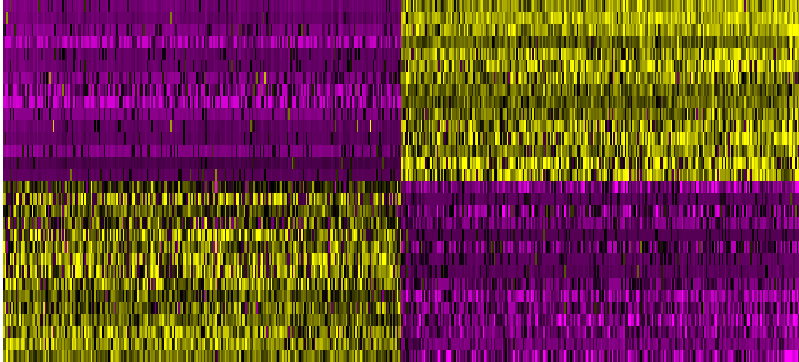
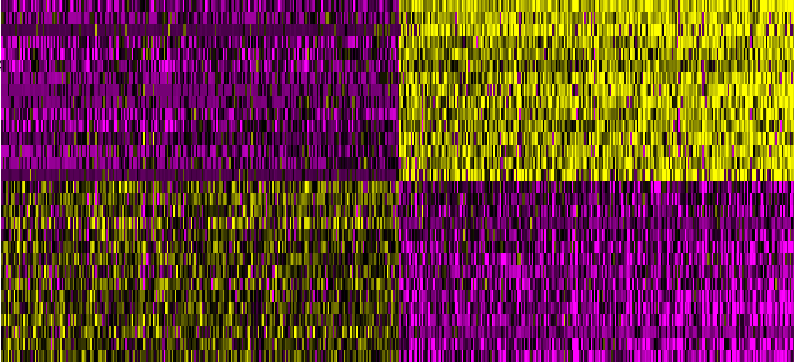


PC_1



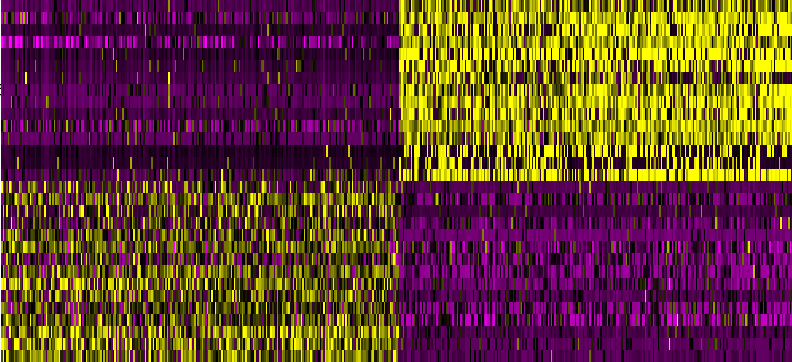
ASTL
EPCAM
ENSGALG000000042
AP1S3
MPZL3
SLC34A2
RNF152
HMG81
NUCKS1
MYL4
CDH2
COTL1
MFAP2
LDHB
STMN1

PC_2



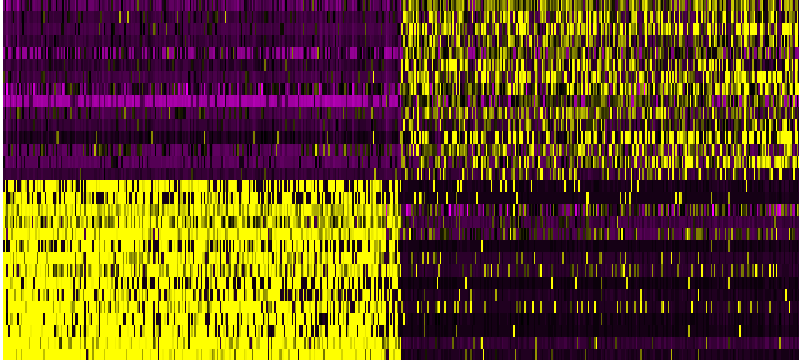
LIN28A
TPR
HMG81
ENSGALG000000025
ZC3H13
FN1
HNRNPM
PGK1
TIMM10
MT-ND1
OTUB1
PGAM1
RPS26
CKB
RPL38

PC_3



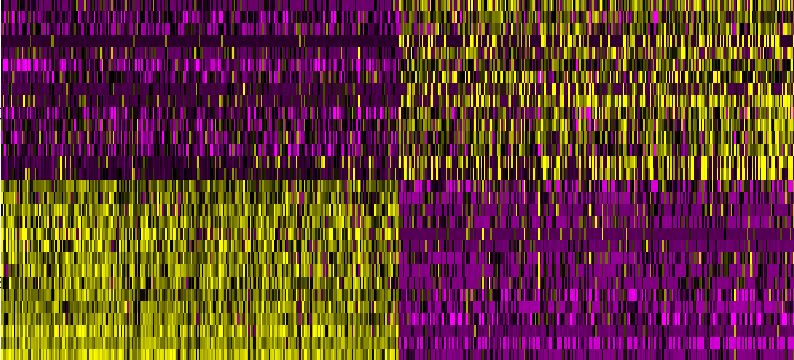
FN1
PRDX1
CSRP2
KRT18
ENSGALG000000000
FABP3
ITGB3
CAPN11
EXFABP
ENSGALG000000000
CKB
USP44
YEATS4
LMO1
SOX21

PC_4



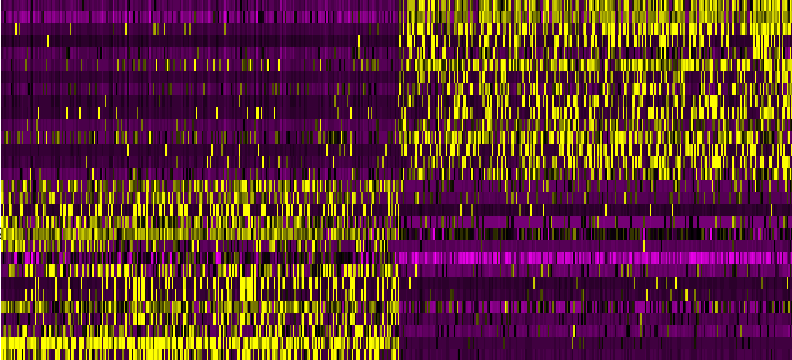
SST
SIX3
PAX6
SEM1
ENSGALG000000030
SIX1
KRT18
ERMN
TMSB4X
LMO4
SOX8
ENSGALG000000053
RASL11B
SOX10
TFAP2B

PC_5



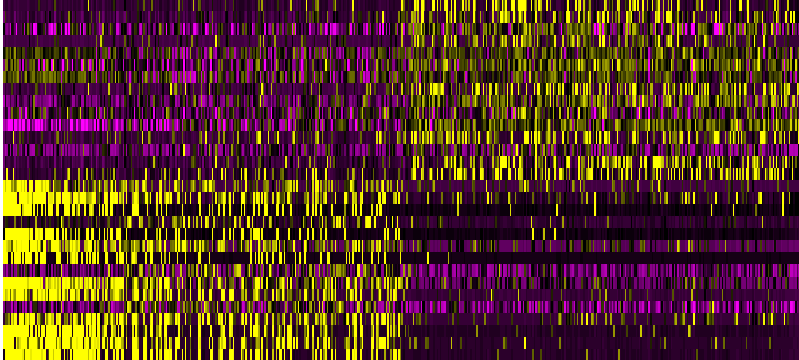
RRM2
CCNE2
NPM1
HIST1H103
THOC7
HSP90B1
SIX3
TOP2A
BUB1B
ENSGALG000000012
ASPM
AURKA
CDC20
CENPF
KPNB2

PC_6



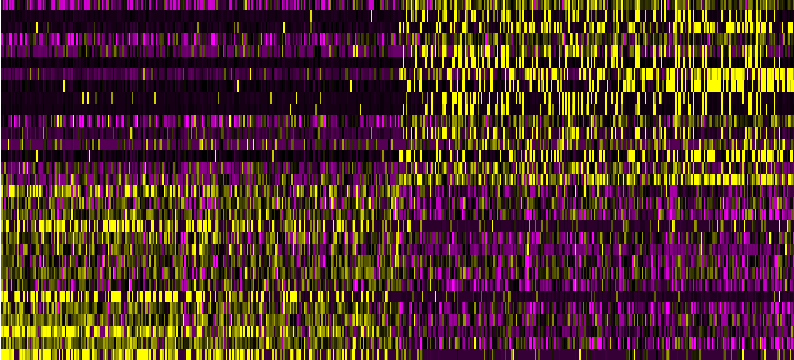
SHISA2
ENSGALG000000000
SOX5
PAX6
FGFR1
CXCR4
TOX
NCAM1
FAM181A
LDHB
AP1S3
PAX2
LFNG
SP5
NKX6-2

PC_7



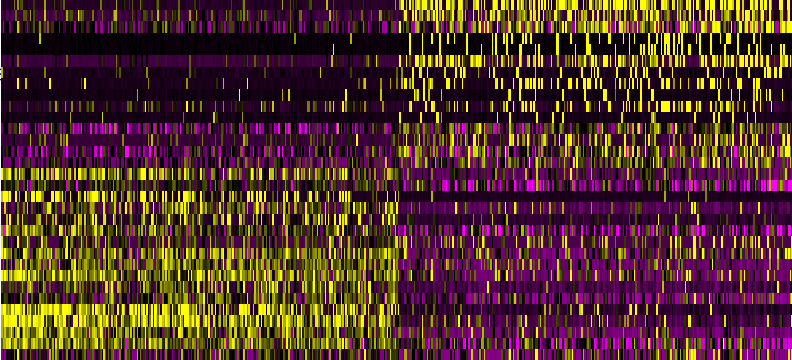
HESX1
CDH11
UCH-L1
SP5
THOC7
EXOC6
SIX3
COL4A1
UPK1B
EPAS1
SLCSA1
RGN
FABP7
FIBIN
PDZK1IP1

PC_8



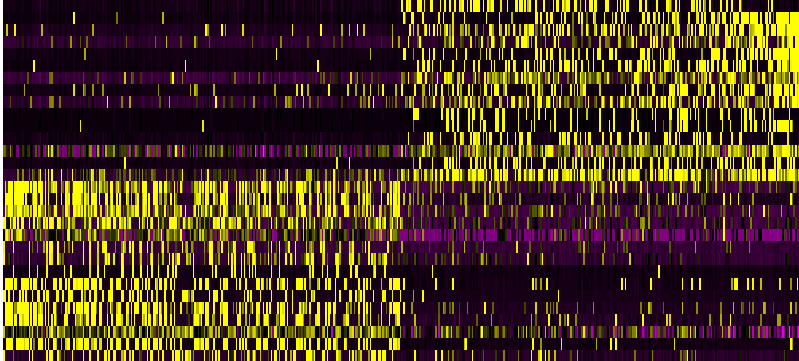
VIM
PITX2
CARHSP1
ENSGALG000000043
EOMES
MT-ND3
BRCA2
LDHA
FDP5
HSPD1
TM7SF2
CDK1
SFRP2
HSPA8
RGN
NDRG1

PC_9



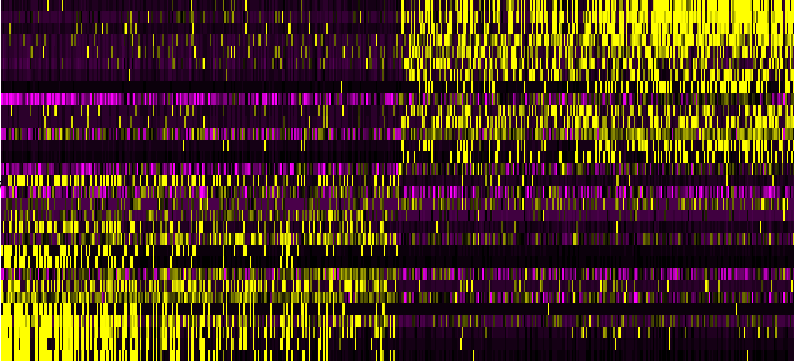
Meis2a.2
PAX2
PDGFA
LAMP5
PTN
HSP90AA1
NCL
FKBP11
WNT1
WNT4
MID1
ENSGALG000000000
OTX1
ZFHX4
HES4
JARID2

PC_10



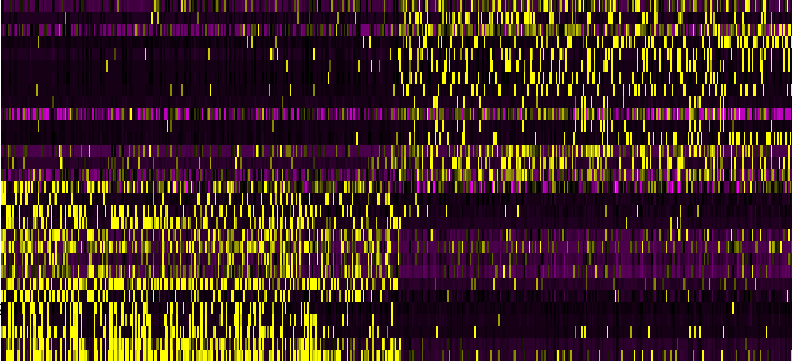
MSX2
NEFM
CDH6
MAFB
RASD2
RSP01
ZNF503
ENSGALG000000032
GFPT2
IFITM3
WNT4
RFTN2
RASL11B
LDHB
NKX6-2

PC_11



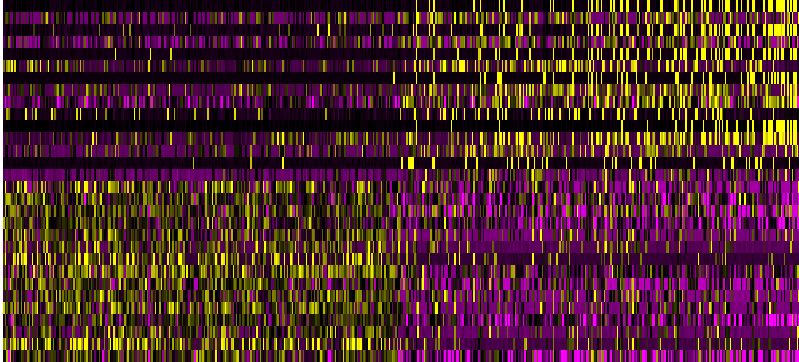
EN1
APCDD1
NR2F2
SP8
NPM3
TNFRSF19
PAX2
NCL
ID1
CDH11
LMO1
NKX2-4
TIMP3
ENSGALG000000042
DBX1
NKX2-1

PC_12



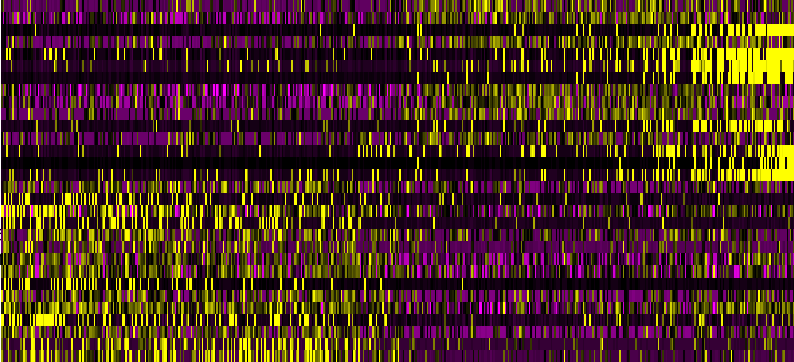
OLIG2
MSX2
GSC
BMP5
ARL4C
EPAS1
TGFB2
SEM1
CRYAB
SIX3
HESX1
GBX2
SP9
FGF8
PTN

PC_13



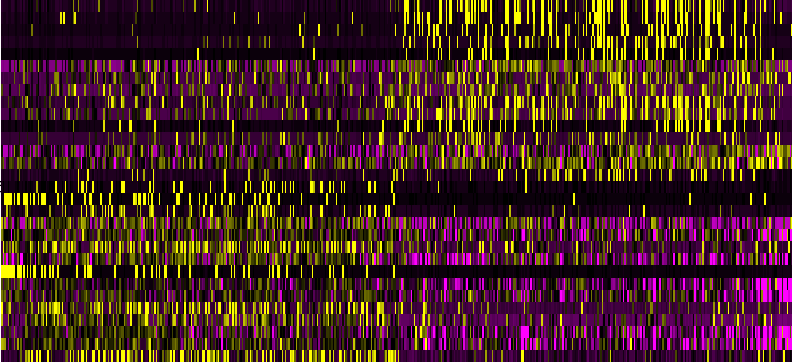
BUB1B
CDC20
DLX5
CDC25B
DLX6
NUAK1
BMP5
GJA1
TOP2A
ENSGALG000000025
ENSGALG000000010
DGUOK
SPC25
FBXO5
RRM2

PC_14



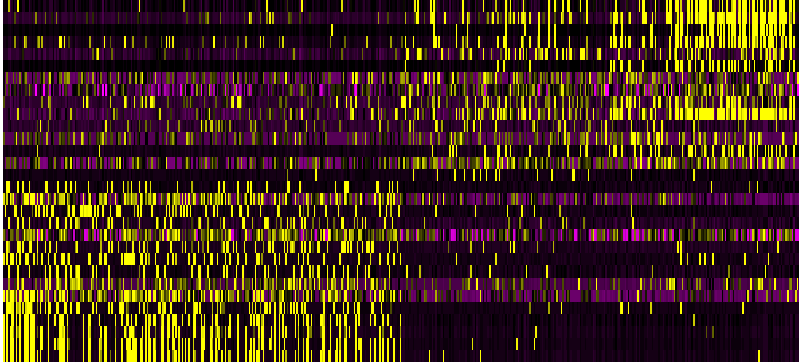
PTTG2
BUB1B
FILIP1
CDK1
PLK1
UNG
TLX1
ENSGALG000000025
SEM1
SPON1
DGUOK
NKX2-1
H3F3B
NUSAP1
HIST1H103

PC_15



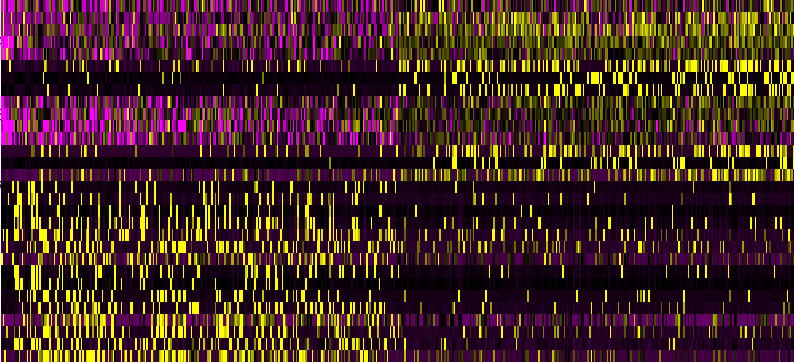
PAX2
WNT1
NUSAP1
SGO1
GALNT6
ENSGALG000000000
H3F3B
EMX2
OLIG2
MT-ND1
CDK1
MT-CYTB
PDGFA
MT-ND4
PAX6

PC_16



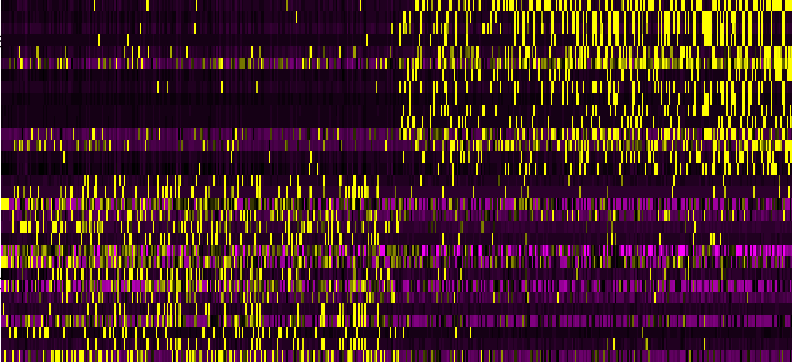
CYTL1
ENSGALG000000042
VEGFD
CALM1
ENSGALG000000043
BRCA2
SPC25
ENSGALG000000042
EPAS1
VIM
LITAF
GALNT6
FGF8
ADMP
EOMES

PC_17



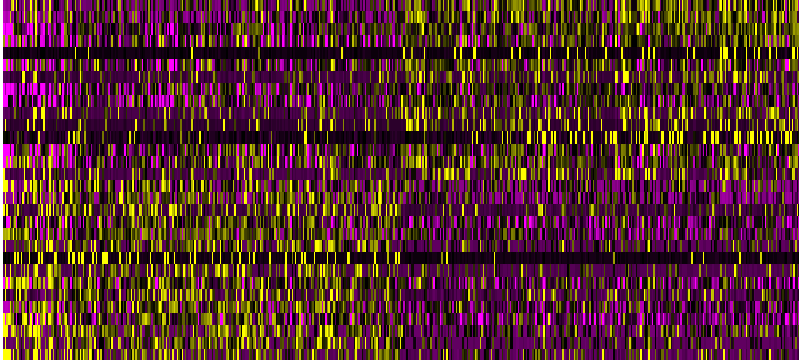
WDR1
ENSGALG000000043
FAM184B
GATA3
MT-ND6
HSPE1
RASD2
PHACTR1
GnRH-I
ANXA2
SP5
ENSGALG000000050
HMP19
FIBIN
PDGFA

PC_18



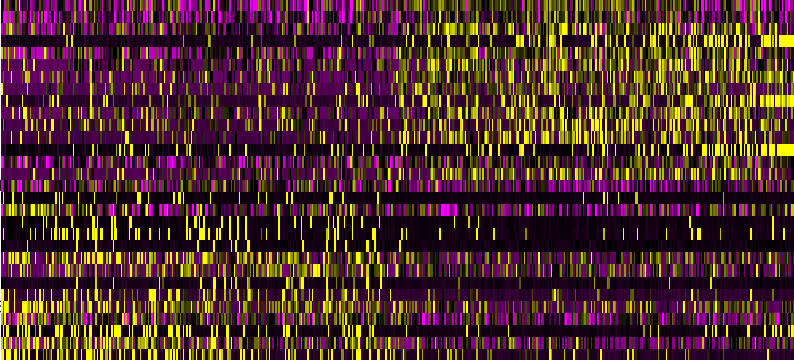
CXCL14
DBX1
ENSGALG000000000
RIPPLY3
NKX2-4
MYO3B
PHLDA2
VEGFD
EPHA1
SST
MBP
JARID2
WDR1
EOMES
RASD2
ENSGALG000000000

PC_19



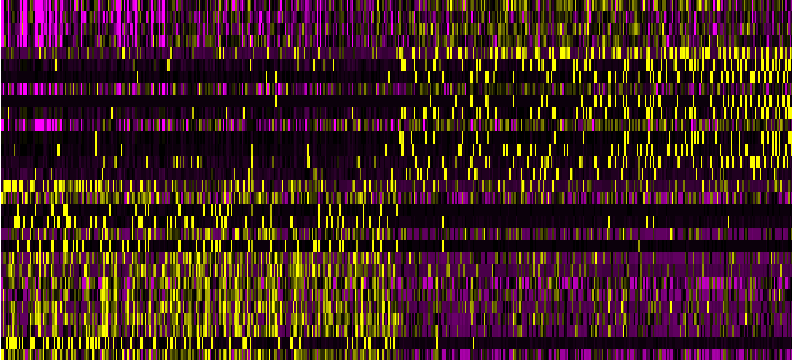
HNRNPM
YWHAE
SRRM1
RPL38
KDM1A
DLX6
EIF3A
ENSGALG000000030
KLF6
CLDN1
CXCL14
HSPA5
ENSGALG000000043
SEMA3F
SLC38A2

PC_20



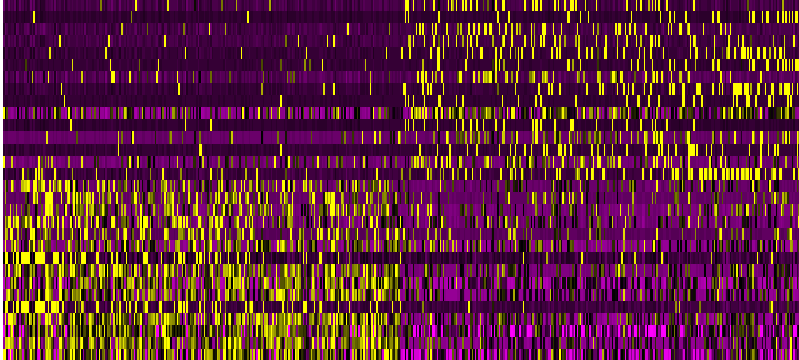
HSPA8
GPC3
MSMO1
SERPINH1
INSIG1
CDH11
UBB
MT-ND1
MT-ND3
CXCL14
NET1
ADMP
SMAD1
GATA3
FAM184B

PC_21



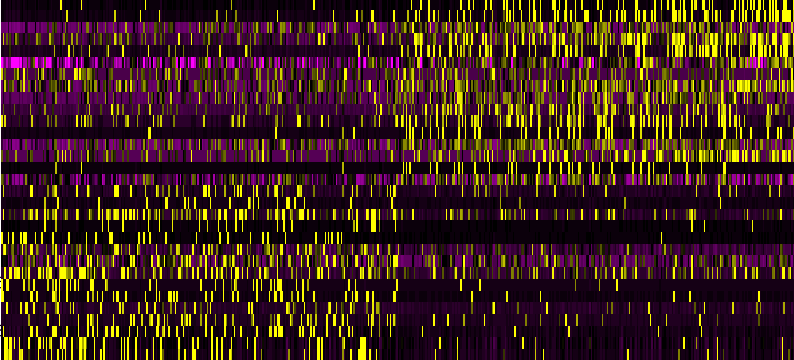
MT-CYTB
MT-ATP6
PITX2
MT-CO1
PITX1
ENSGALG000000000
DBX1
PHLDA2
KROX20
INSIG1
SQLE
ENSGALG000000000
FDTF1
MSMO1
FDP5

PC_22



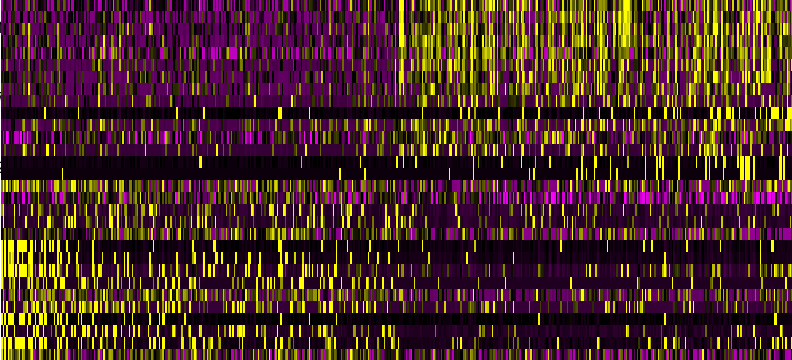
RASD2
APCDD1
MAFB
DHRS3
DMD
ELAVL4
ITGA6
PDGFA
DHCR24
HHLA2
PITX1
ENSGALG000000000
PITX2
ENSGALG000000043
FDP5

PC_23



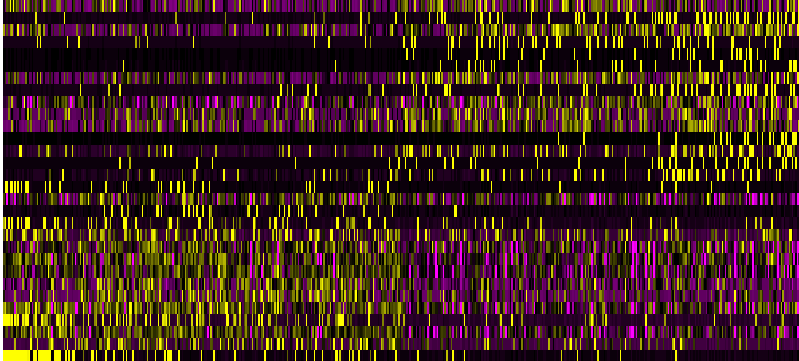
RIPPLY3
ENSGALG000000000
HMP19
MID1
ENSGALG000000022
OPTC
HES5
ENSGALG000000002
ZIC1
PTN
FOXA1
FKBP11
RSP02
SGK1
RSP01
PITX2

PC_24



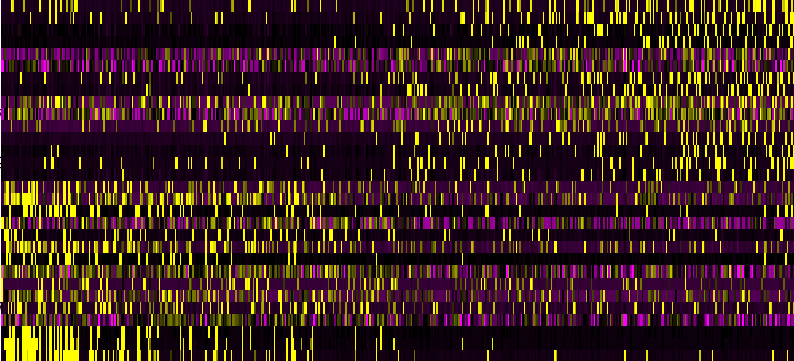
ENSGALG000000000
CYP51A1
FDTF1
SQLE
NKX2-5
EIF4G2
NKX2-1
ATP1B1
MYC
ZC3H6
CYP26C1
HAS2
NDRG1
SGK1
CCND1

PC_25



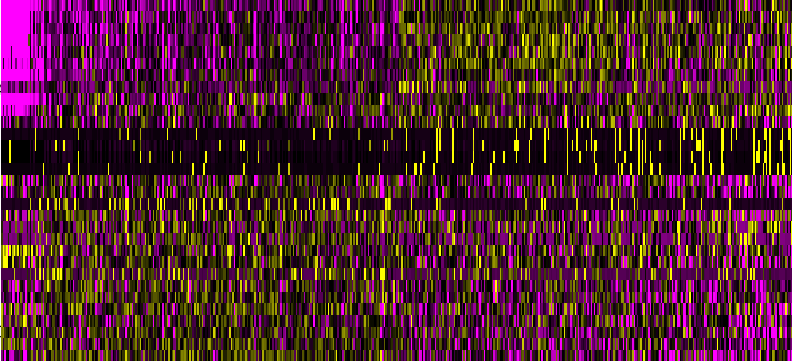
PHACTR1
RSP02
EBF2
JAG2
ENSGALG000000022
VAX1
ENSGALG000000043
EN2
SP9
Meis2a.2
RPS26
LDHA
ENSGALG000000032
RPS29
MSX2

PC_26



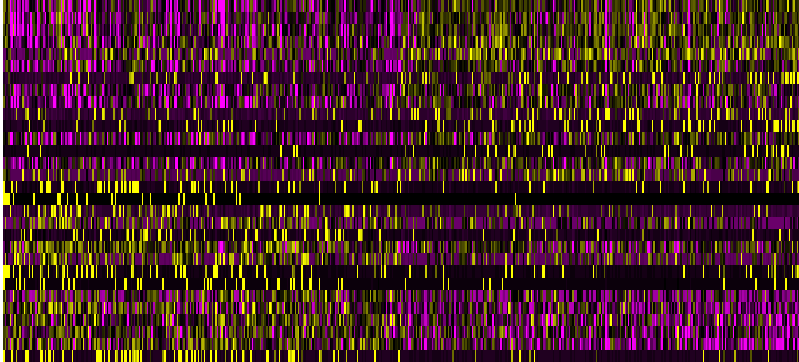
DLL1
EBF2
ENO1
ENSGALG000000054
CXXC5
DNASE2B
JAG2
CRABP-I
RASD2
CXCL14
HOXB2
LRP1B
CLEC19A
ENSGALG000000050
MAFB

PC_27



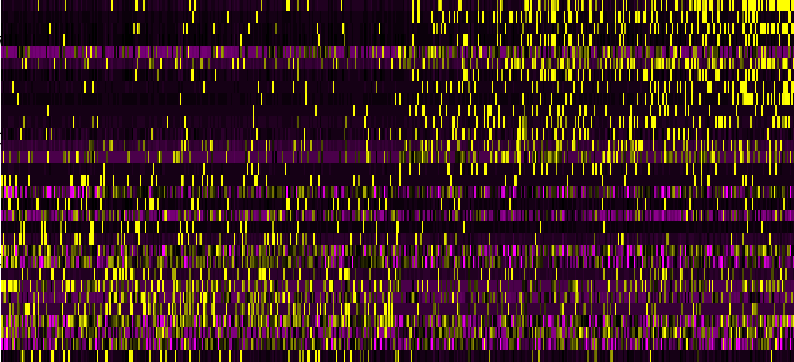
MT-ATP6
MT-ND1
MT-ND5
MT-CYTB
MT-ND6
ENO1
ENSGALG000000000
CDH6
UQCRI1
ID2
ENSGALG000000000
UCH-L1
HSPA8
TCF12
SRF7
EIF4G2

PC_28



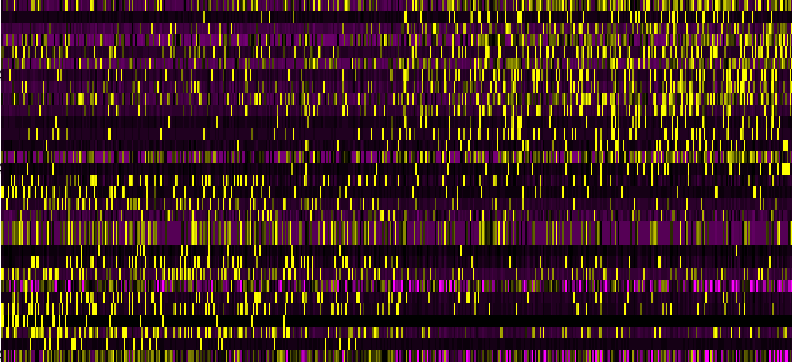
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ENSGALG000000032
RPL38
UQCRI1
CYP26A1
ENSGALG000000030
NDUFA13
WNT6
LINGO1
GPC3
ENSGALG000000043
WNT3A
ARL4C
PGAM1
WNT1

PC_29



EPHA7
MSX2
ORAI1
ENSGALG000000054
CDH6
GBE
BP1FB3
ENSGALG000000032
MT-ND1
LDHA
SGK1
EIF4G2
FAT1
RBM38
JARID2
ENSGALG000000043

PC_30



MSX1
MERTK
GBX2
SPAM1
SP5
EOMES
KAZALD1
NKX2-3
MAFB
CA2
BRCA2
ENSGALG000000000
HSPB1
HMP19
LHFPL6
HSP90AA1