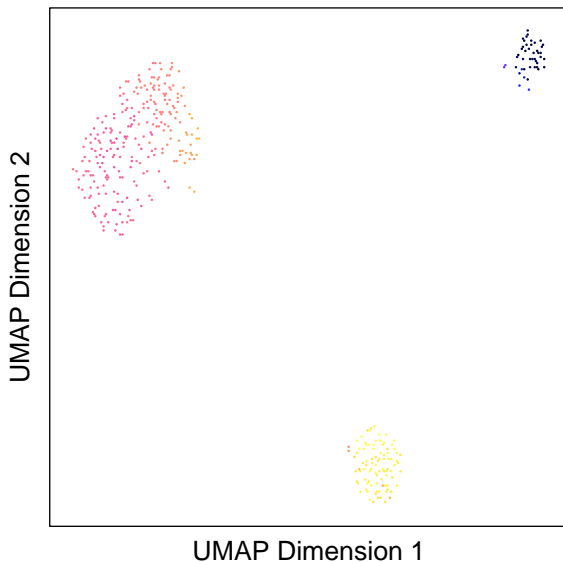
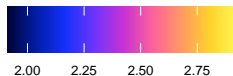


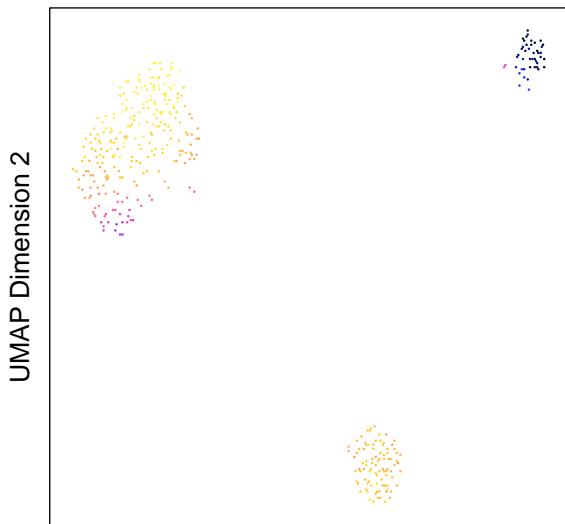
# UMAP\_IterativeLSI of IterativeLSI colored by GeneScoreMatrix : FOS



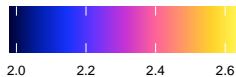
Log2(NormCounts + 1)



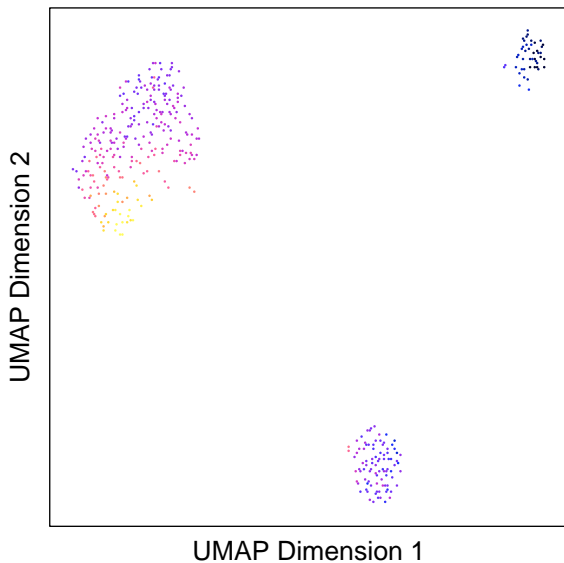
# UMAP\_IterativeLSI of IterativeLSI colored by GeneScoreMatrix : FOSB



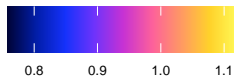
Log2(NormCounts + 1)



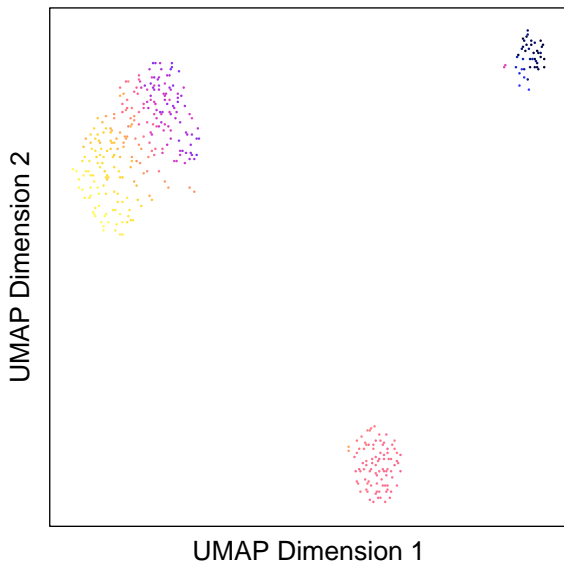
# UMAP\_IterativeLSI of IterativeLSI colored by GeneScoreMatrix : FOSL1



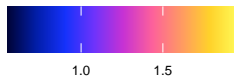
Log2(NormCounts + 1)



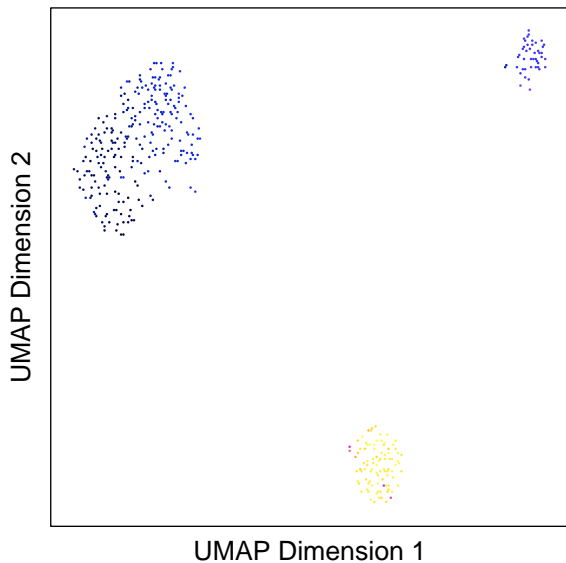
# UMAP\_IterativeLSI of IterativeLSI colored by GeneScoreMatrix : FOSL2



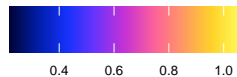
Log2(NormCounts + 1)



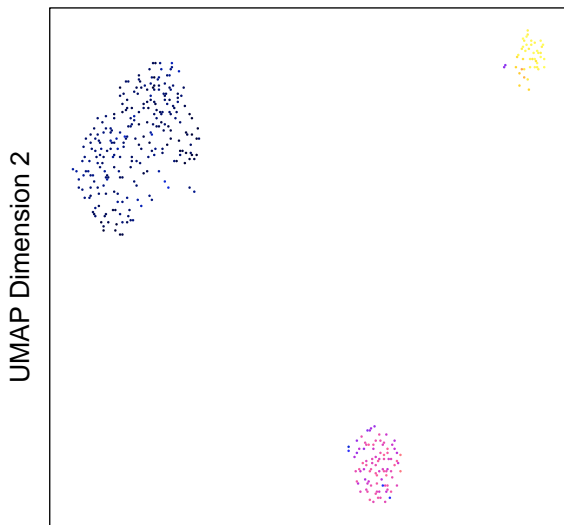
UMAP\_IterativeLSI of IterativeLSI colored by  
GeneScoreMatrix : SPI1



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



# UMAP IterativeLSI of IterativeLSI colored by GeneScoreMatrix : SPIB



UMAP Dimension 1

Log2(NormCounts + 1)

