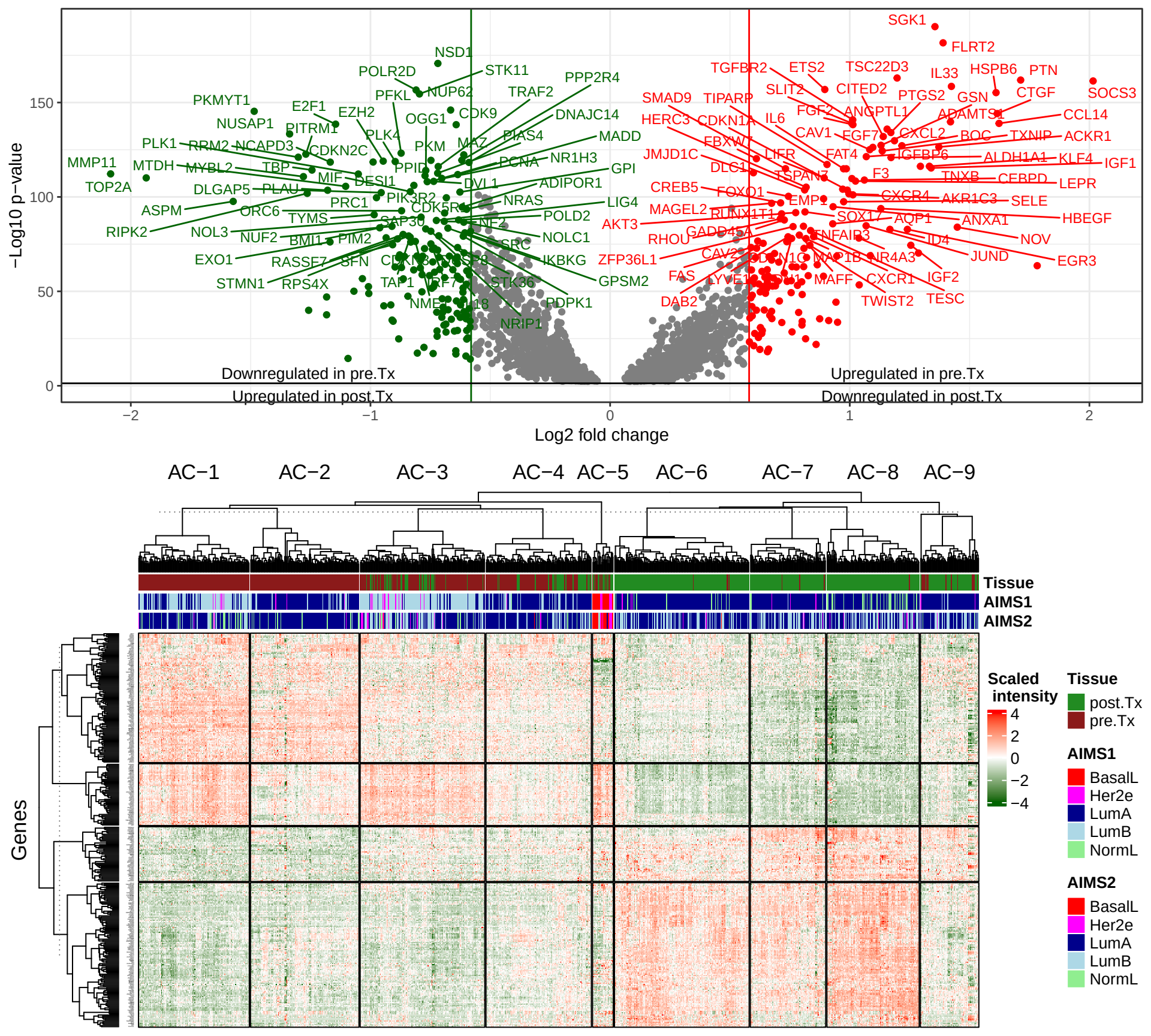
−Log10 p−value
Log2 fold change
Downregulated in pre.Tx
Upregulated in post.Tx
Upregulated in pre.Tx
Downregulated in post.Tx
SGK1
FLRT2
TGFBR2
ETS2
TSC22D3
IL33
HSPB6
PTN
SOCS3
SLIT2
CITED2
PTGS2
GSN
CTGF
SMAD9
TIPARP
FGF2
ANGPTL1
ADAMTS1
CCL14
HERC3
CDKN1A
IL6
CAV1
FGF7
CXCL2
BOC
TXNIP
ACKR1
FBXW7
JMJD1C
LIFR
FAT4
IGFBP6
ALDH1A1
KLF4
IGF1
DLC1
TSPAN7
F3
TNXB
CEBPD
LEPR
CREB5
FOXO1
EMP1
CXCR4
AKR1C3
SELE
MAGEL2
RUNX1T1
SOX17
AQP1
ANXA1
HBEGF
AKT3
GADD45A
TNFAIP3
RHOU
ID4
NOV
CAV2
ZFP36L1
MAP1B
NR4A3
JUND
EGR3
FAS
LYVE1
MAFF
CXCR1
IGF2
DAB2
TWIST2
TESC
NSD1
POLR2D
STK11
PPP2R4
PKMYT1
PFKL
NUP62
TRAF2
E2F1
EZH2
CDK9
DNAJC14
NUSAP1
PITRM1
OGG1
PLK4
PIAS4
MADD
PLK1
RRM2
NCAPD3
CDKN2C
PKM
MAZ
NR1H3
MMP11
MTDH
MYBL2
TBP
PPID
PCNA
GPI
TOP2A
MIF
DESI1
DVL1
ADIPOR1
DLGAP5
PLAU
PIK3R2
NRAS
LIG4
ASPM
PRC1
ORC6
CDK5R1
POLD2
TYMS
SAP30
ENF2
RIPK2
NOL3
NUF2
BMI1
PIM2
NOLC1
SRC
EXO1
RASSF7
SFN
CDK13
SP8
IKBKG
STMN1
RPS4X
TAP1
IRF7
STK36
GPSM2
NME1
IL18
RDPK1
NRIP1
150
100
50
0
−2
−1
0
1
2
AC−1
AC−2
AC−3
AC−4
AC−5
AC−6
AC−7
AC−8
AC−9
Tissue
AIMS1
AIMS2
Genes
Scaled intensity
4
2
0
−2
−4
Tissue
post.Tx
pre.Tx
AIMS1
BasalL
Her2e
LumA
LumB
NormL
AIMS2
BasalL
Her2e
LumA
LumB
NormL