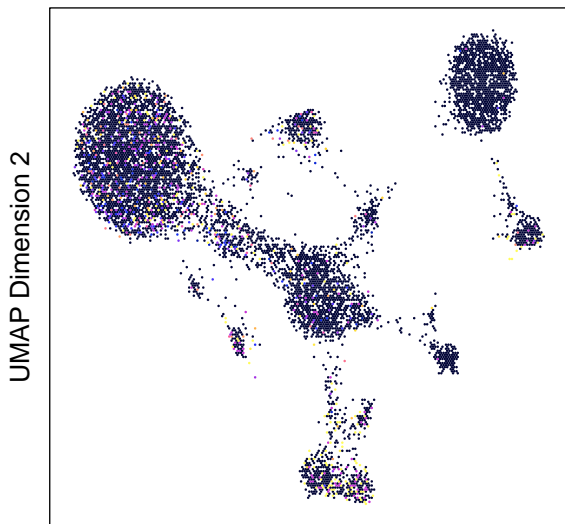
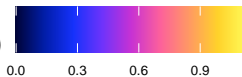


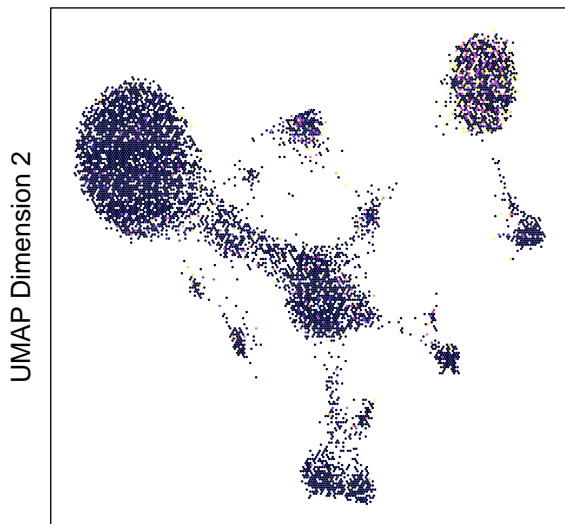
UMAP of IterativeLSI colored by
GeneScoreMatrix : Aqp4



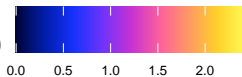
$\text{Log}_2(\text{NormCounts} + 1)$



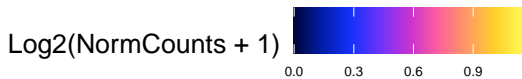
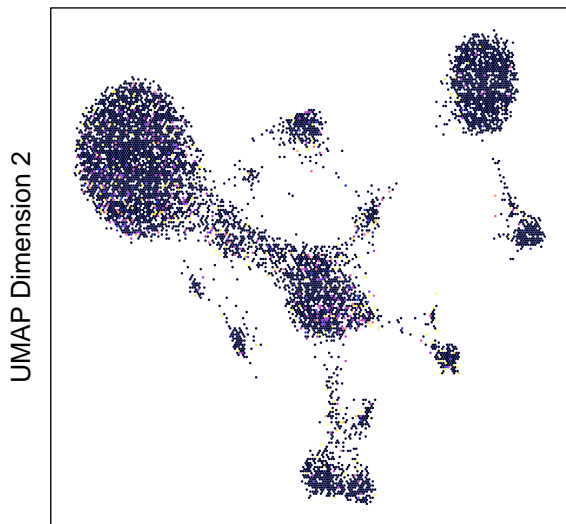
UMAP of IterativeLSI colored by
GeneScoreMatrix : Mog



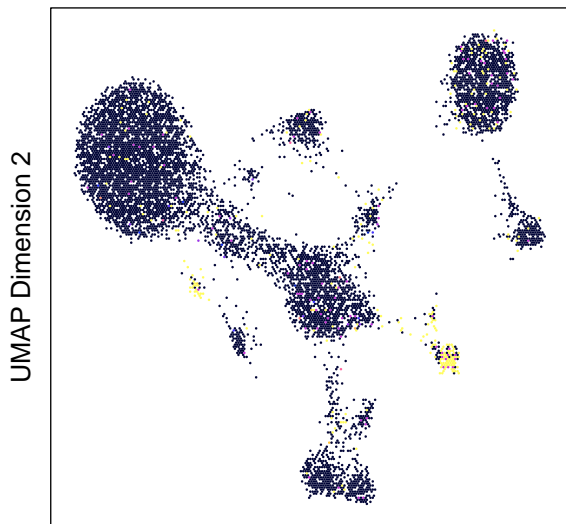
Log2(NormCounts + 1)



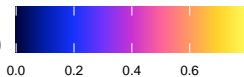
UMAP of IterativeLSI colored by
GeneScoreMatrix : Flt1



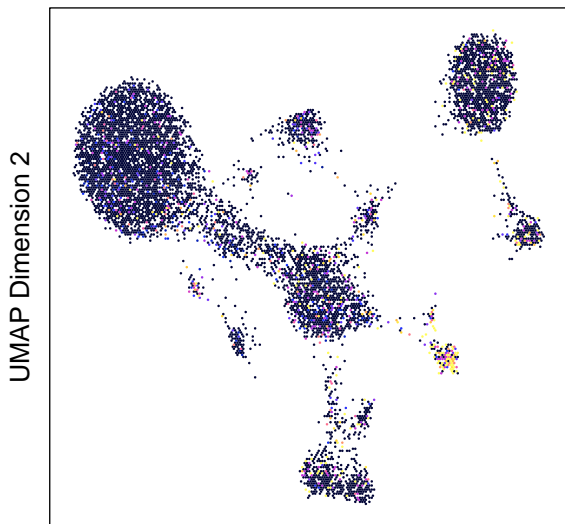
UMAP of IterativeLSI colored by
GeneScoreMatrix : Cx3cr1



Log2(NormCounts + 1)

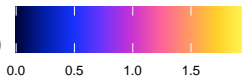


UMAP of IterativeLSI colored by
GeneScoreMatrix : Tgfb β 1

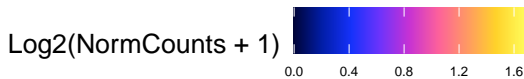
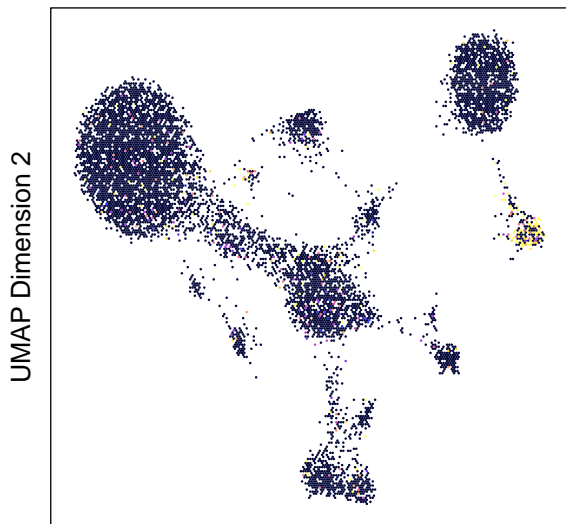


UMAP Dimension 1

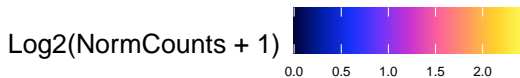
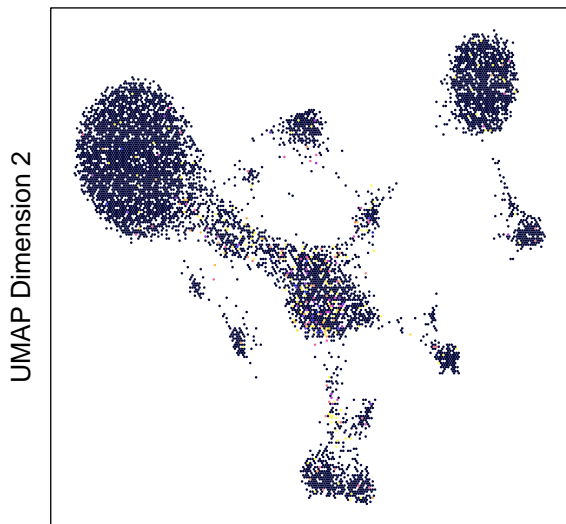
Log2(NormCounts + 1)



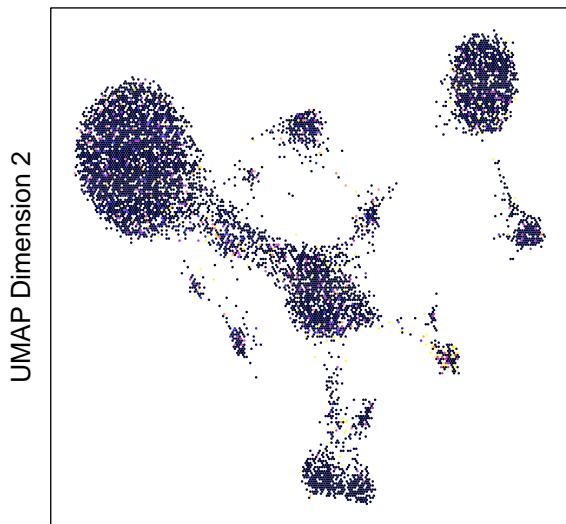
UMAP of IterativeLSI colored by
GeneScoreMatrix : Pdgfra



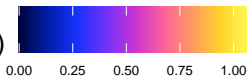
UMAP of IterativeLSI colored by
GeneScoreMatrix : Dlx1



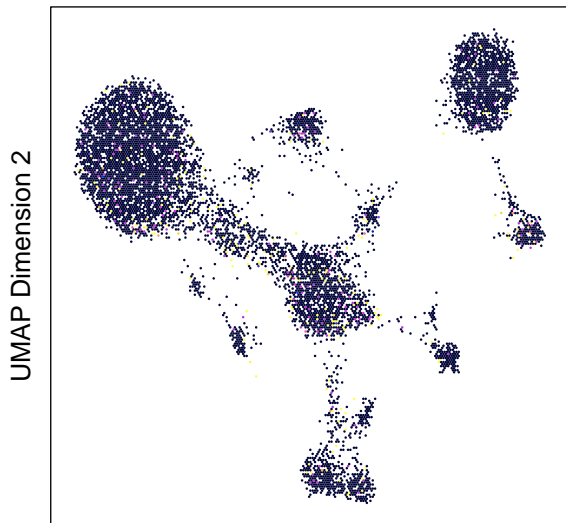
UMAP of IterativeLSI colored by
GeneScoreMatrix : Tmem72



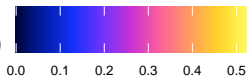
Log2(NormCounts + 1)



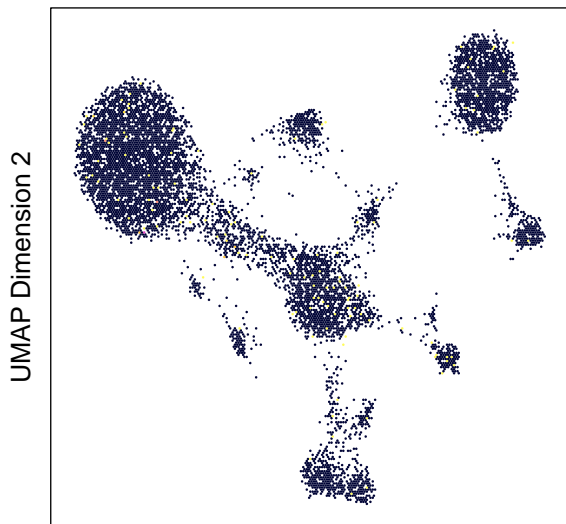
UMAP of IterativeLSI colored by
GeneScoreMatrix : Tmem212



Log2(NormCounts + 1)

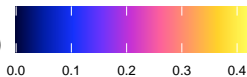


UMAP of IterativeLSI colored by
GeneScoreMatrix : Mrc1

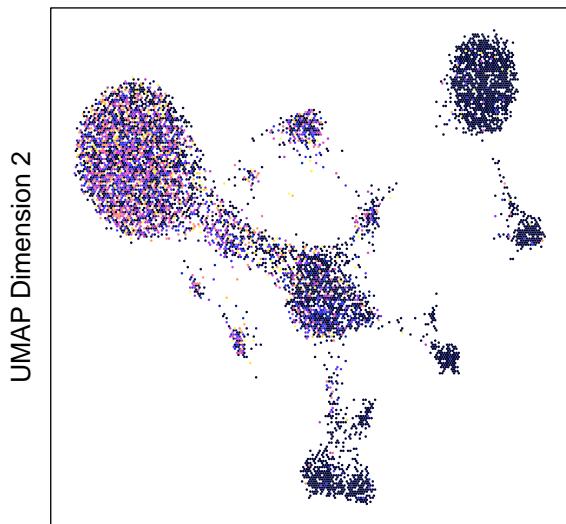


UMAP Dimension 1

$\text{Log}_2(\text{NormCounts} + 1)$



UMAP of IterativeLSI colored by
GeneScoreMatrix : Trank1



Log2(NormCounts + 1)

