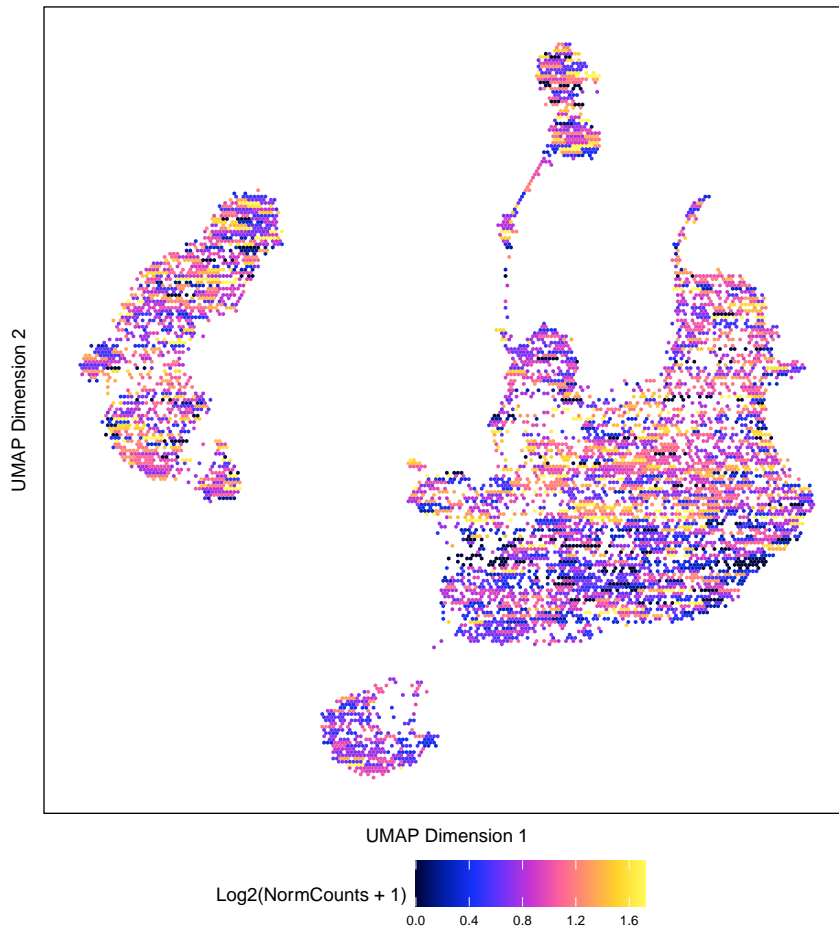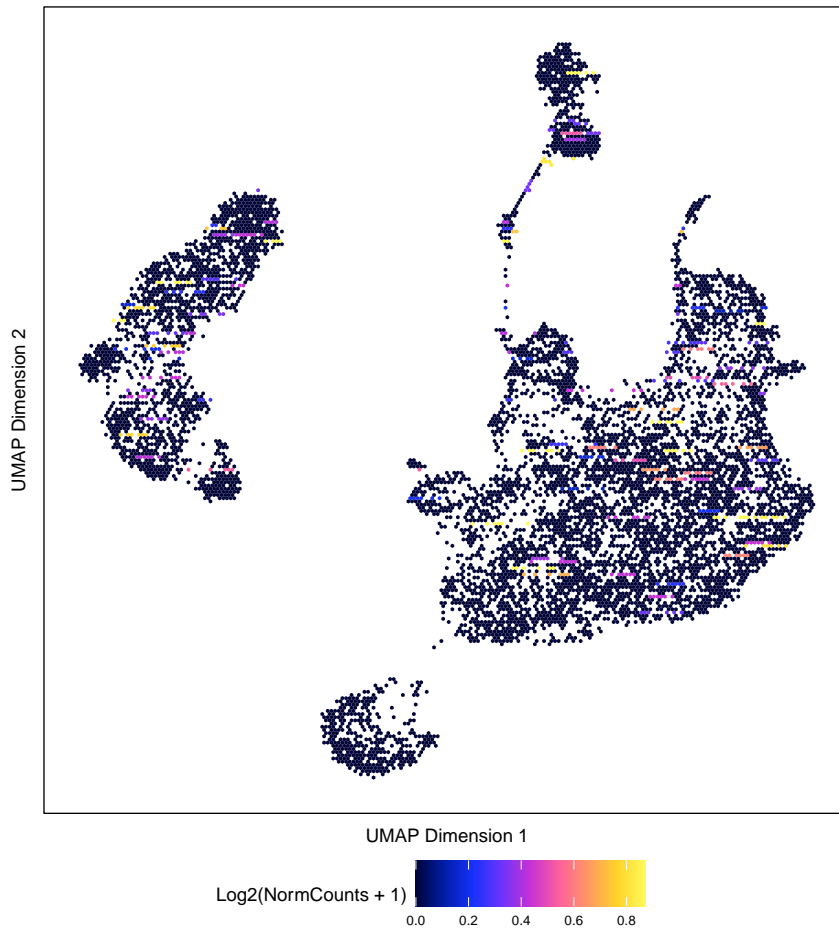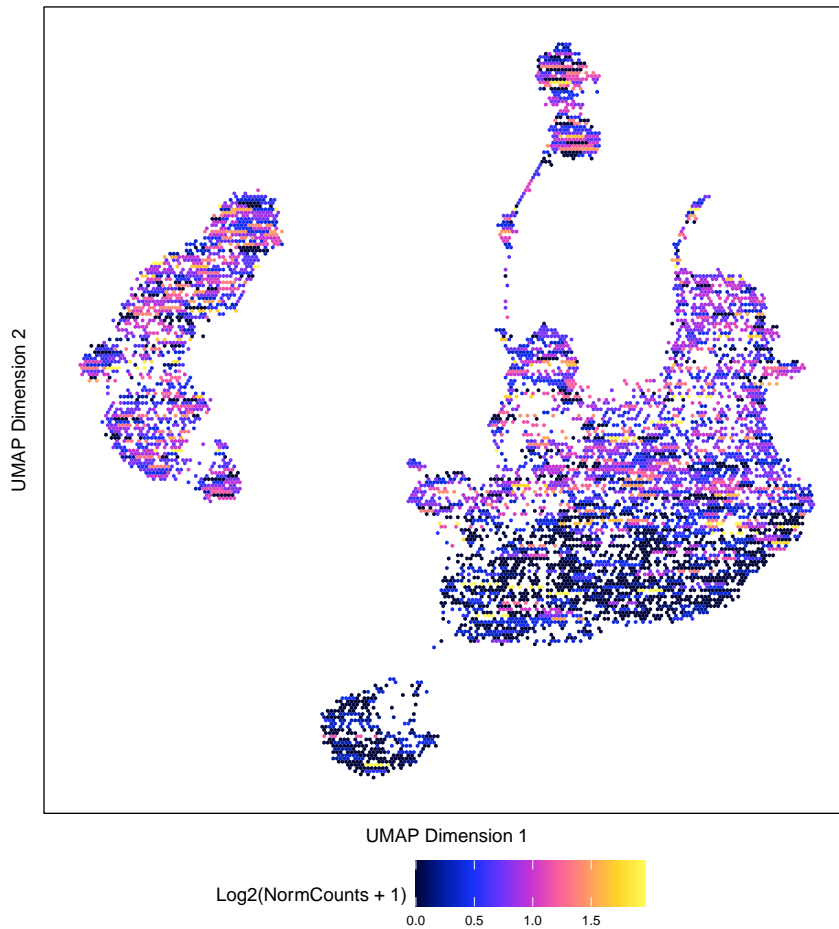UMAP of Iterative−LSI−1.0 colored by
GeneScoreMatrix : CD34

UMAP of Iterative−LSI−1.0 colored by
GeneScoreMatrix : GATA1

UMAP Dimension 2

UMAP Dimension 1

Log2(NormCounts + 1)

0.0   0.2   0.4   0.6   0.8

UMAP of Iterative−LSI−1.0 colored by
GeneScoreMatrix : PAX5

UMAP of Iterative−LSI−1.0 colored by
GeneScoreMatrix : MS4A1

UMAP Dimension 2

UMAP Dimension 1

Log2(NormCounts + 1)

0.0   0.2   0.4   0.6

UMAP of Iterative−LSI−1.0 colored by
GeneScoreMatrix : MME

UMAP Dimension 2
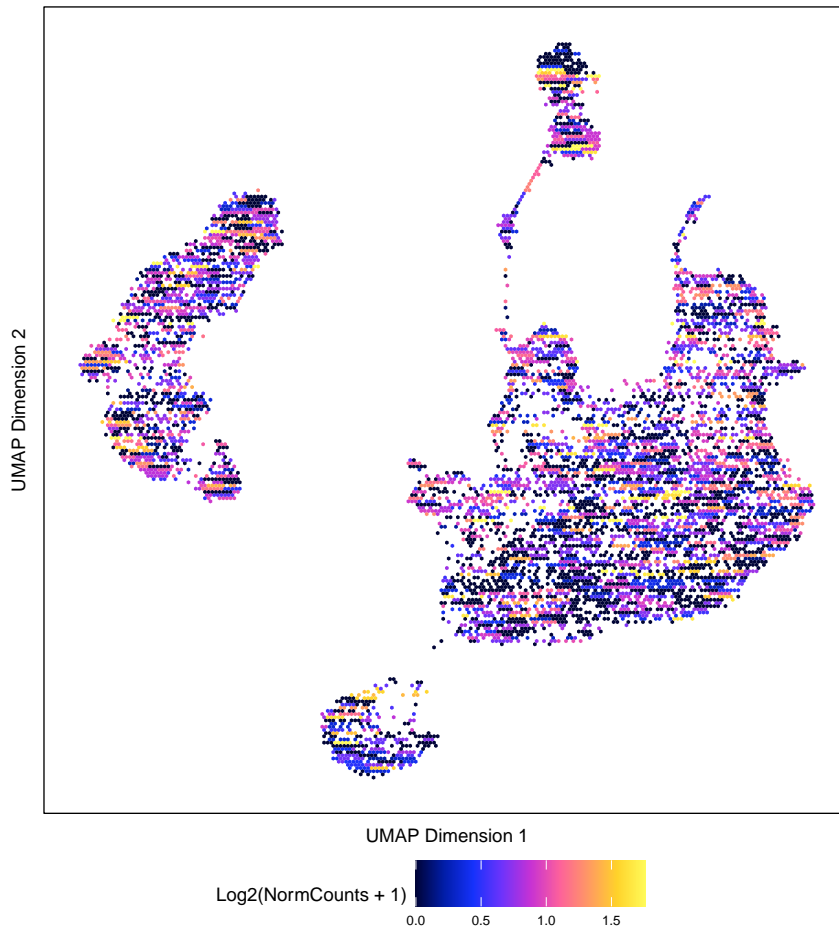
UMAP Dimension 1

Log2(NormCounts + 1)

0.00   0.25   0.50   0.75

UMAP of Iterative−LSI−1.0 colored by
GeneScoreMatrix : CD19

UMAP Dimension 2

UMAP Dimension 1

Log2(NormCounts + 1)

0.0   0.5   1.0   1.5
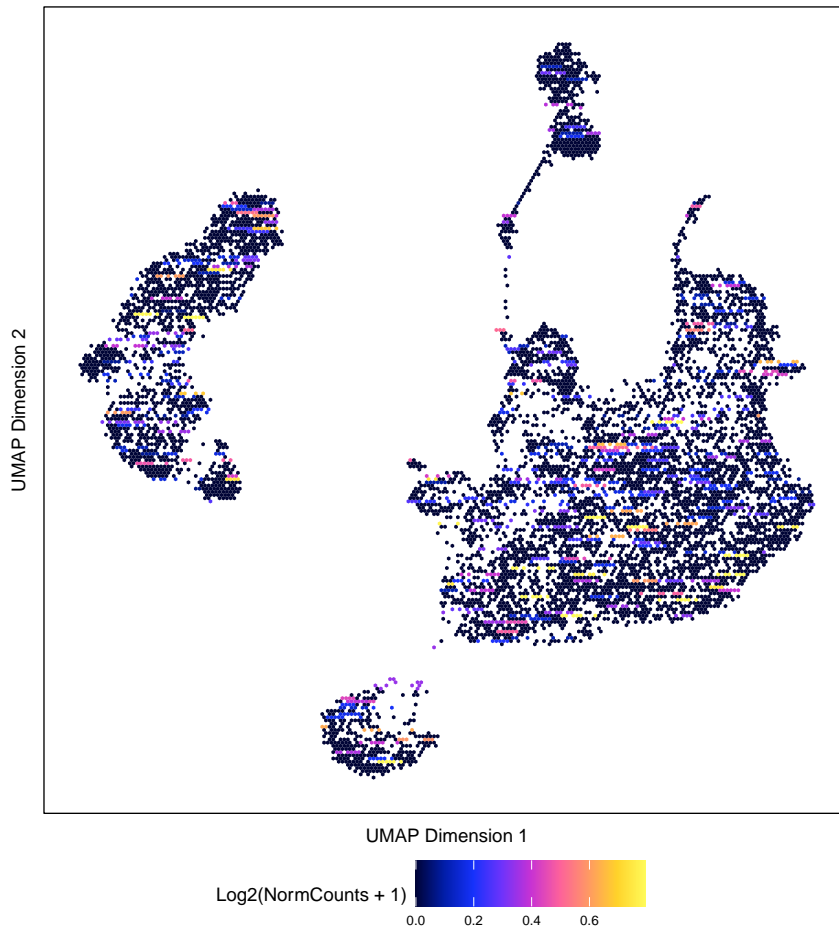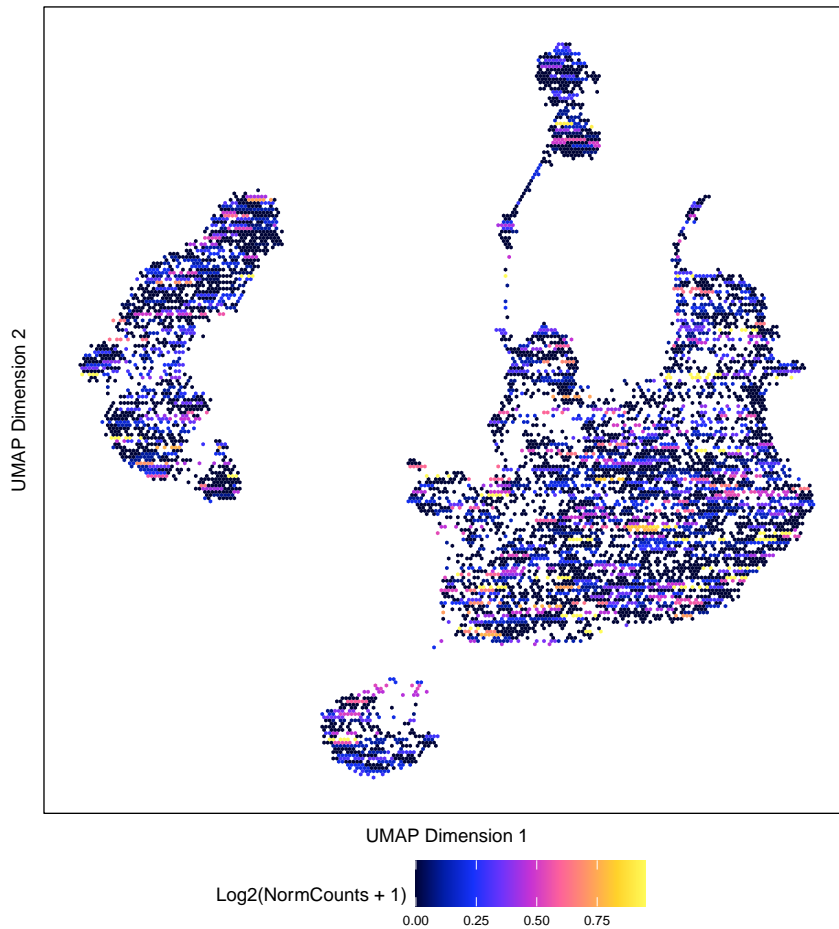
UMAP of Iterative−LSI−1.0 colored by
GeneScoreMatrix : CD3D

UMAP Dimension 2

UMAP Dimension 1

Log2(NormCounts + 1)

0.0   0.4   0.8   1.2   1.6

UMAP of Iterative−LSI−1.0 colored by
GeneScoreMatrix : CD8A

UMAP Dimension 2

UMAP Dimension 1

Log2(NormCounts + 1)

0.0   0.4   0.8   1.2   1.6

UMAP of Iterative−LSI−1.0 colored by
GeneScoreMatrix : CD4

UMAP Dimension 2

UMAP Dimension 1
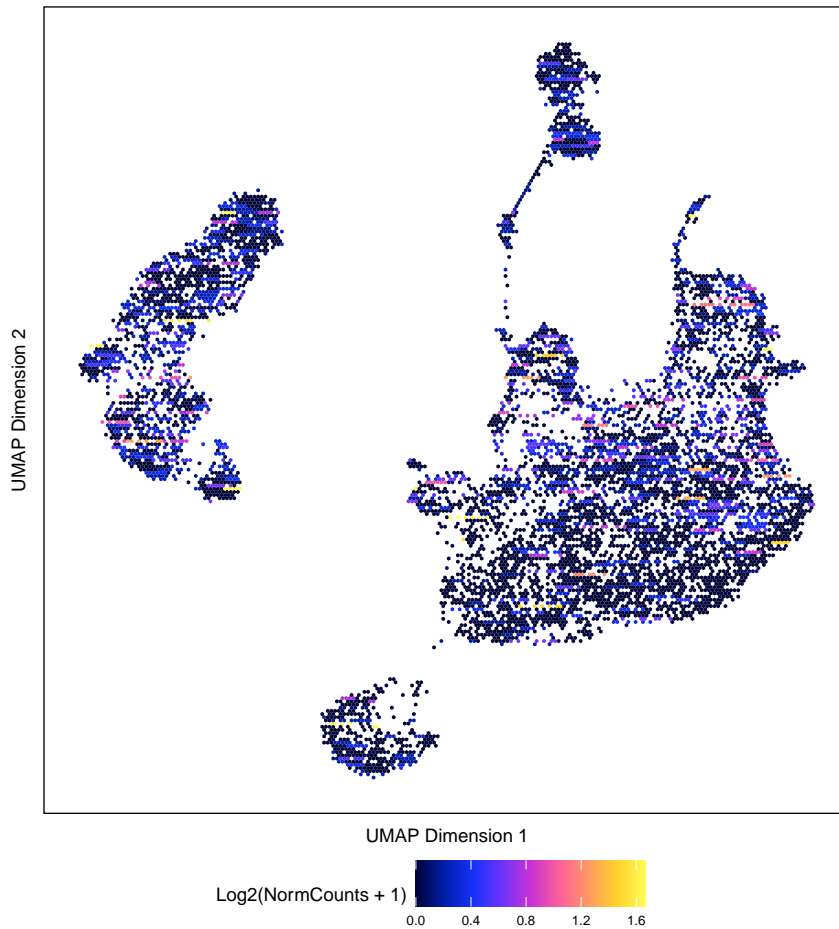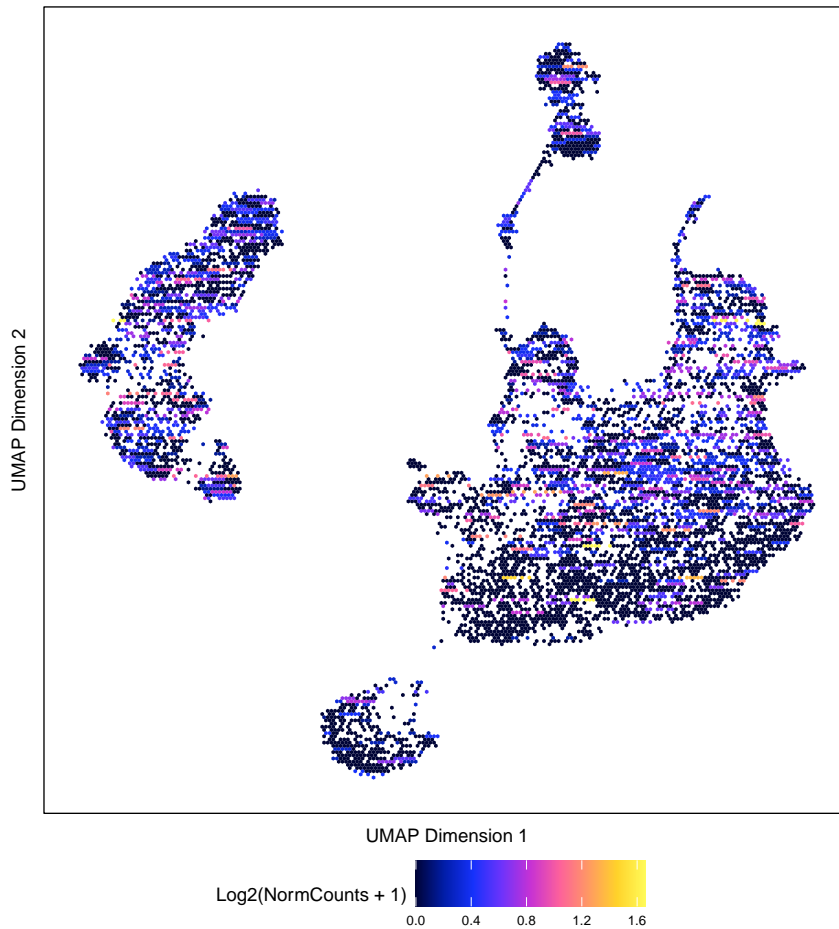
Log2(NormCounts + 1)

0.00   0.25   0.50   0.75   1.00   1.25
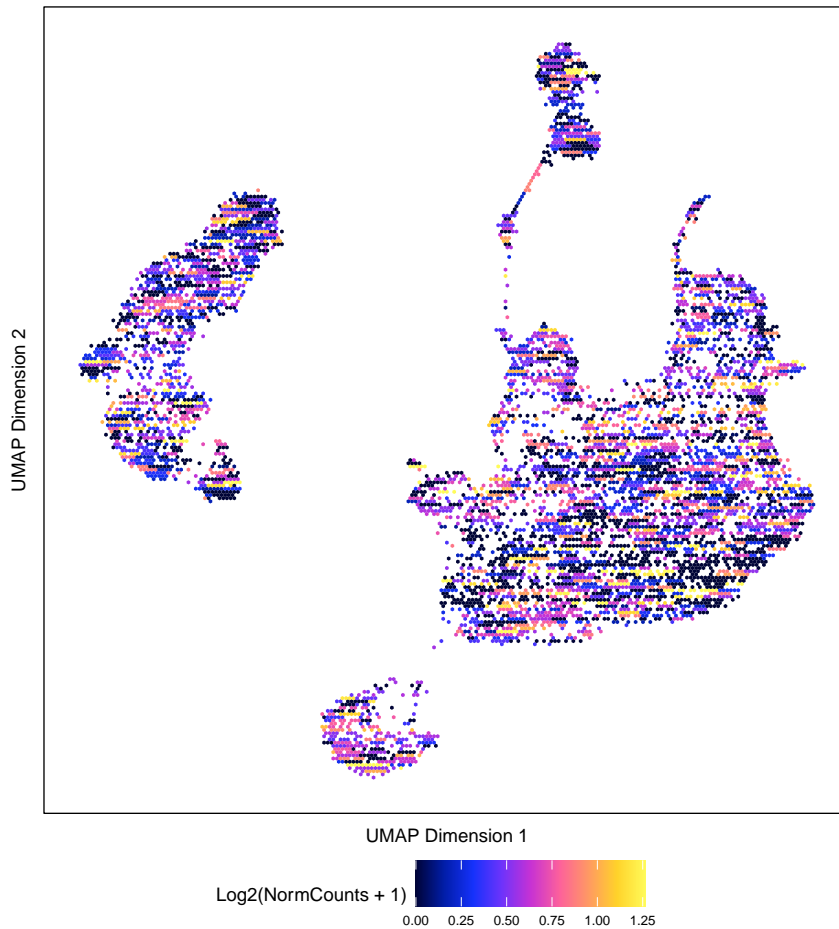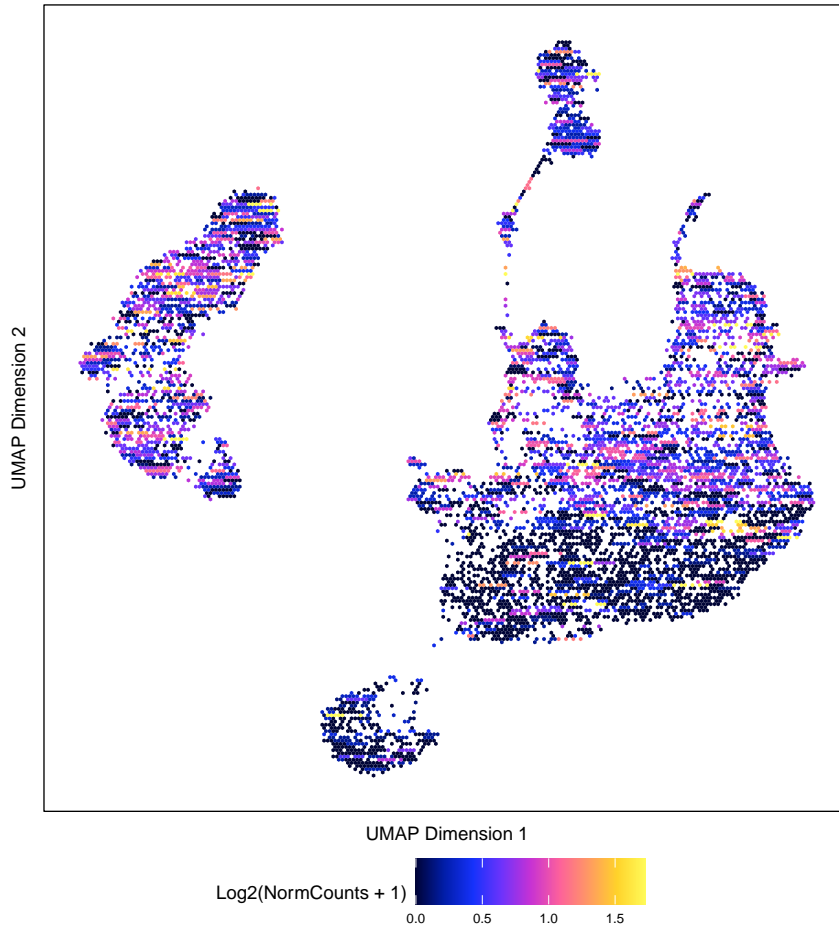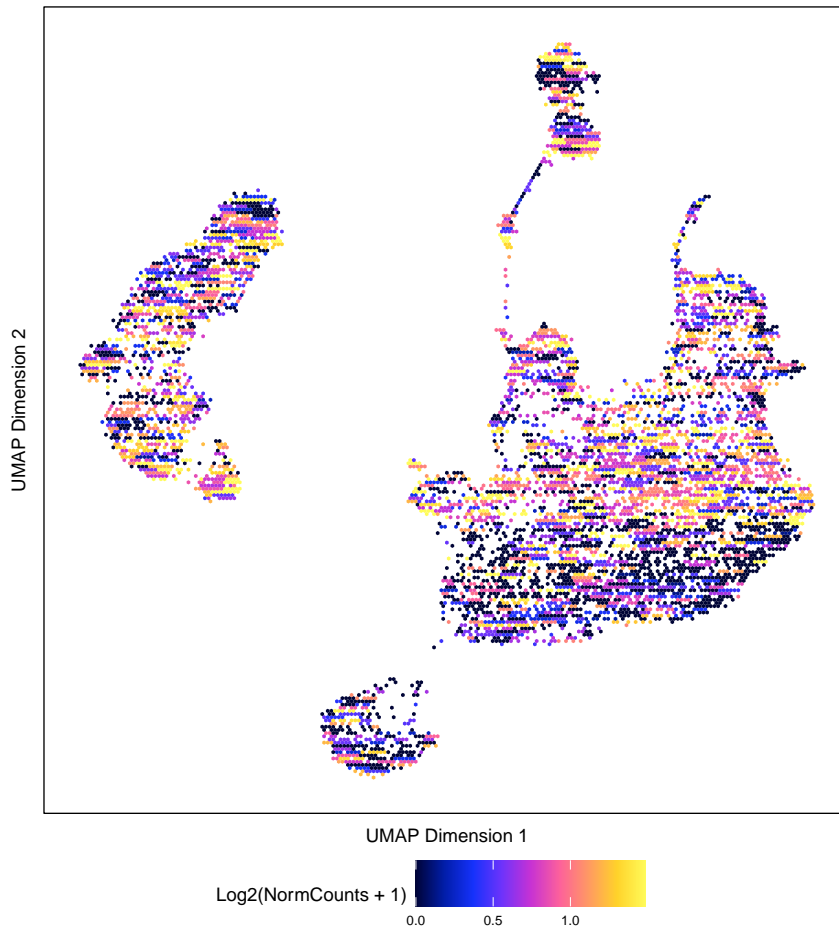
UMAP of Iterative−LSI−1.0 colored by
GeneScoreMatrix : TBX21

UMAP of Iterative−LSI−1.0 colored by
GeneScoreMatrix : CD14

UMAP Dimension 2

UMAP Dimension 1

Log2(NormCounts + 1)

0.0   0.5   1.0   1.5   2.0
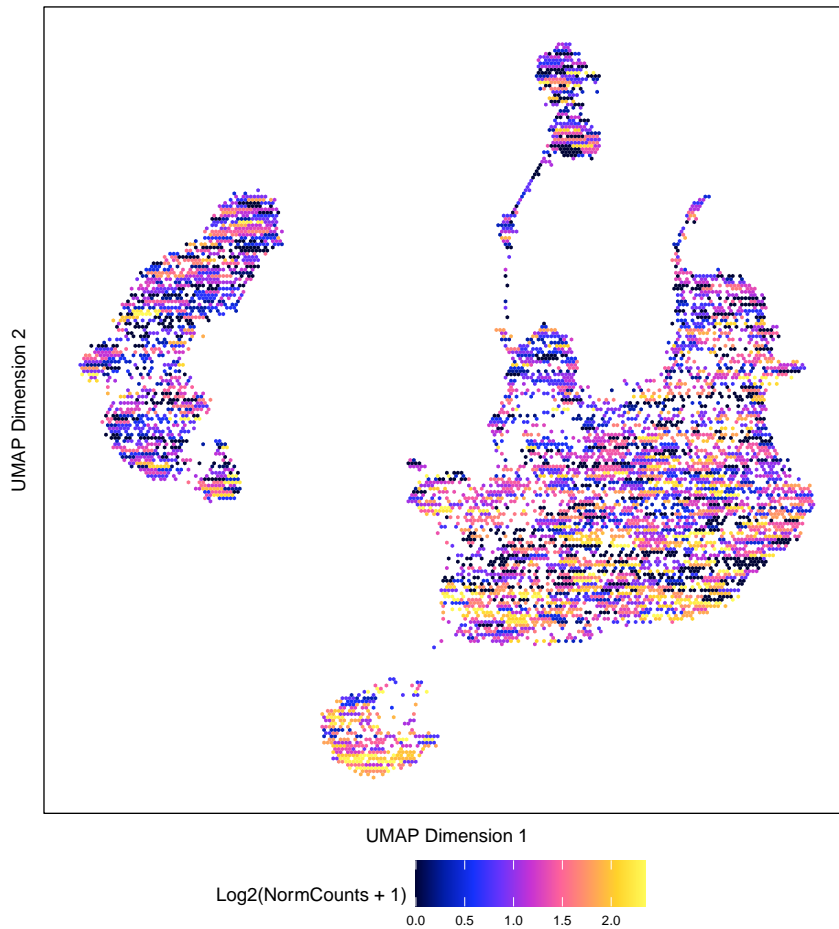
UMAP of Iterative−LSI−1.0 colored by
GeneScoreMatrix : MPO

UMAP Dimension 2

UMAP Dimension 1

Log2(NormCounts + 1)

0.0   0.5   1.0

UMAP of Iterative−LSI−1.0 colored by
GeneScoreMatrix : CEBPB

UMAP Dimension 2

UMAP Dimension 1

Log2(NormCounts + 1)

0.0   0.5   1.0   1.5   2.0

UMAP of Iterative−LSI−1.0 colored by
GeneScoreMatrix : TBX21
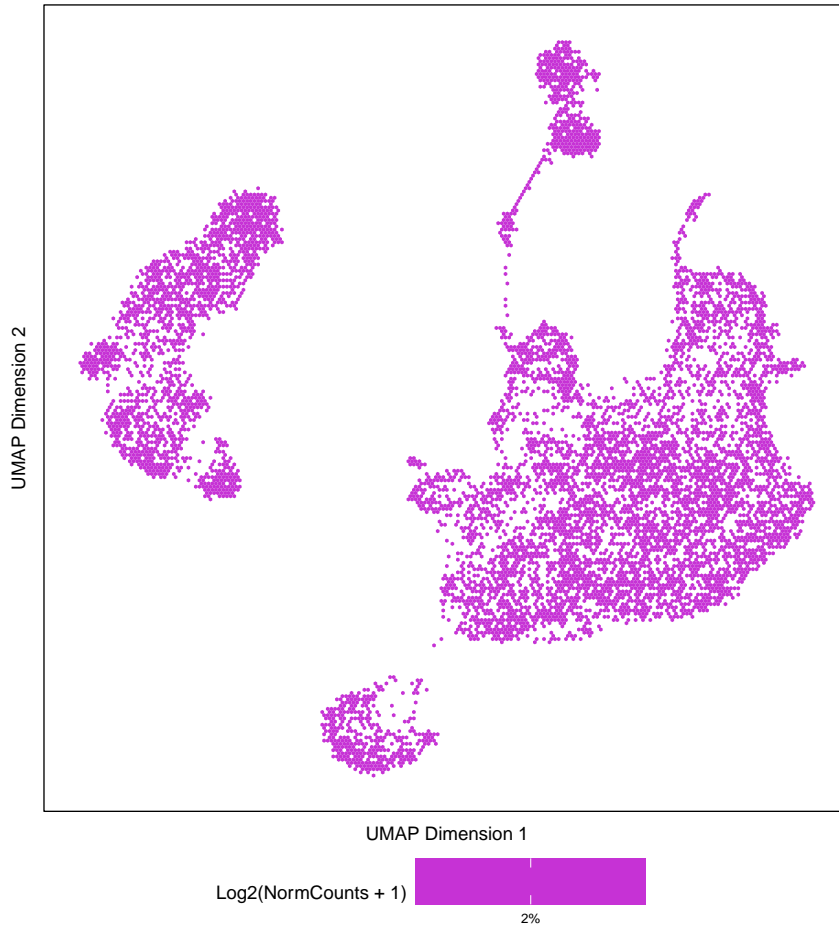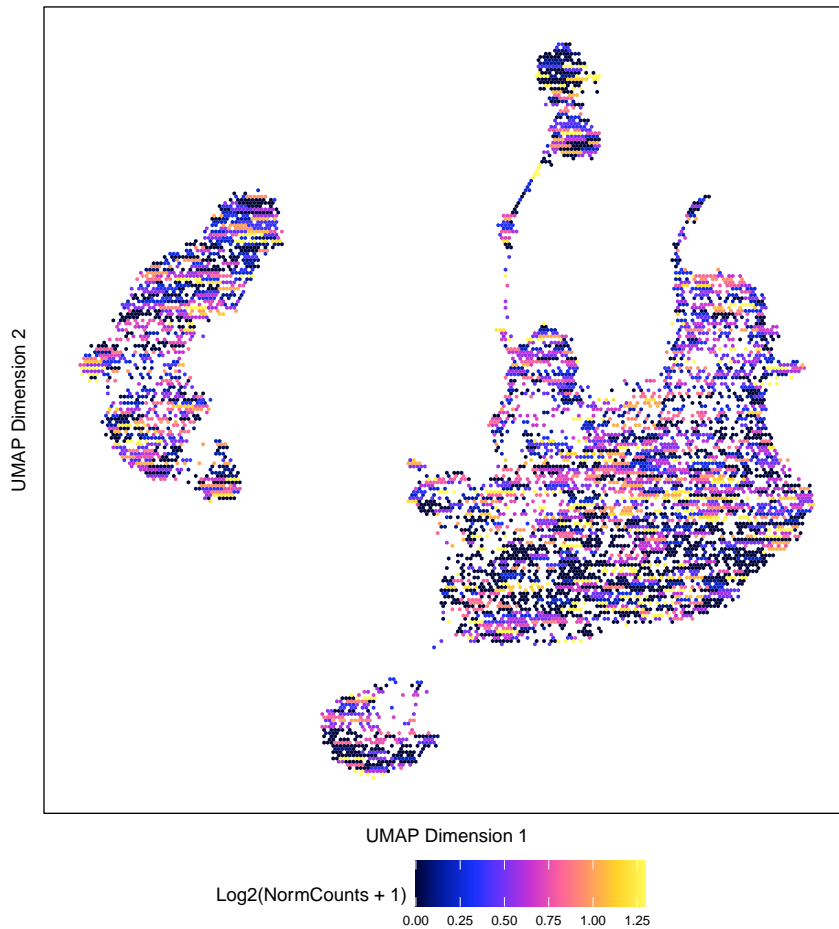
UMAP Dimension 2

UMAP Dimension 1

Log2(NormCounts + 1)

2%
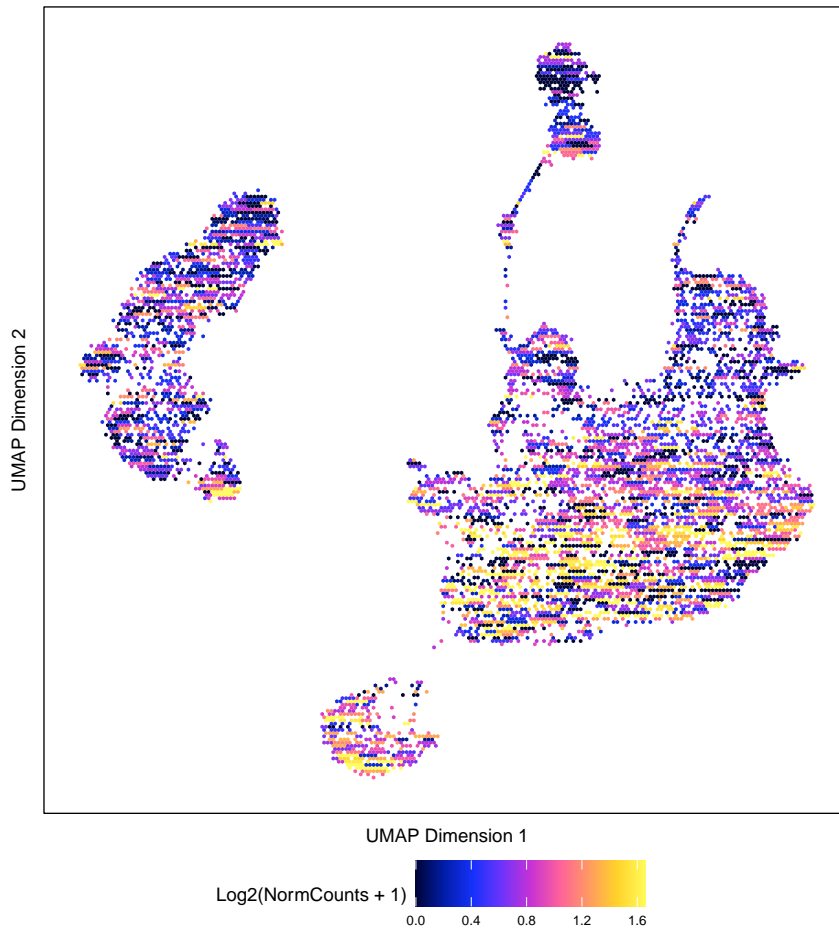
UMAP of Iterative−LSI−1.0 colored by
GeneScoreMatrix : CD38

UMAP of Iterative−LSI−1.0 colored by
GeneScoreMatrix : CD52

UMAP Dimension 2

UMAP Dimension 1

Log2(NormCounts + 1)

0.0   0.5   1.0   1.5
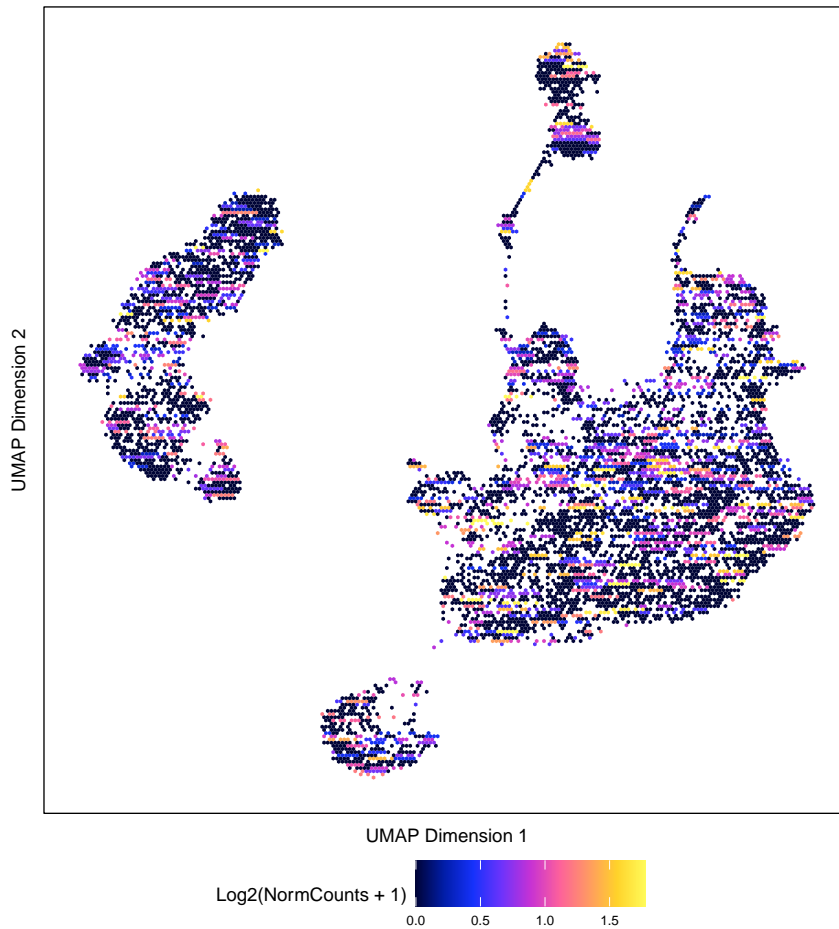
UMAP of Iterative−LSI−1.0 colored by
GeneScoreMatrix : CSF3R

UMAP Dimension 2

UMAP Dimension 1

Log2(NormCounts + 1)

0.00 0.25 0.50 0.75 1.00 1.25

UMAP of Iterative−LSI−1.0 colored by
GeneScoreMatrix : ANXA1

UMAP Dimension 2

UMAP Dimension 1

Log2(NormCounts + 1)

0.0   0.4   0.8   1.2   1.6

UMAP of Iterative−LSI−1.0 colored by
GeneScoreMatrix : APLP2

UMAP Dimension 2

UMAP Dimension 1

Log2(NormCounts + 1)

0.0   0.4   0.8   1.2   1.6

UMAP of Iterative−LSI−1.0 colored by
GeneScoreMatrix : CD19

UMAP Dimension 2

UMAP Dimension 1

Log2(NormCounts + 1)

2%

UMAP of Iterative−LSI−1.0 colored by
GeneScoreMatrix : CD34

UMAP Dimension 2

UMAP Dimension 1

Log2(NormCounts + 1)

2%