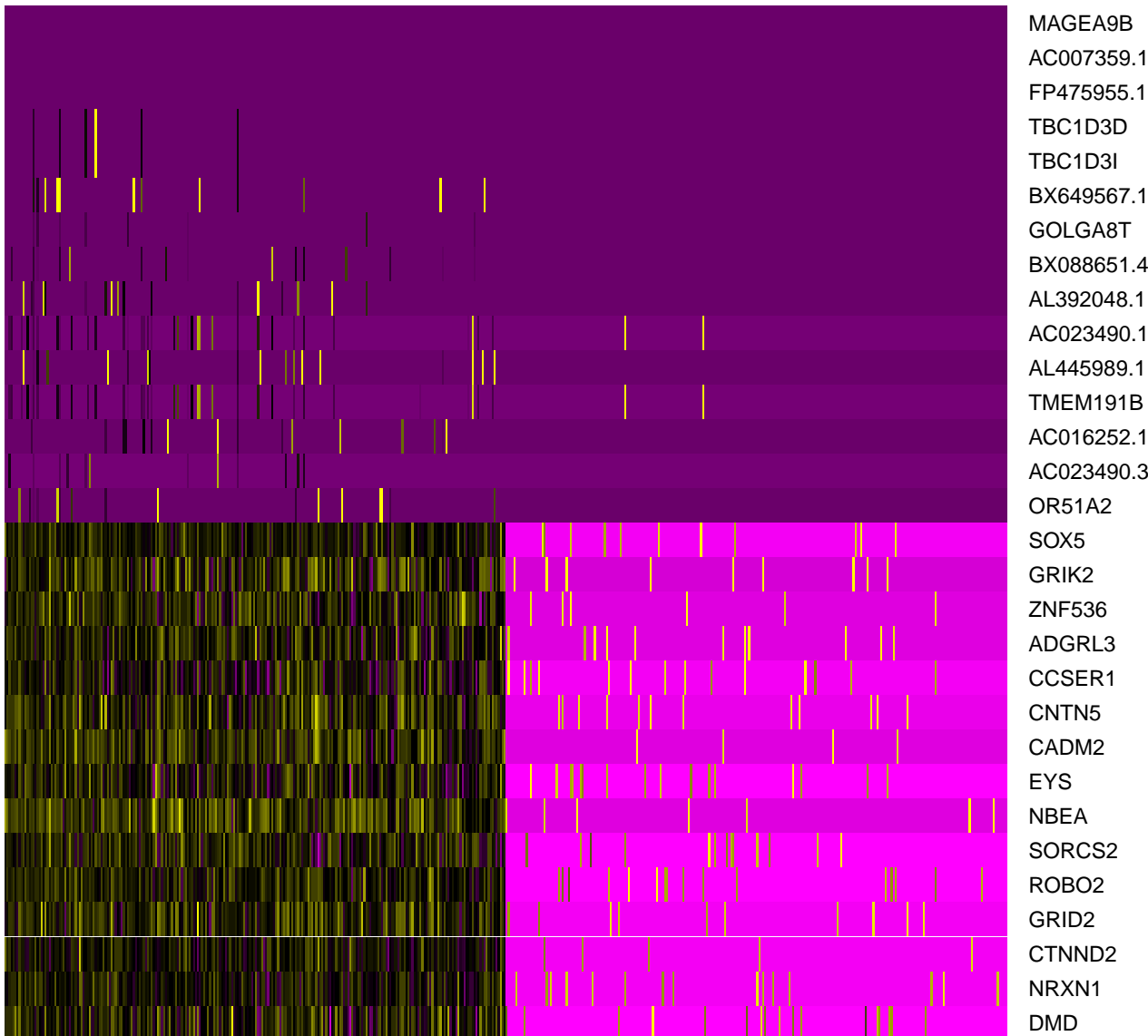
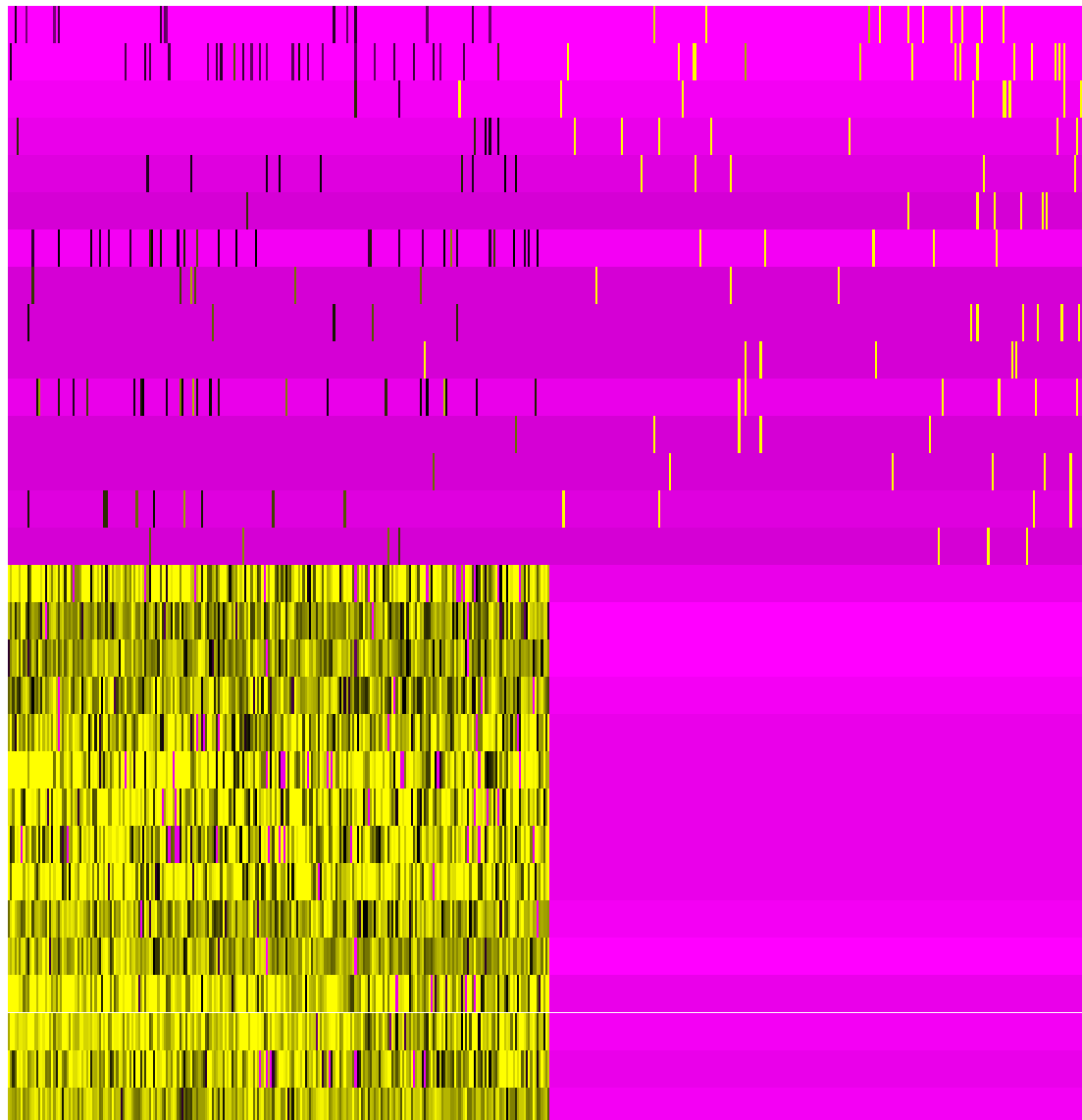


PC_1

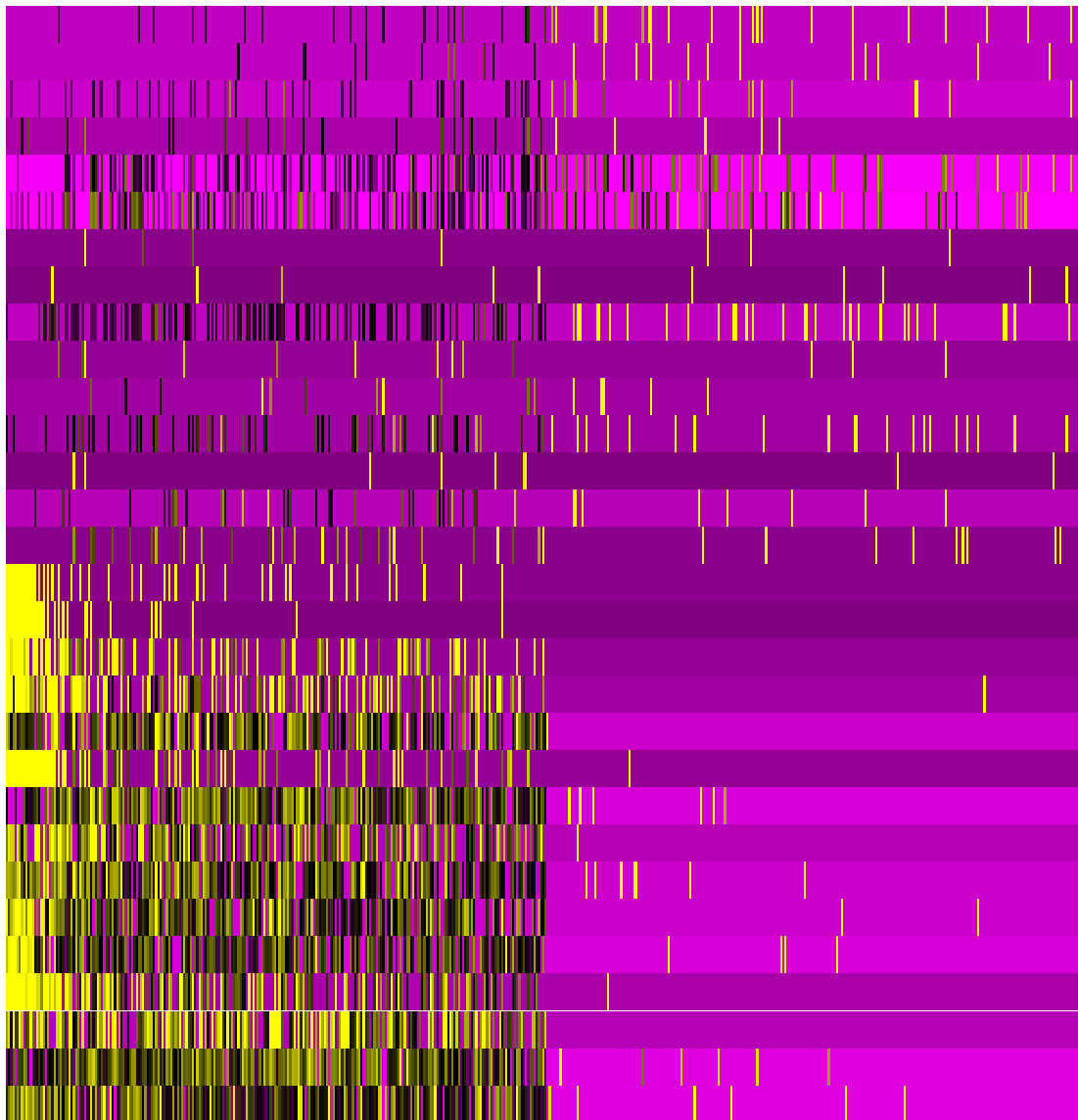


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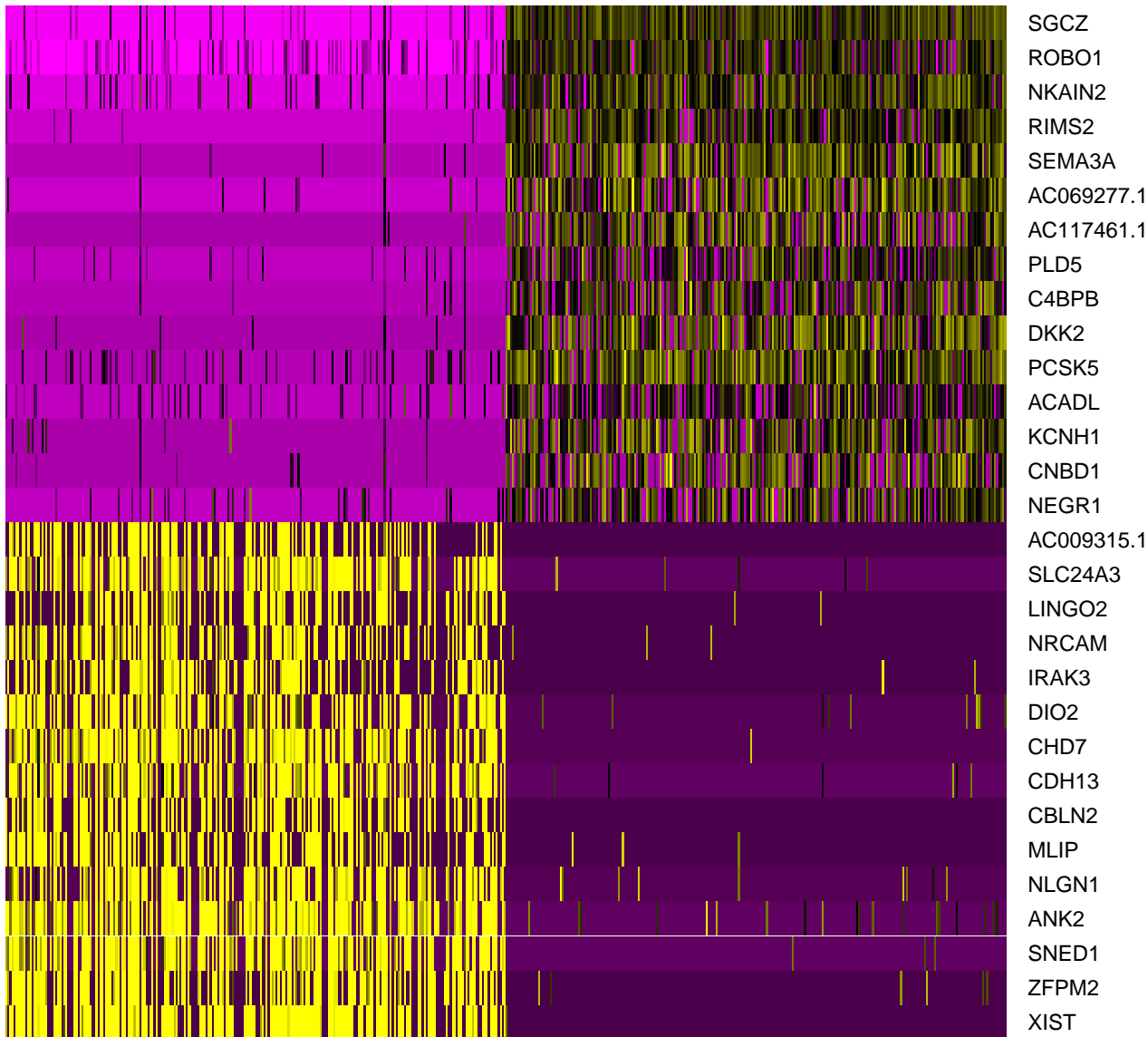
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LRFN5
ITIH5
HMCN2
ASIC2
SNTG2
MYO16
RRM2
ALDH1A2
LINC01250
LINC00861
DGKB
NTRK3
ADAM12
AC092957.1
RORA
HLTF
SGCD
EYA4
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CALCR
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RIMS2
RBMS3
WWTR1
SGCZ
ROBO1
SERPINE2
LAMA2

PC_1

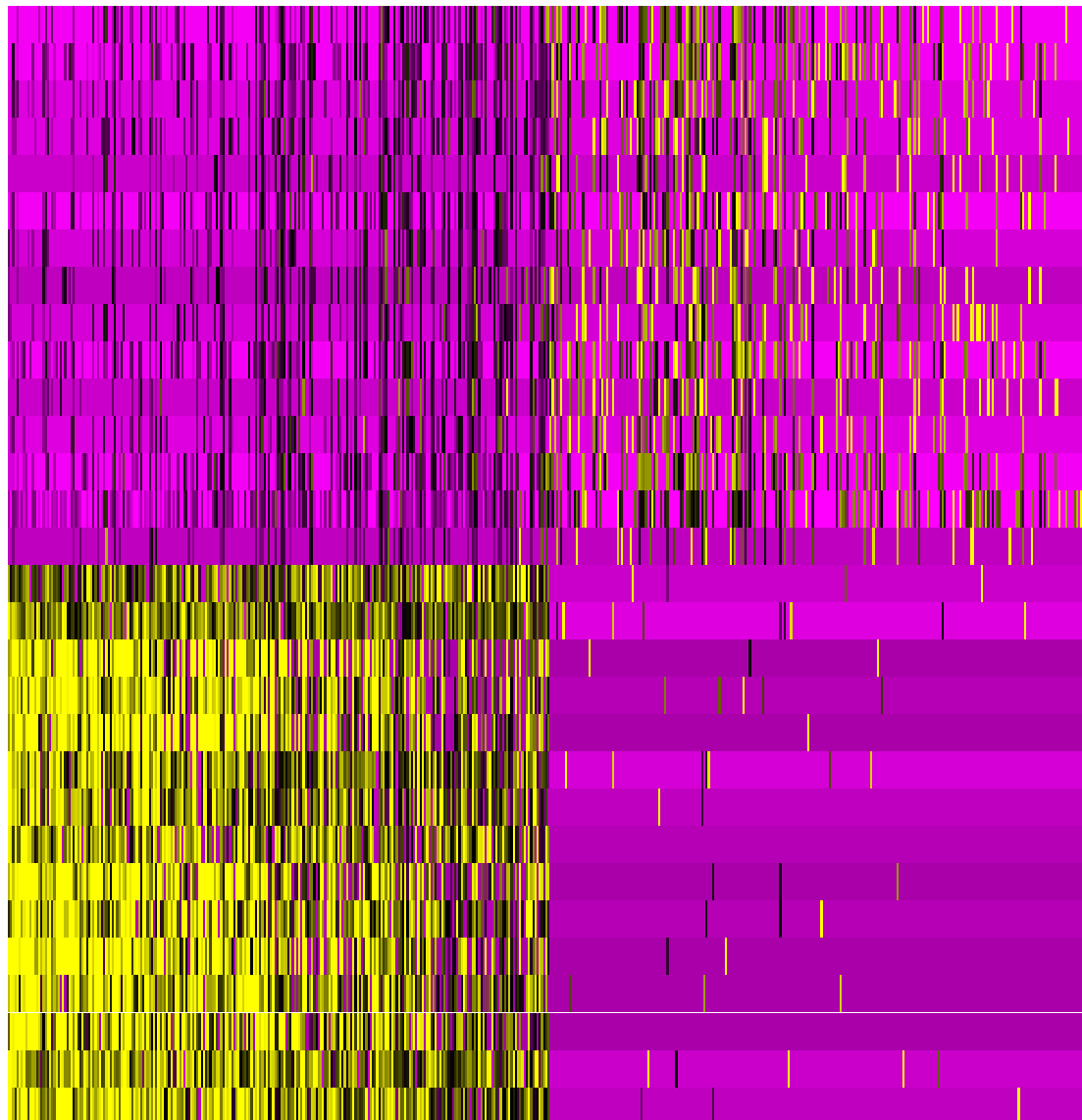


ALG1L9P
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NES
CSMD1
TTN
RRM2
IGHG4
IGKC
POLQ
SRRM4
RPS15
MKI67
TMEM132D
RPL27
ANGPT2
MIR181A1HG
PTPRK
NRP1
MEF2A
MIR4435-2HG
ARHGAP24
MBNL1
FNDC3B
PTPRM
TCF4
RASAL2
RBMS1
ZBTB20
NEAT1

PC_1



PC_1



AGLB1
RYR3
MEGF10
NOS1
MIR1-1HG
FRAS1
AF279873.3
MTMR7
LDB3
CNR1
AC007846.1
LINGO2
DIO2
NEB
MYPN
TSHZ2
RASAL2
MECOM
ADGRL4
ANO2
COL4A2
COL4A1
RBPMS
VWF
RAI14
PTPRB
CADPS2
ERG
IGFBP7
LDB2

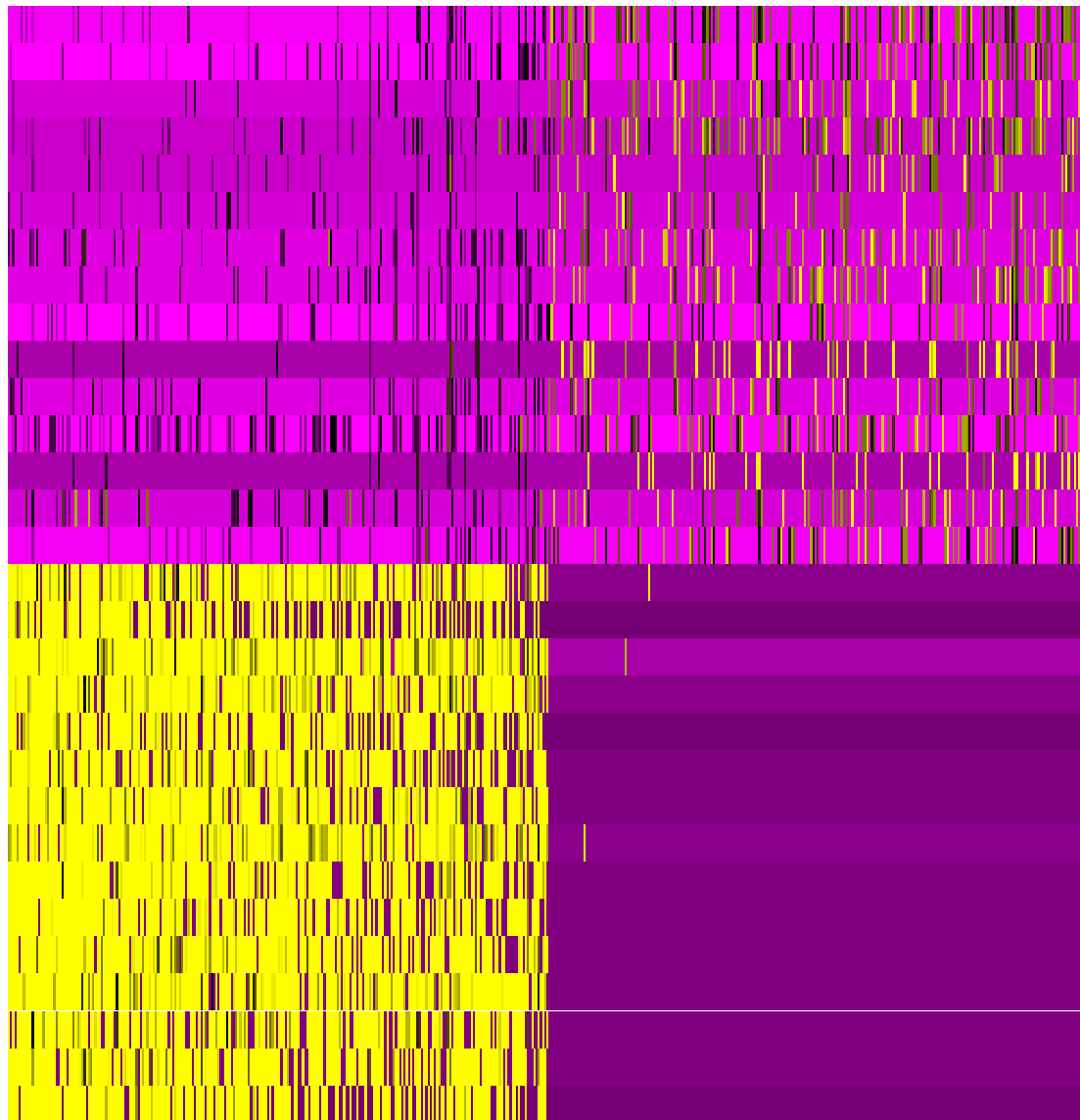
PC_1



PC_1

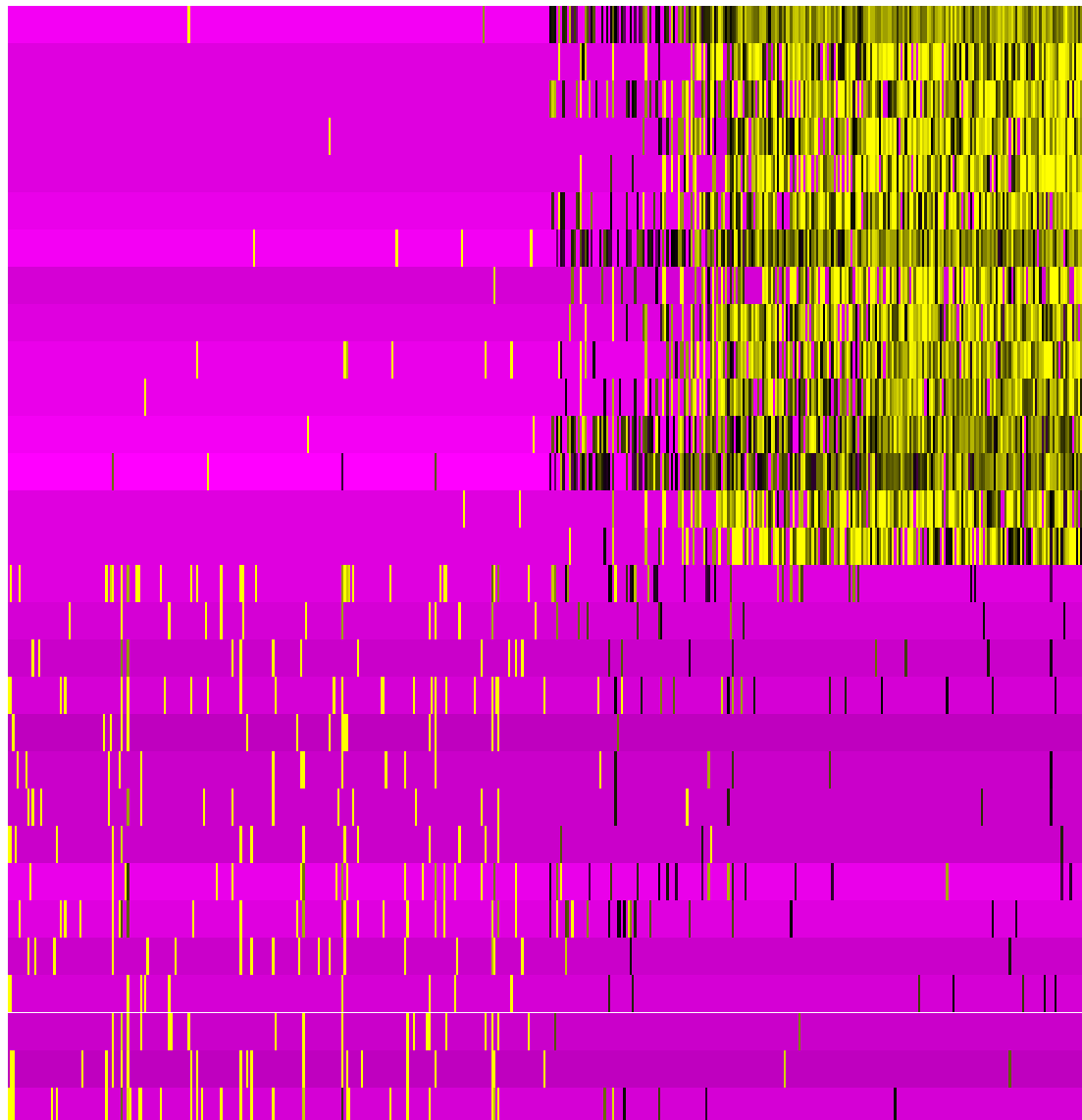


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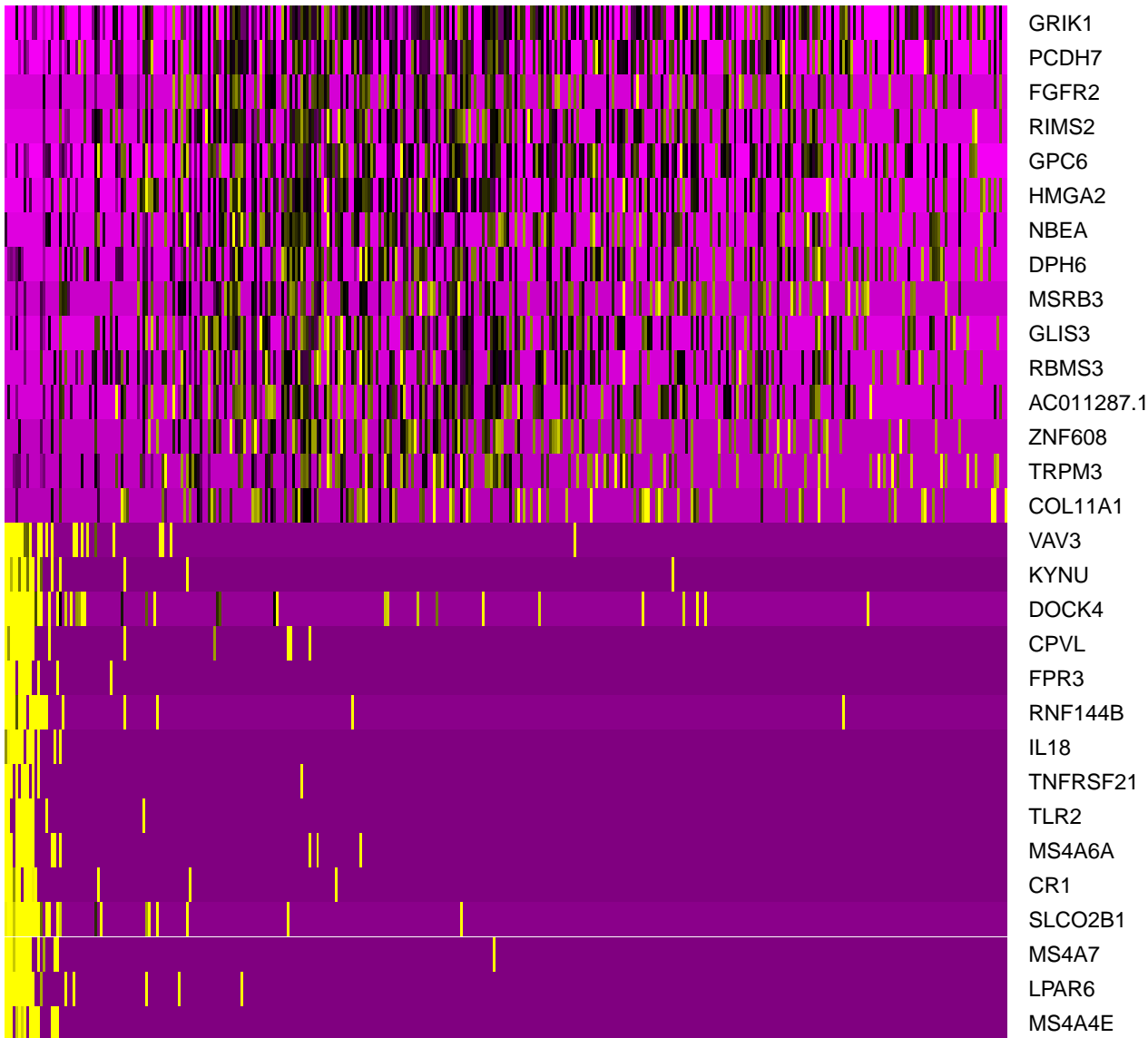
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POSTN
CSMD3
DPP10
MIR3681HG
LRRC4C
ADAMTS17
COL25A1
MEG8
AGBL4
PCDH9
FN1
LINC00377
FGF13
COL3A1
ARRDC3
PPP1R16B
GRB10
CADPS2
C22orf34
PCDH17
PECAM1
XIST
RASGRF2
PTPRB
SHANK3
FLT1
VEGFC
EGFL7
DIPK2B

PC_1

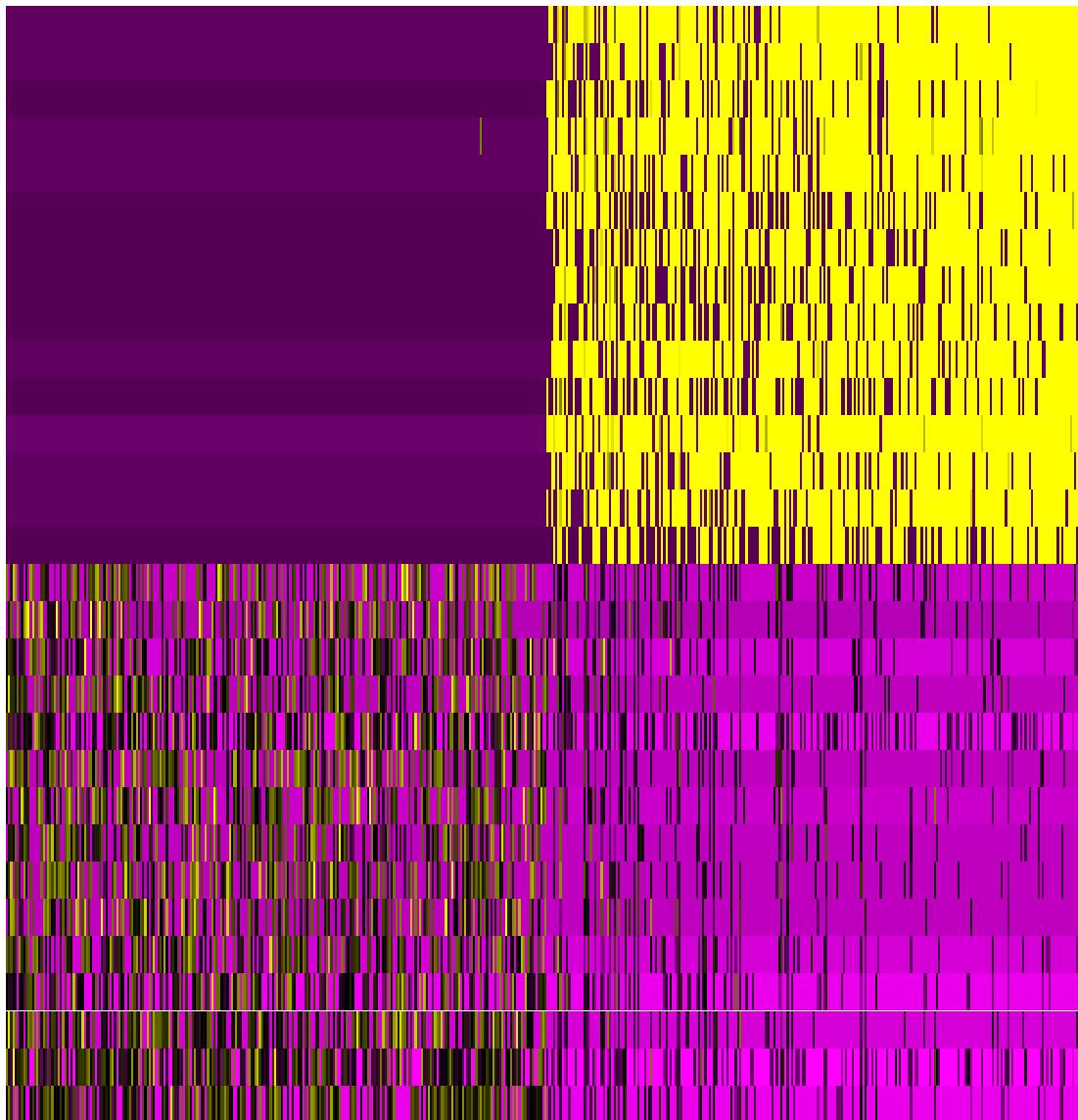


PTPRM
MECOM
EPAS1
PTPRB
EMCN
GRB10
LDB2
CALCRL
VWF
ST6GALNAC
PRKCH
SPARCL1
NFIB
ANO2
FLT1
COL27A1
TENM3
FAM20C
GALNT17
SCN3B
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ST6GALNAC
MEG8
RUNX1
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LINC02511
AP001924.1
PAX7
CLMP

PC_1

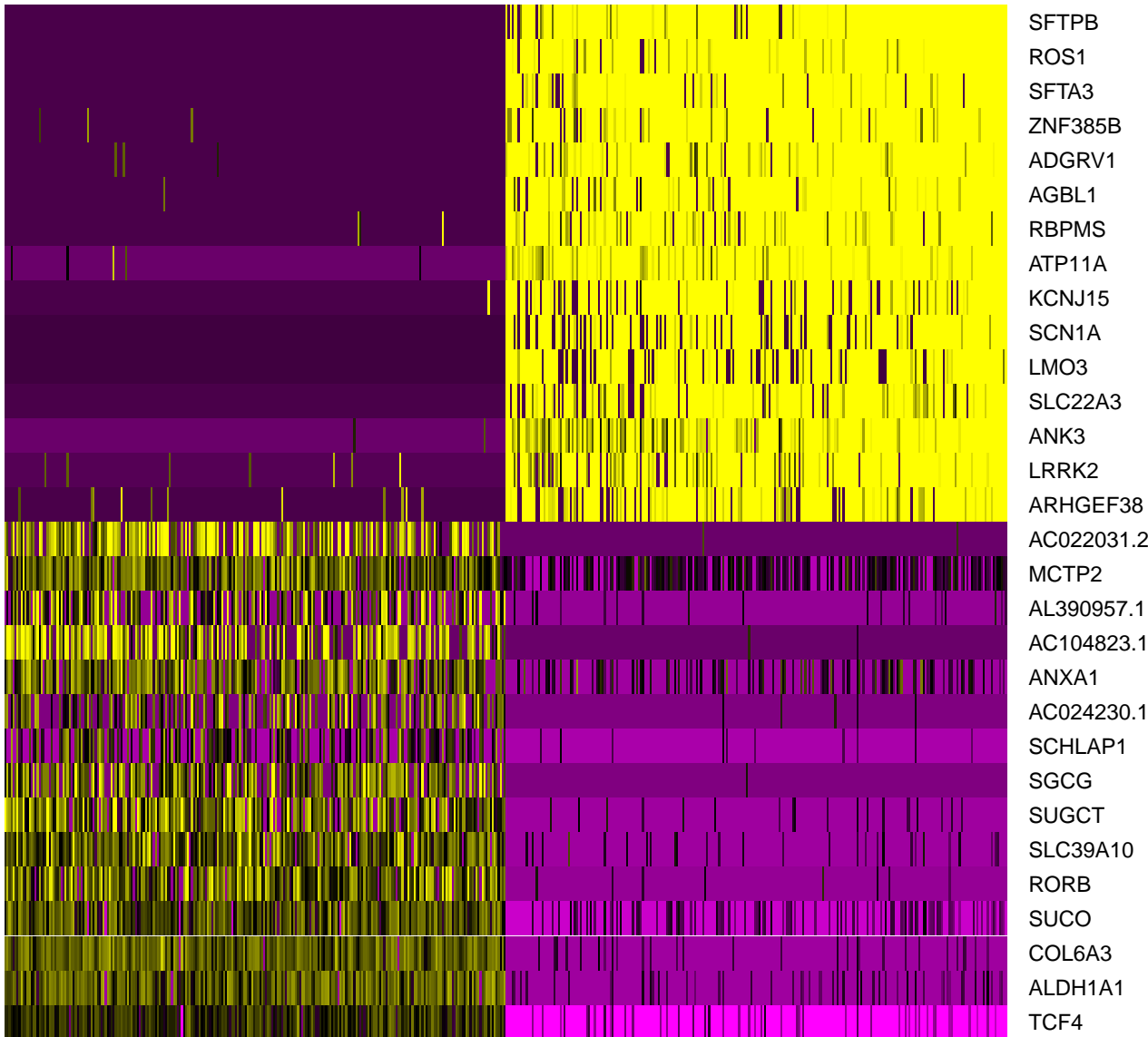


PC_1

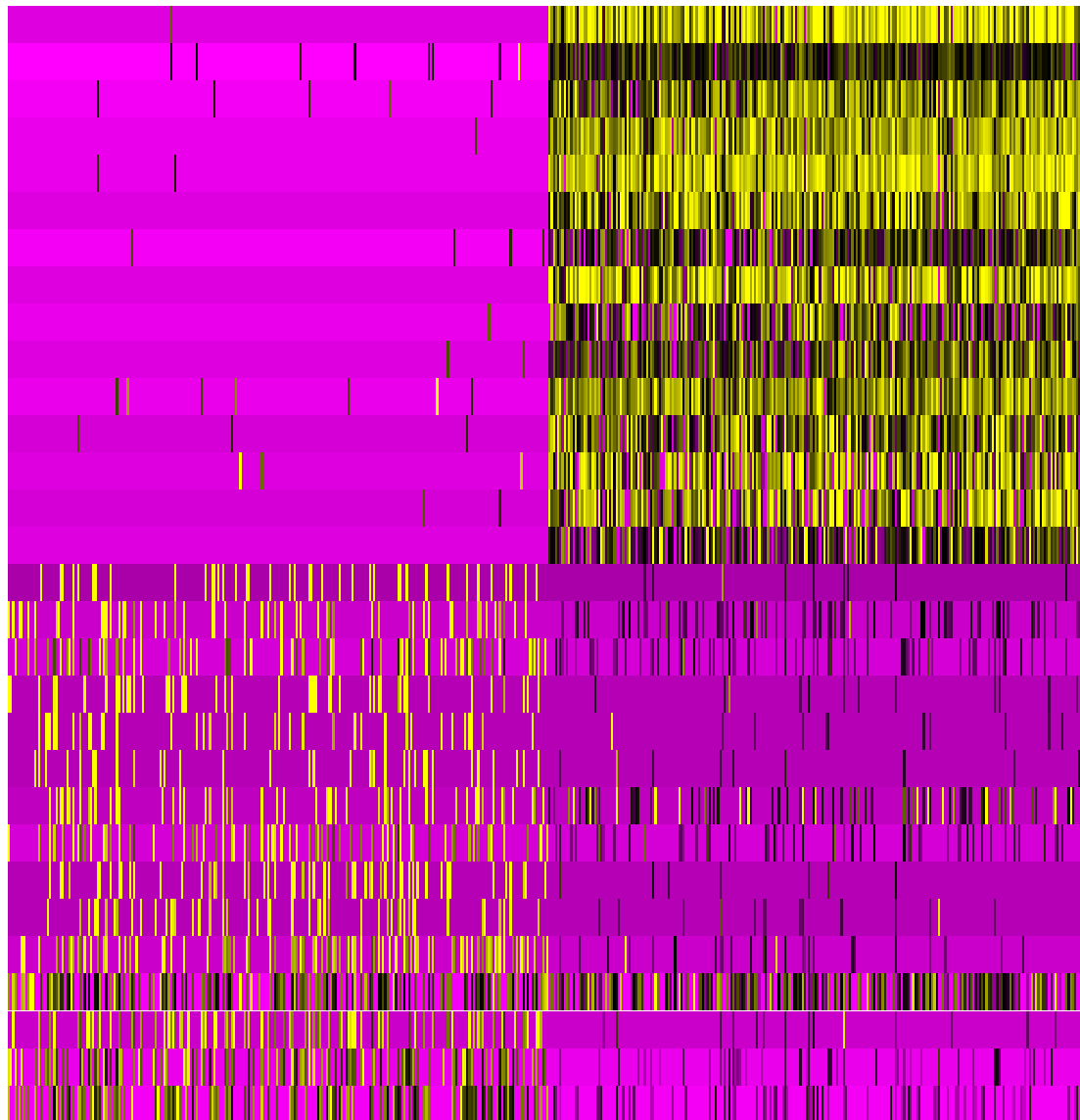


ATP8B4
ARHGAP18
MRC1
F13A1
CD163L1
MS4A6A
LYN
SLCO2B1
FYB1
SYK
LINC01374
ITPR2
DOCK8
PTPRC
MS4A7
CDK14
SCHLAP1
PALLD
PDE10A
INPP4B
TSHZ2
BMPER
LRIG3
COL21A1
ZFHX4-AS1
RBFOX1
KIF26B
CCBE1
NPAS3
BMP5

PC_1



PC_1

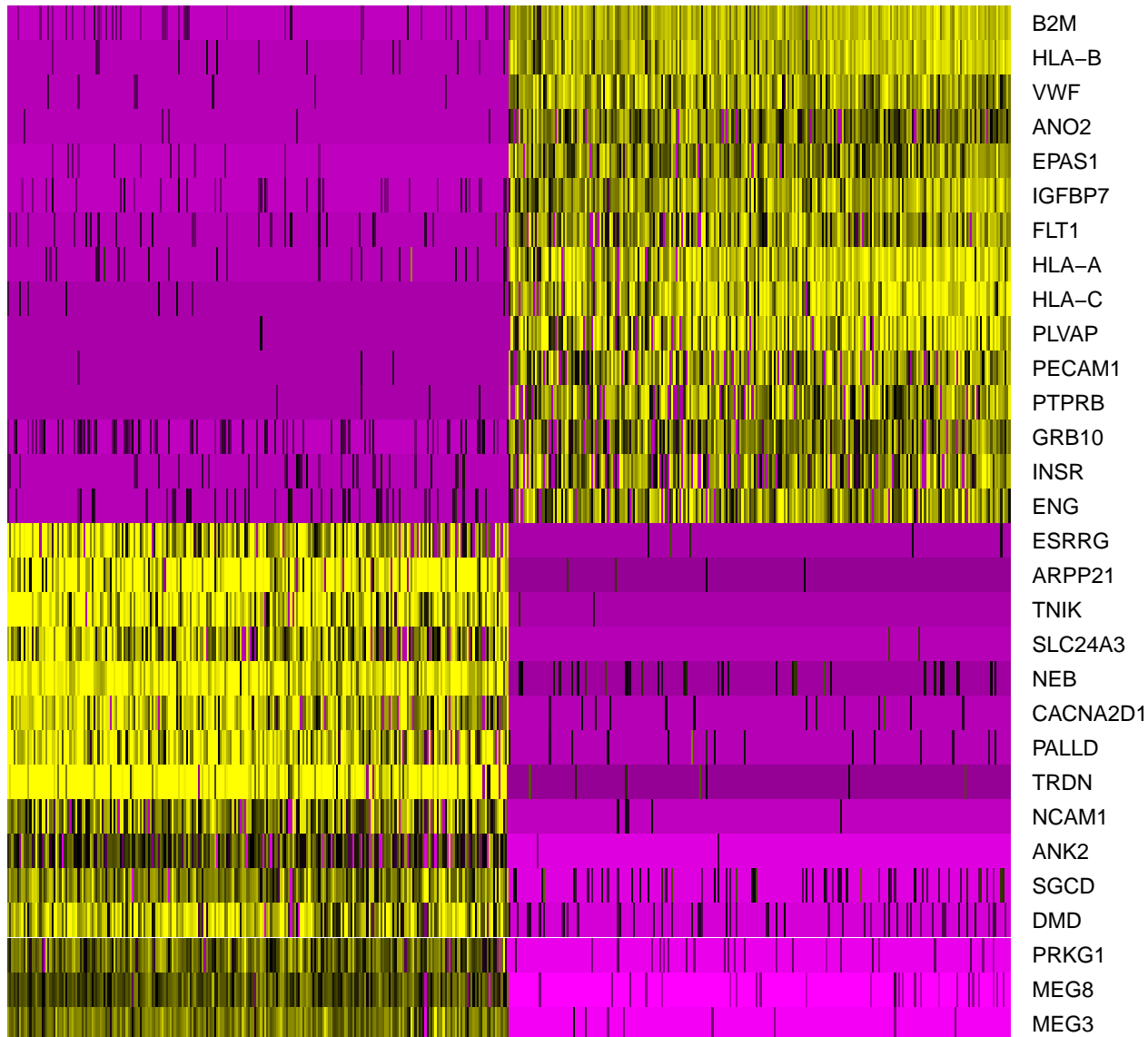


BBOX1-AS1
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COBL
SNTB1
LINC02391
AKAP6
MEGF10
TMEM108
EPHA3
MTSS1
CADM2
MPPED2
RGS7
AP001924.1
AC068633.1
ANKUB1
TNC
LRRC4C
APOLD1
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TMEM132B
MAGI2
AL691420.1
OPCML
CA10
NELL1
SLC8A1
ZFPM2-AS1
CACNA1C

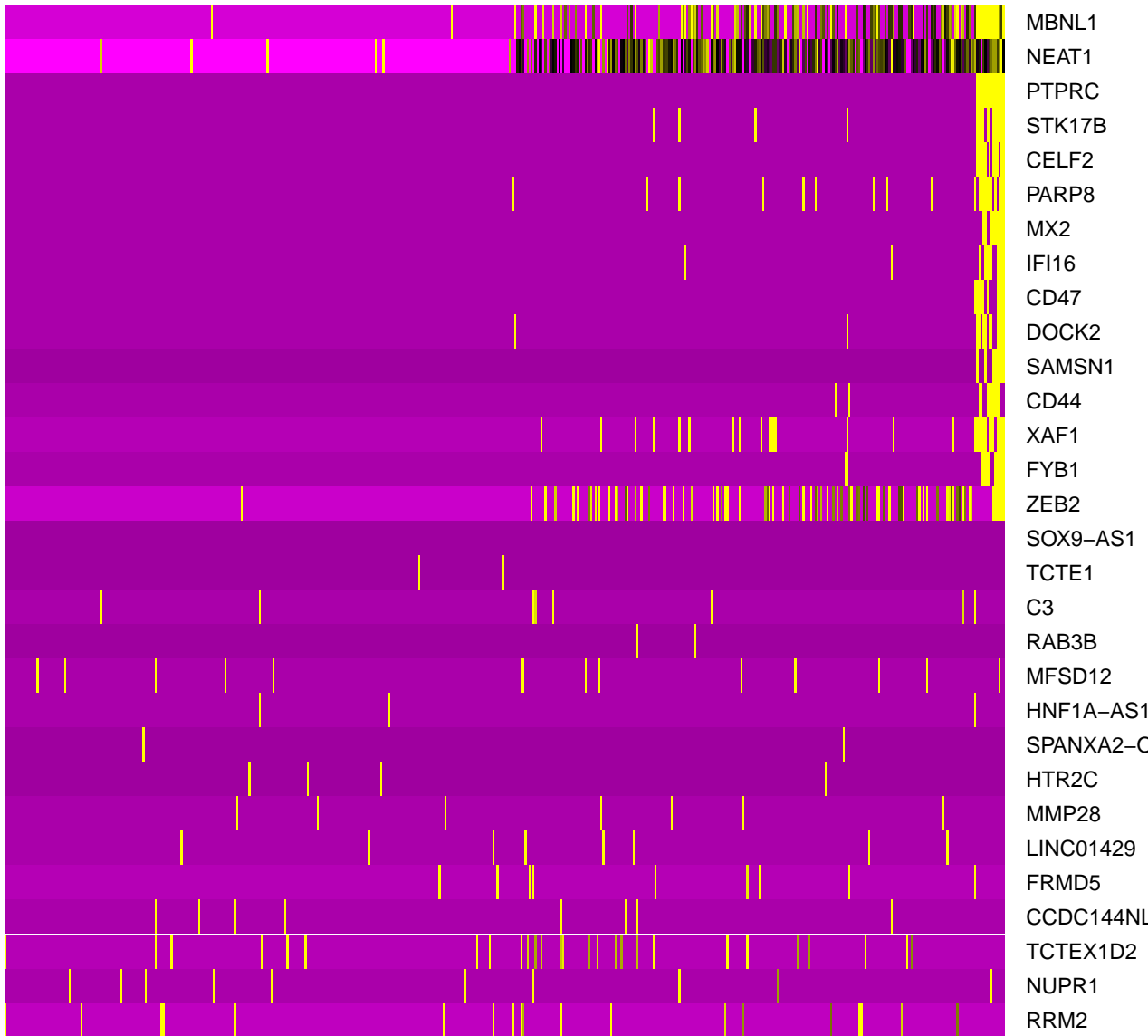
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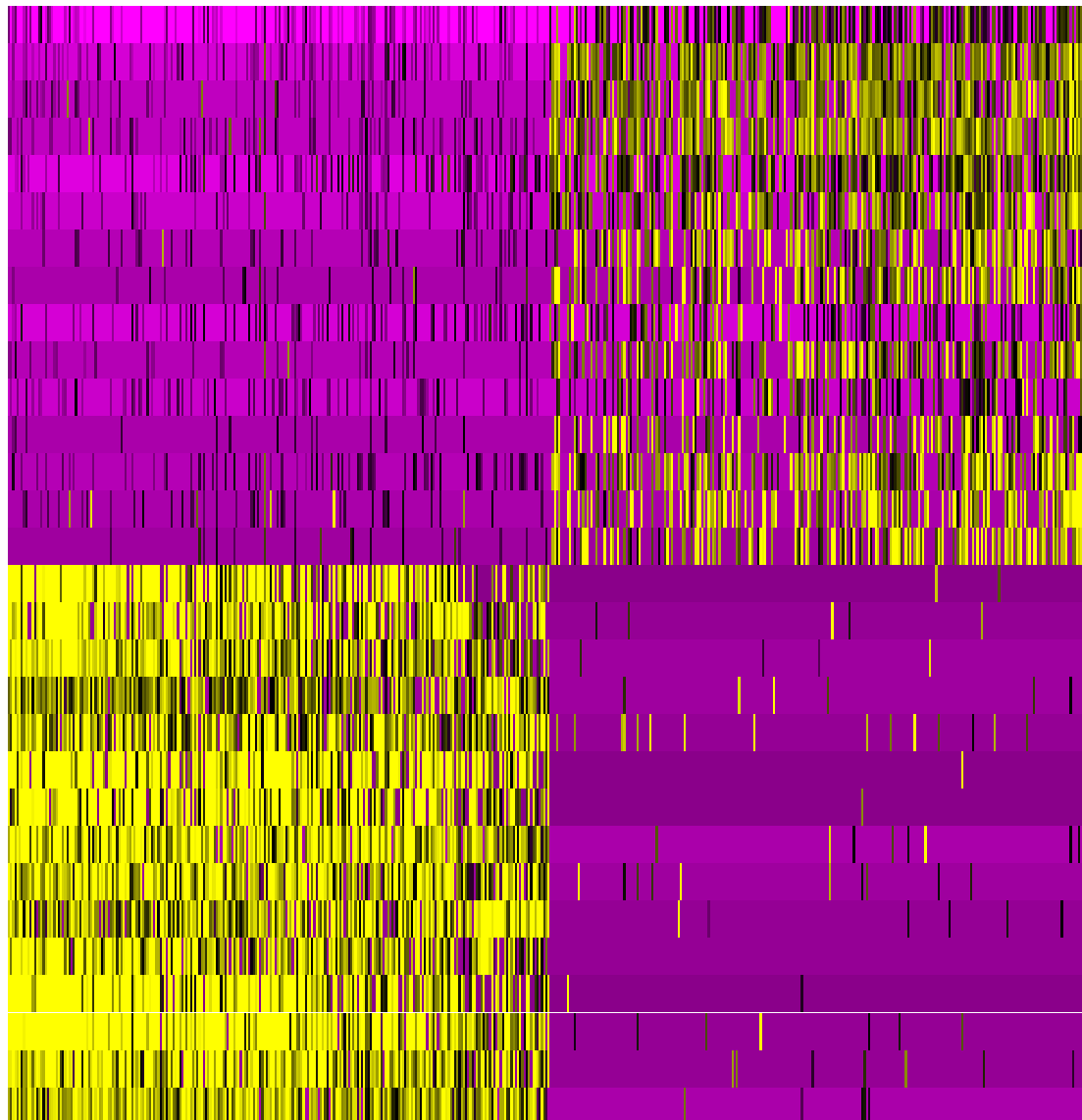
PC_1



PC_1

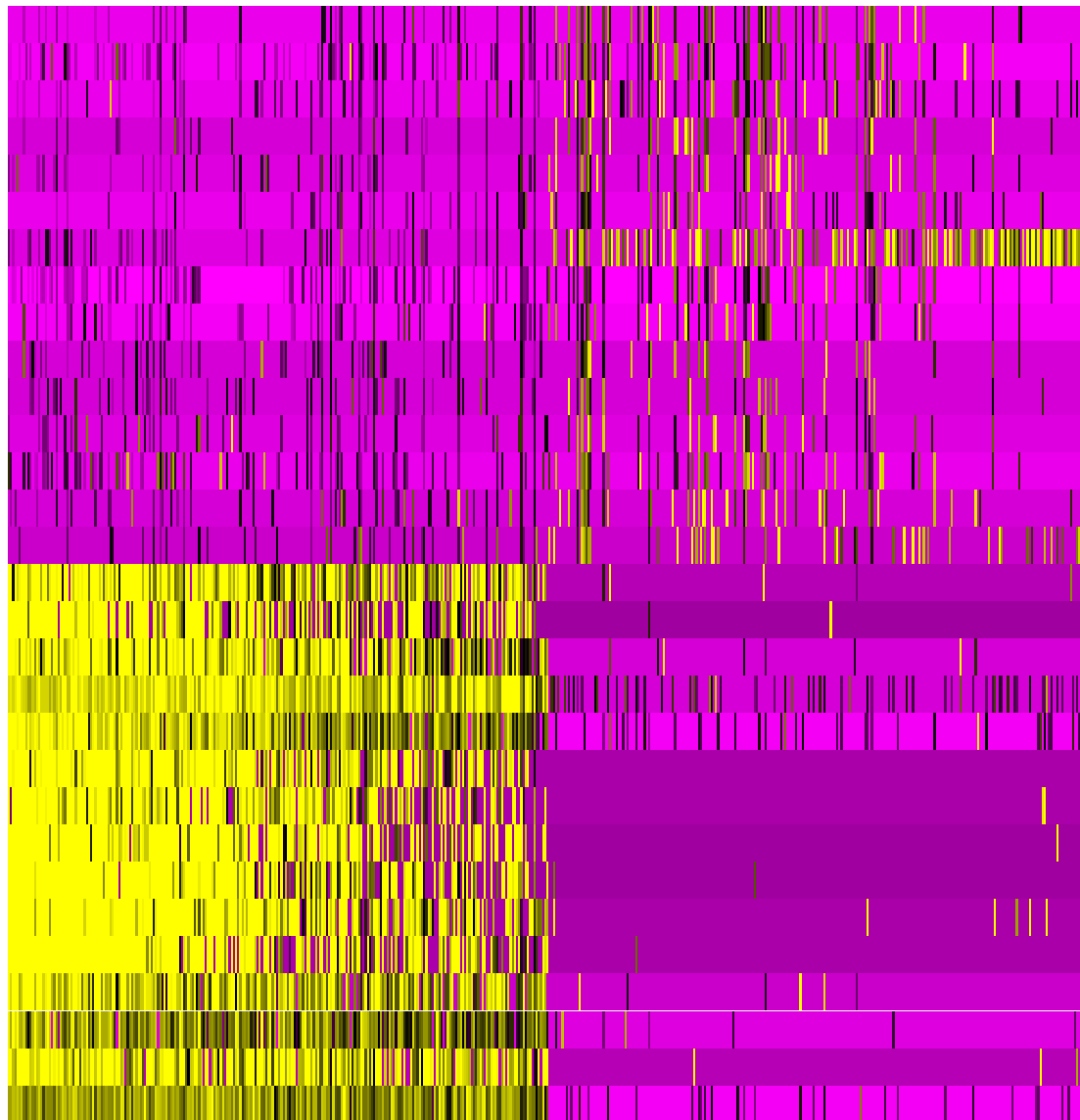


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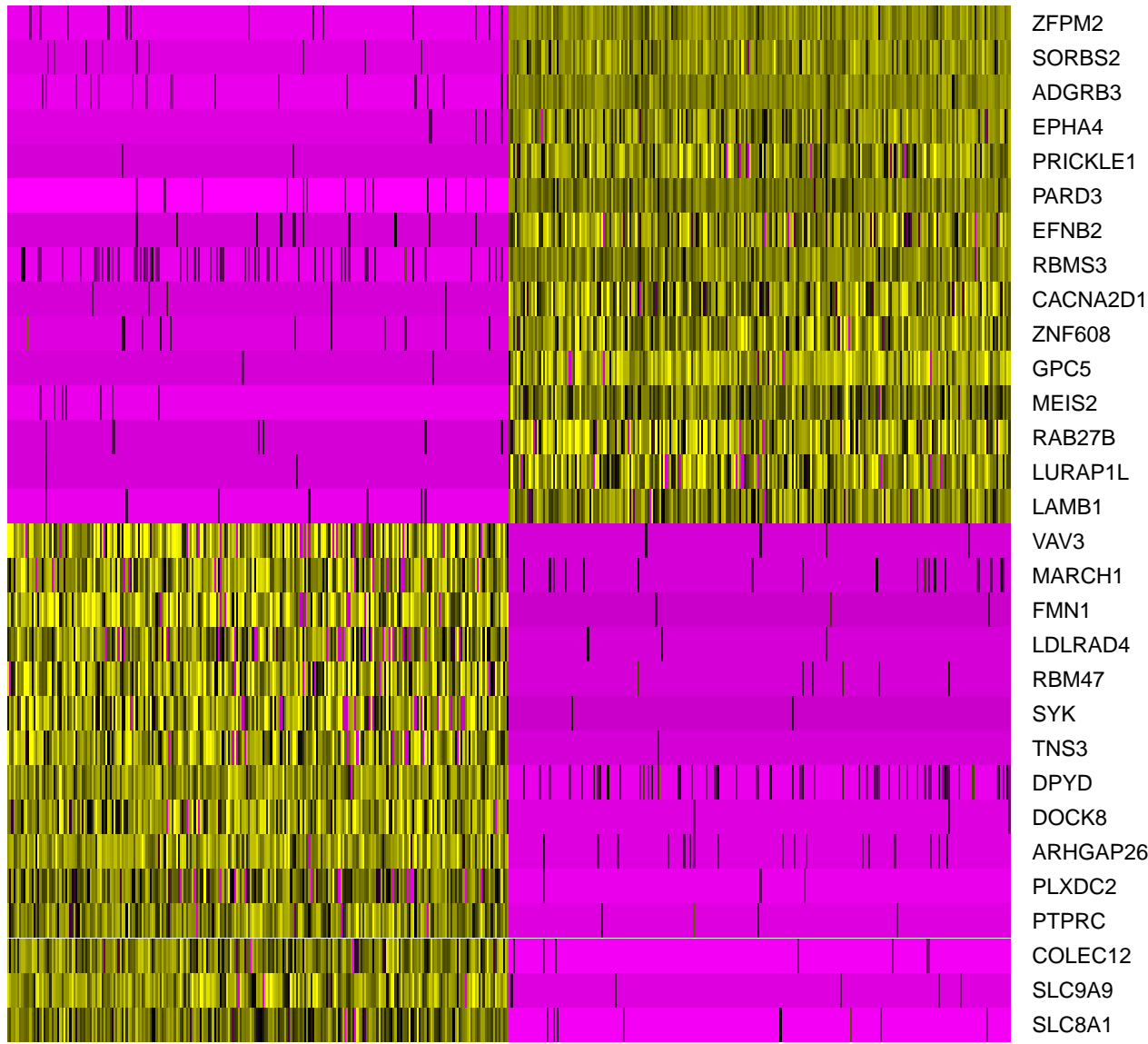
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CTNNA3
DKK2
NCAM2
TENM3
AC138627.1
AC124854.1
NLGN1
ERBB4
DLGAP1
PCLO
LRRC4C
GPM6A
AP001977.1
CSGALNAC1
NRIP1
DOCK4
ATXN1
PDLIM5
ERG
EPAS1
PPP3CA
COL4A1
EBF1
PTPRE
GRB10
LDB2
COL4A2
NRP1

PC_1

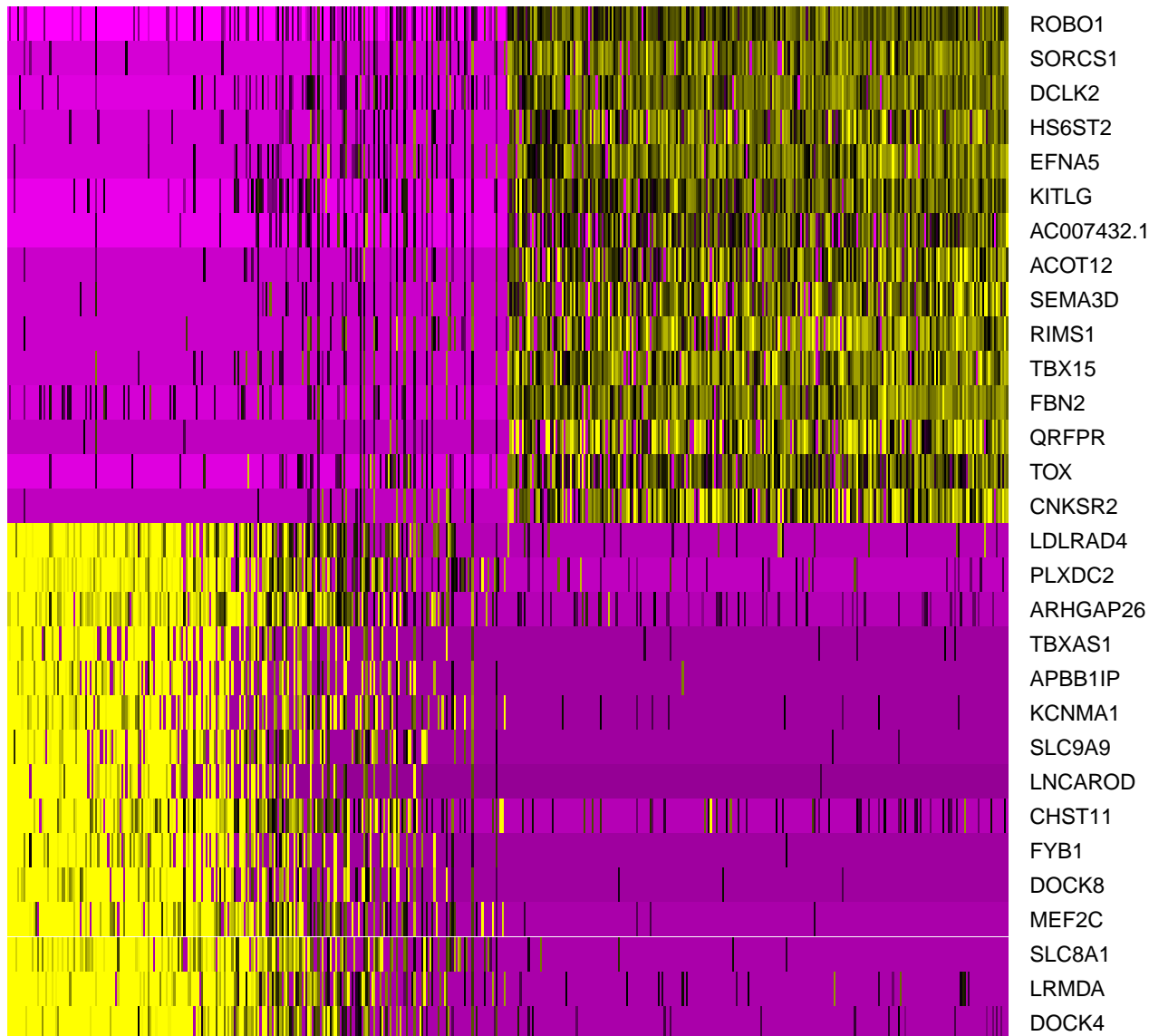


ADAMTSL3
DLGAP1
CACNA2D3
ENPP2
TENM1
PDE7B
MSR1
MMP16
COL11A1
EPHA3
DLGAP2
ANGPT1
ABI3BP
CPA6
ROR1
ACTN2
TRDN
FGF13
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DMD
REEP1
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LRRN1
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TRIM55
NEB

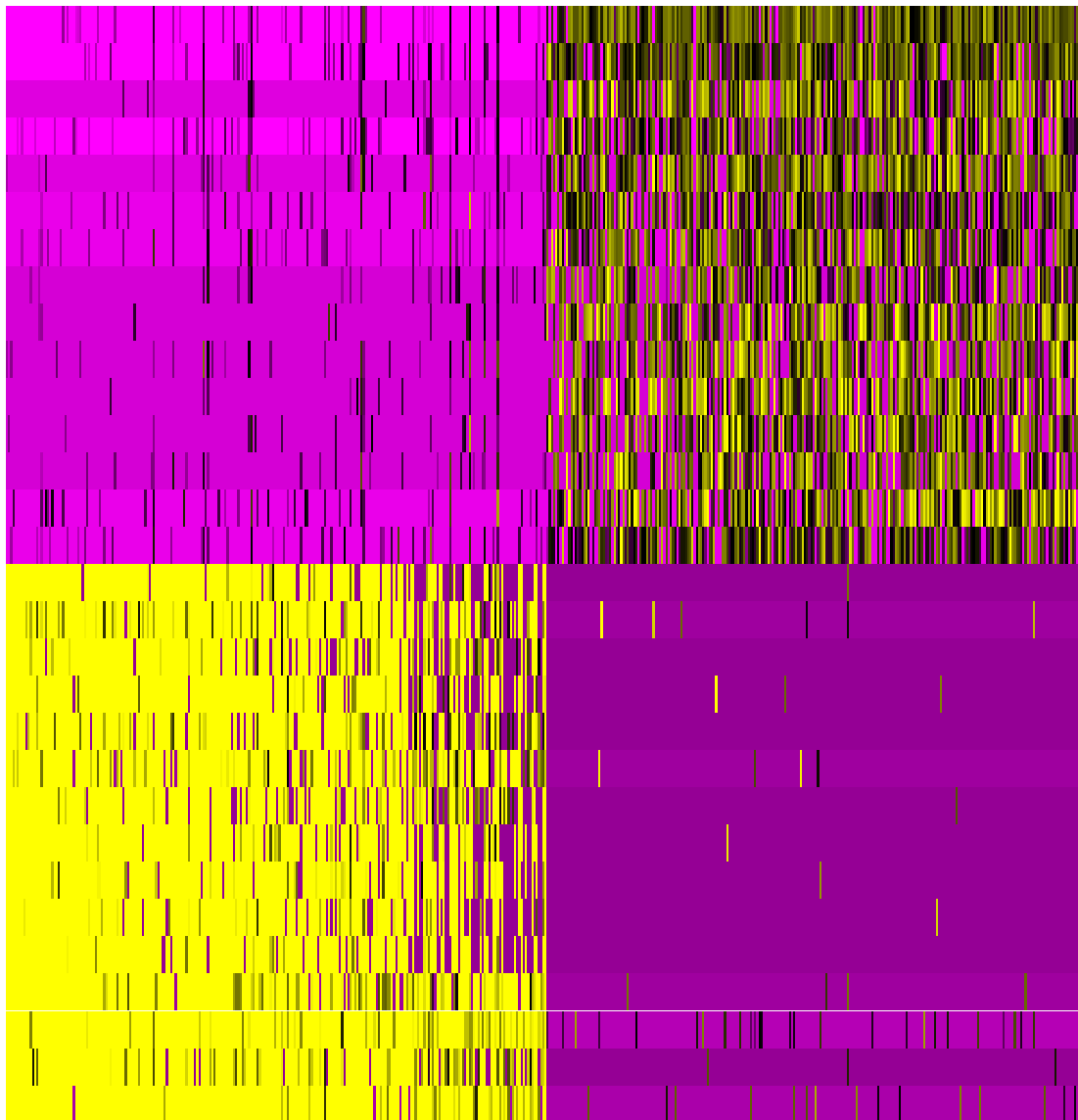
PC_1



PC_1

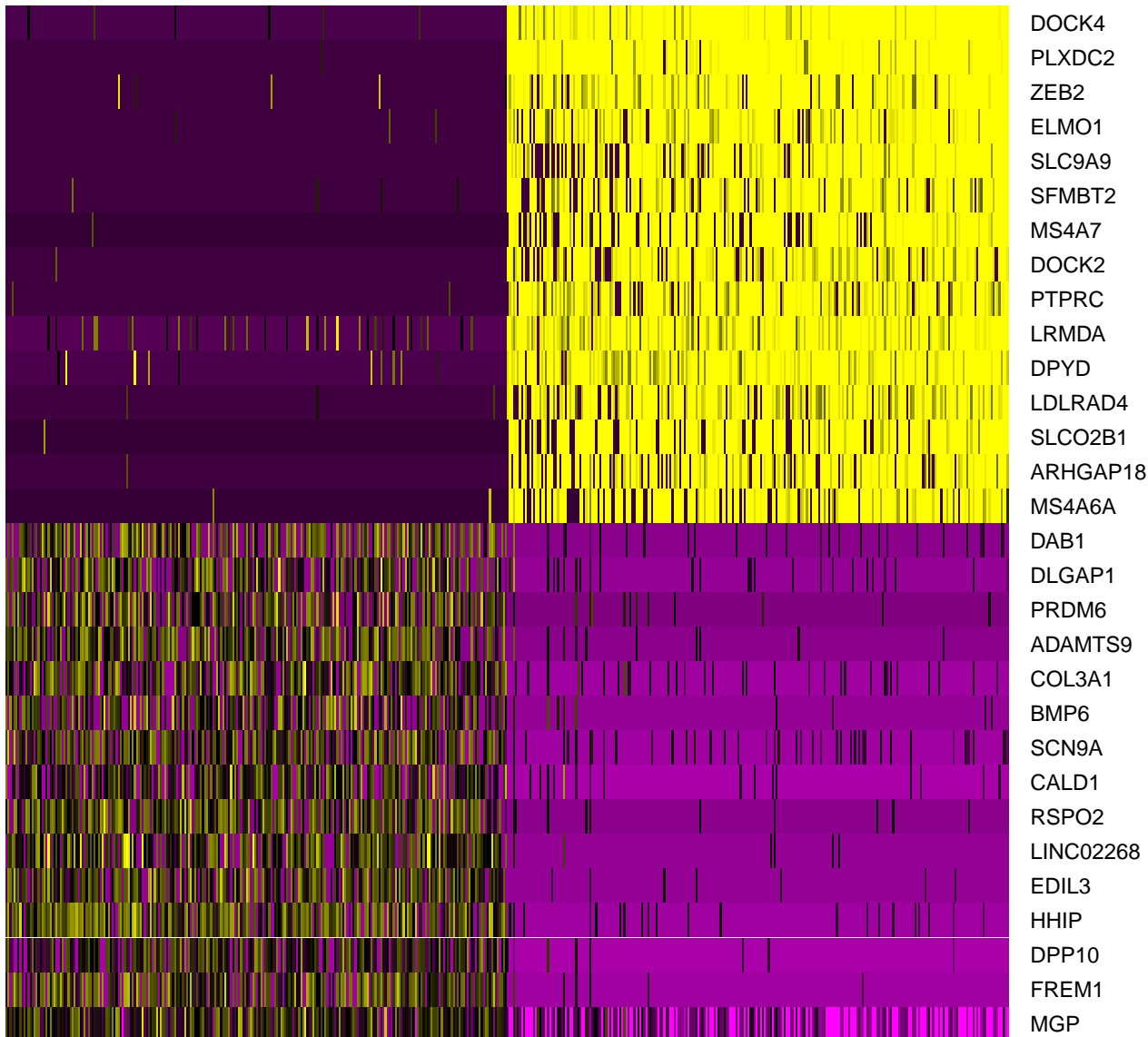


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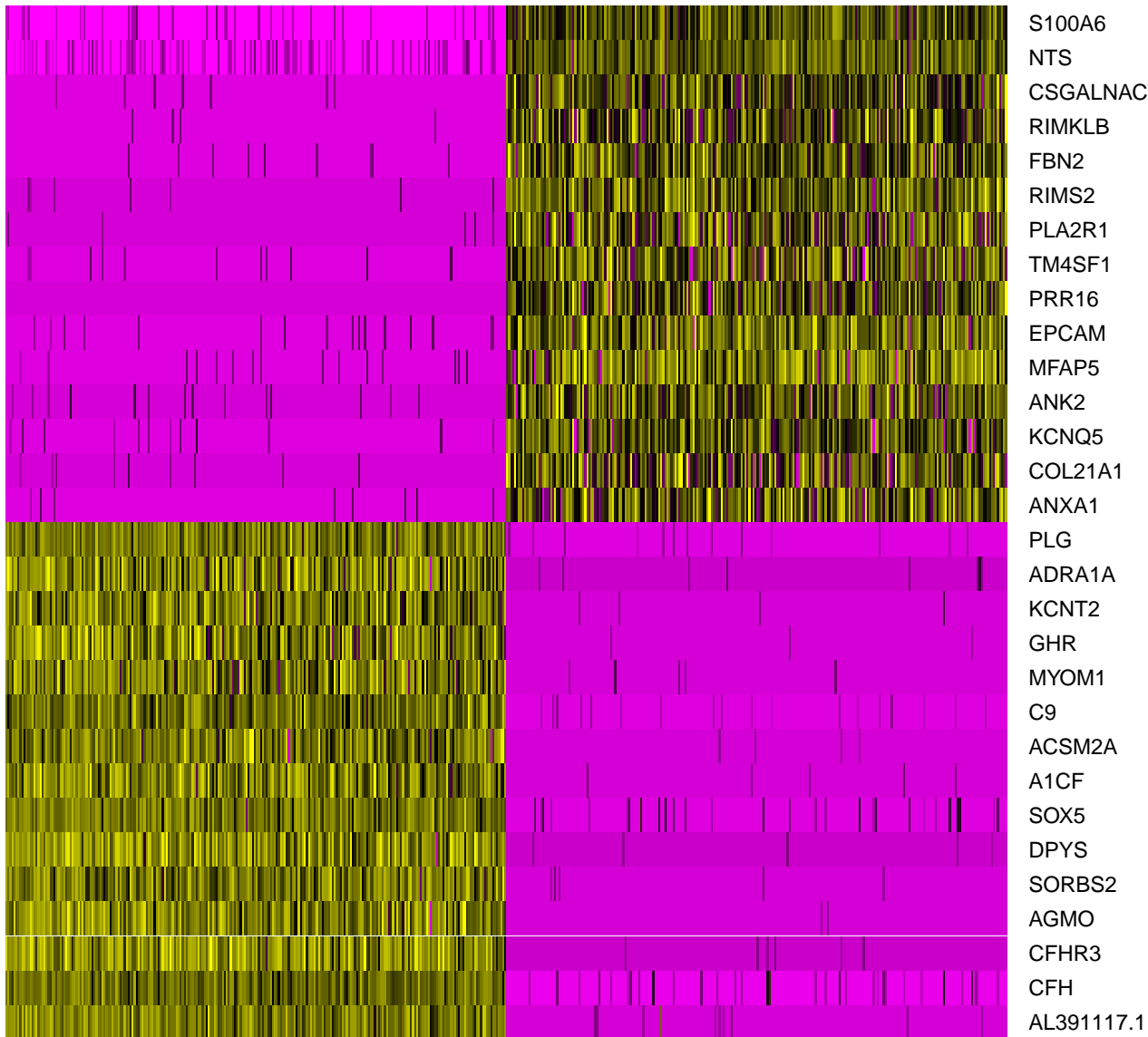


HTR2C
CADM2
OTX2-AS1
TENM3
SOX2-OT
PTPRD
LRP1B
MTUS2
DCDC2
RYR3
NTNG1
EYA4
AC109466.1
AGBL4
NPAS3
DOCK2
ELL2
SRGN
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SYK
ELMO1
DOCK4
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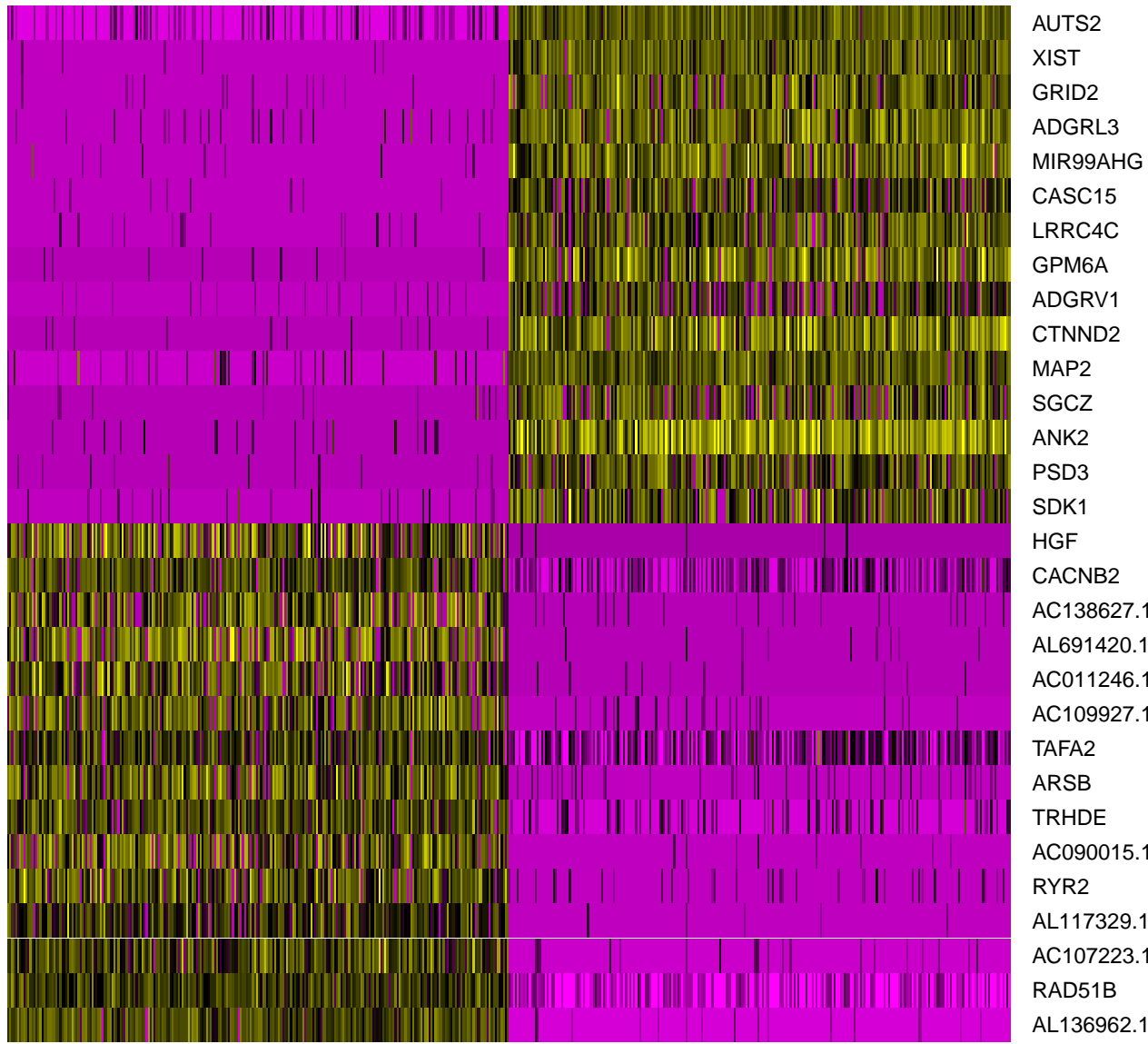
PC_1



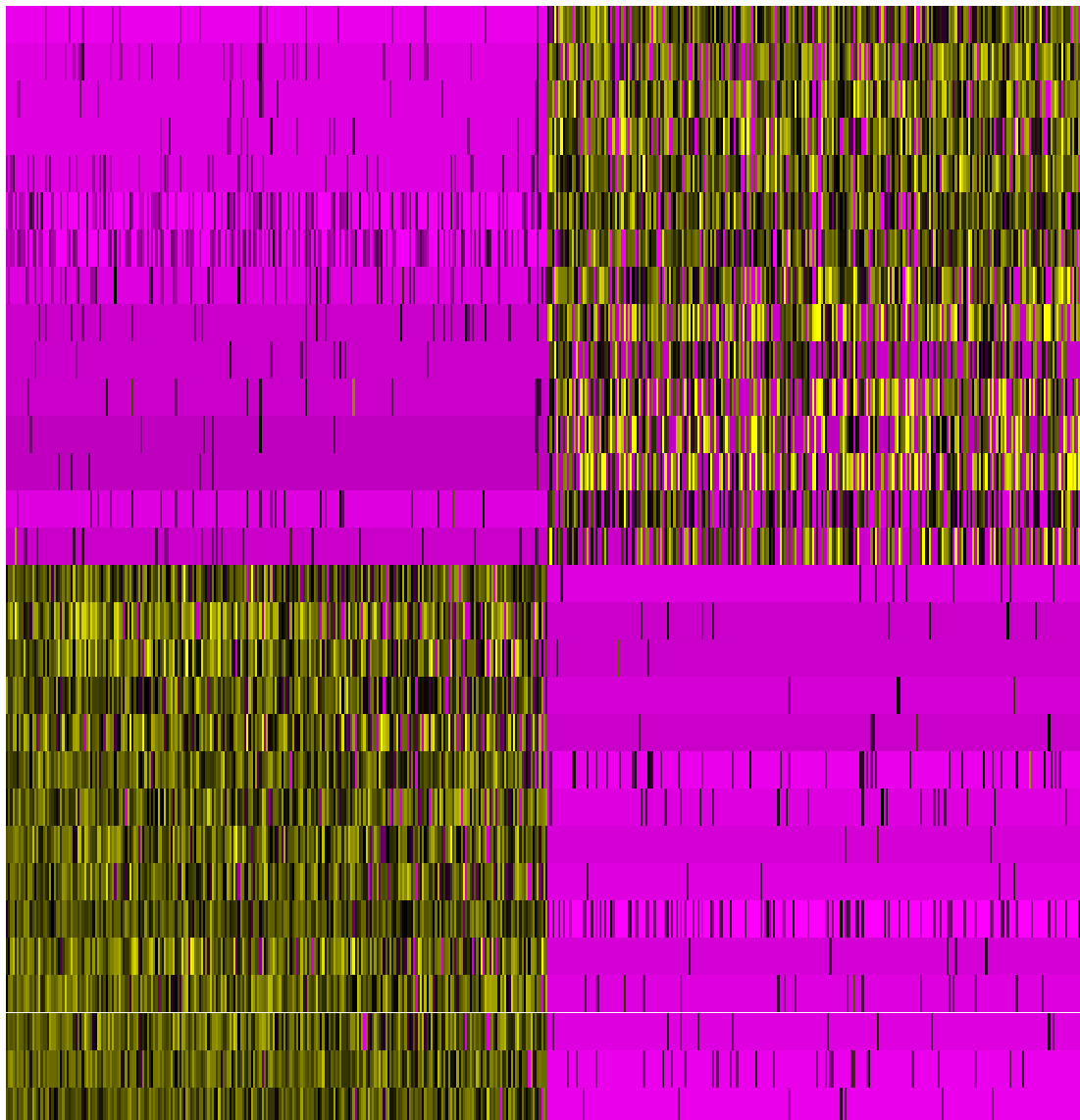
PC_1



PC_1

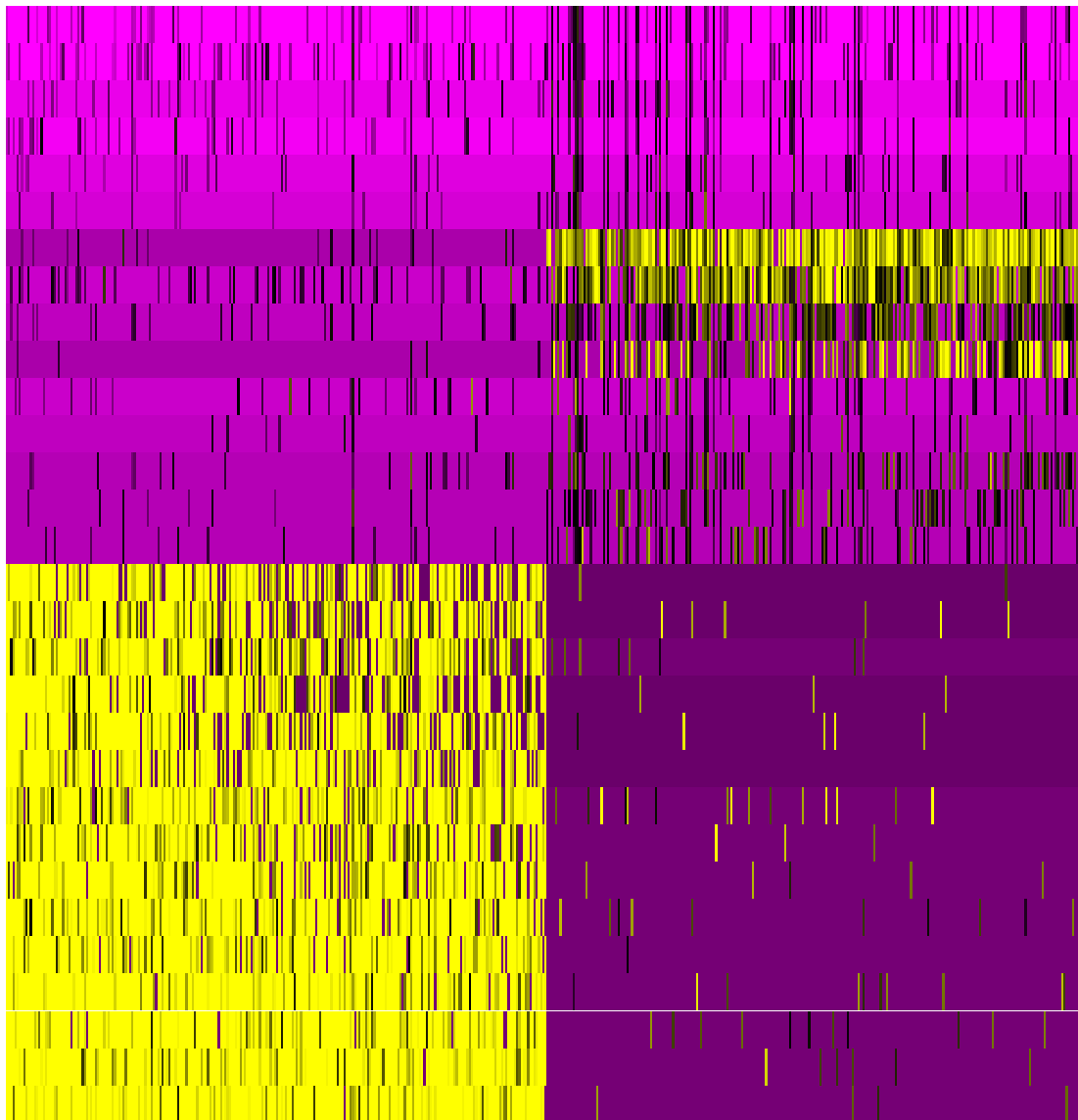


PC_1

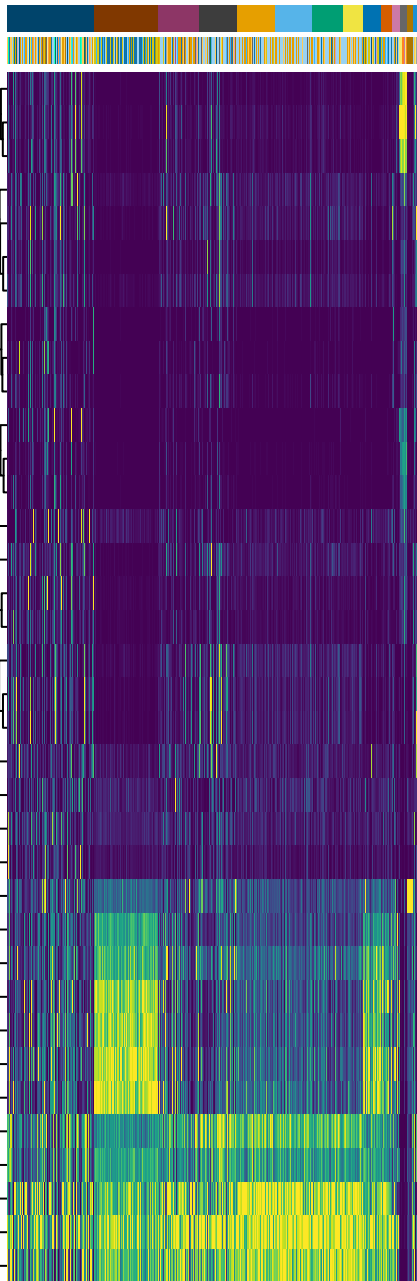
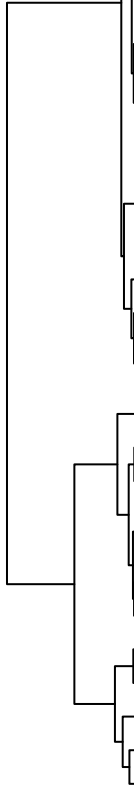


MEOX2
GRIA4
LINC02253
ZFHX4
COL21A1
TAFA2
SUCO
MCTP2
DLGAP2
SCHLAP1
SMOC2
RBFox3
ZFHX4-AS1
MECOM
CREB5
SLIT2
DPP6
OTX2-AS1
MYOF
AC092691.1
DISP1
MAP2
AL033504.1
GLIS3
RMST
CDH2
HTR2C
SAMD5
TRPM3
LSAMP

PC_1



Scores



Clusters
Labels

Monocyte
Macrophage
DC
HSC_CD34+
Pro-B_cell_CD34+
GMP
CMP
BM
Myelocyte
Pro-Myelocyte
Neutrophils
HSC_-G-CSF
Pre-B_cell_CD34-
Platelets
B_cell
NK_cell
T_cells
MEP
Erythroblast
BM & Prog.
Gametocytes
Hepatocytes
Epithelial_cells
Keratinocytes
Endothelial_cells
Tissue_stem_cells
MSC
Osteoblasts
Chondrocytes
Smooth_muscle_cells
Fibroblasts
Embryonic_stem_cells
iPS_cells
Astrocyte
Neurons
Neuroepithelial_cell



Clusters

0
1
2
3
4
5
6
7
8
9
10
11
12
13
14
15

Labels

Astrocyte
B_cell
BM
BM & Prog.
Chondrocytes
CMP
DC
Embryonic_stem_cells
Endothelial_cells
Epithelial_cells
Erythroblast
Fibroblasts
Gametocytes
GMP
Hepatocytes
HSC_-G-CSF