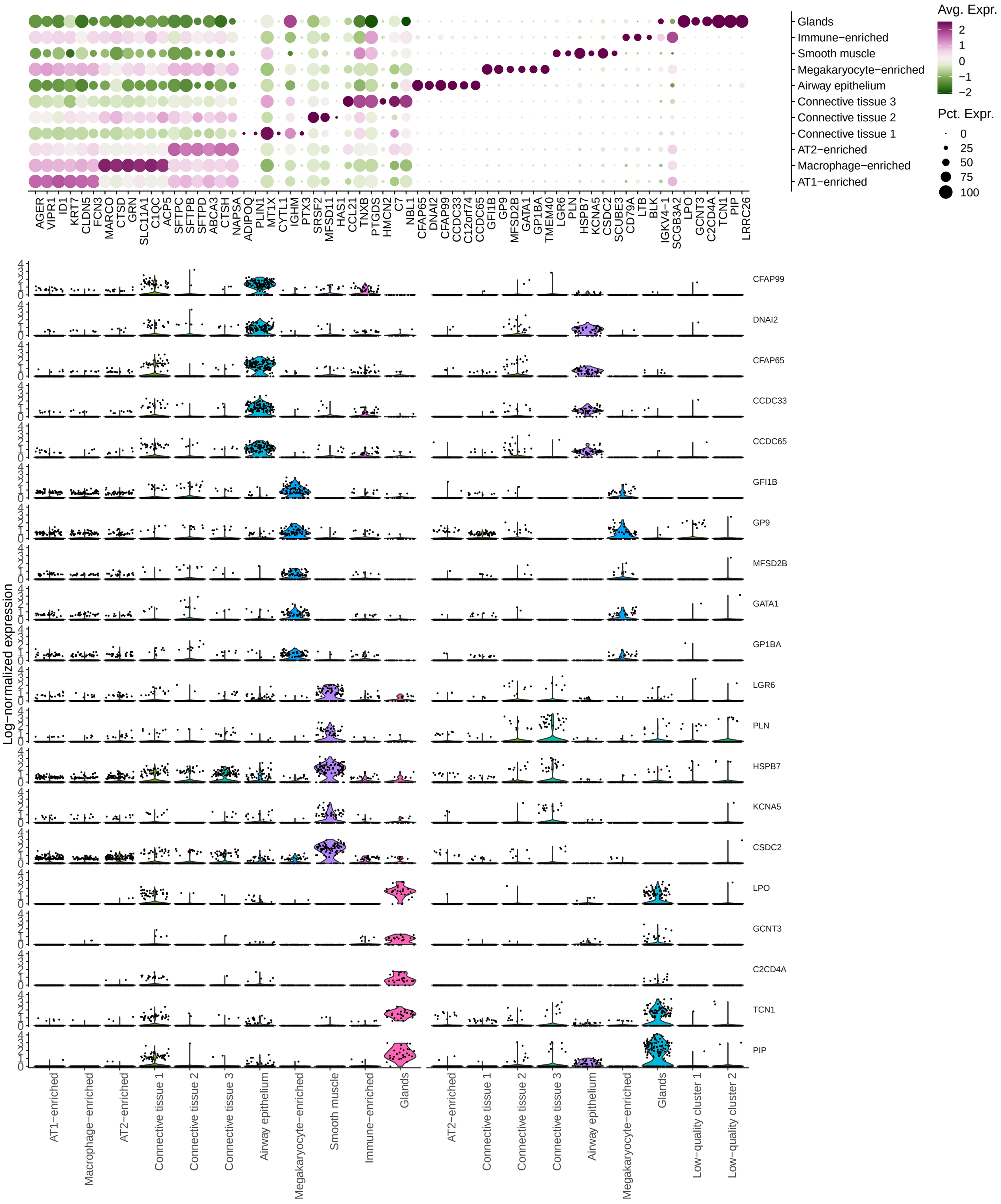
Glands
Immune−enriched
Smooth muscle
Megakaryocyte−enriched
Airway epithelium
Connective tissue 3
Connective tissue 2
Connective tissue 1
AT2−enriched
Macrophage−enriched
AT1−enriched
Avg. Expr.
2
1
0
−1
−2
Pct. Expr.
0
25
50
75
100
AGER VIPR1 ID1 KRT7 CLDN5 FCN3 MARCO CTSD GRN SLC11A1 C1QC ACP5 SFTPC SFTPB SFTPD ABCA3 CTSH NAPSA ADIPOQ PLIN1 MT1X CYTL1 IGHM PTX3 SRSF2 MFSD11 HAS1 CCL21 TNXB PTGDS HMCN2 C7 NBL1 CFAP65 DNAI2 CFAP99 CCDC33 C12orf74 CCDC65 GFI1B GP9 MFSD2B GATA1 GP1BA TMEM40 LGR6 PLN HSPB7 KCNA5 CSDC2 SCUBE3 CD79A LTB BLK IGKV4−1 SCGB3A2 LPO GCNT3 C2CD4A TCN1 PIP LRRC26
Log−normalized expression
CFAP99
DNAI2
CFAP65
CCDC33
CCDC65
GFI1B
GP9
MFSD2B
GATA1
GP1BA
LGR6
PLN
HSPB7
KCNA5
CSDC2
LPO
GCNT3
C2CD4A
TCN1
PIP
AT1−enriched
Macrophage−enriched
AT2−enriched
Connective tissue 1
Connective tissue 2
Connective tissue 3
Airway epithelium
Megakaryocyte−enriched
Smooth muscle
Immune−enriched
Glands
AT2−enriched
Connective tissue 1
Connective tissue 2
Connective tissue 3
Airway epithelium
Megakaryocyte−enriched
Glands
Low−quality cluster 1
Low−quality cluster 2