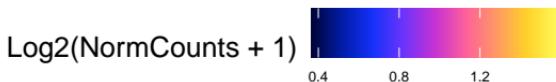
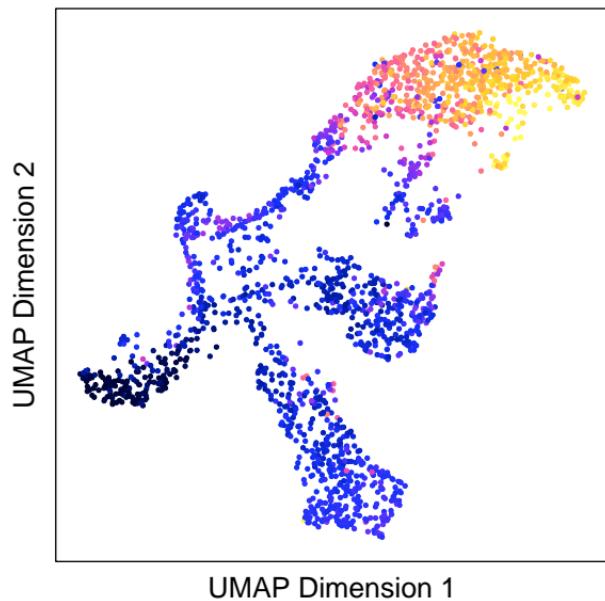
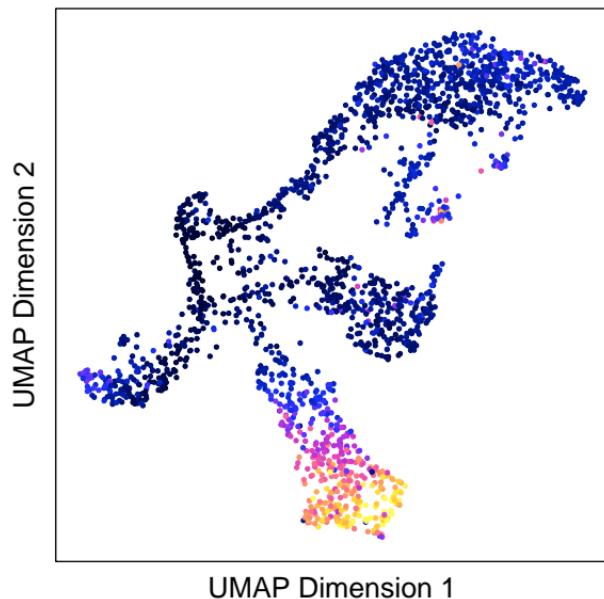


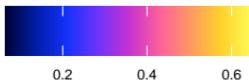
UMAP of IterativeLSI–Endo colored by
GeneScoreMatrix : INS



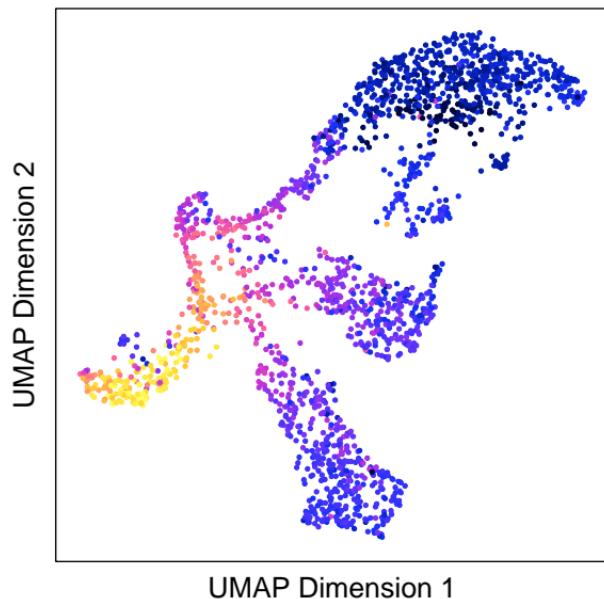
UMAP of IterativeLSI–Endo colored by
GeneScoreMatrix : GCG



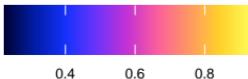
Log2(NormCounts + 1)



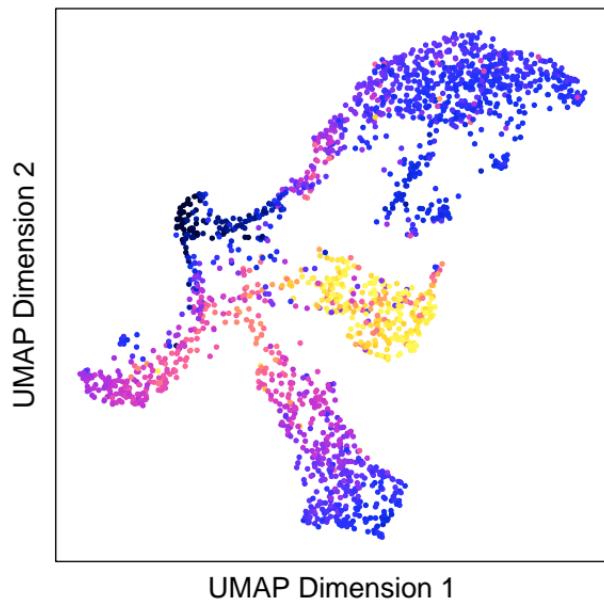
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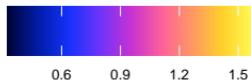
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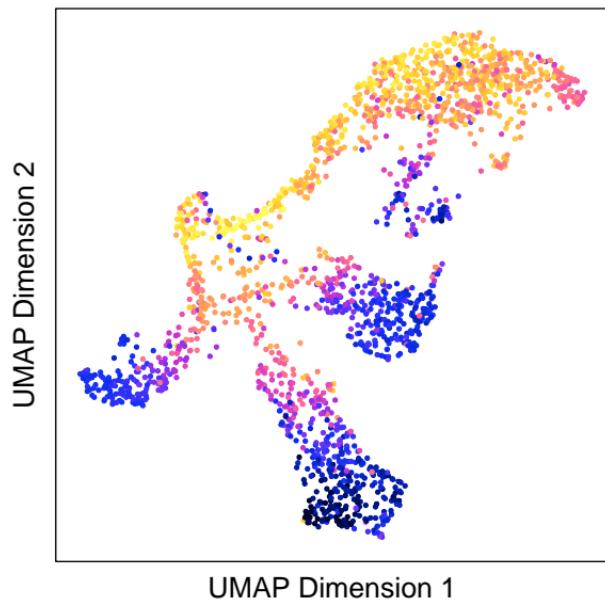
UMAP of IterativeLSI–Endo colored by
GeneScoreMatrix : SST



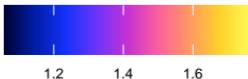
Log2(NormCounts + 1)



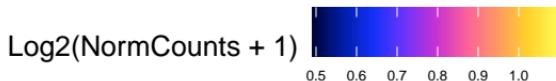
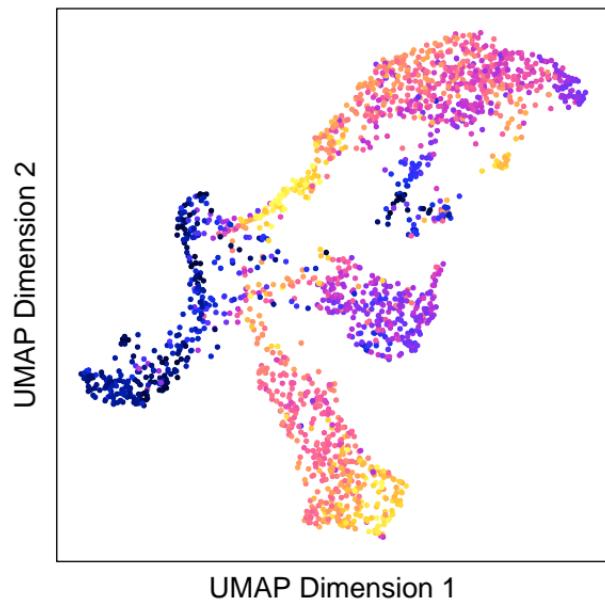
UMAP of IterativeLSI–Endo colored by
GeneScoreMatrix : NEUROG3



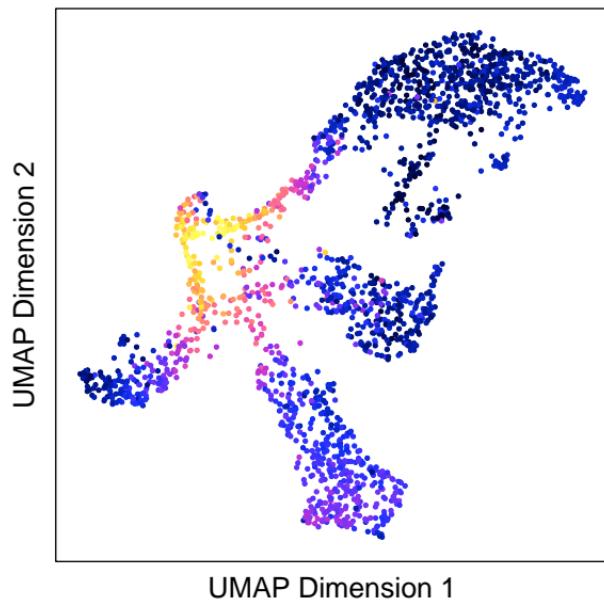
Log2(NormCounts + 1)



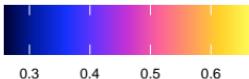
UMAP of IterativeLSI–Endo colored by
GeneScoreMatrix : FEV



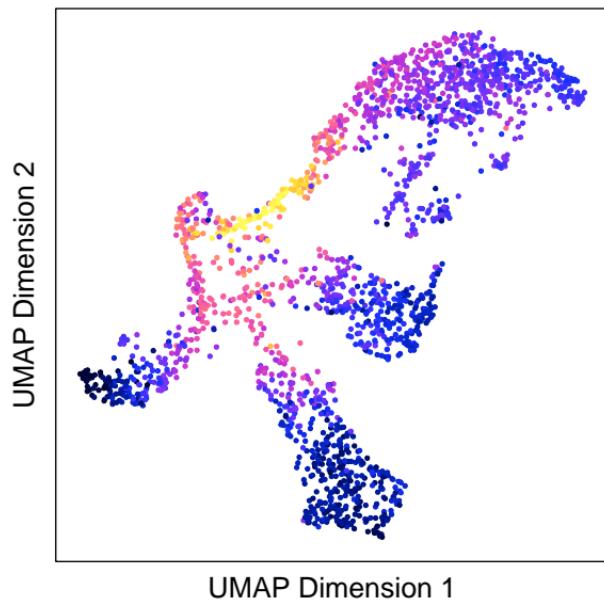
UMAP of IterativeLSI–Endo colored by
GeneScoreMatrix : SUSD2



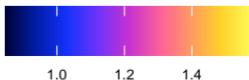
Log2(NormCounts + 1)



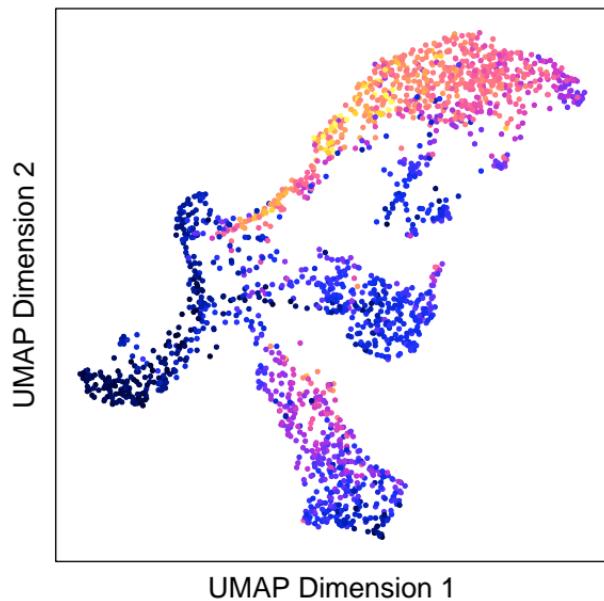
UMAP of IterativeLSI–Endo colored by
GeneScoreMatrix : LMX1B



Log2(NormCounts + 1)



UMAP of IterativeLSI–Endo colored by
GeneScoreMatrix : PRPH



Log2(NormCounts + 1) 
0.4 0.5 0.6 0.7 0.8