TREML2\ APBB1IP AIRE DOCK2, FUT7 CD8B CD1C WNT1 CD3G BCL11B
CD7 GPR18 RUNX2-TCF7 SPI1 NLRC3 CD7 \ GPR18, \ CASP8 JAML ITGB1 CCR9 CCR6 CD3D PIK3CD CD79B FLT3

LCP1 CLEC7A SLAMF6 TNFSF8 RHOH BATF PTK2B

ISA1 SLA2 GSN NLRP3 CR1 TNFRSF13B TCF3 ROU2F2 PIK3CG WAS BTN3A1 SIT1 IL23R SOC\$1 IRF4 TNFRSF18 DCLRE1C CPR183 C17orf9 CCDC88B CD300A PTPN22 PTPRC VAV1 ABL1 GPR183 C17orf99 NOD2 CD5 BTLA HLA-G 1L23A CCR2FANCD2 FANCA CR2 BLNK CSK XCL1 CLECHIIIactivation NCKAP RUNXBOXXIPPhocyte differentiation TNFSF11 HLA-DPBRIPOR CARD11 LCK SPN CD74 -HLA-DOA IKZF1

TIGIT PDCD1 TNFAIRBATA3 IL10 CD83 JAK3 CD2-LGALS9 LYL1

KLRK1 RAEGILLABON OF T CEIZACRIVATION IL7 IL12RB1 PLA2G2D PLCG2

GRAP2 regulation of T ceizacrivation IL7 IL12RB1 PLA2G2D PLCG2

HHLA2 TNFRSELIGG INVARIANCE CONTRACTOR INTERPRETATION INTERPRETA 103 116 128 HHLA2 TNFRSFIA HIMB DEMEM131L7 ADAMSCRICTLES LAMES CD79A 141 SKAP1 SELL ITGB7/ DOCK8 / CD6 ITGA4 ITGA5 ITGB2 LST1\ HLA-F STK10 IGLU1 SAMSN1 FERMT3 IGLL5 CD22 TNFAID BANK1 TNFAIP3 MZB1 ADTRP-SEMA4D TNF SELPLG