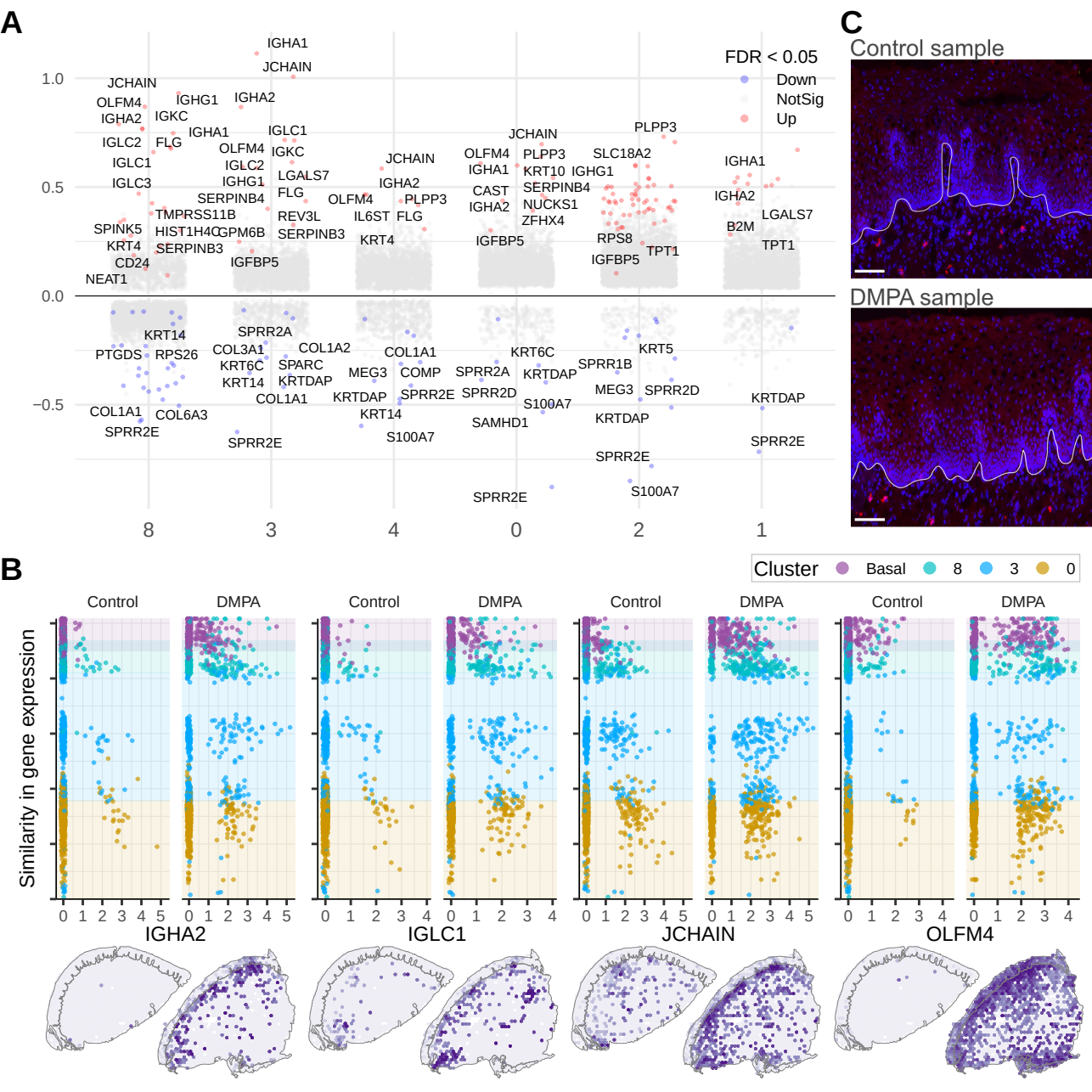
A
FDR < 0.05
Down
NotSig
Up
1.0
0.5
0.0
−0.5
IGHA1
JCHAIN
JCHAIN
OLFM4
IGHG1
IGHA2
IGHA2
IGKC
IGHA1
IGLC1
IGLC2
FLG
OLFM4
IGKC
IGLC1
IGLC2
IGLC3
IGHG1
LGALS7
SERPINB4
FLG
TMPRSS11B
REV3L
SPINK5
HIST1H4C
GPM6B
SERPINB3
KRT4
SERPINB3
CD24
IGFBP5
NEAT1
JCHAIN
OLFM4
IGHA2
PLPP3
IL6ST
FLG
KRT4
OLFM4
PLPP3
IGHA1
KRT10
CAST
SERPINB4
IGHA2
NUCKS1
ZFHX4
IGFBP5
JCHAIN
PLPP3
SLC18A2
IGHG1
RPS8
TPT1
IGFBP5
IGHA1
IGHA2
LGALS7
B2M
TPT1
KRT14
PTGDS
RPS26
SPRR2A
COL3A1
COL1A2
KRT6C
SPARC
KRT14
KRTDAP
COL1A1
COL6A3
SPRR2E
SPRR2E
MEG3
KRTDAP
KRT14
S100A7
COL1A1
COMP
SPRR2E
KRT6C
SPRR2A
KRTDAP
SPRR2D
S100A7
SAMHD1
SPRR2E
KRT5
SPRR1B
MEG3
SPRR2D
KRTDAP
SPRR2E
S100A7
KRTDAP
SPRR2E
8
3
4
0
2
1
C
Control sample
DMPA sample
B
Cluster
Basal
8
3
0
Control
DMPA
Similarity in gene expression
0 1 2 3 4 5
0 1 2 3 4
IGHA2
IGLC1
JCHAIN
OLFM4