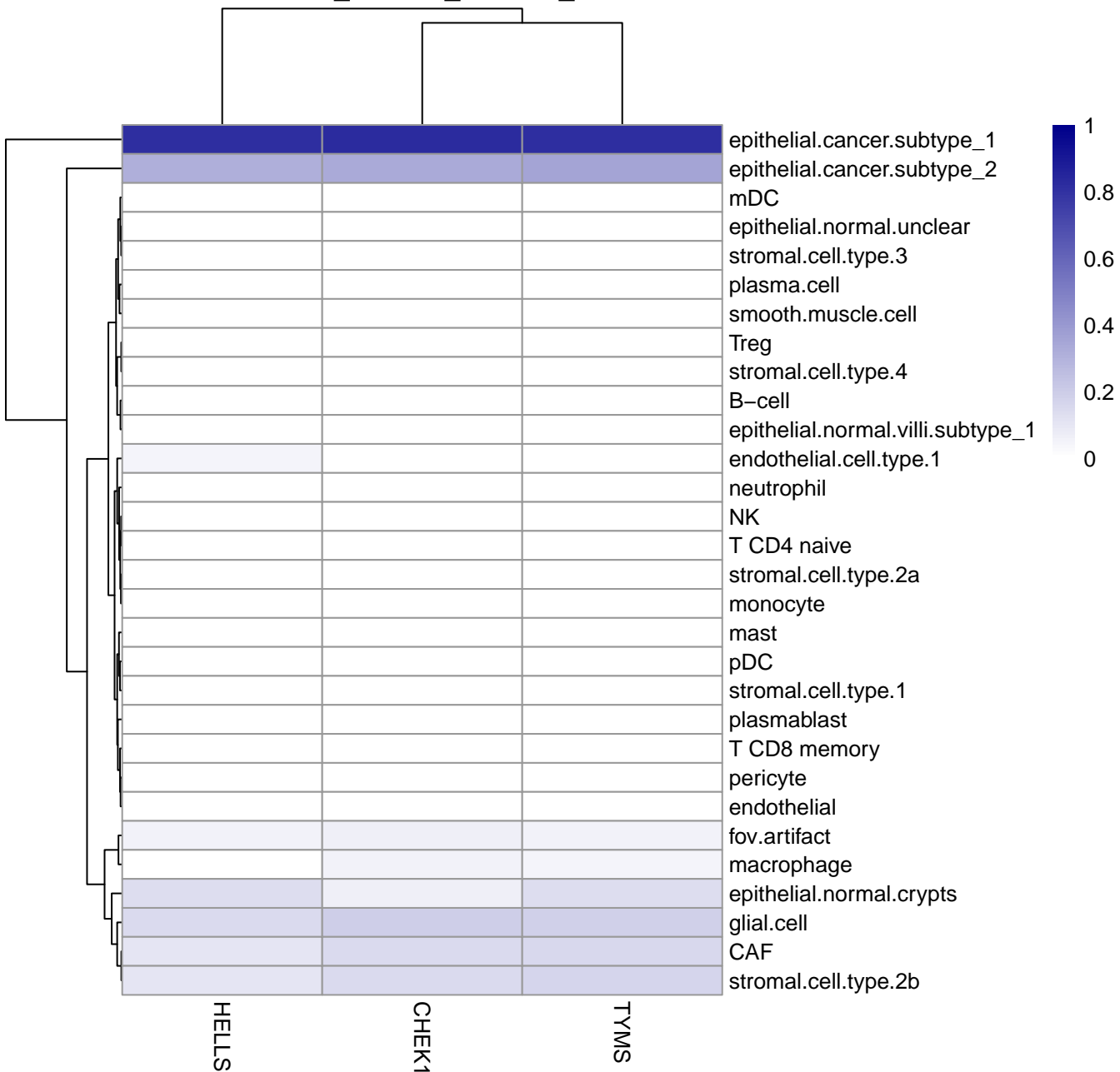
CDK1_SMC4_NUSAP1_3



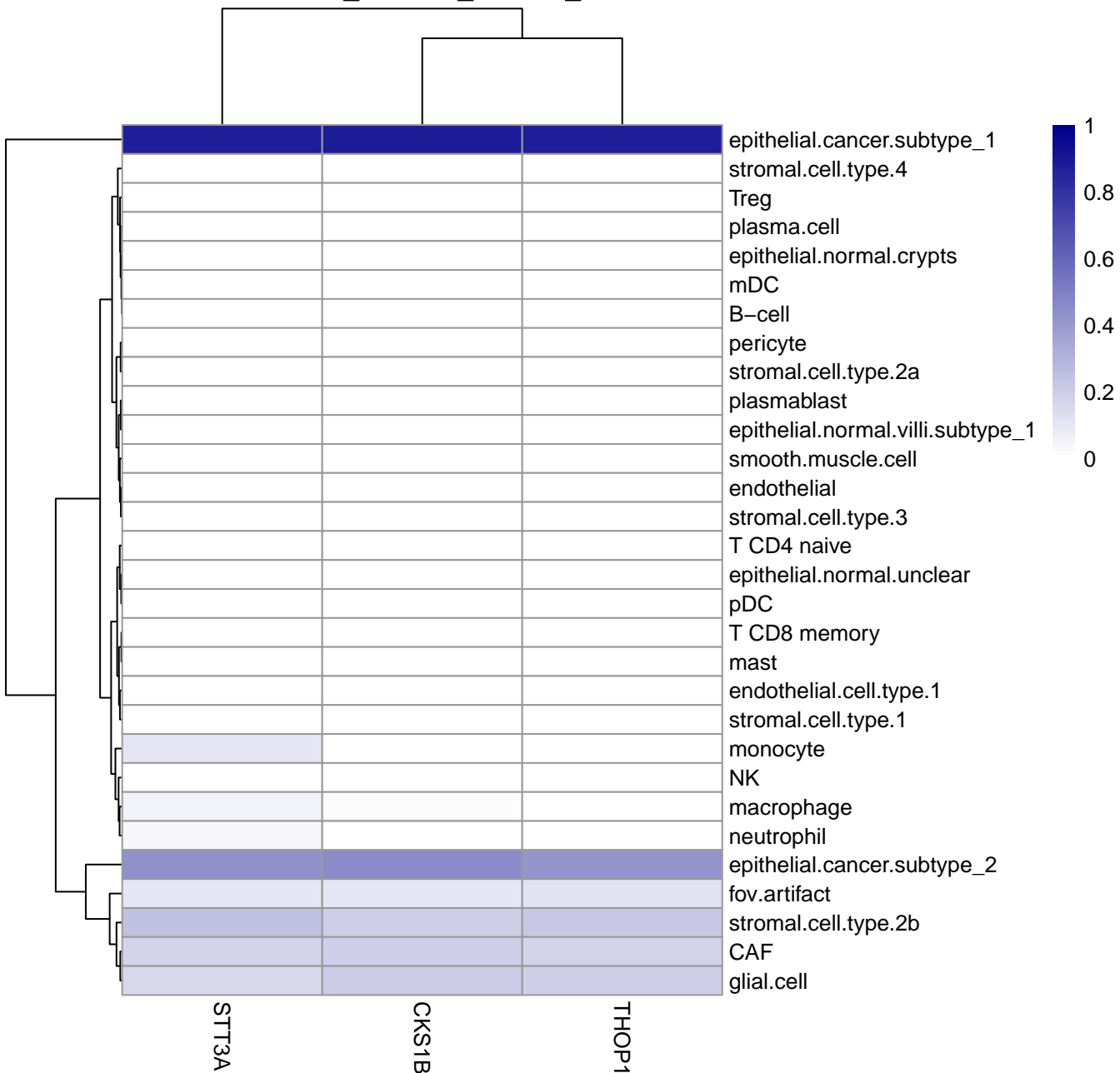
CEBPB_CEBPD_MAF_3



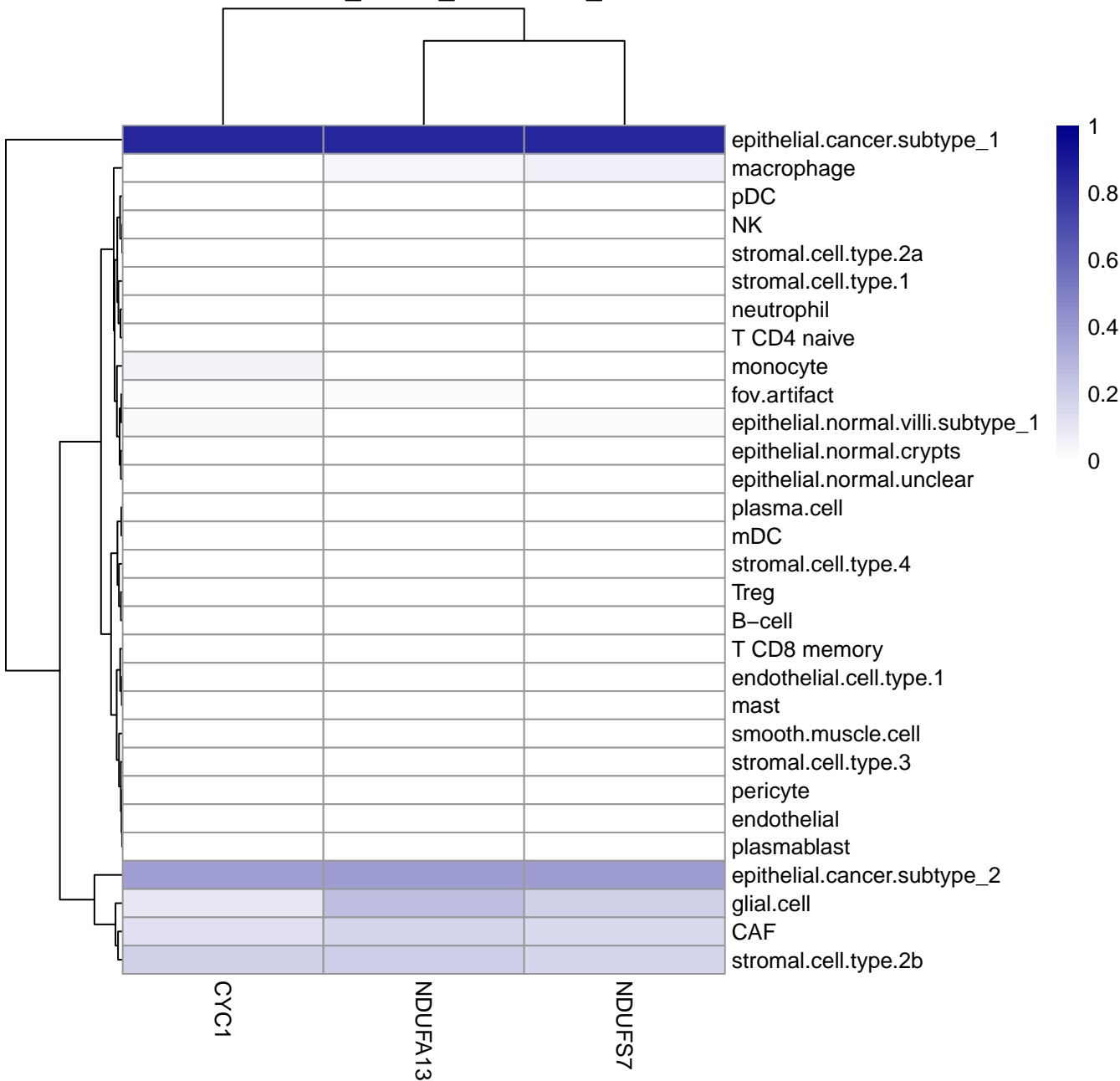
TYMS_HELLS_CHEK1_3



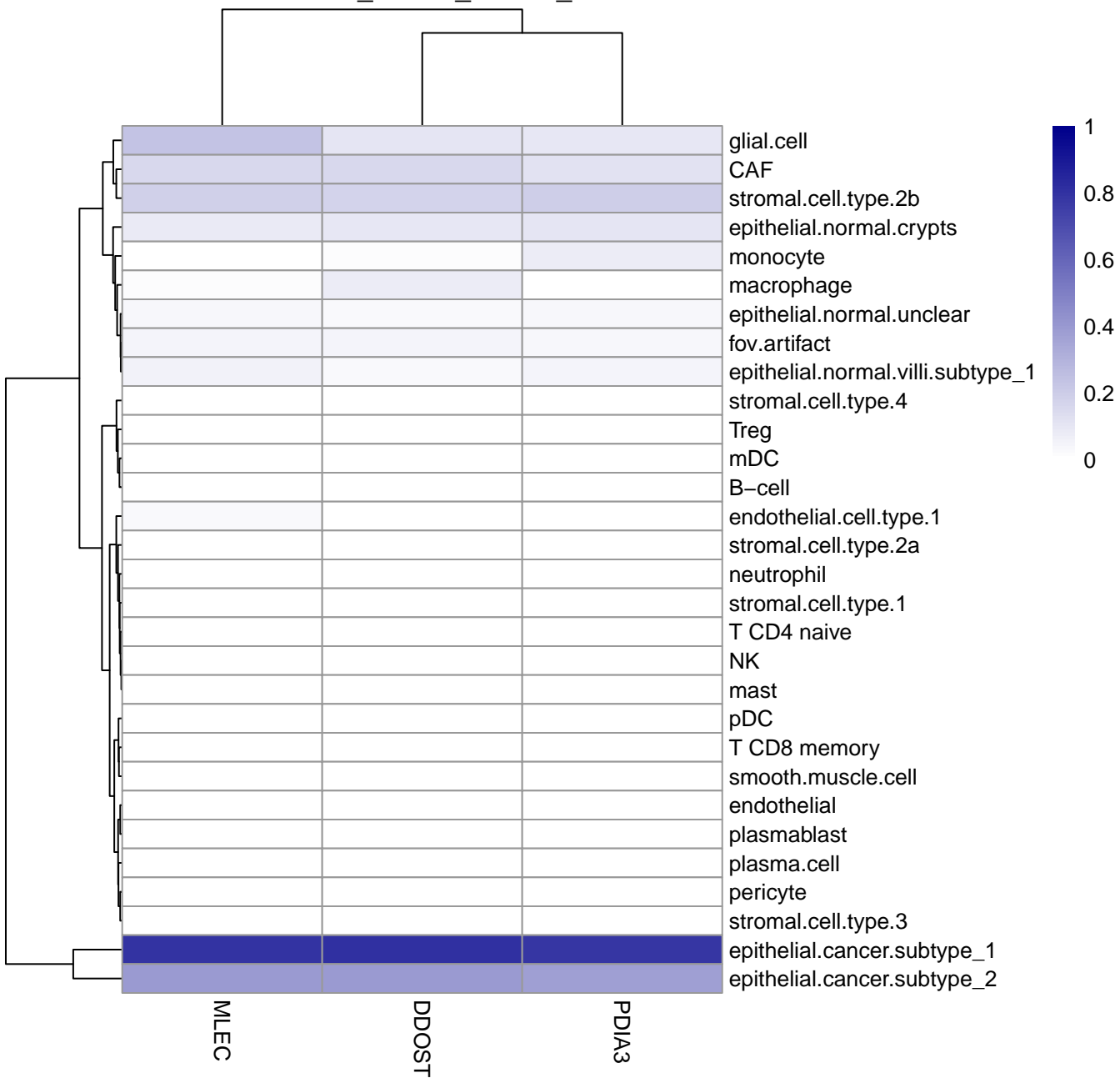
THOP1_CKS1B_STT3A_3



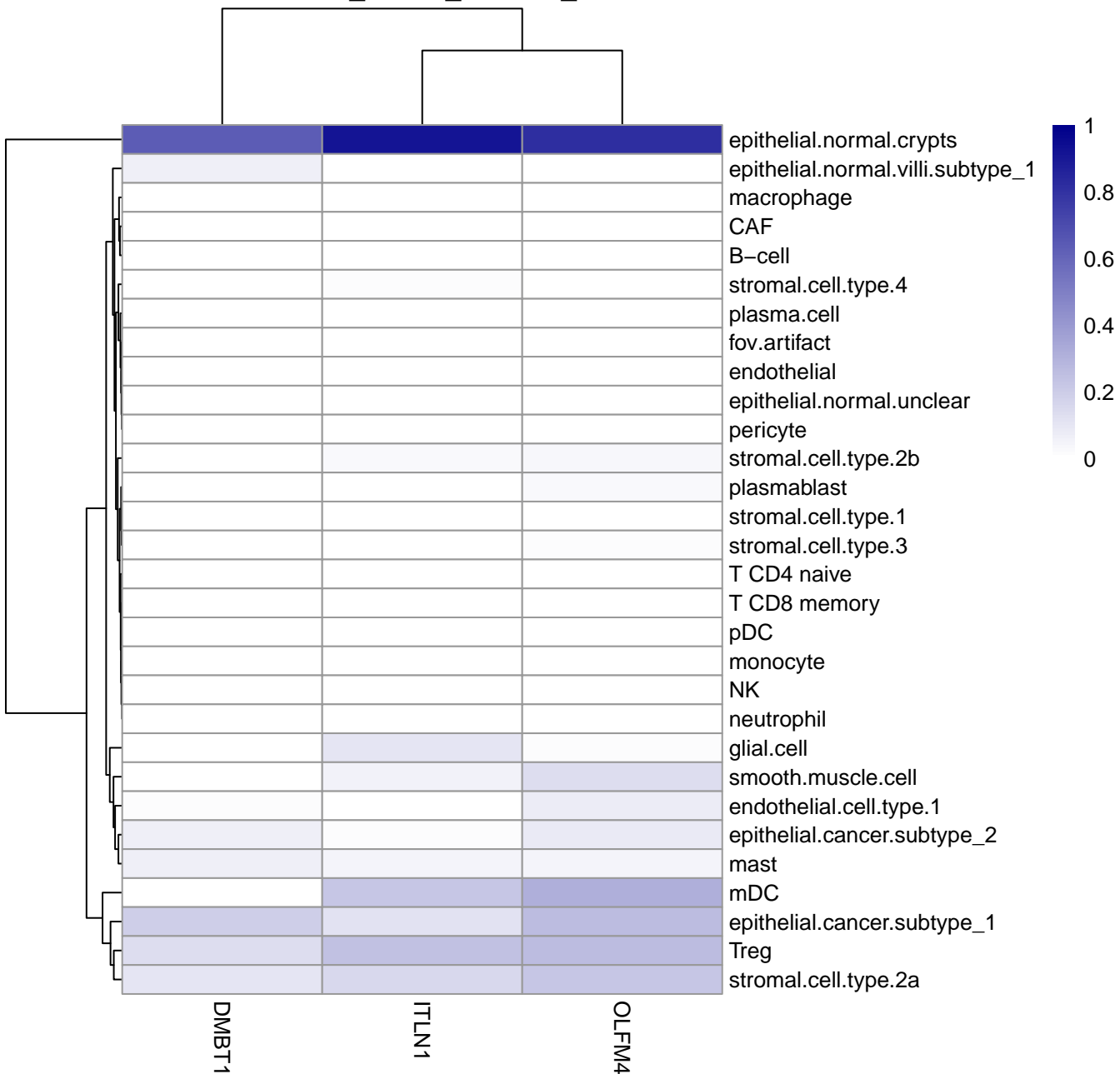
NDUFS7_CYC1_NDUFA13_3



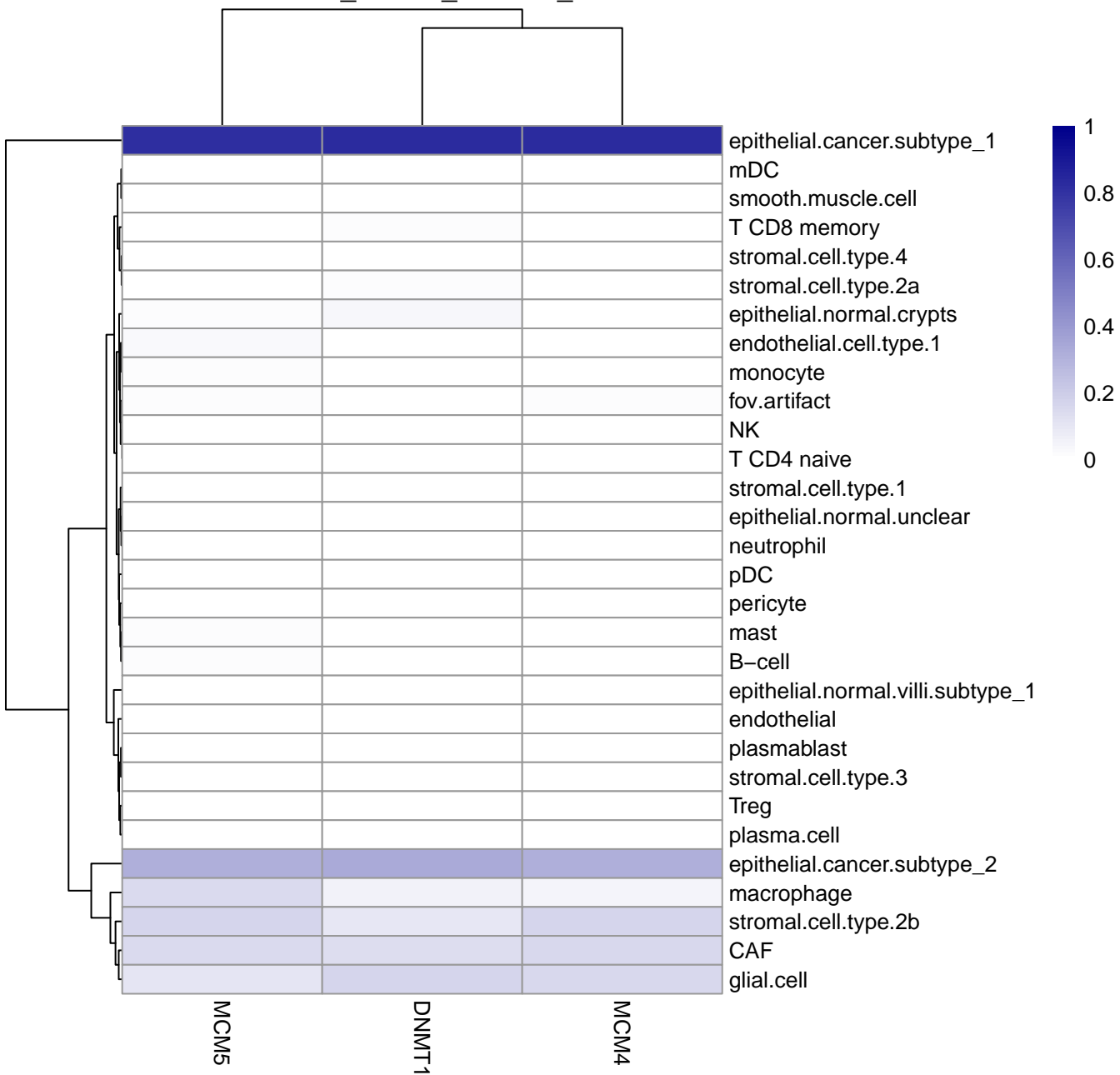
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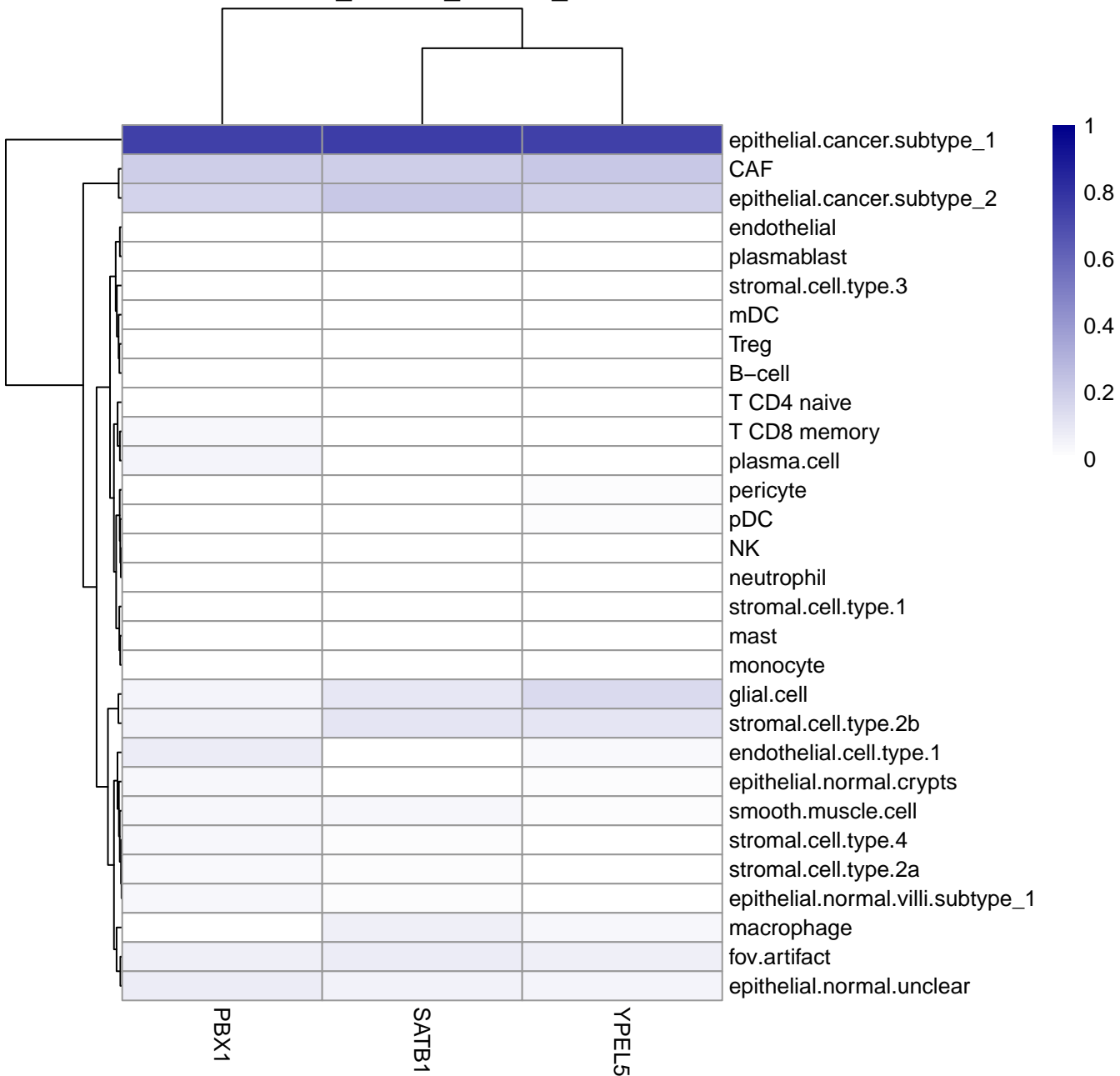
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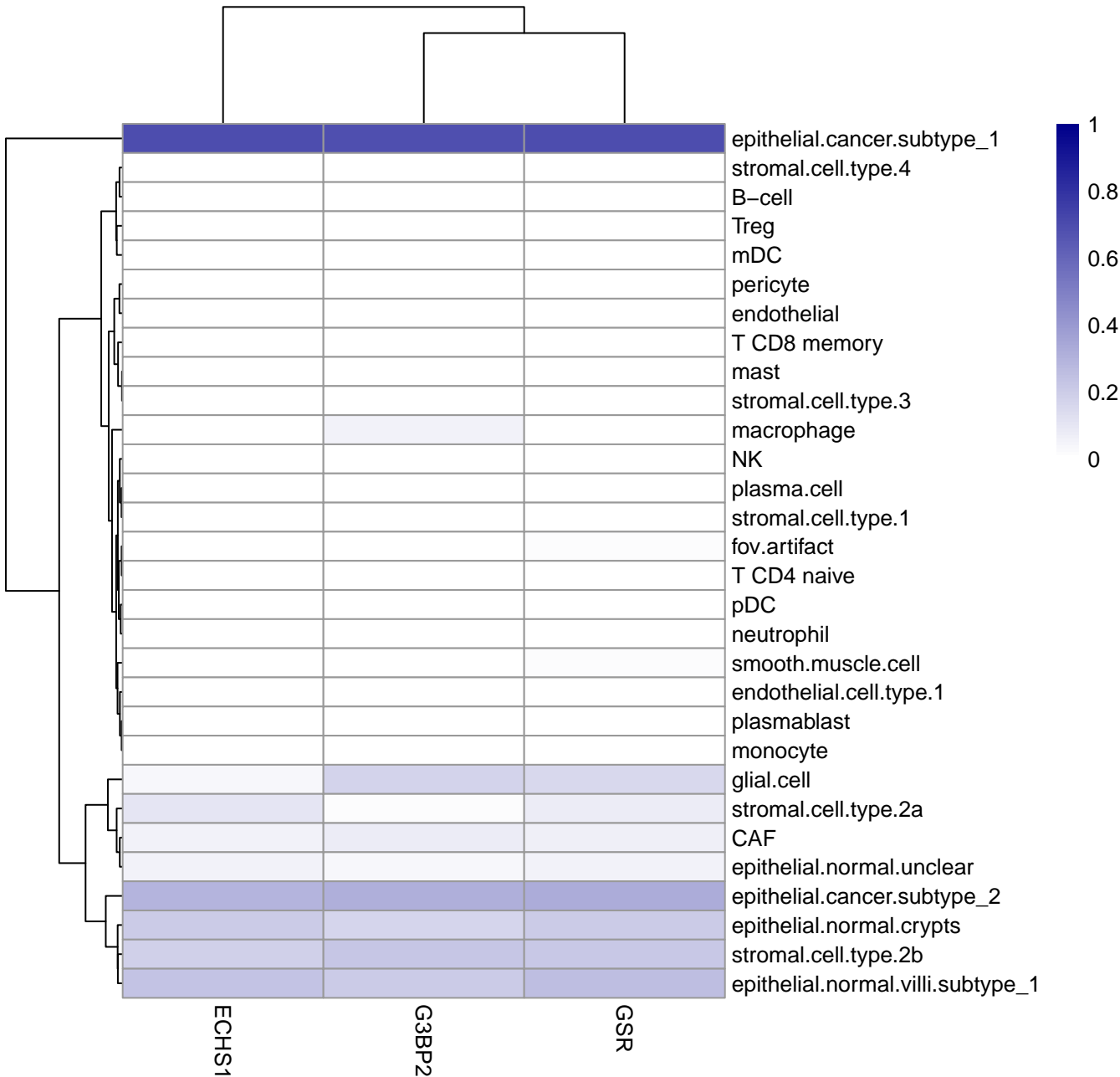
MCM4_MCM5_DNMT1_3



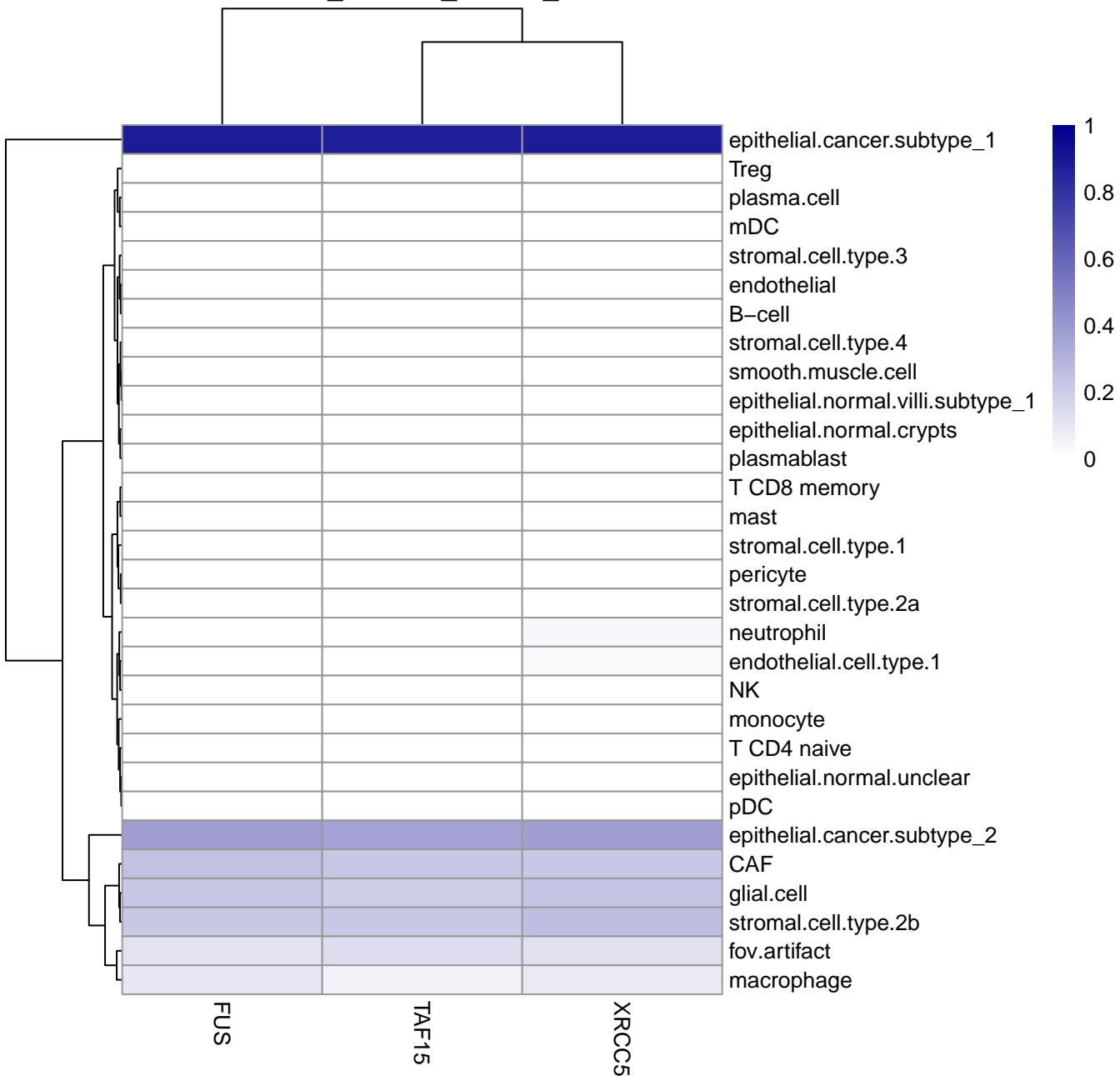
PBX1_SATB1_YPEL5_3



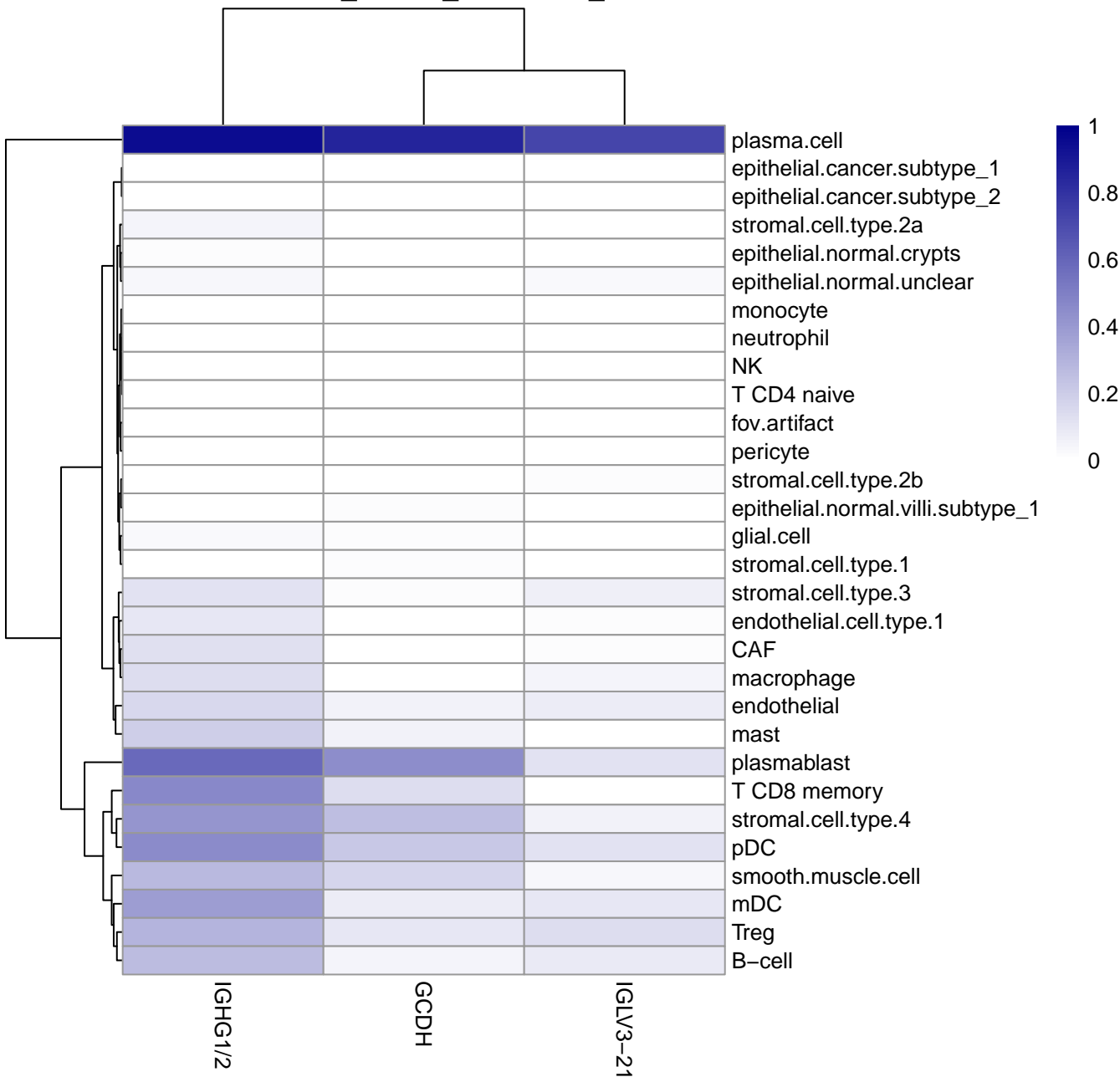
GSR_ECHS1_G3BP2_3



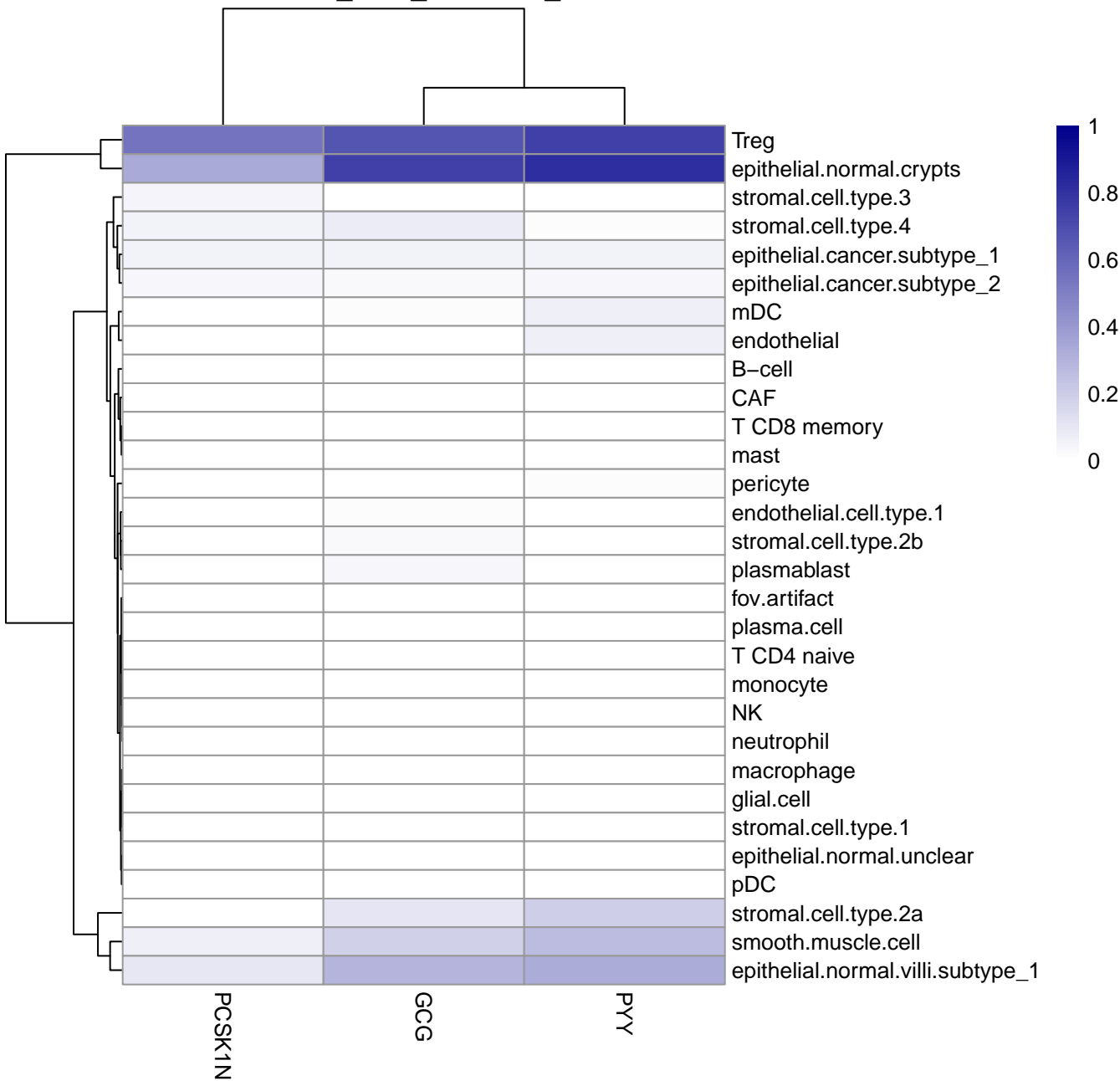
FUS_XRCC5_TAF15_3



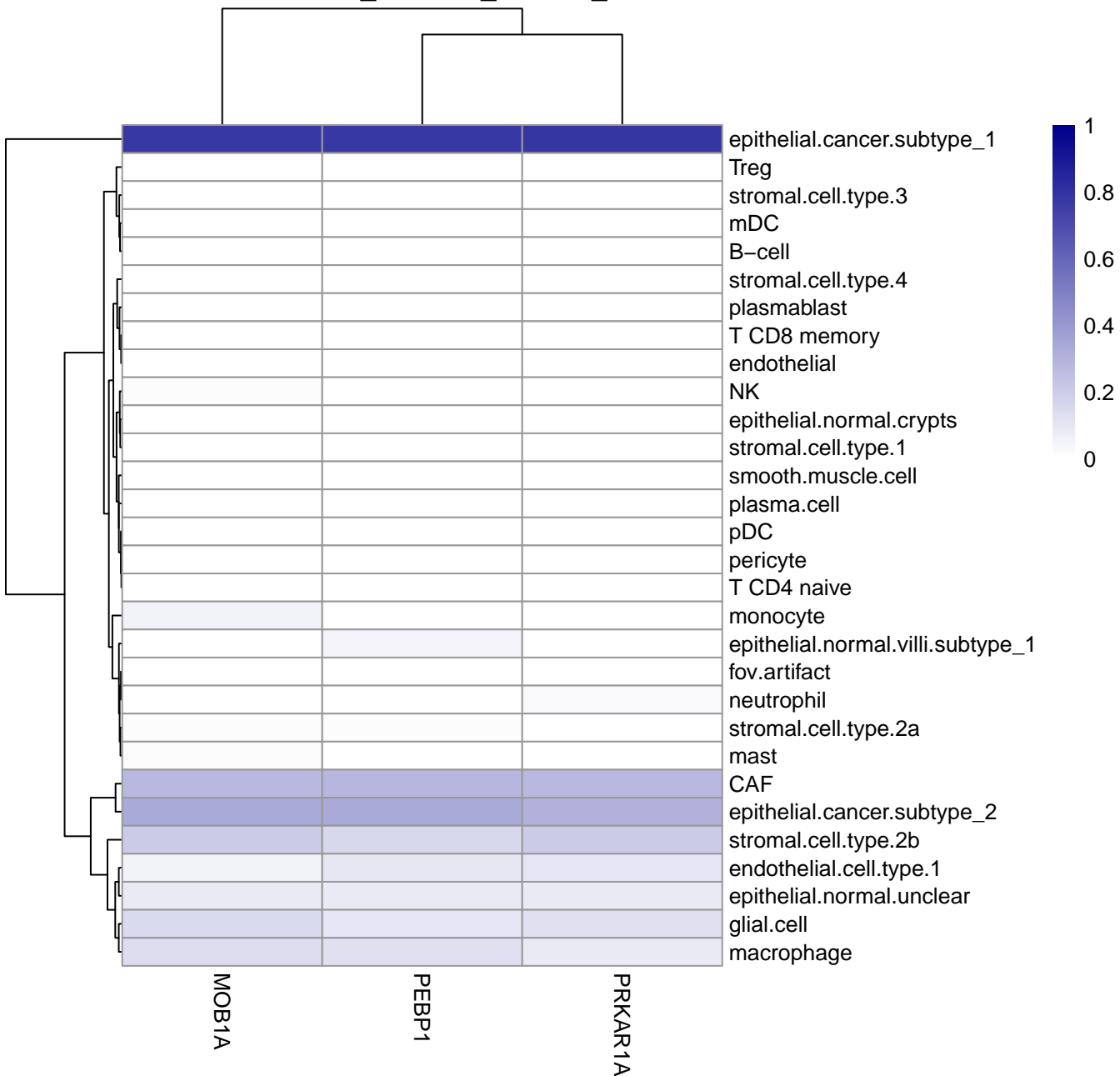
IGHG1/2_GCDH_IGLV3-21_3



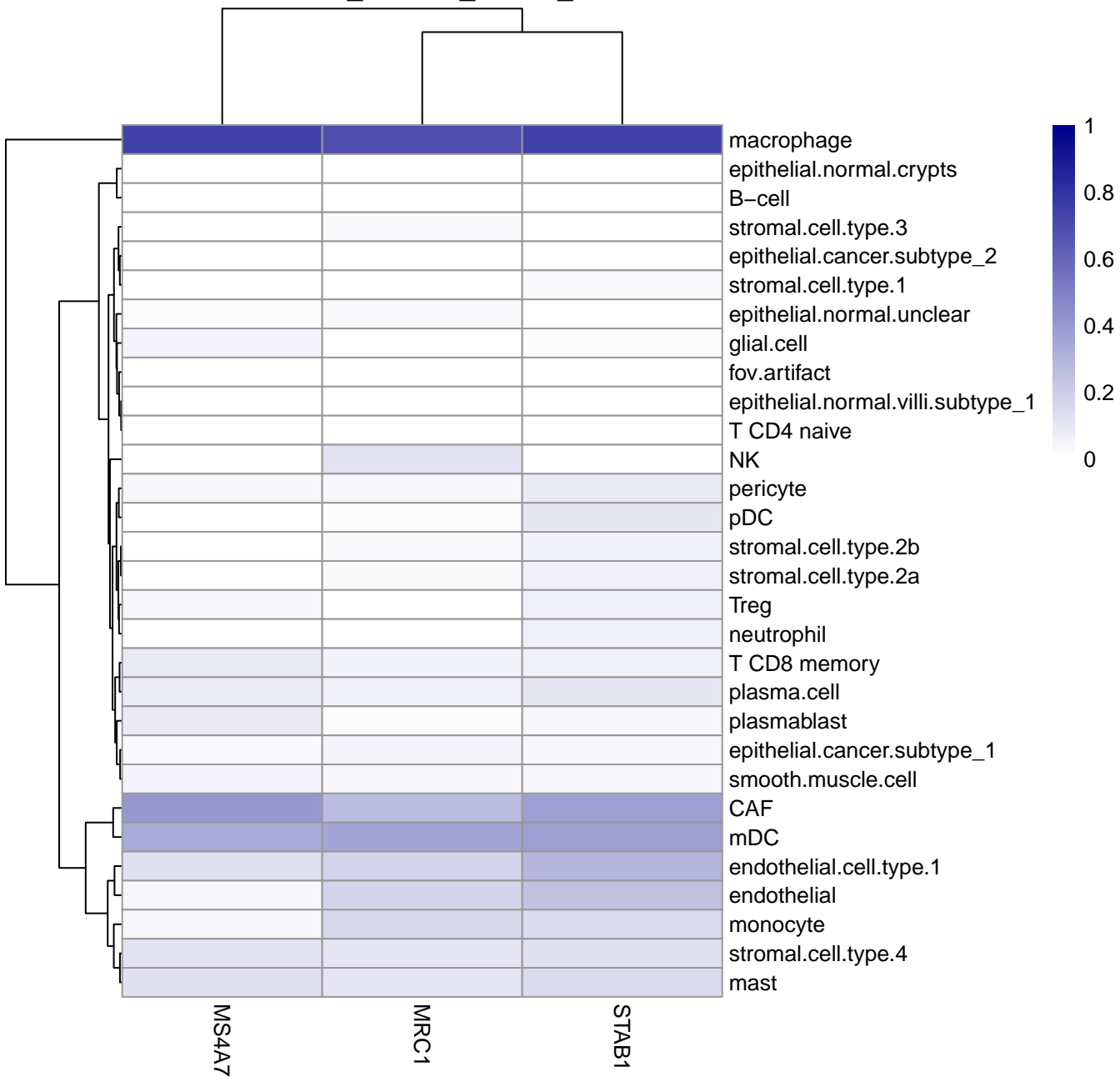
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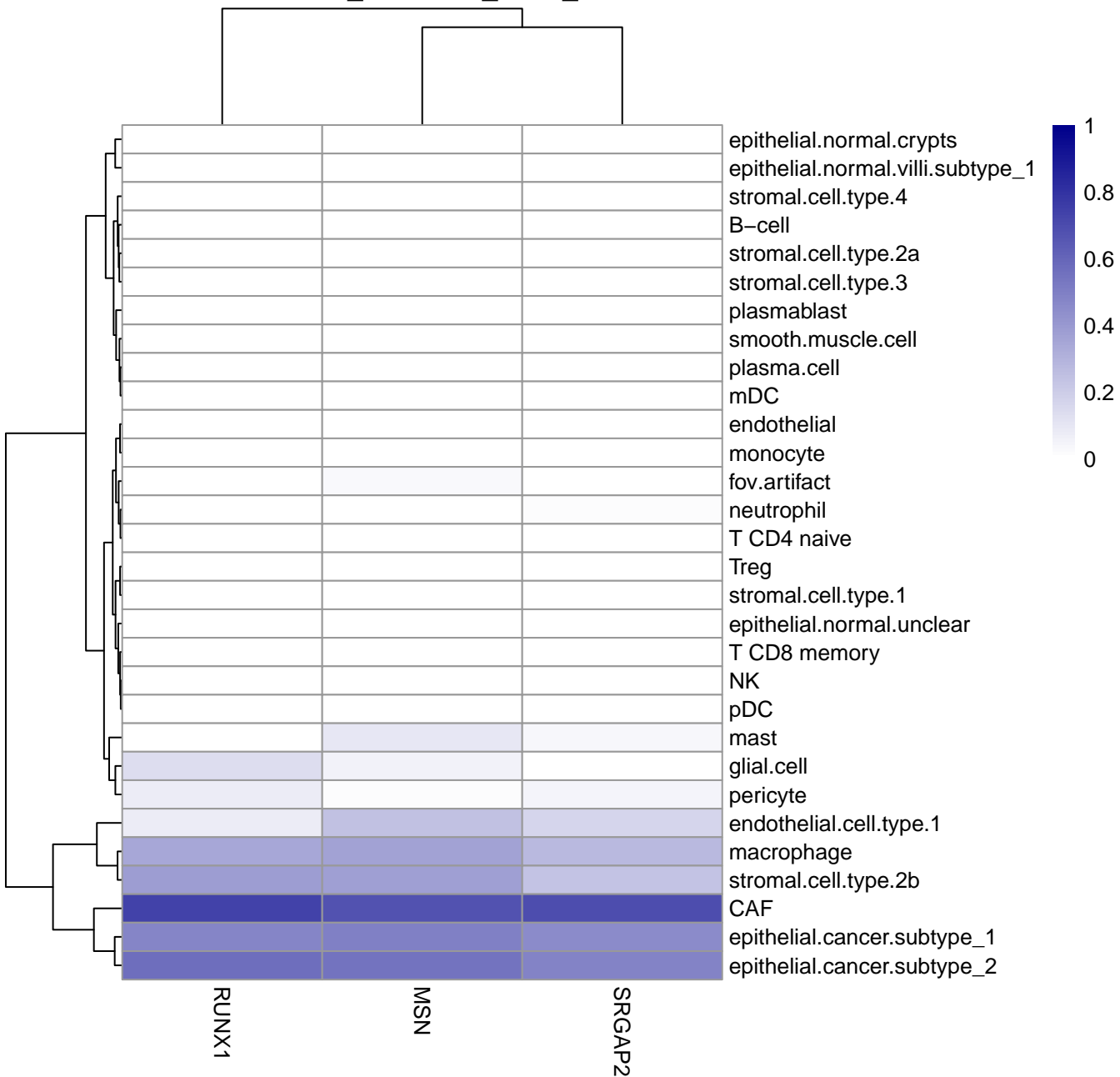
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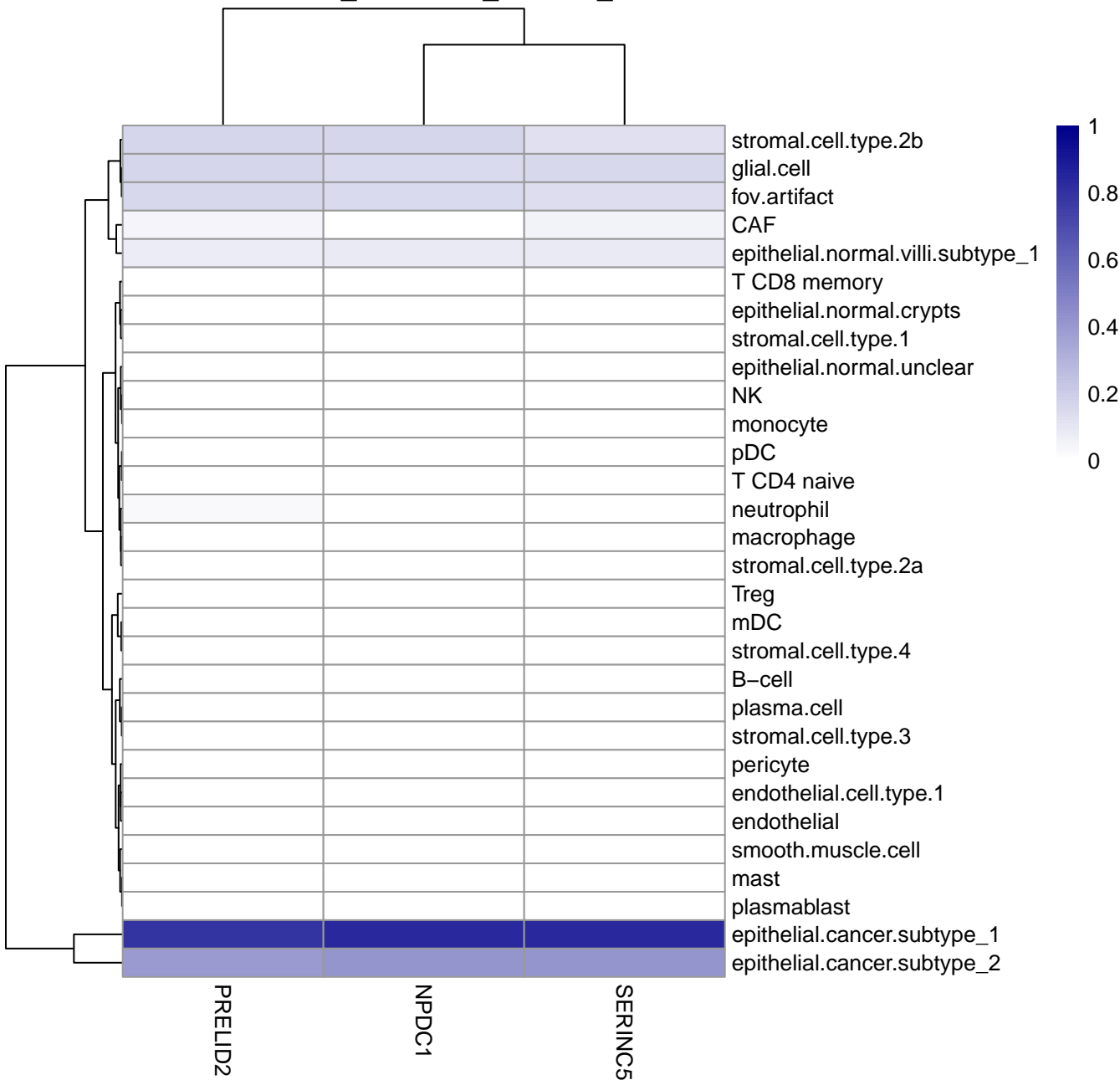
STAB1_MS4A7_MRC1_3



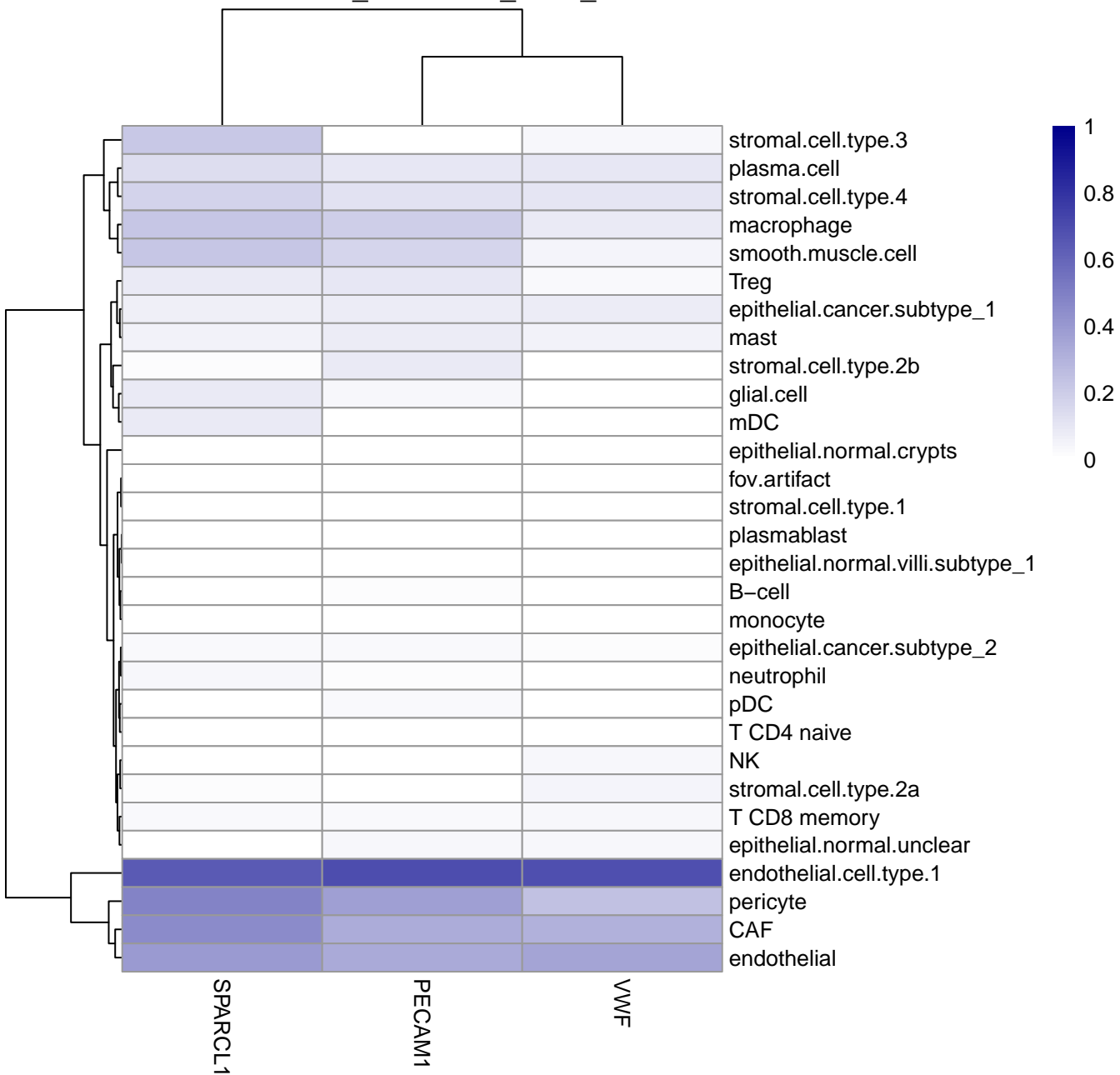
RUNX1_SRGAP2_MSN_3



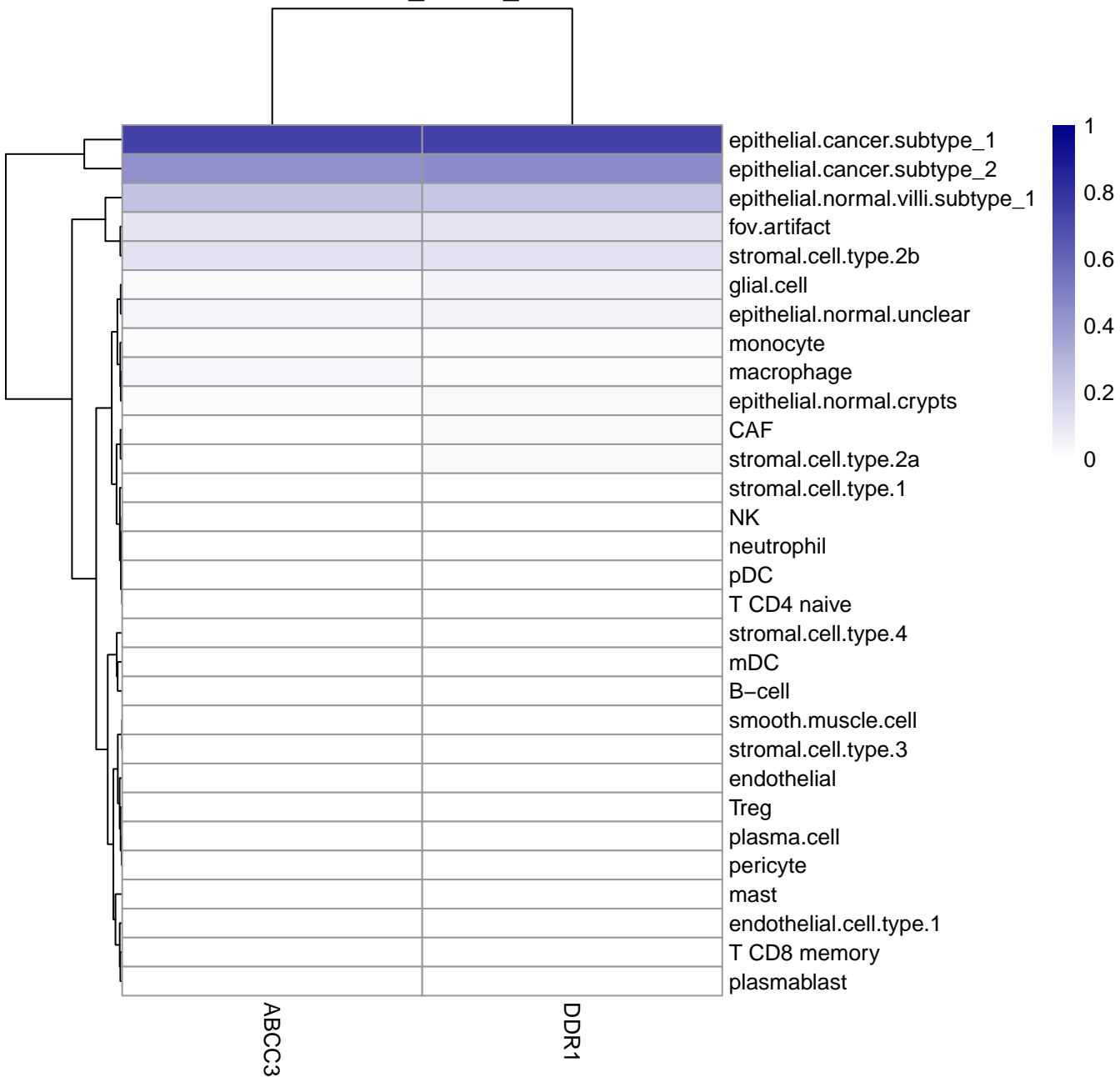
PRELID2_SERINC5_NPDC1_3



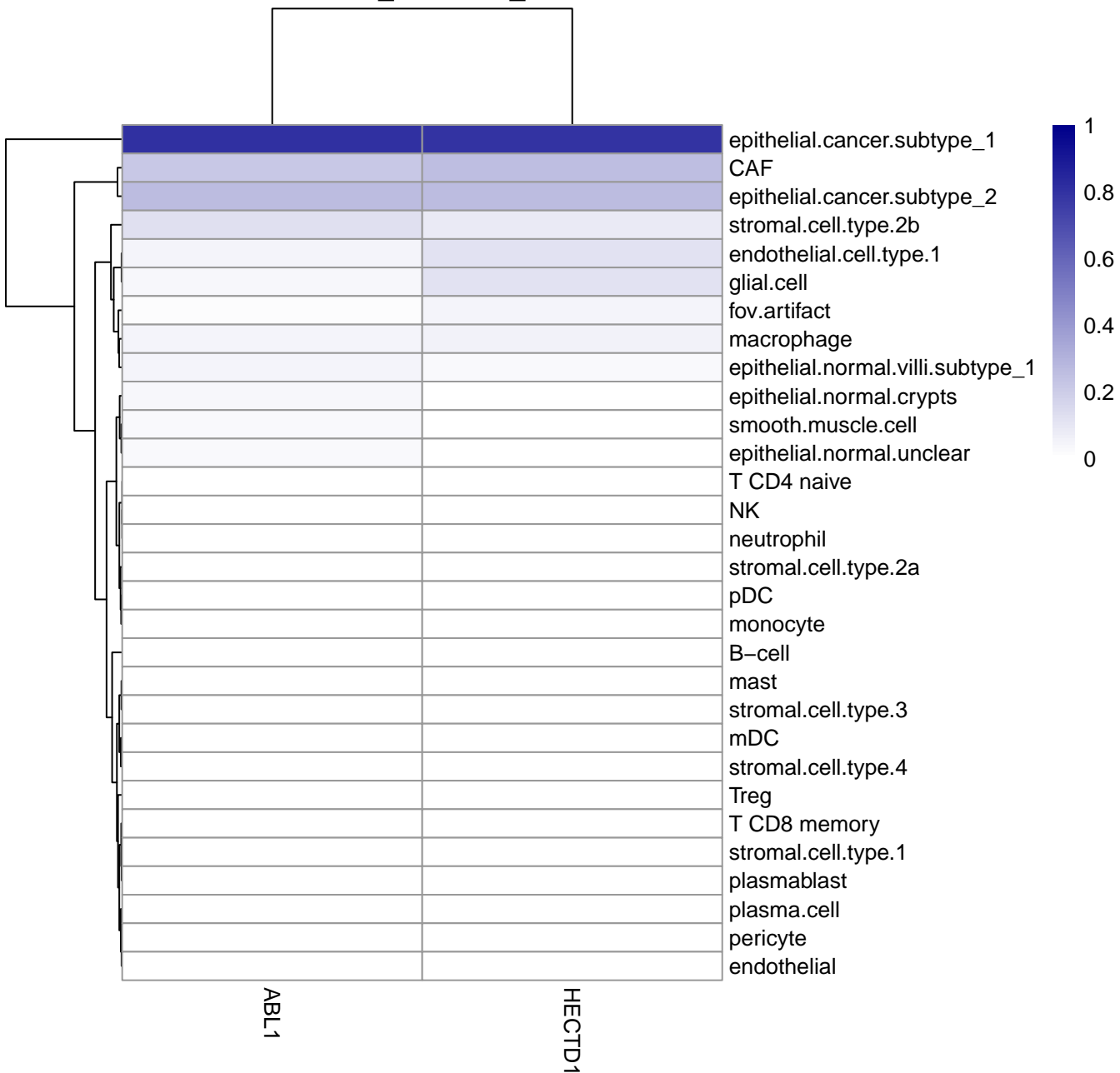
PECAM1_SPARCL1_VWF_3



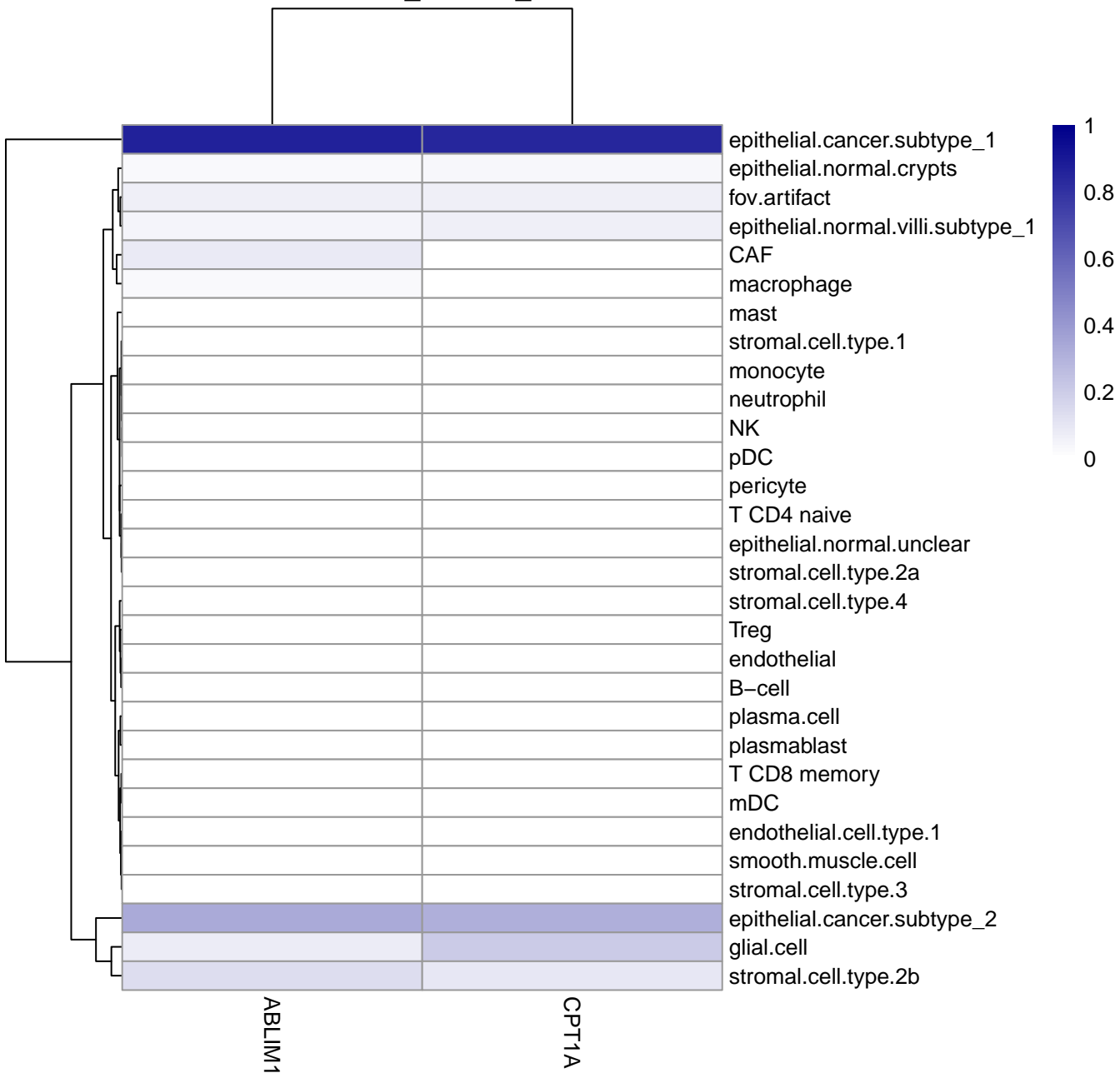
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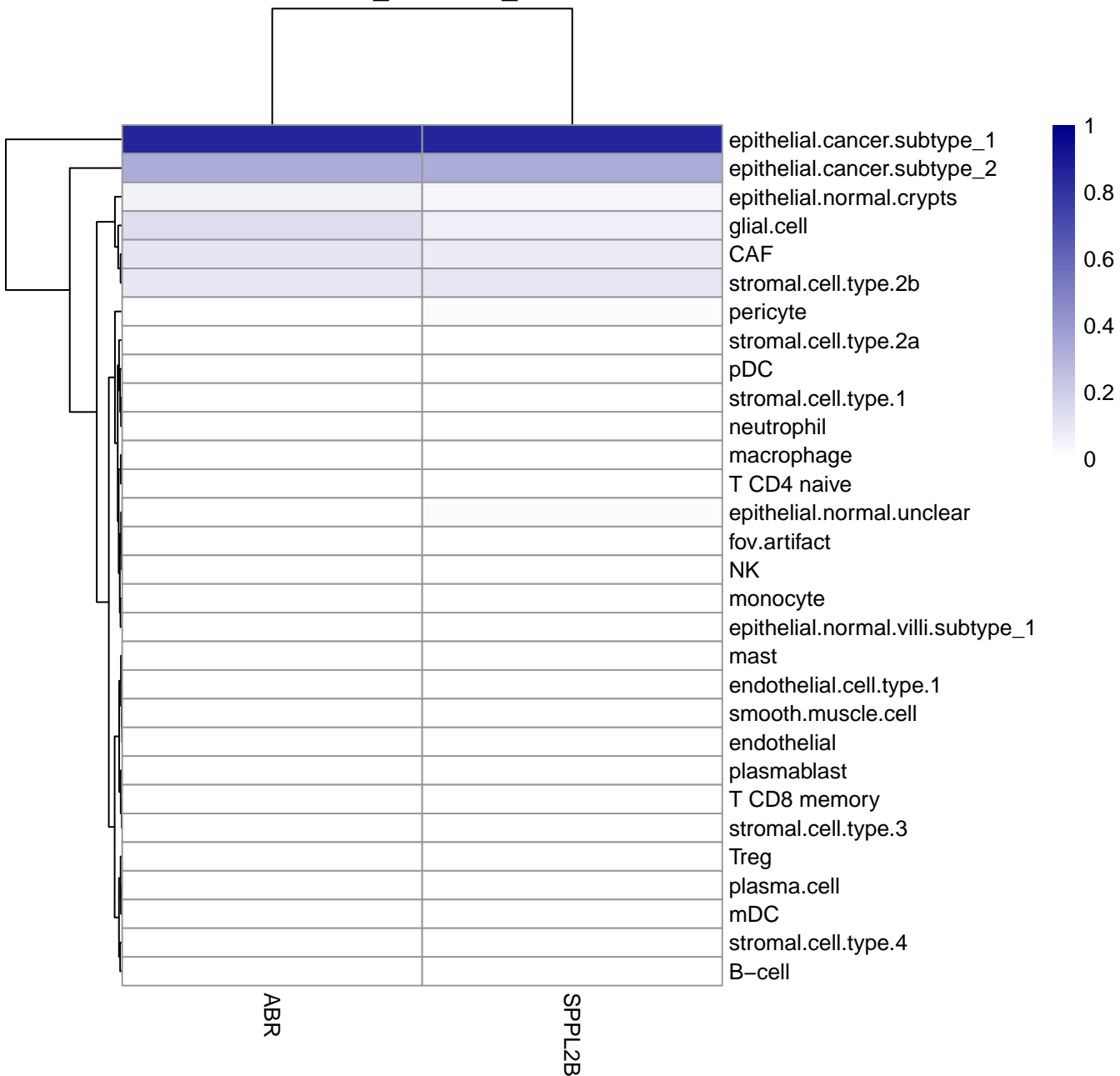
ABL1_HECTD1_2



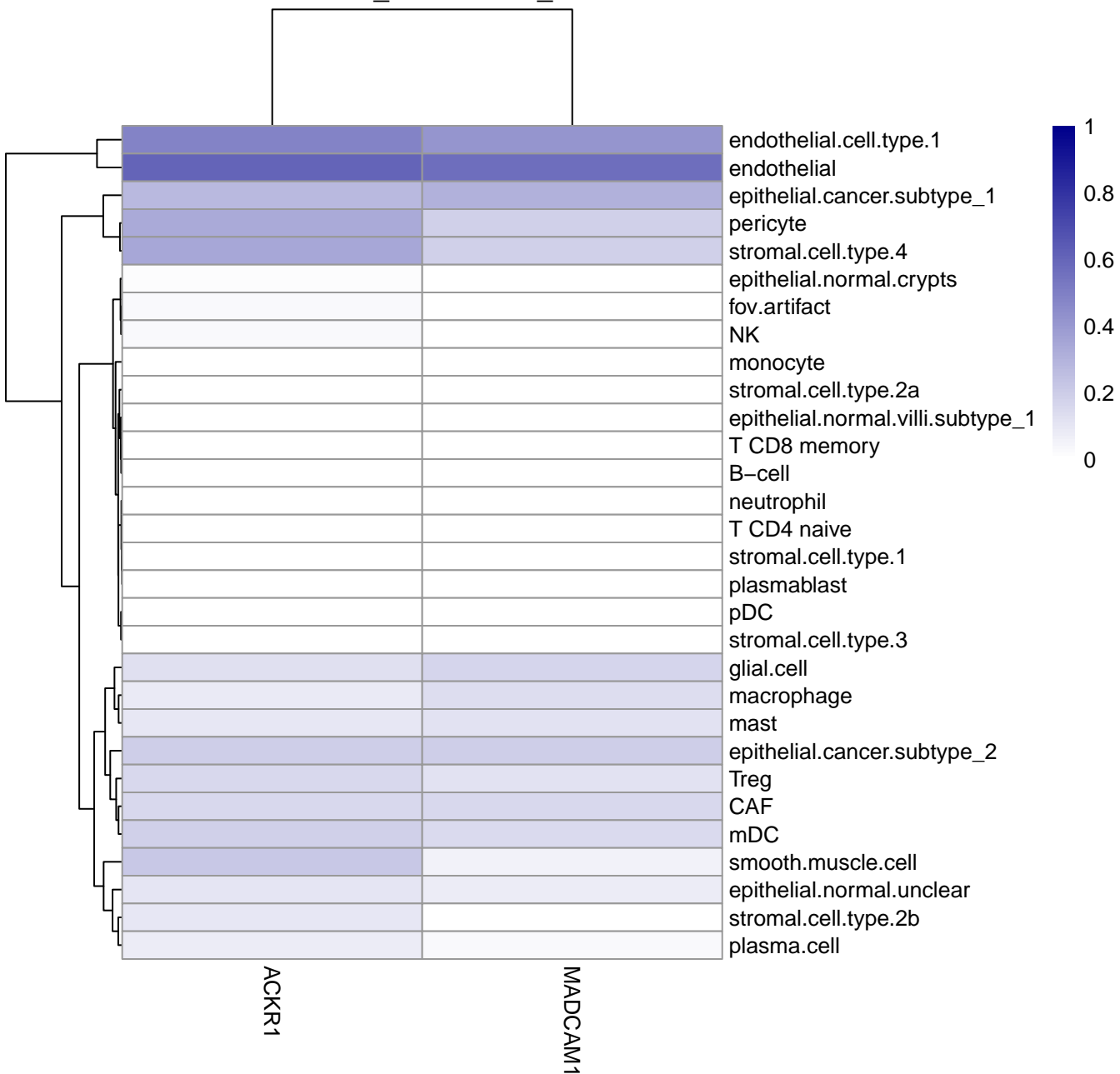
ABLIM1_CPT1A_2



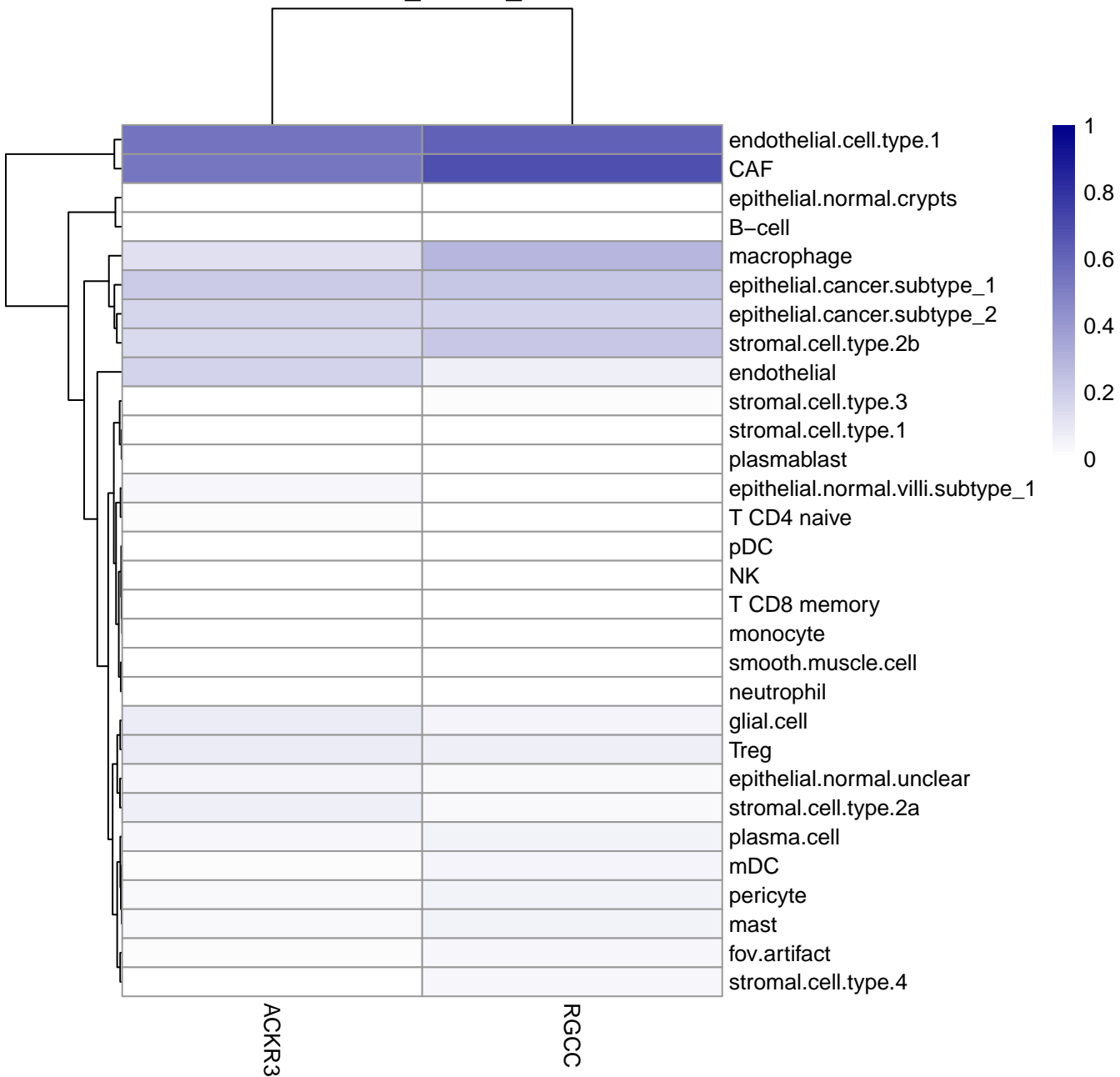
ABR_SPPL2B_2



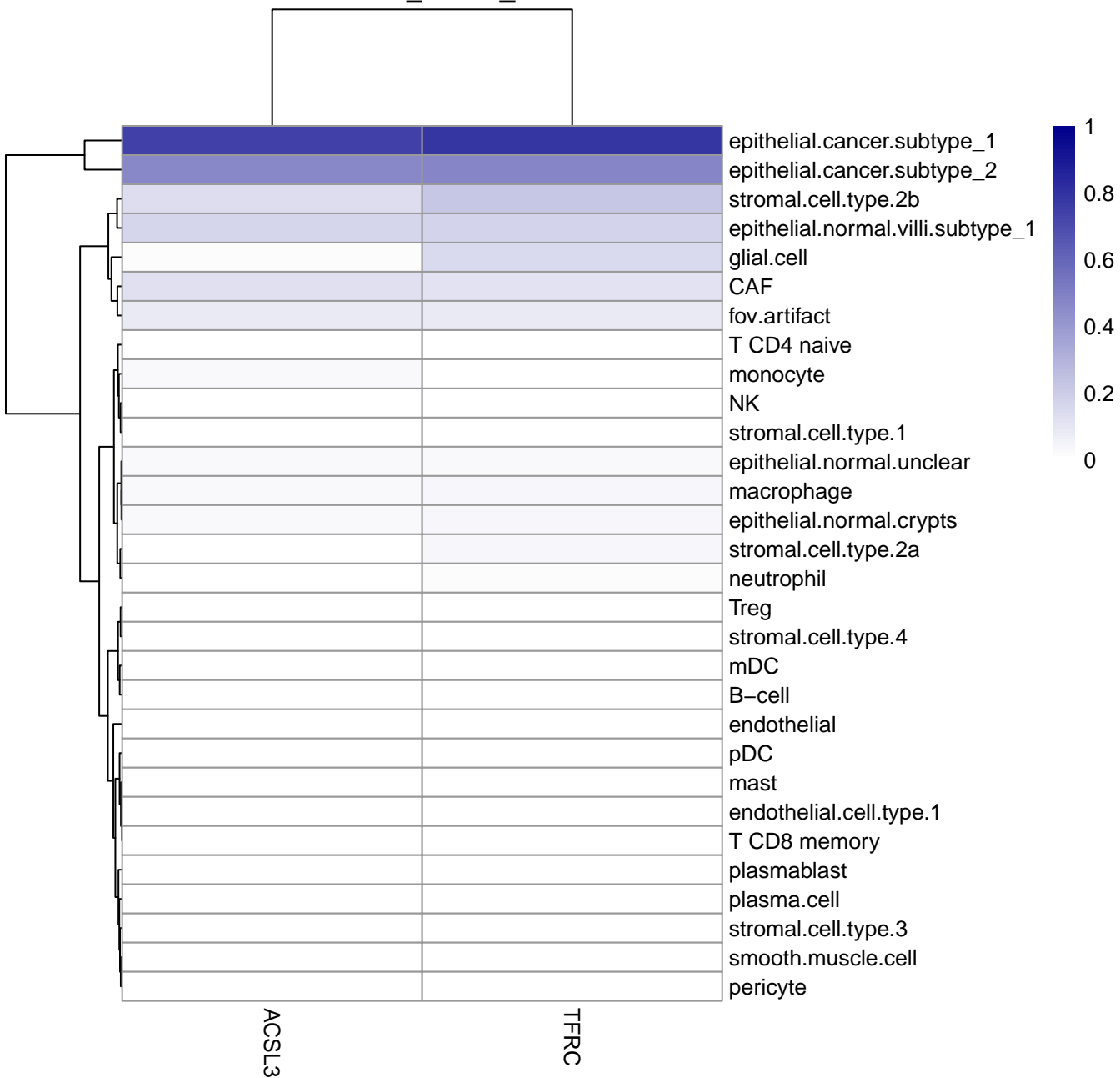
ACKR1_MADCAM1_2



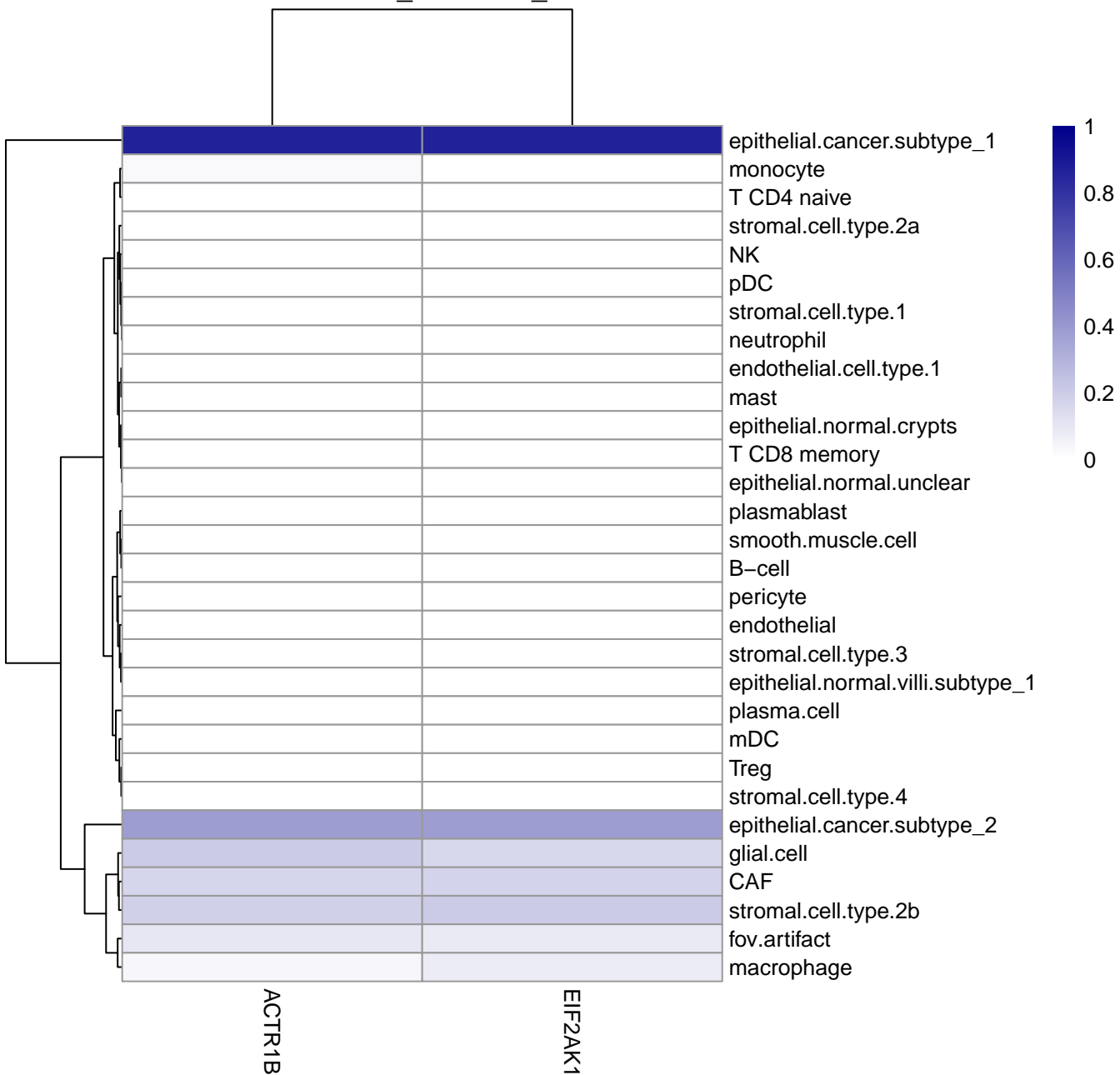
ACKR3_RGCC_2



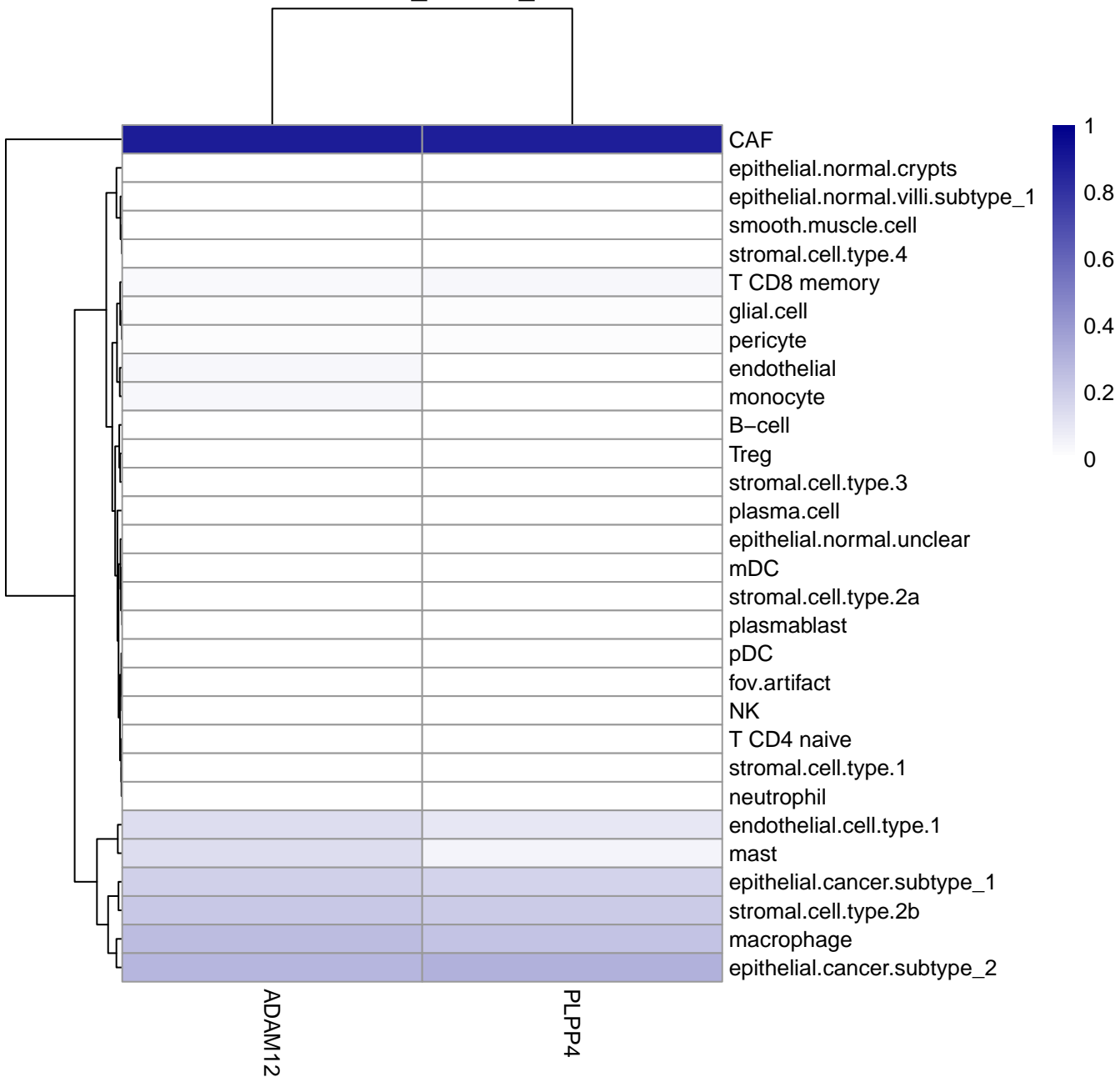
ACSL3_TFRC_2



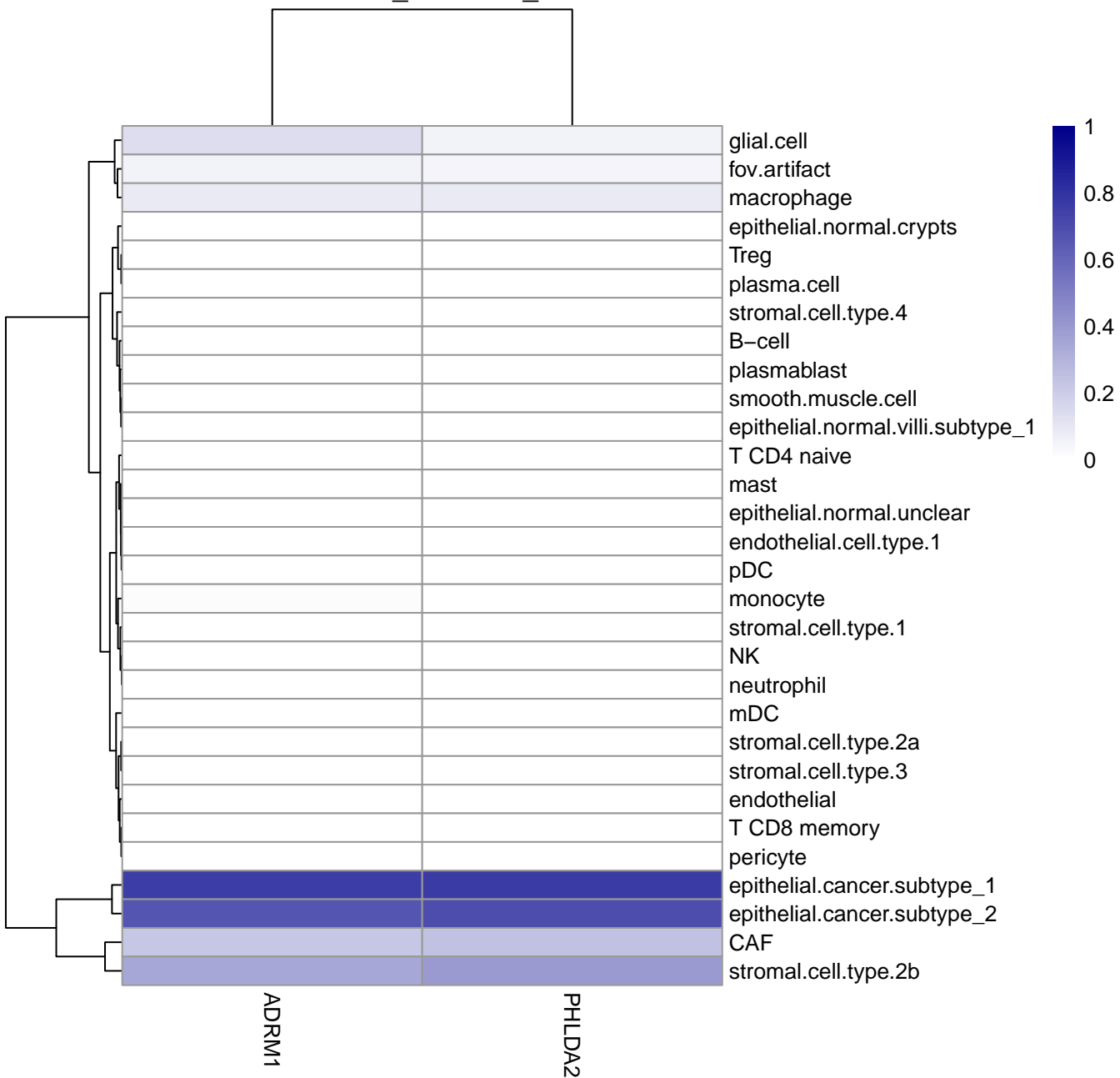
ACTR1B EIF2AK1_2



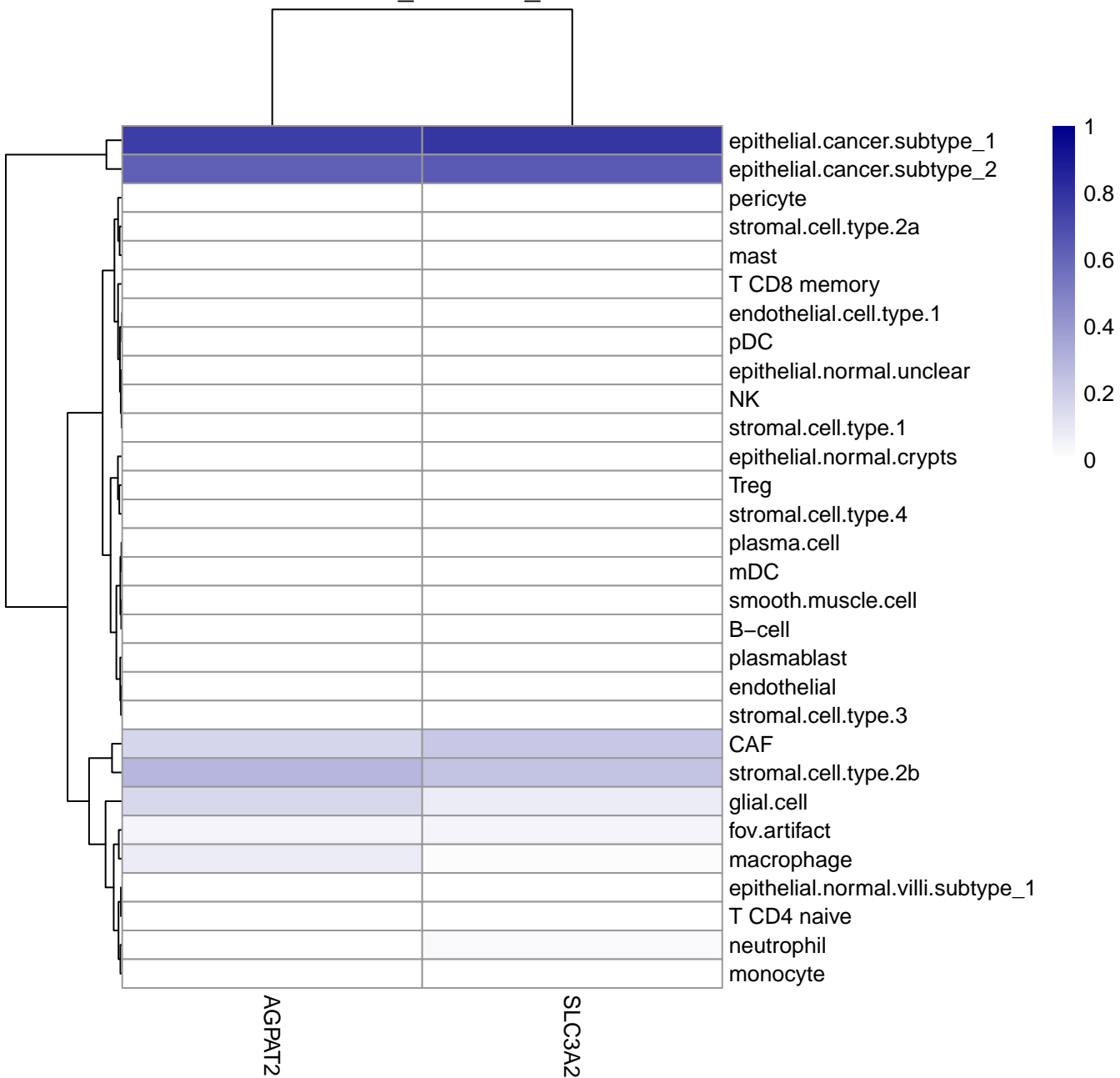
ADAM12_PLPP4_2



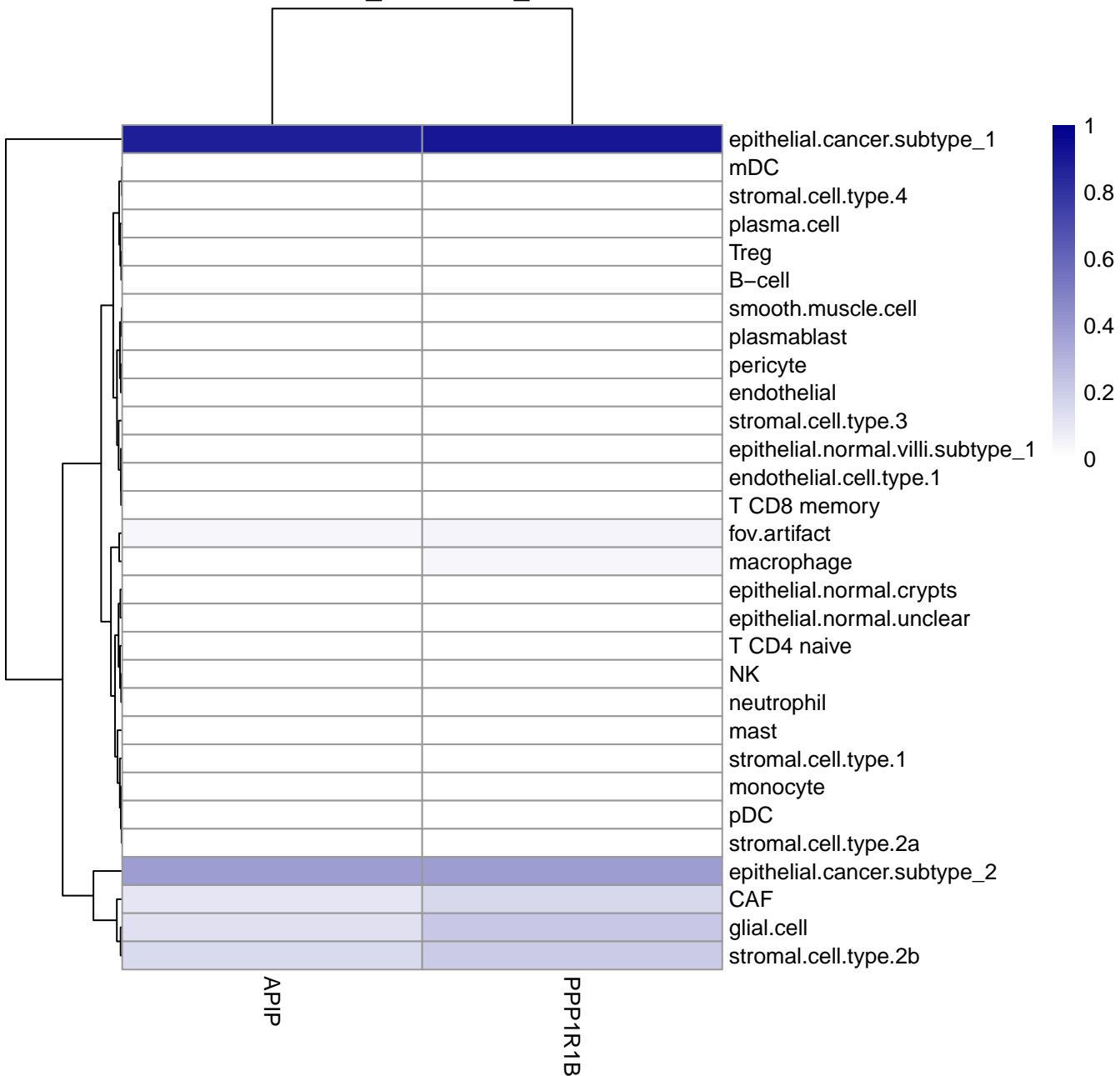
ADRM1_PHLDA2_2



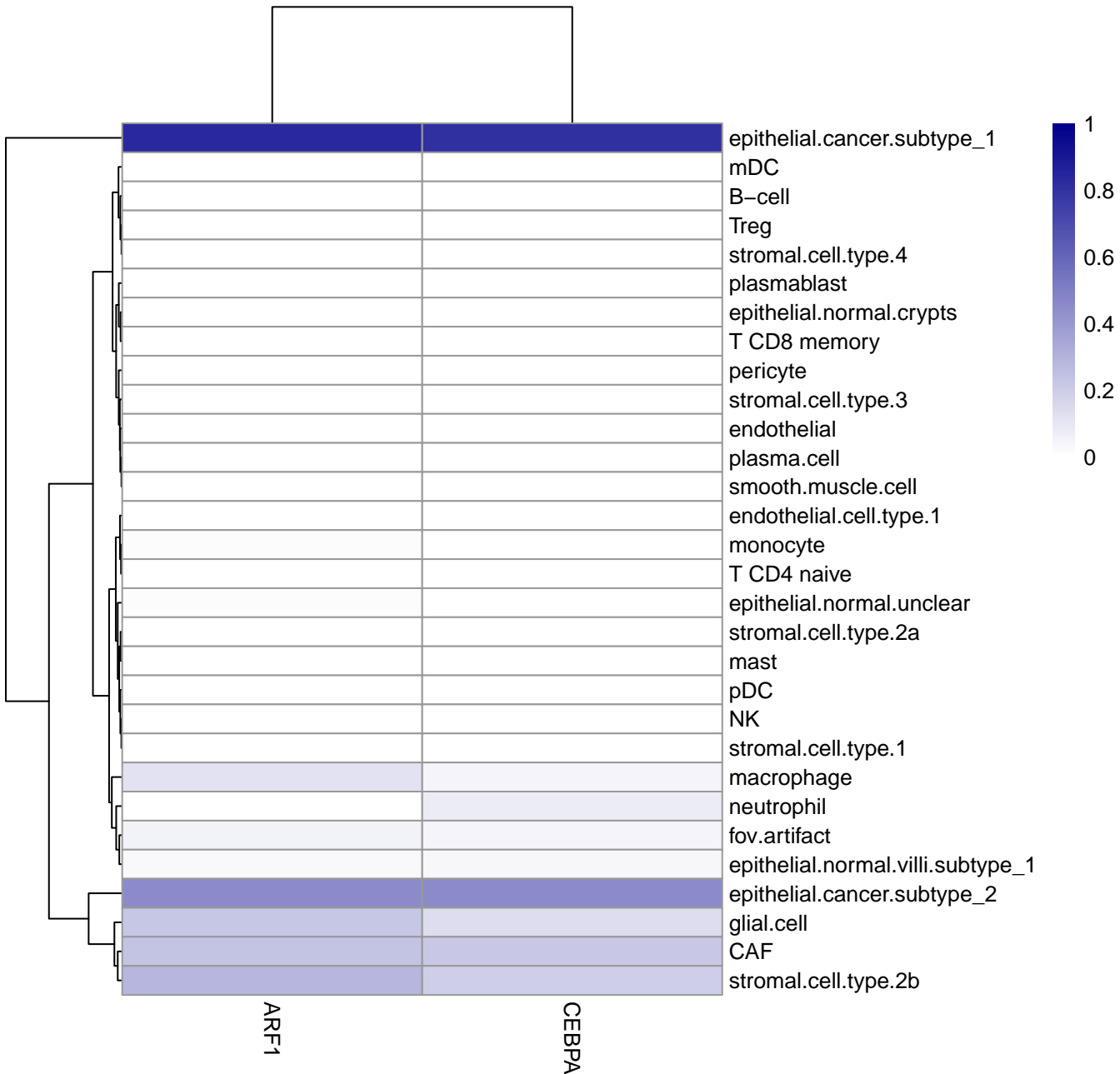
AGPAT2_SLC3A2_2



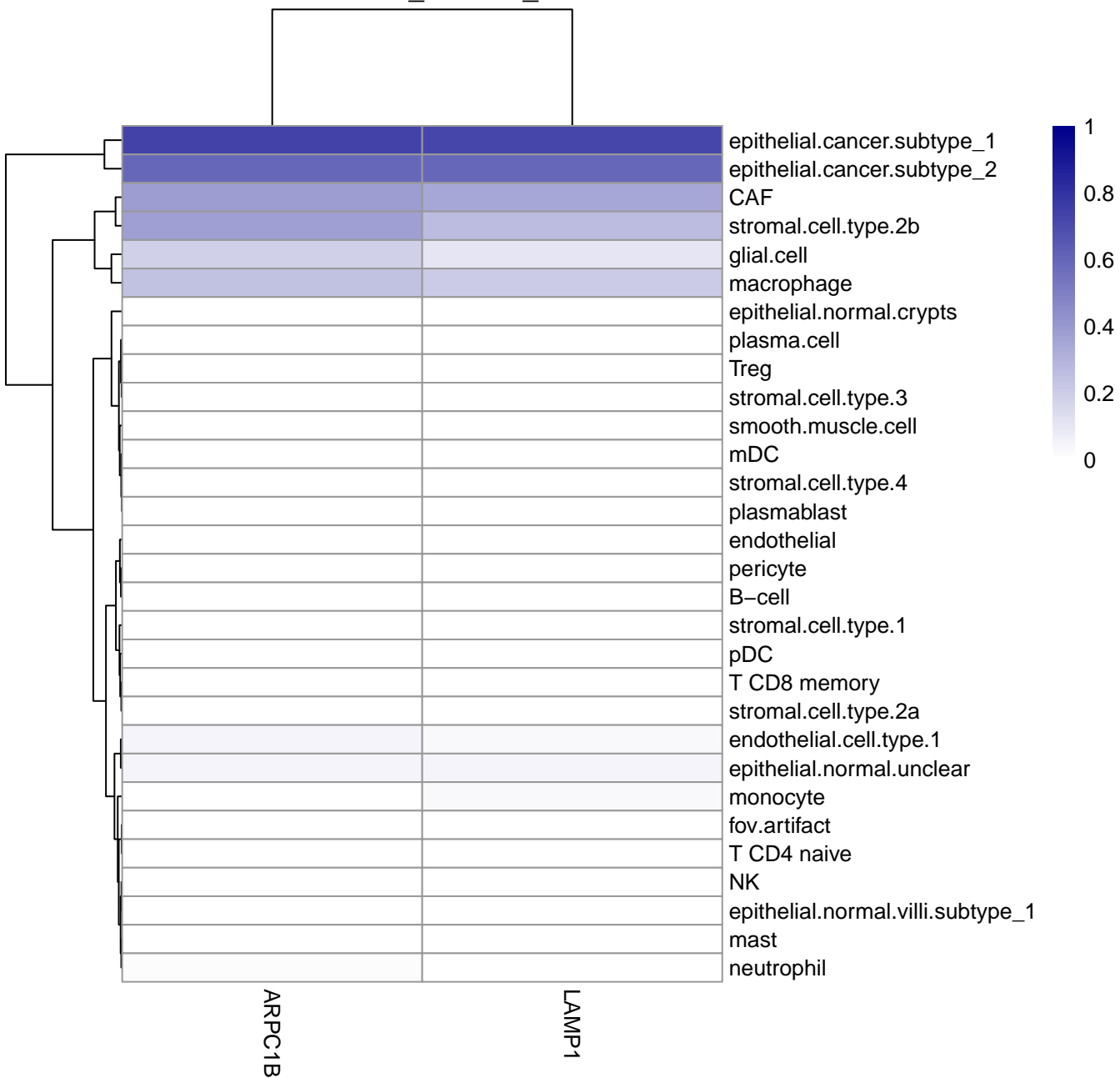
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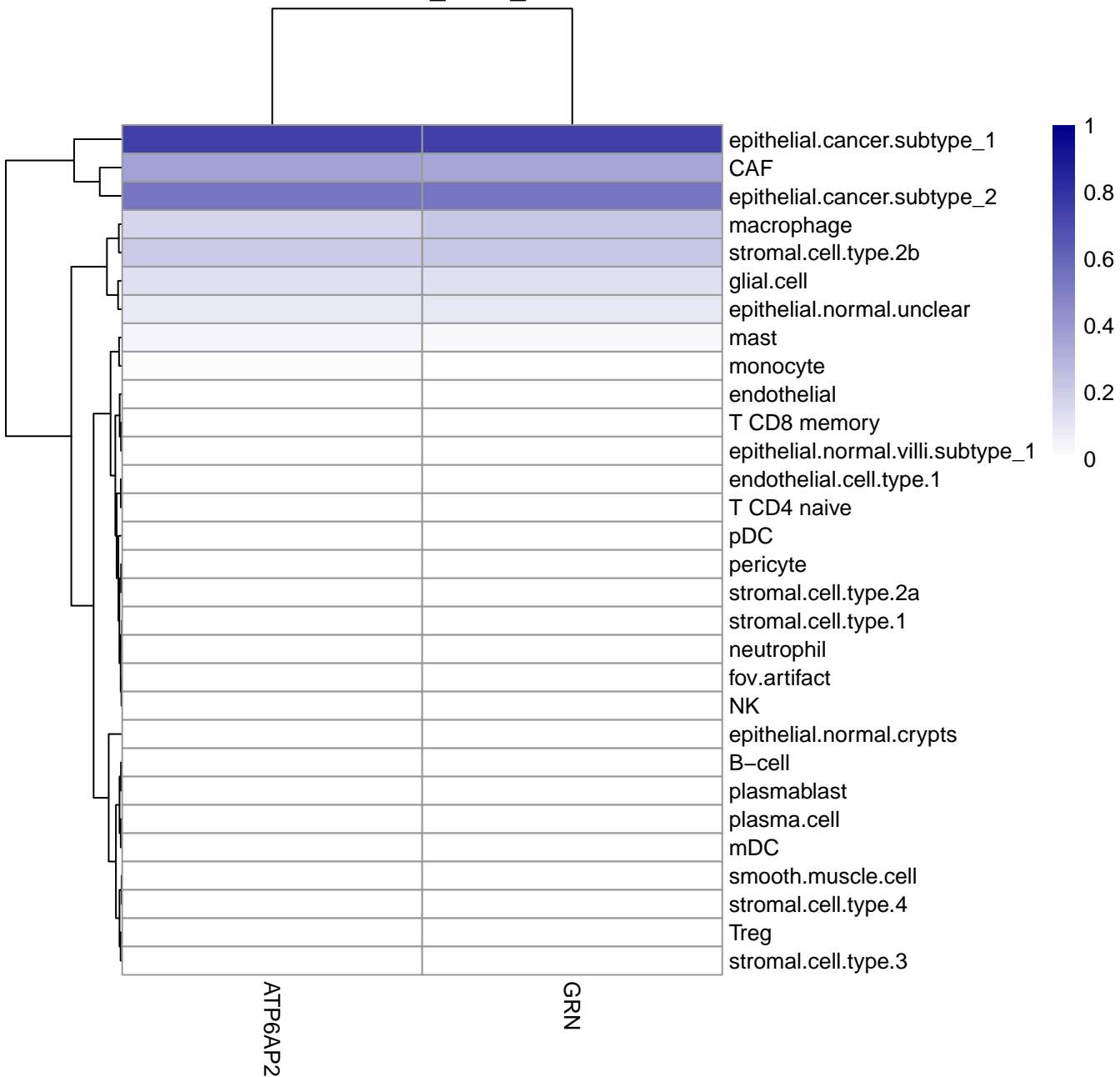
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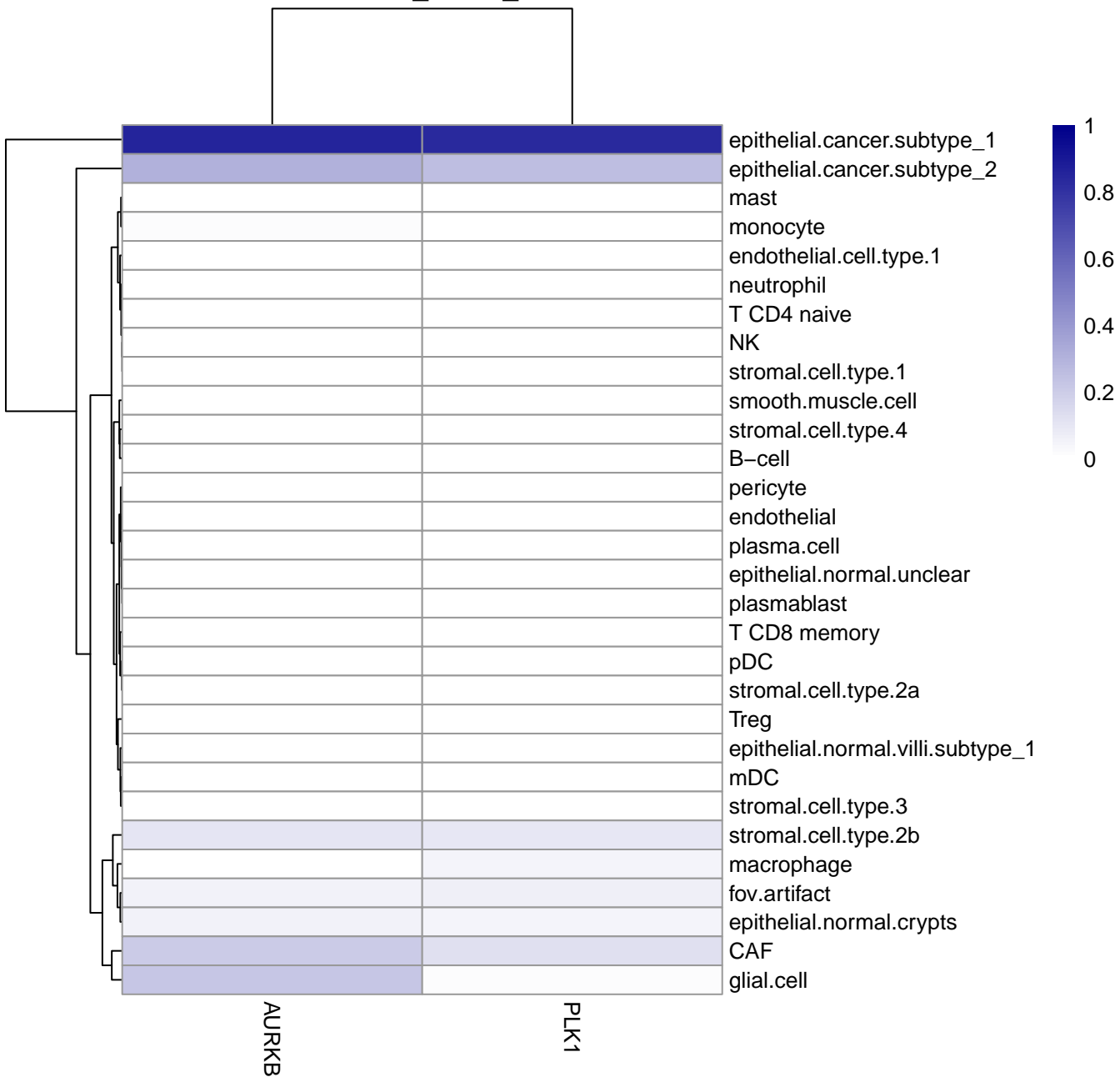
ARPC1B_LAMP1_2



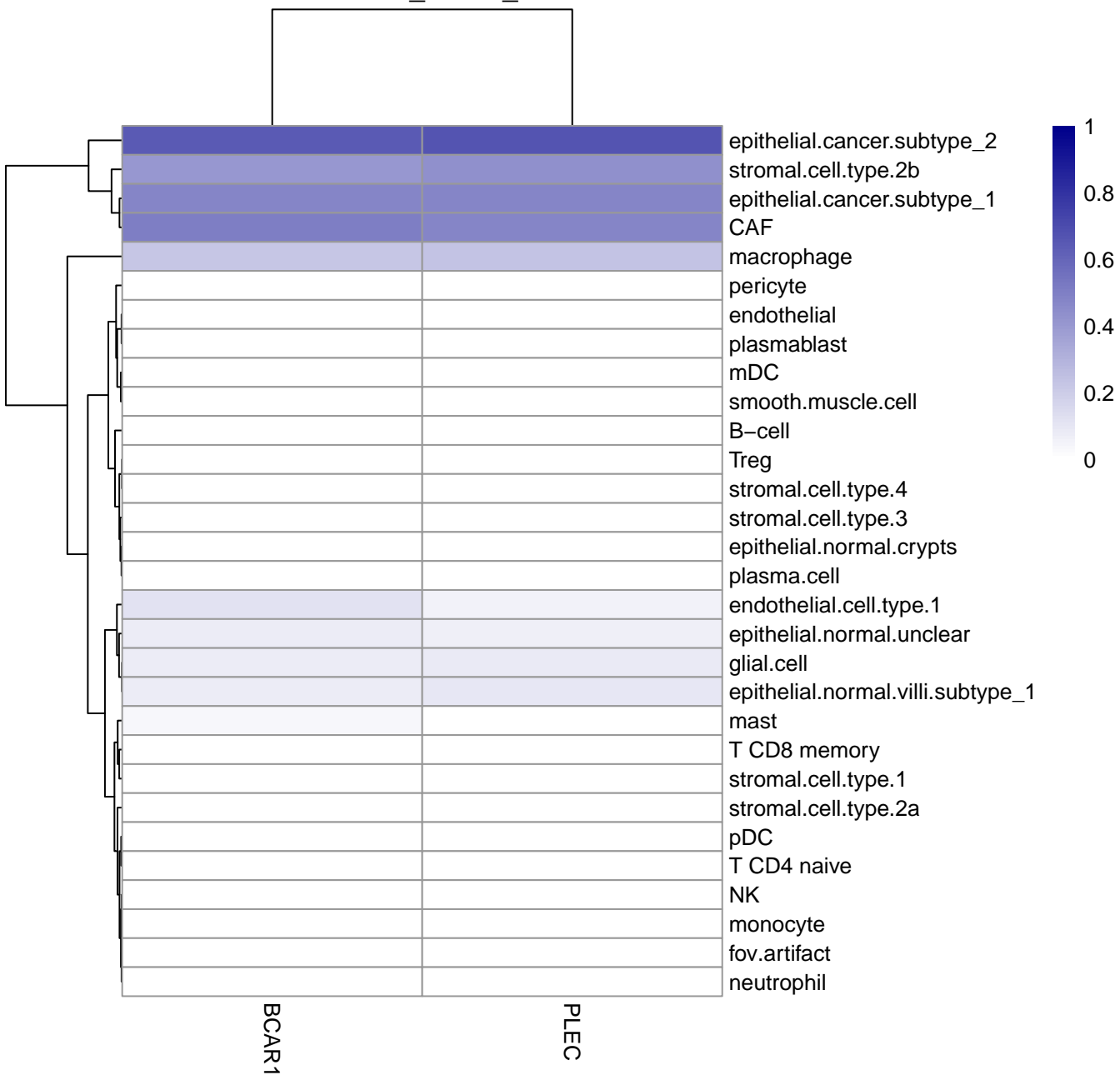
ATP6AP2_GRN_2



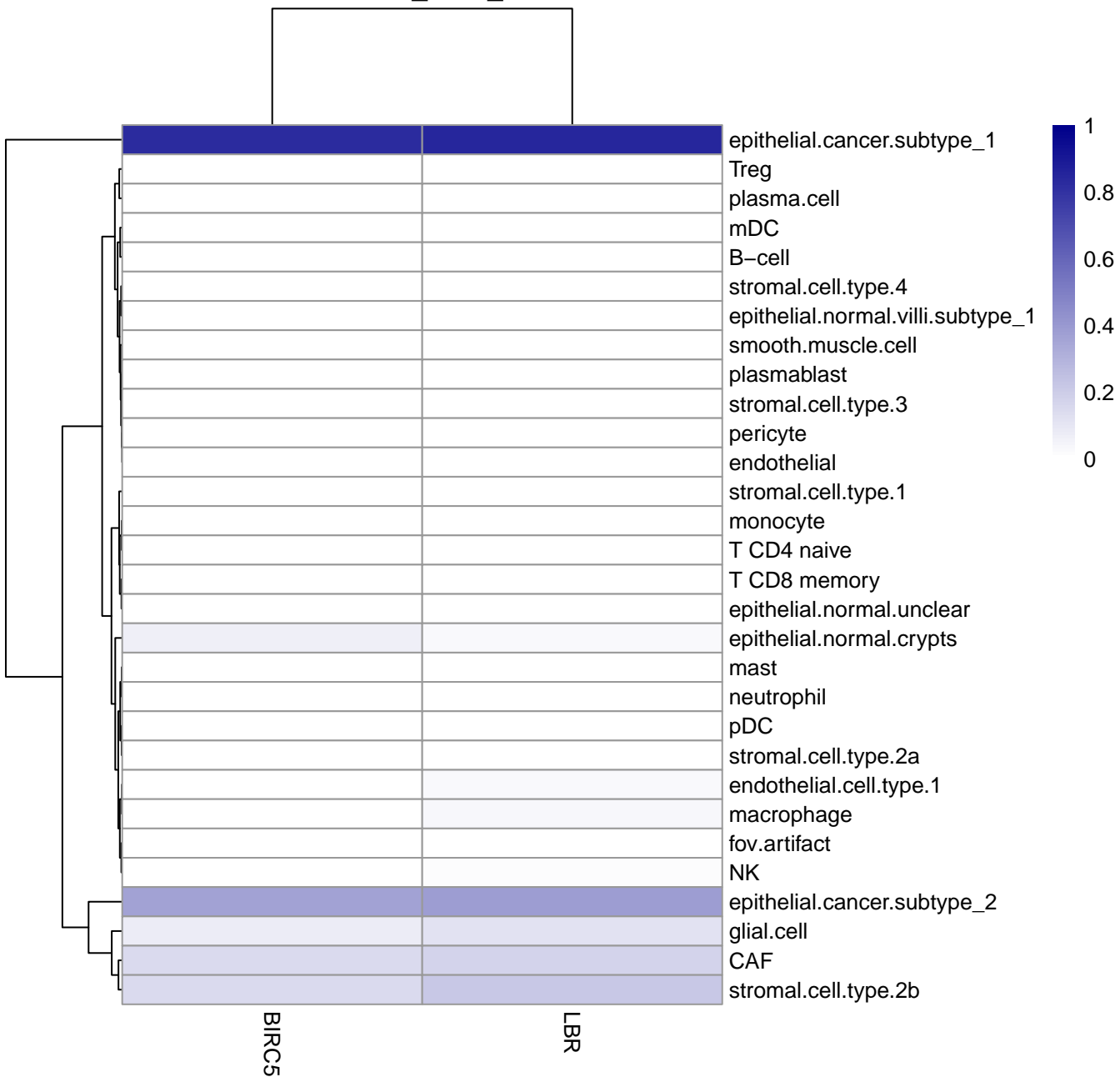
AURKB_PLK1_2



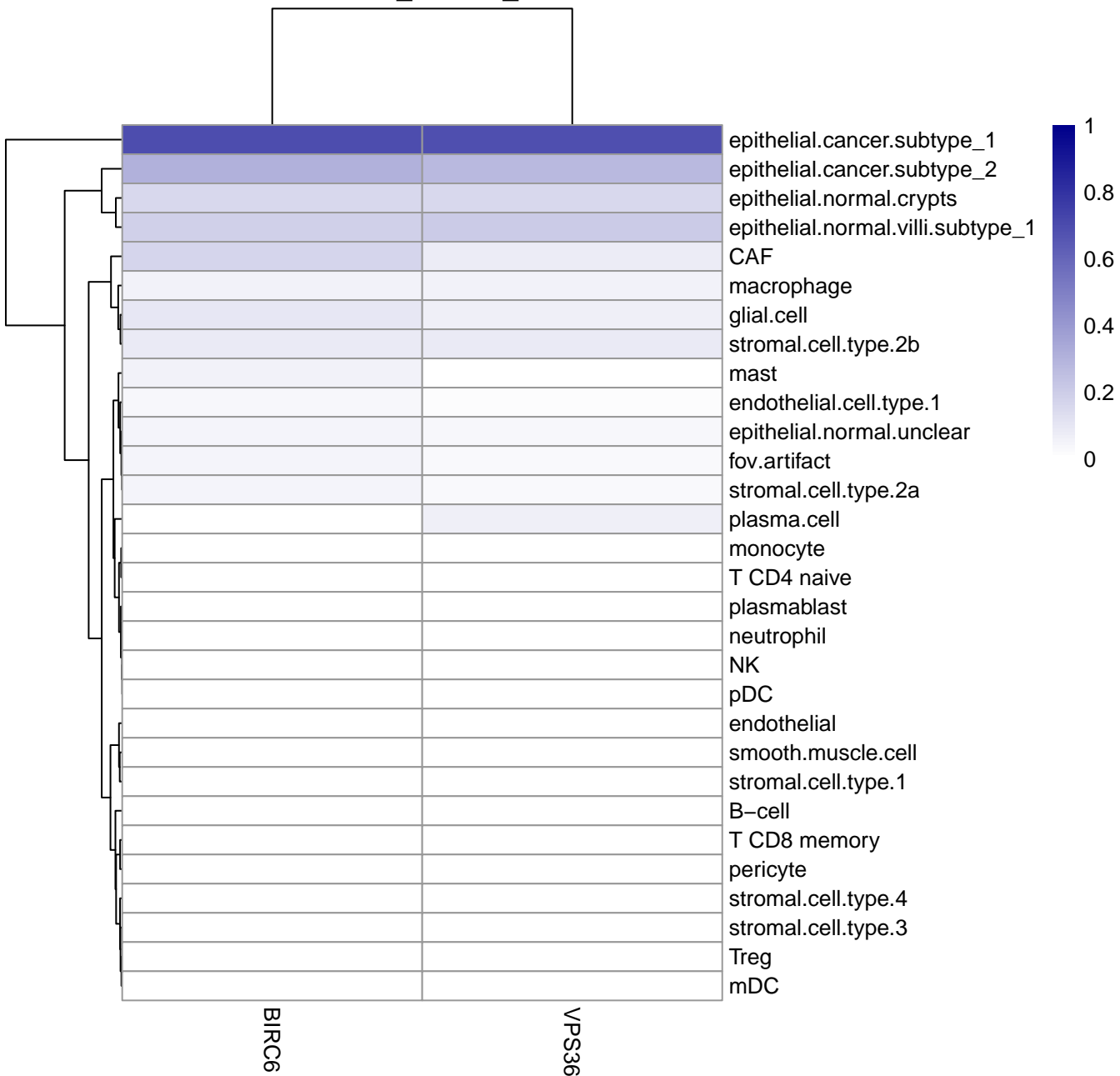
BCAR1_PLEC_2



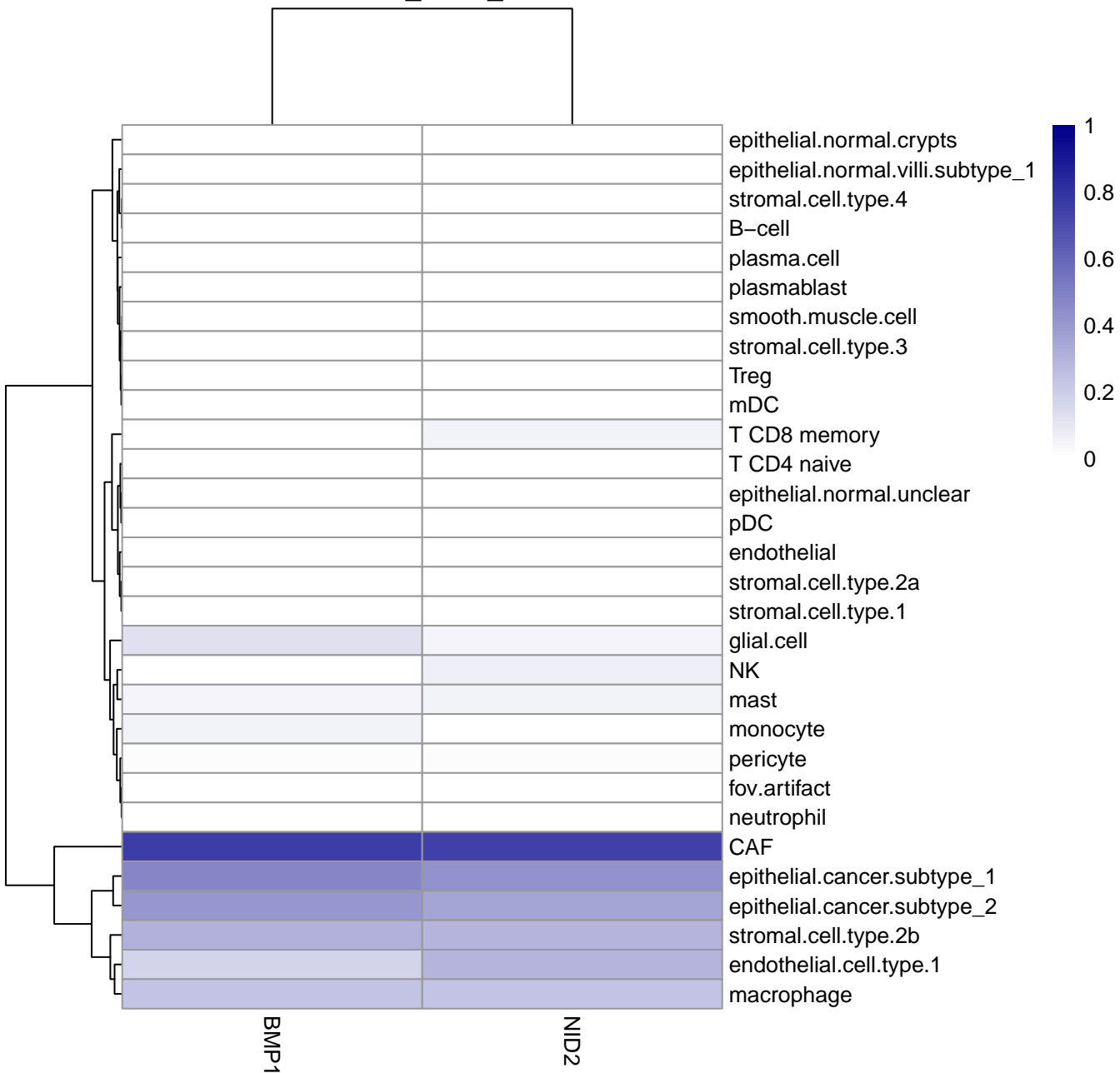
BIRC5_LBR_2



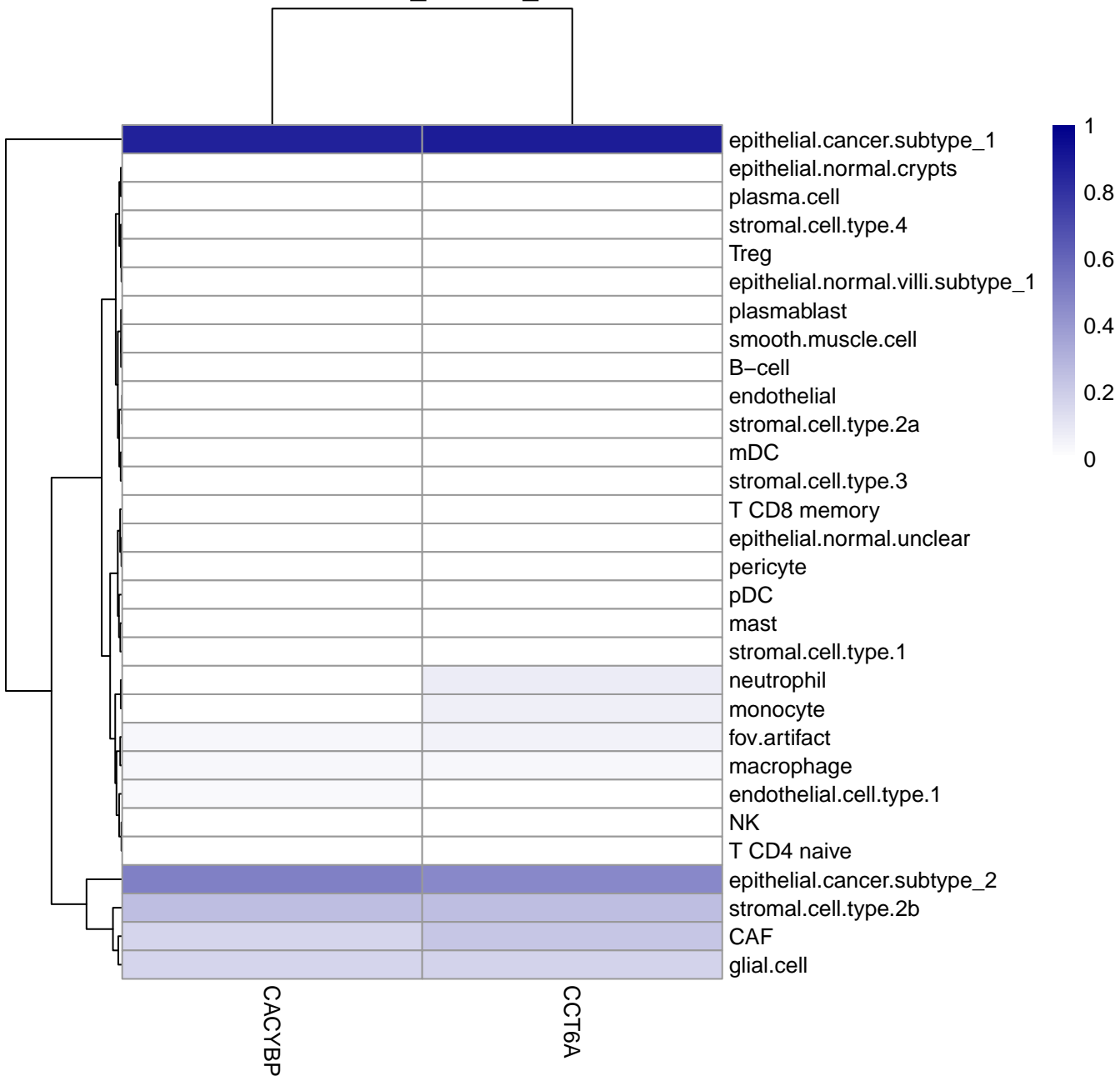
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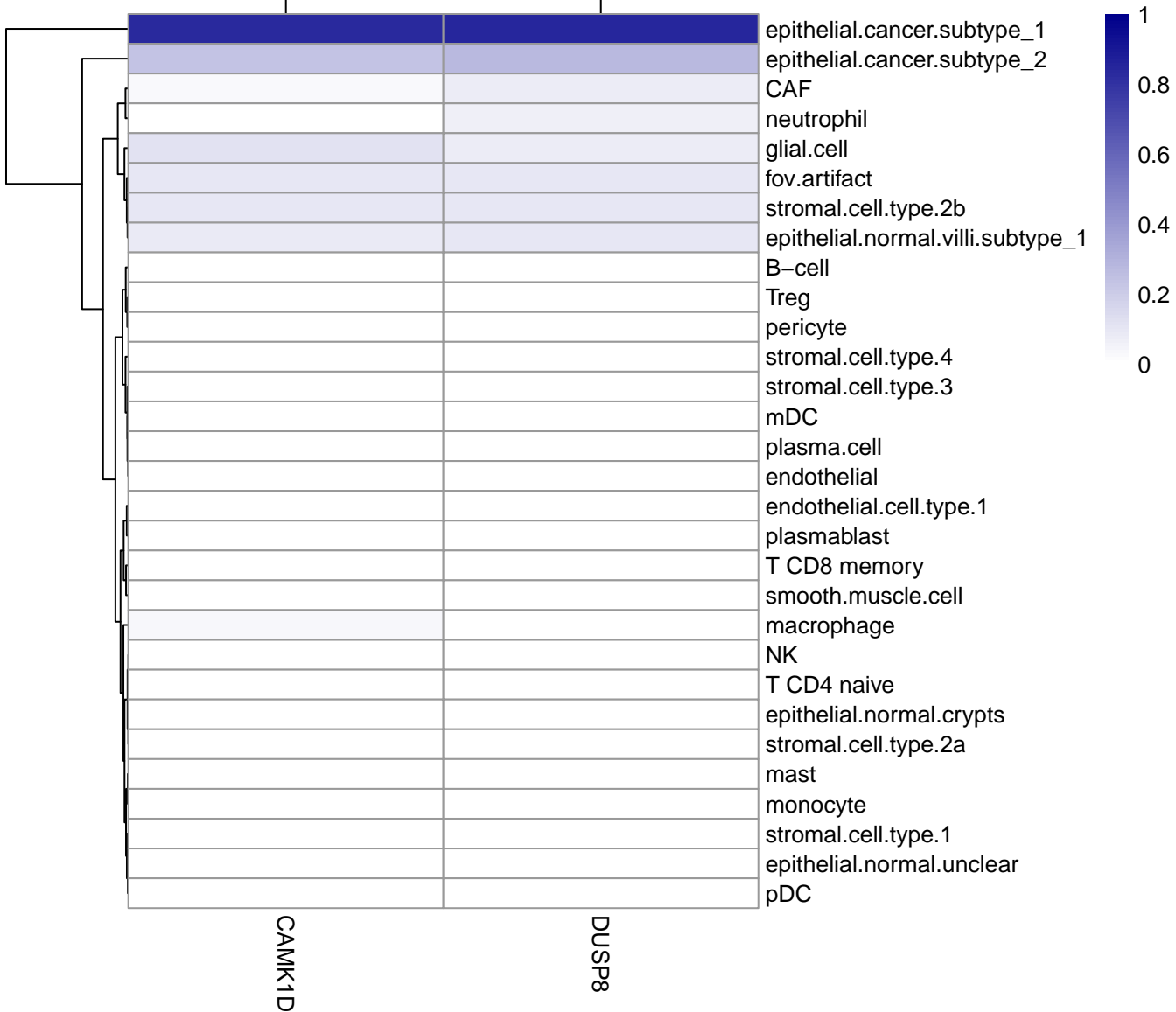
BMP1_NID2_2



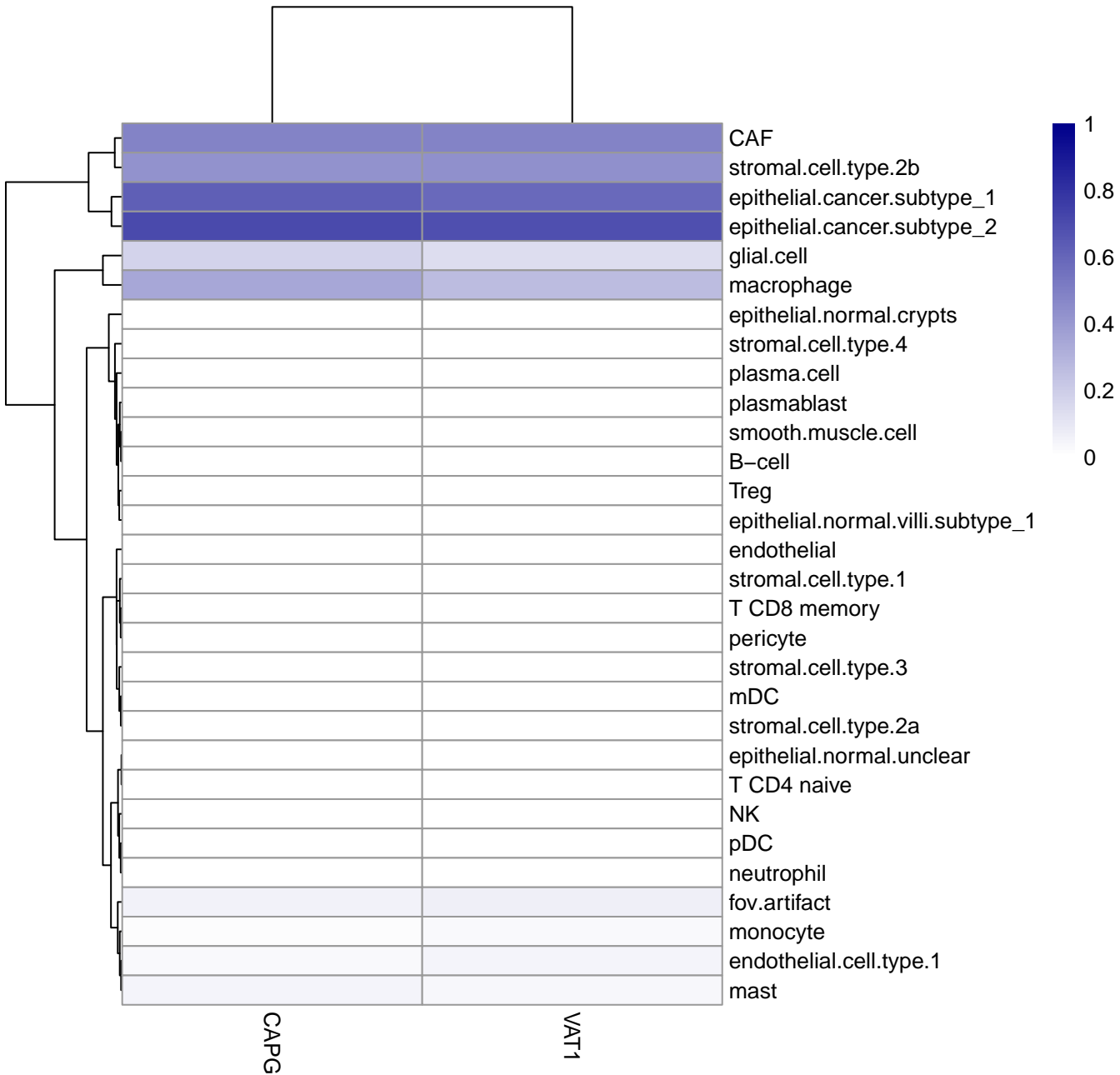
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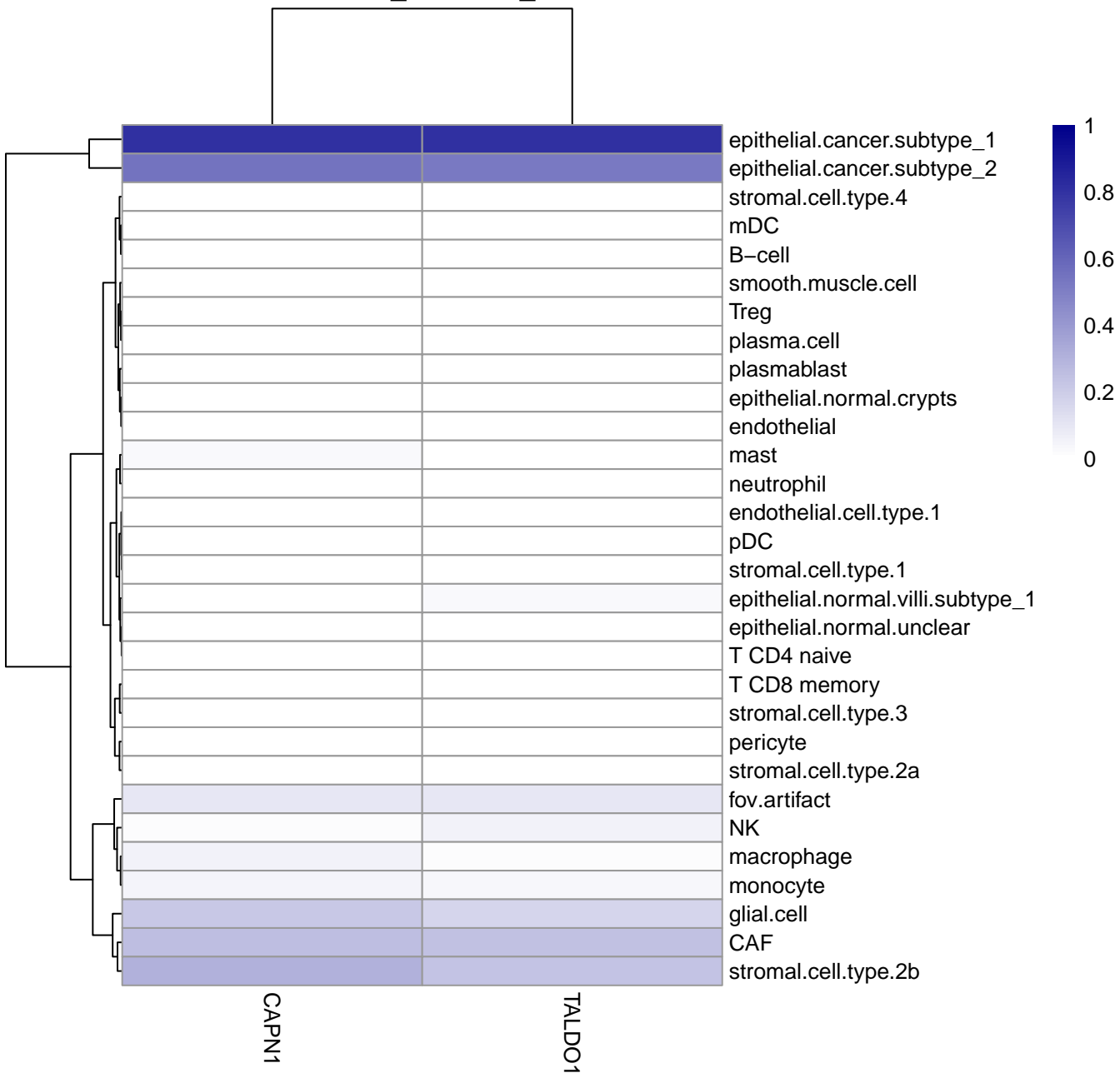
CAMK1D_DUSP8_2



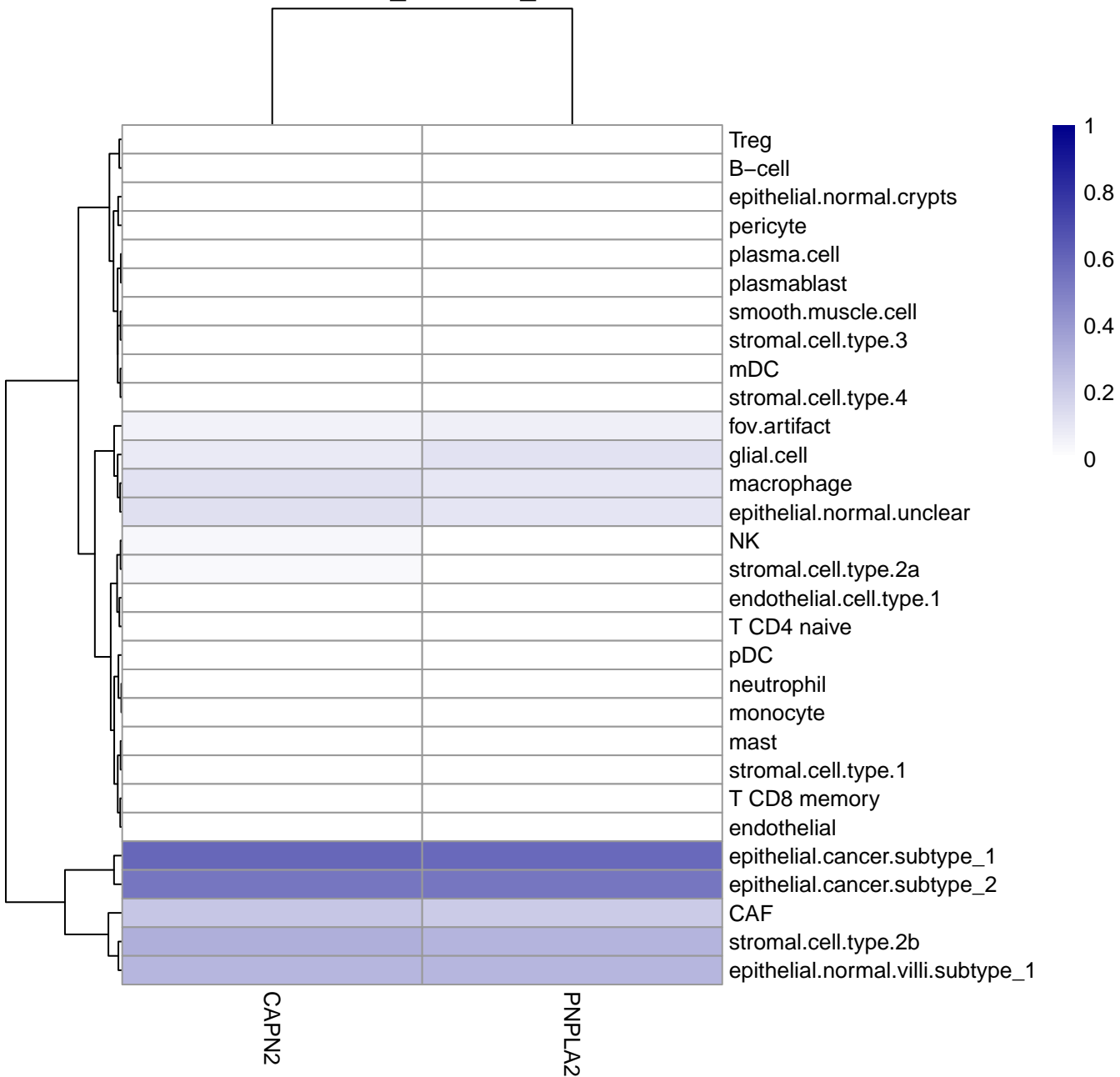
CAPG_VAT1_2



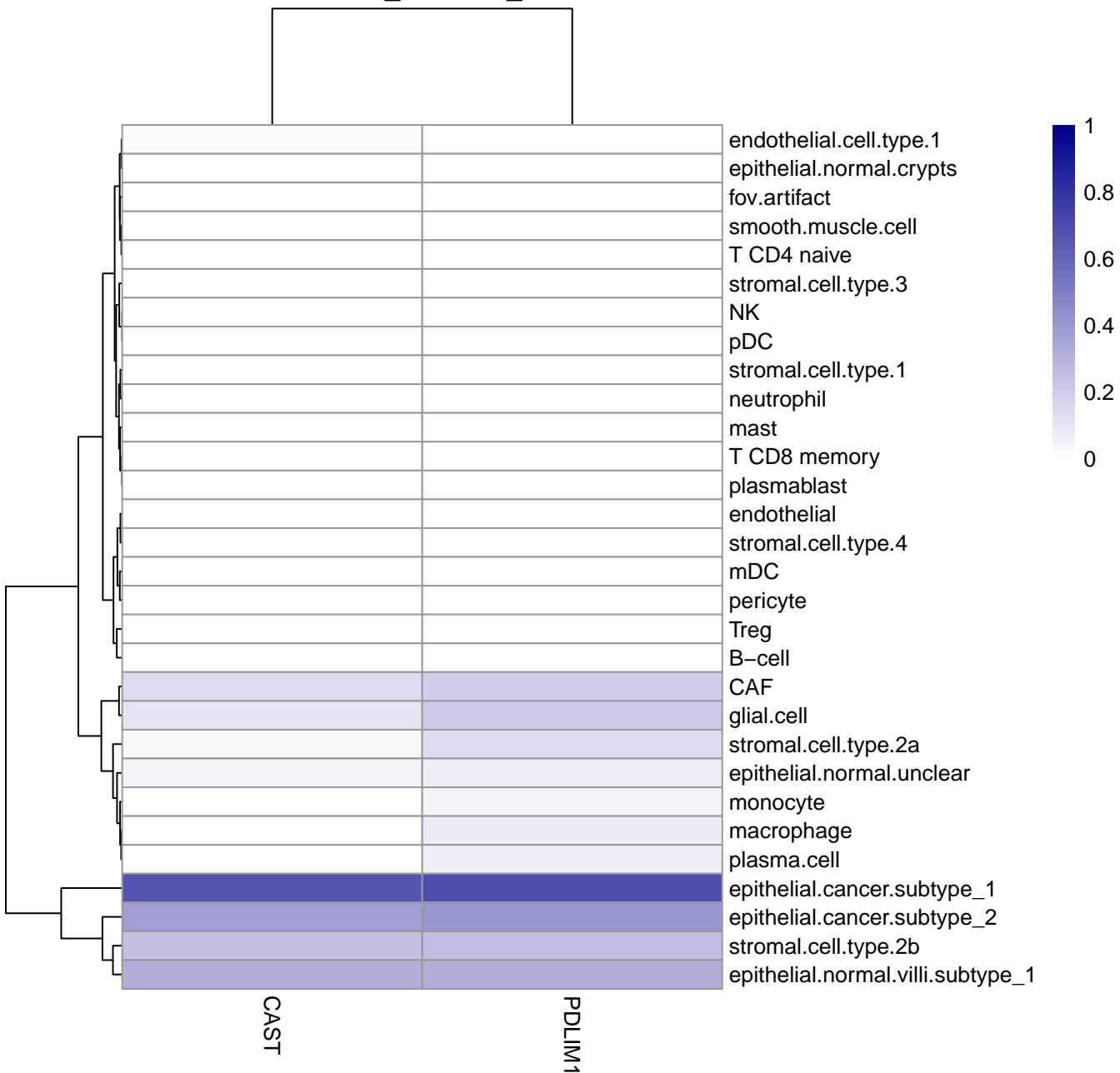
CAPN1_TALDO1_2



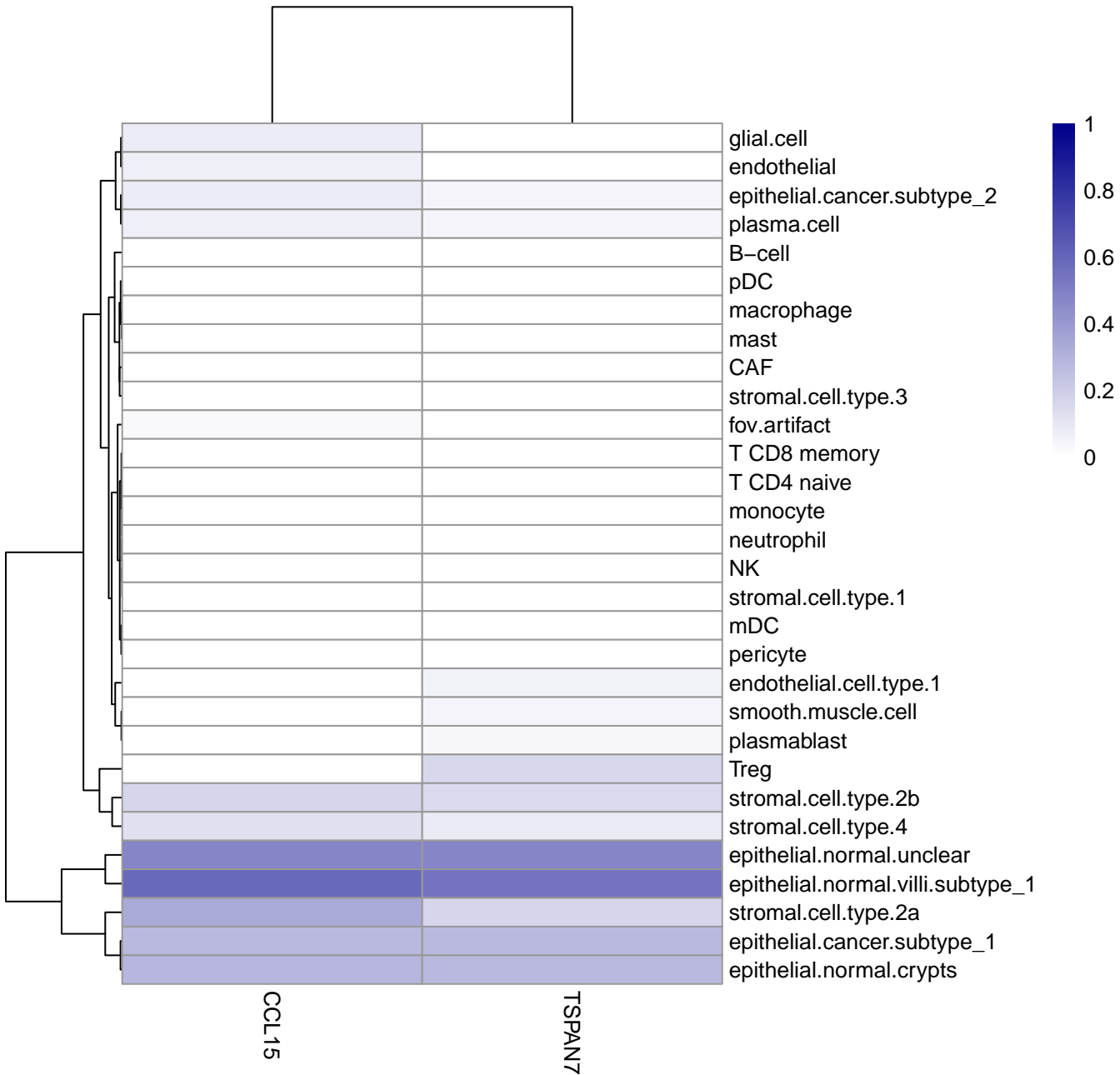
CAPN2_PNPLA2_2



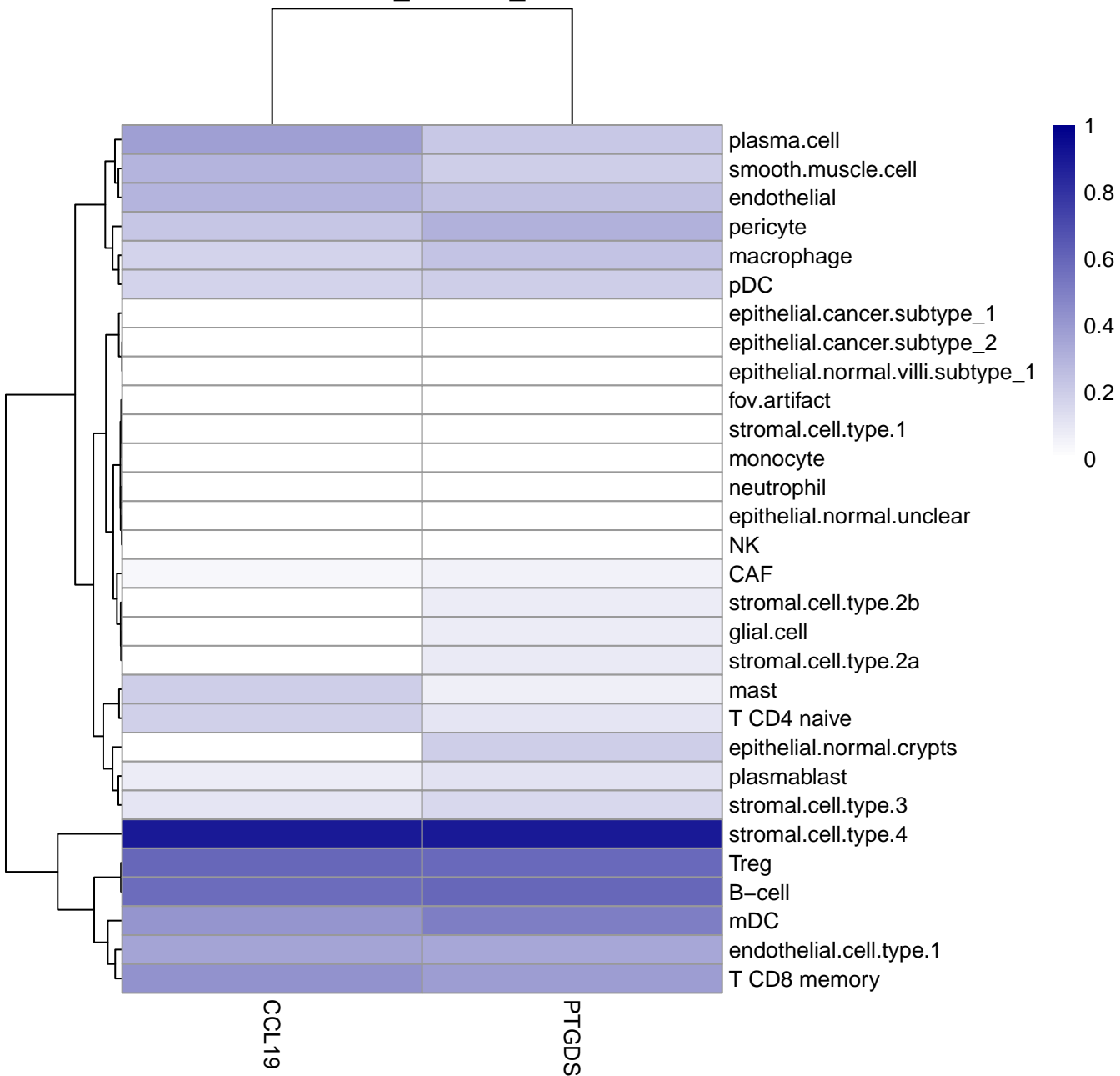
CAST_PDLIM1_2



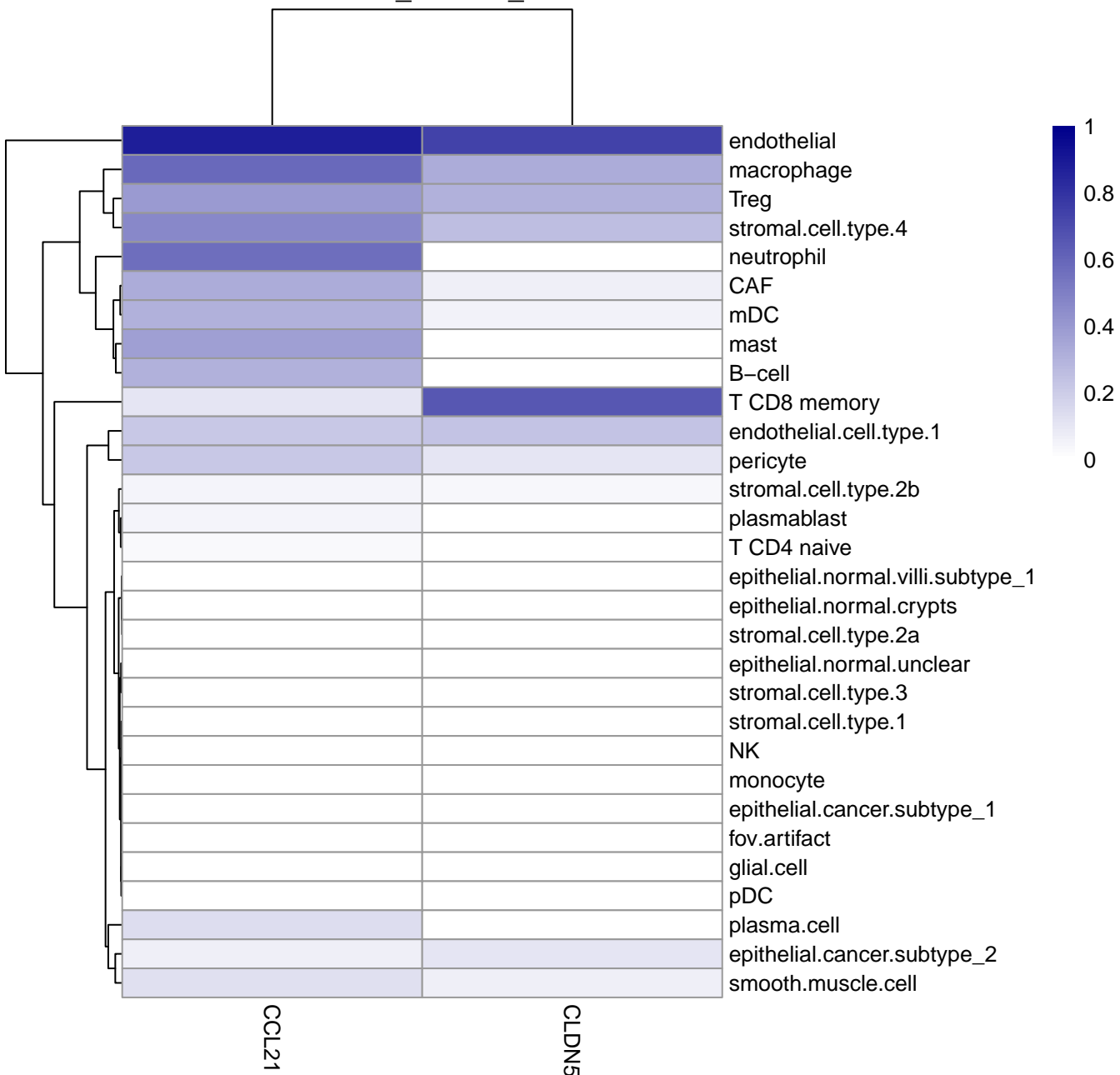
CCL15_TSPAN7_2



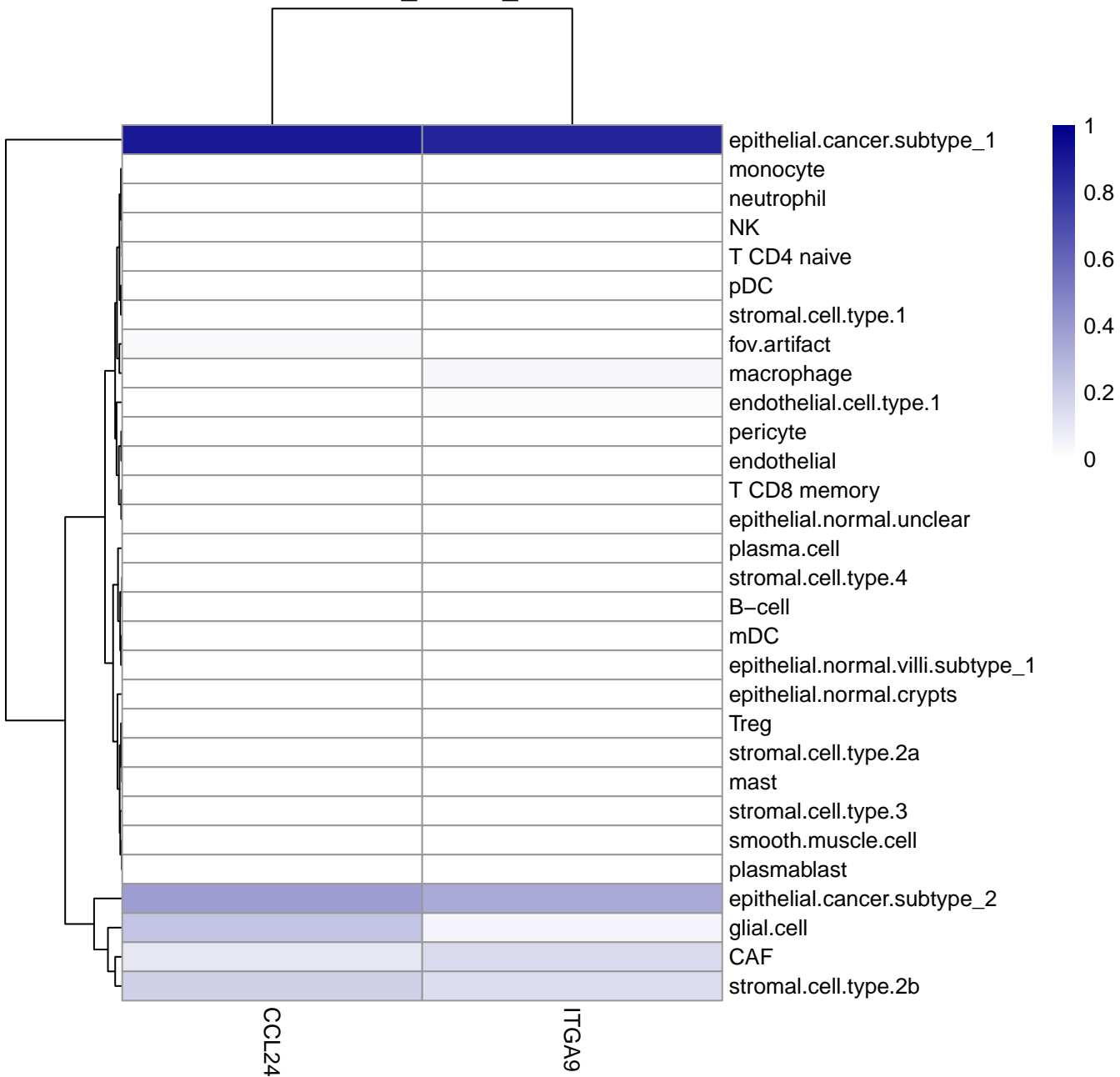
CCL19_PTGDS_2



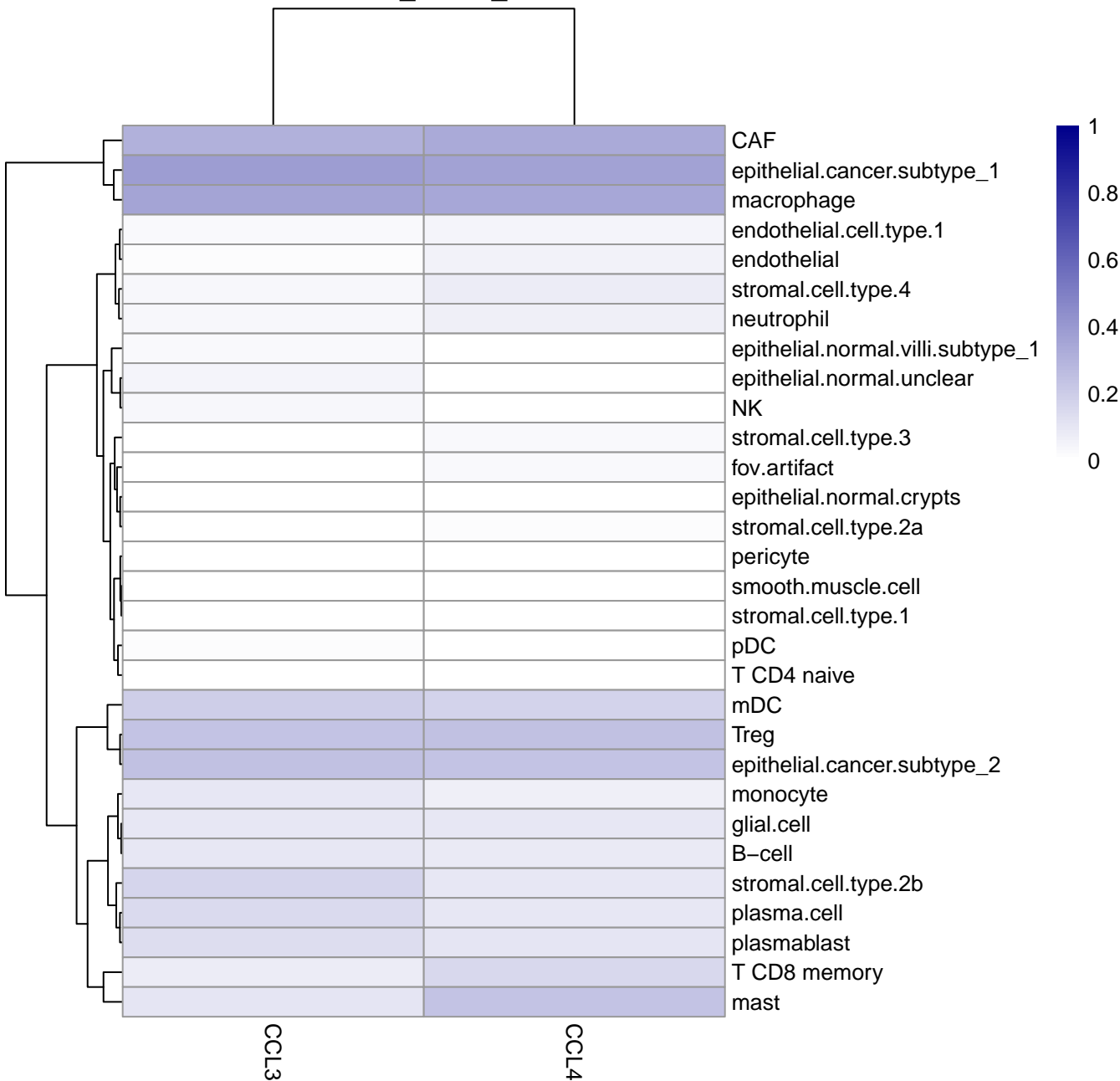
CCL21_CLDN5_2



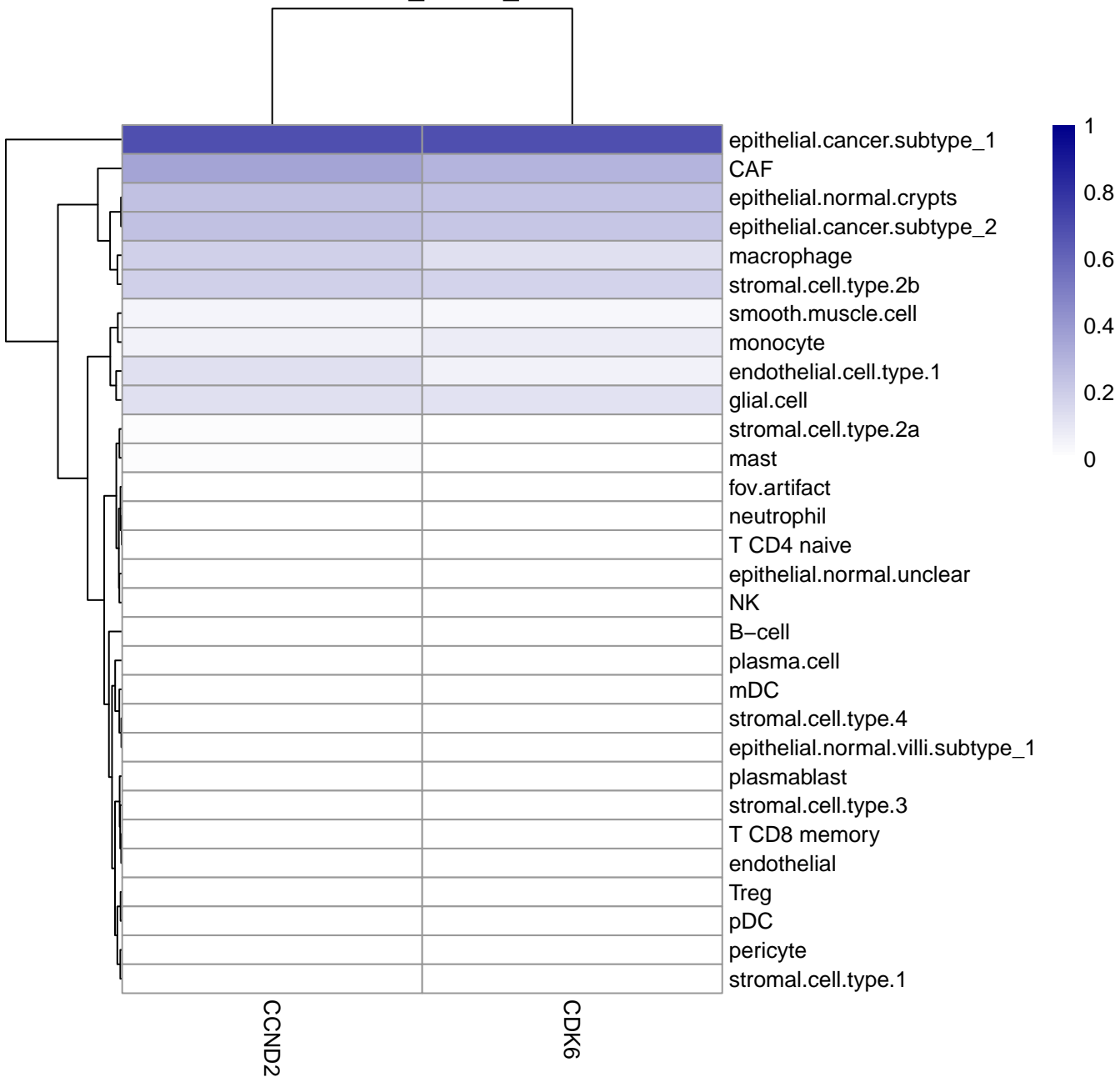
CCL24_ITGA9_2



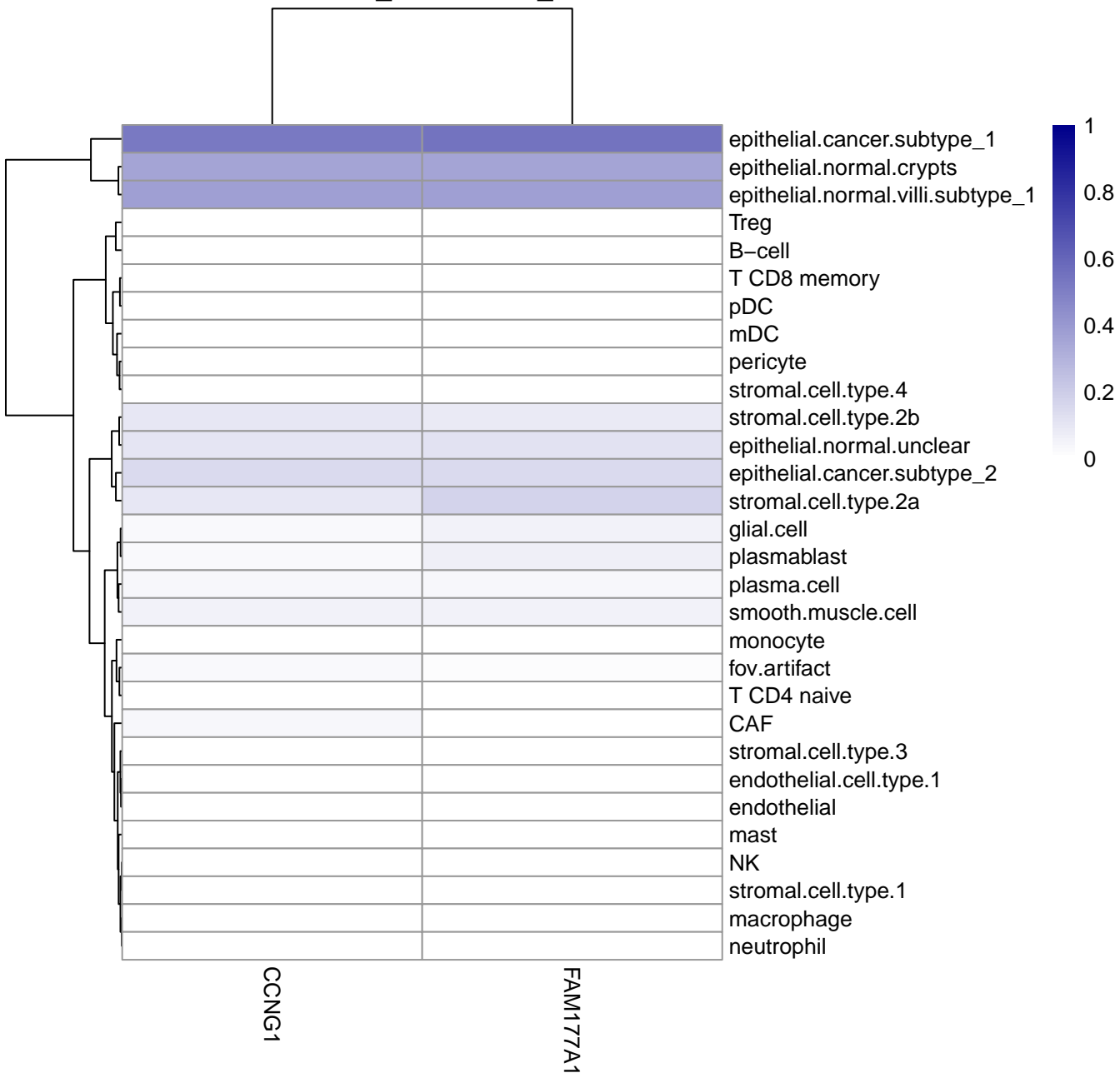
CCL3_CCL4_2



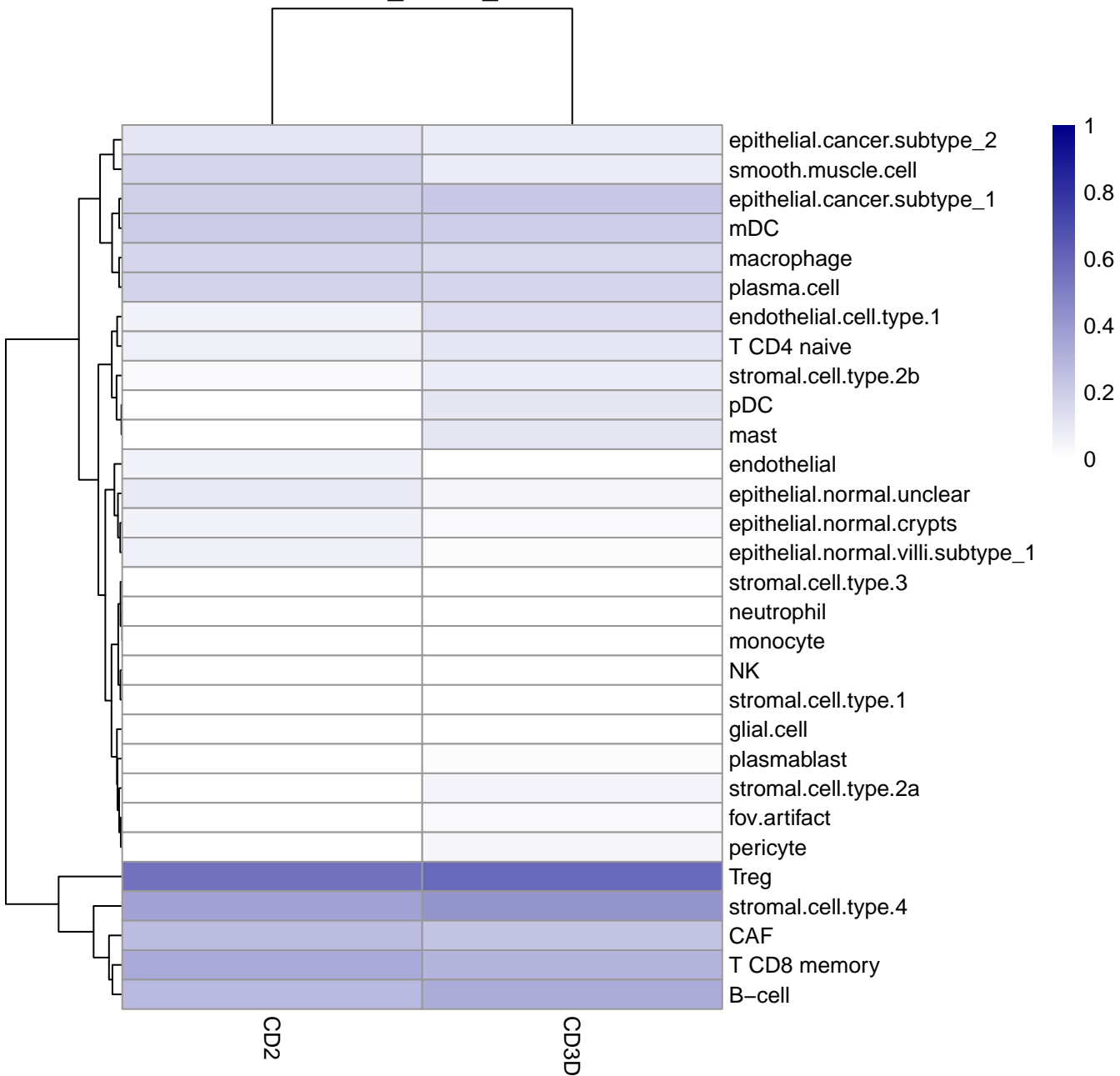
CCND2_CDK6_2



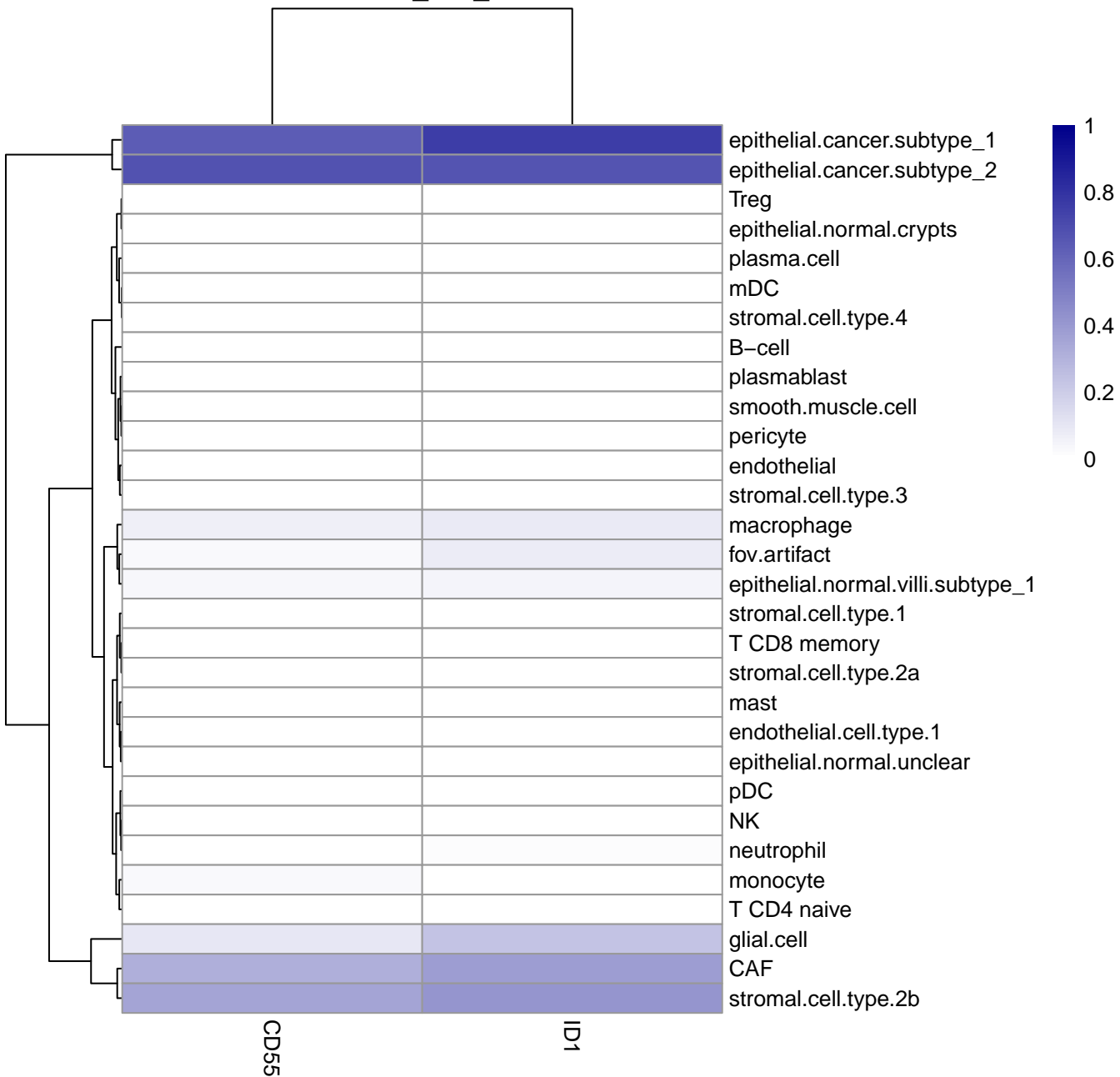
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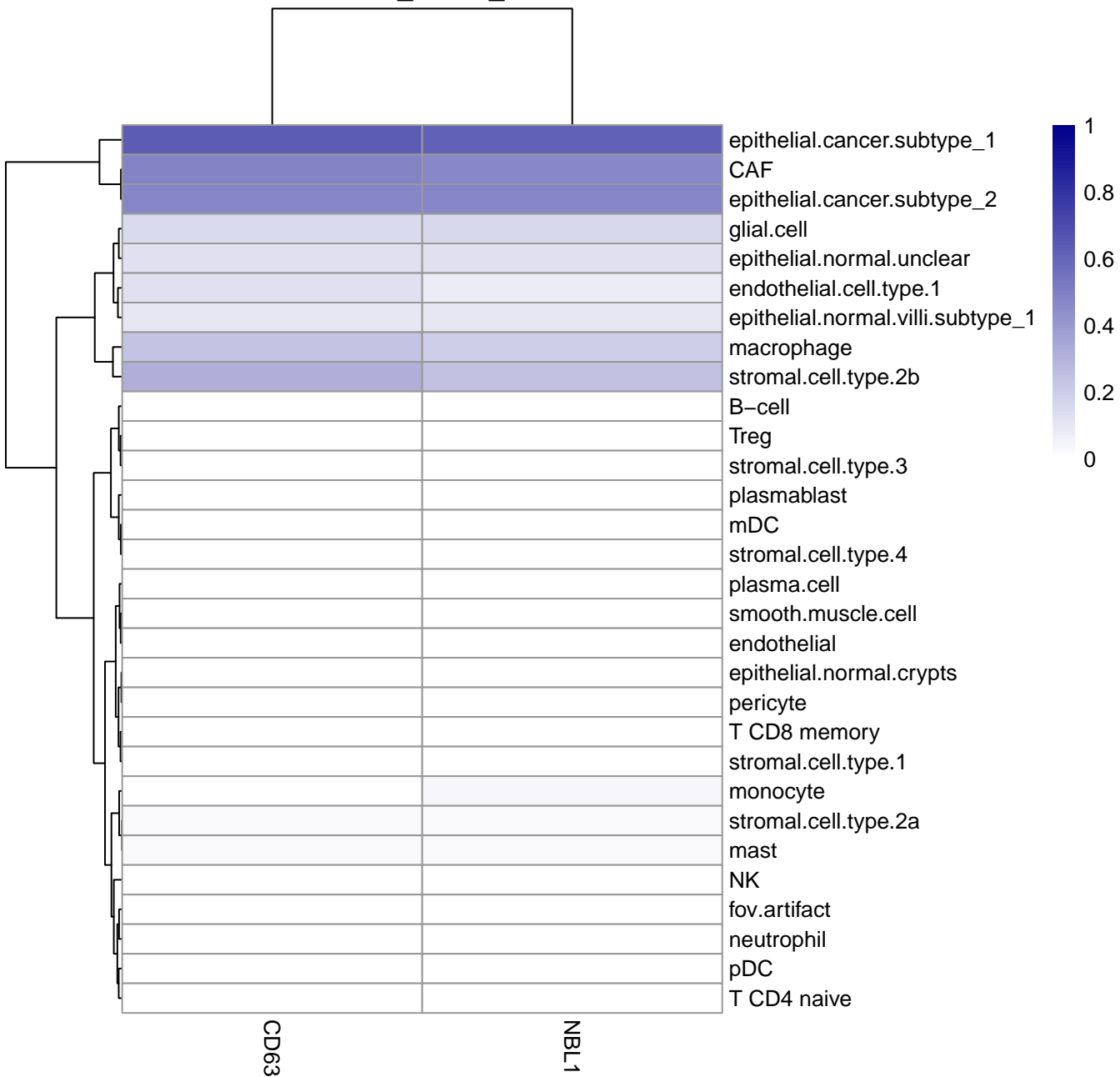
CD2_CD3D_2



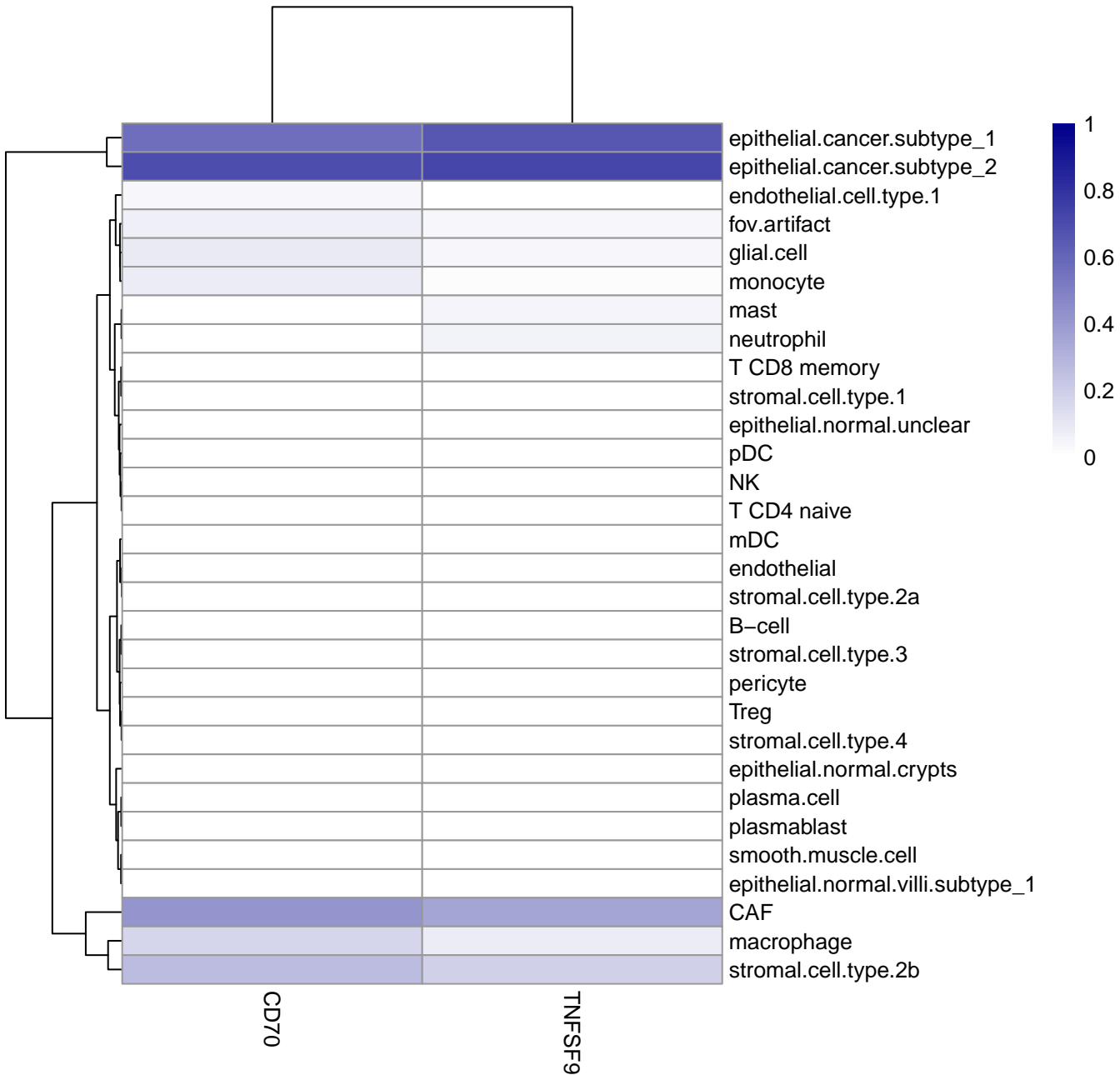
CD55_ID1_2



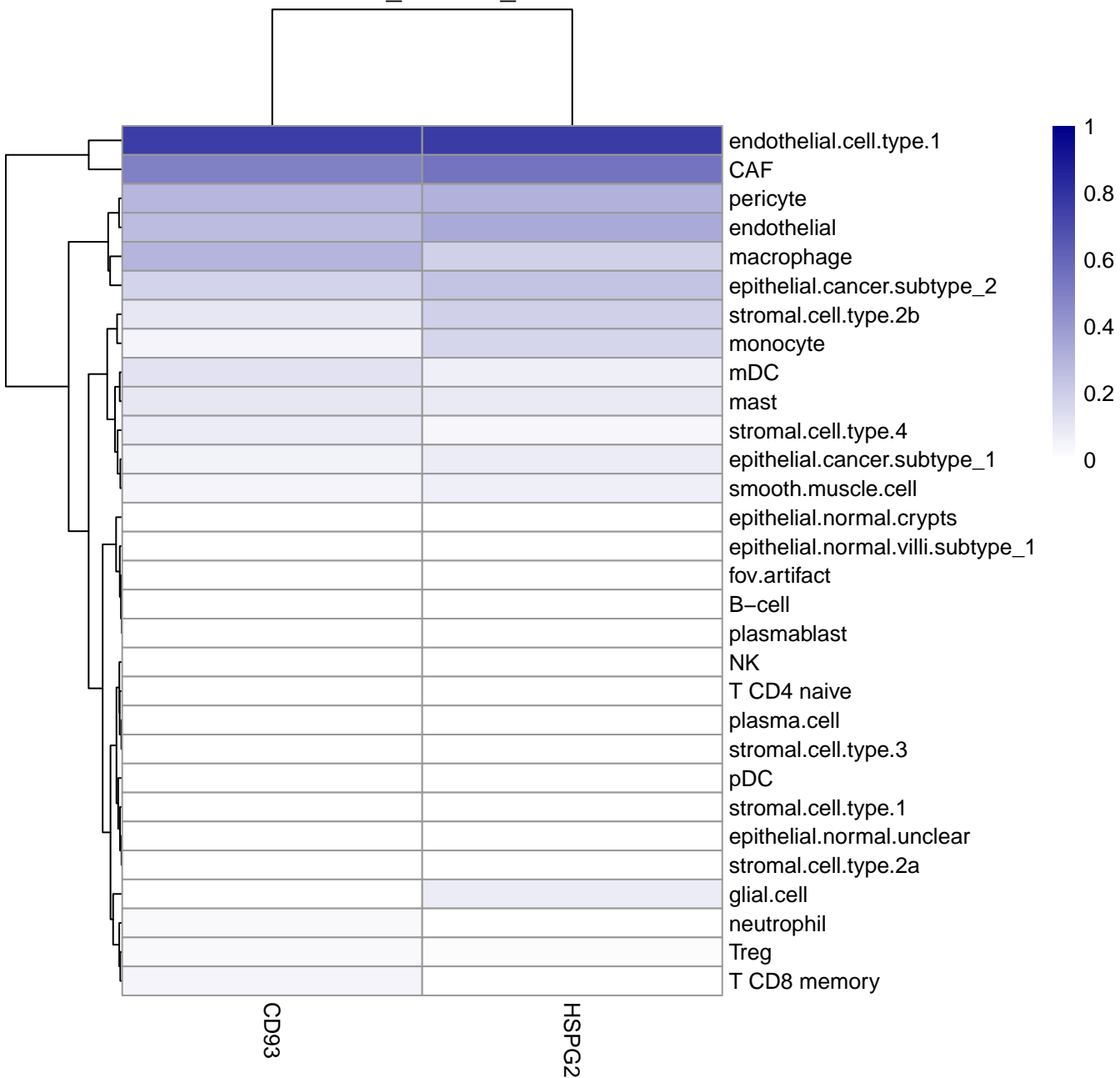
CD63_NBL1_2



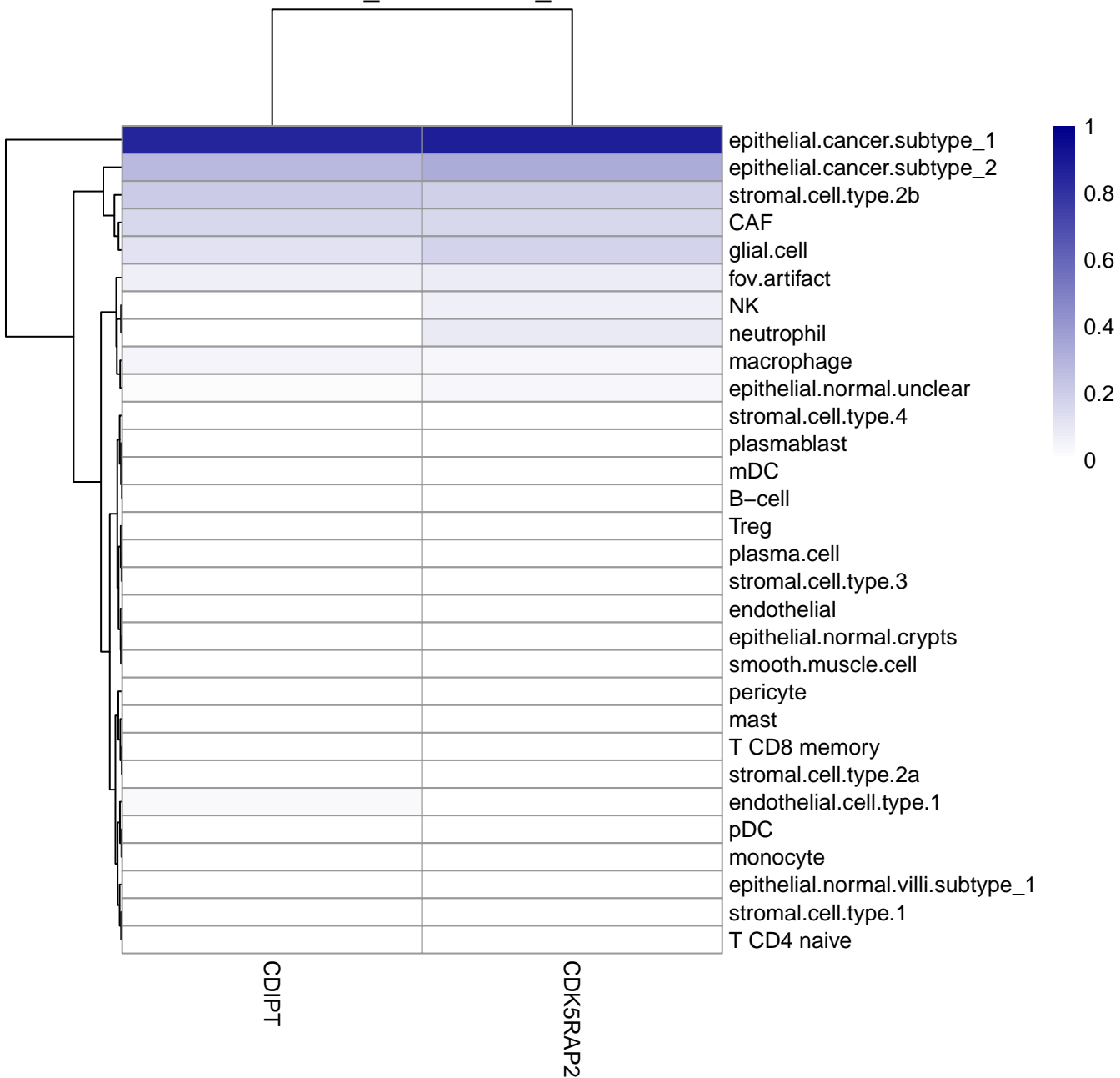
CD70_TNFSF9_2



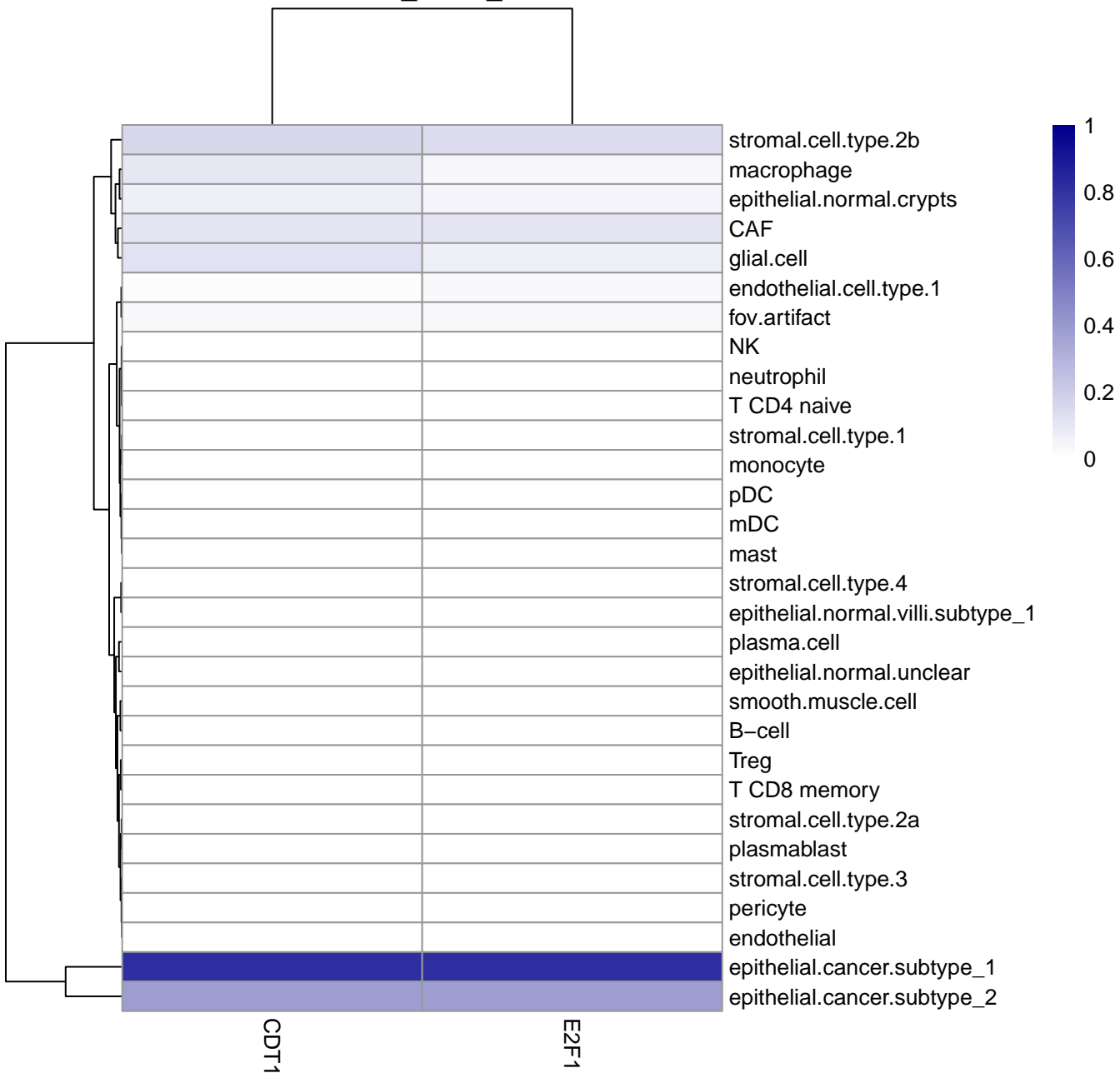
CD93_HSPG2_2



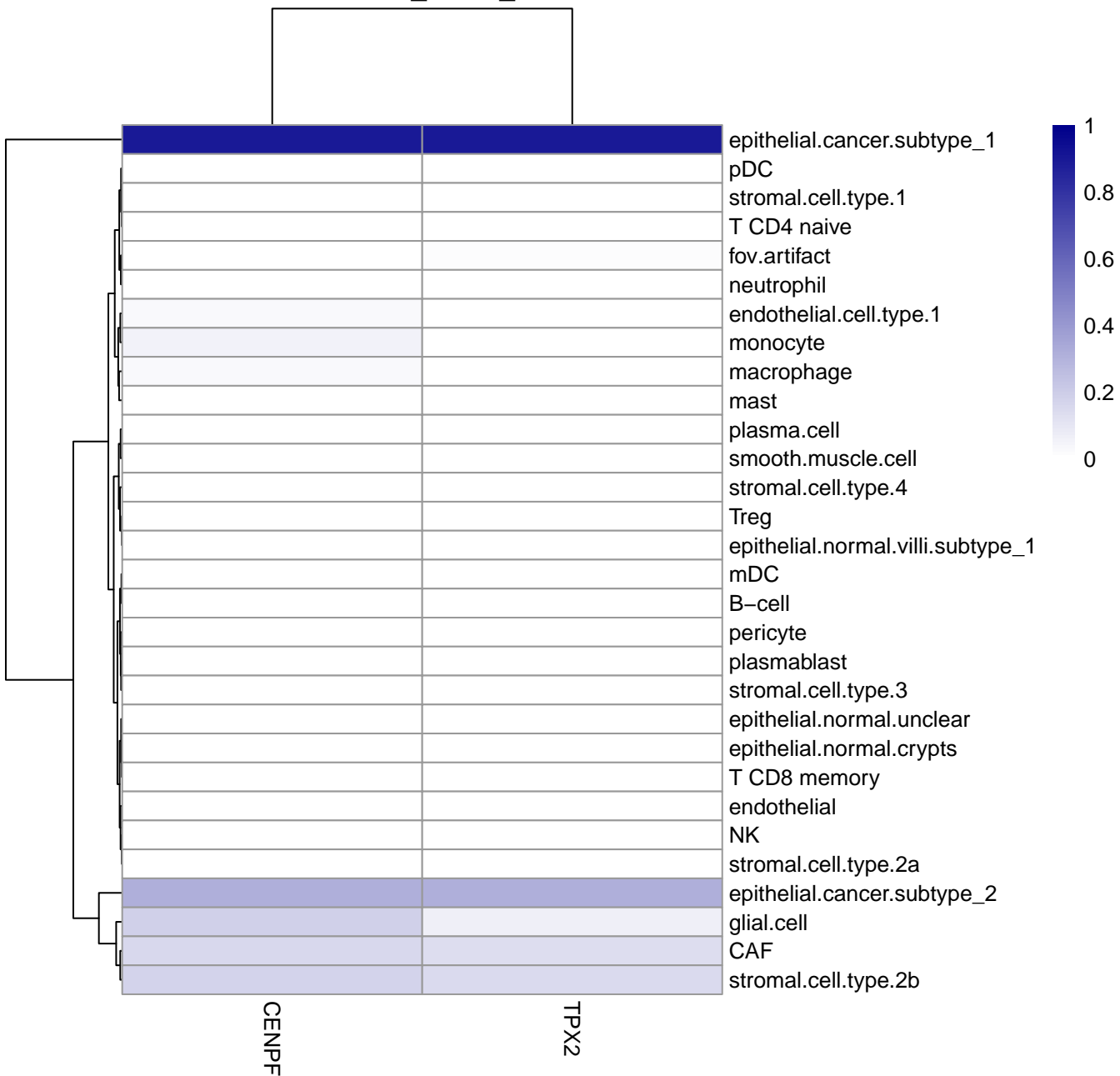
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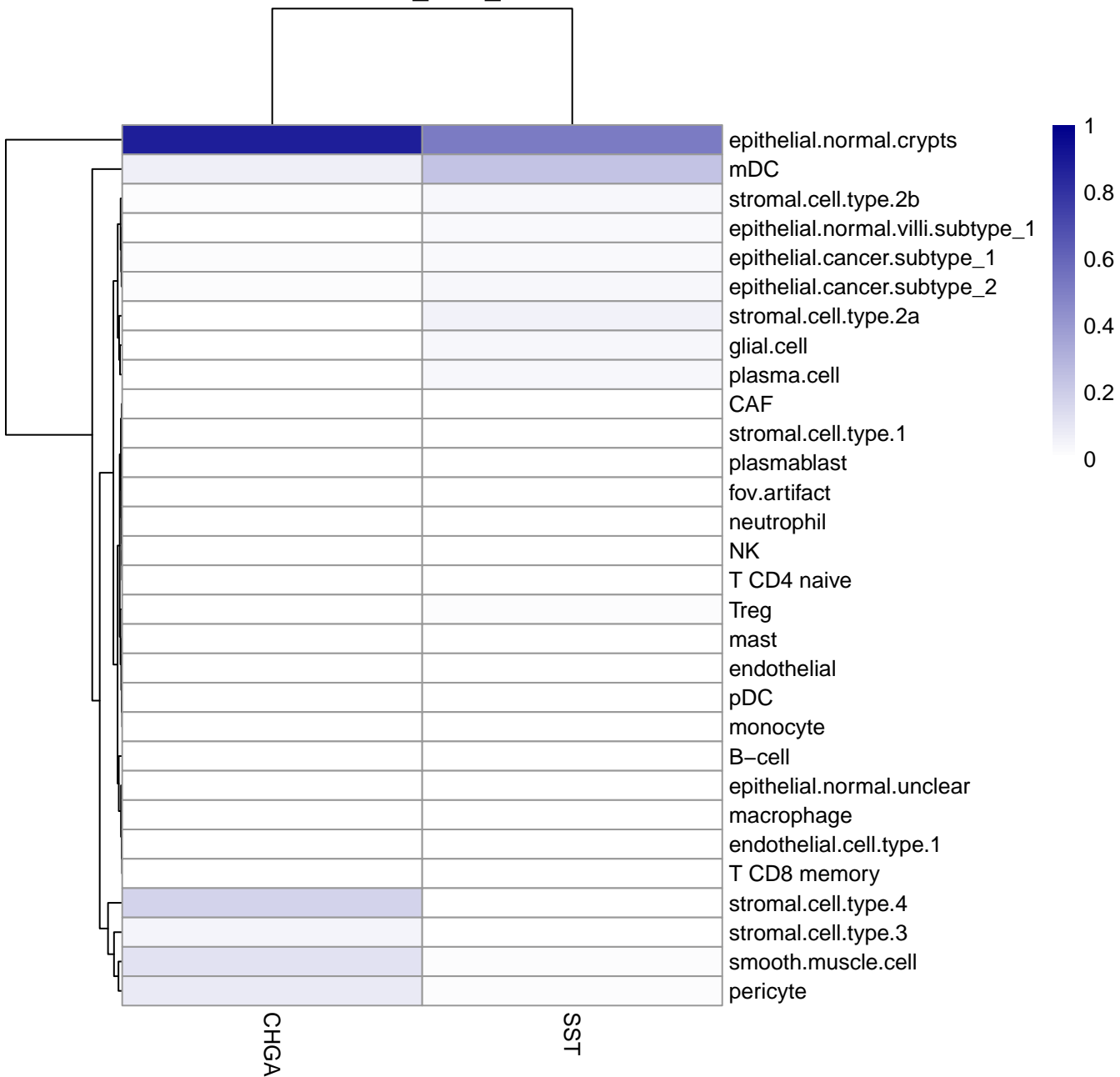
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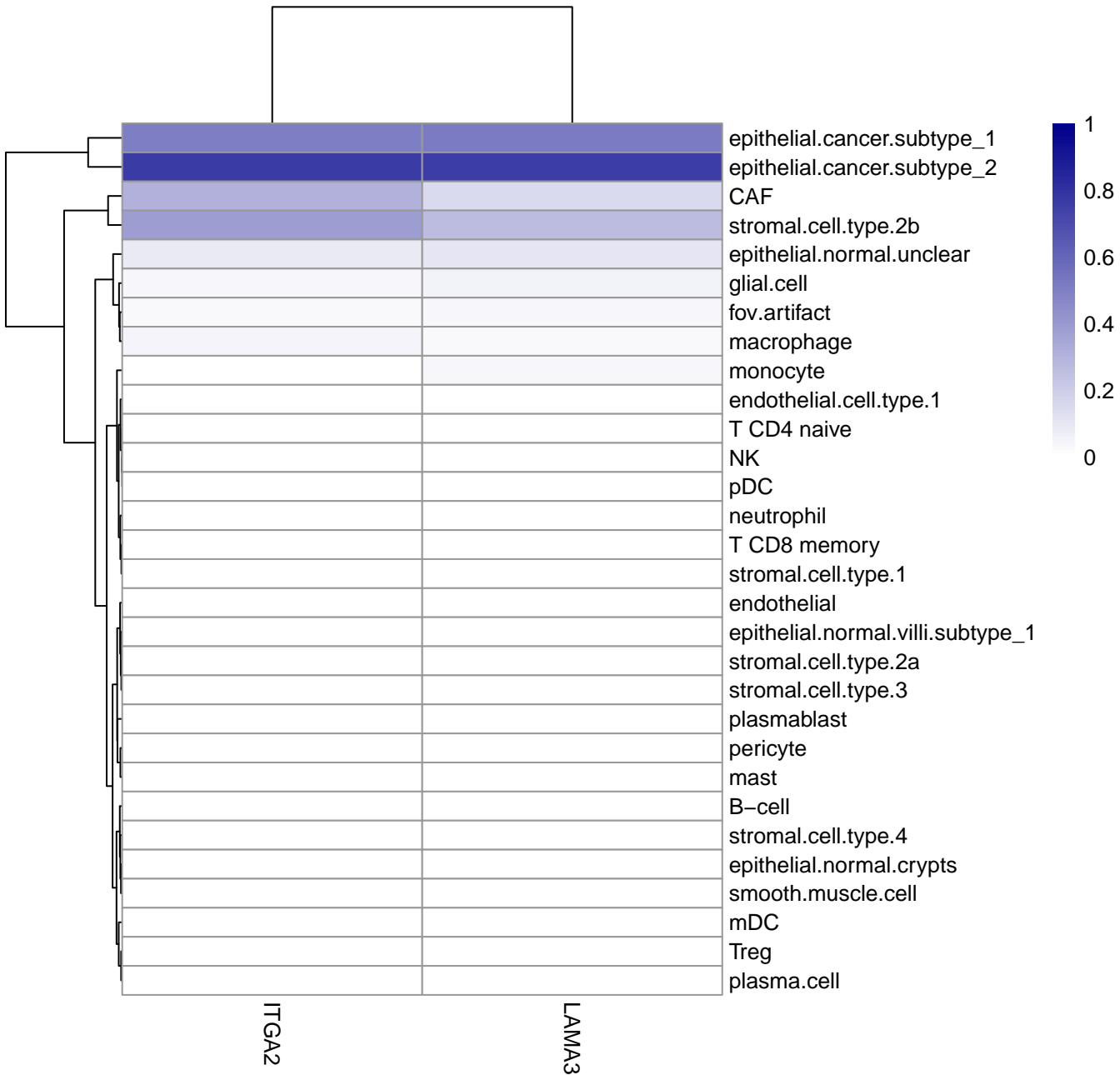
CENPF_TPX2_2



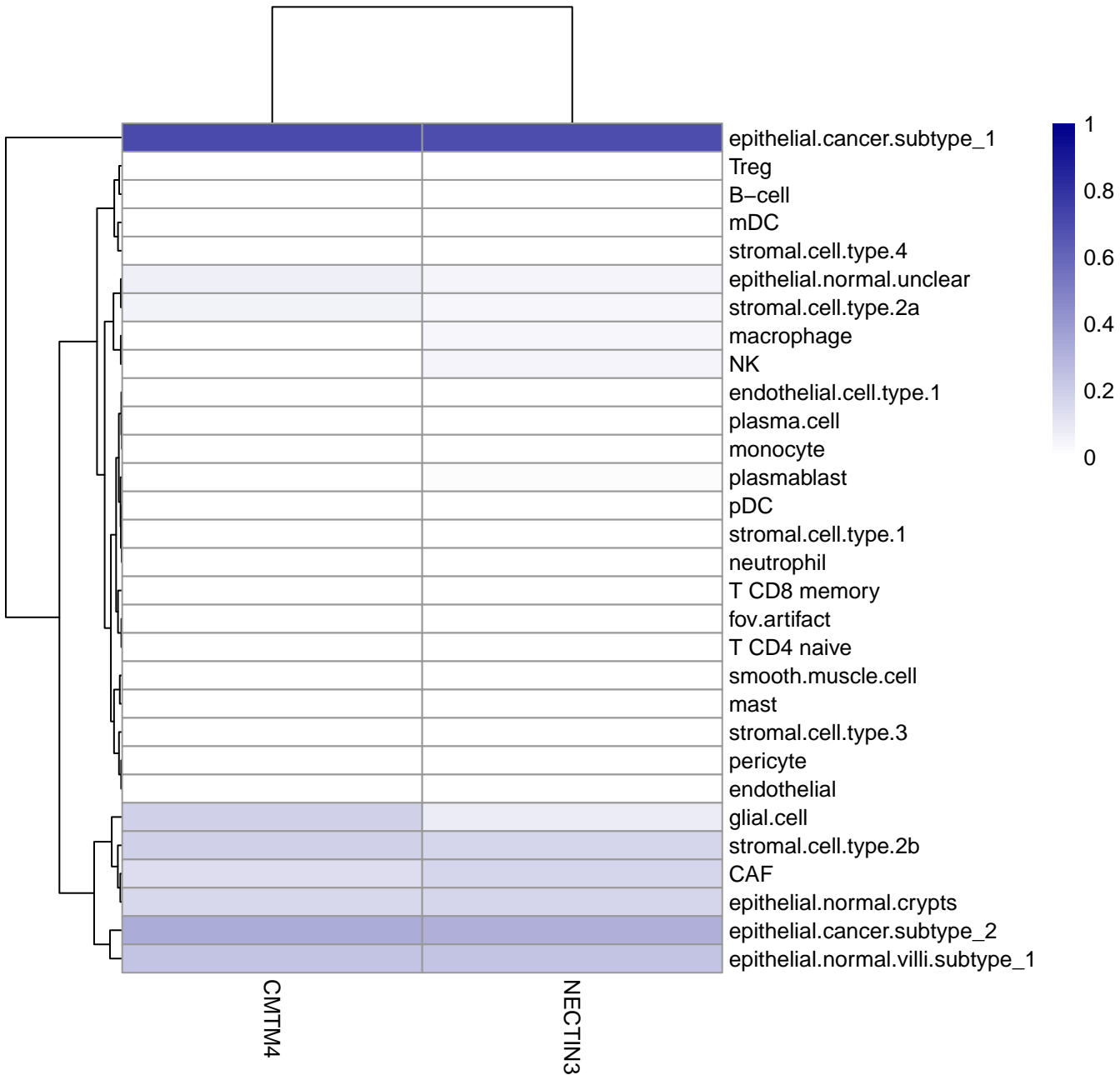
CHGA_SST_2



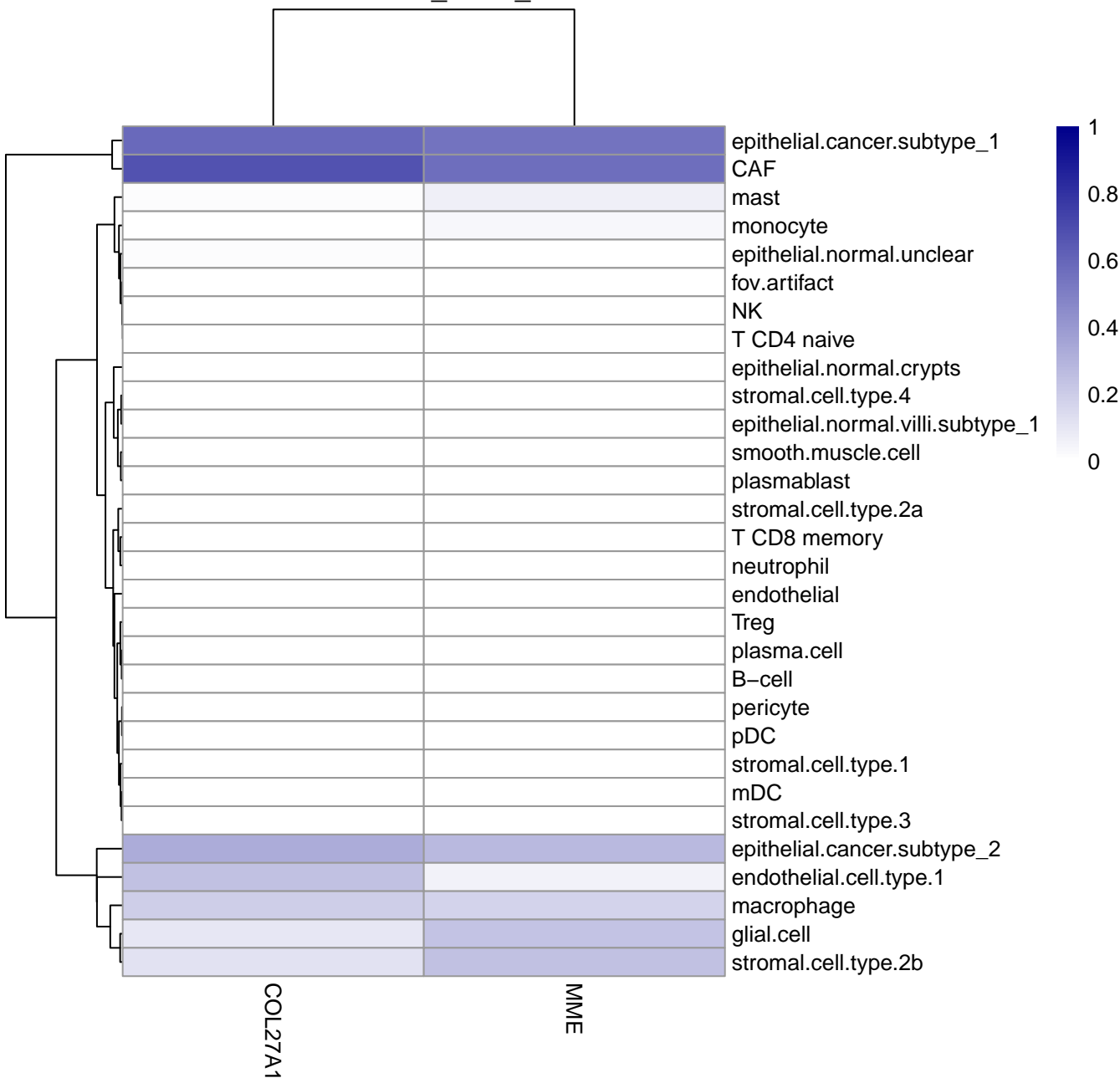
ITGA2_LAMA3_2



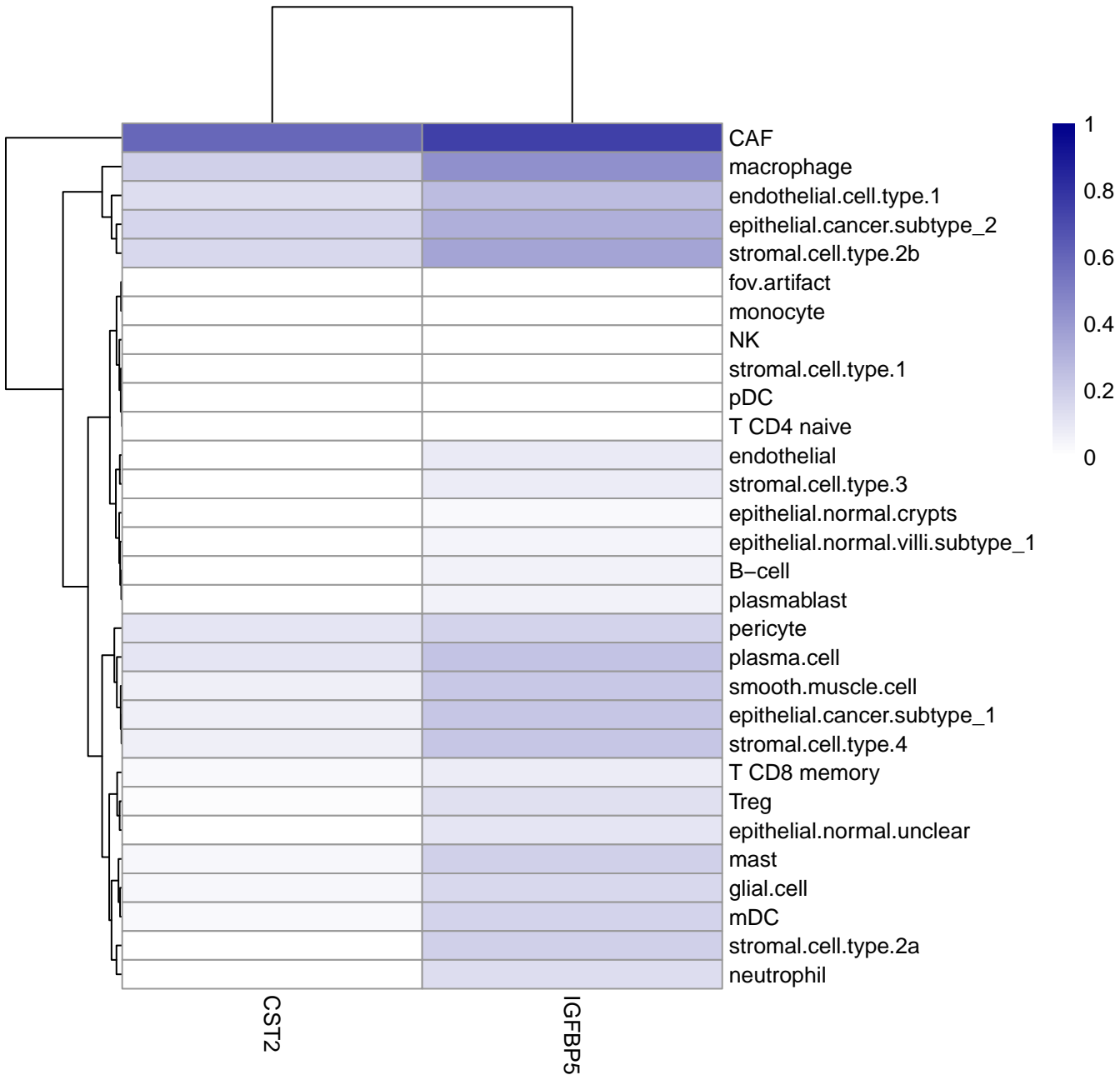
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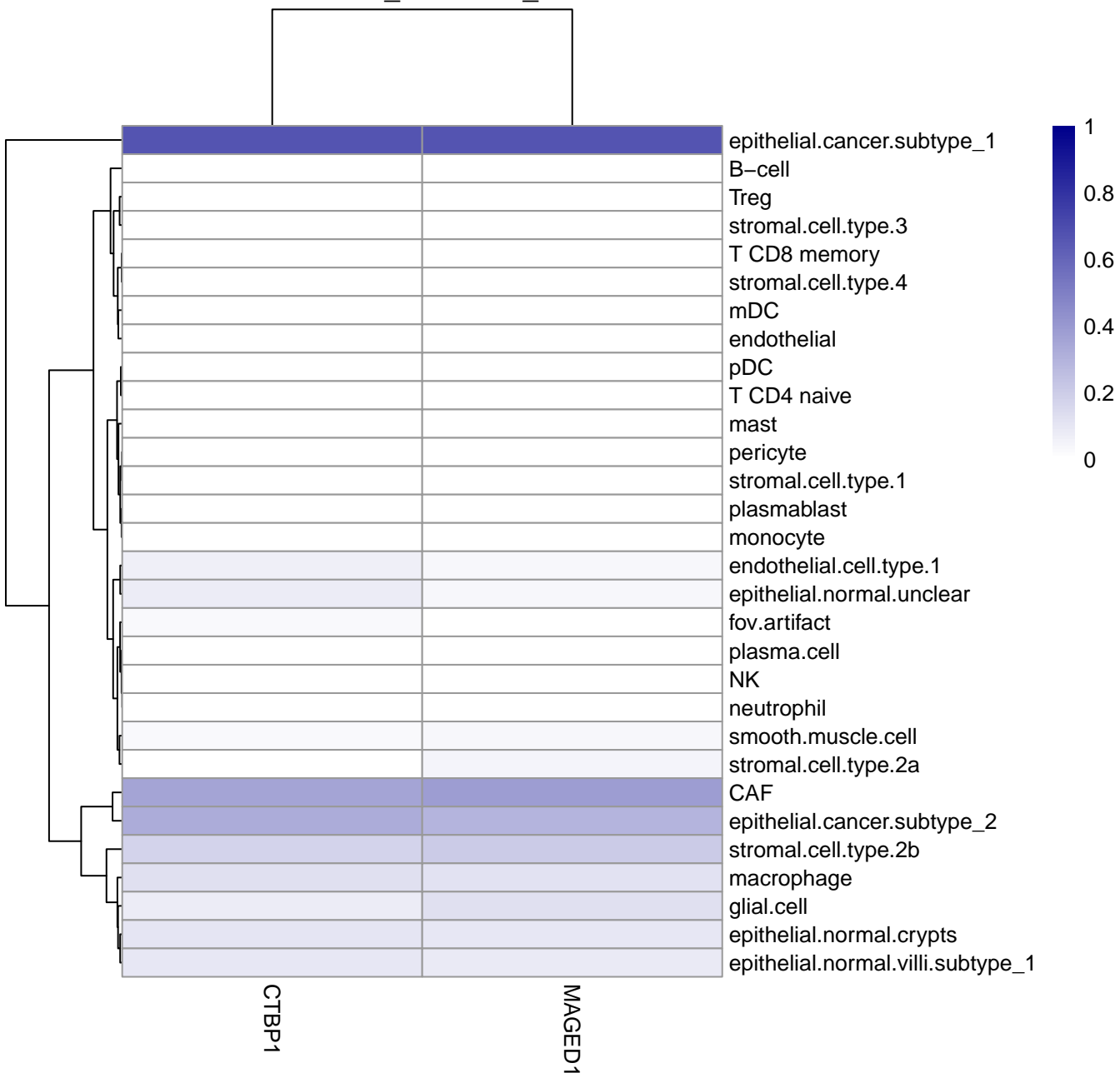
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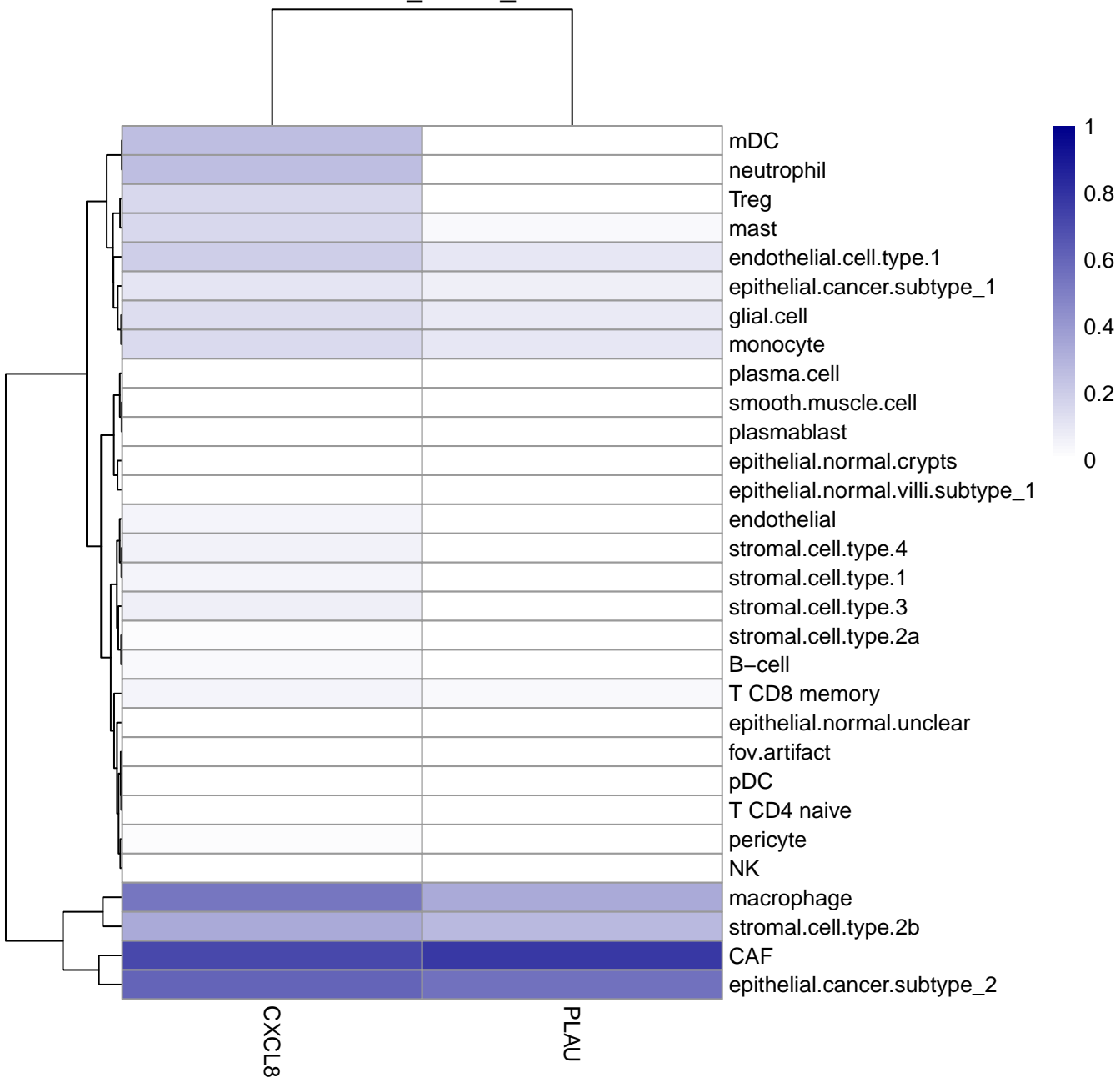
CST2_IGFBP5_2



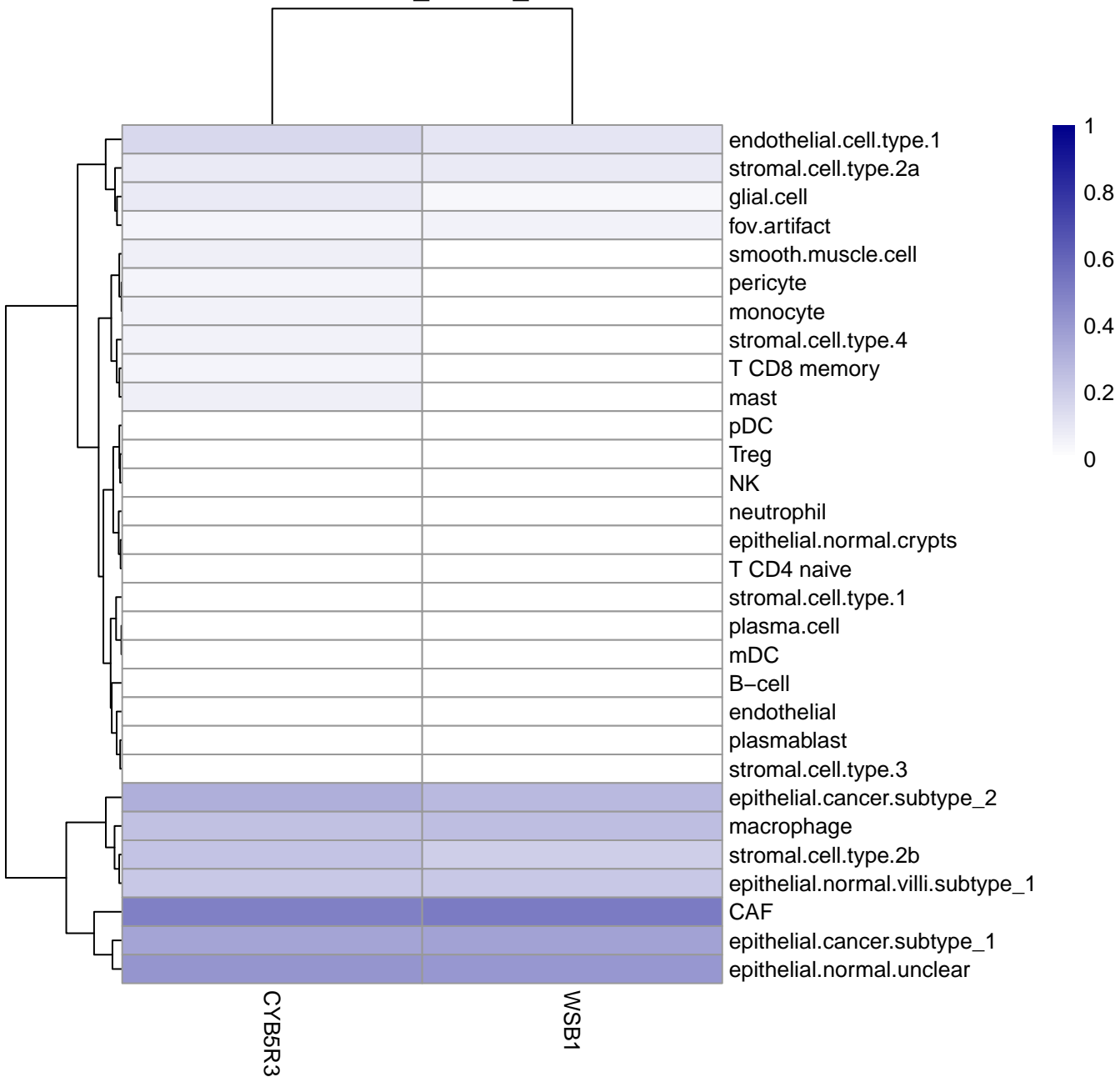
CTBP1_MAGED1_2



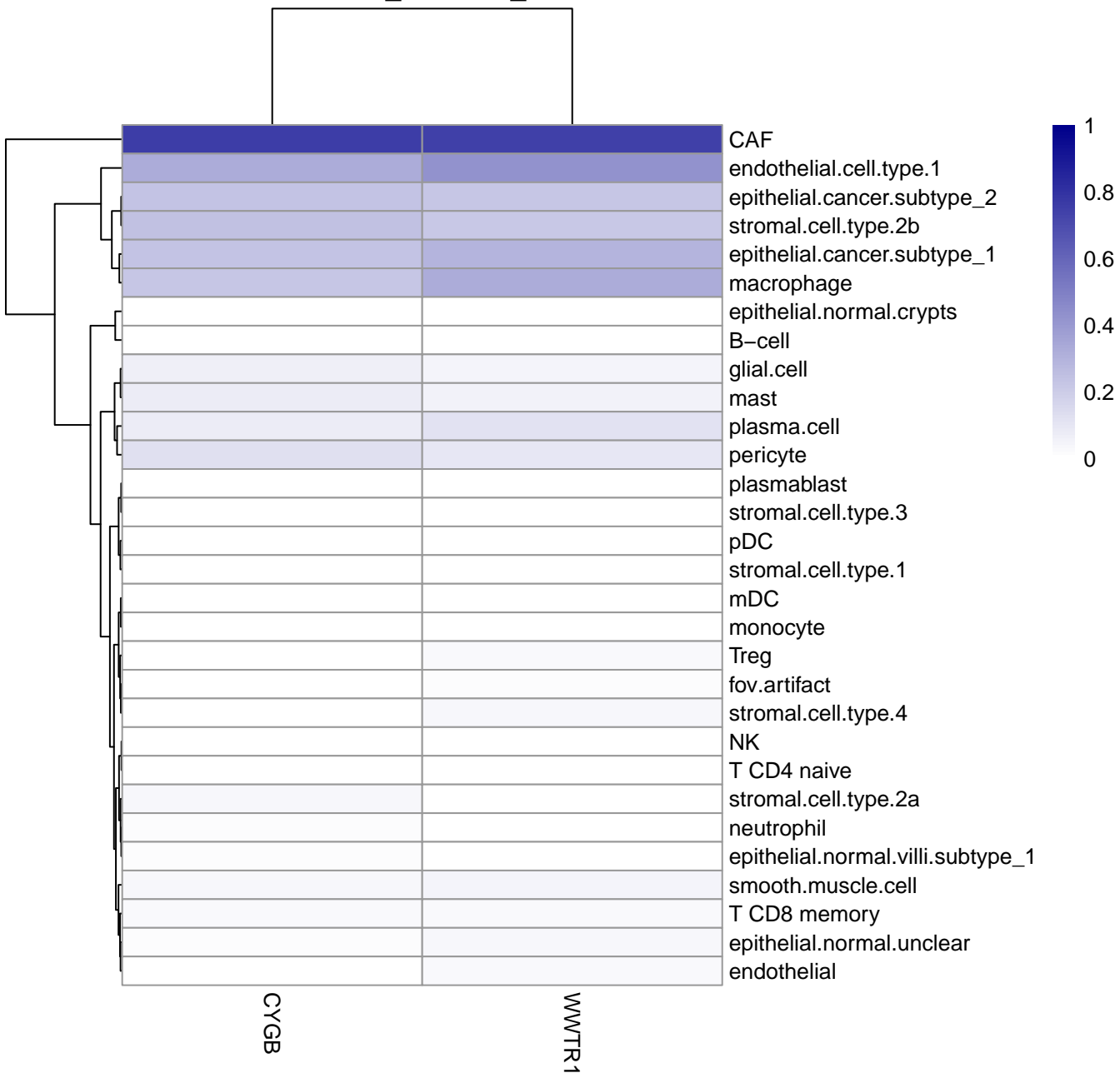
CXCL8_PLAU_2



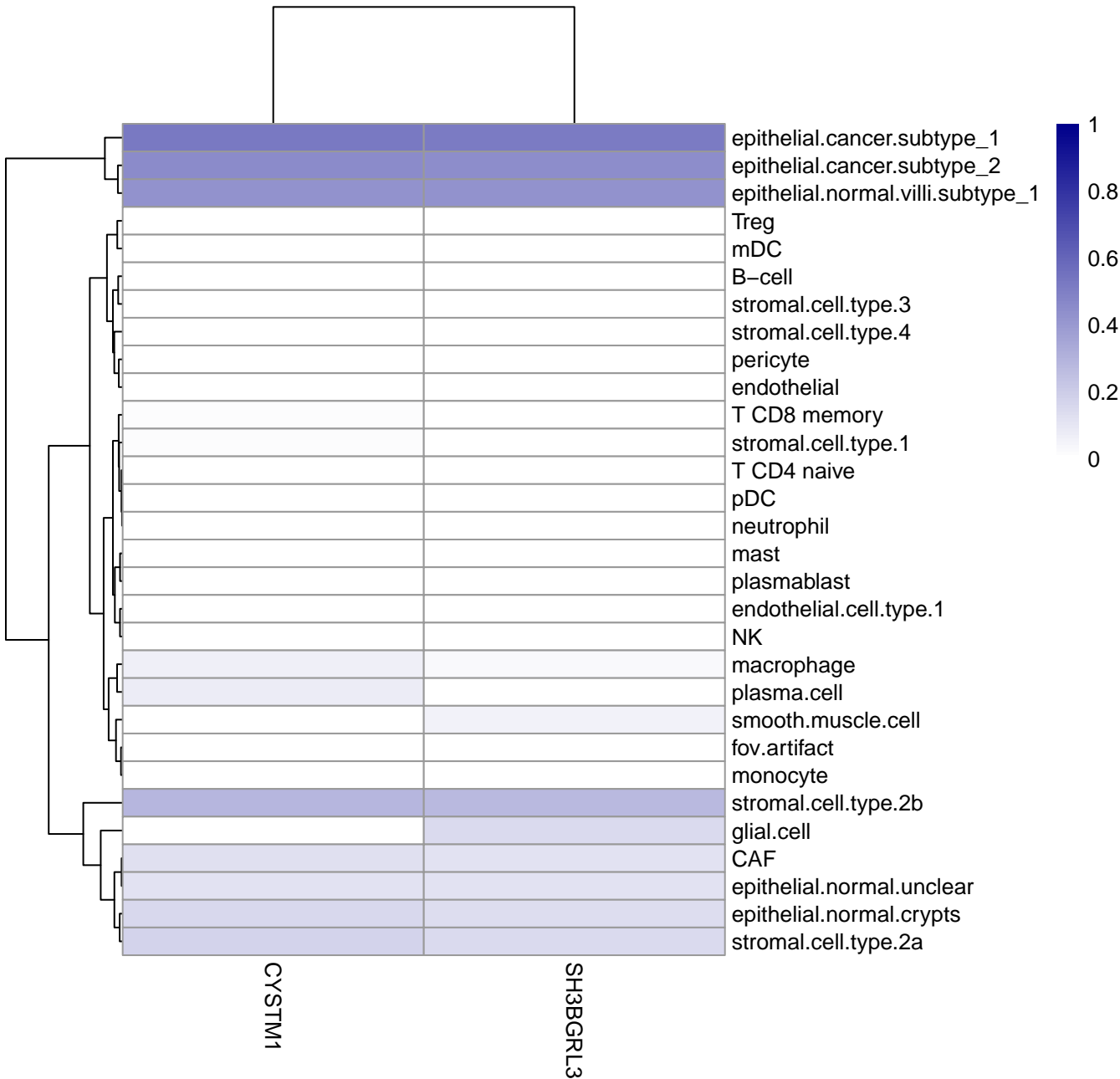
CYB5R3_WSB1_2



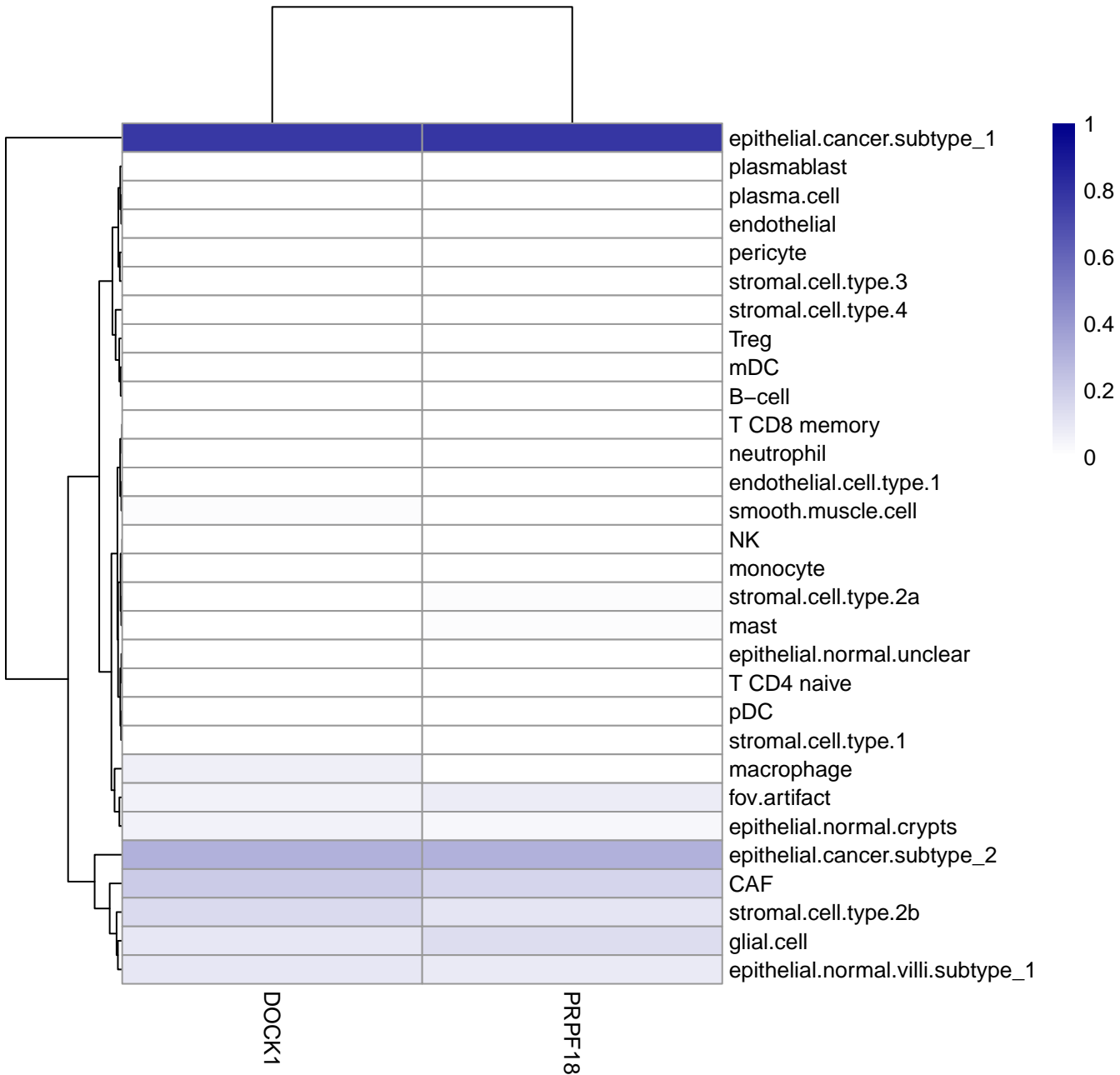
CYGB_WWTR1_2



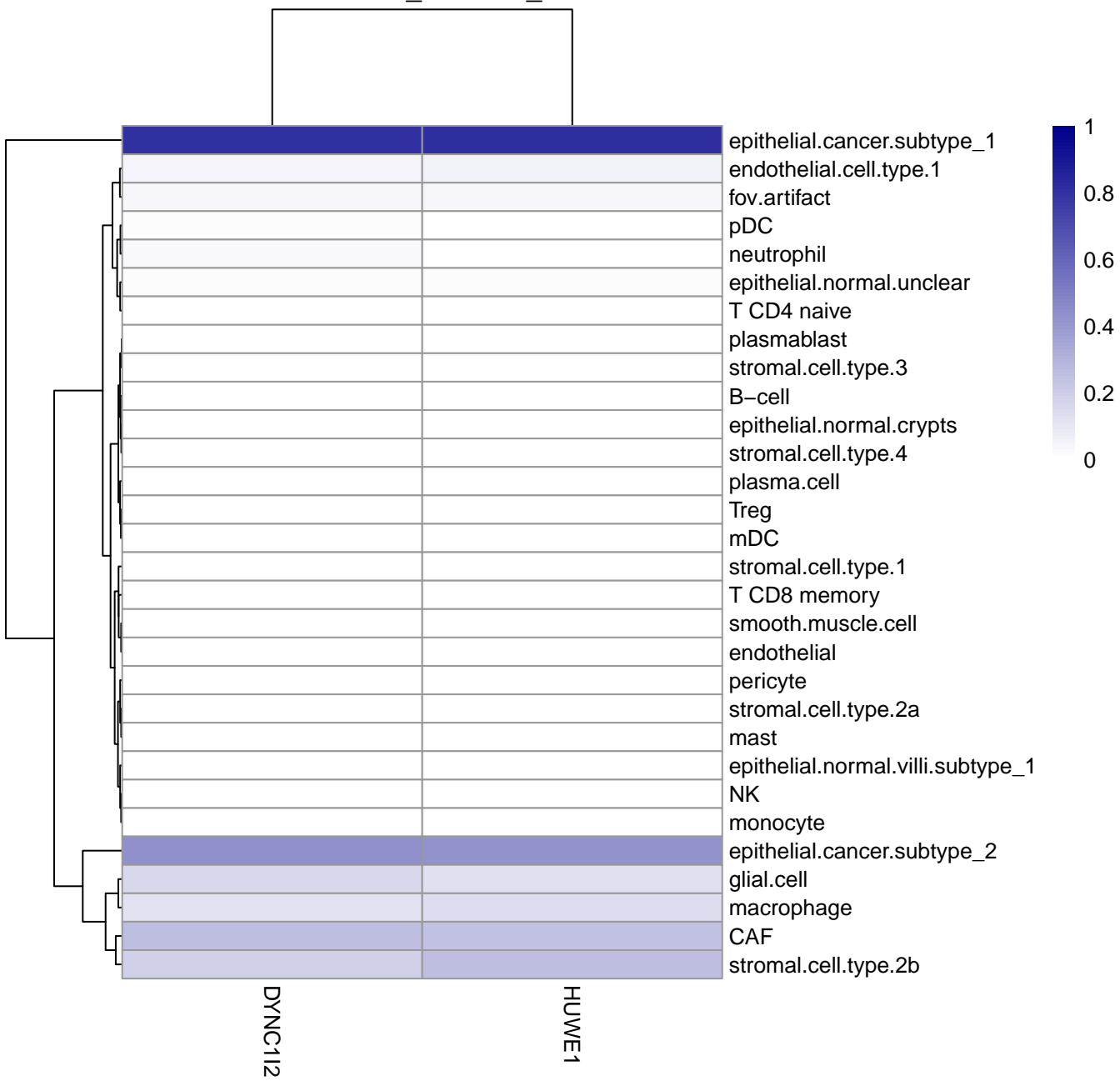
CYSTM1_SH3BGRL3_2



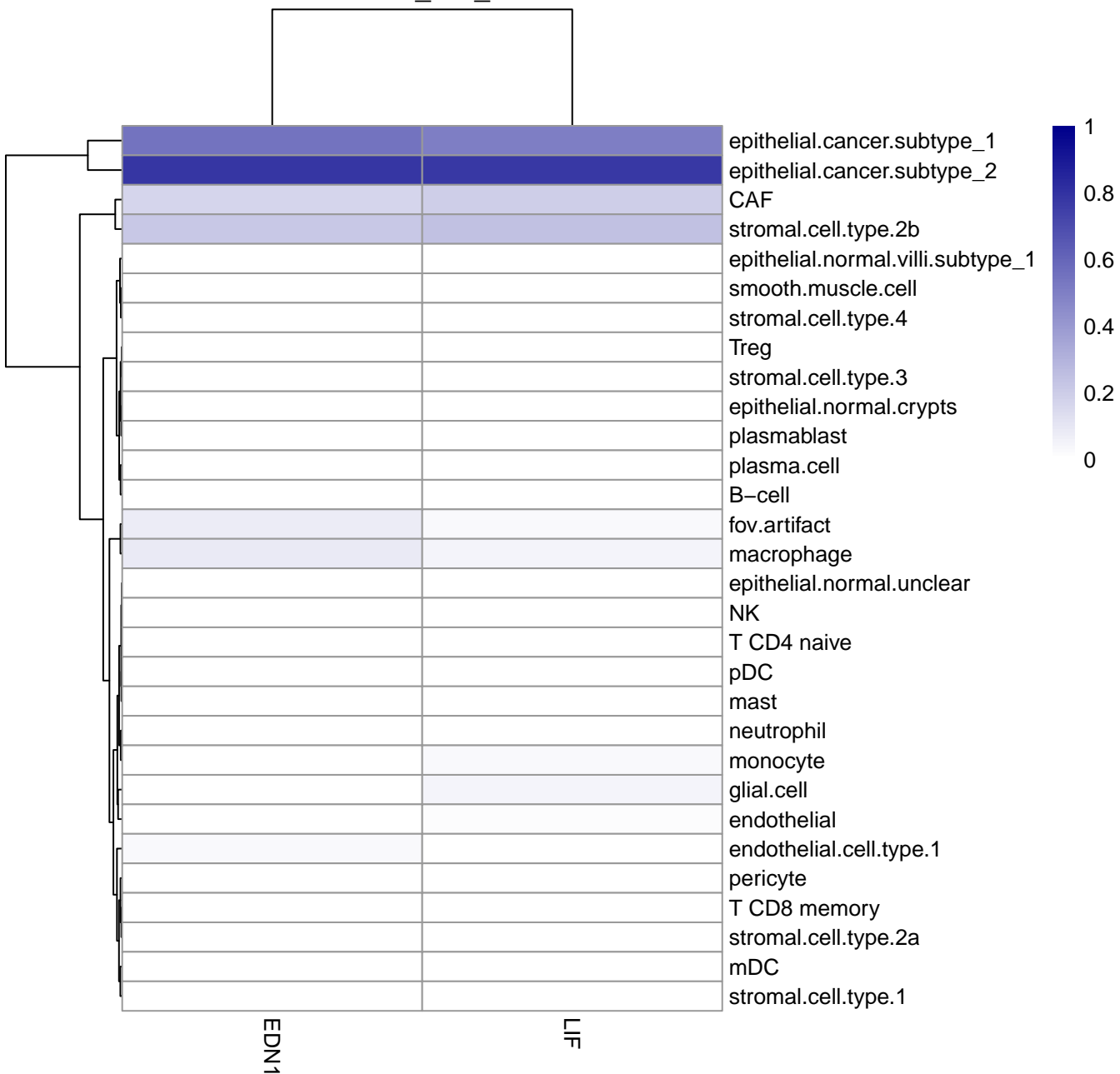
DOCK1_PRPF18_2



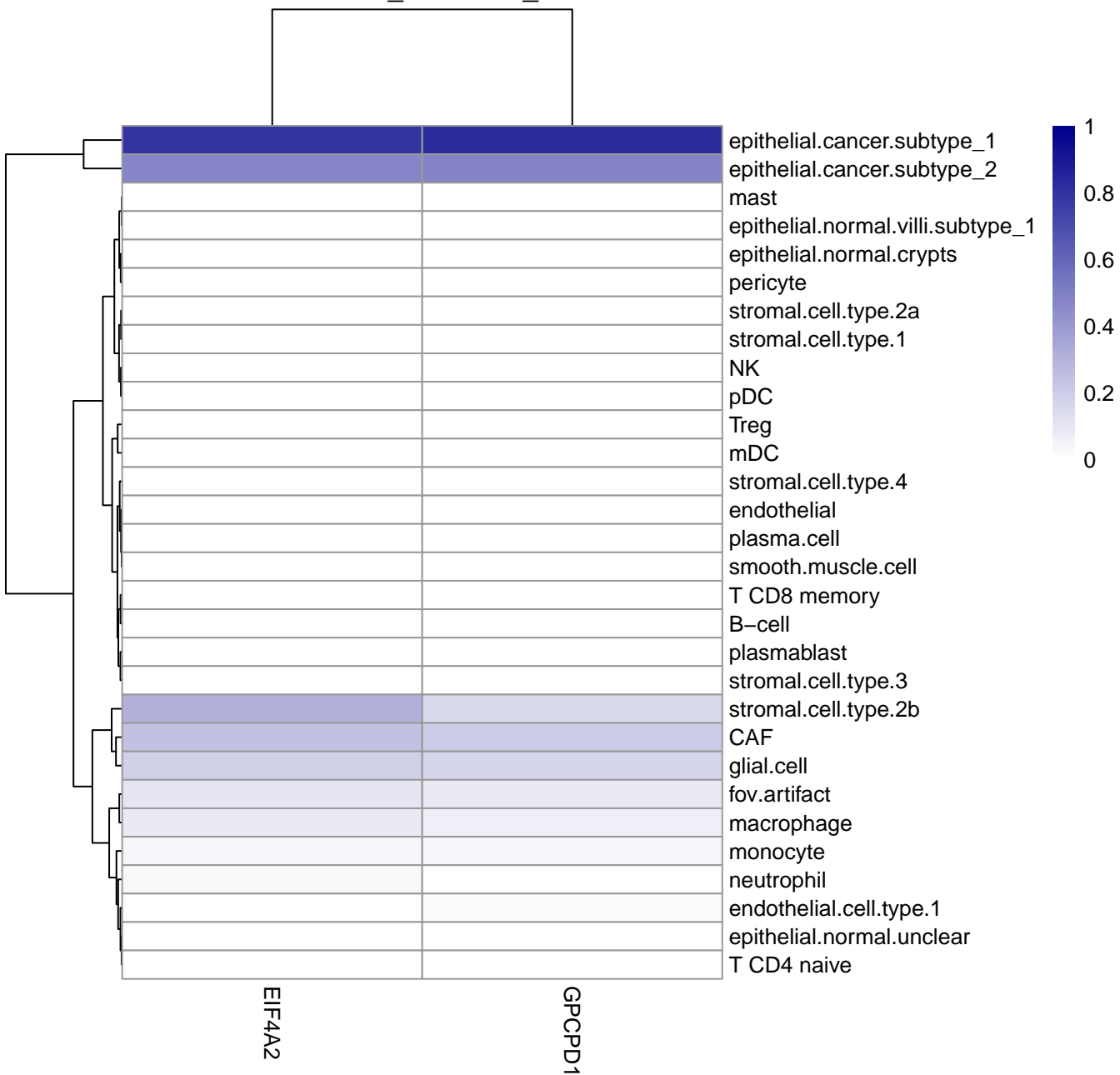
DYNC112_HUWE1_2



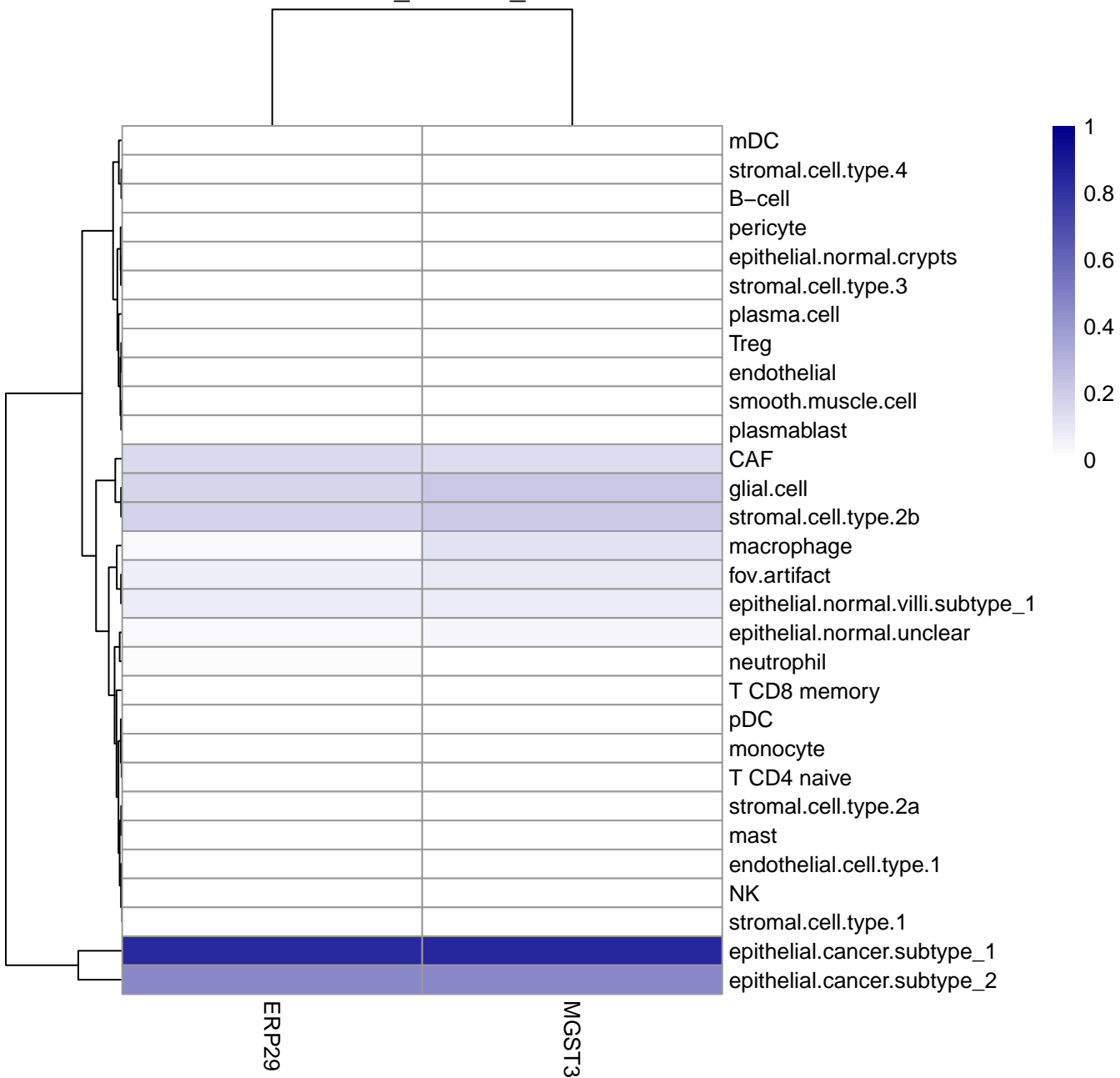
EDN1_LIF_2



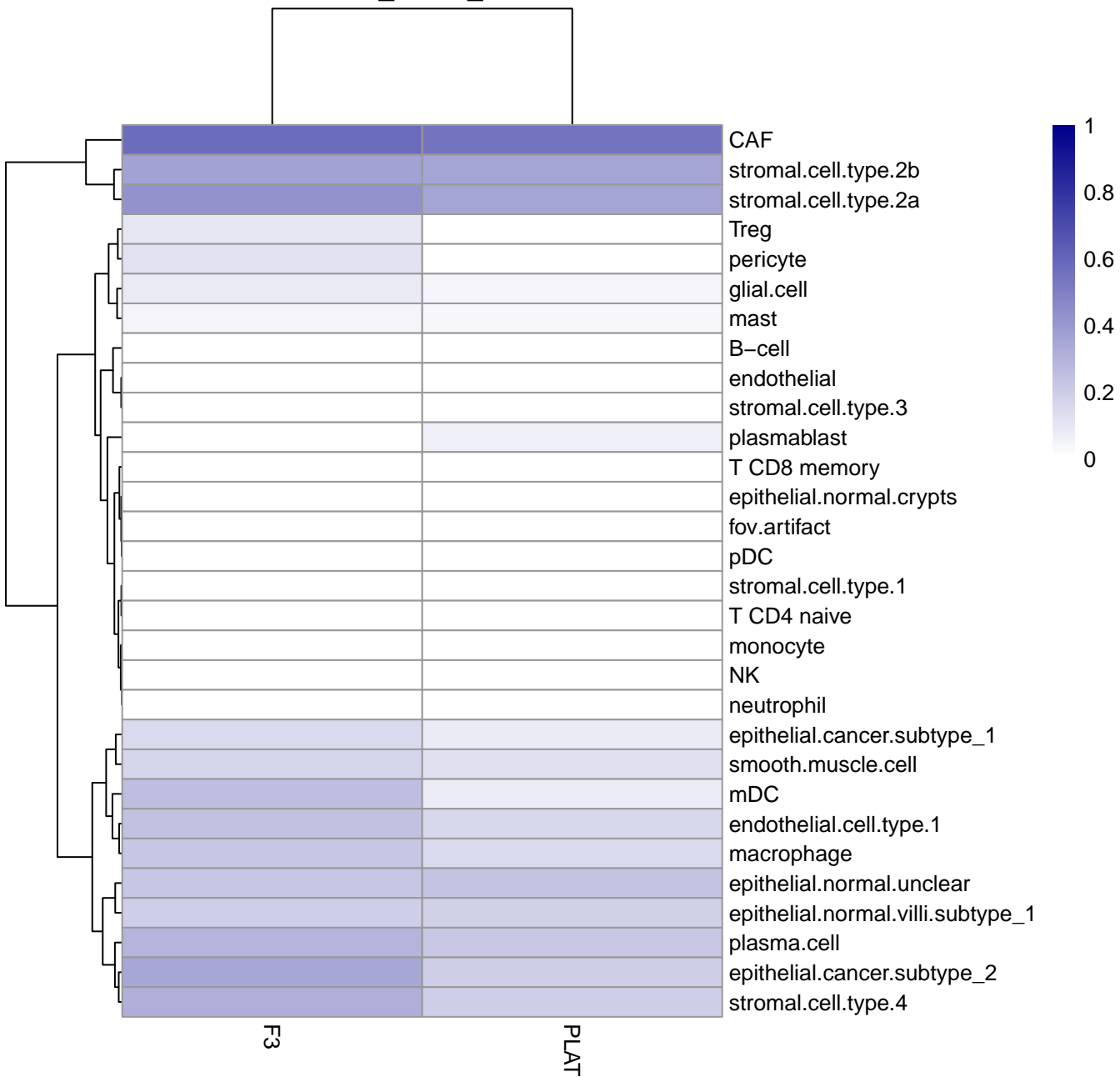
EIF4A2_GPCPD1_2



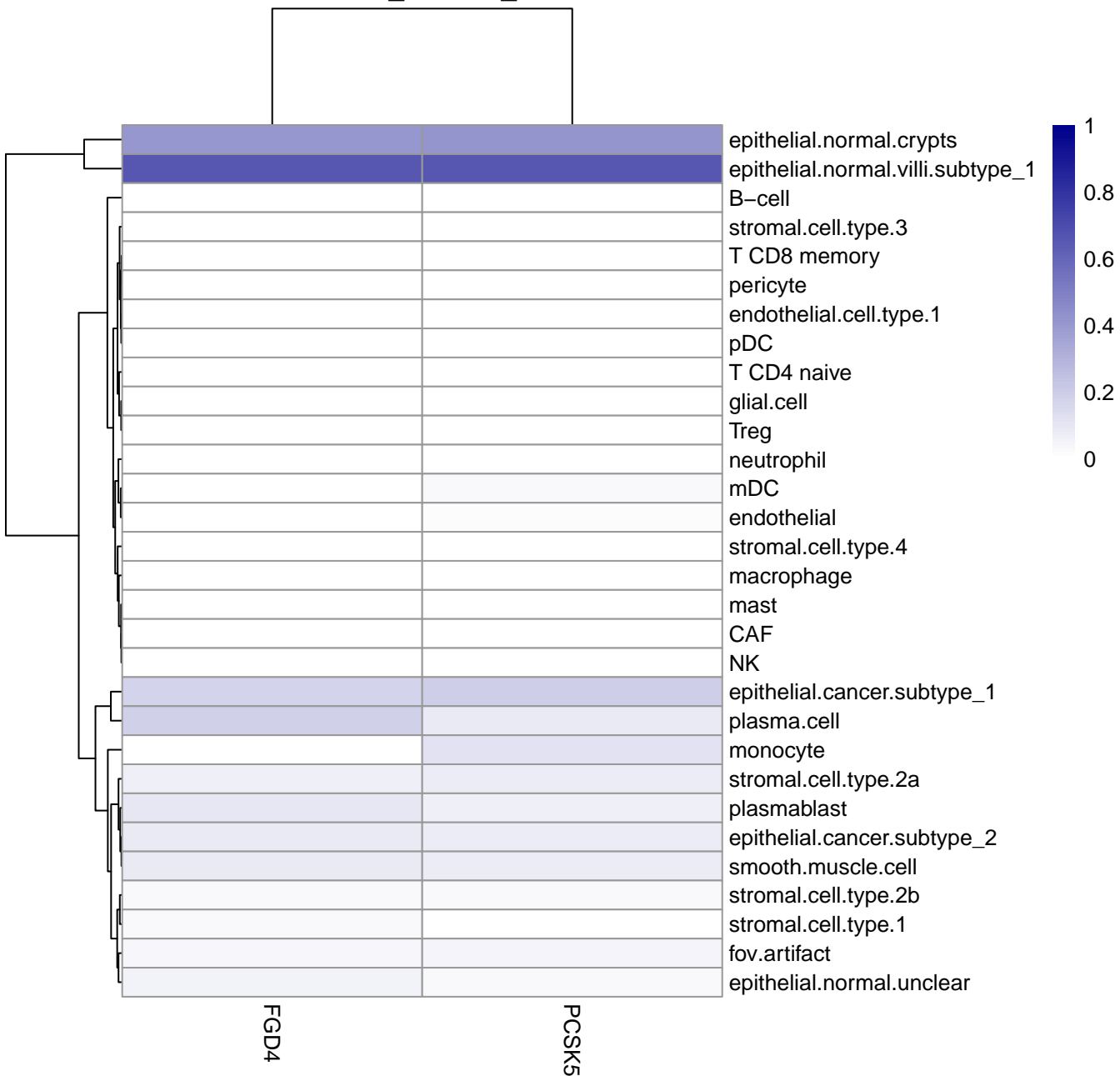
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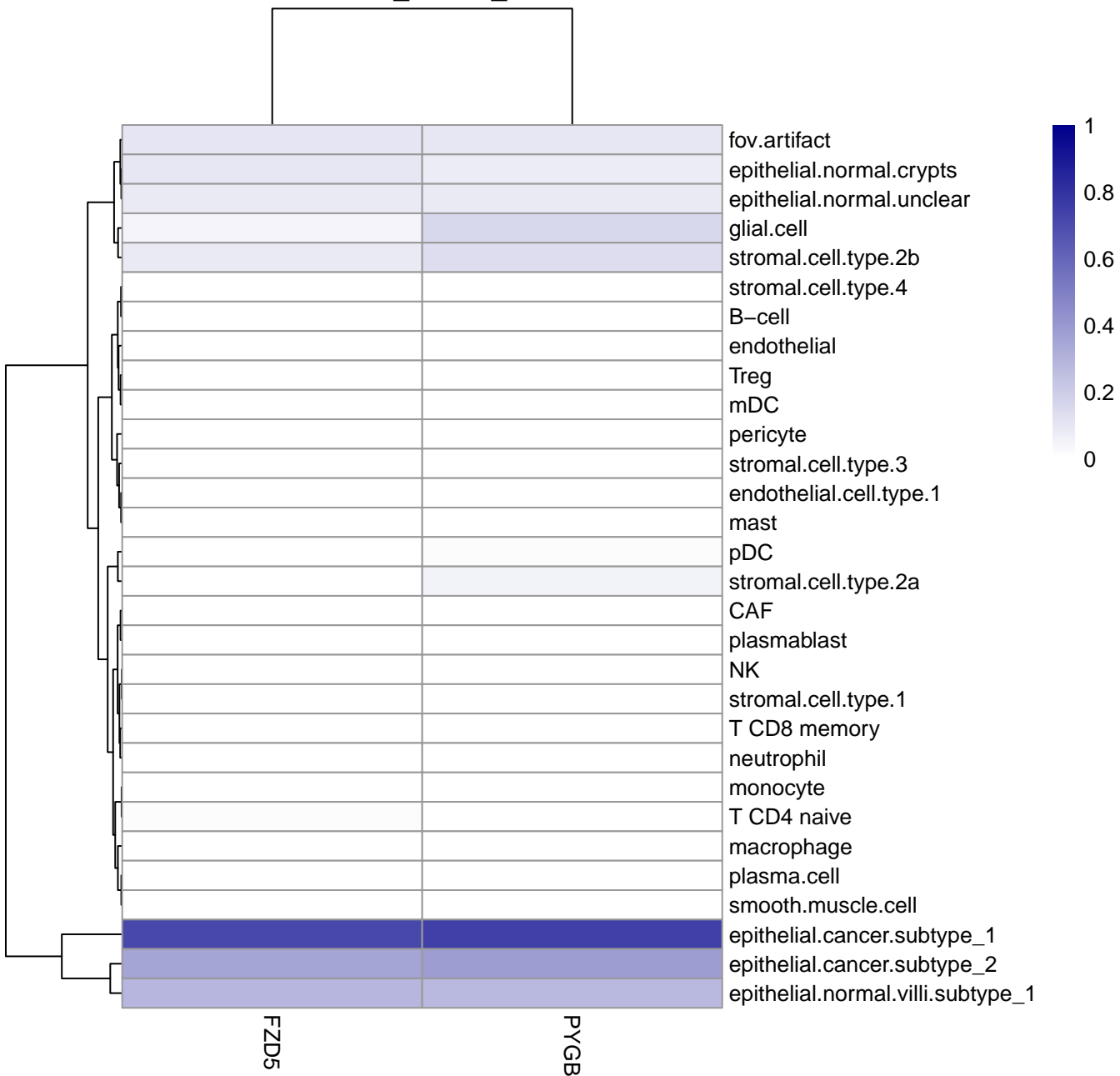
F3_PLAT_2



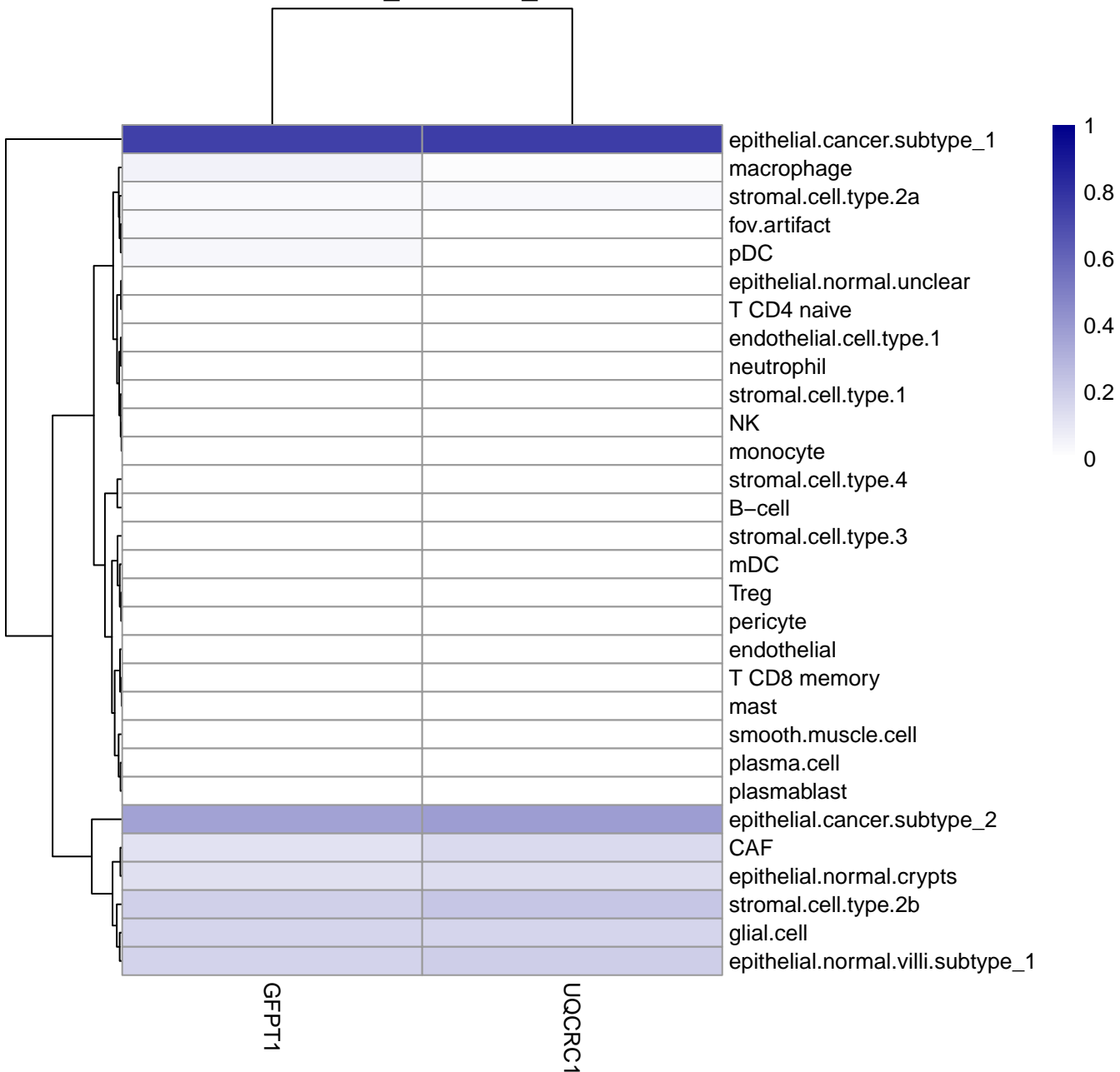
FGD4_PCSK5_2



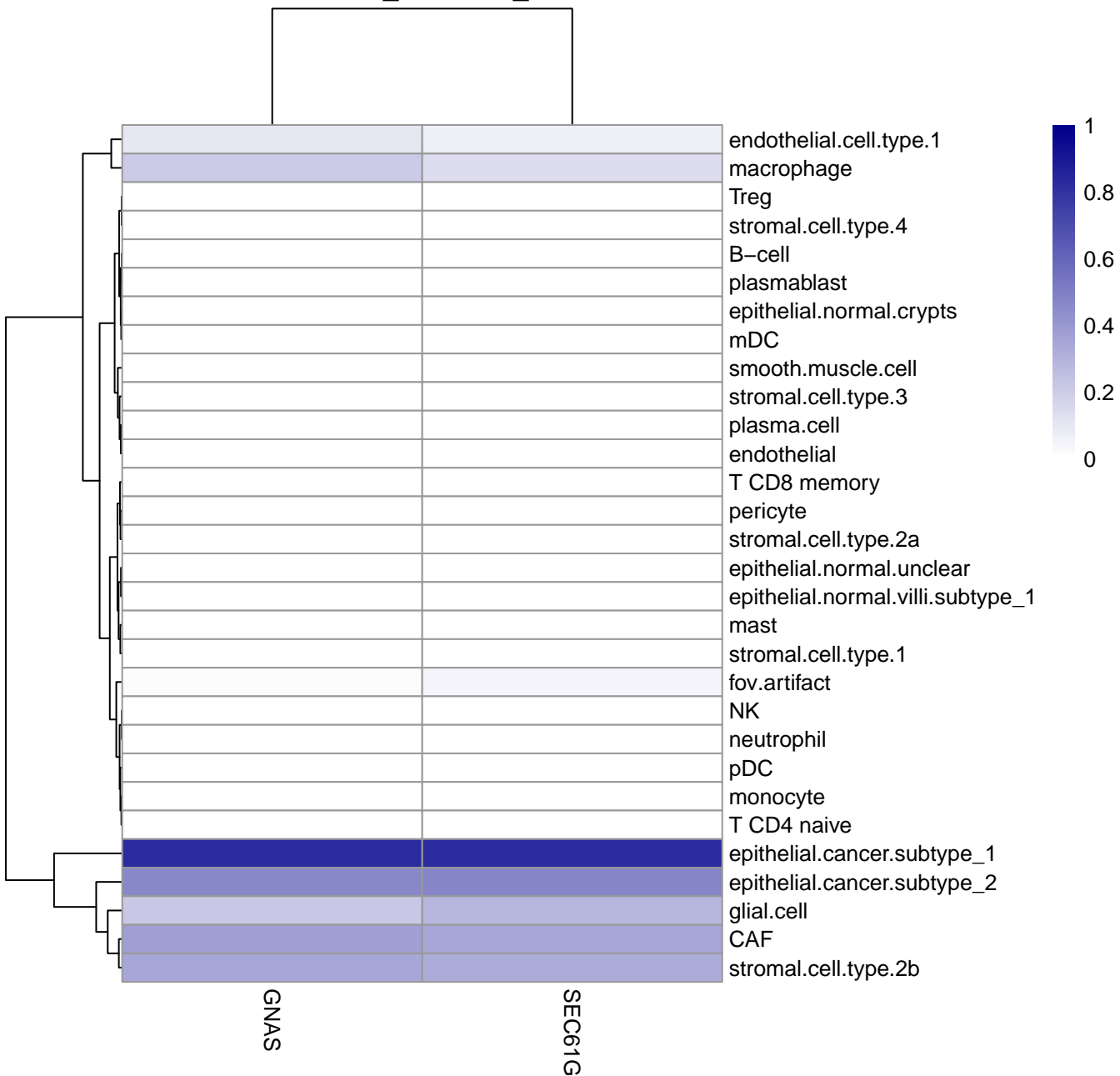
FZD5_PYGB_2



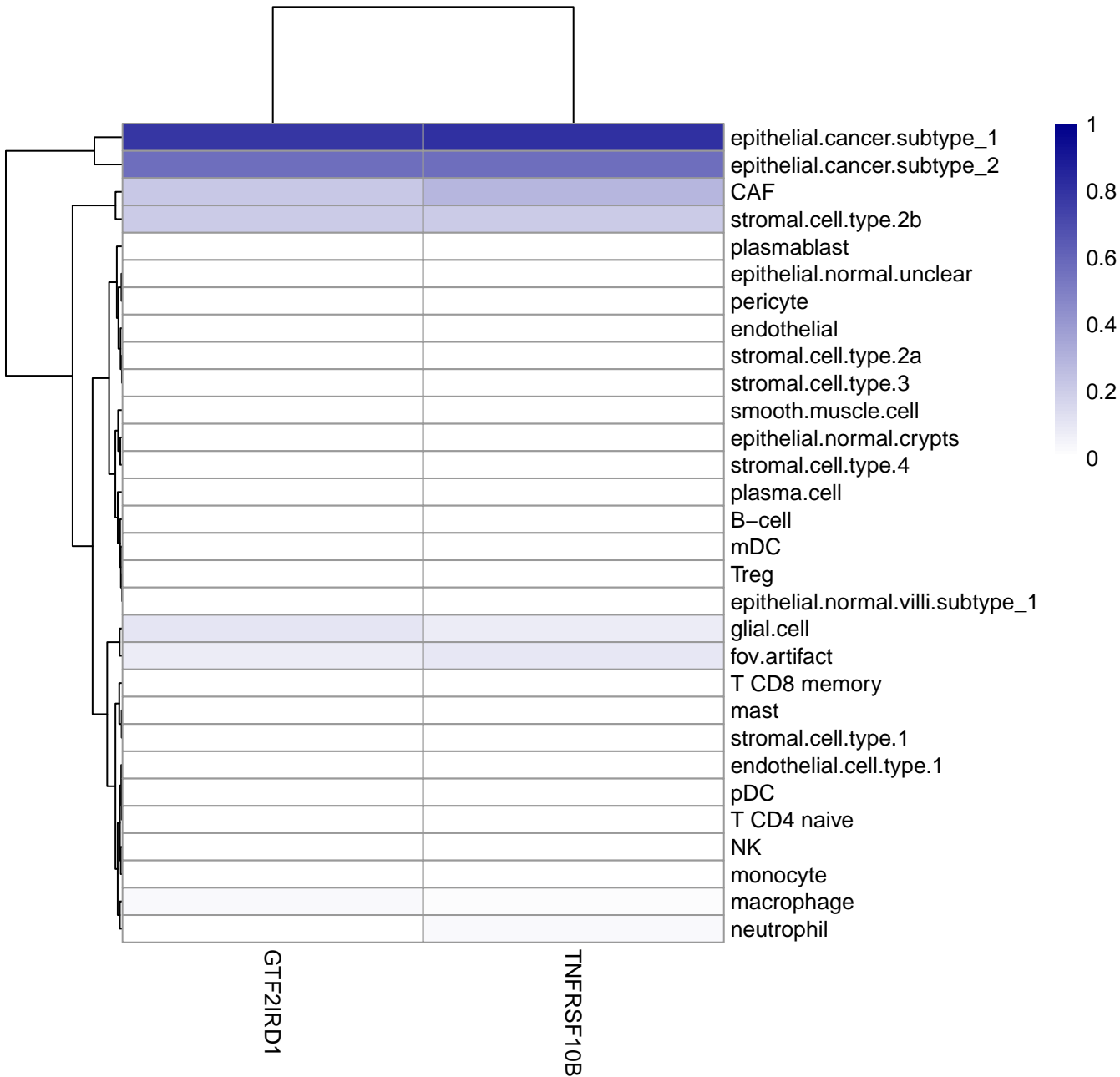
GFPT1_UQCRC1_2



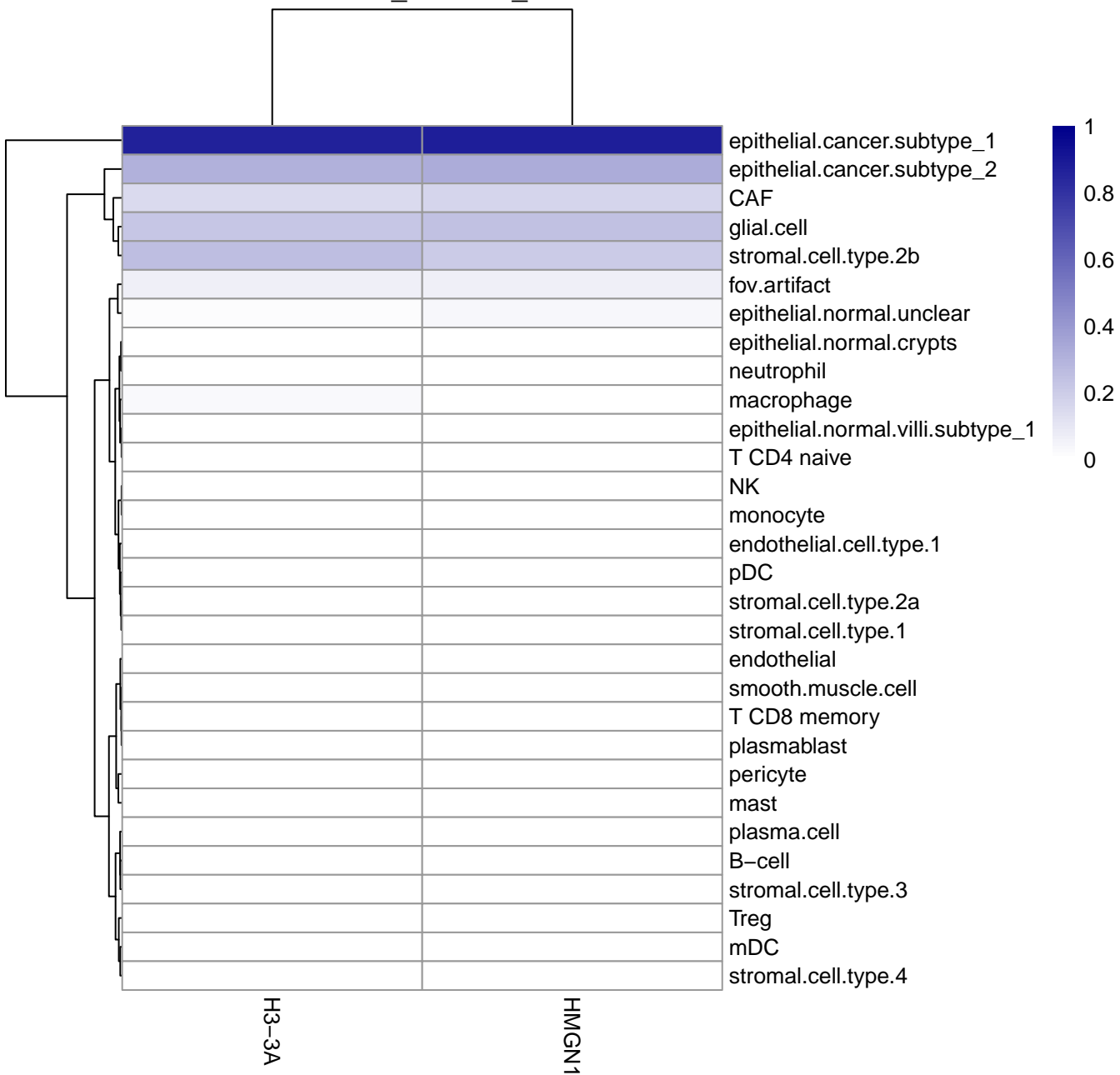
GNAS_SEC61G_2



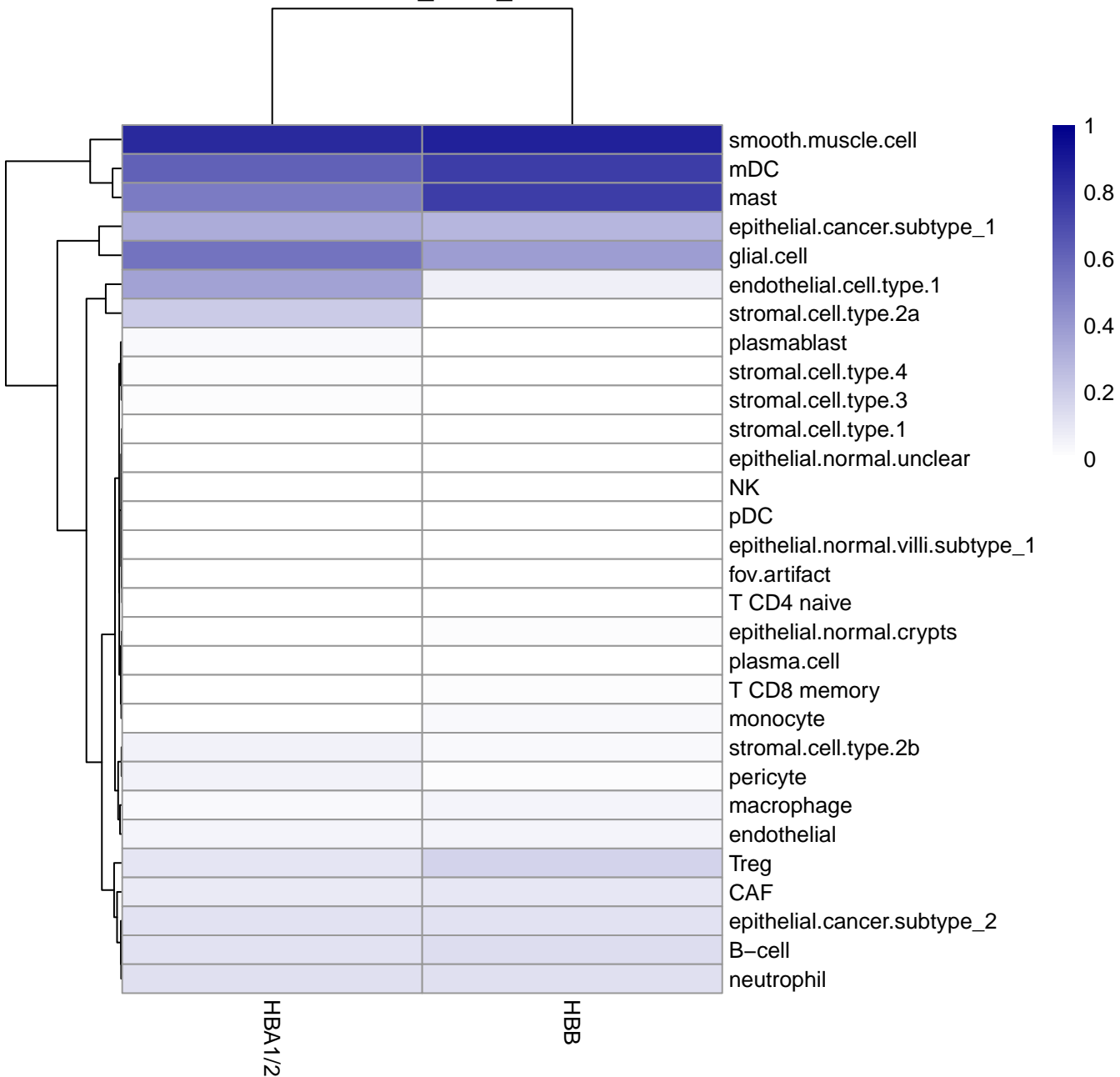
GTF2IRD1_TNFRSF10B_2



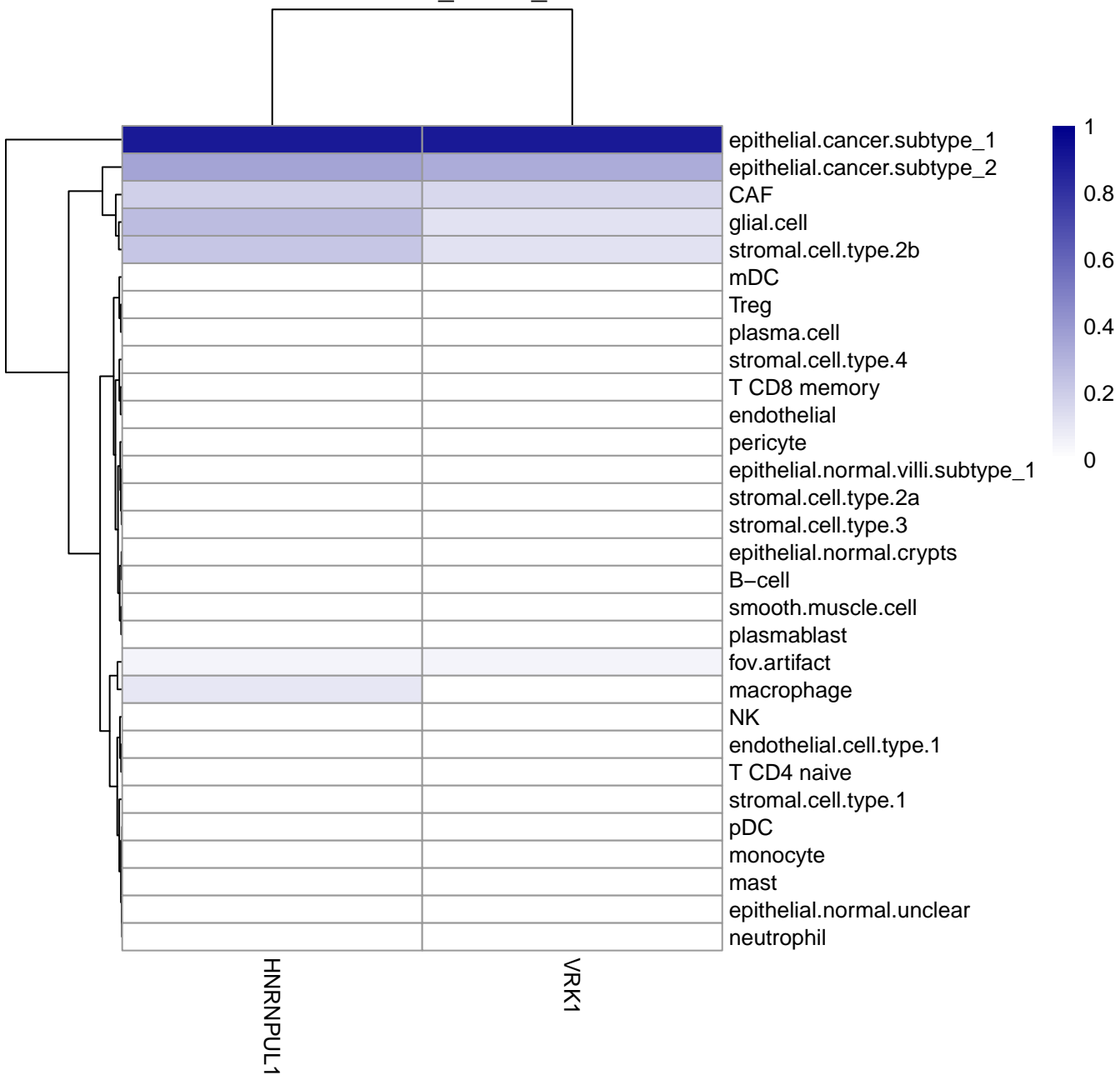
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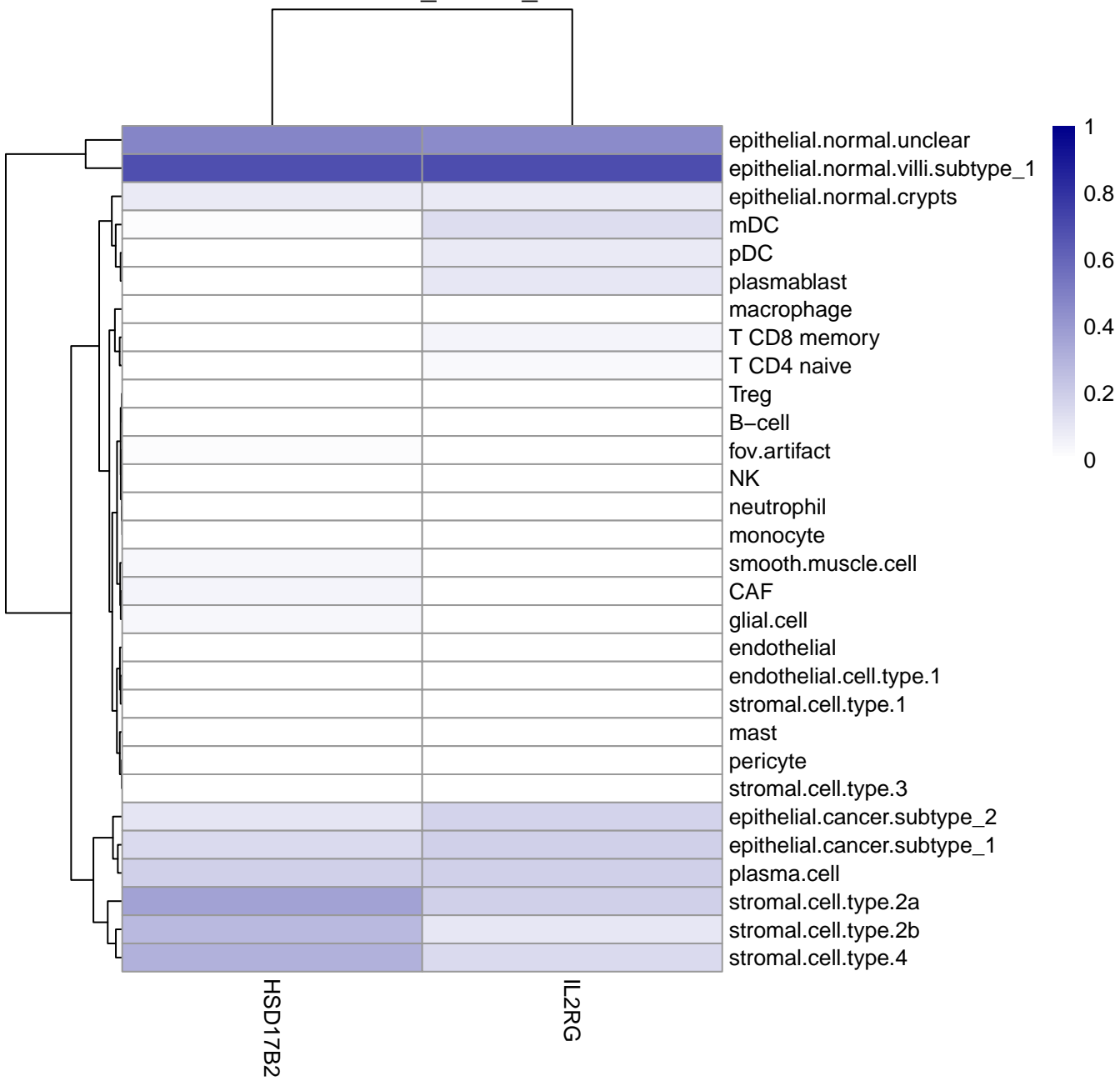
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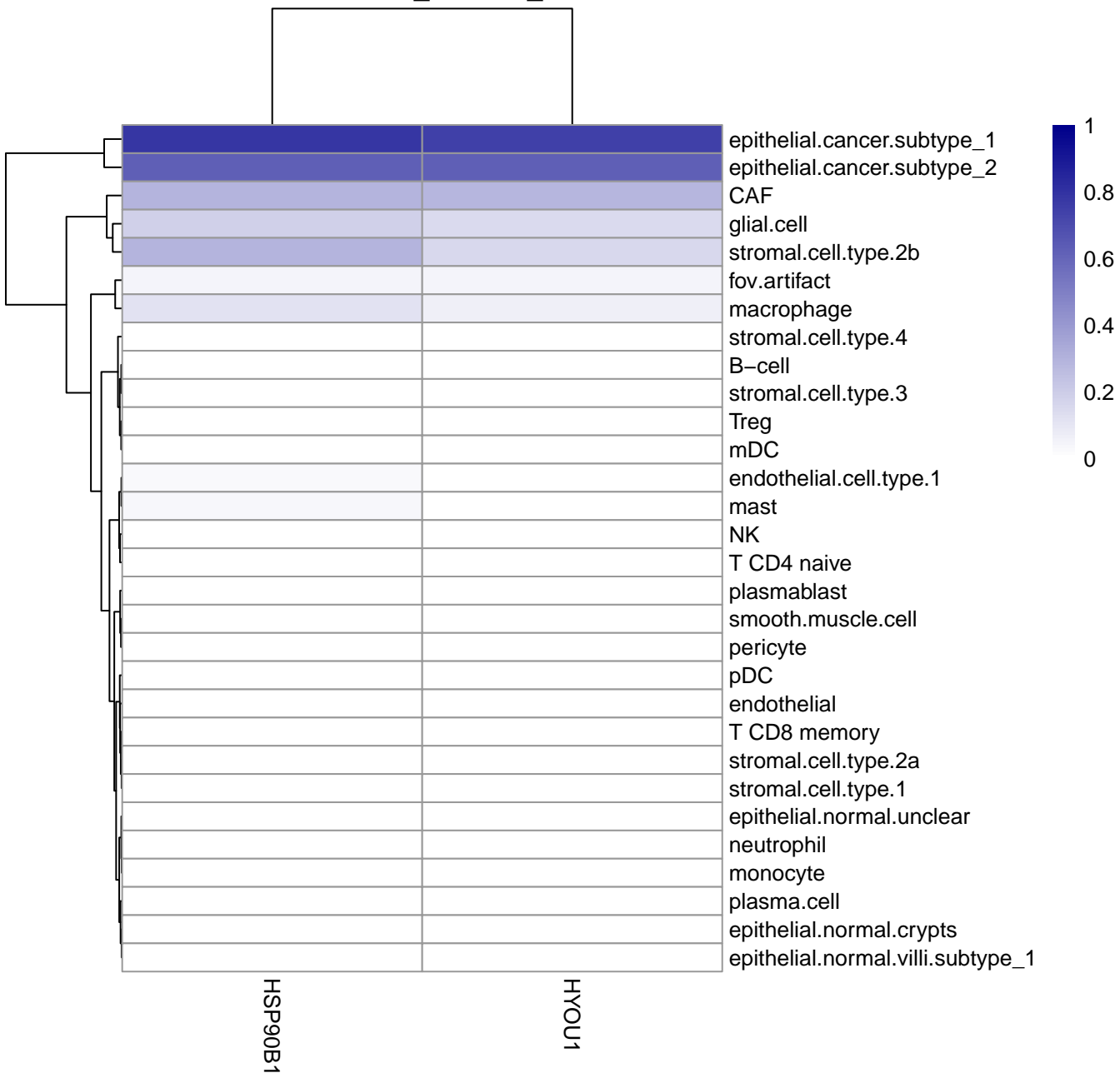
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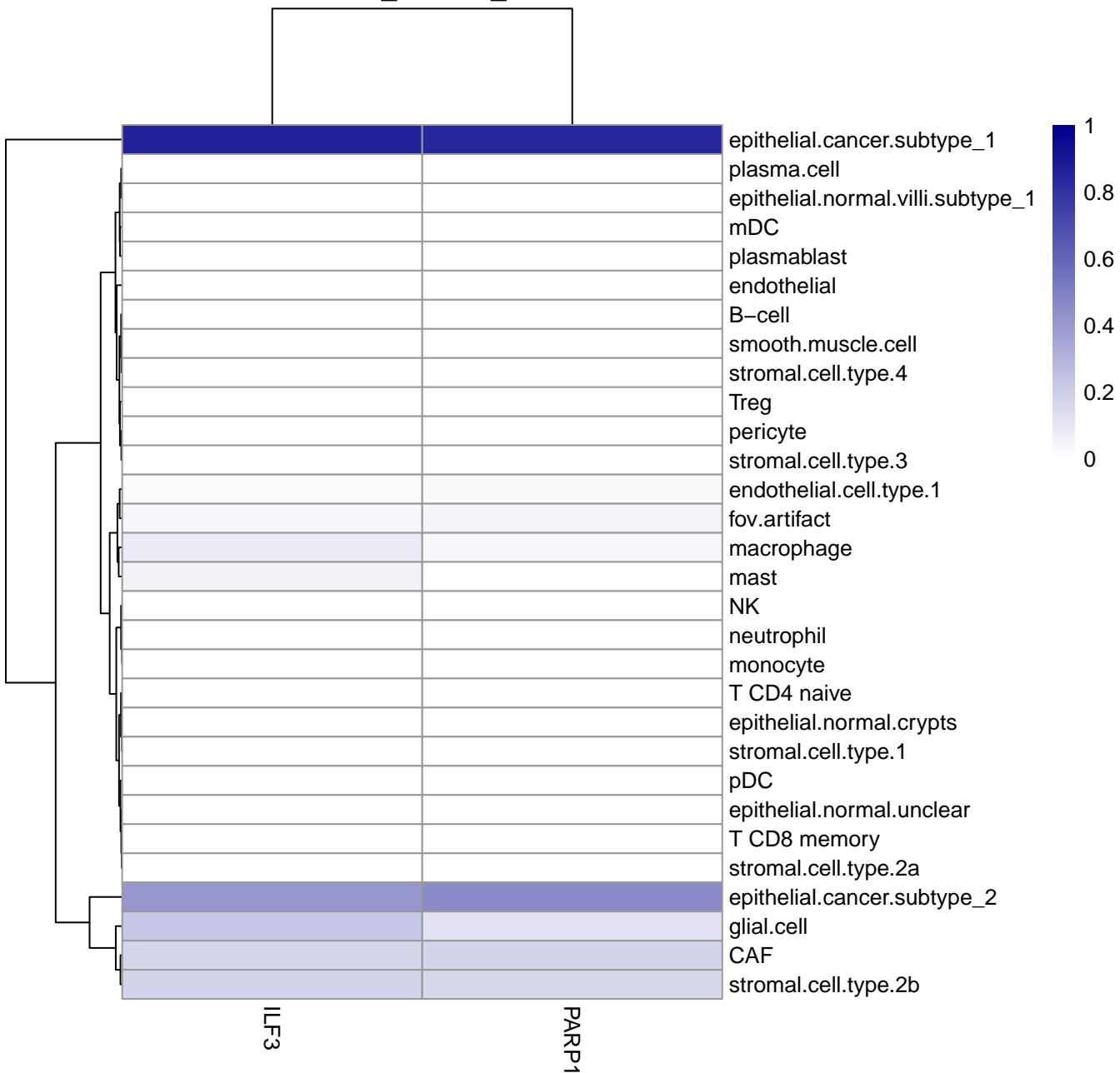
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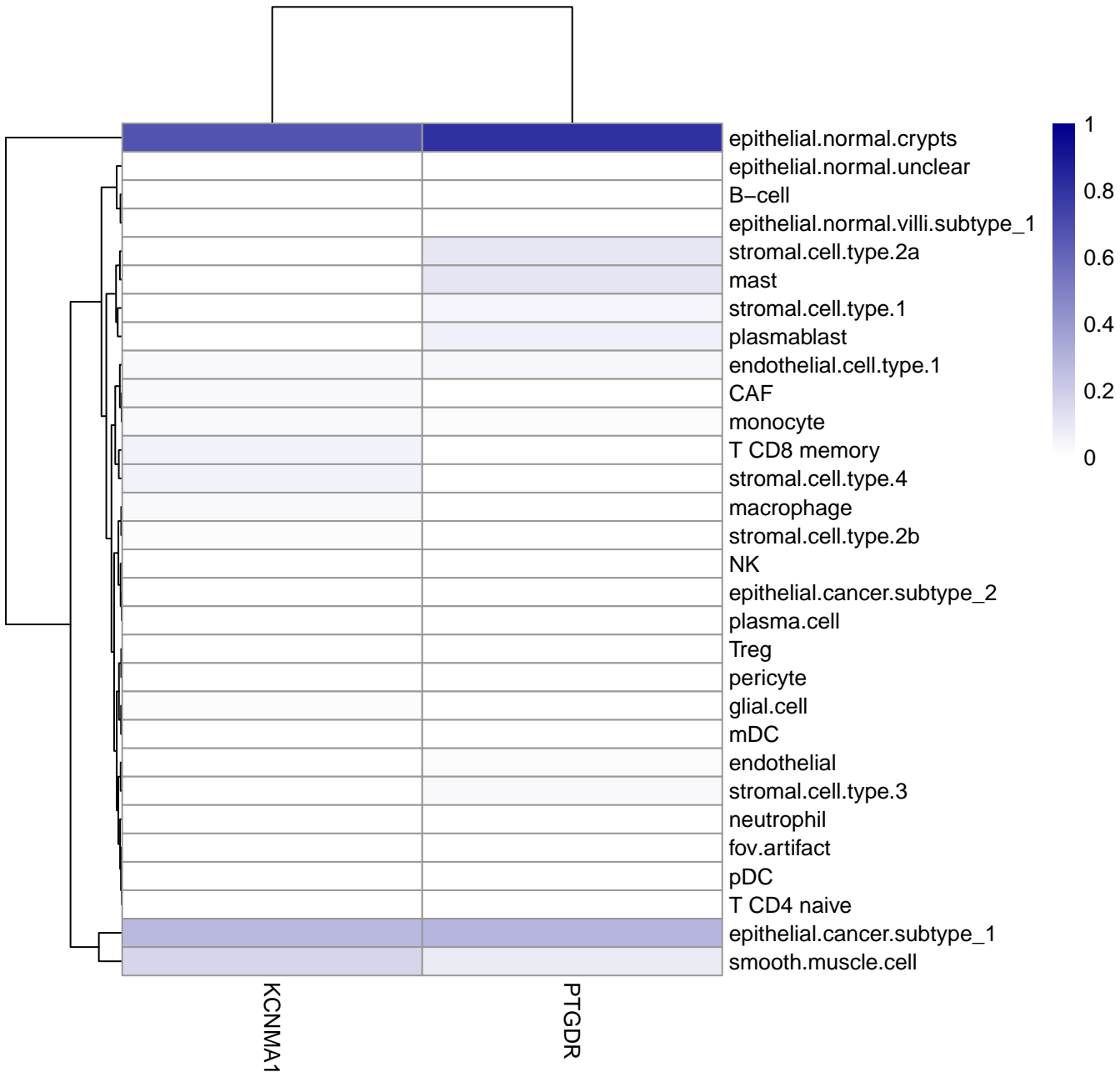
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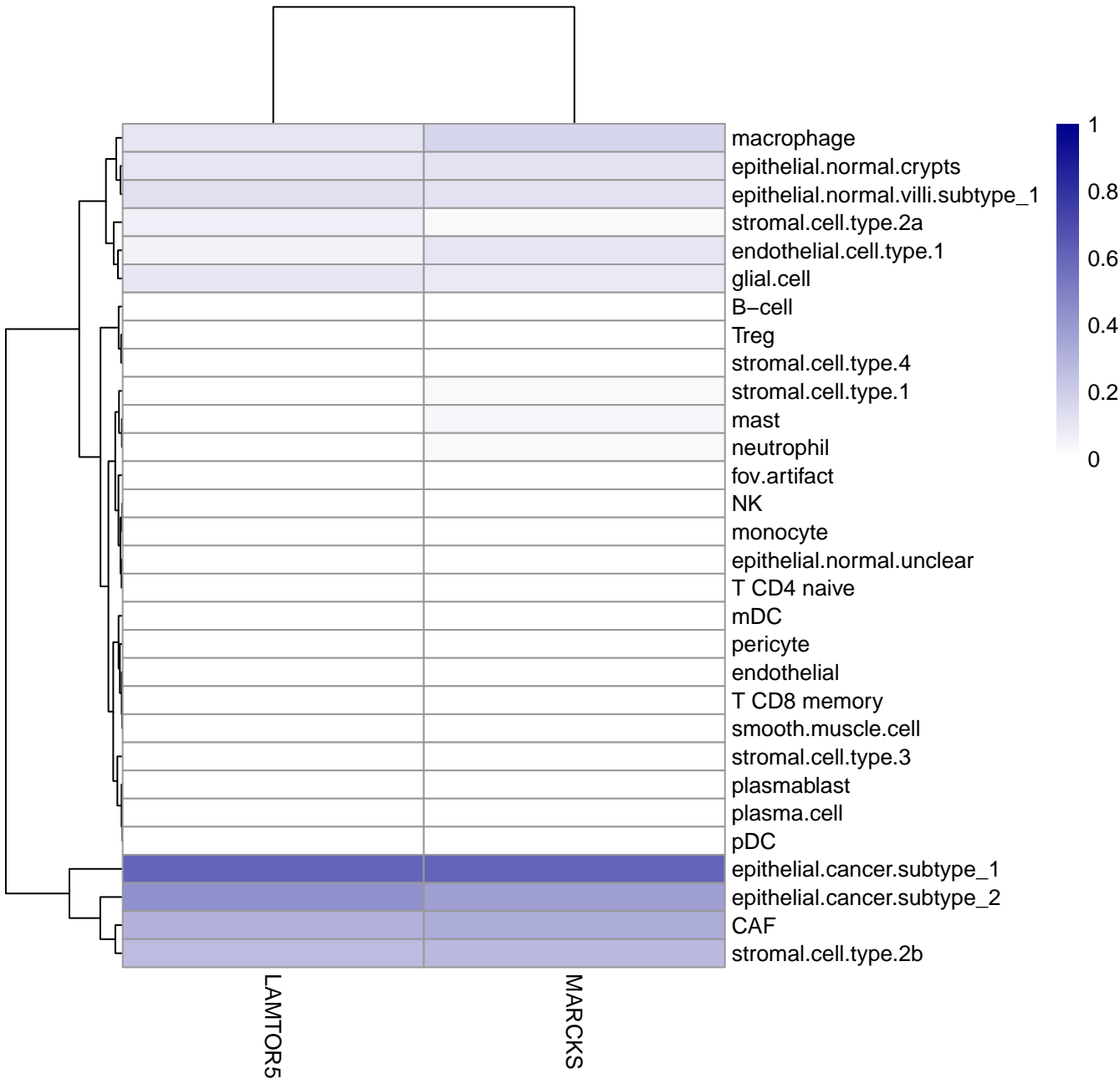
ILF3_PARP1_2



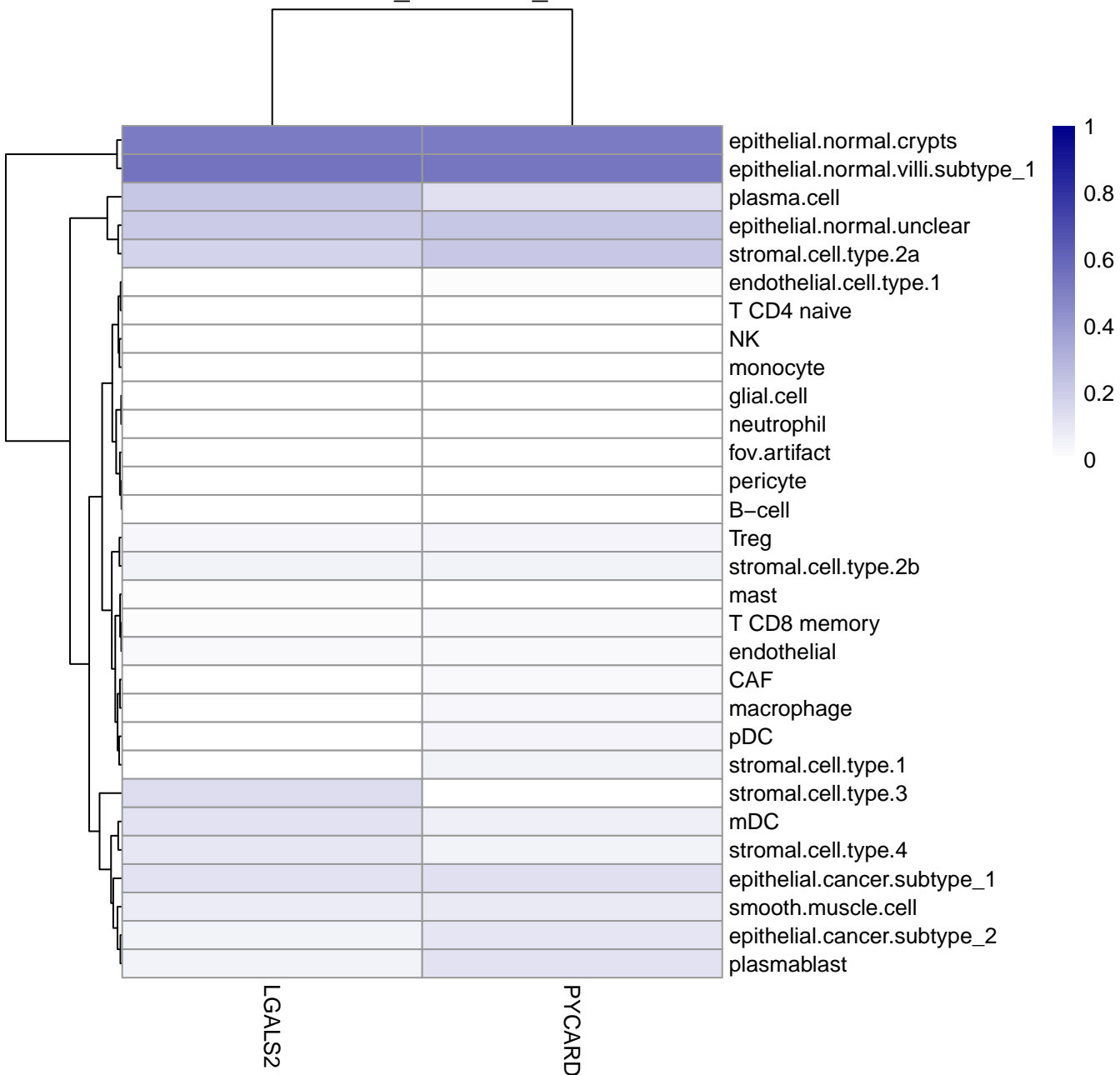
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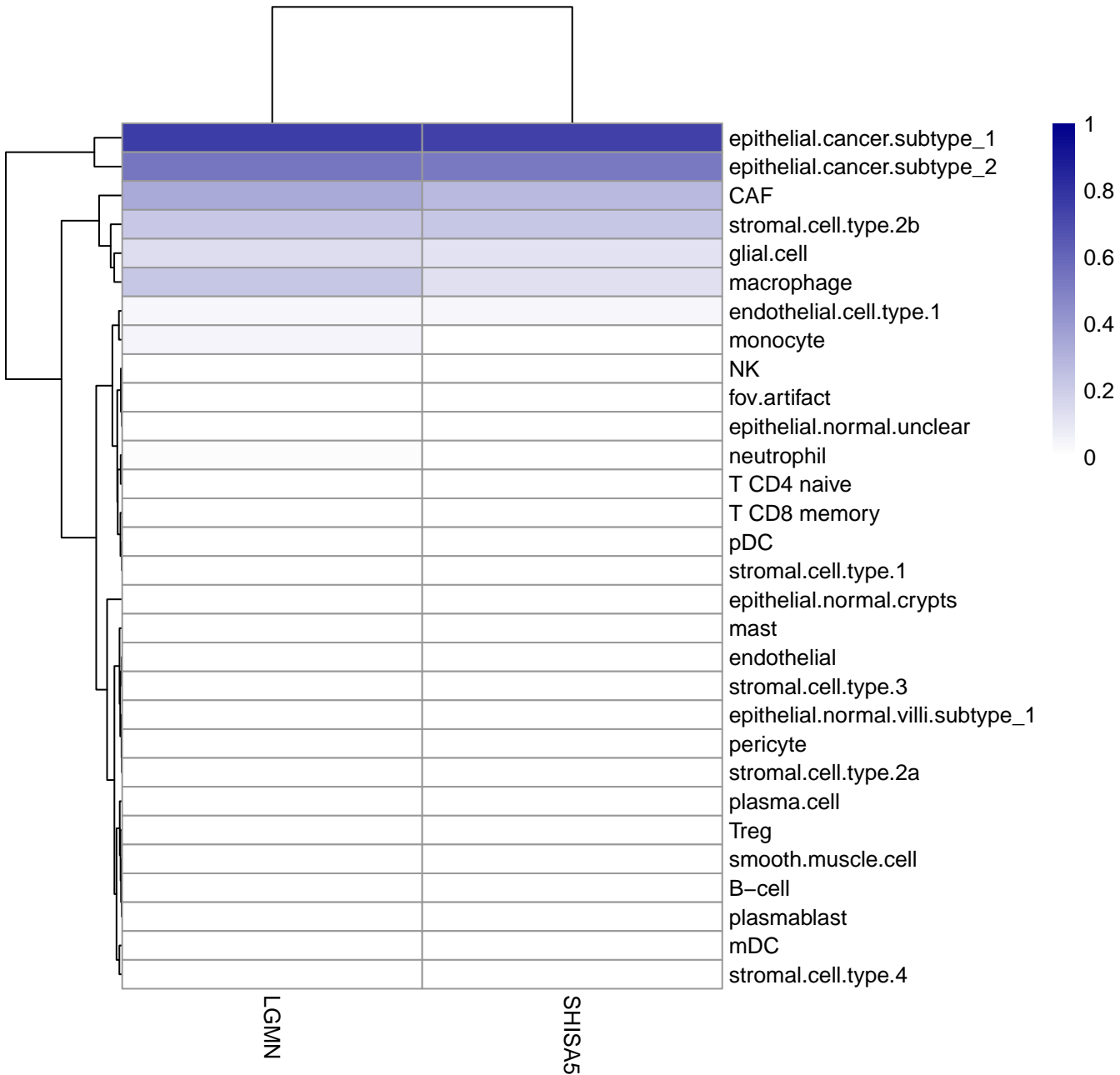
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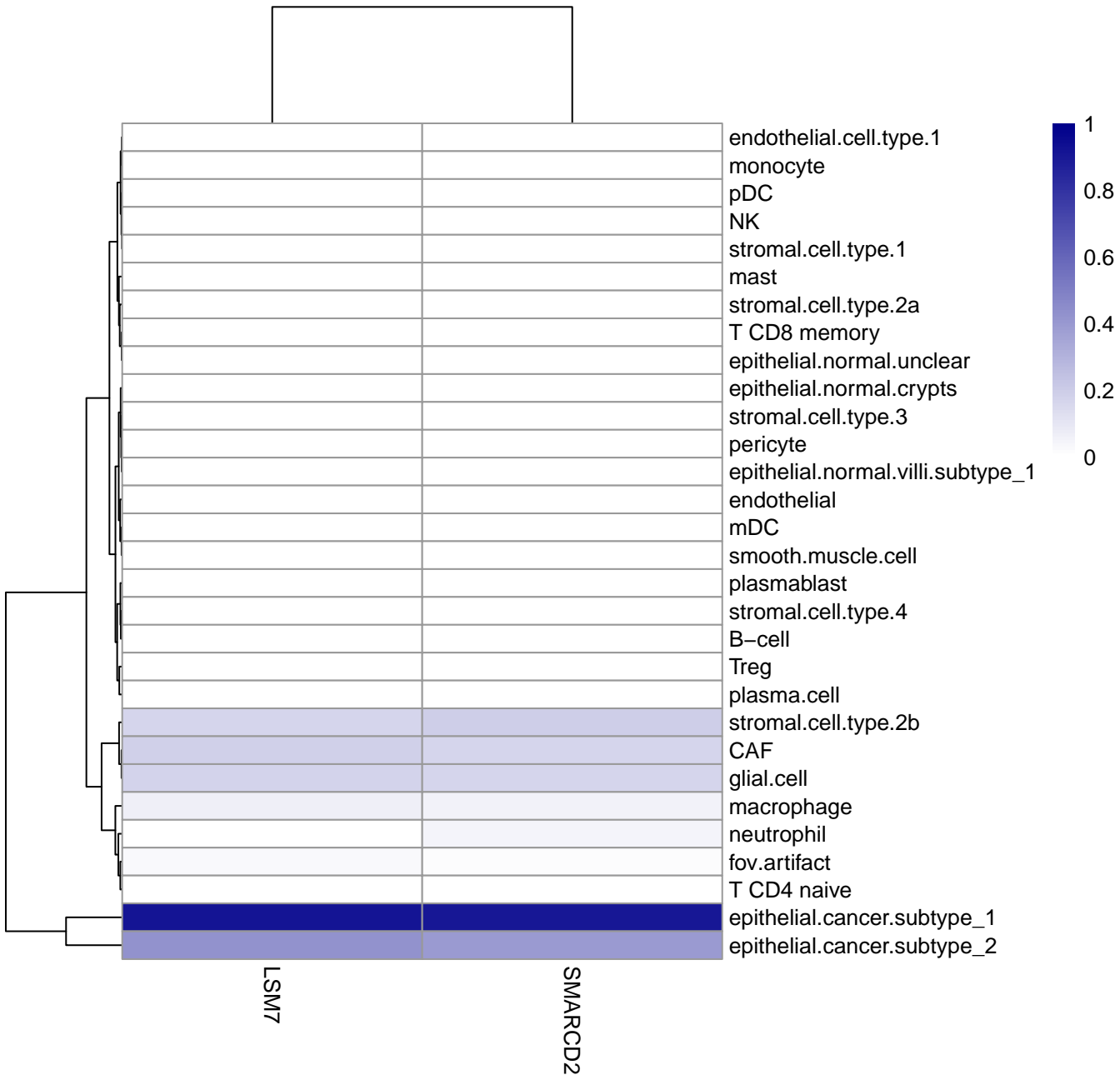
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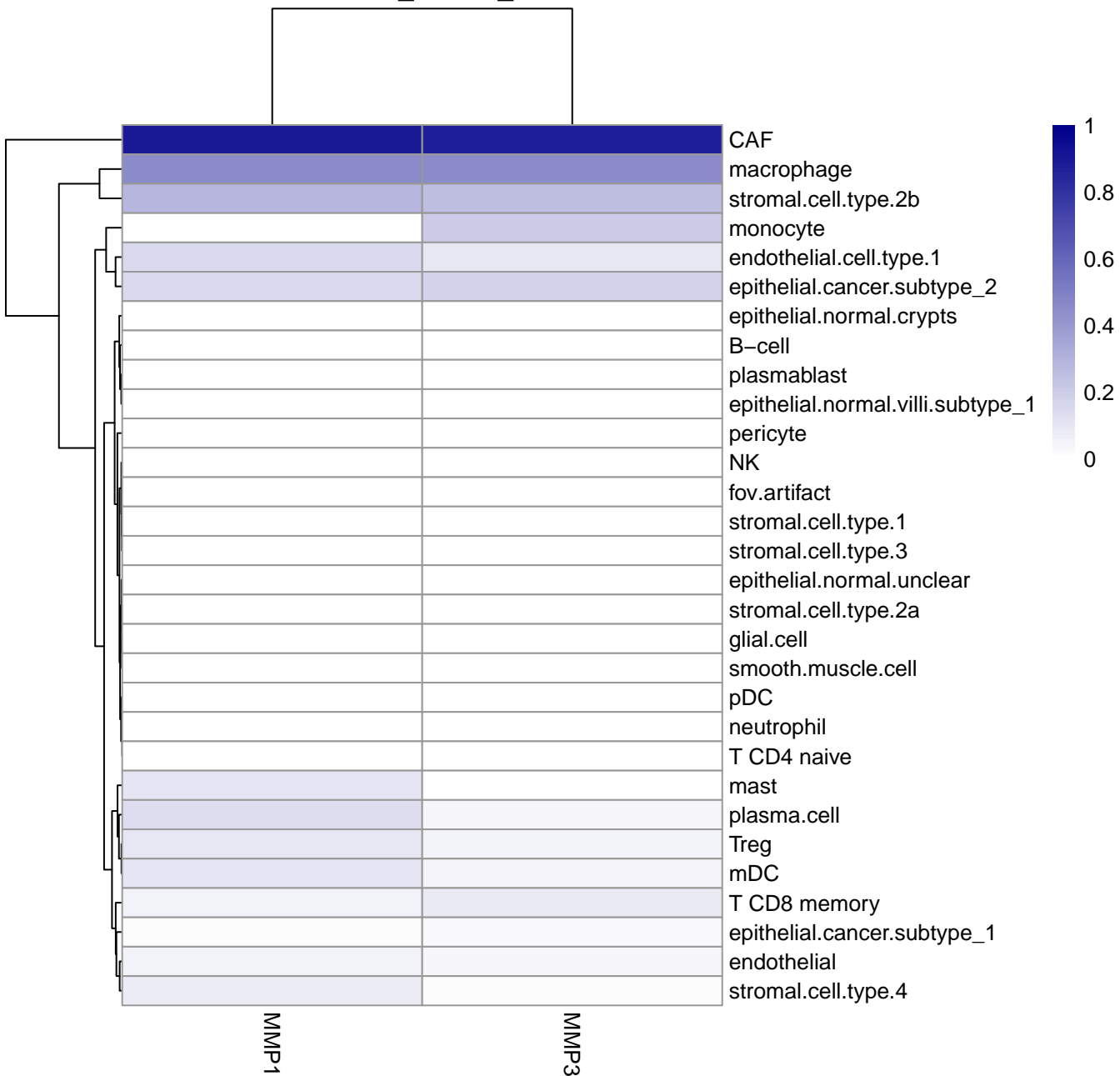
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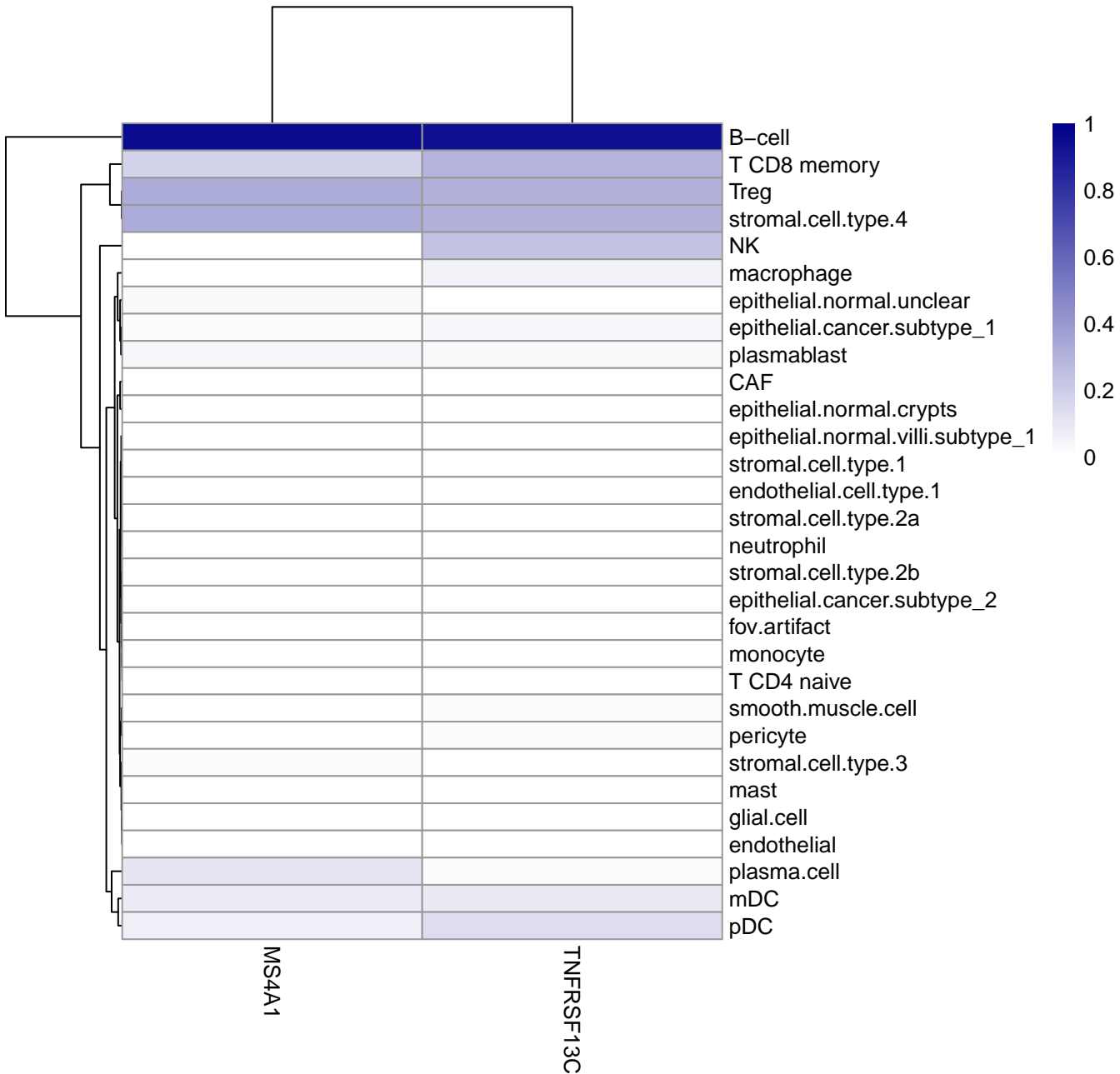
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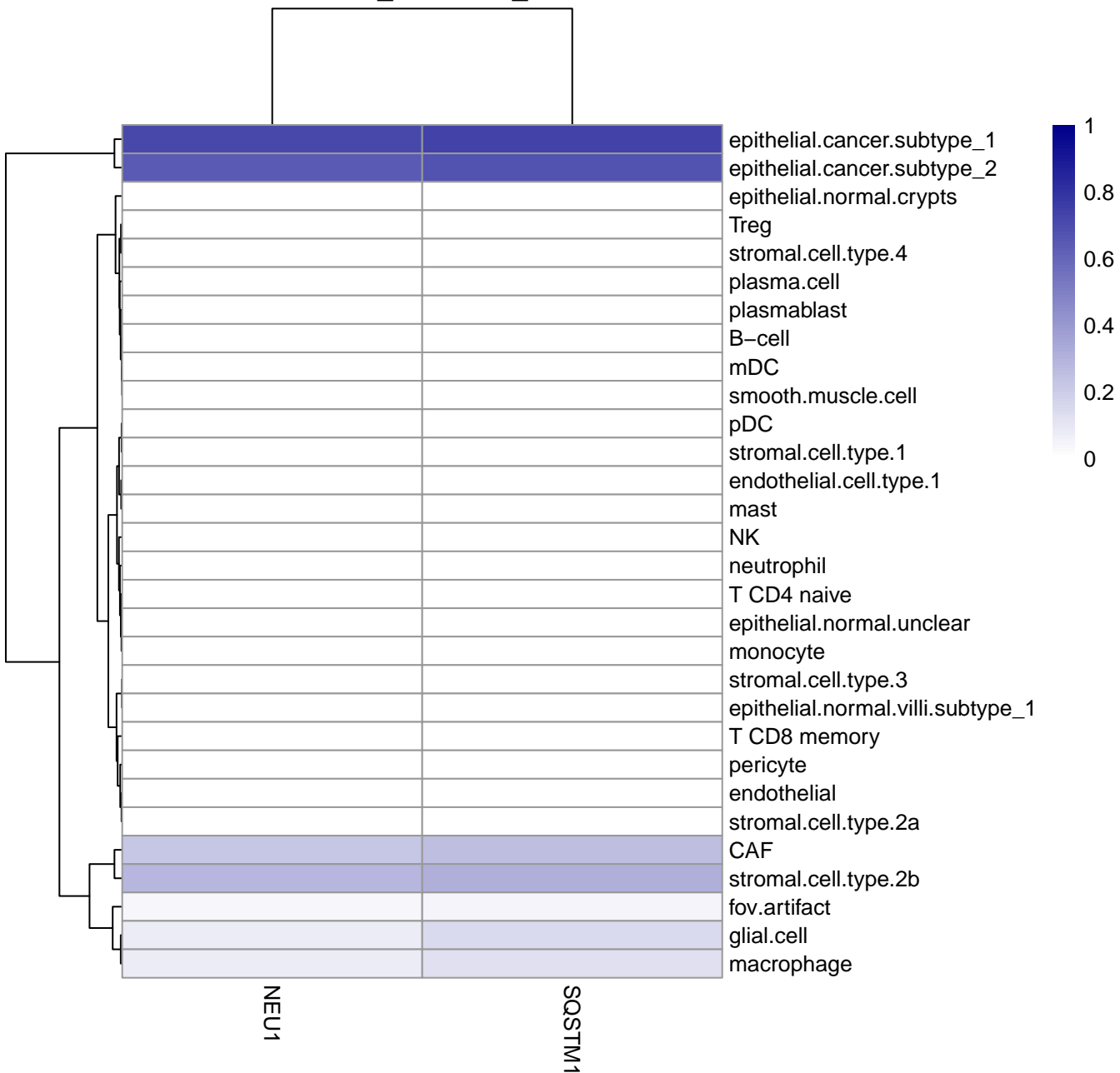
MMP1_MMP3_2



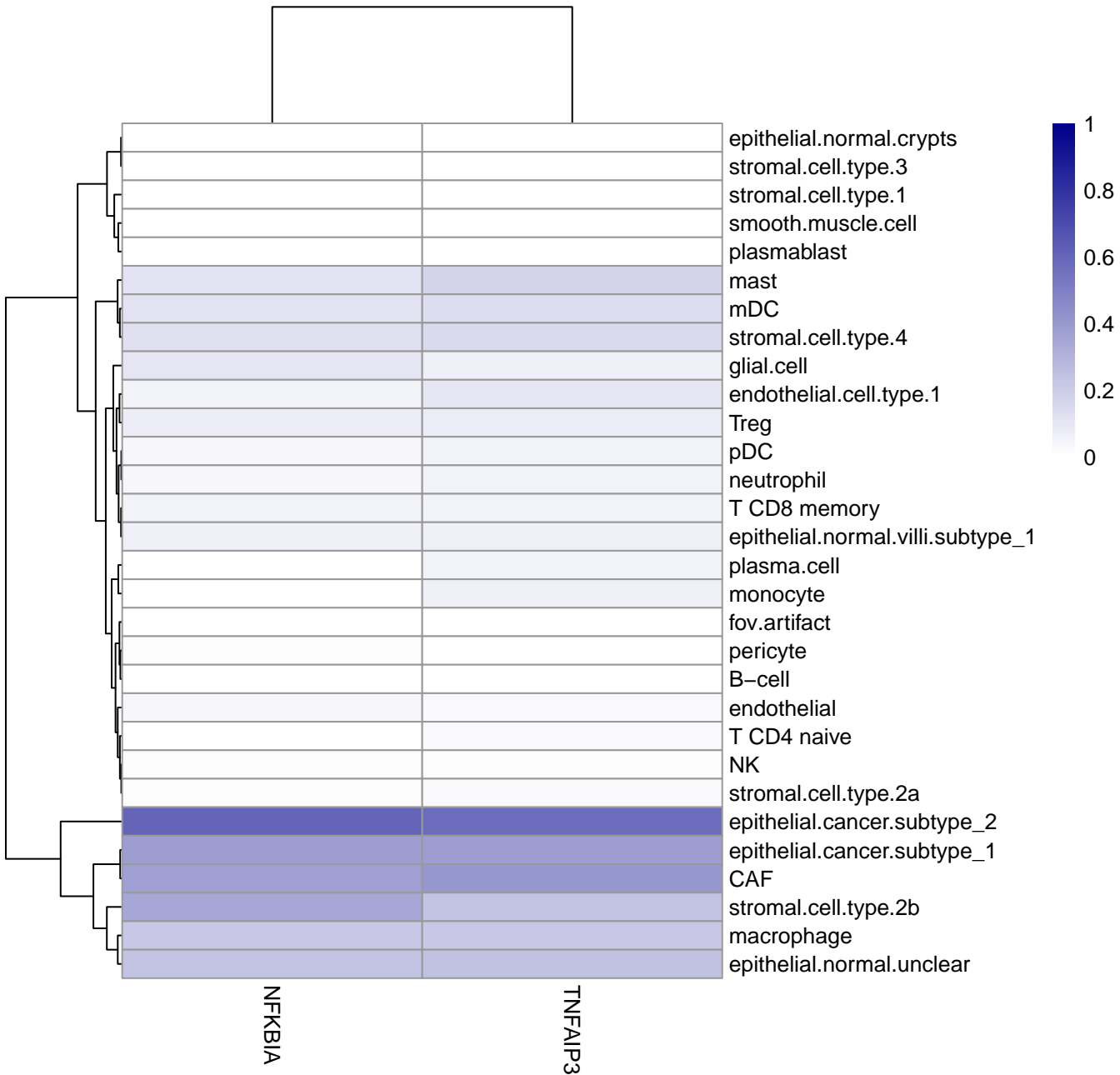
MS4A1_TNFRSF13C_2



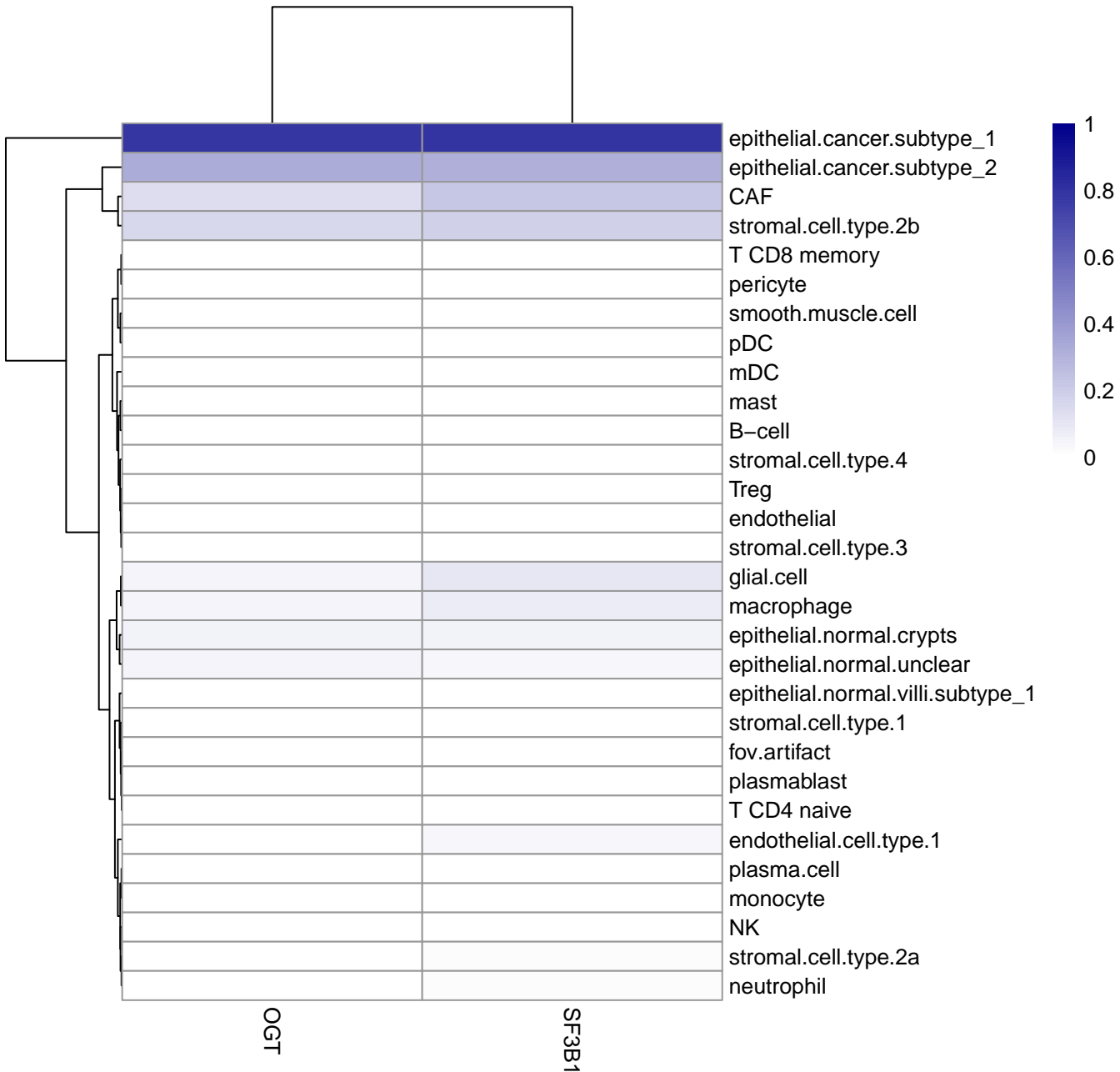
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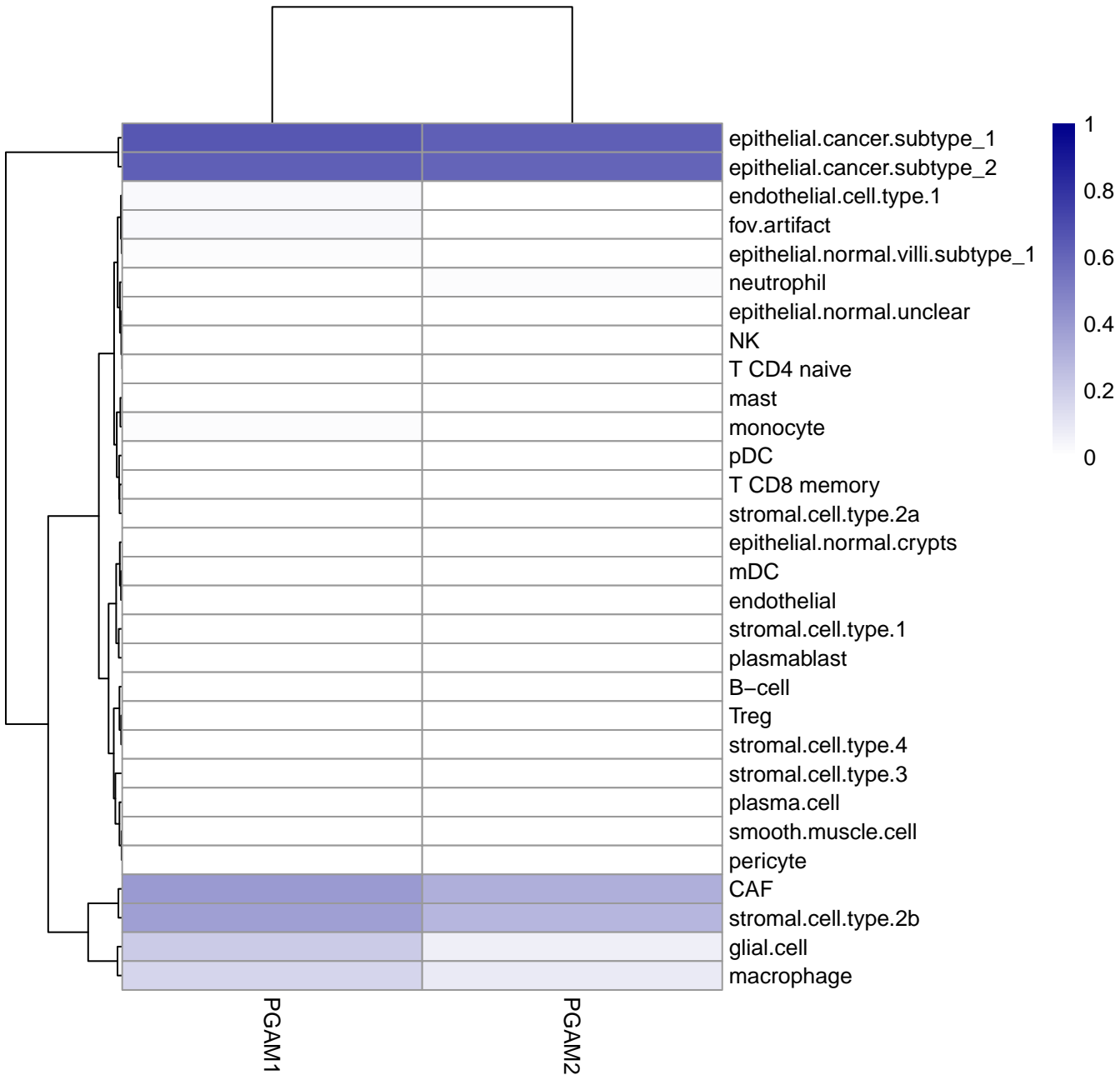
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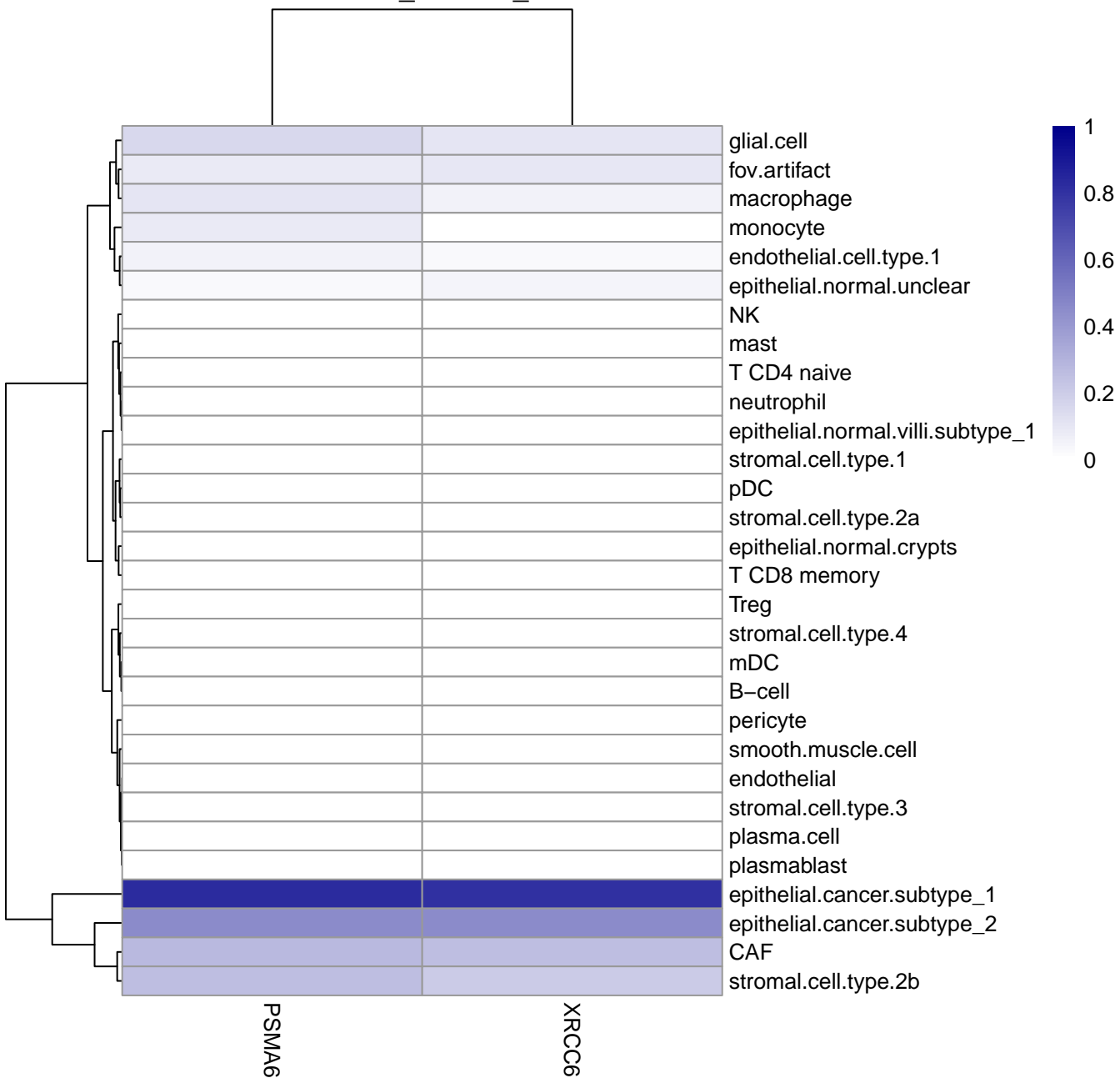
OGT_SF3B1_2



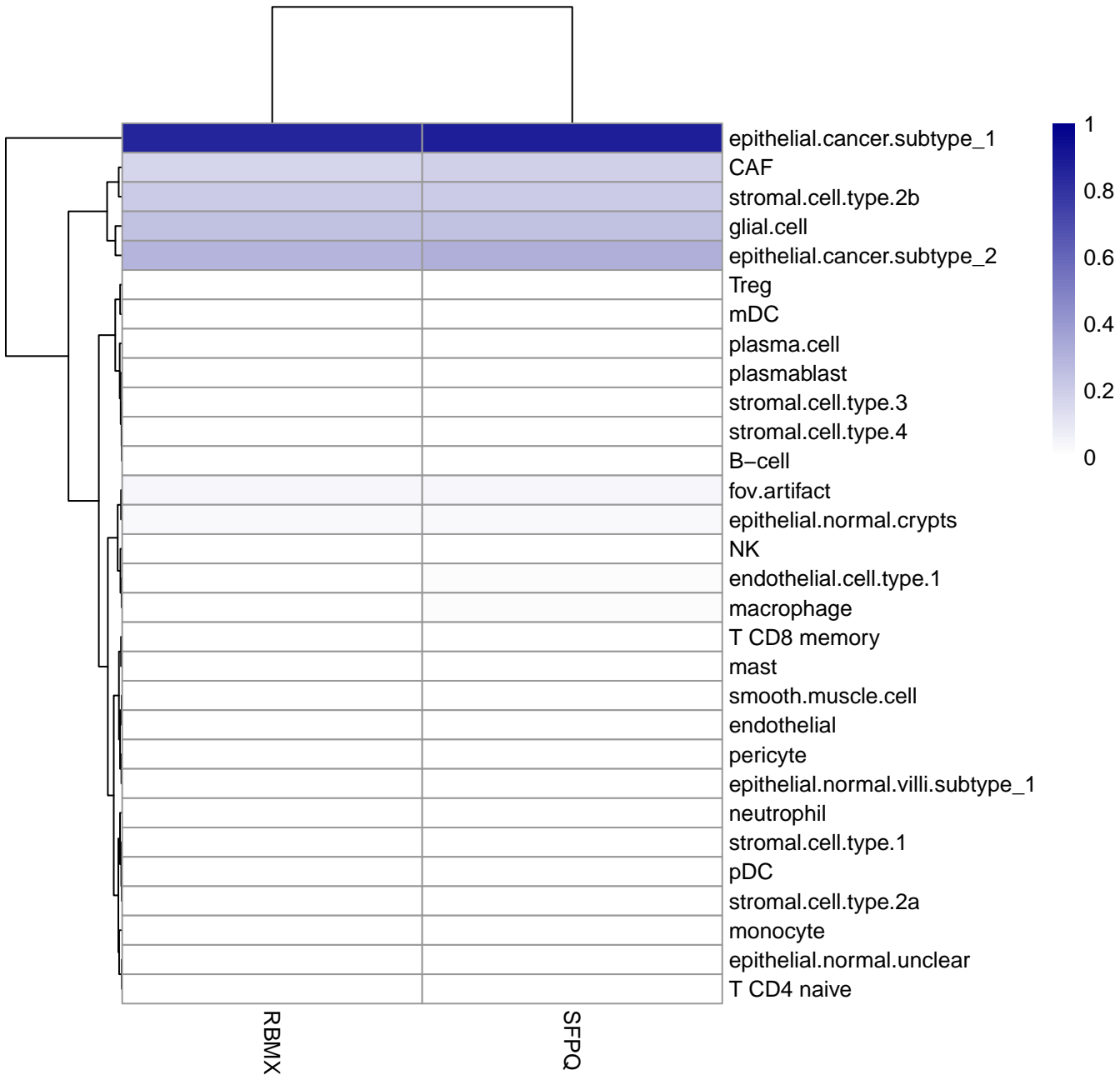
PGAM1_PGAM2_2



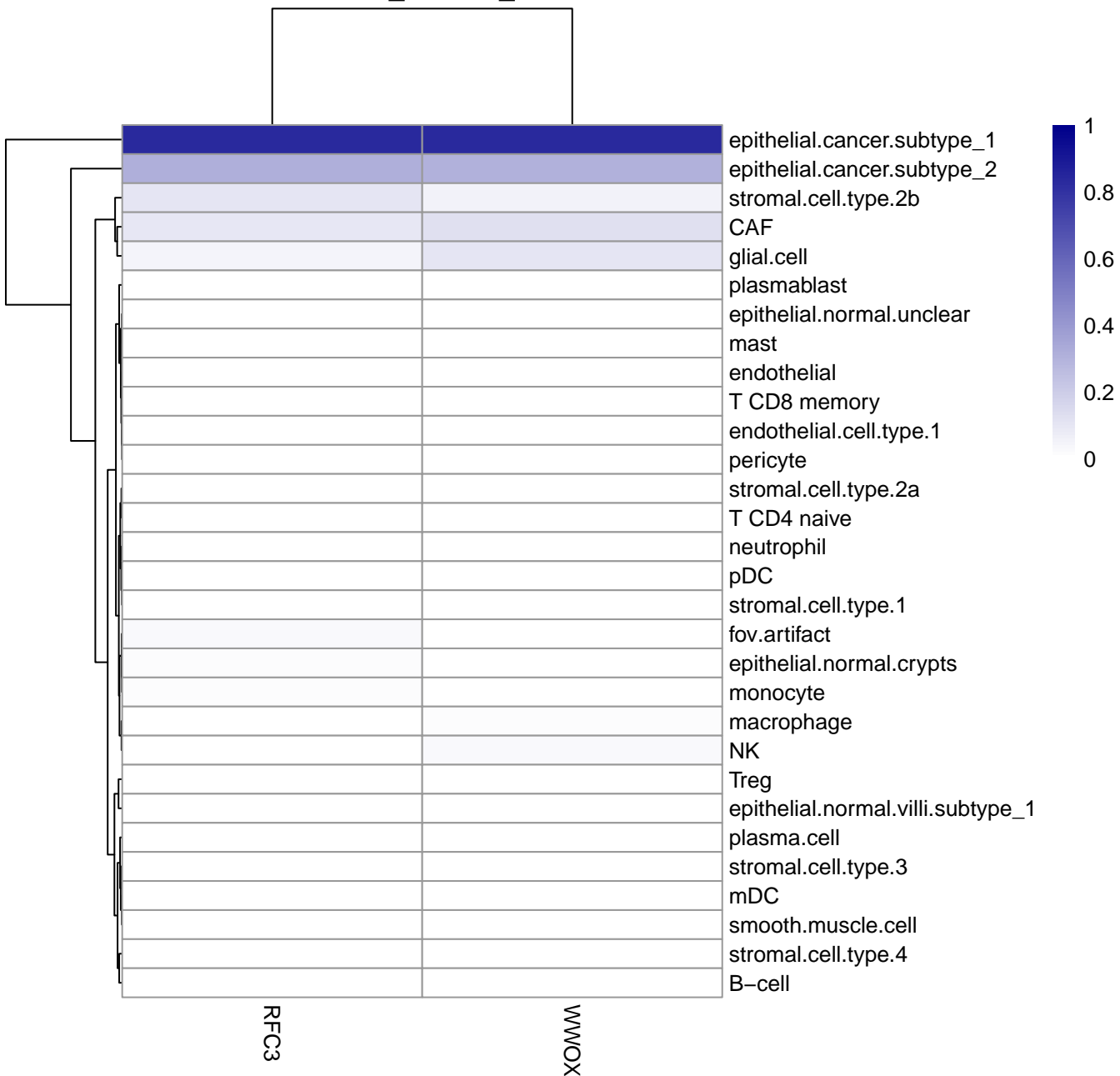
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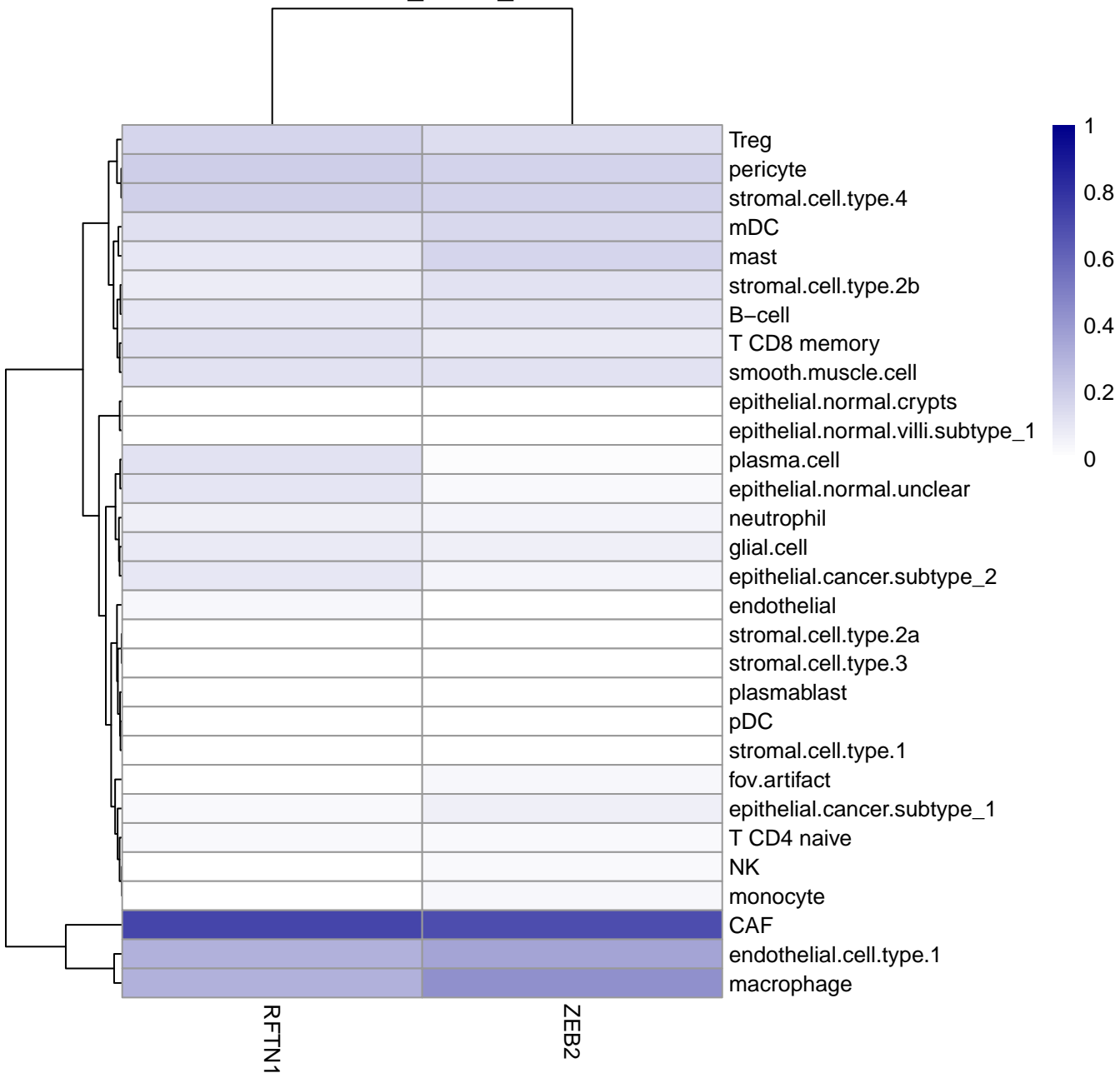
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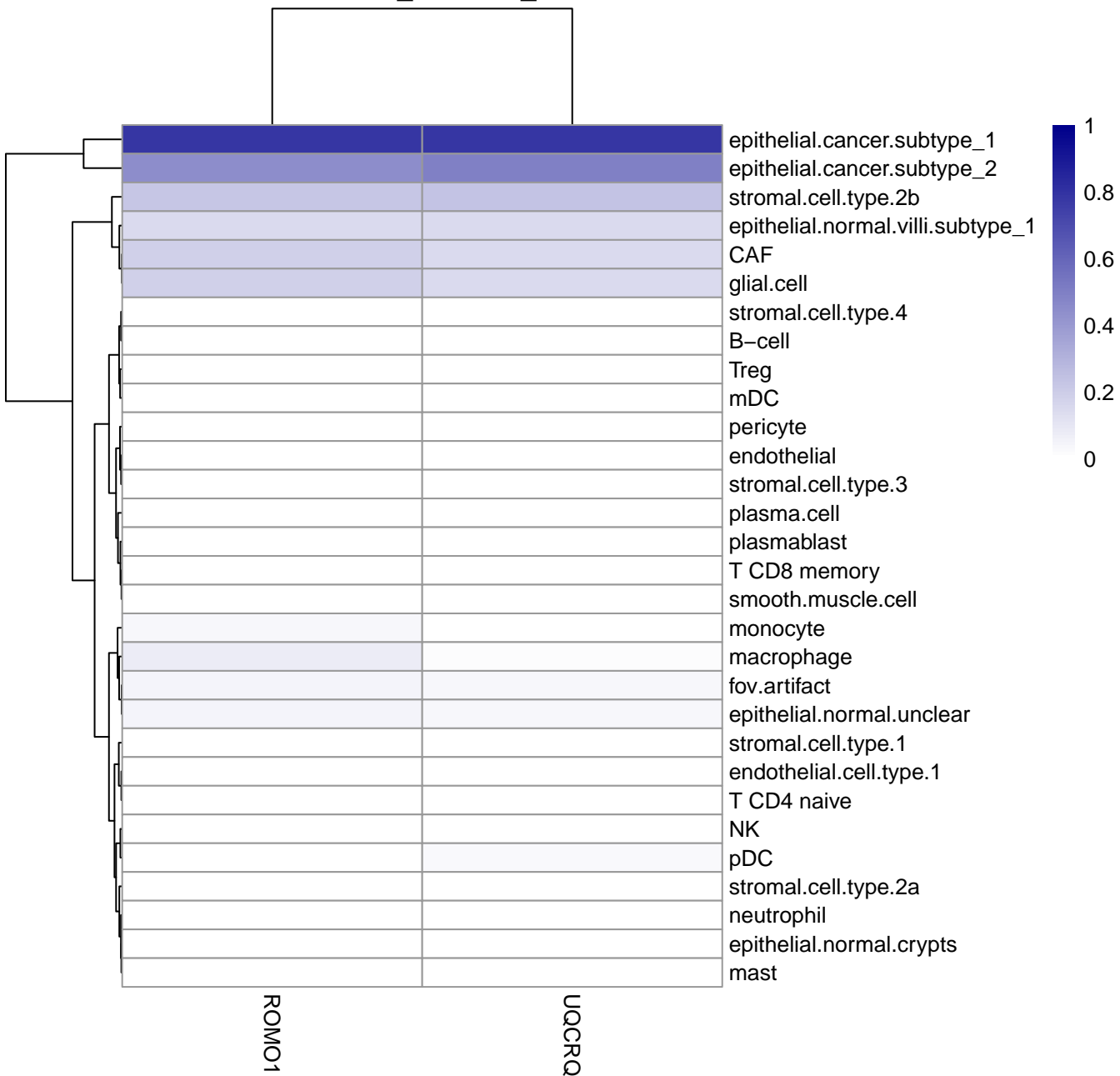
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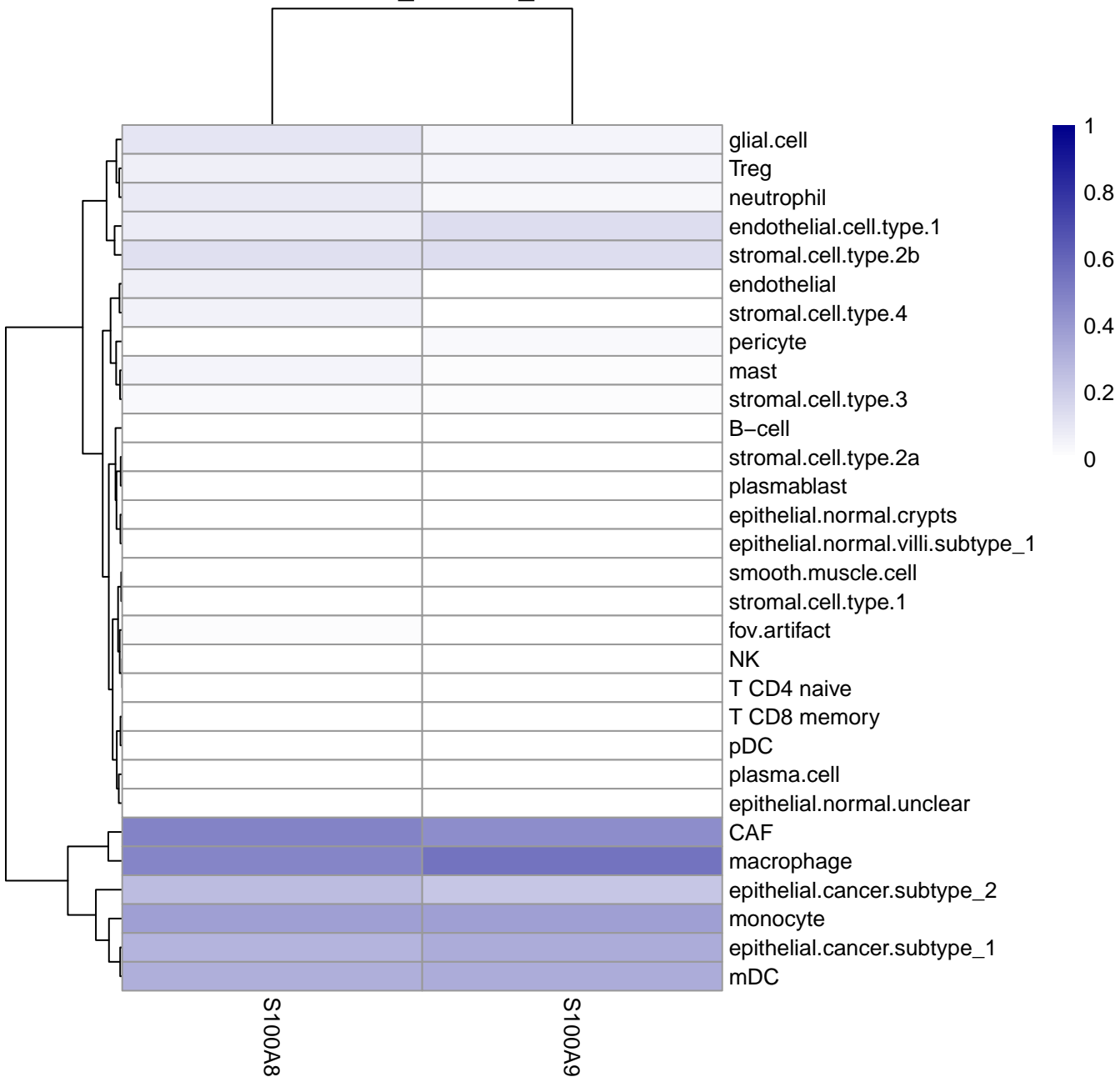
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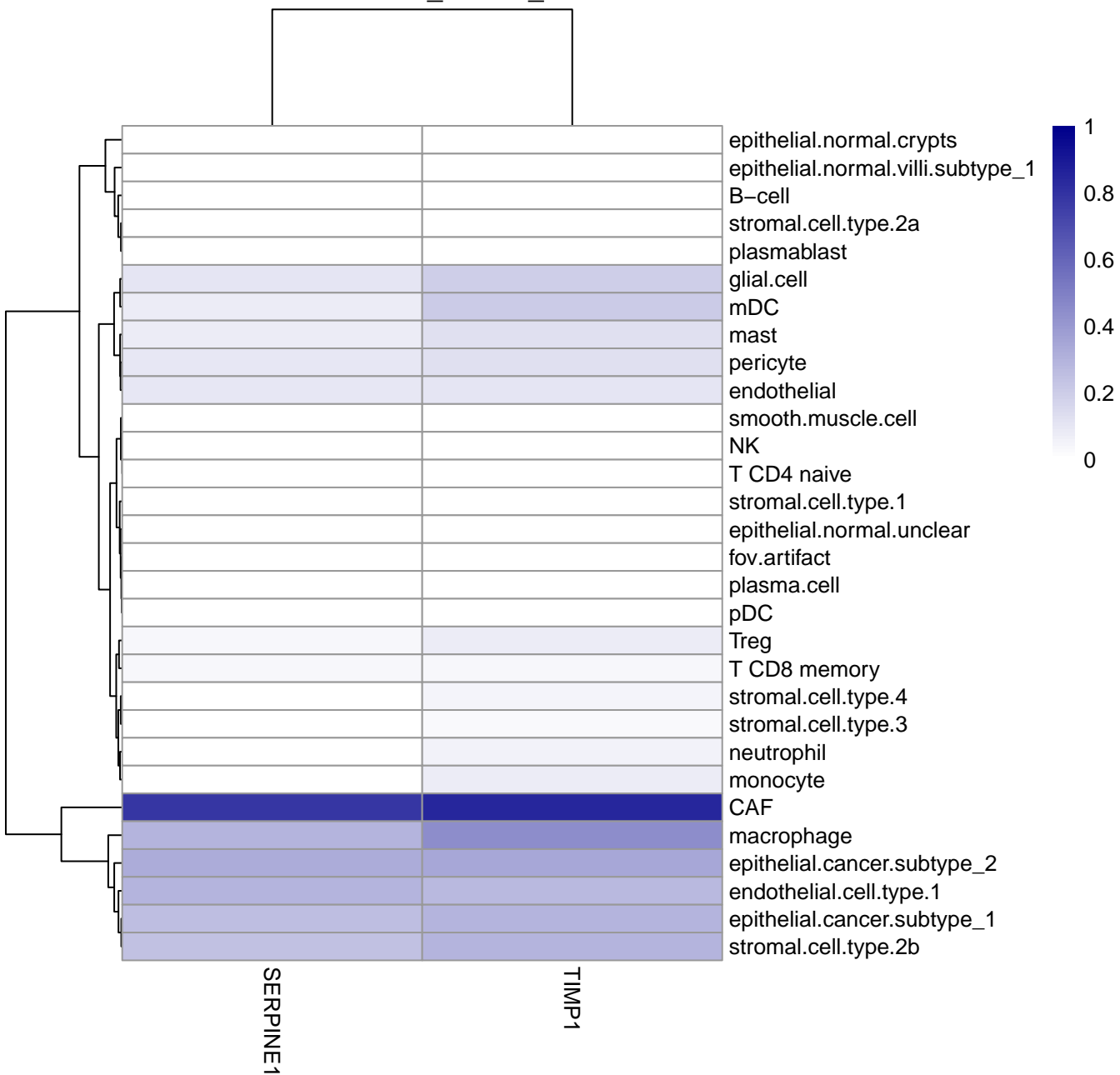
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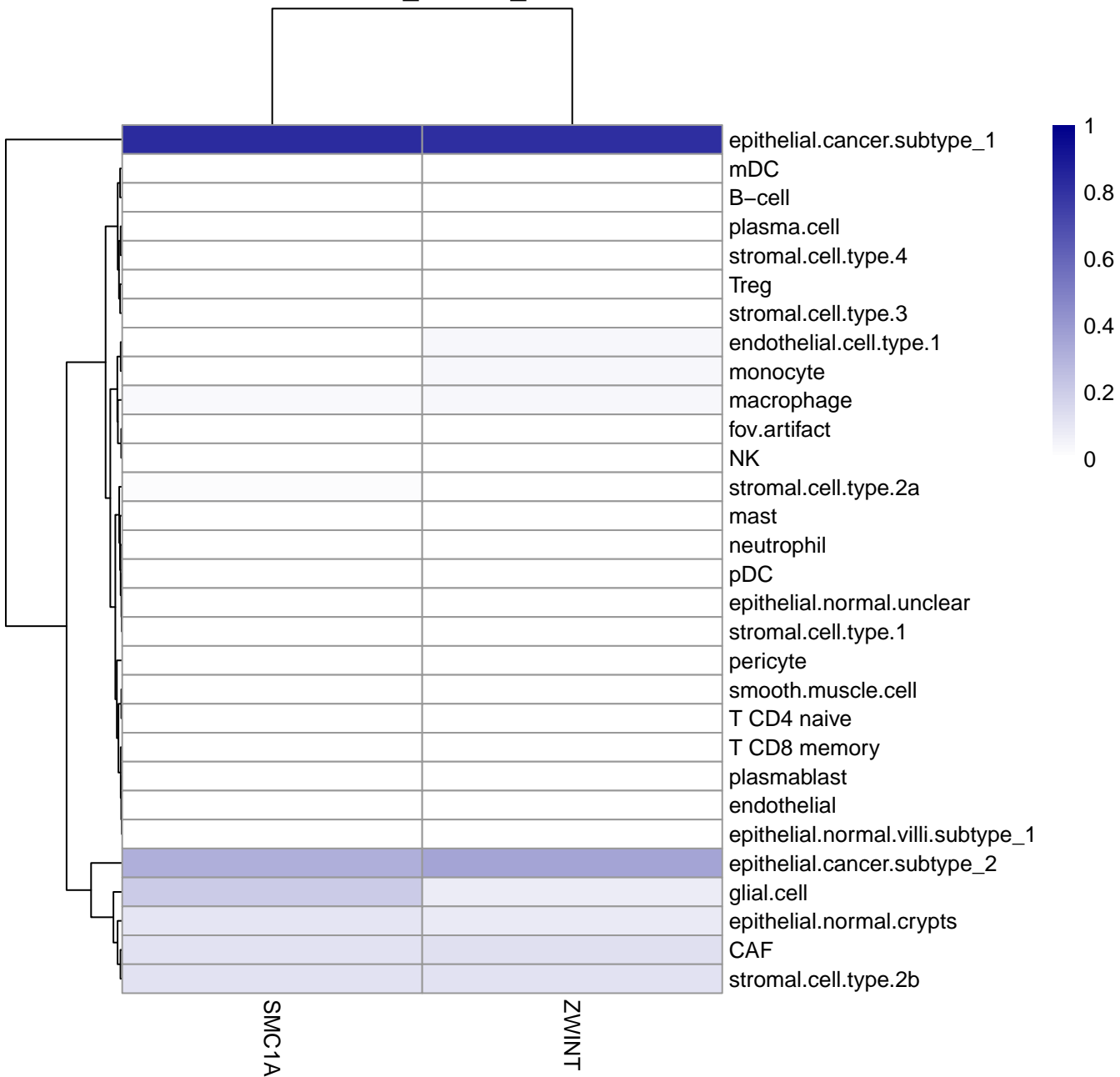
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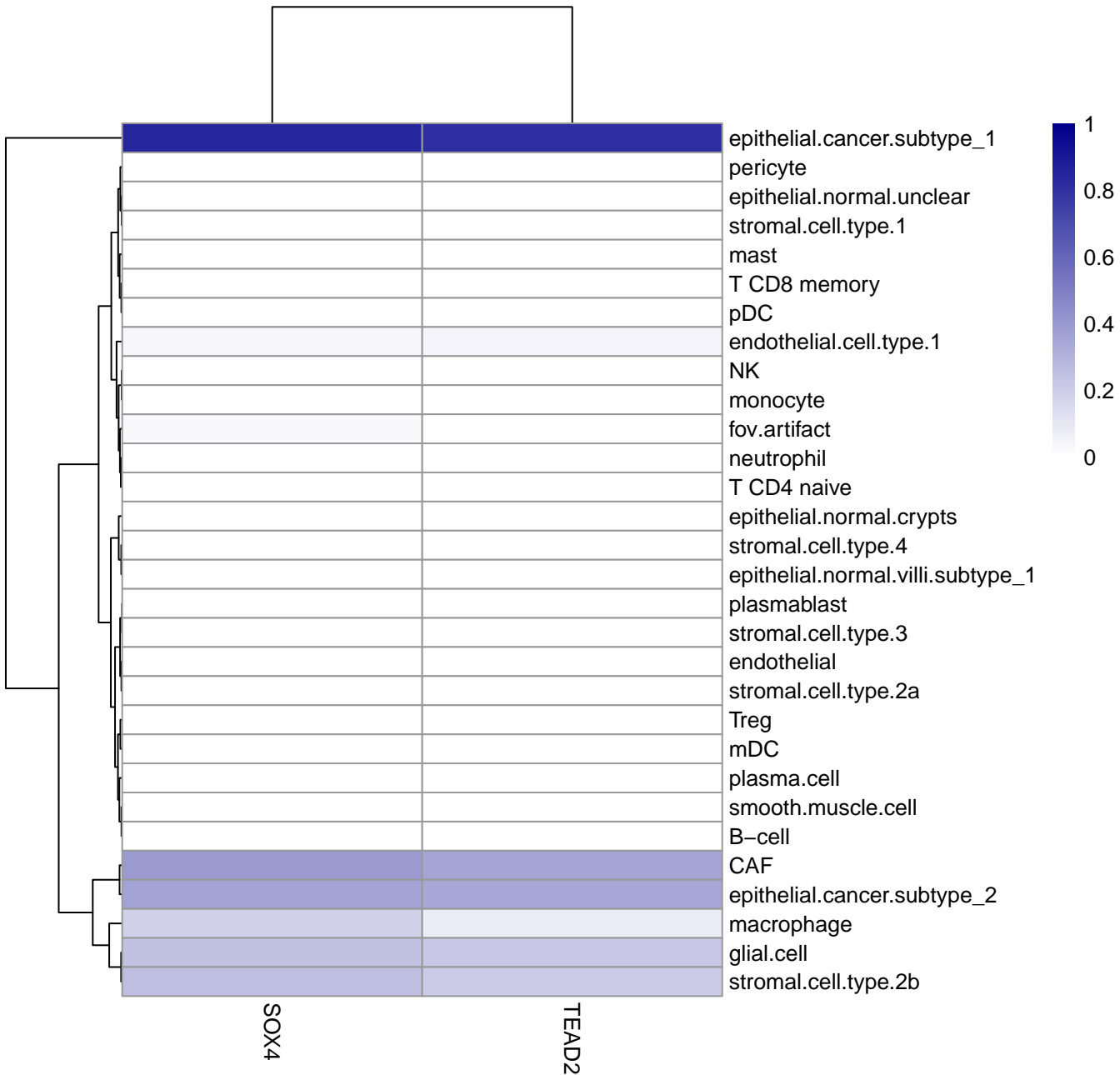
SERPINE1_TIMP1_2



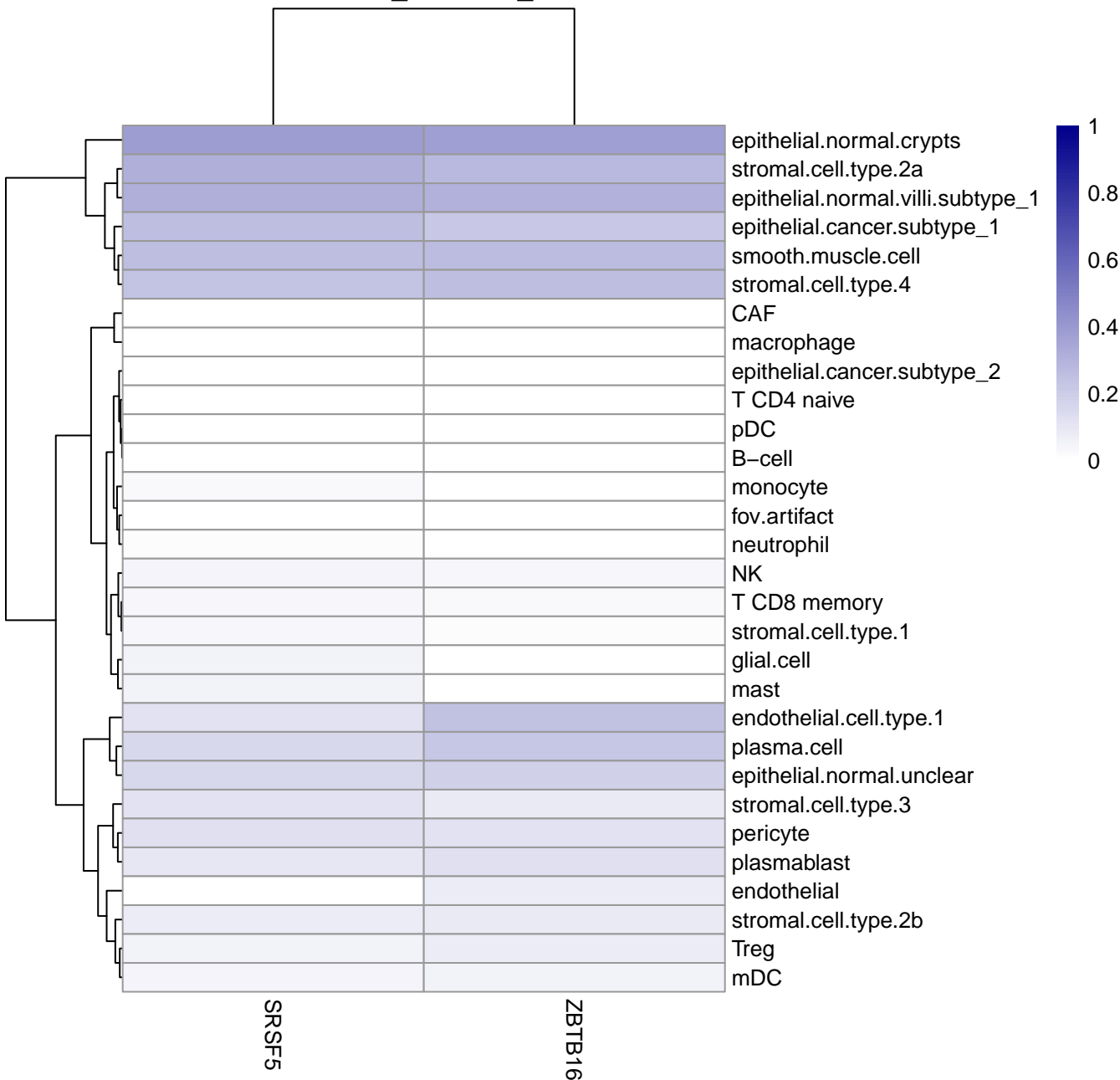
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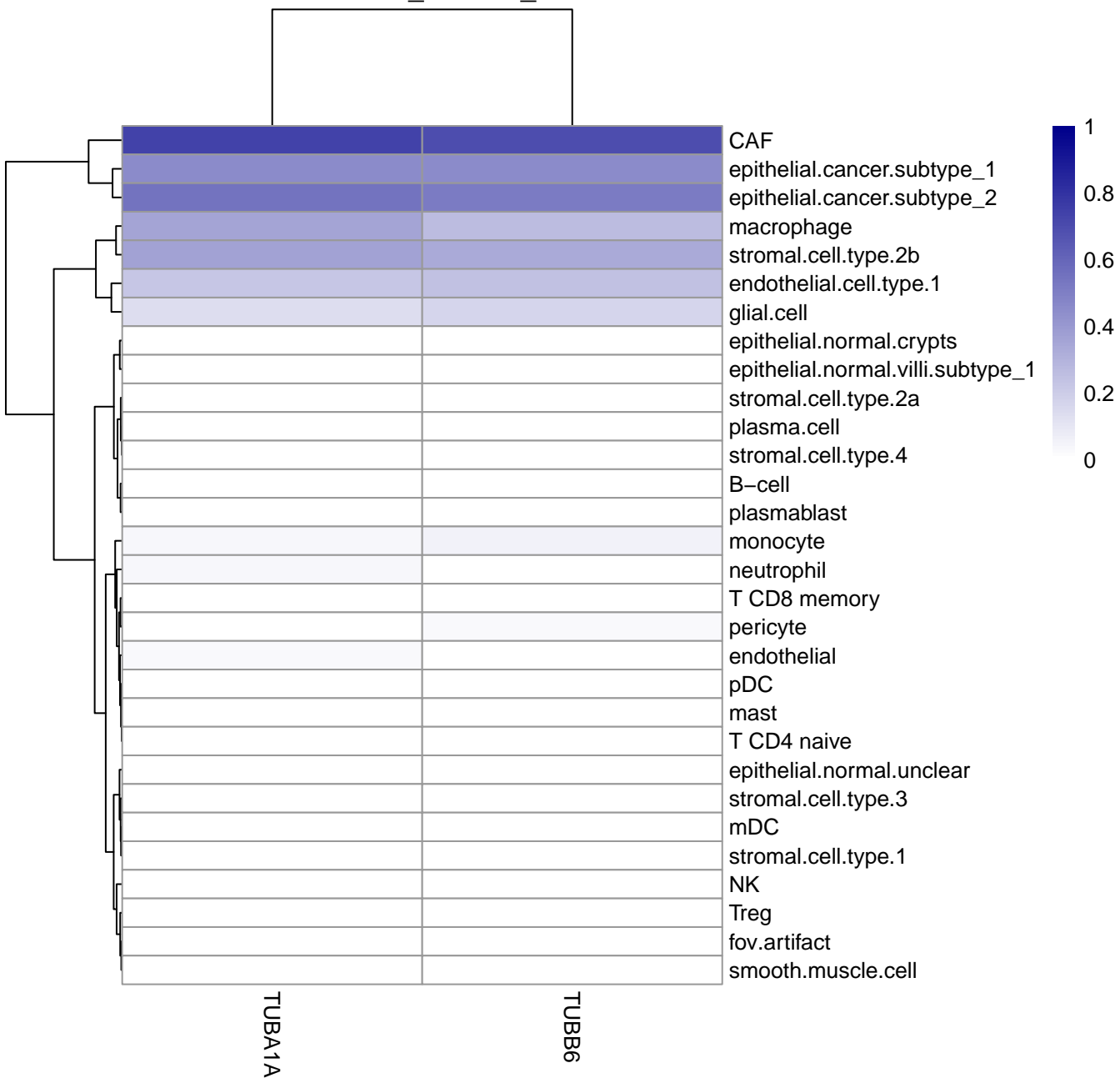
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