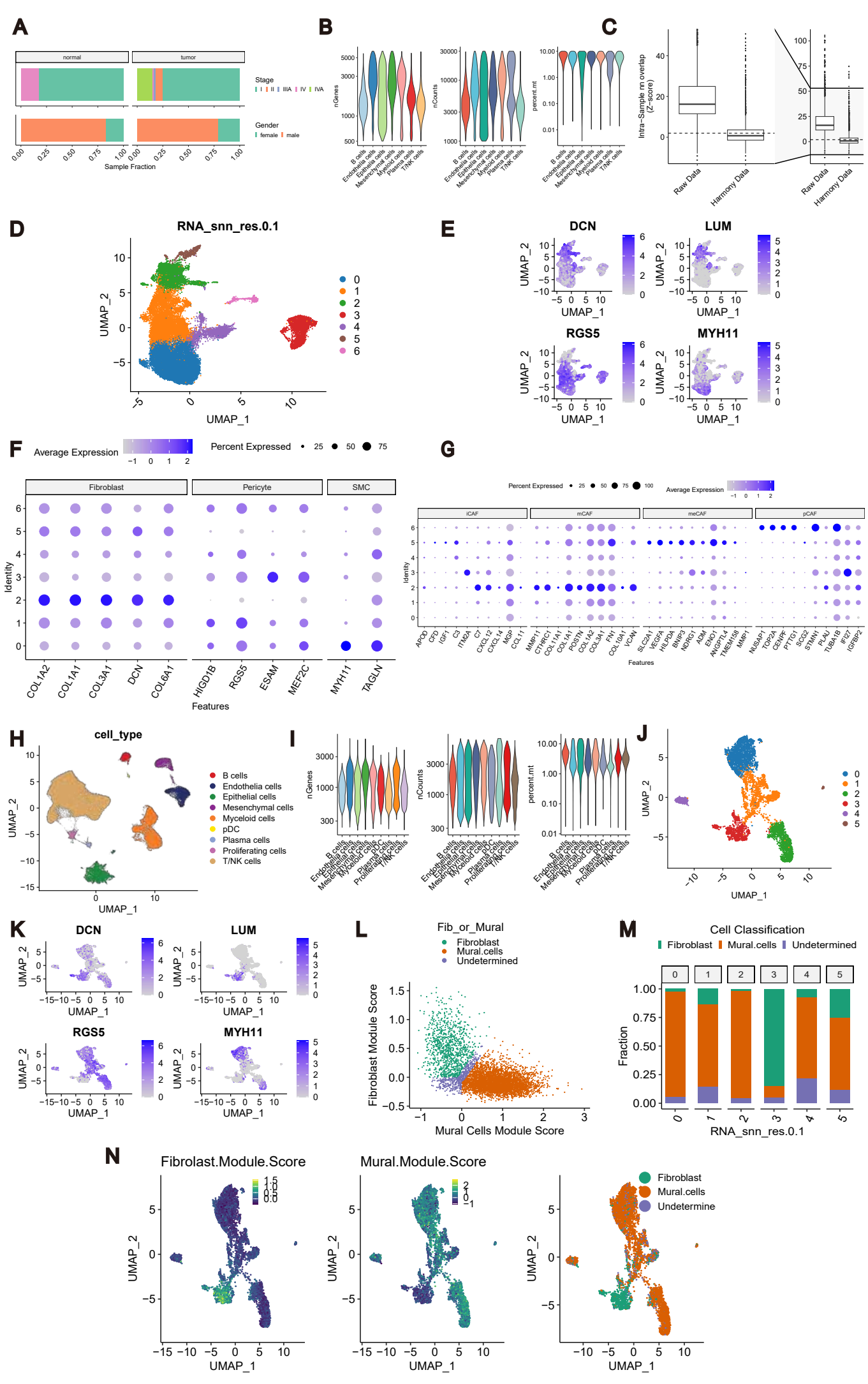
A
normal
tumor
Stage
I I II IIIA IV IVA
Gender
female male
Sample Fraction
B
nGenes
nCounts
percent.mt
B cells
Endothelia cells
Epithelia cells
Mesenchymal cells
Myeloid cells
Plasma cells
T/NK cells
C
Intra−Sample nn overlap (Z−score)
Raw Data
Harmony Data
D
RNA_snn_res.0.1
UMAP_1
UMAP_2
E
DCN
LUM
RGS5
MYH11
F
Average Expression
Percent Expressed
Fibroblast
Pericyte
SMC
Identity
COL1A2
COL1A1
COL3A1
DCN
COL6A1
HIGD1B
RGS5
ESAM
MEF2C
MYH11
TAGLN
Features
G
Percent Expressed
Average Expression
iCAF
mCAF
meCAF
pCAF
APOD
CFD
IGF1
C3
ITM2A
C7
CXCL12
CXCL14
MGP
CCL11
MMP11
CTHRC1
COL11A1
POSTN
COL4A2
COL3A1
FN1
COL10A1
VCAN
SLC2A1
VEGFA
HILPDA
BNIP3
NDRG1
ADM
ENO1
ANGPTL4
TMEM158
MMP1
NUSAP1
TOP2A
CENPF
PTTG1
SCG2
STMN1
PLAU
TUBA1B
IFI27
IGFBP2
H
cell_type
B cells
Endothelia cells
Epithelial cells
Mesenchymal cells
Myeloid cells
pDC
Plasma cells
Proliferating cells
T/NK cells
I
nGenes
nCounts
percent.mt
J
K
DCN
LUM
RGS5
MYH11
L
Fib_or_Mural
Fibroblast
Mural.cells
Undetermined
Fibroblast Module Score
Mural Cells Module Score
M
Cell Classification
Fibroblast
Mural.cells
Undetermined
Fraction
RNA_snn_res.0.1
N
Fibrolast.Module.Score
Mural.Module.Score
Fibroblast
Mural.cells
Undetermine