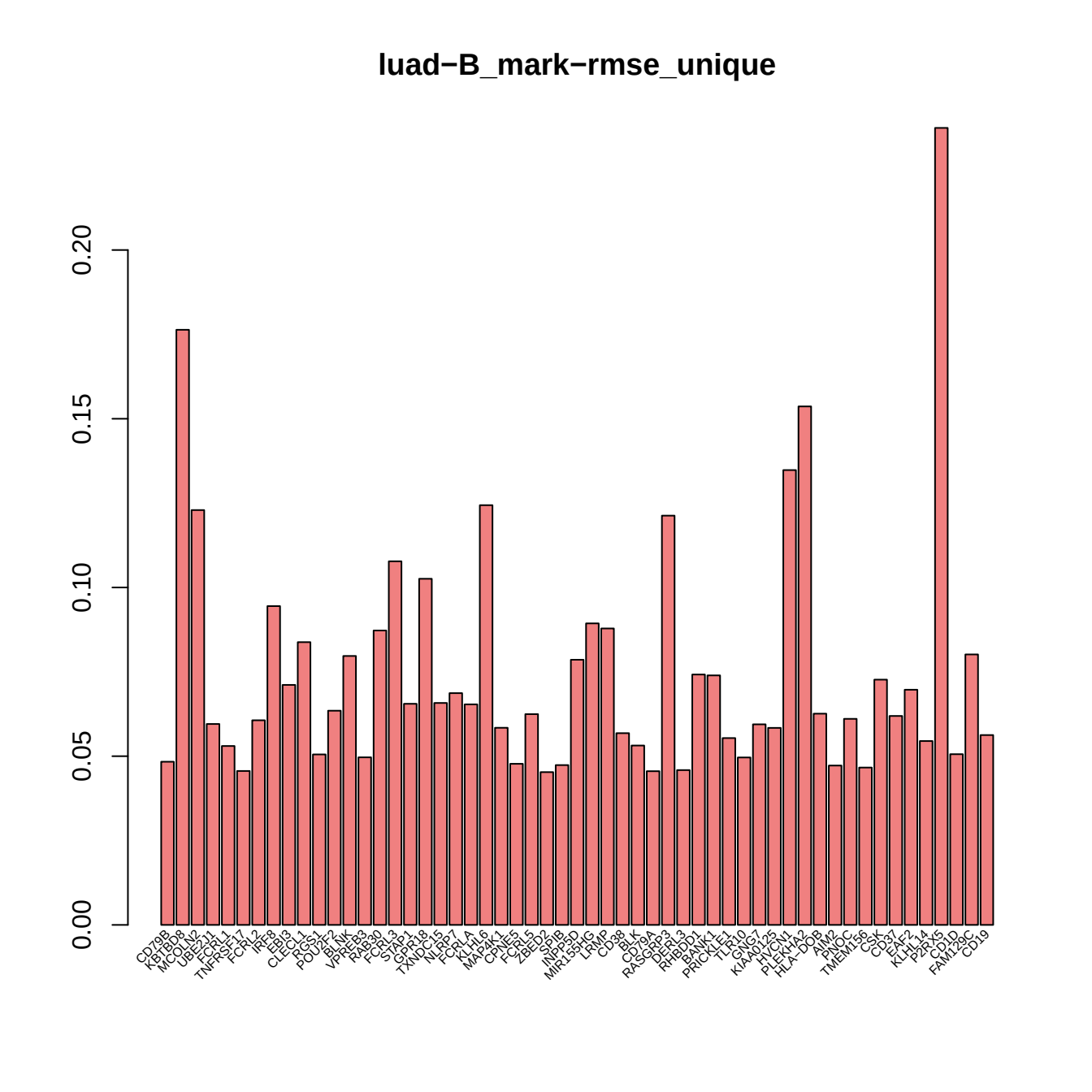
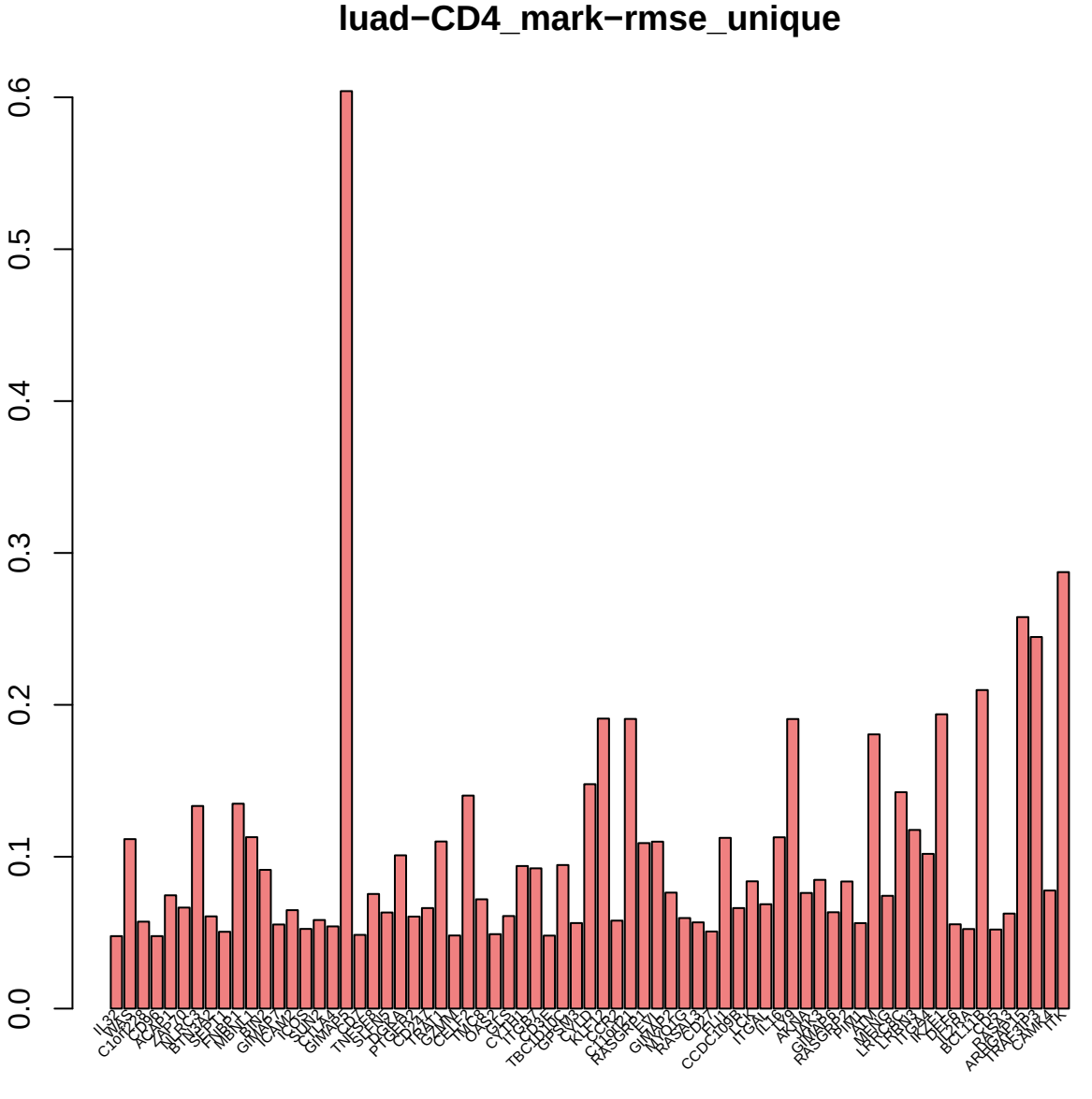
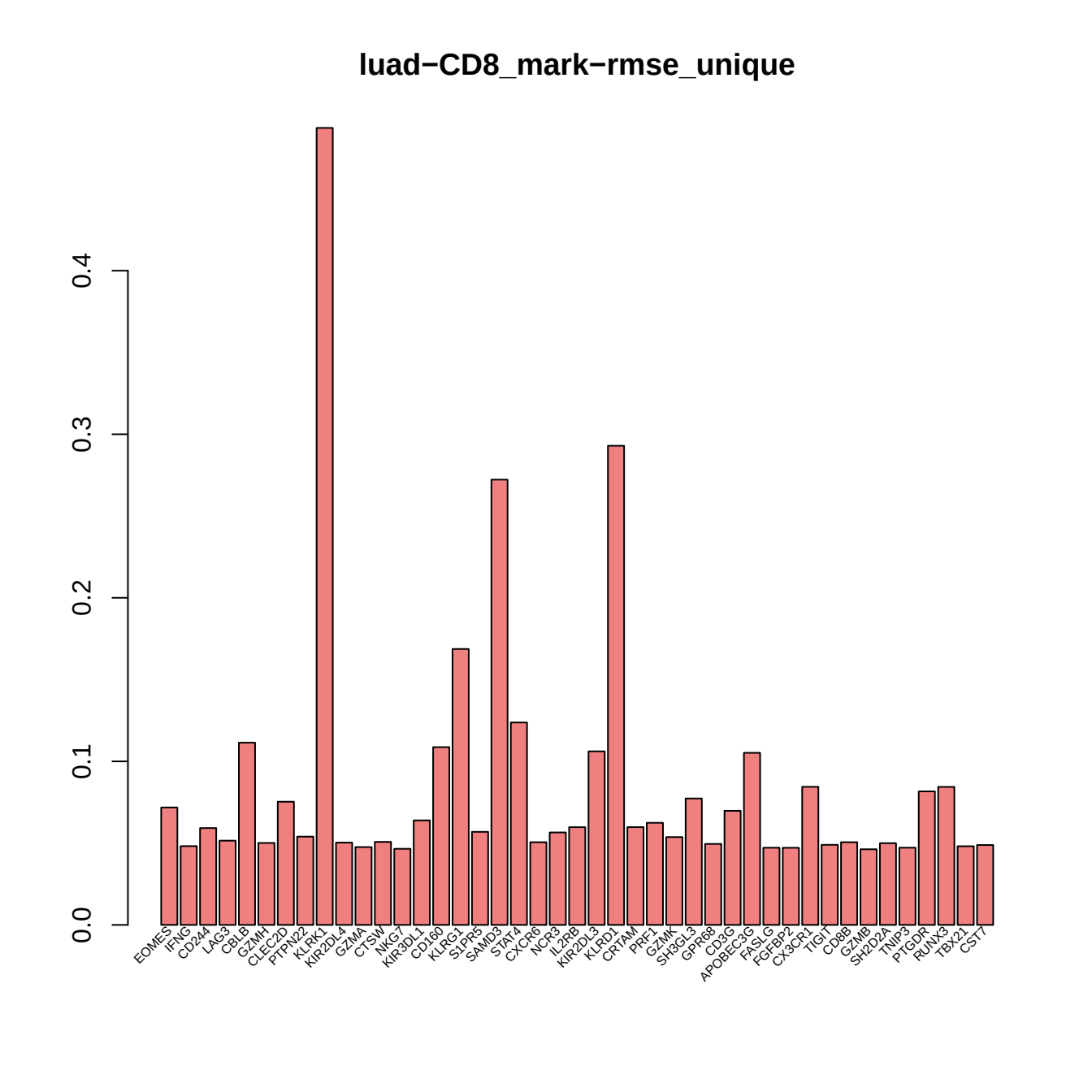
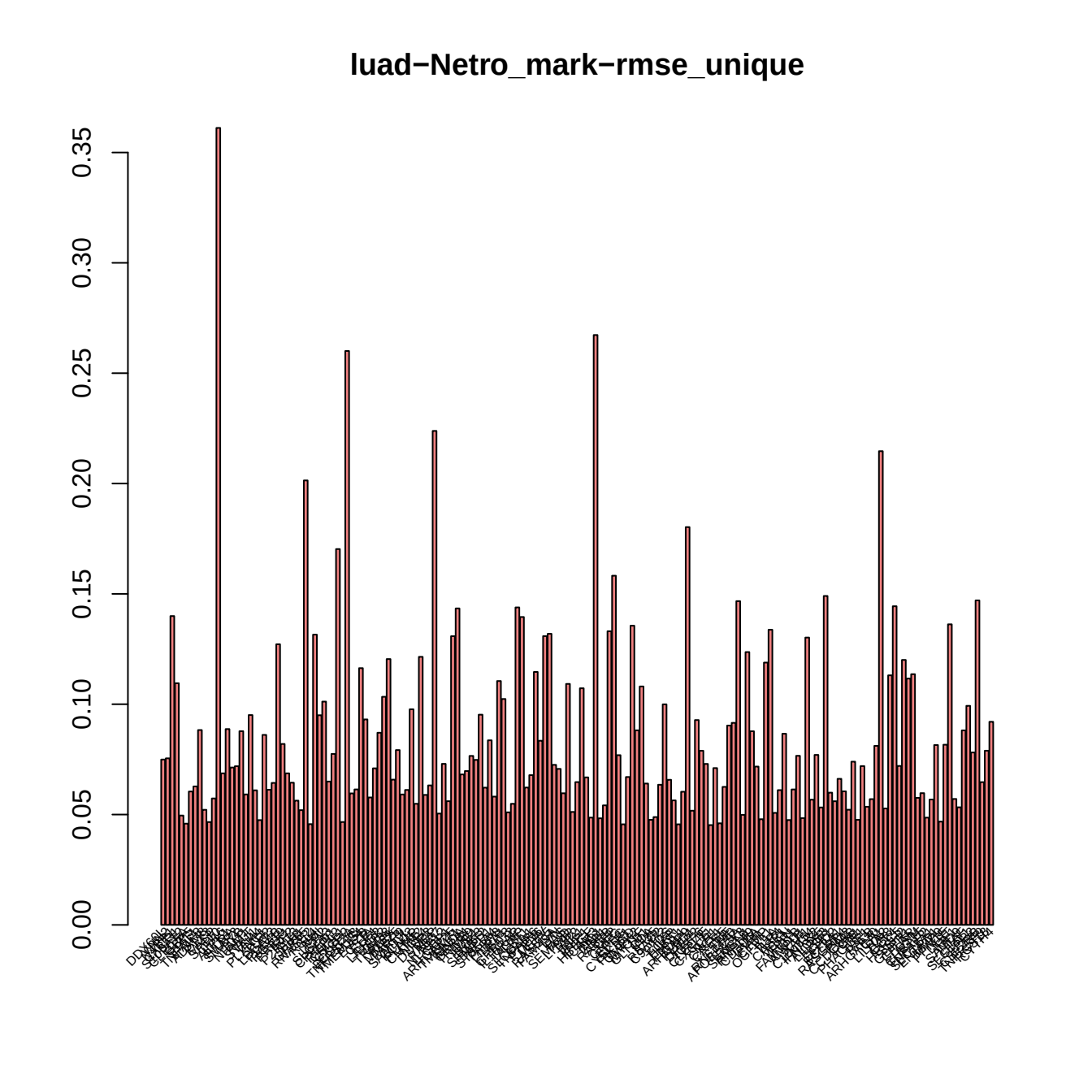
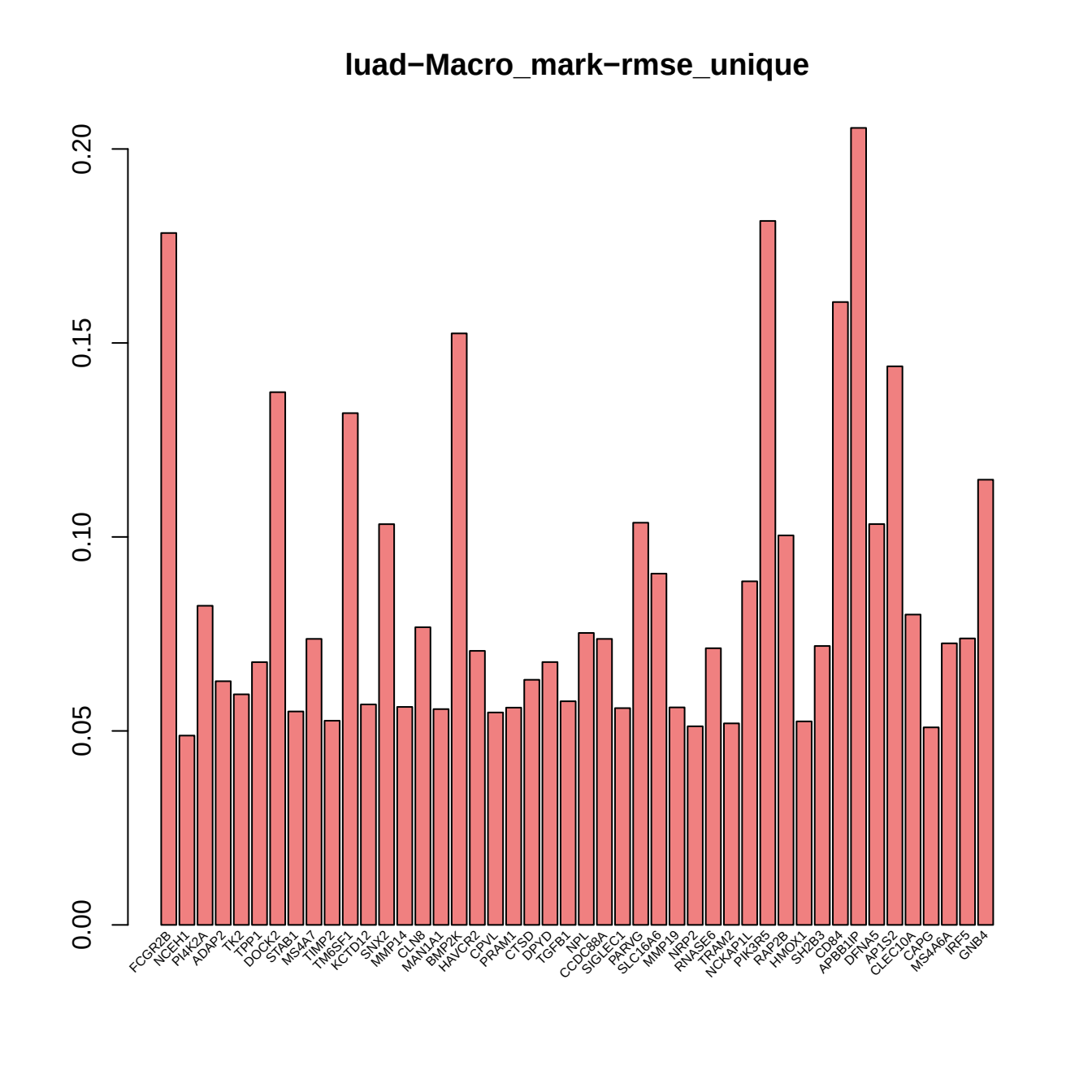
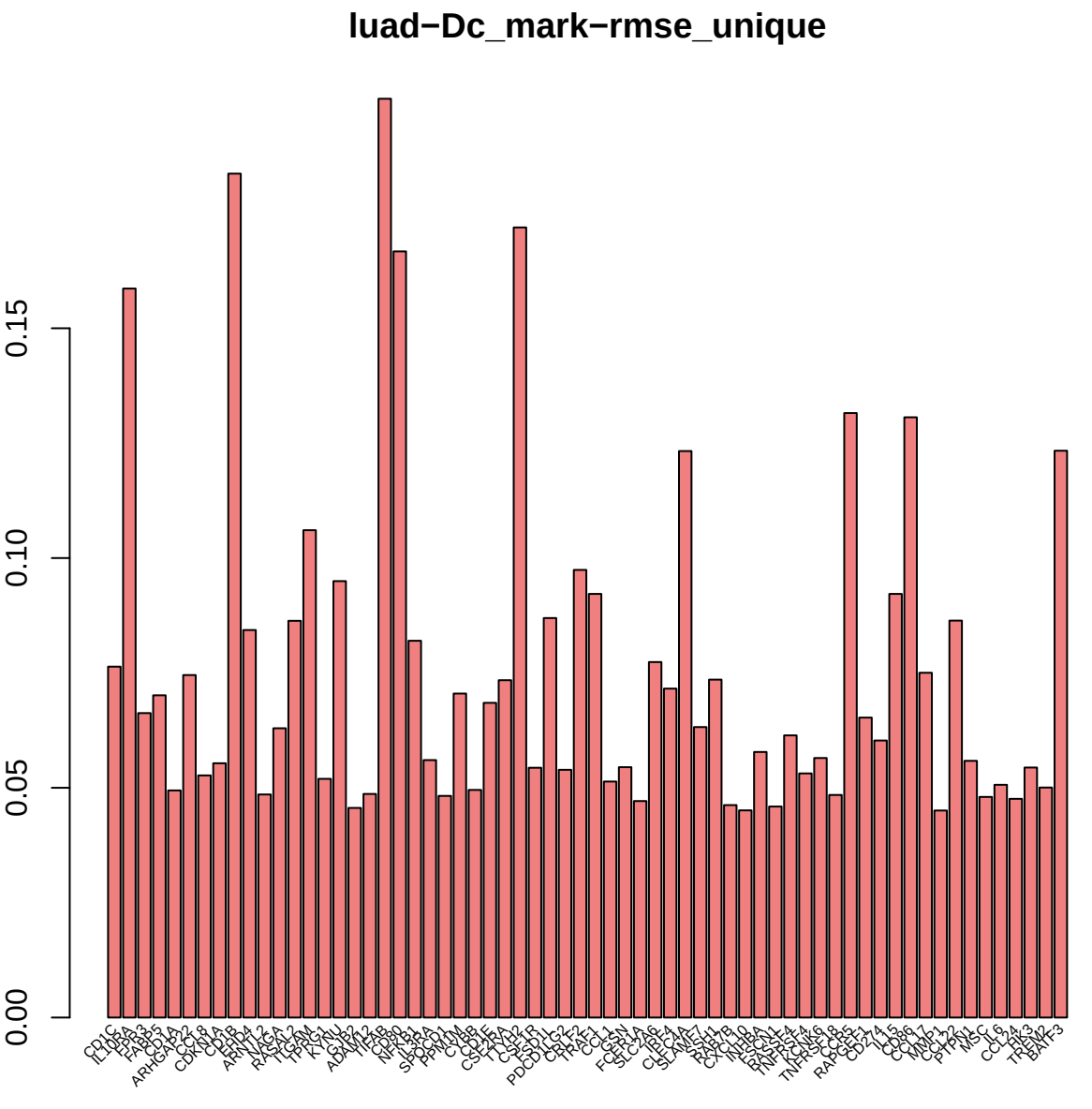

luad−B_mark−rmse_unique
0.00
0.05
0.10
0.15
0.20
CD79B
KBTBD8
MCOLN2
UBE2J1
FCRLA
TNFRSF17
FCRL2
IRF8
EBI3
CLECL1
RGS1
POU2F2
BLNK
VPREB3
RAB30
FCRL3
STAP1
GPR18
TXNDC15
NLRP7
FCRLA
KLHL6
MAPAK1
CPNE5
FCRLS
ZBED2
SPIB
INPP5D
MIR155HG
LRMP
CD38
BLK
CD79A
RASGRP3
DERL3
RHBDD1
BANK1
PRICKLE1
TLR10
GNG7
KIAA0125
HVCN1
PLEKHA2
HLA−DOB
AIM2
PNOC
TMEM156
CSK
CD37
EAF2
KLHL14
PXK
CD1D
FAM129C
CD19


luad−CD4_mark−rmse_unique
0.0
0.1
0.2
0.3
0.4
0.5
0.6
IL32
LY9
C16orf28
FCRL6
ADAP1
ZAP70
BTN3A2
SPOCK2
FYB1
MBNL1
PTPN7
GIMAP7
ICOS
SLAMF1
GIMAP4
CD7
TNFSF8
SELPLG
PDCD1
PTGER2
CD247
TRAT1
GZMA
CTLA4
CTSC
UACA
CYTH1
ITGB7
CD5
TBC1D10C
GPR171
PVT1
KLHL6
LCP2
C16orf54
RASGRF1
EVL
GIMAP2
MYO1G
RASAL3
CD27
CCDC109B
PCK1
ITGAL
IL18R1
AKNA
ANXA3
GIMAP8
RASGRP2
PIM1
MBNL1
LRRC32
ITM2A
IKZF1
IFI6
DUSP6
BCL11B
CD5
RASA3
ARHGAP15
ITGA4
CAMK4
ITK


luad−CD8_mark−rmse_unique
0.0
0.1
0.2
0.3
0.4
EOMES
IFNG
CD244
LAG3
CBLB
GZMH
CLEC2D
PTPN22
KLRK1
KIR2DL4
GZMA
CTSW
NKG7
KIR3DL1
CD160
KLRG1
S1PR5
SAMD3
STAT4
CXCR6
NCR3
IL2RB
KIR2DL3
KLRD1
CRTAM
PRF1
GZMK
SH3GL3
GPR68
CD3G
APOBEC3G
FASLG
FGFBP2
CX3CR1
TIGIT
CD8B
GZMB
SH2D2A
TNIP3
PTGDR
RUNX3
TBX21
CST7


luad−Netro_mark−rmse_unique
0.00
0.05
0.10
0.15
0.20
0.25
0.30
0.35


luad−Macro_mark−rmse_unique
0.00
0.05
0.10
0.15
0.20
FCGR2B
NCEH1
PI4K2A
ADAP2
TK2
TPP1
DOCK2
STAB1
MS4A7
TIMP2
TM6SF1
KCTD12
SNX2
MMP14
CLN8
MAN1A1
BMP2K
HAVCR2
CPVL
PRAM1
CTSD
DPYD
TGFB1
NPL
CCDC88A
SIGLEC1
PARVG
SLC16A6
MMP19
NRP2
RNASE6
TRAM2
NCKAP1L
PIK3R5
RAP2B
HMOX1
SH2B3
CD84
APBB1IP
DFNA5
AP1S2
CLEC10A
CAPG
MS4A6A
IRF5
GNB4


luad−Dc_mark−rmse_unique
0.00
0.05
0.10
0.15
CD1C
IL10RA
FAB3
FAB5
CD1A
ARHGAP22
CCR8
CDKN1A
CD1B
ERD4
ARNT2
NAGA
RASAL2
TGAM
TFRM
KYNU
GJB2
ADAM12
TIGAB
CD80
NFKB1
SP3RA
SPCD1
PPMIM
CYBB
CDPE
CSF2RA
TYK2
CSF2R
FSD1L
PDCD1LG2
CRLF2
TRAF1
CCL1
GSN
FCER1A
SLC2A6
IRF4
CLEC4A
SLAMF7
RSH1
RAB7B
CXCL10
INHBA
FSCN1
RASSF4
TNFRSF4
KCNK6
TNFRSF18
CCR5
RAPGEF1
CD274
IL15
CD86
CCN7
MMP1
CCL2
PTPN1
MSC
IL6
CCL4
HK3
TREM2
BATF3