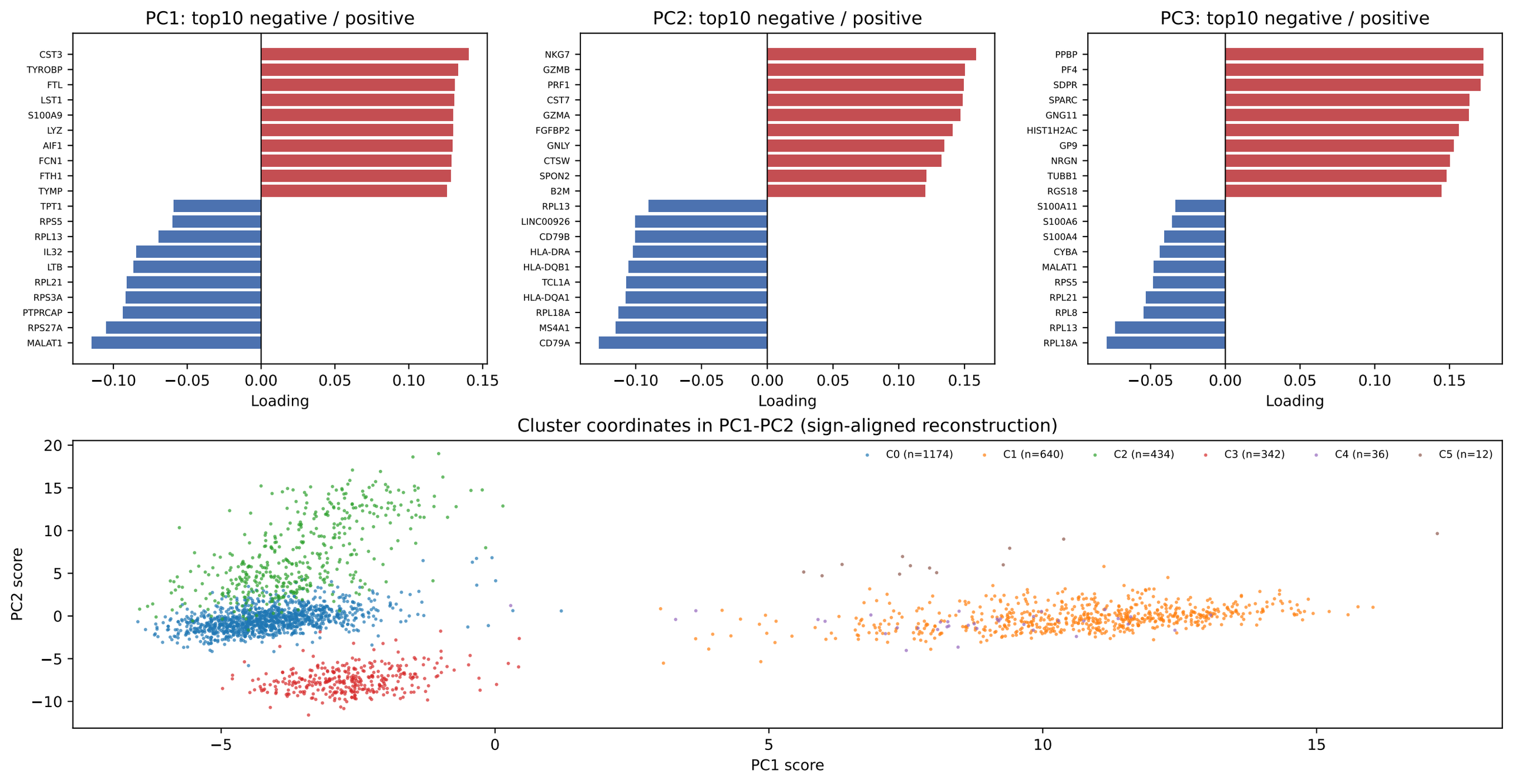
PC1: top10 negative / positive
CST3
TYROBP
FTL
LST1
S100A9
LYZ
AIF1
FCN1
FTH1
TYMP
TPT1
RPS5
RPL13
IL32
LTB
RPL21
RPS3A
PTPRCAP
RPS27A
MALAT1
−0.10
−0.05
0.00
0.05
0.10
0.15
Loading
PC2: top10 negative / positive
NKG7
GZMB
PRF1
CST7
GZMA
FGFBP2
GNLY
CTSW
SPON2
B2M
RPL13
LINC00926
CD79B
HLA-DRA
HLA-DQB1
TCL1A
HLA-DQA1
RPL18A
MS4A1
CD79A
−0.10
−0.05
0.00
0.05
0.10
0.15
Loading
PC3: top10 negative / positive
PPBP
PF4
SDPR
SPARC
GNG11
HIST1H2AC
GP9
NRGN
TUBB1
RGS18
S100A11
S100A6
S100A4
CYBA
MALAT1
RPS5
RPL21
RPL8
RPL13
RPL18A
−0.05
0.00
0.05
0.10
0.15
Loading
Cluster coordinates in PC1-PC2 (sign-aligned reconstruction)
C0 (n=1174)
C1 (n=640)
C2 (n=434)
C3 (n=342)
C4 (n=36)
C5 (n=12)
PC2 score
20
15
10
5
0
−5
−10
−5
0
5
10
15
PC1 score