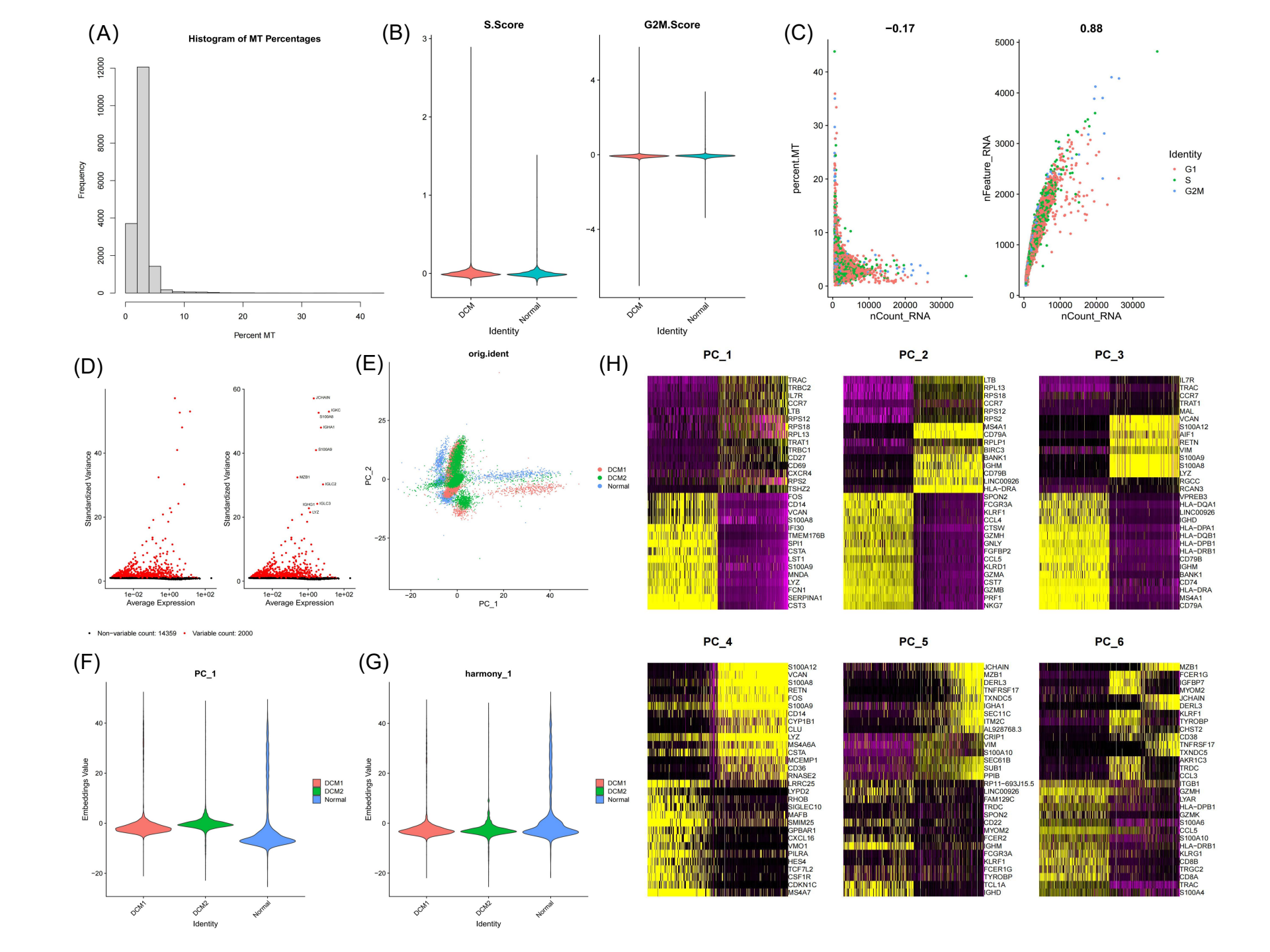
(A)
Histogram of MT Percentages
Frequency
Percent MT
(B)
S.Score
G2M.Score
DCM
Normal
Identity
(C)
−0.17
0.88
percent.MT
nFeature_RNA
nCount_RNA
Identity
G1
S
G2M
(D)
Standardized Variance
Average Expression
Non−variable count: 14359
Variable count: 2000
(E)
orig.ident
PC_1
PC_2
DCM1
DCM2
Normal
(F)
PC_1
Embeddings Value
(G)
harmony_1
(H)
PC_1
PC_2
PC_3
PC_4
PC_5
PC_6