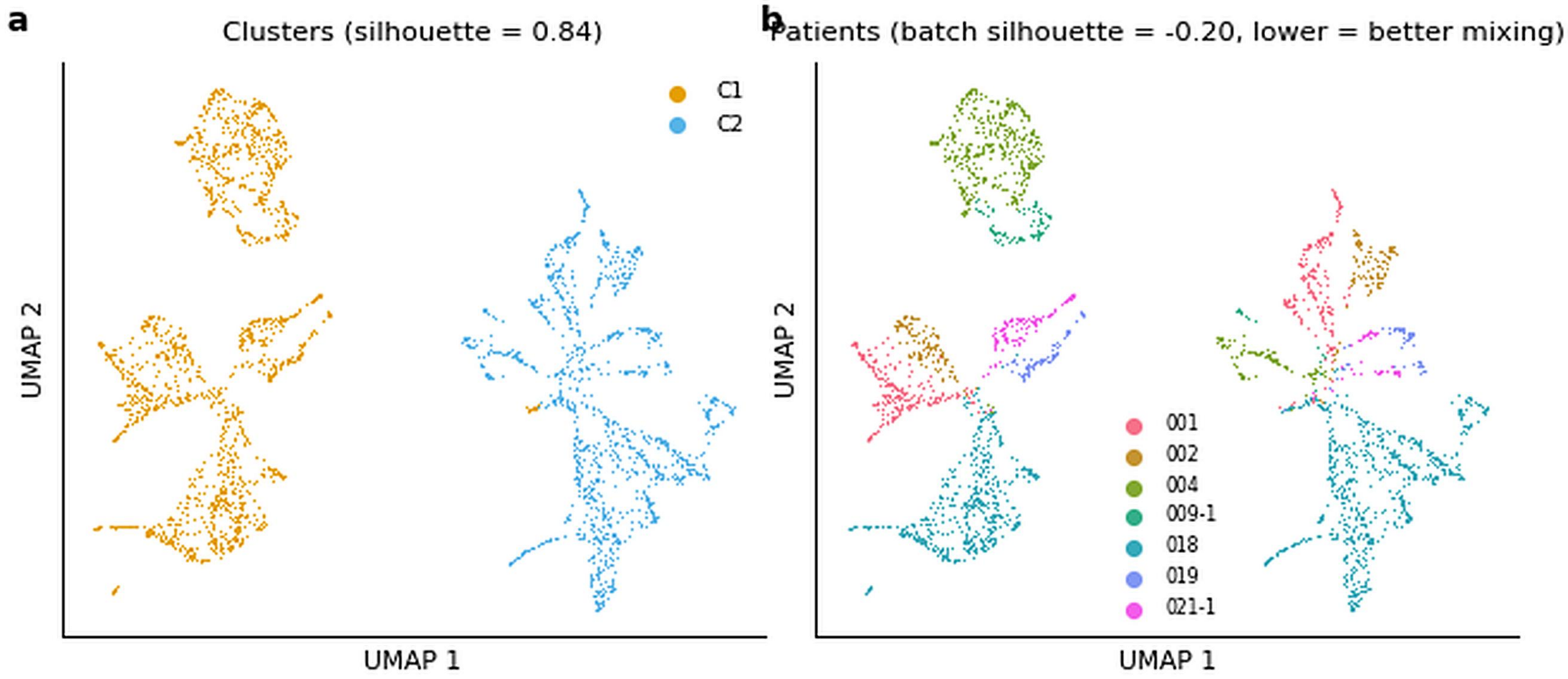

a
Clusters (silhouette = 0.84)
C1
C2
UMAP 2
UMAP 1
b
Patients (batch silhouette = -0.20, lower = better mixing)
001
002
004
009-1
018
019
021-1
UMAP 2
UMAP 1