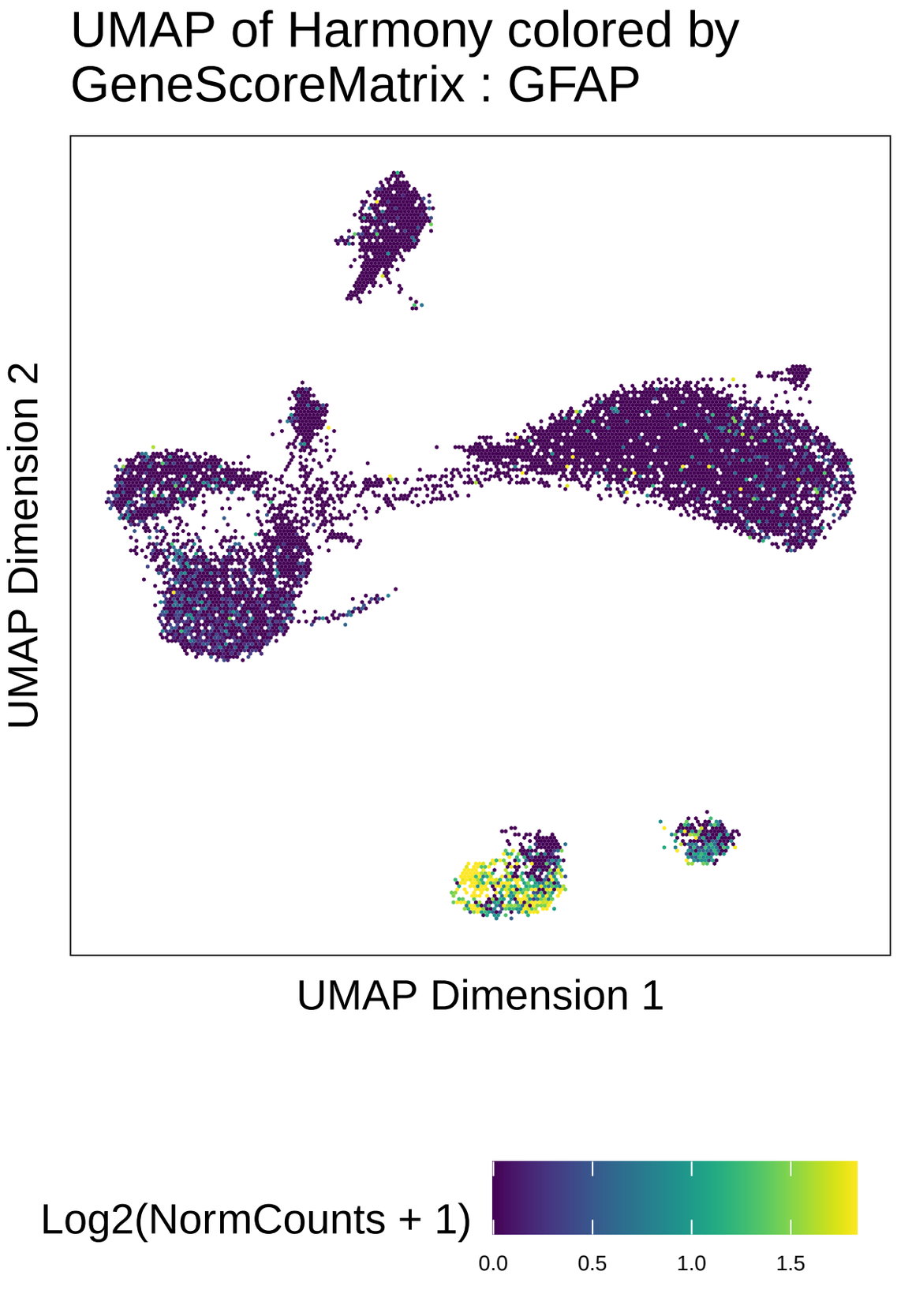
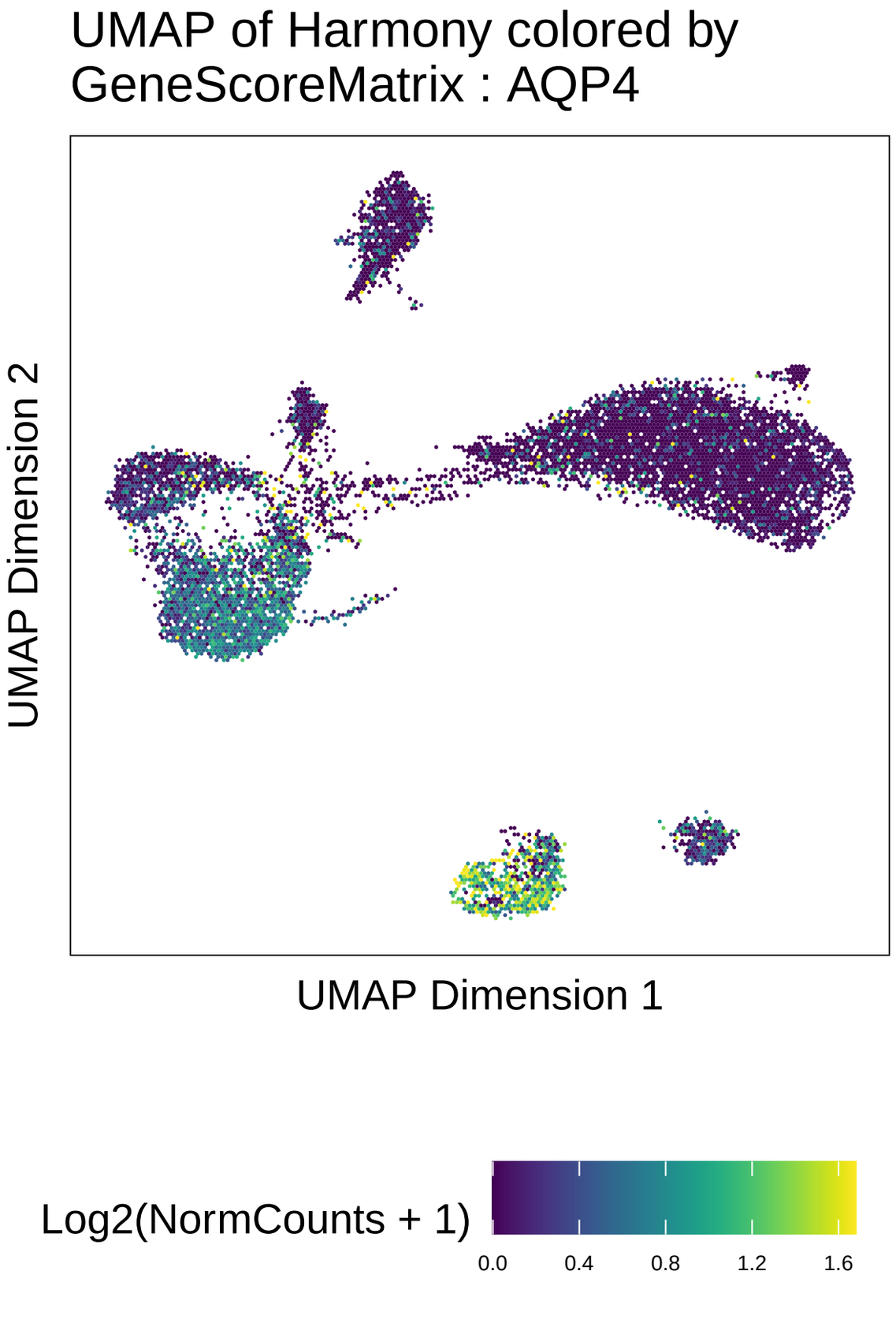
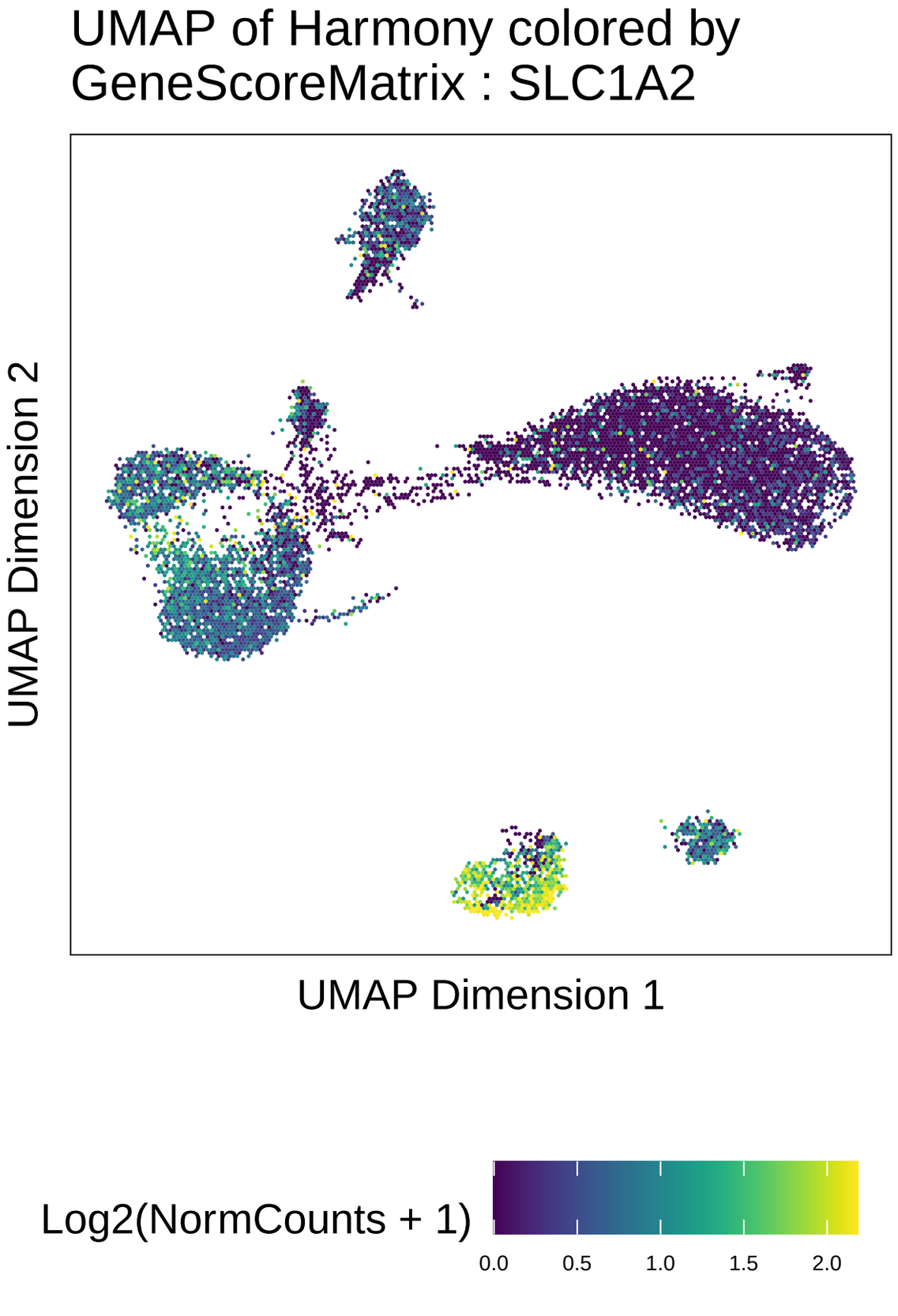
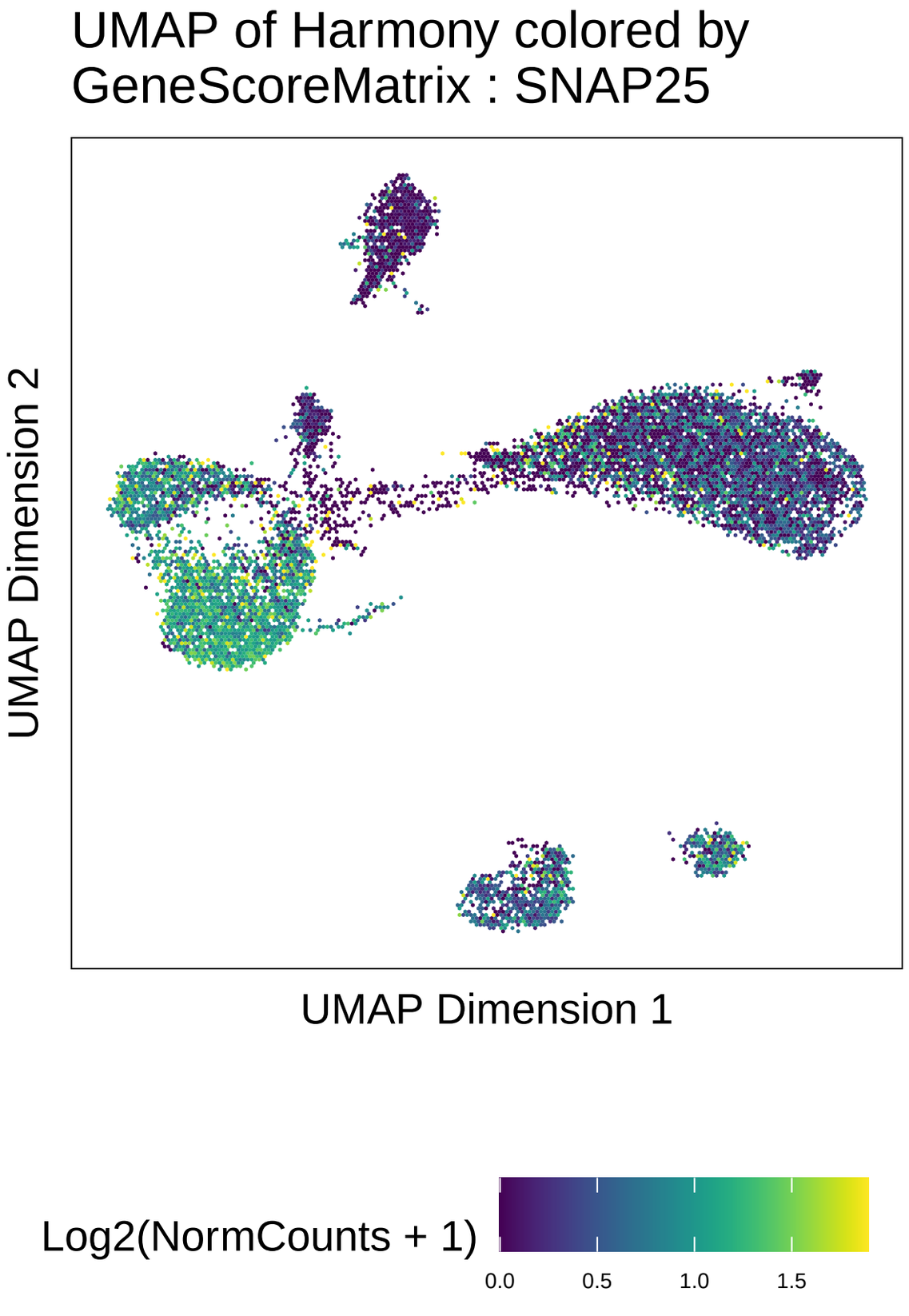
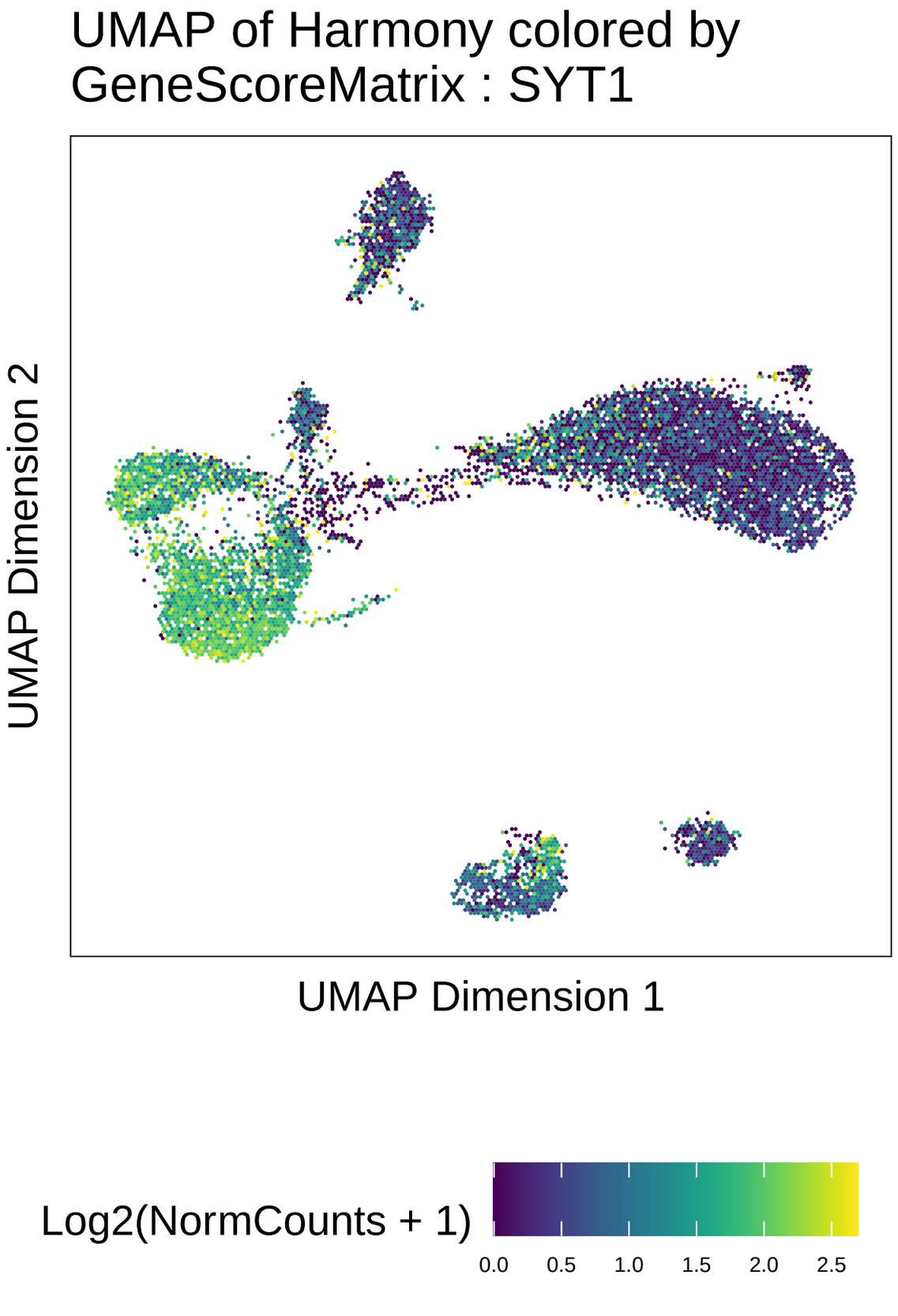
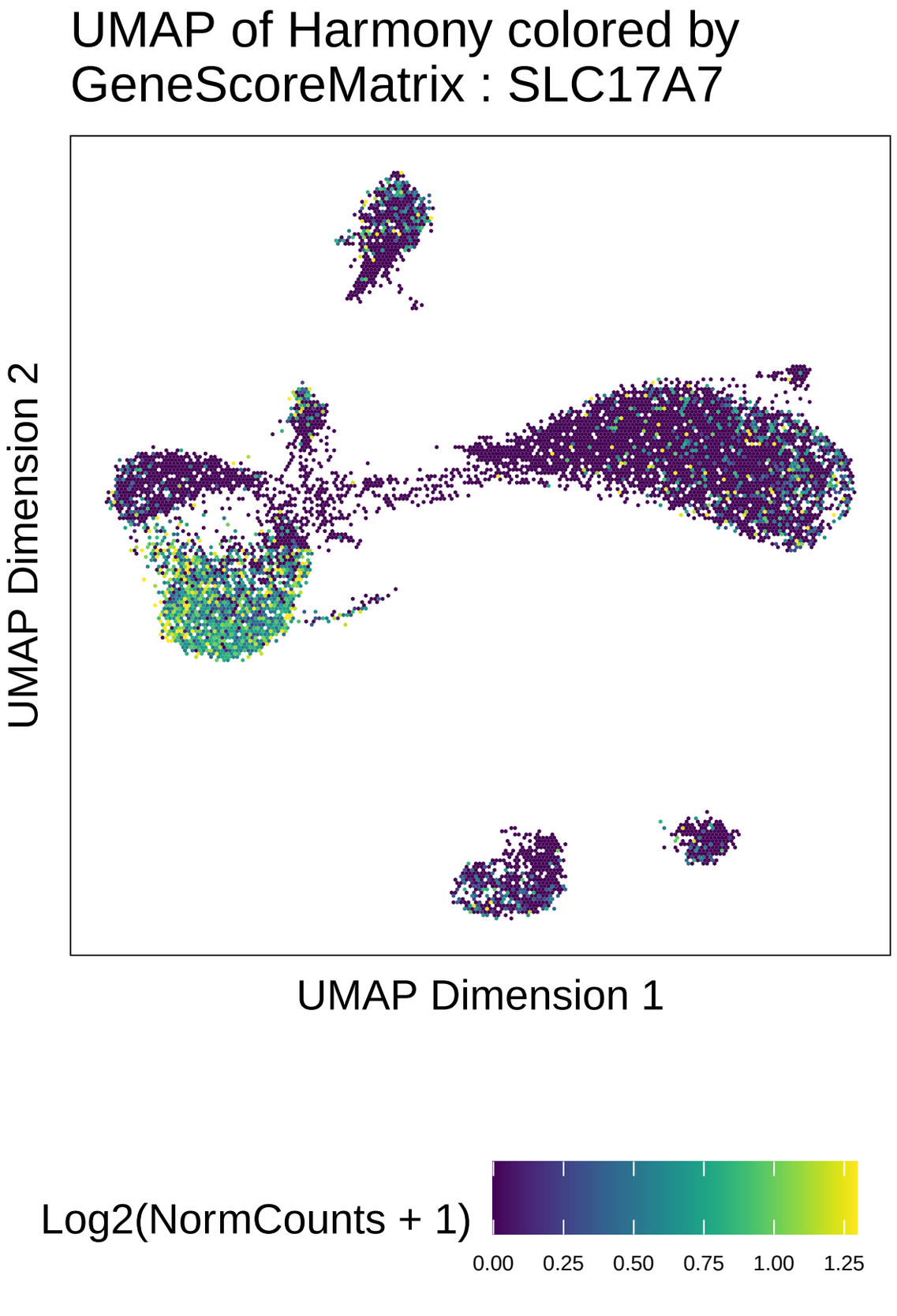

UMAP of Harmony colored by
GeneScoreMatrix : GFAP
UMAP Dimension 2
UMAP Dimension 1
Log2(NormCounts + 1)
0.0
0.5
1.0
1.5

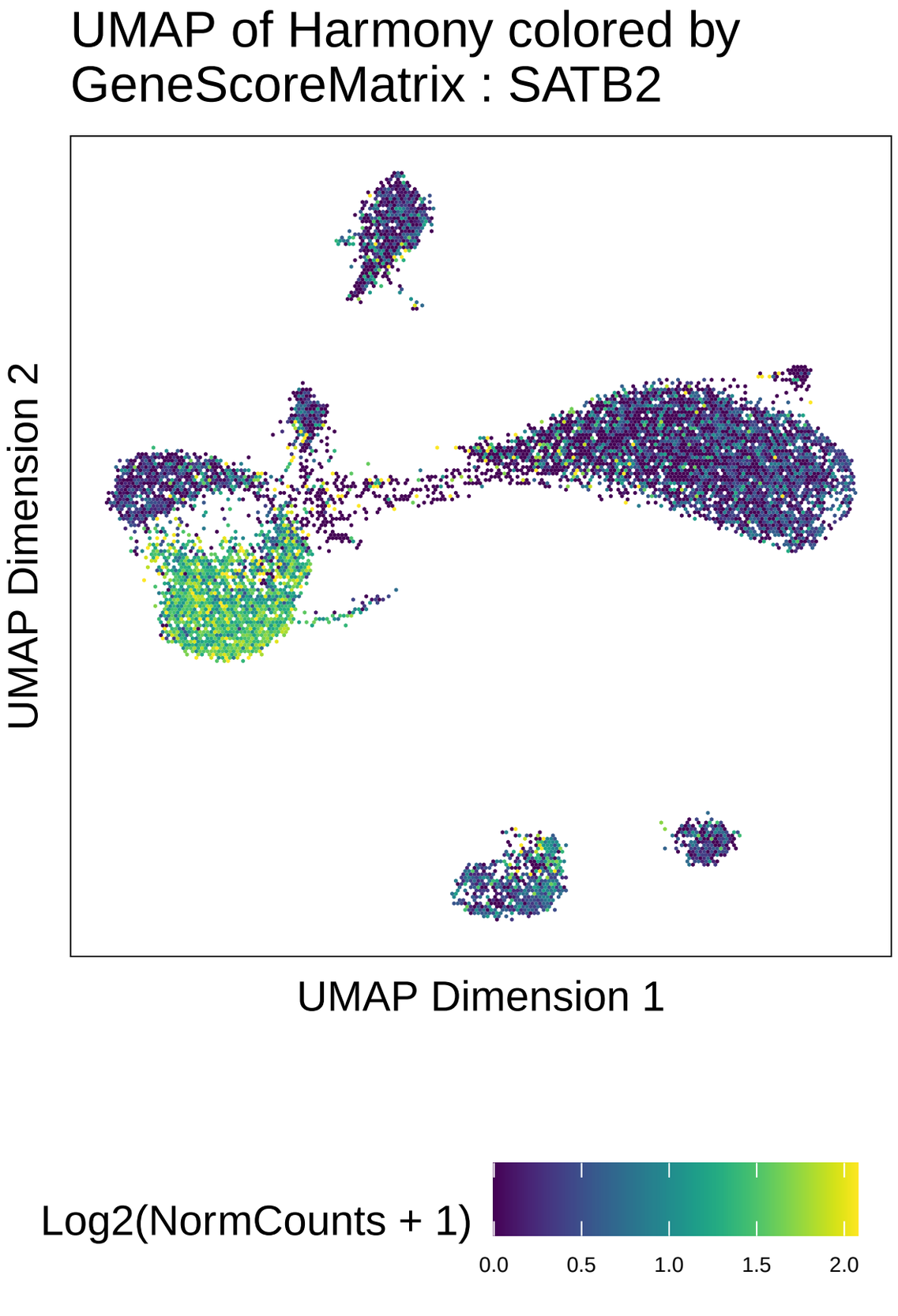

UMAP of Harmony colored by
GeneScoreMatrix : AQP4
UMAP Dimension 2
UMAP Dimension 1
Log2(NormCounts + 1)
0.0
0.4
0.8
1.2
1.6

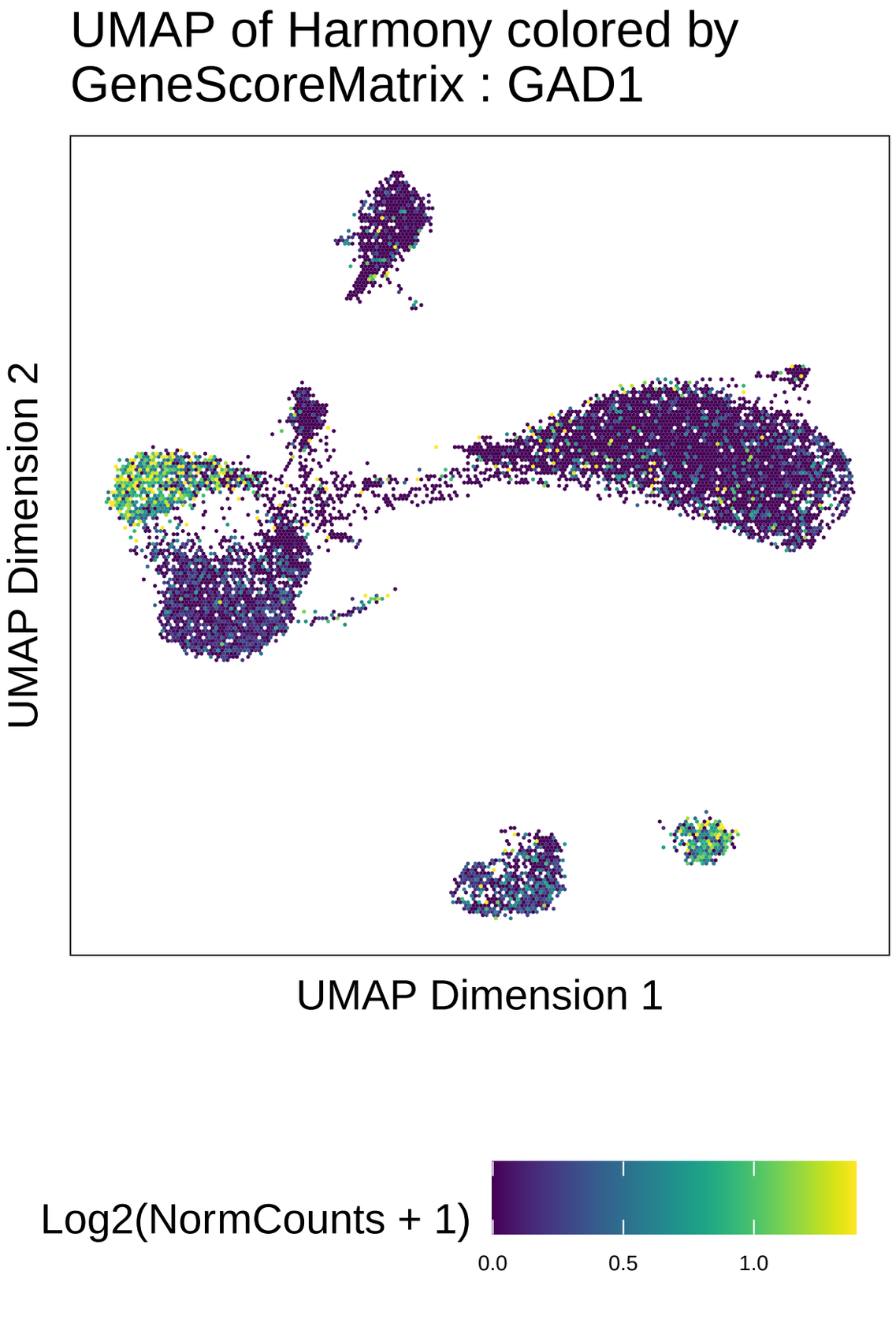

UMAP of Harmony colored by
GeneScoreMatrix : SLC1A2
UMAP Dimension 2
UMAP Dimension 1
Log2(NormCounts + 1)
0.0
0.5
1.0
1.5
2.0

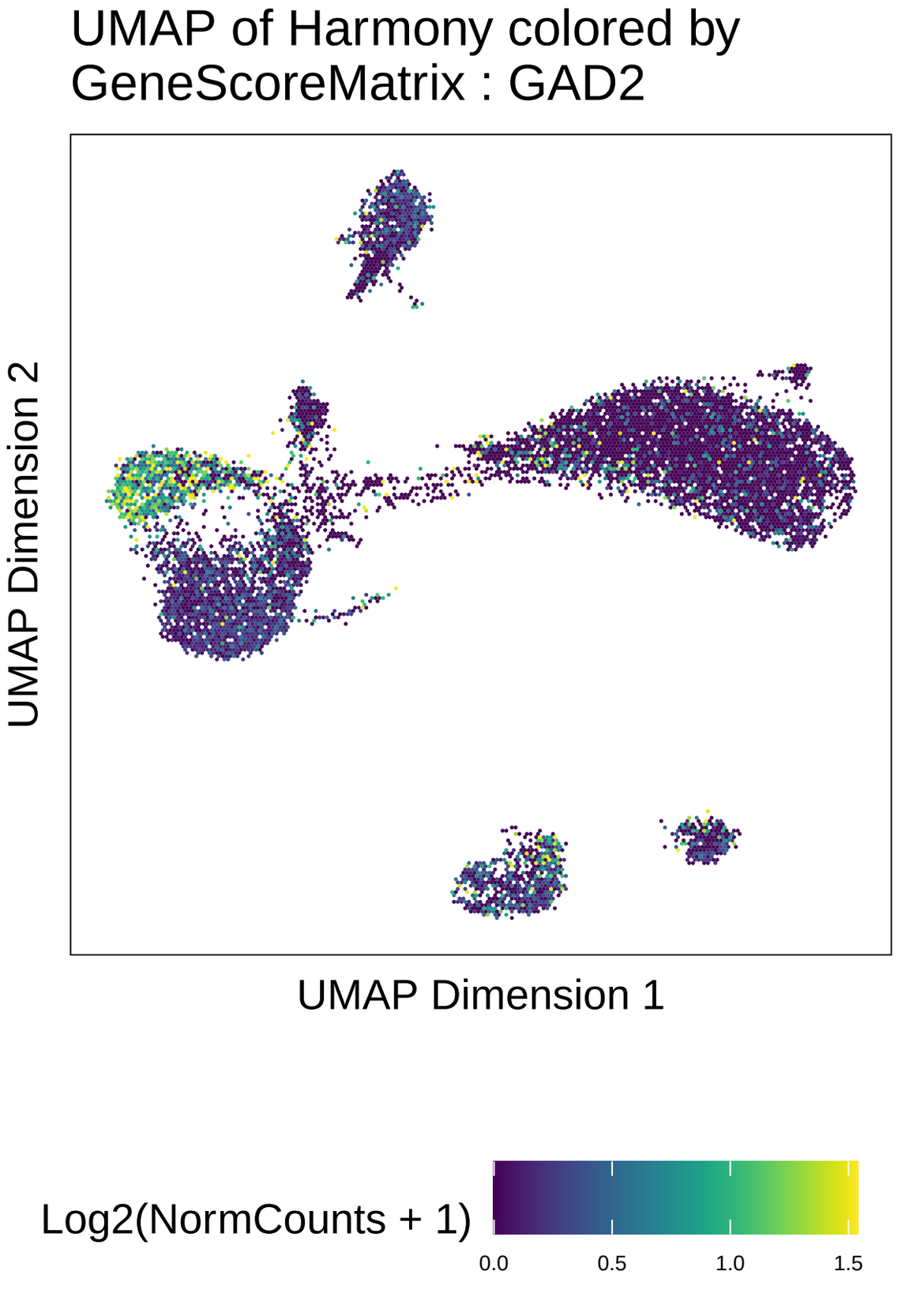

UMAP of Harmony colored by
GeneScoreMatrix : SNAP25
UMAP Dimension 2
UMAP Dimension 1
Log2(NormCounts + 1)
0.0
0.5
1.0
1.5

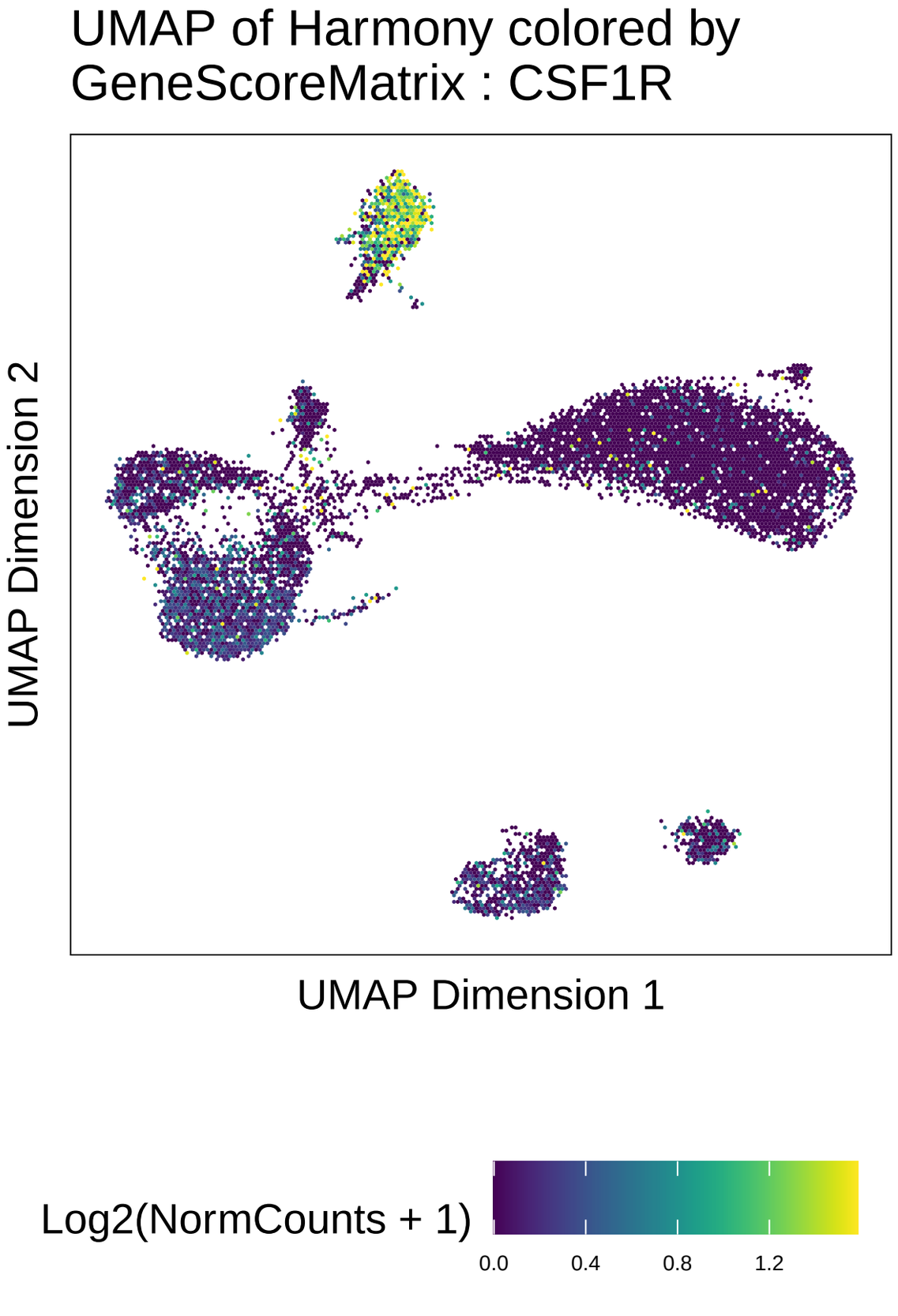

UMAP of Harmony colored by
GeneScoreMatrix : SYT1
UMAP Dimension 2
UMAP Dimension 1
Log2(NormCounts + 1)
0.0
0.5
1.0
1.5
2.0
2.5

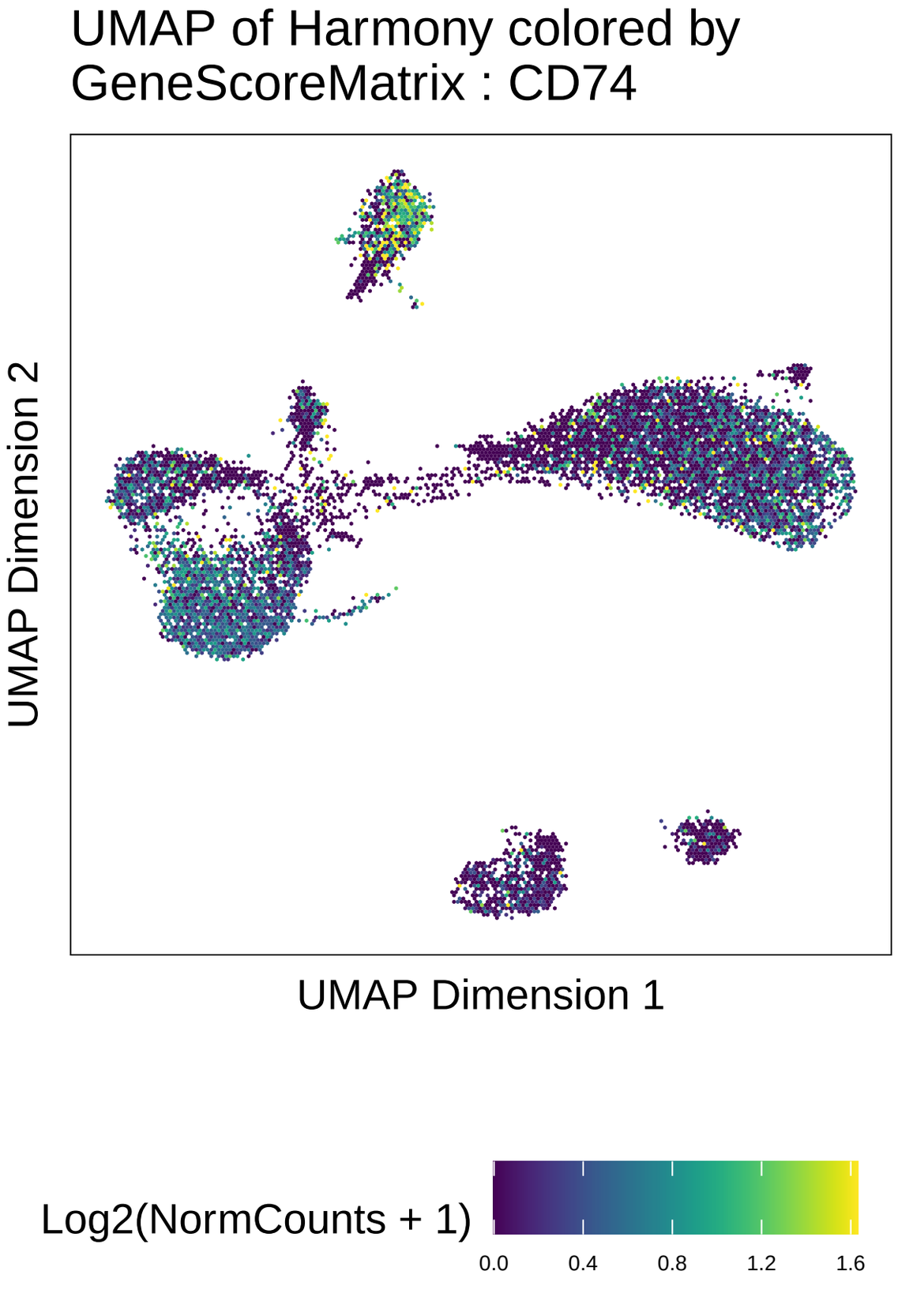

UMAP of Harmony colored by
GeneScoreMatrix : SLC17A7
UMAP Dimension 2
UMAP Dimension 1
Log2(NormCounts + 1)
0.00
0.25
0.50
0.75
1.00
1.25

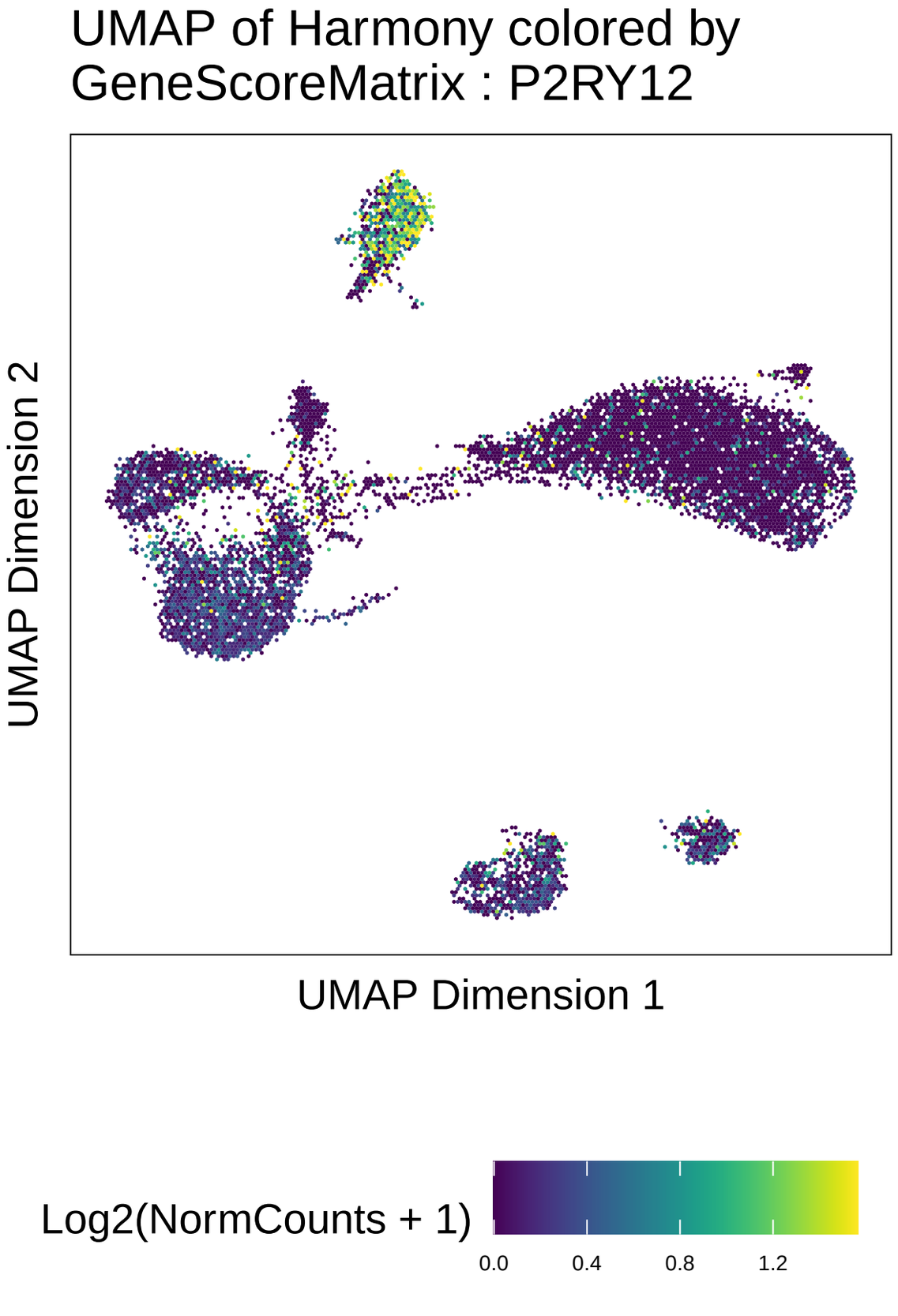

UMAP of Harmony colored by
GeneScoreMatrix : SATB2
UMAP Dimension 2
UMAP Dimension 1
Log2(NormCounts + 1)
0.0
0.5
1.0
1.5
2.0

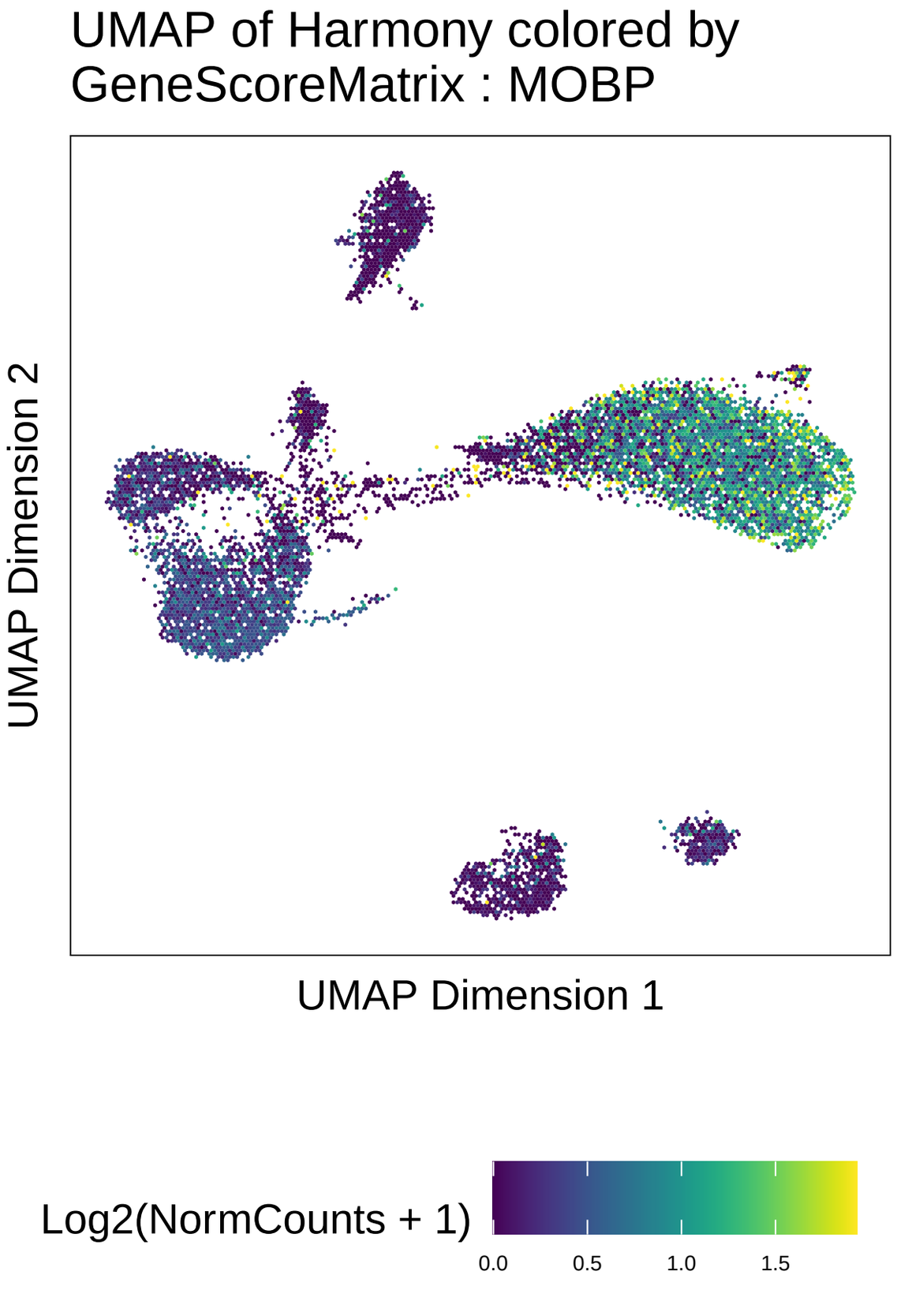

UMAP of Harmony colored by
GeneScoreMatrix : GAD1
UMAP Dimension 2
UMAP Dimension 1
Log2(NormCounts + 1)
0.0
0.5
1.0

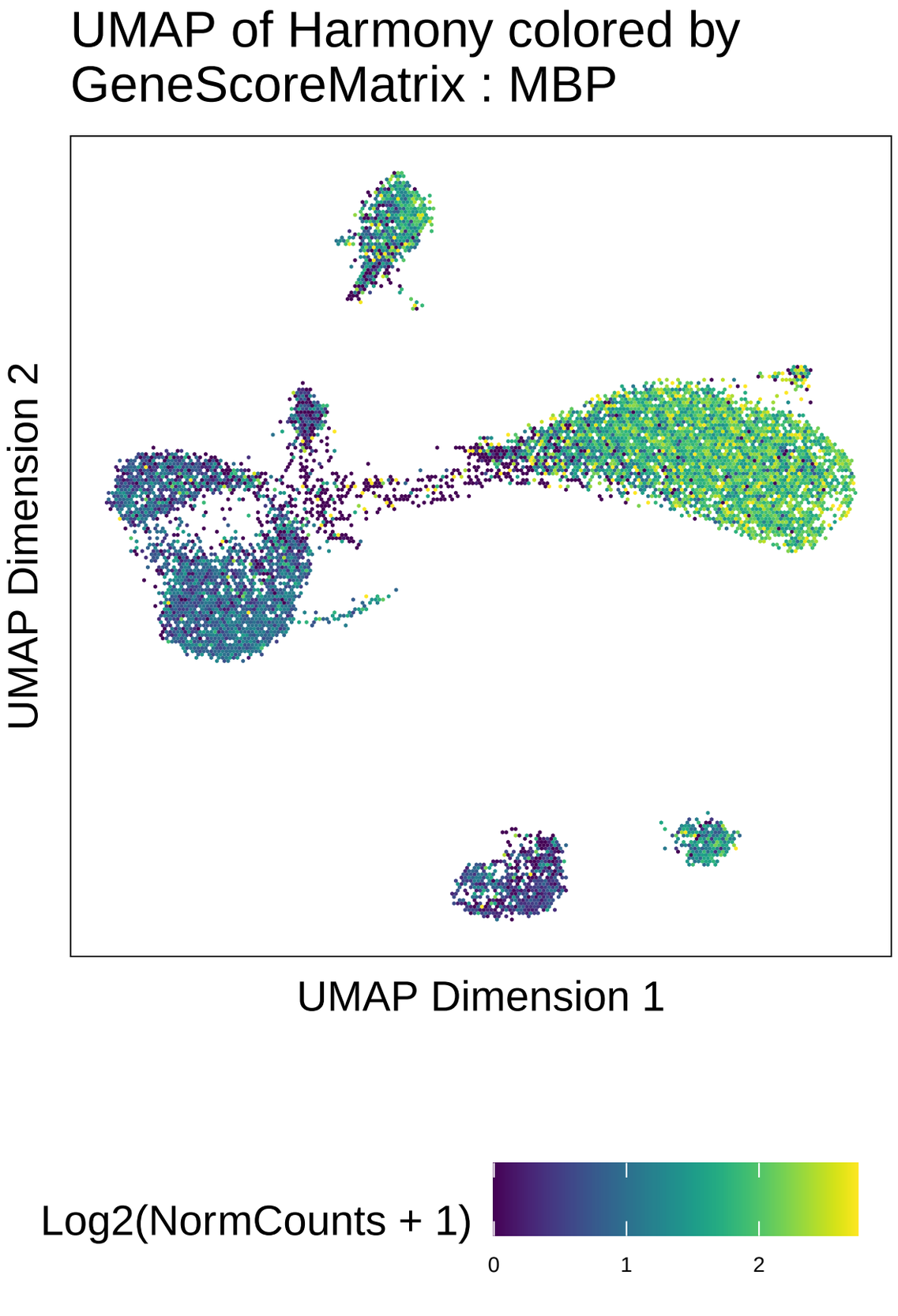

UMAP of Harmony colored by
GeneScoreMatrix : GAD2
UMAP Dimension 2
UMAP Dimension 1
Log2(NormCounts + 1)
0.0
0.5
1.0
1.5

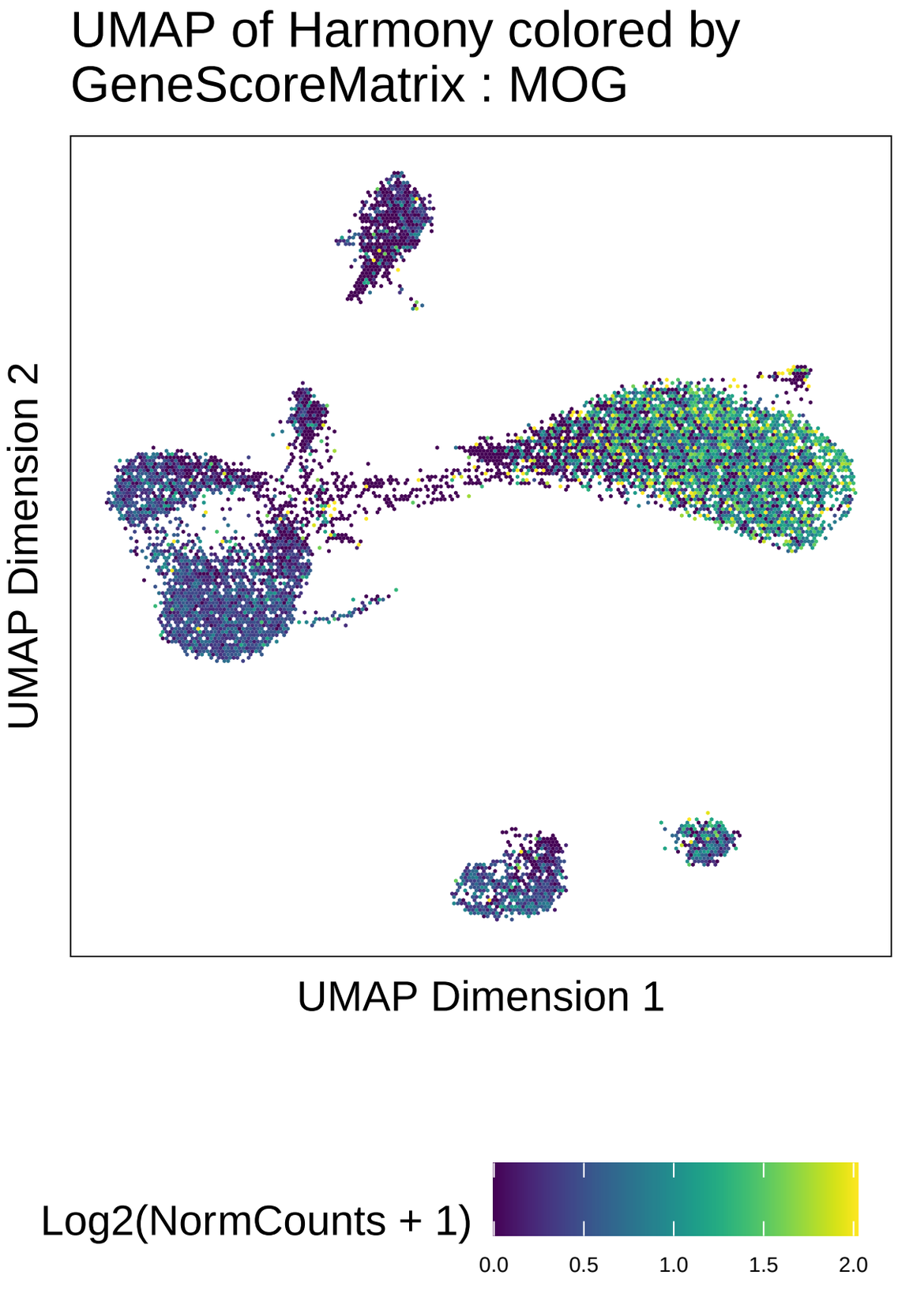

UMAP of Harmony colored by
GeneScoreMatrix : CSF1R
UMAP Dimension 2
UMAP Dimension 1
Log2(NormCounts + 1)
0.0
0.4
0.8
1.2

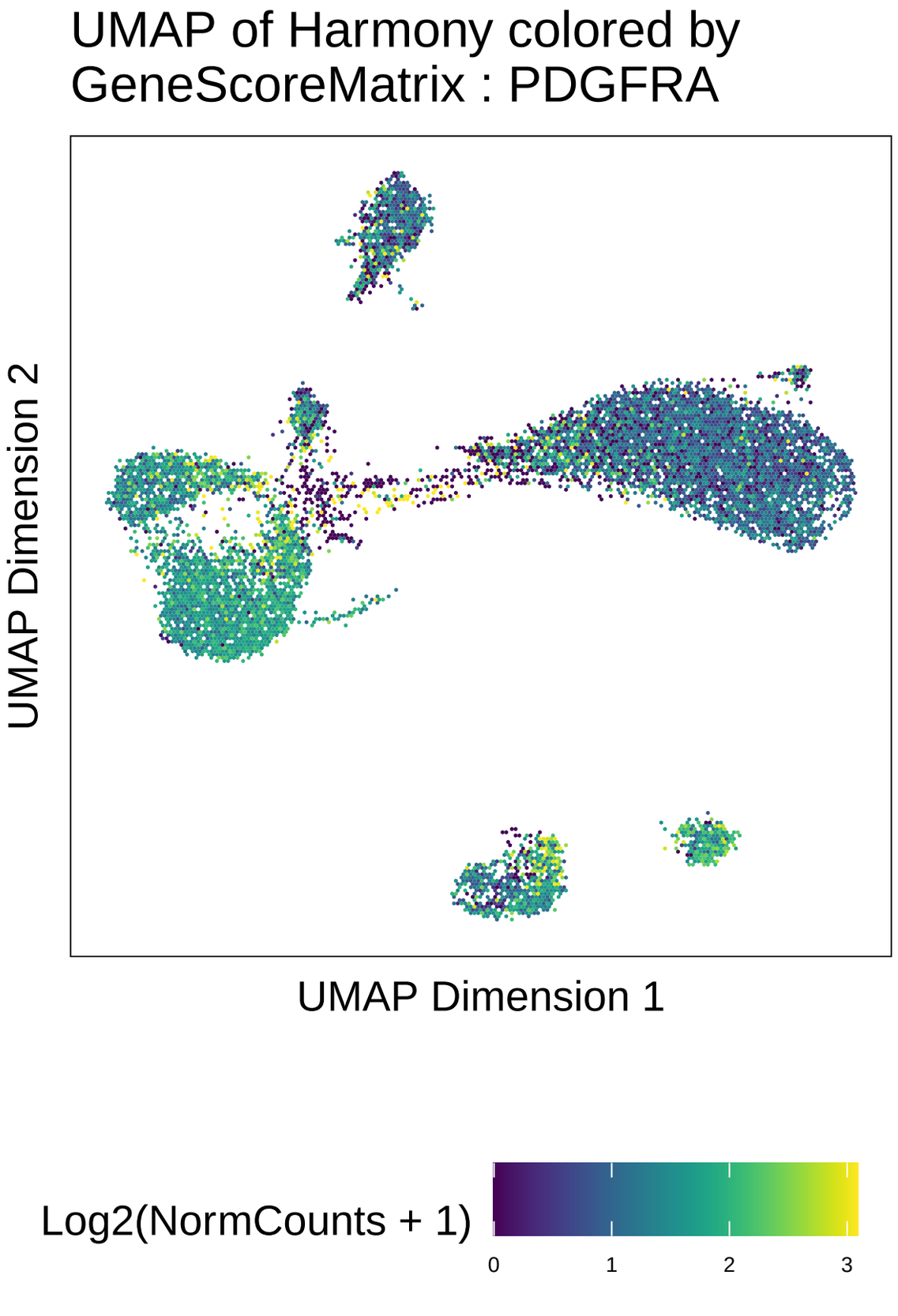

UMAP of Harmony colored by
GeneScoreMatrix : CD74
UMAP Dimension 2
UMAP Dimension 1
Log2(NormCounts + 1)
0.0
0.4
0.8
1.2
1.6

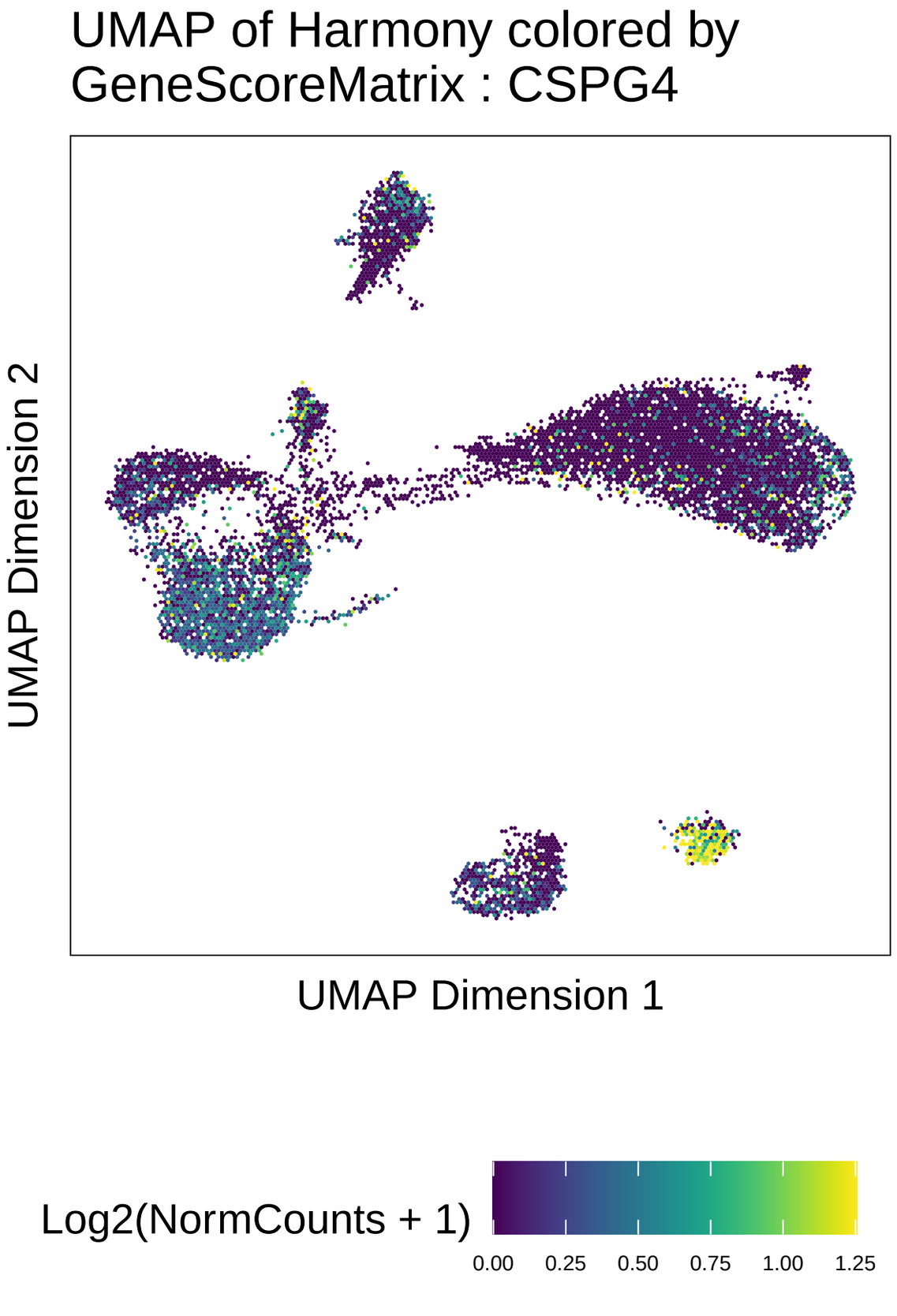

UMAP of Harmony colored by
GeneScoreMatrix : P2RY12
UMAP Dimension 2
UMAP Dimension 1
Log2(NormCounts + 1)
0.0
0.4
0.8
1.2


UMAP of Harmony colored by
GeneScoreMatrix : MOBP
UMAP Dimension 2
UMAP Dimension 1
Log2(NormCounts + 1)
0.0
0.5
1.0
1.5


UMAP of Harmony colored by
GeneScoreMatrix : MBP
UMAP Dimension 2
UMAP Dimension 1
Log2(NormCounts + 1)
0
1
2


UMAP of Harmony colored by
GeneScoreMatrix : MOG
UMAP Dimension 2
UMAP Dimension 1
Log2(NormCounts + 1)
0.0
0.5
1.0
1.5
2.0


UMAP of Harmony colored by
GeneScoreMatrix : PDGFRA
UMAP Dimension 2
UMAP Dimension 1
Log2(NormCounts + 1)
0
1
2
3


UMAP of Harmony colored by
GeneScoreMatrix : CSPG4
UMAP Dimension 2
UMAP Dimension 1
Log2(NormCounts + 1)
0.00
0.25
0.50
0.75
1.00
1.25