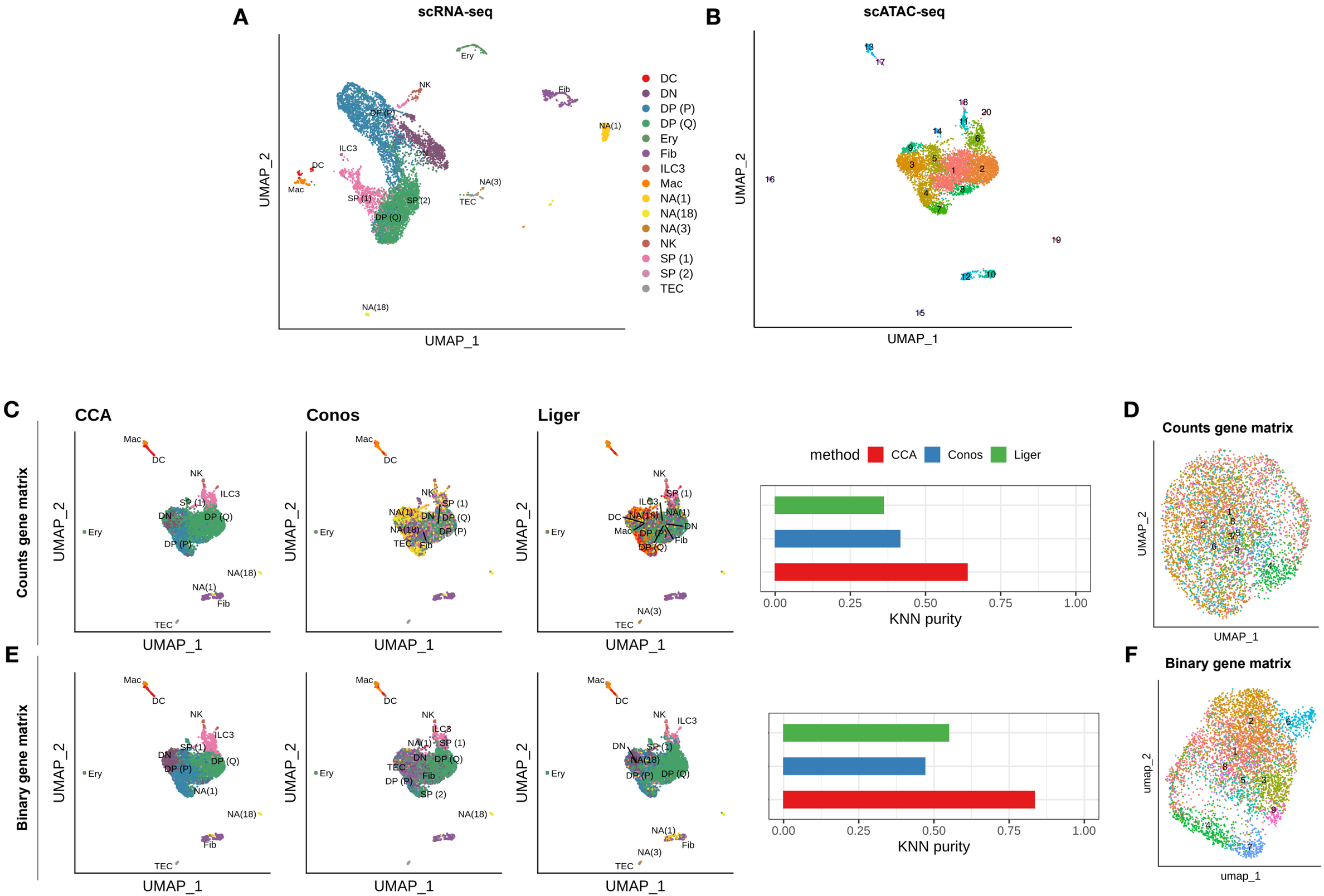
A
scRNA-seq
UMAP_2
UMAP_1
Ery
NK
Fib
NA(1)
DP (P)
ILC3
DN
DC
Mac
NA(3)
SP (1)
SP (2)
TEC
DP (Q)
NA(18)
DC
DN
DP (P)
DP (Q)
Ery
Fib
ILC3
Mac
NA(1)
NA(18)
NA(3)
NK
SP (1)
SP (2)
TEC
B
scATAC-seq
UMAP_2
UMAP_1
C
CCA
Conos
Liger
Counts gene matrix
method
CCA
Conos
Liger
0.00
0.25
0.50
0.75
1.00
KNN purity
D
Counts gene matrix
UMAP_2
UMAP_1
E
Binary gene matrix
F
Binary gene matrix
umap_2
umap_1