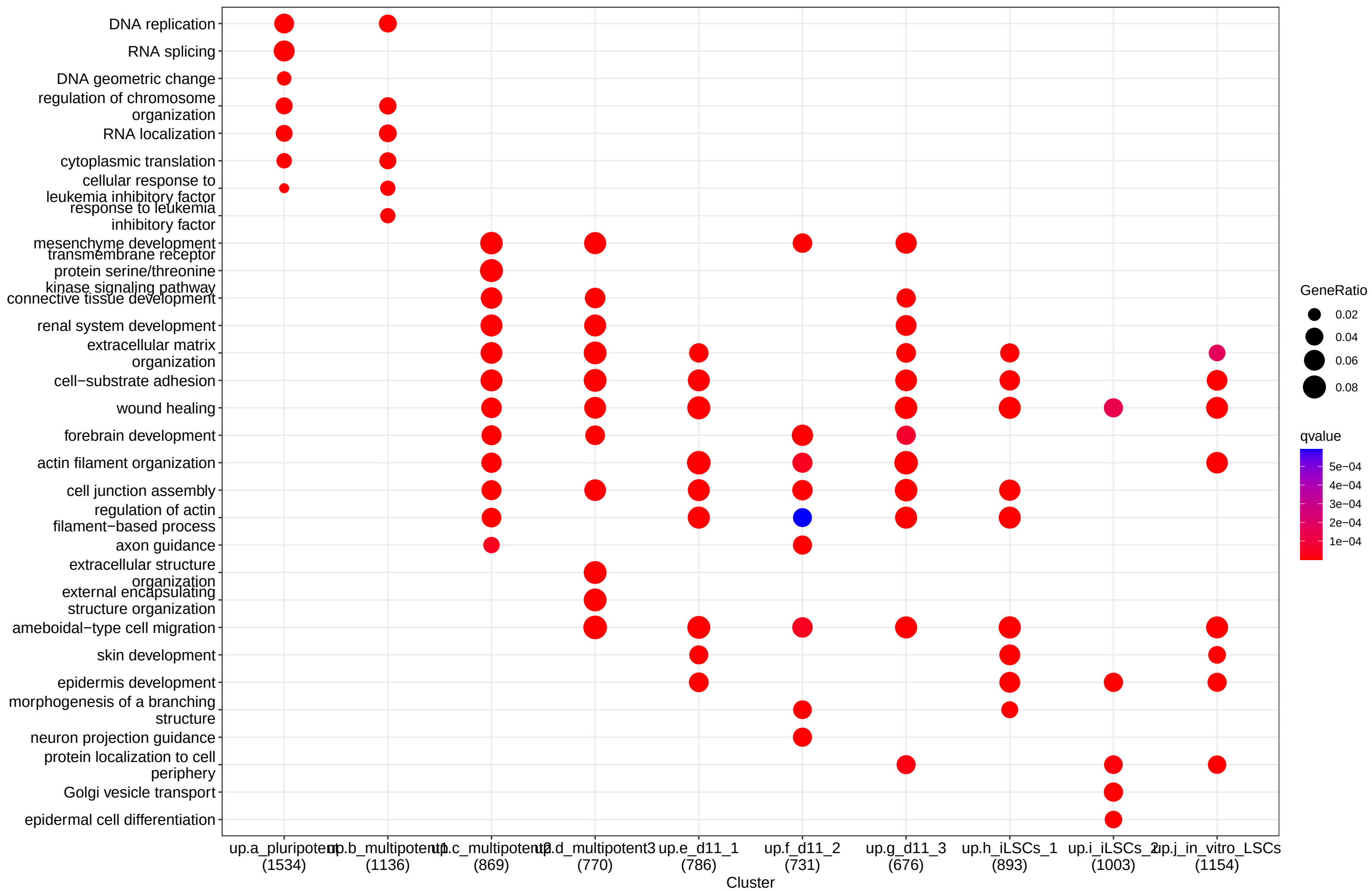
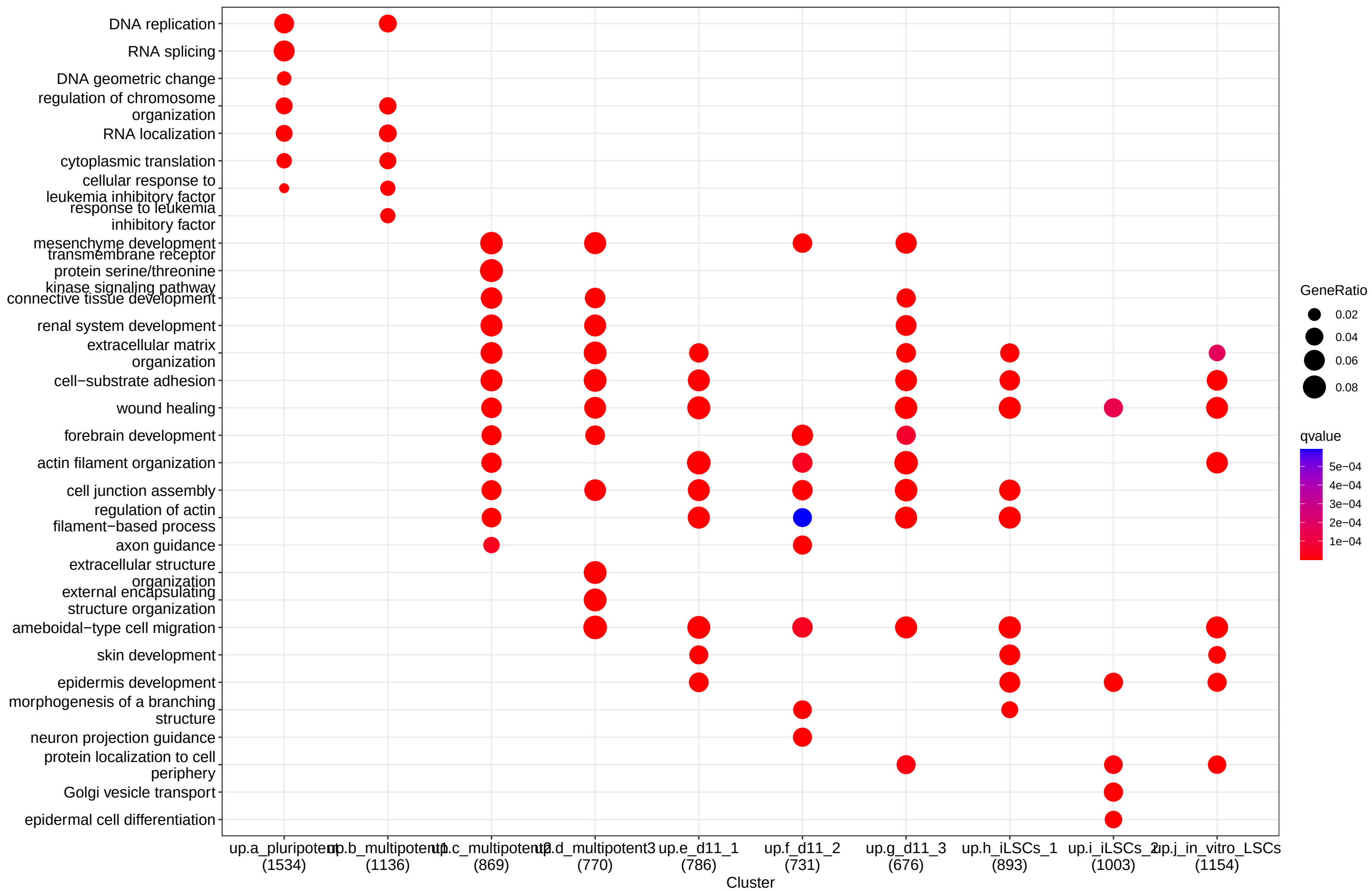
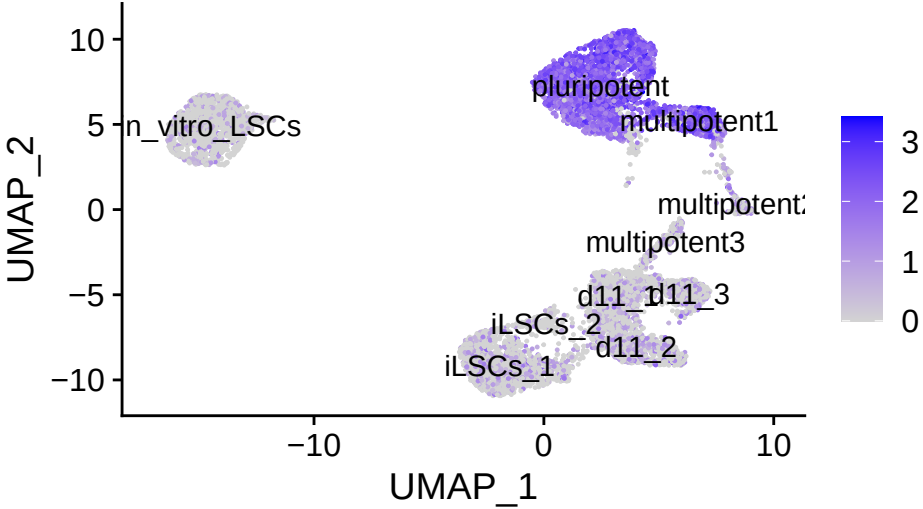




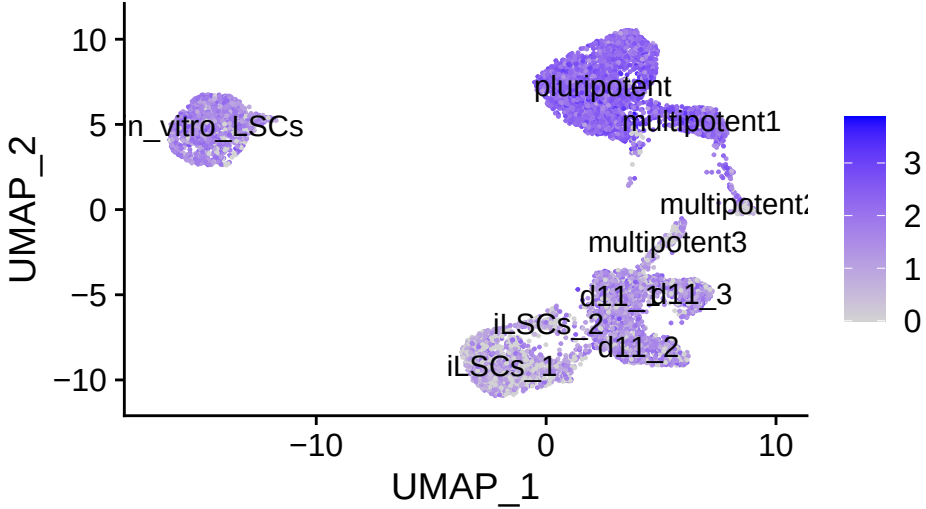


enriched go term: DNA replication from cluster: a_pluripotent

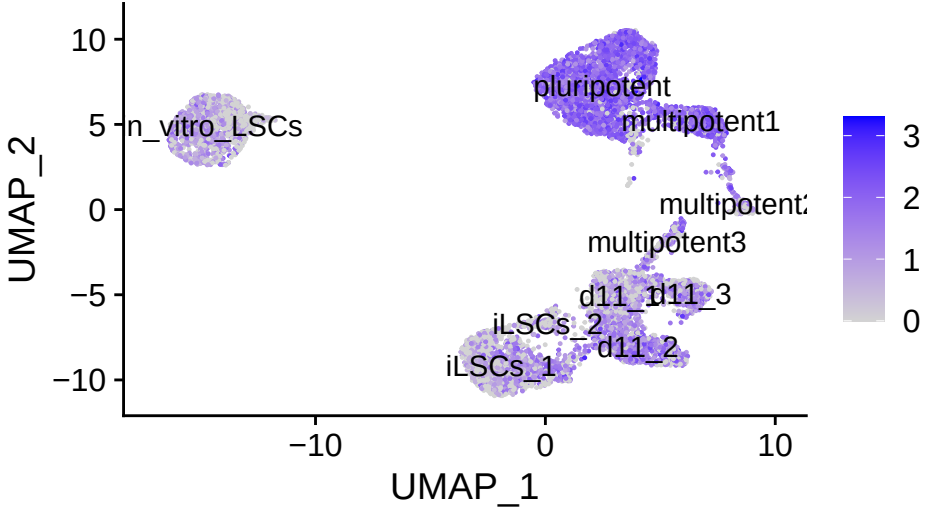
TERF1



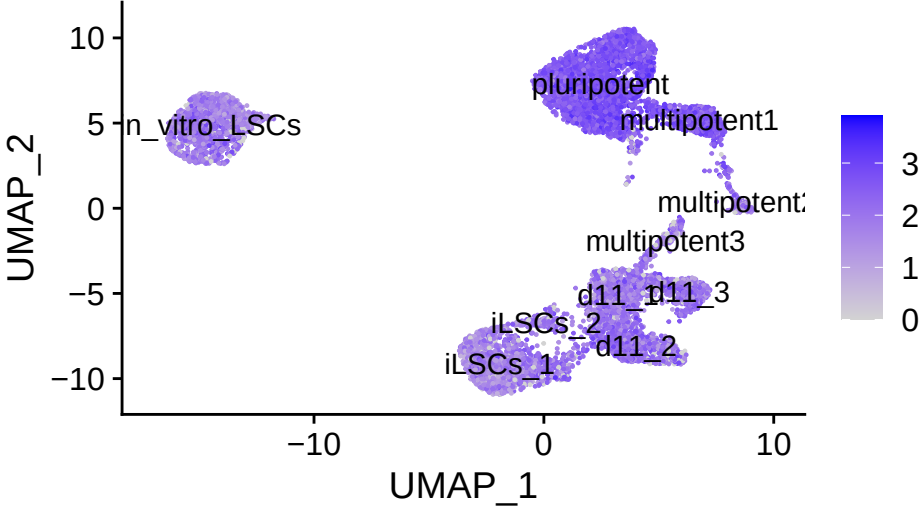
HMGA1



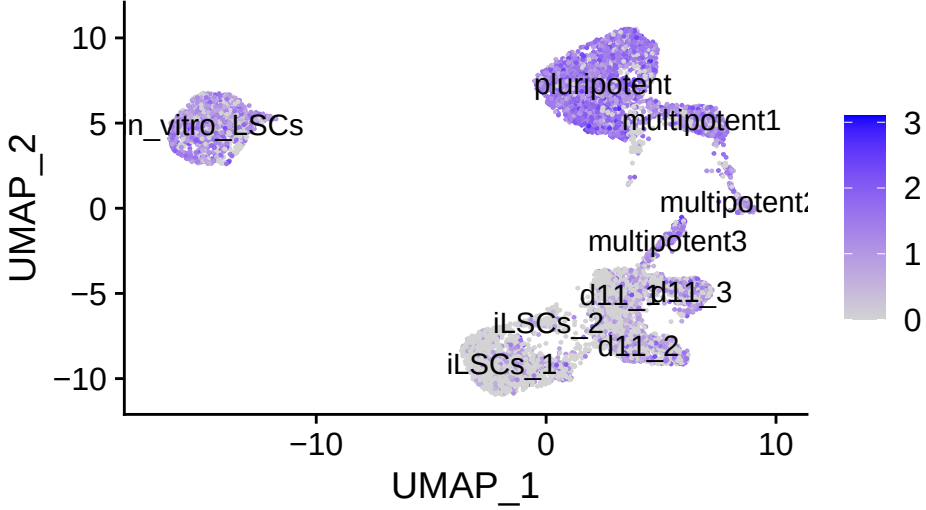
NASP



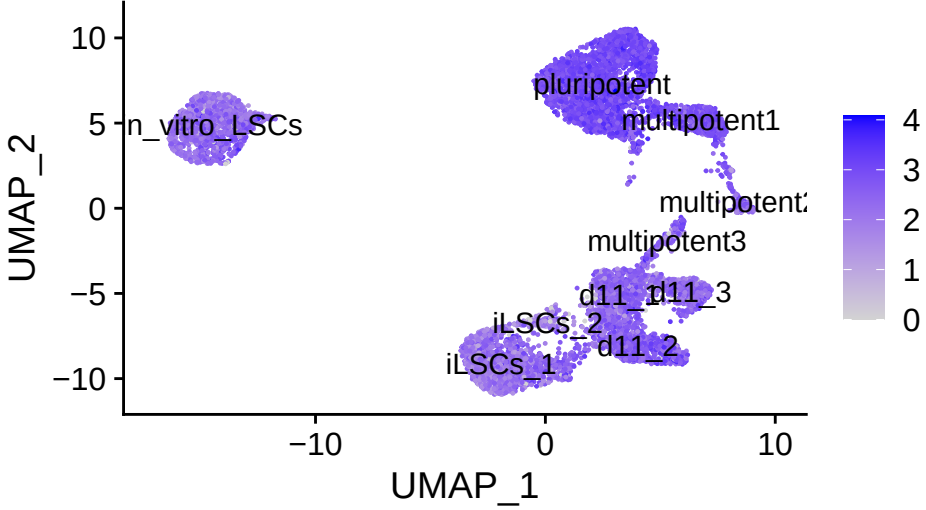
SET



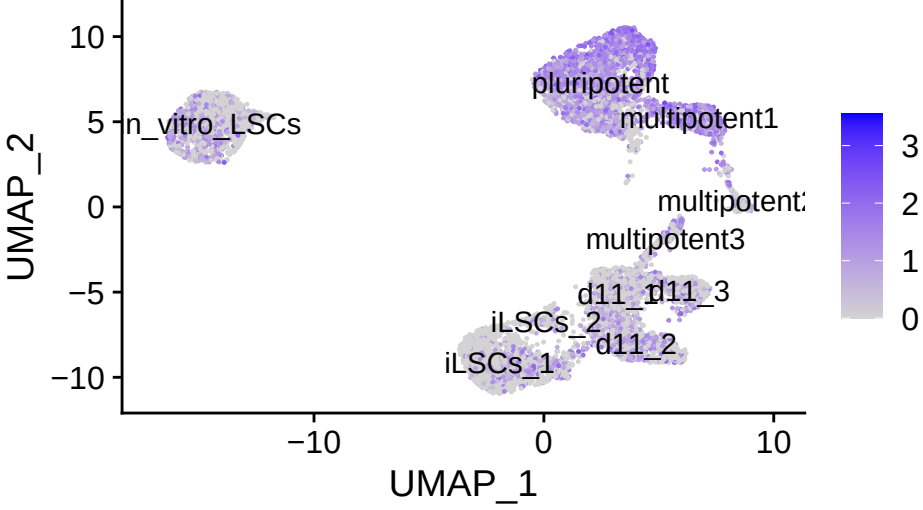
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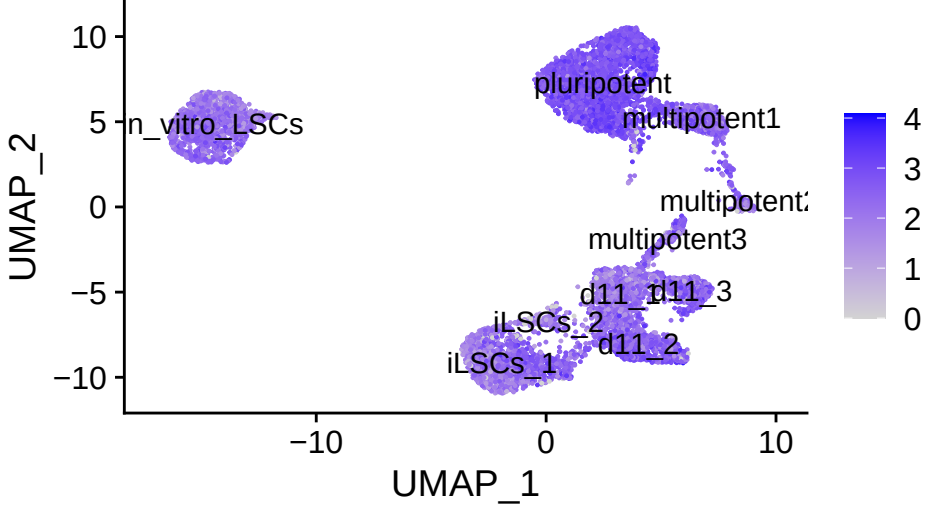
NAP1L1



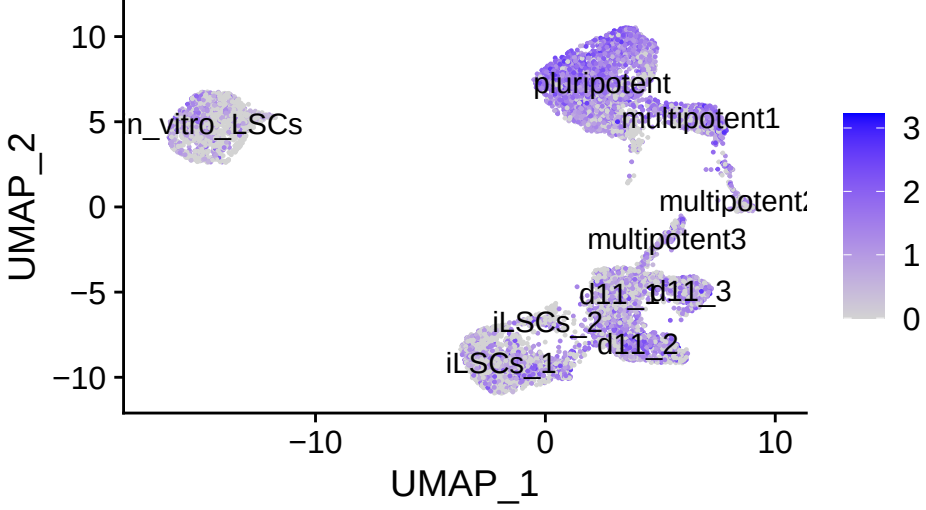
WDHD1



NUCKS1

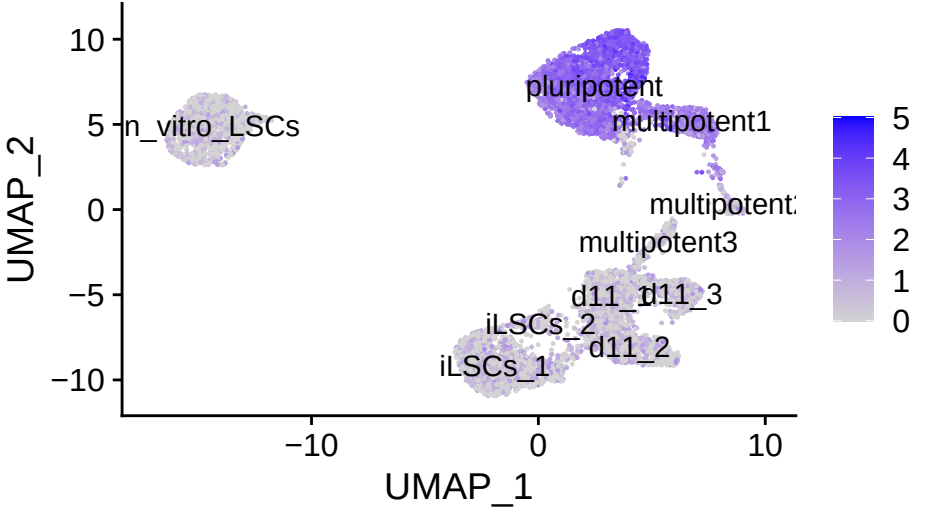


POLA1

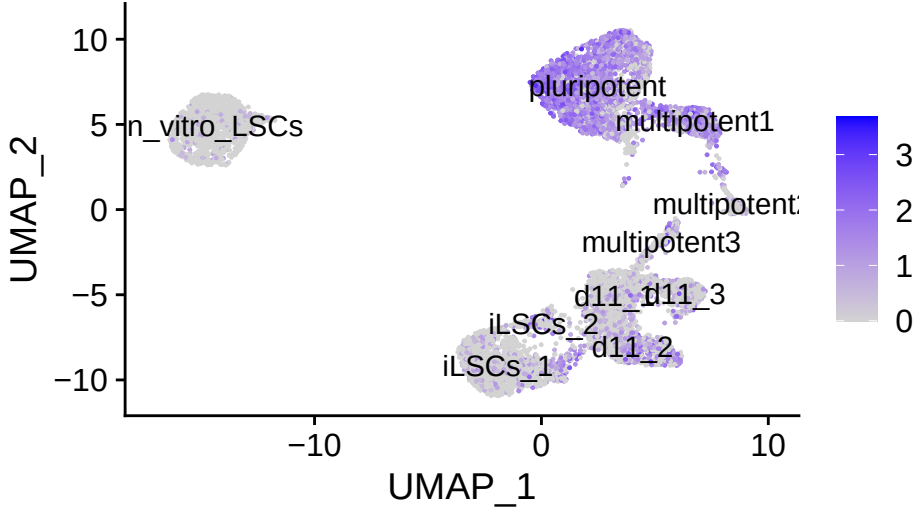


enriched go term: RNA splicing from cluster: a_pluripotent

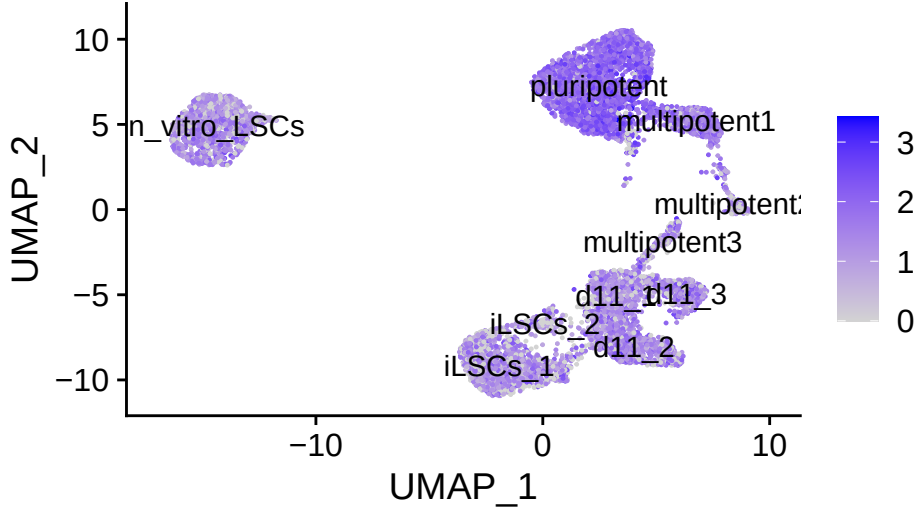
LARP7



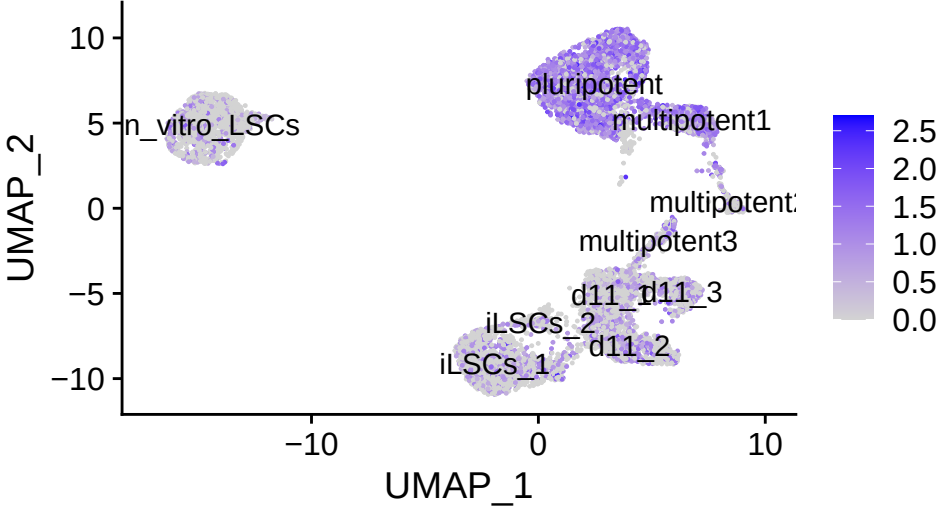
KHDRBS2



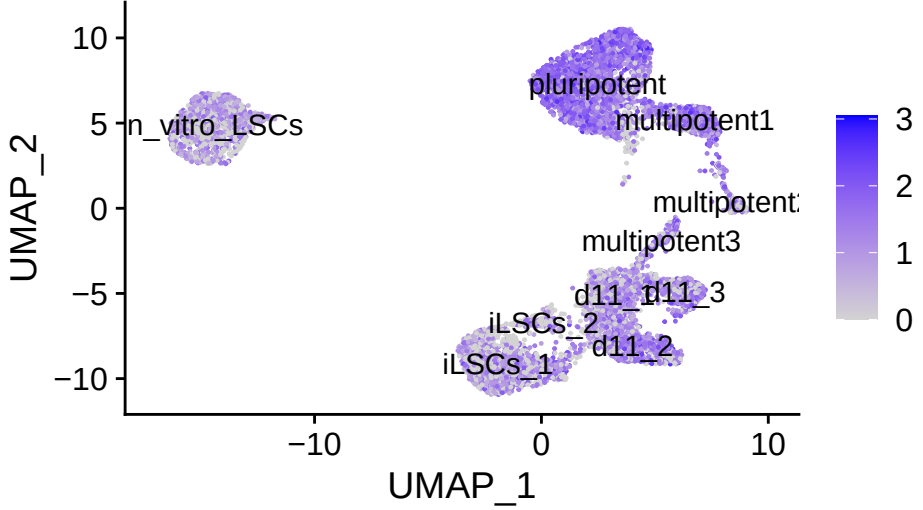
NPM1



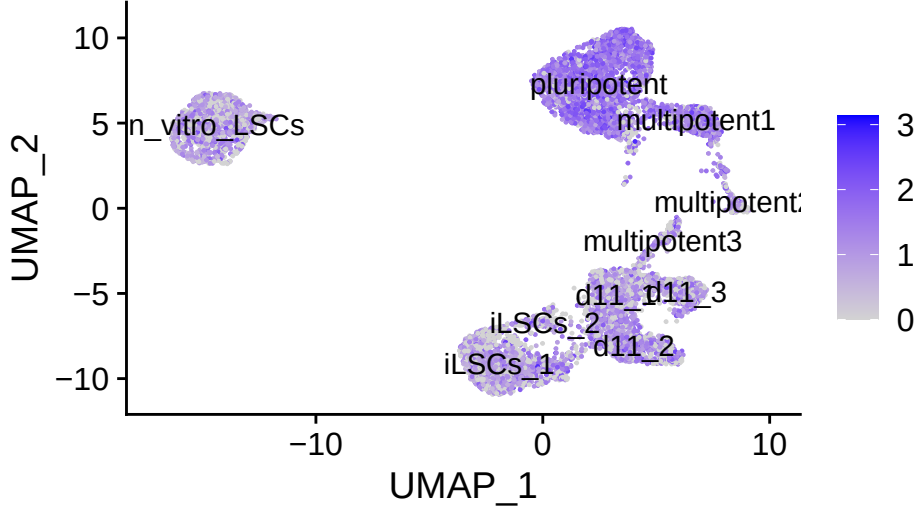
PSIP1



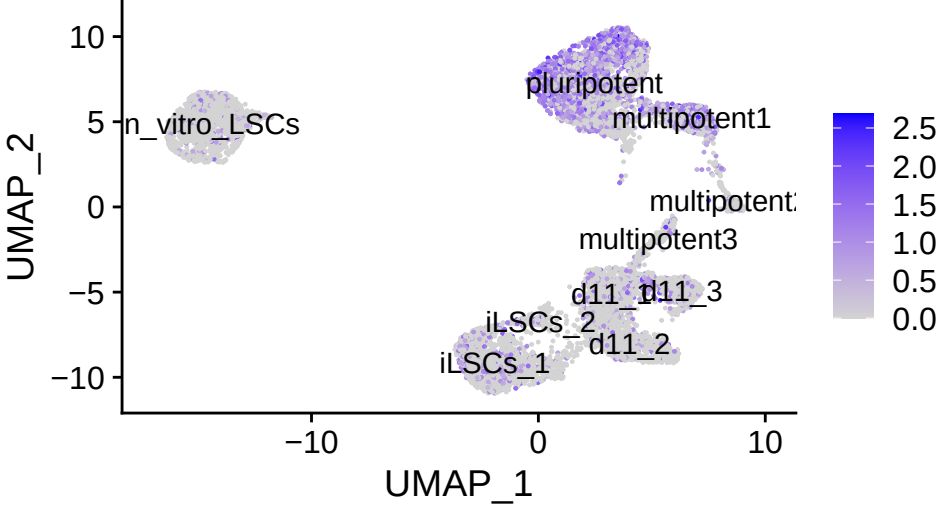
PTBP2



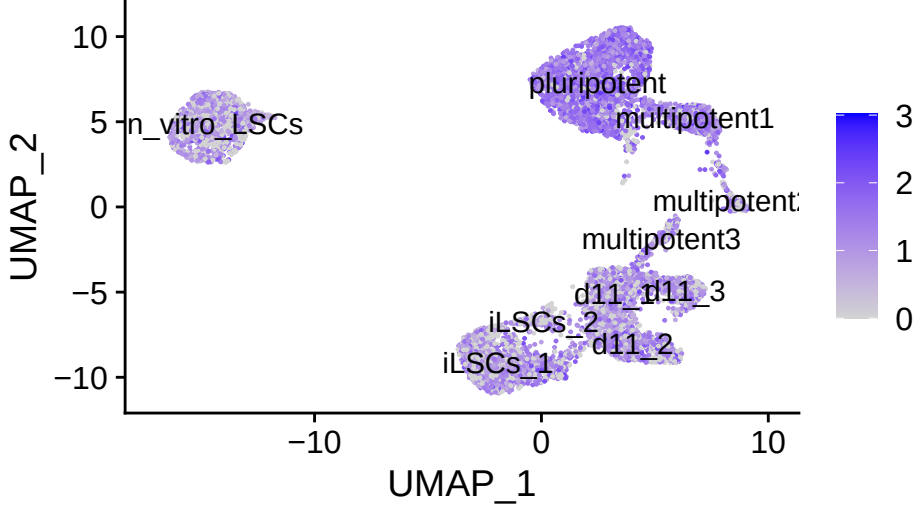
PNN



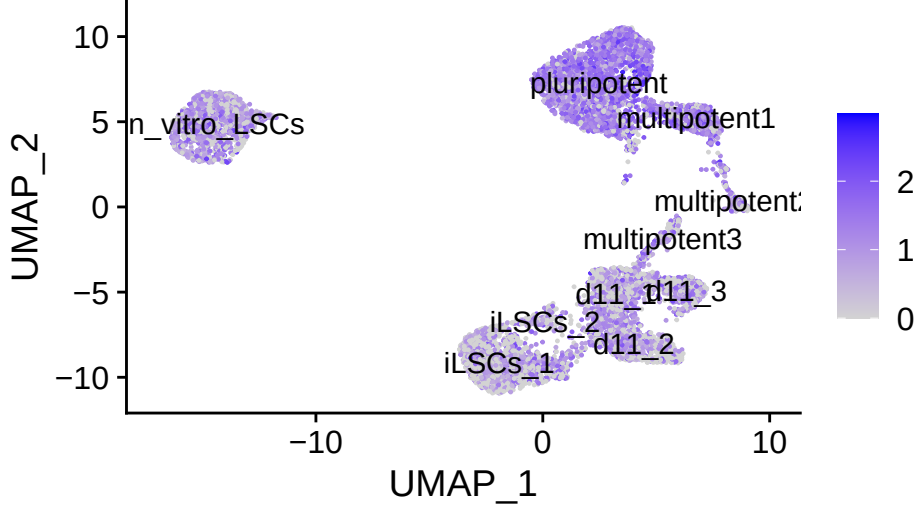
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FXR1

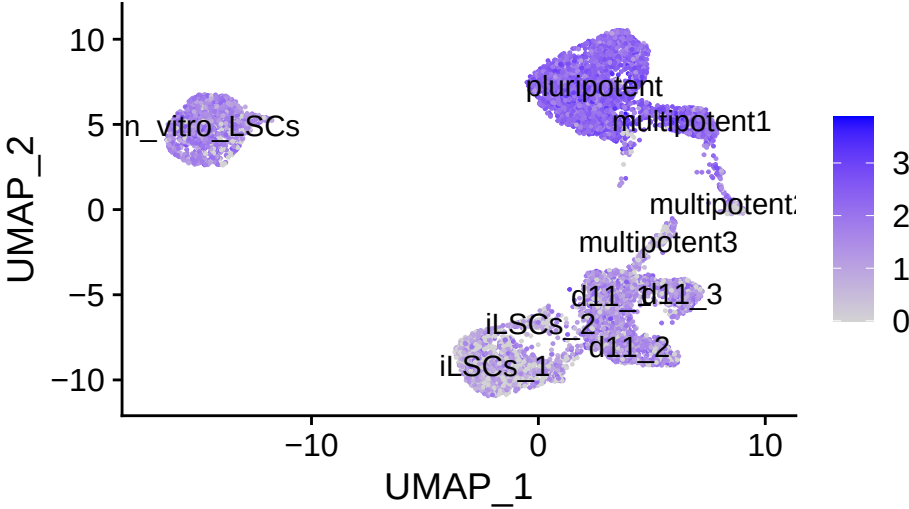


SNRPF

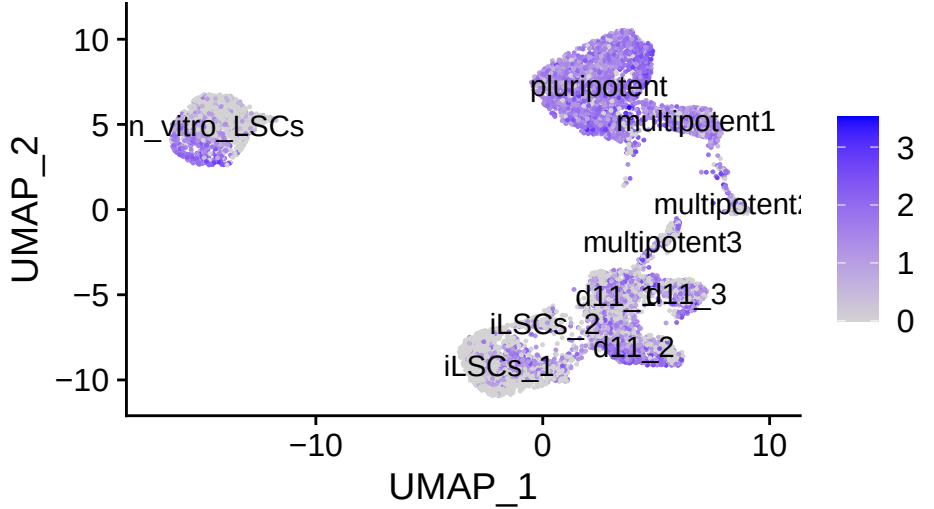


enriched go term: DNA geometric change from cluster: a_pluripotent

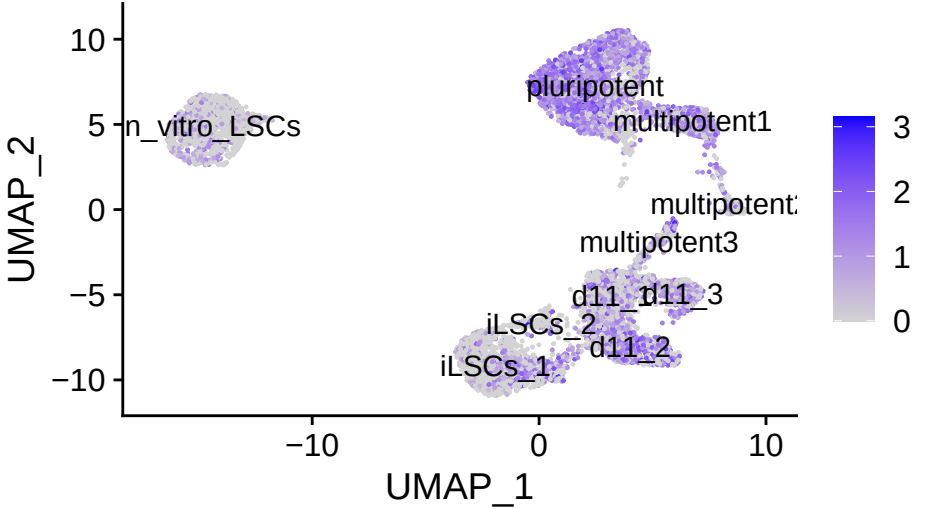
HMGA1



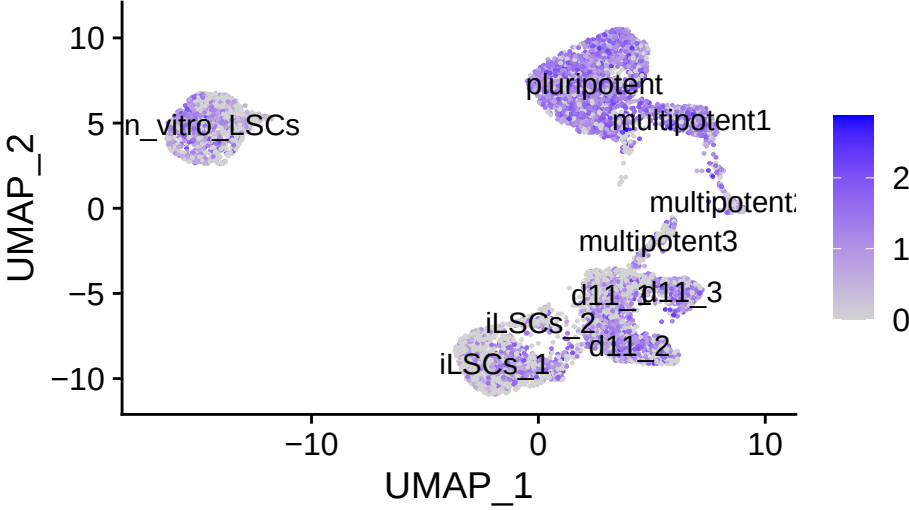
HMGB2



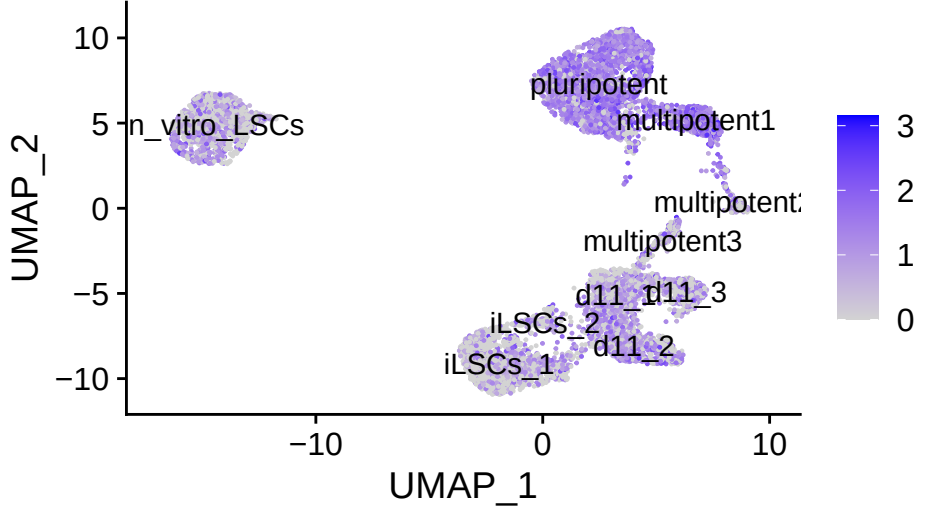
RFC3



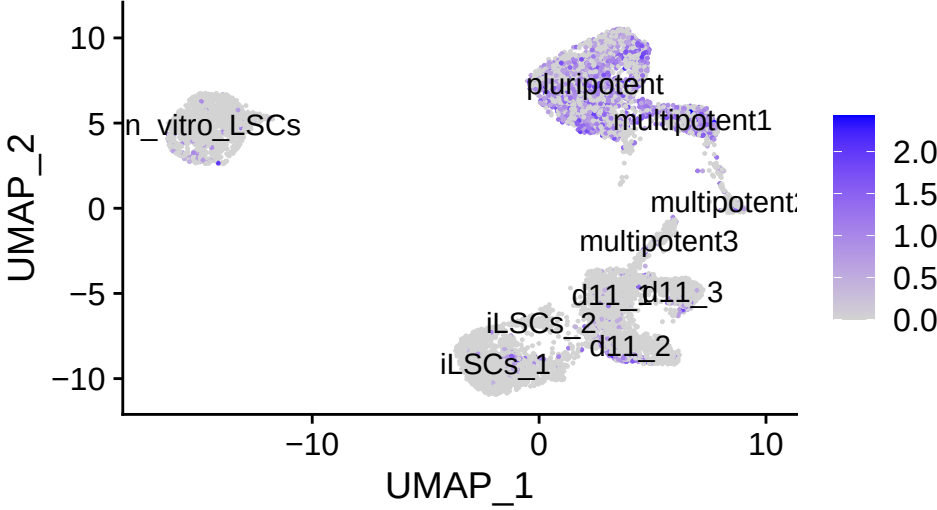
MCM4



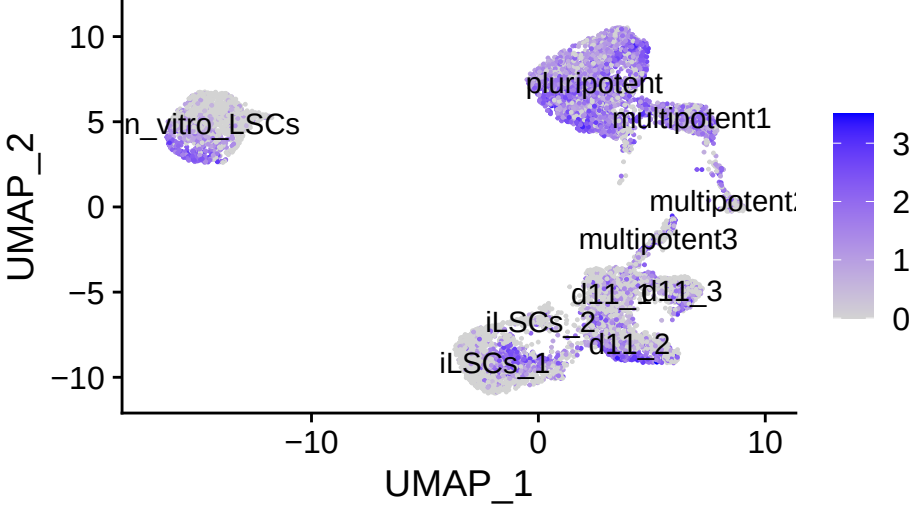
HMGB3



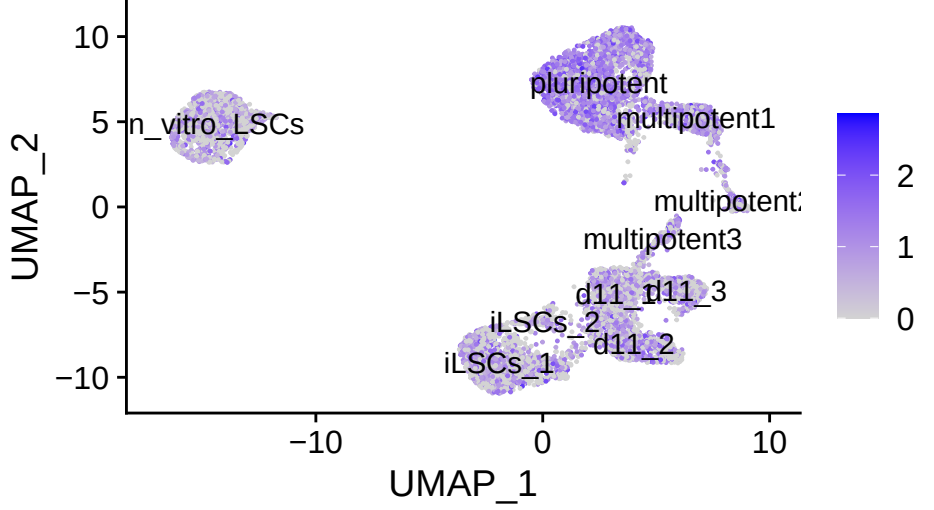
PIF1



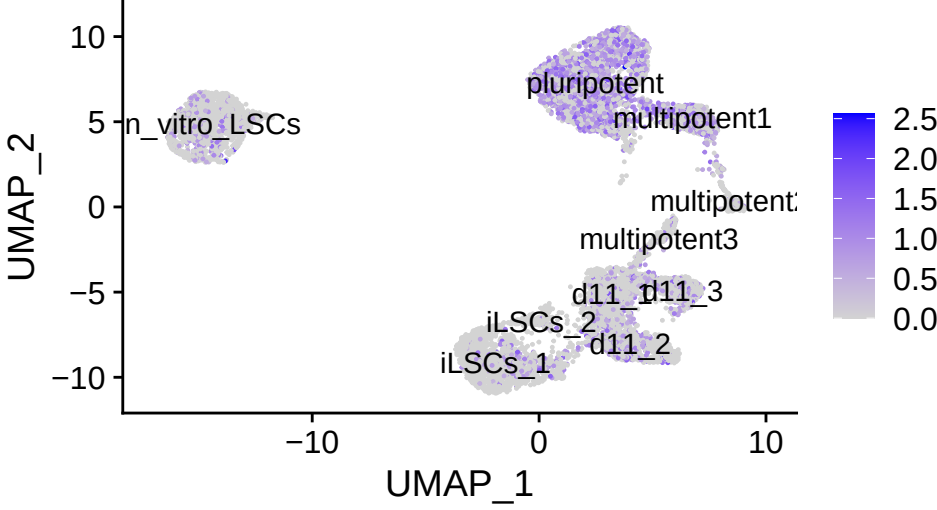
TOP2A



WRN

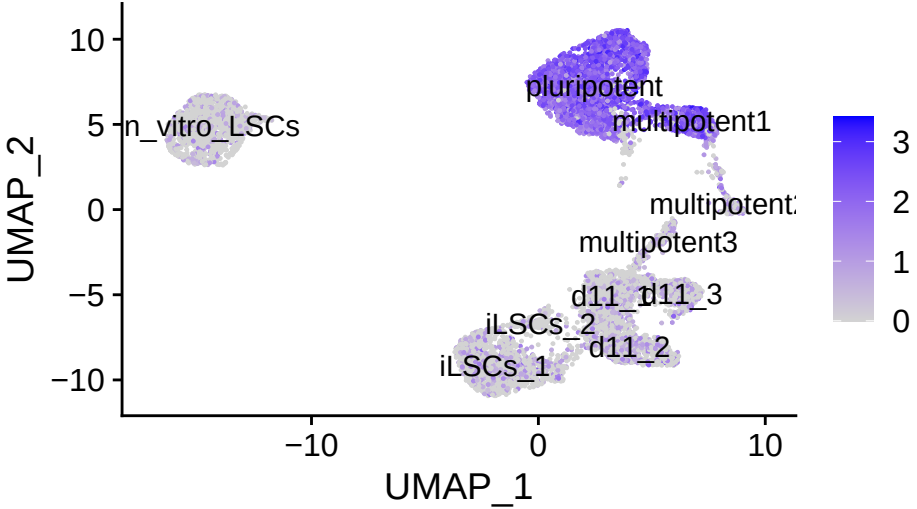


DSCC1

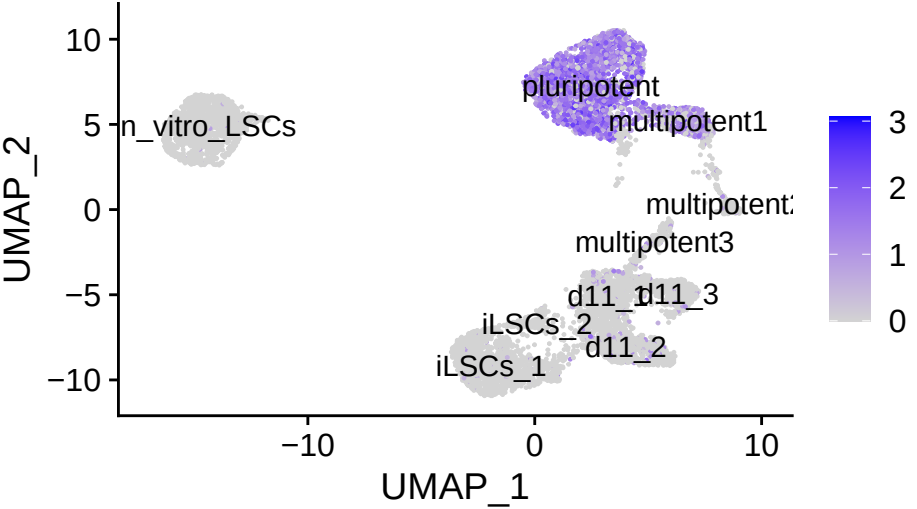


enriched go term: regulation of chromosome organization from cluster: a_pluripotent

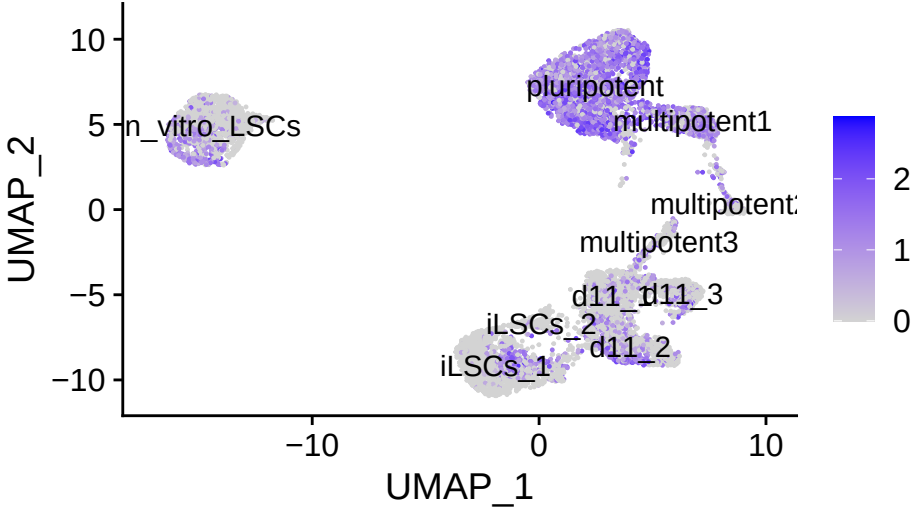
TERF1



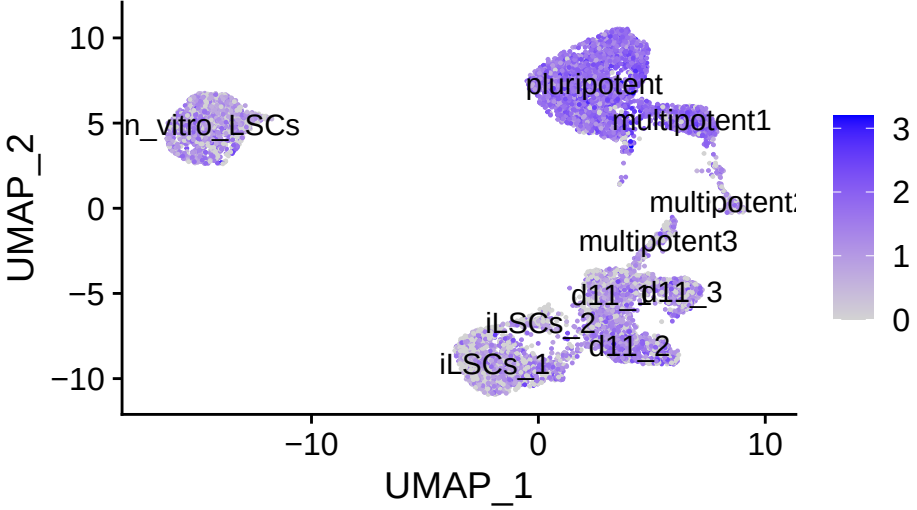
USP44



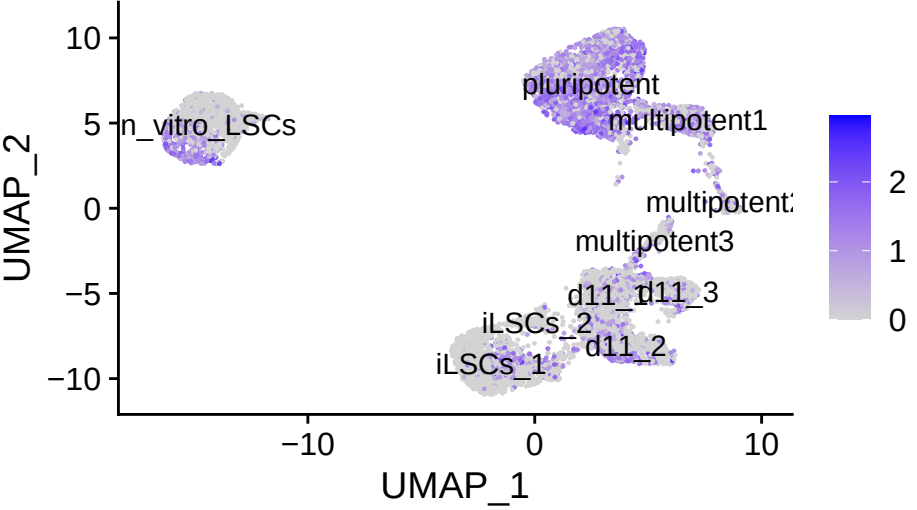
DLGAP5



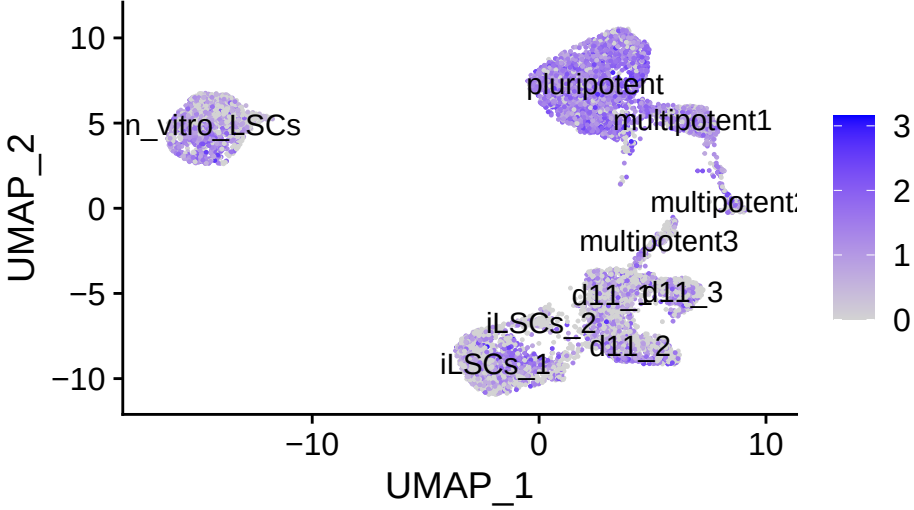
PARP1



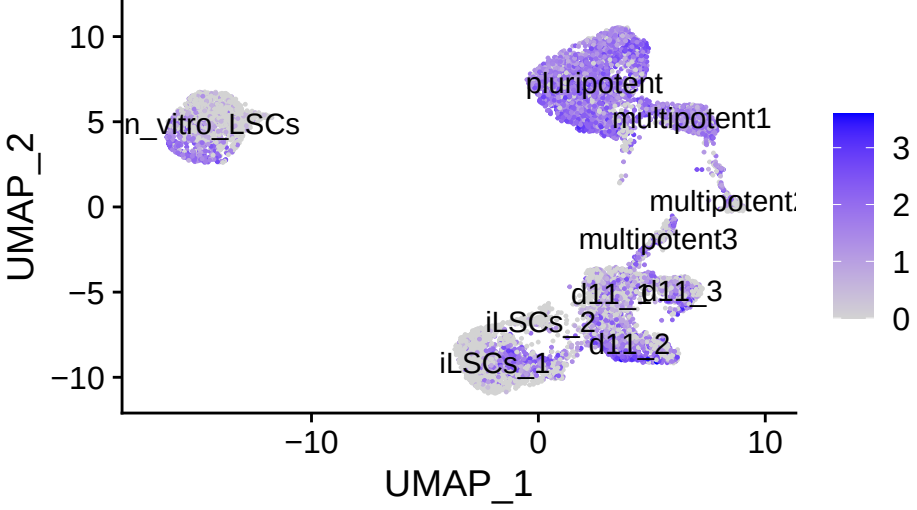
BUB1



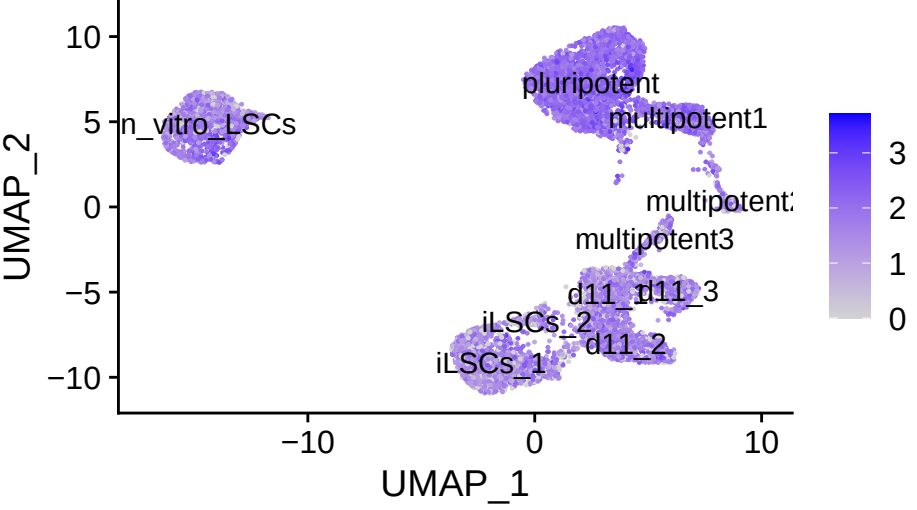
PTTG1



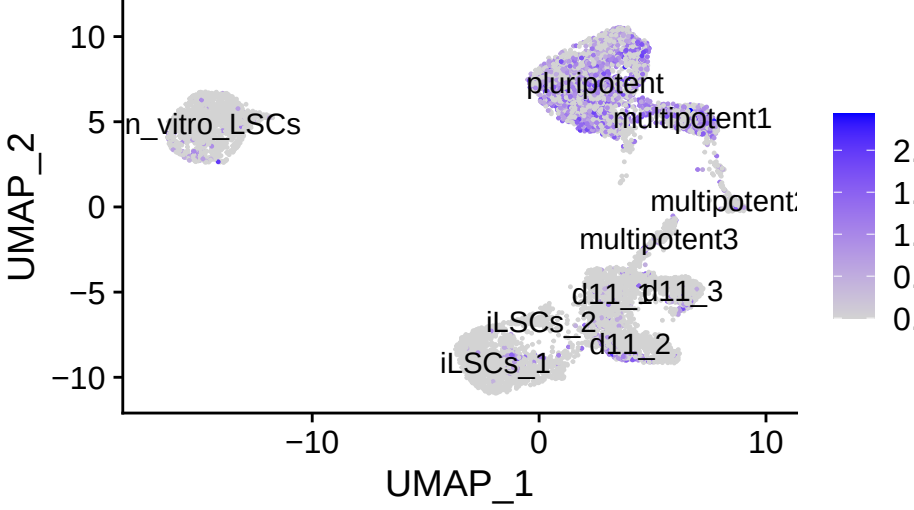
CENPF



CCT4

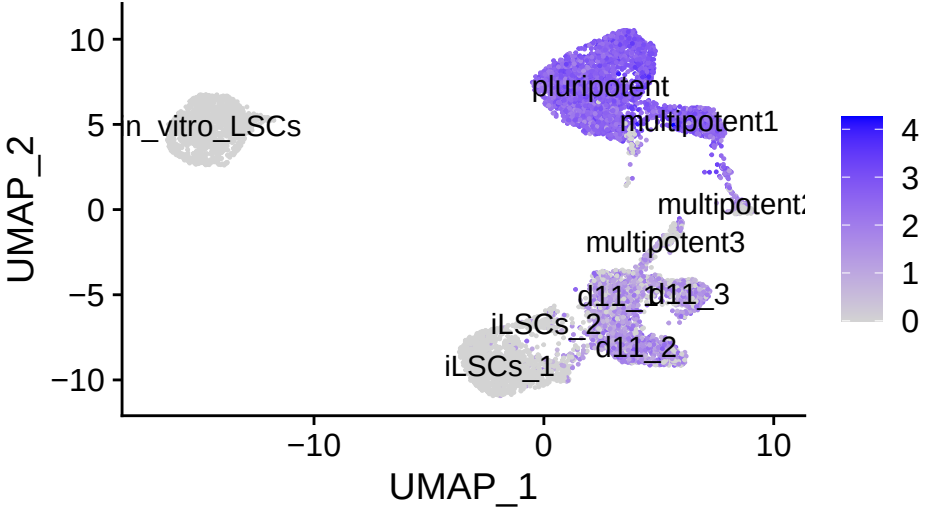


PIF1

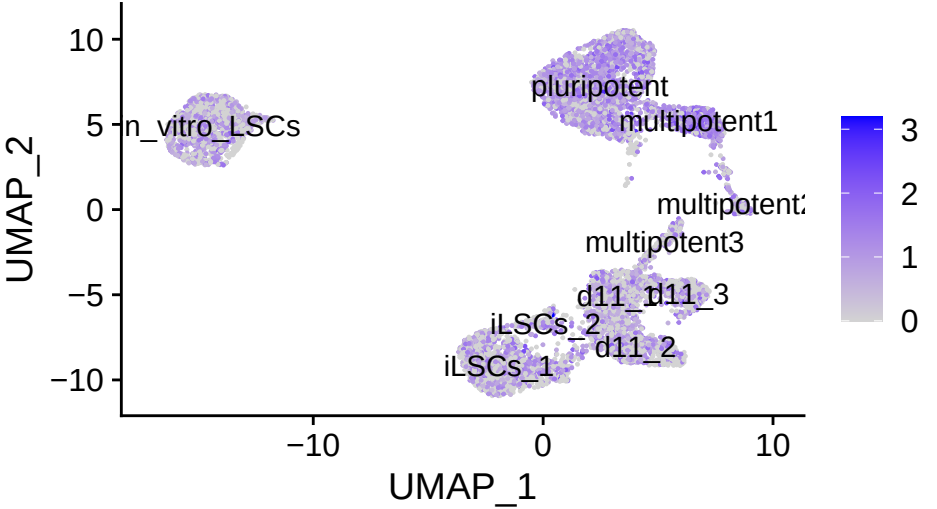


enriched go term: cytoplasmic translation from cluster: b_multipotent1

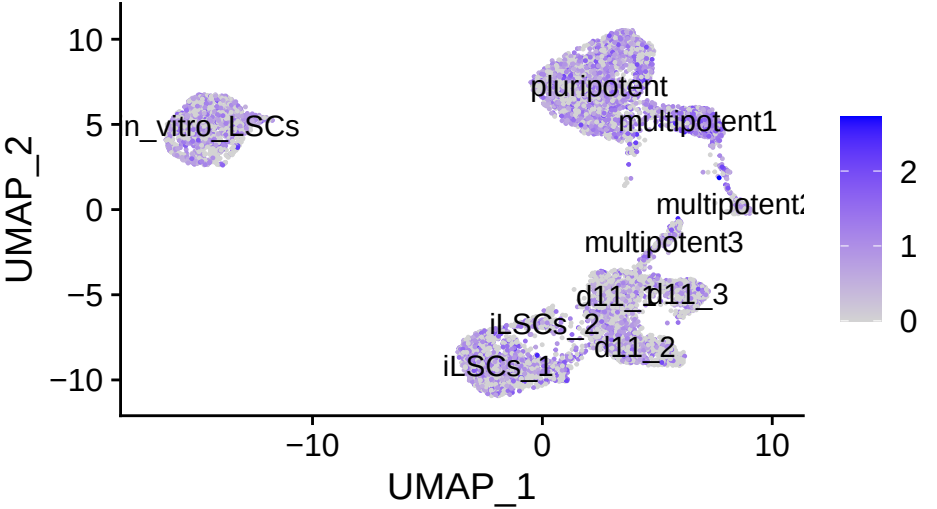
LIN28A



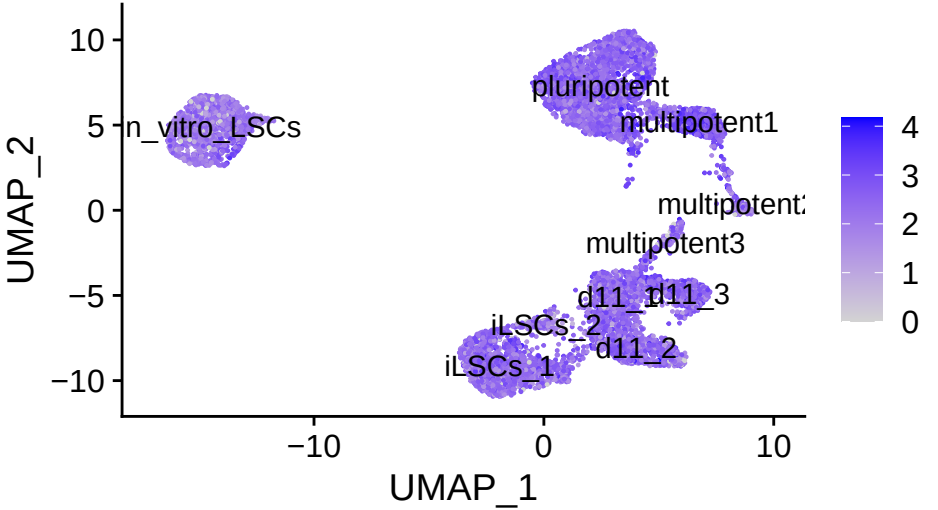
AARS1



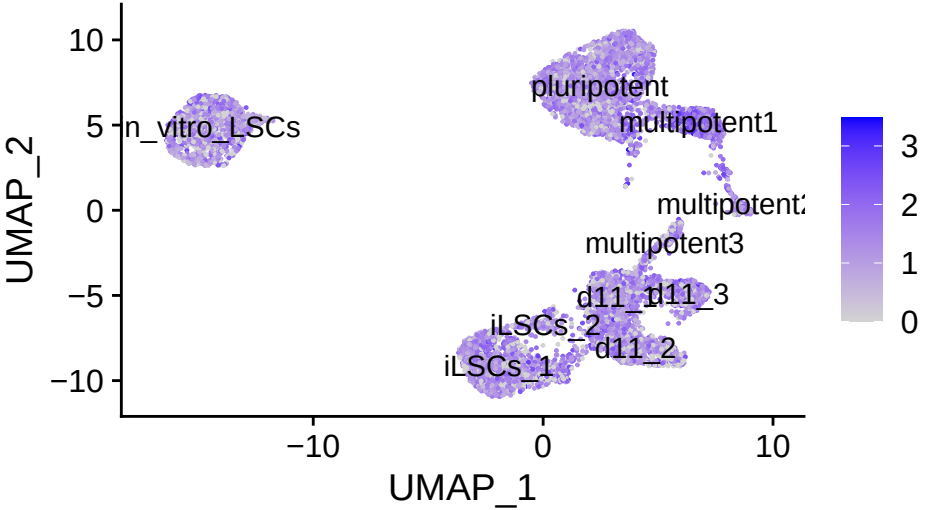
EIF3K



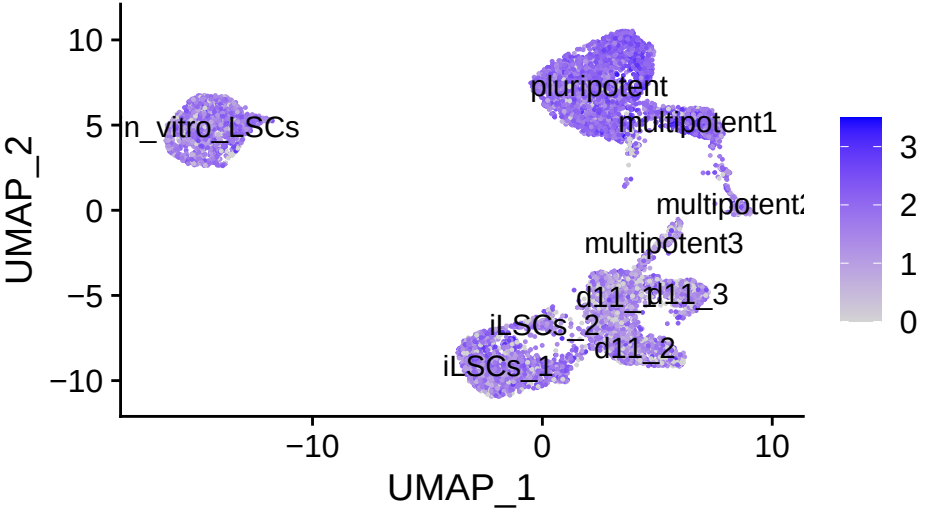
RPS15



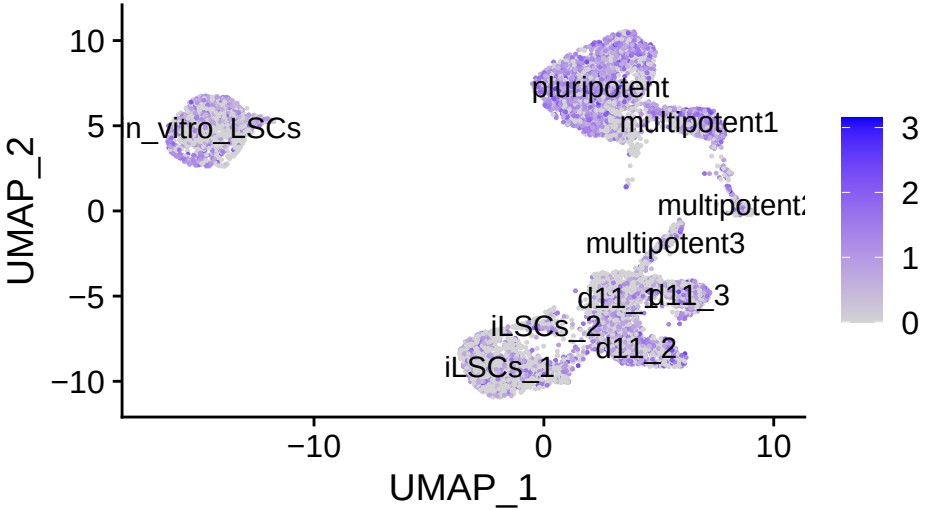
RPS28



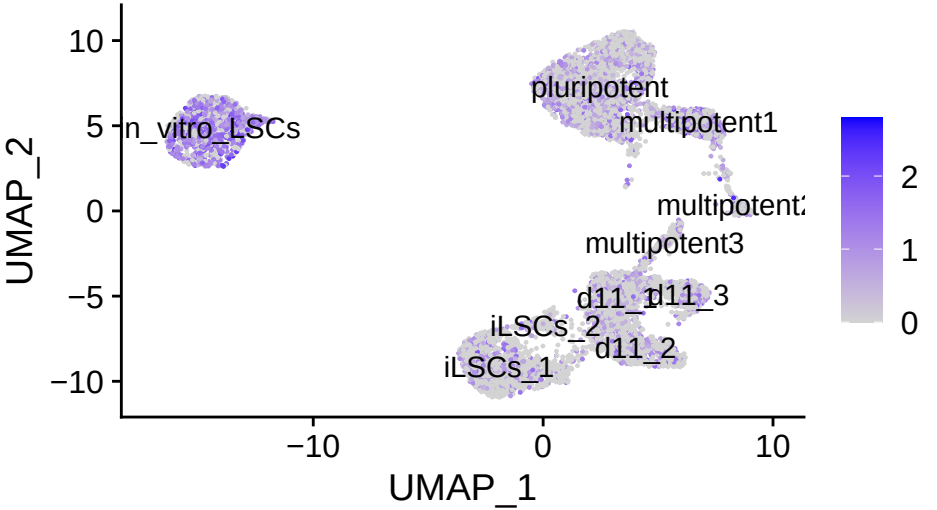
RPL13A



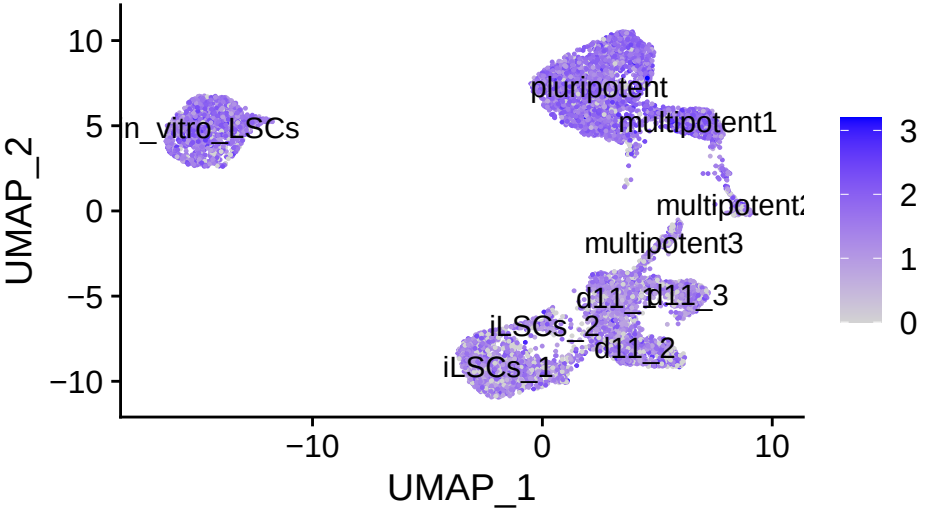
DPH6



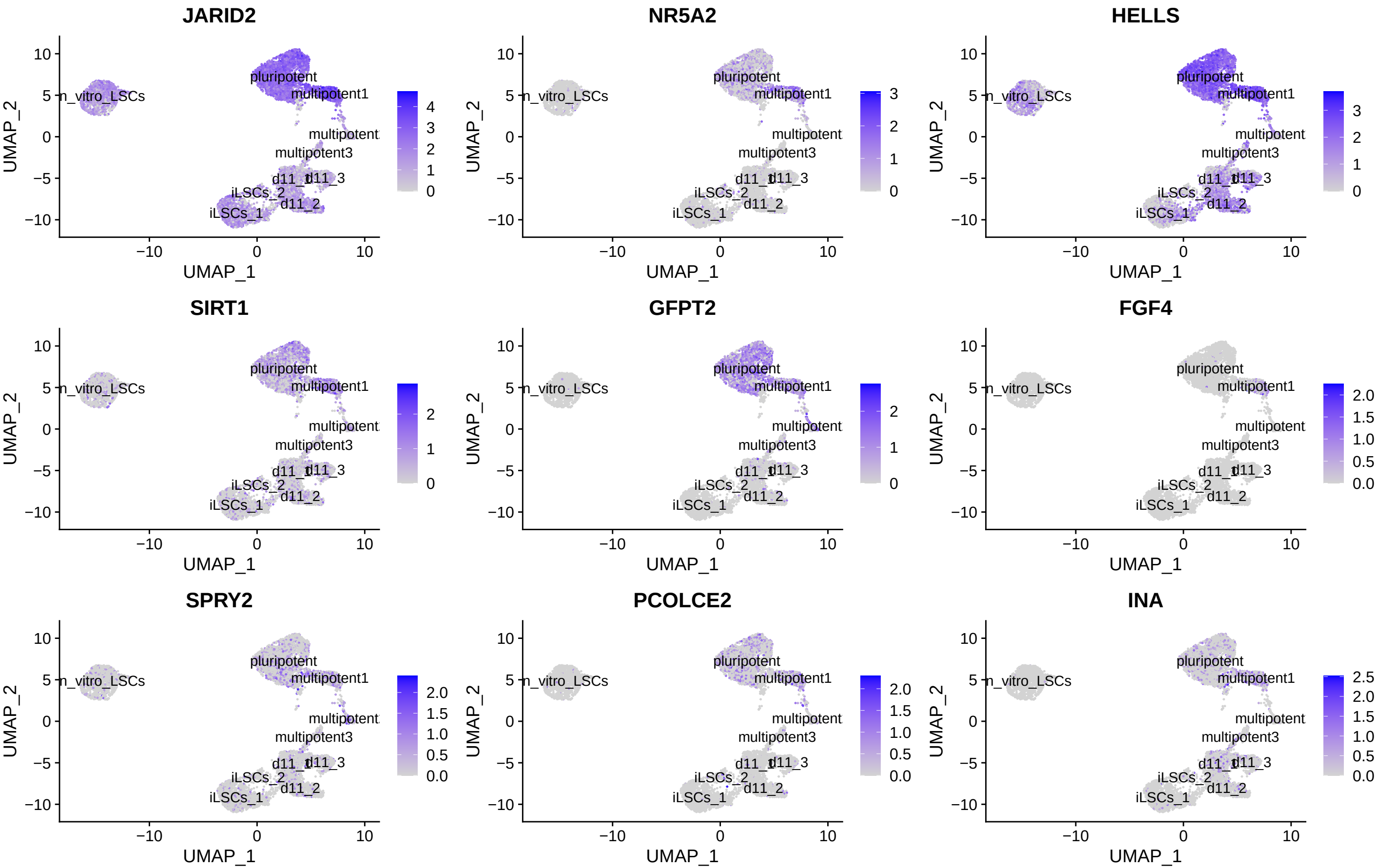
RPL9



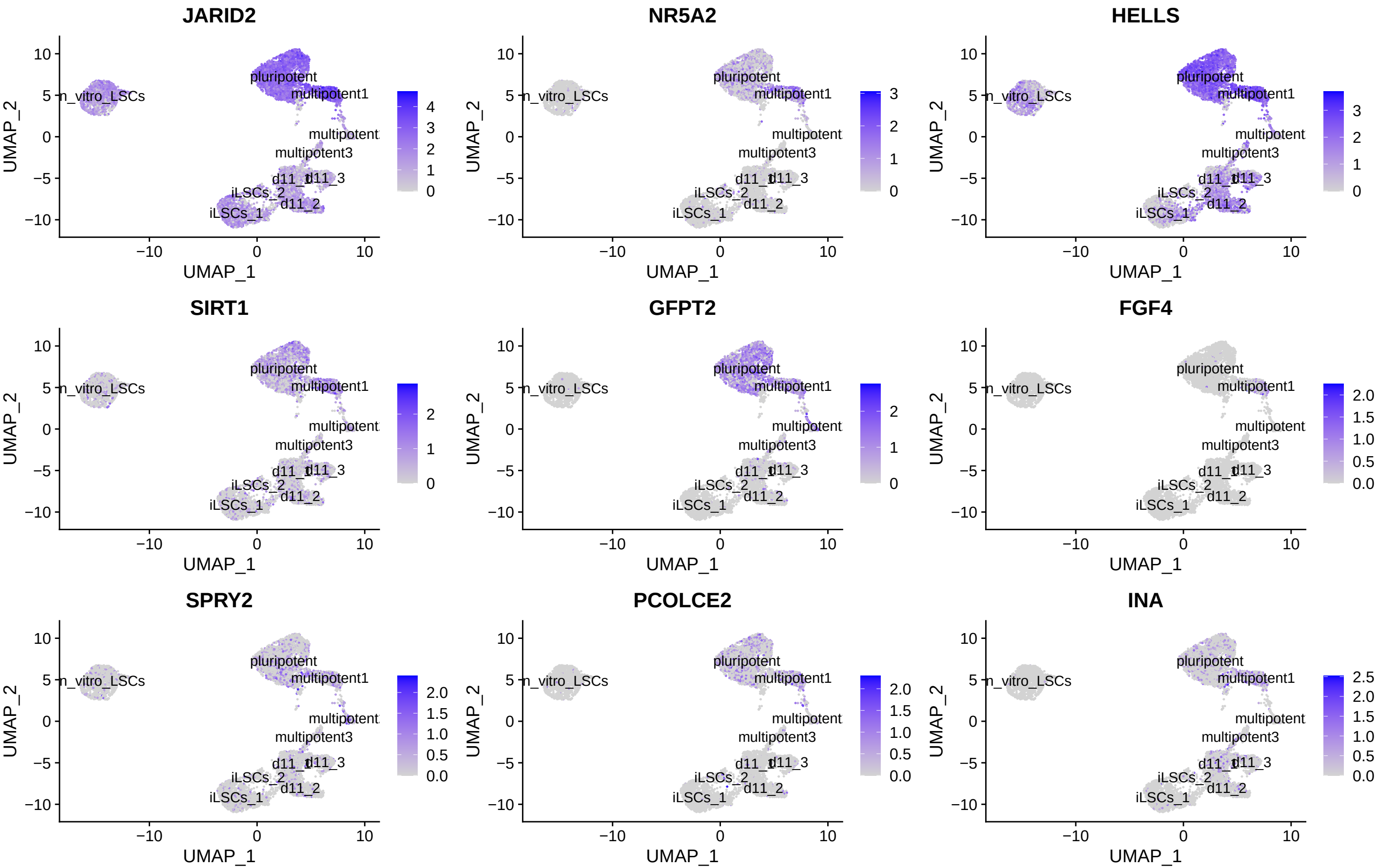
EIF4B



enriched go term: cellular response to leukemia inhibitory factor from cluster: b_multipotent1

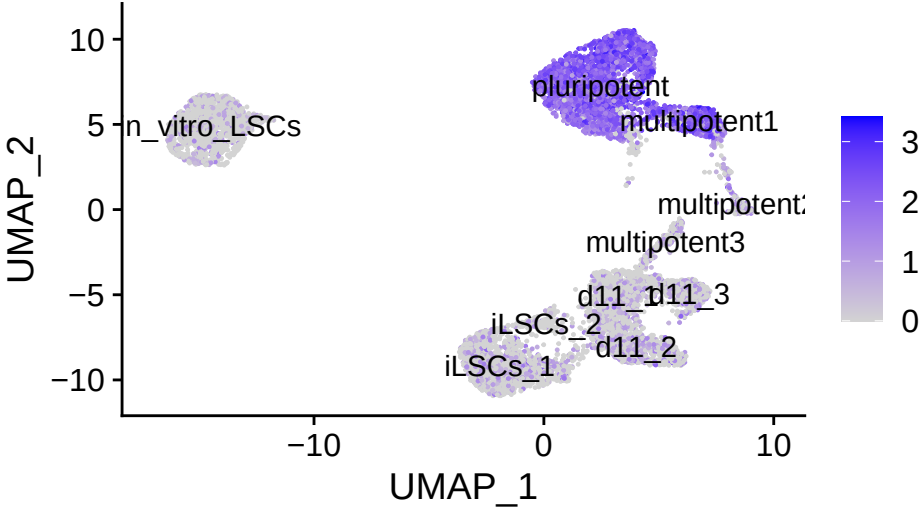


enriched go term: response to leukemia inhibitory factor from cluster: b_multipotent1

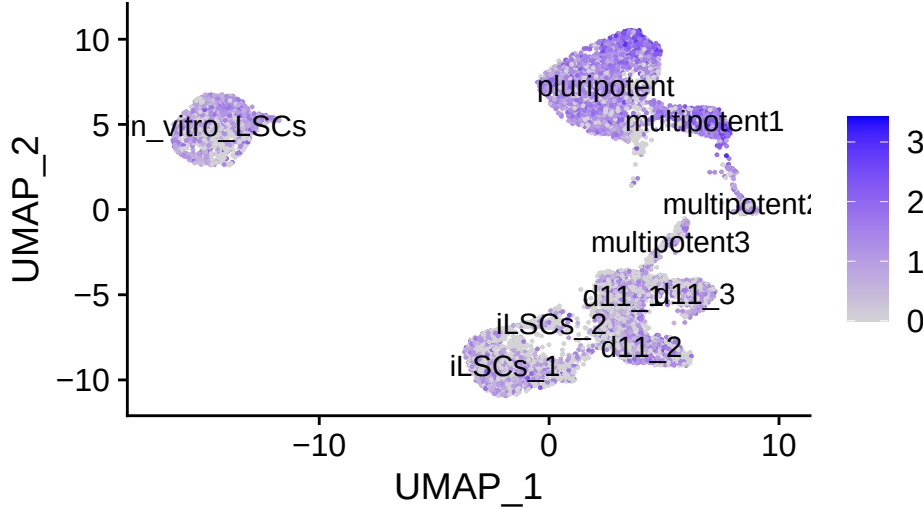


enriched go term: RNA localization from cluster: b_multipotent1

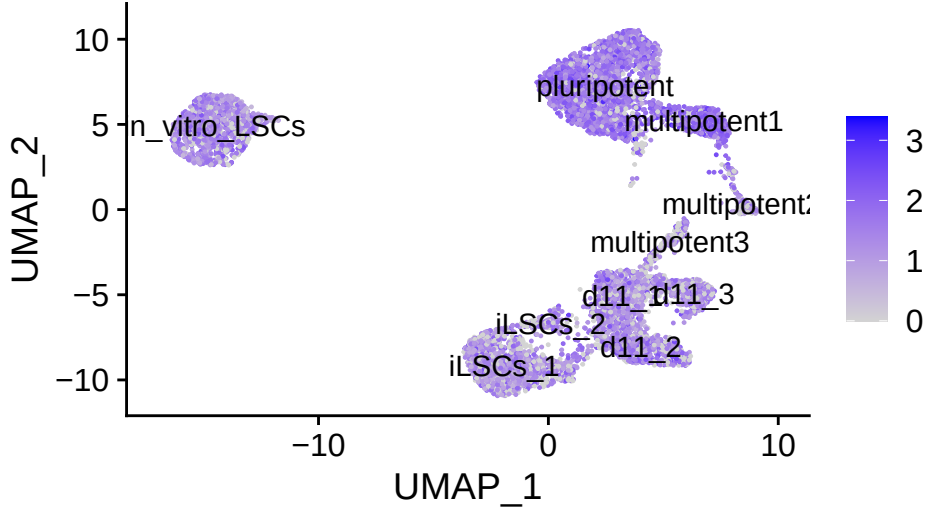
TERF1



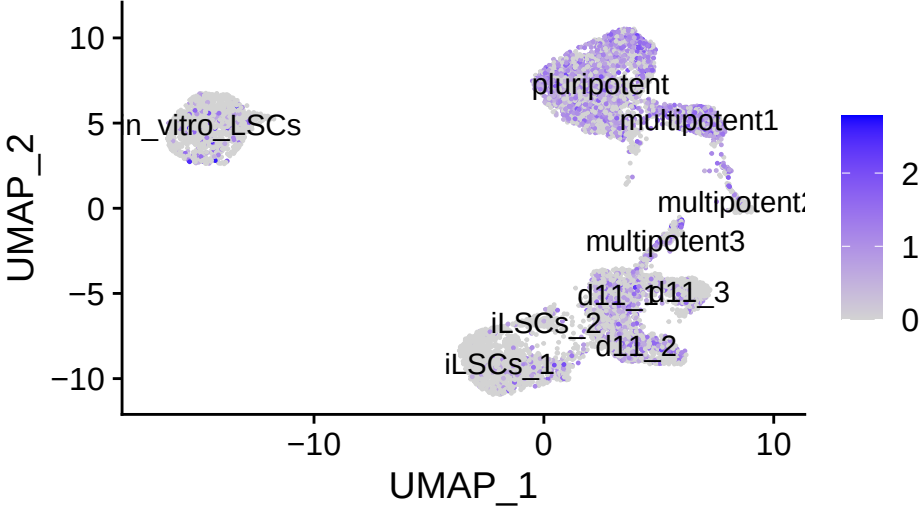
BICD1



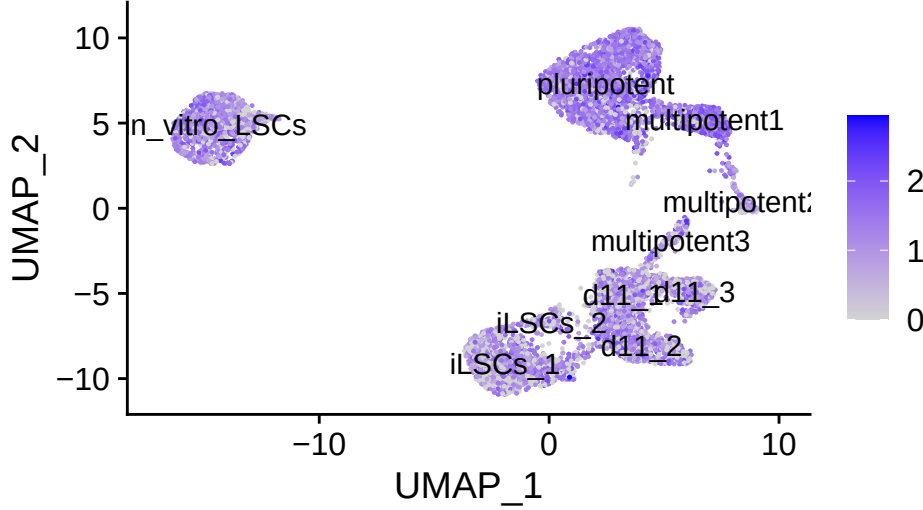
LRPPRC



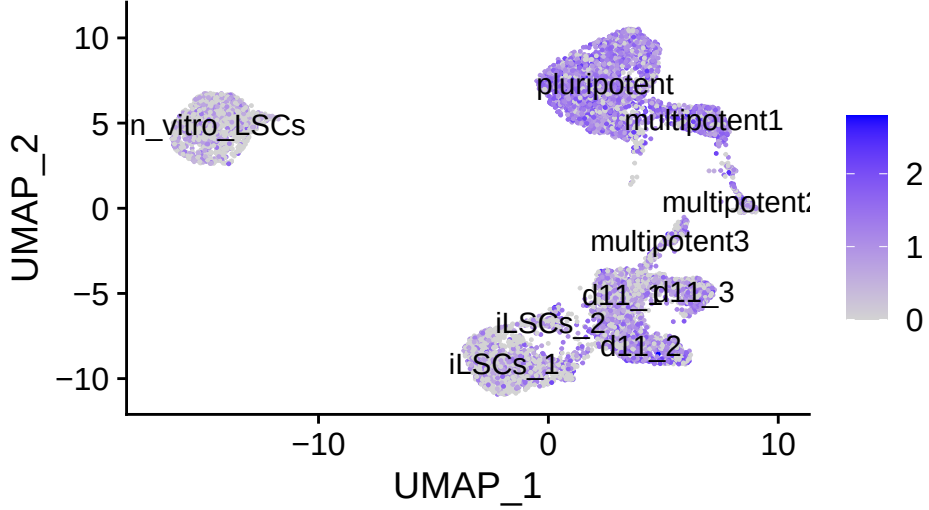
KIF5C



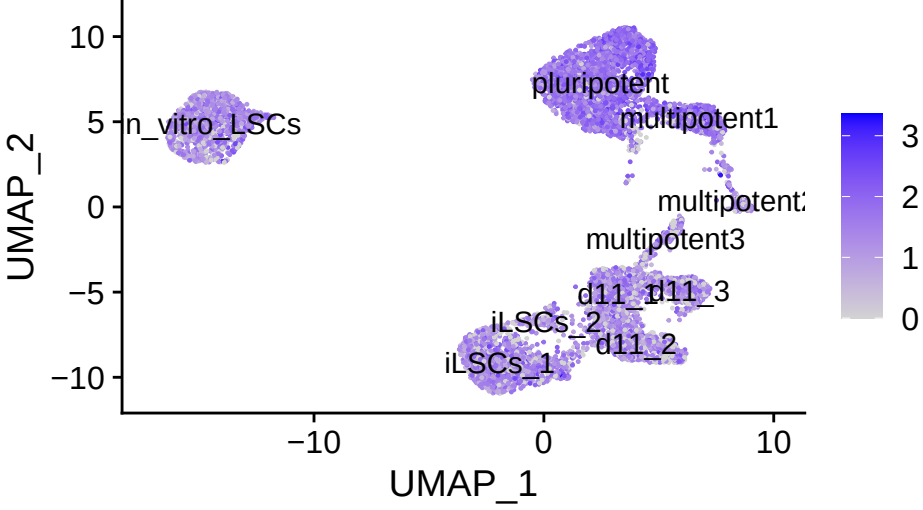
NOP58



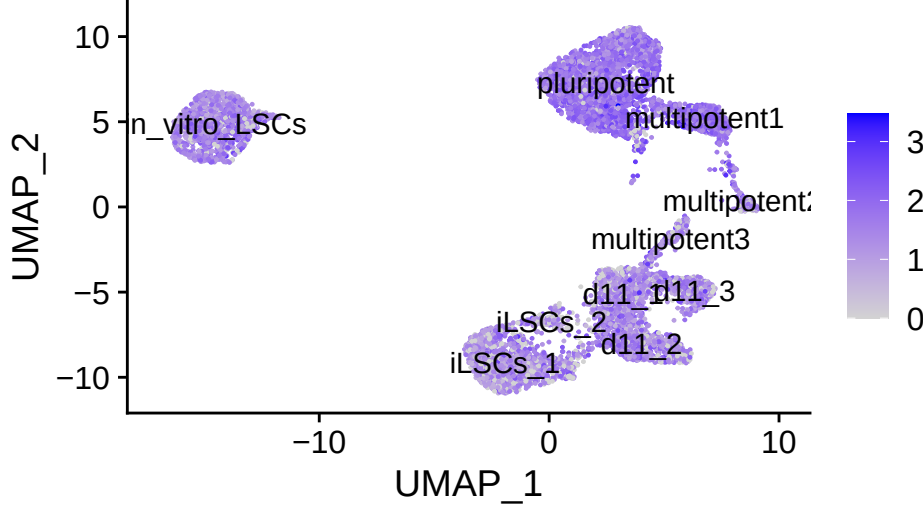
DCP2



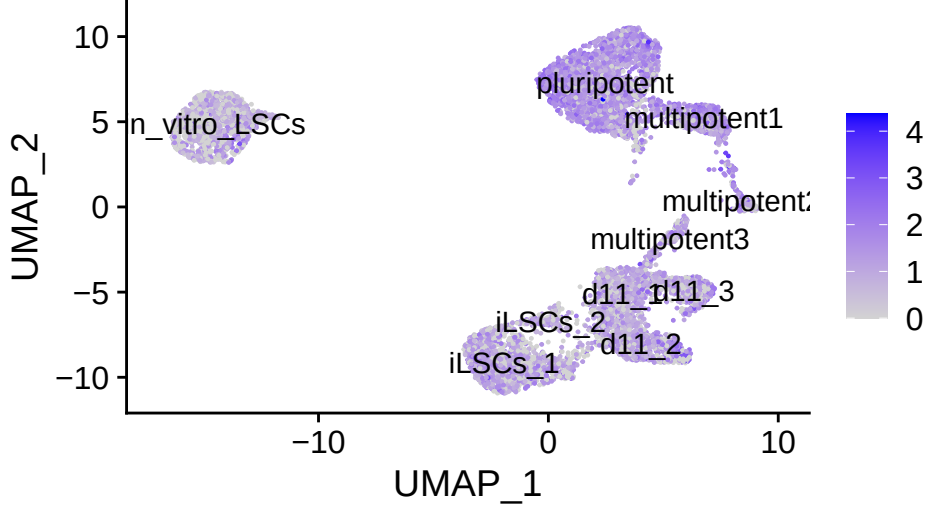
G3BP2



CCT3

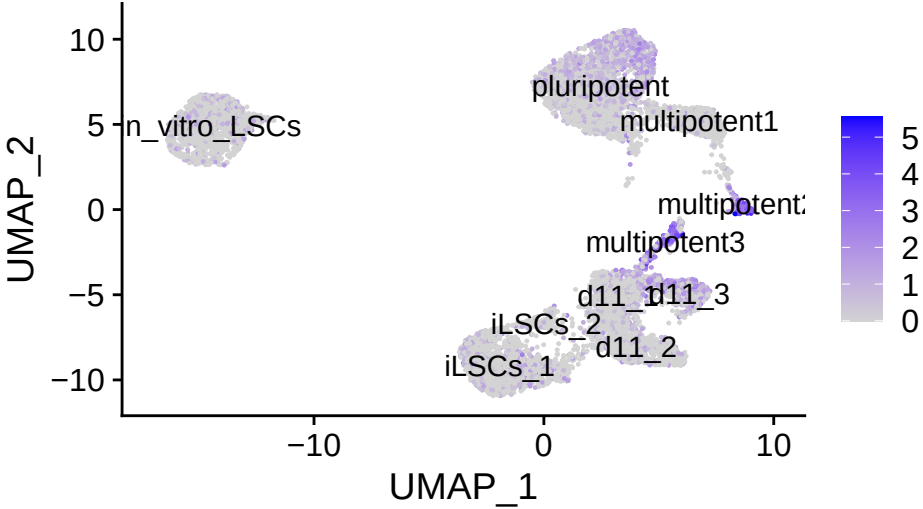


RBFOX1

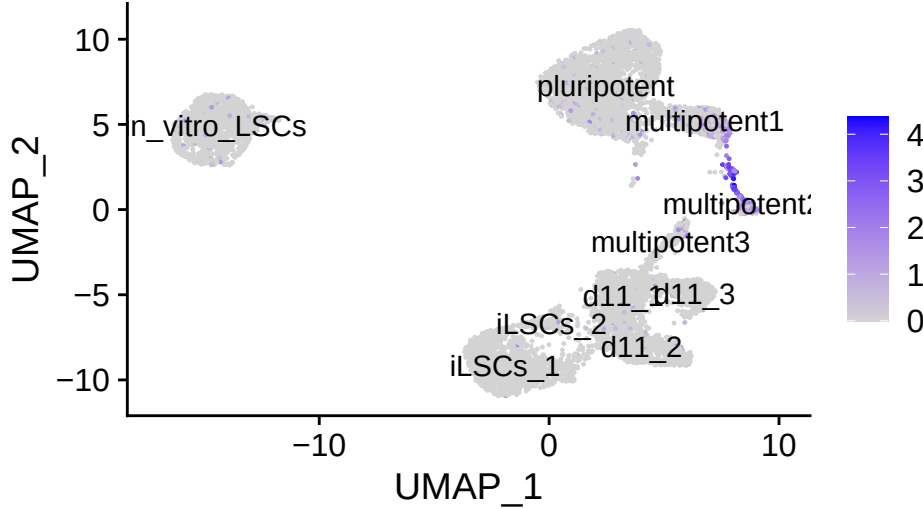


enriched go term: mesenchyme development from cluster: c_multipotent2

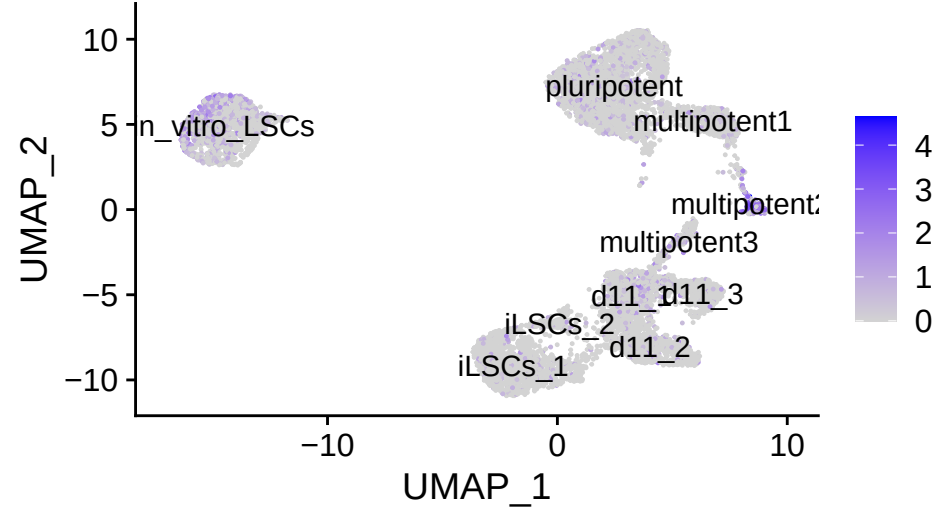
COL1A1



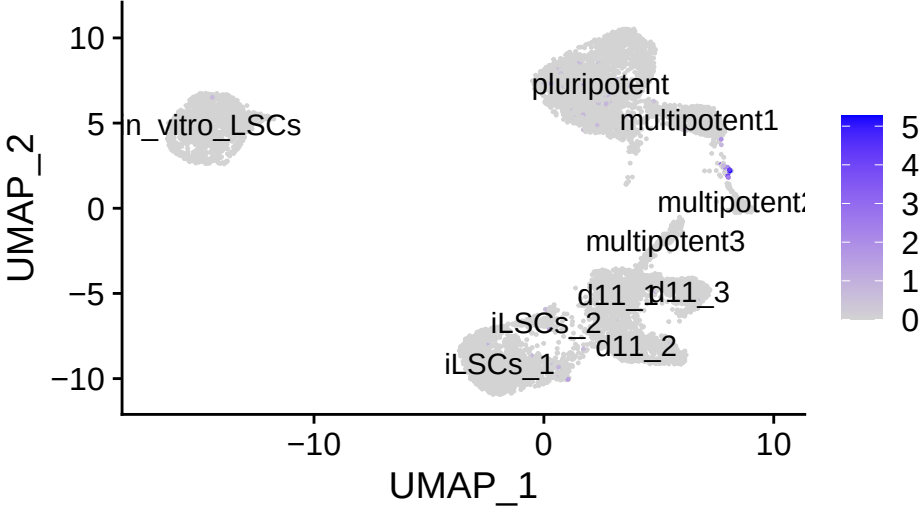
HAS2



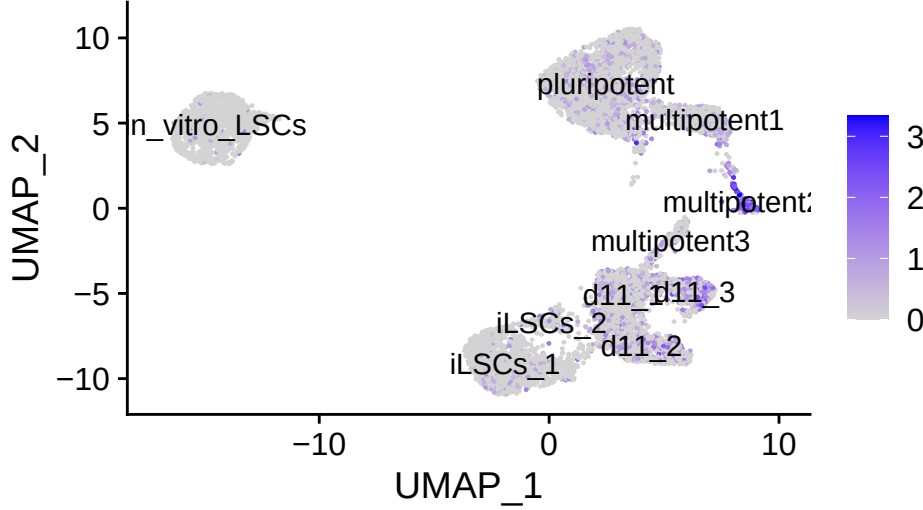
ZFPM2



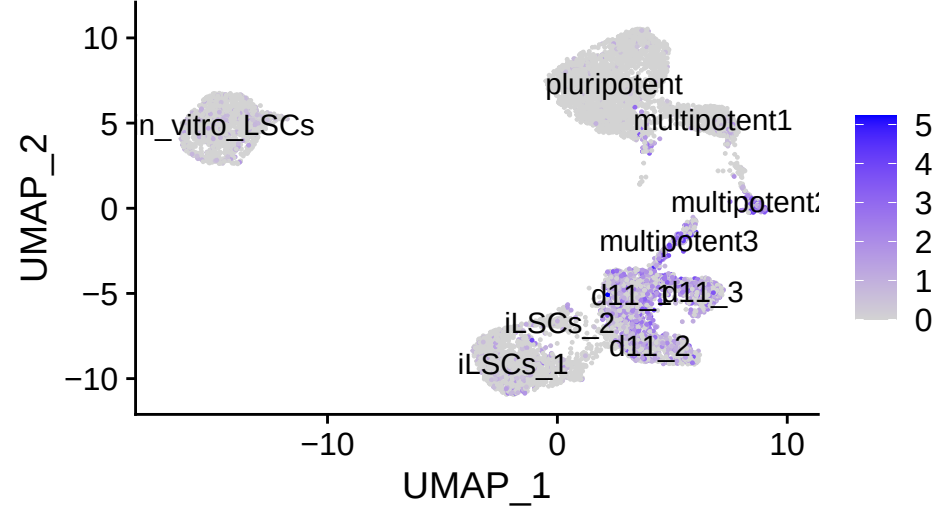
CER1



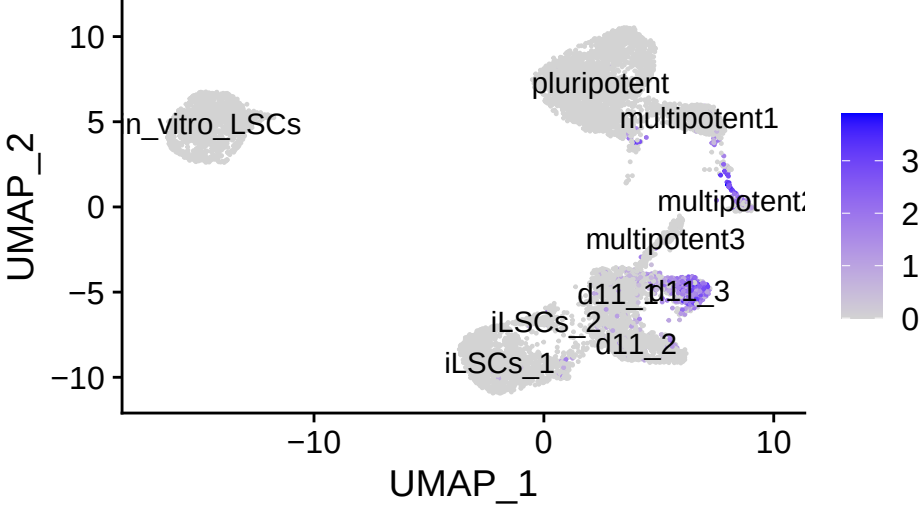
BAMBI



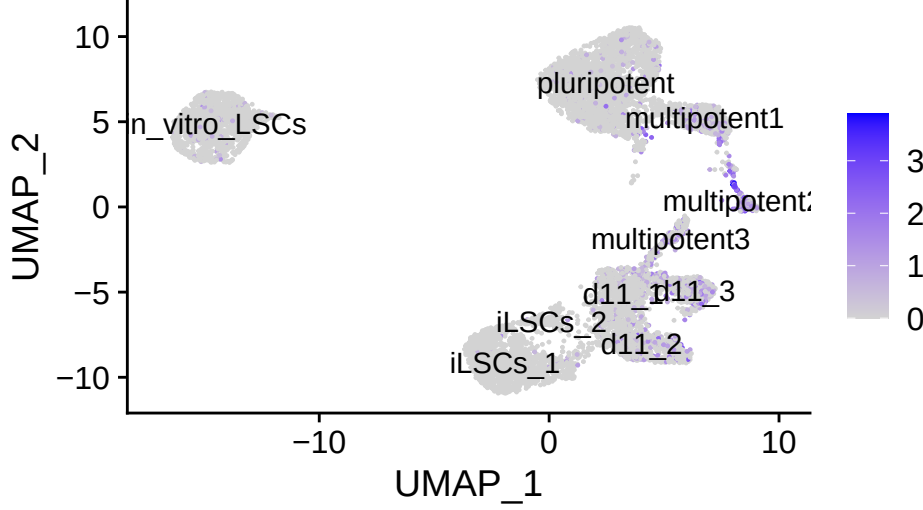
ACTA2



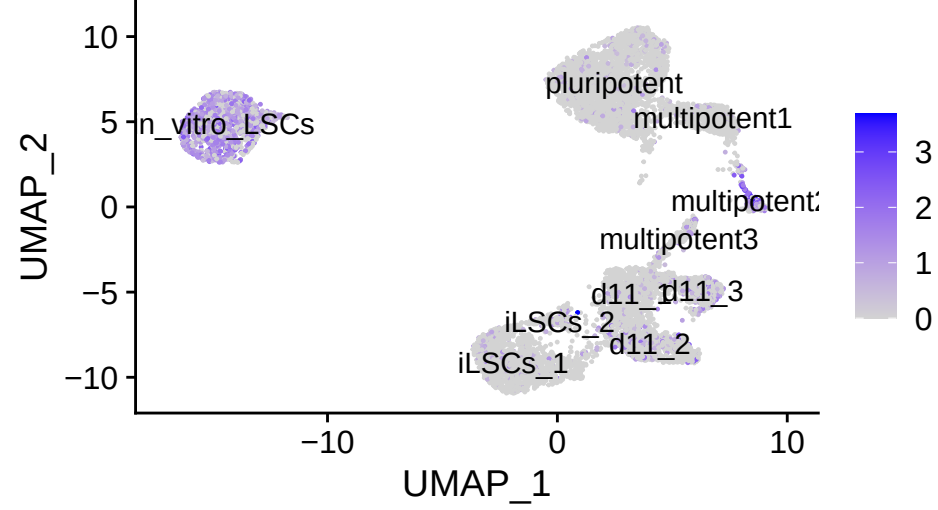
HAND1



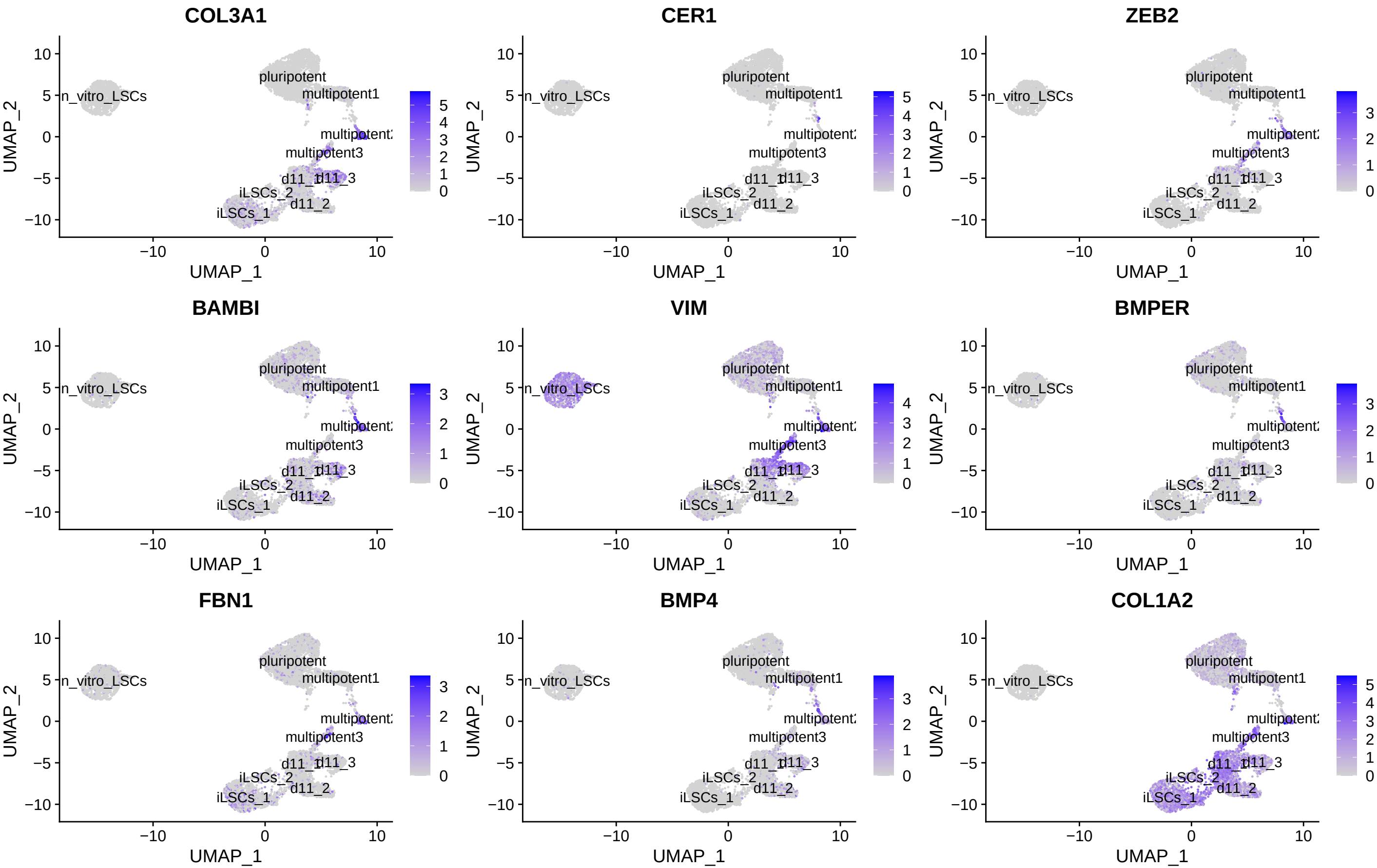
BMP4



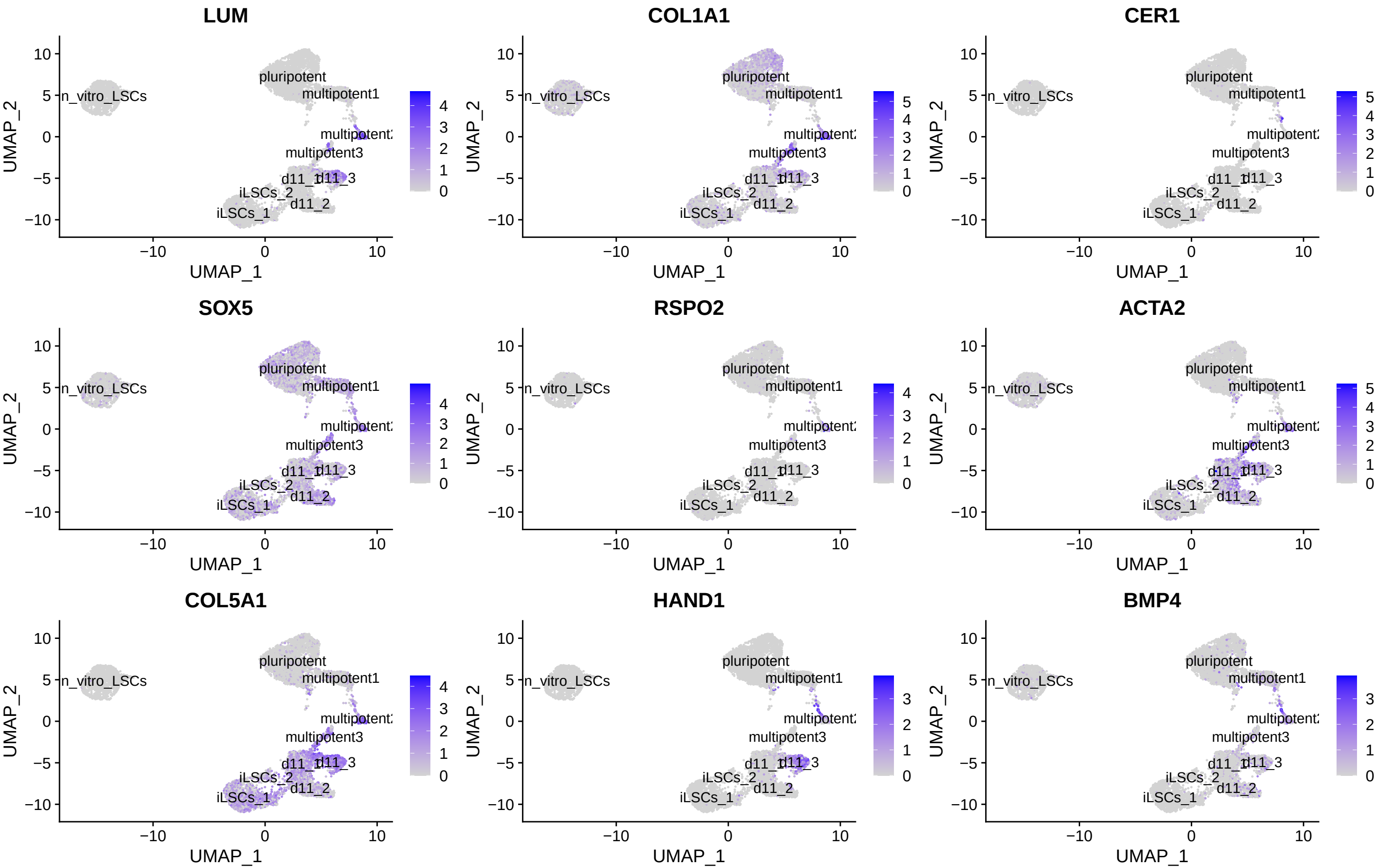
NRP1



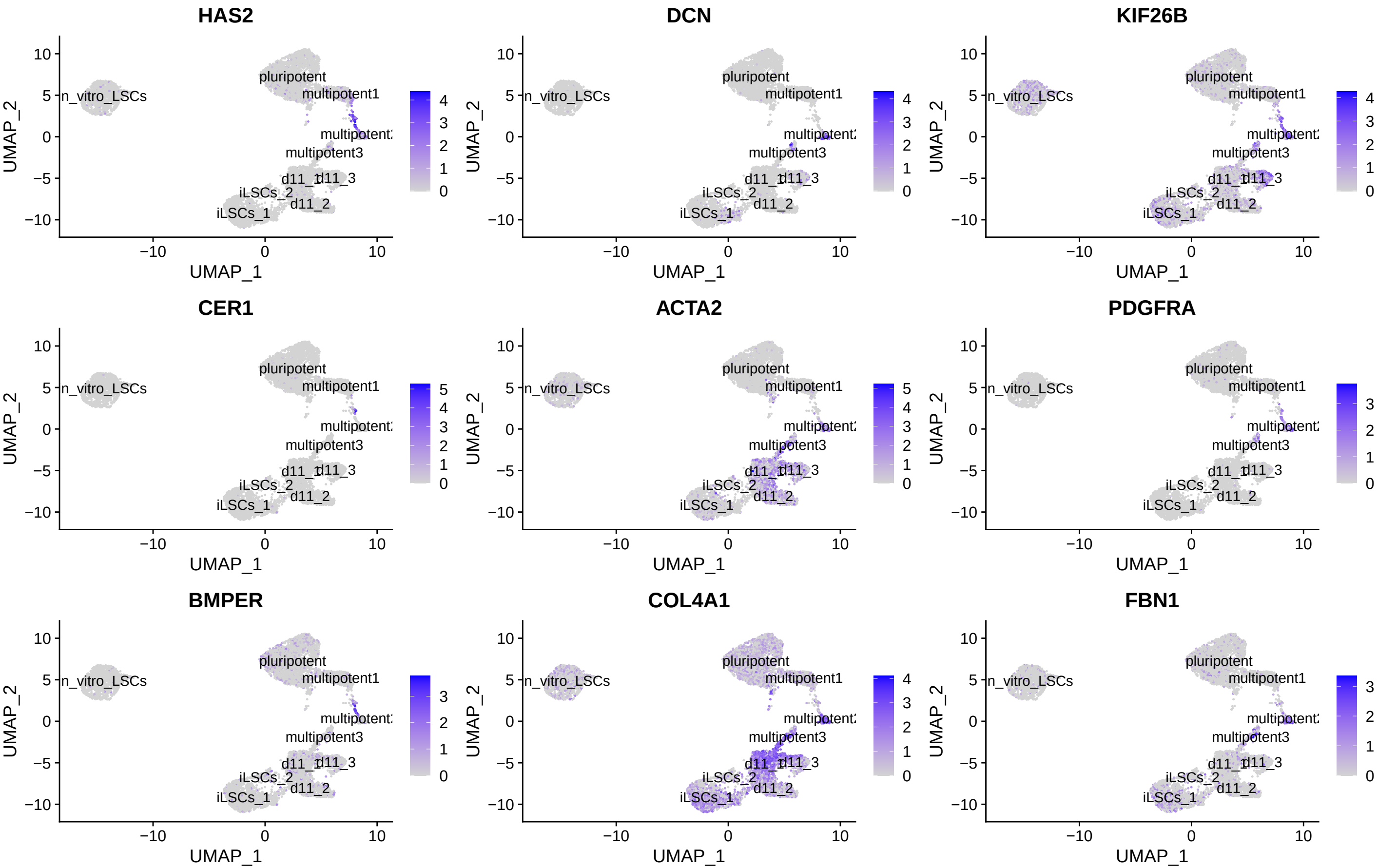
enriched go term: transmembrane receptor protein serine/threonine kinase signaling pathway from cluster: c_multipotent2



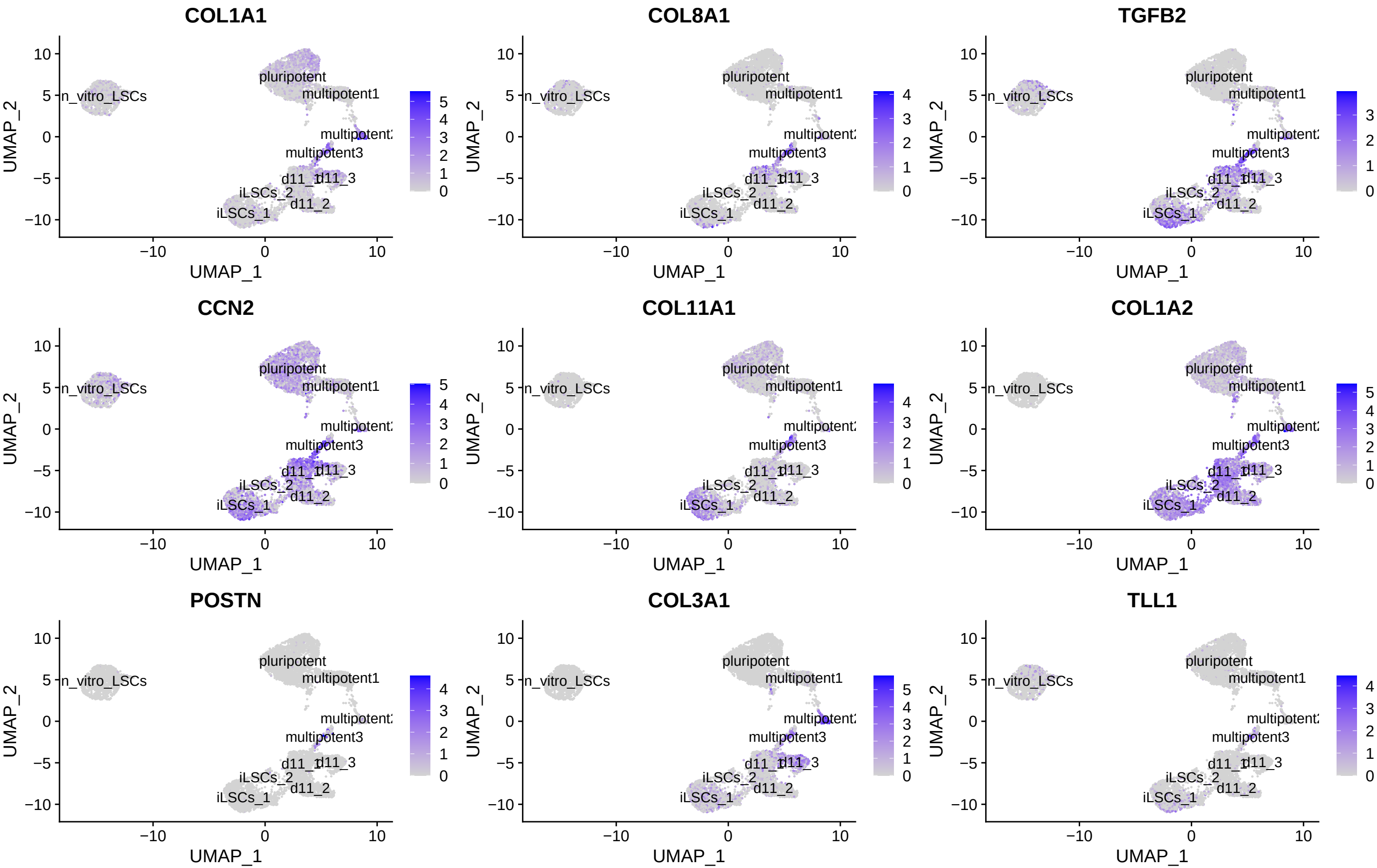
enriched go term: connective tissue development from cluster: c_multipotent2



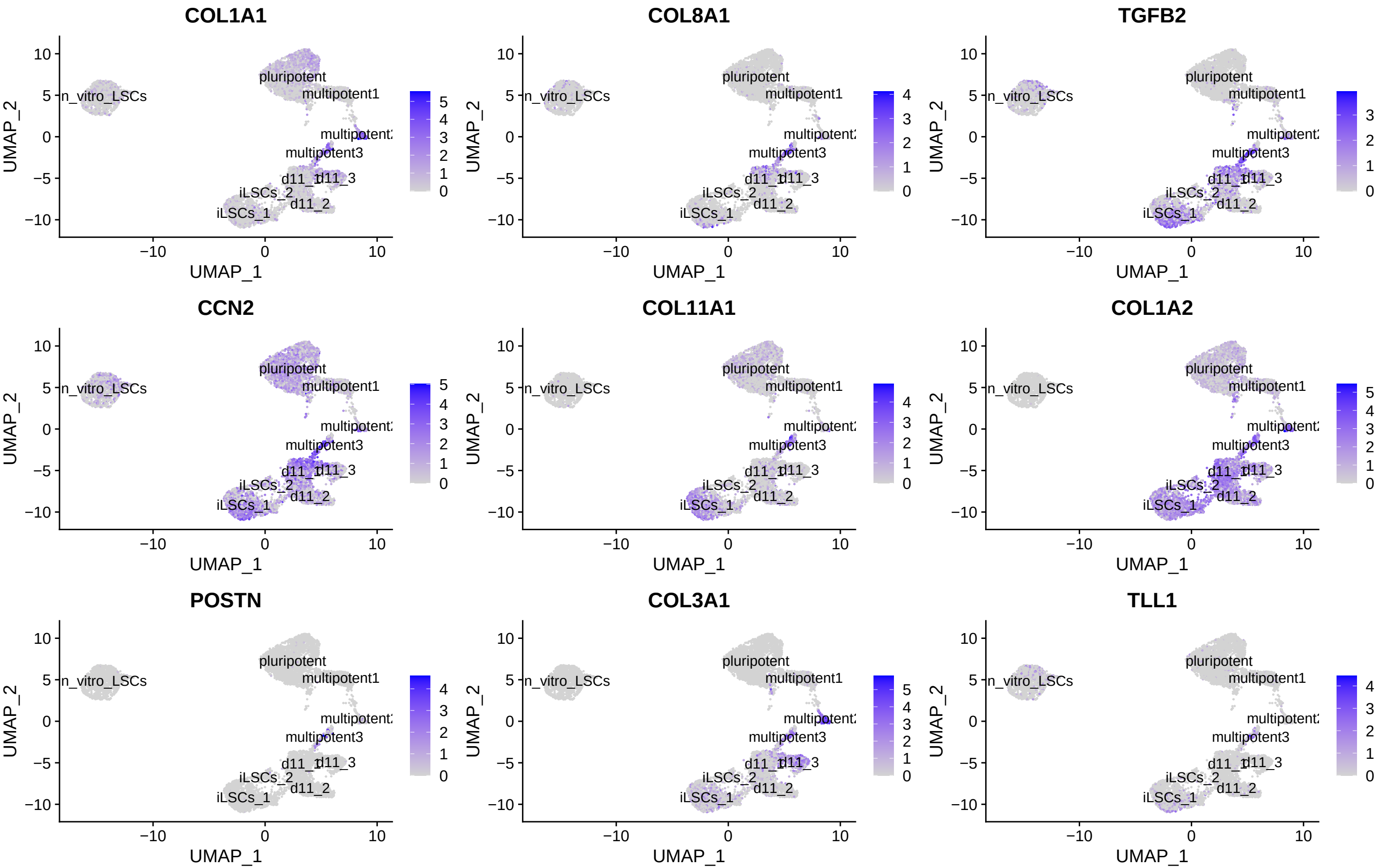
enriched go term: renal system development from cluster: c_multipotent2



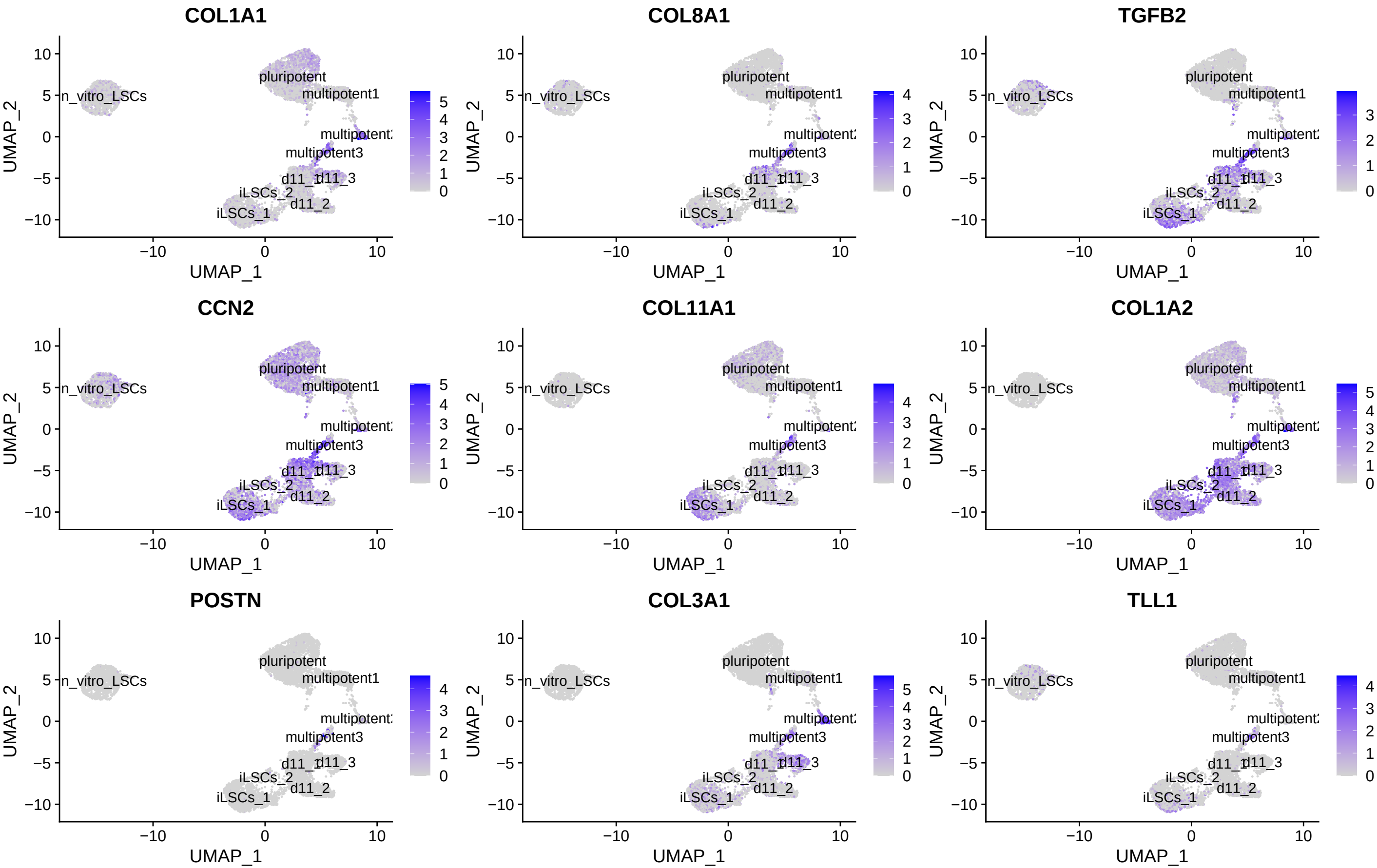
enriched go term: extracellular matrix organization from cluster: d_multipotent3



enriched go term: extracellular structure organization from cluster: d_multipotent3

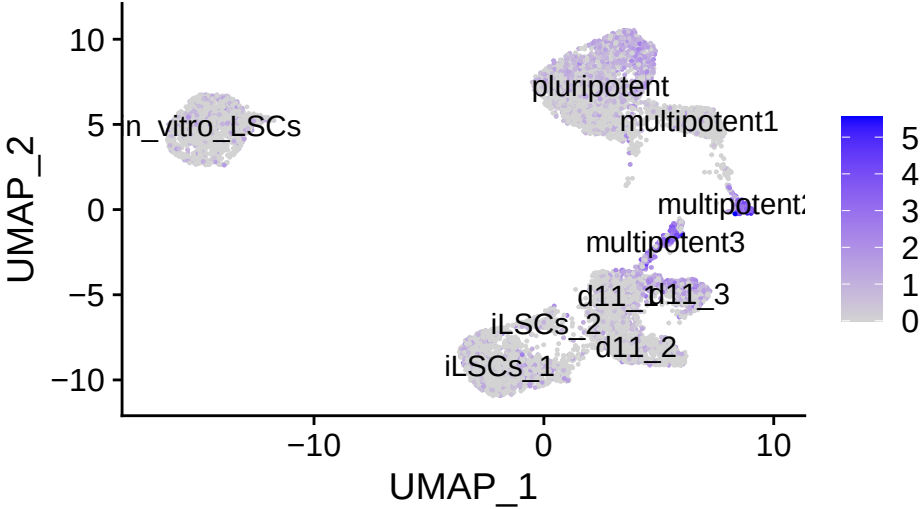


enriched go term: external encapsulating structure organization from cluster: d_multipotent3

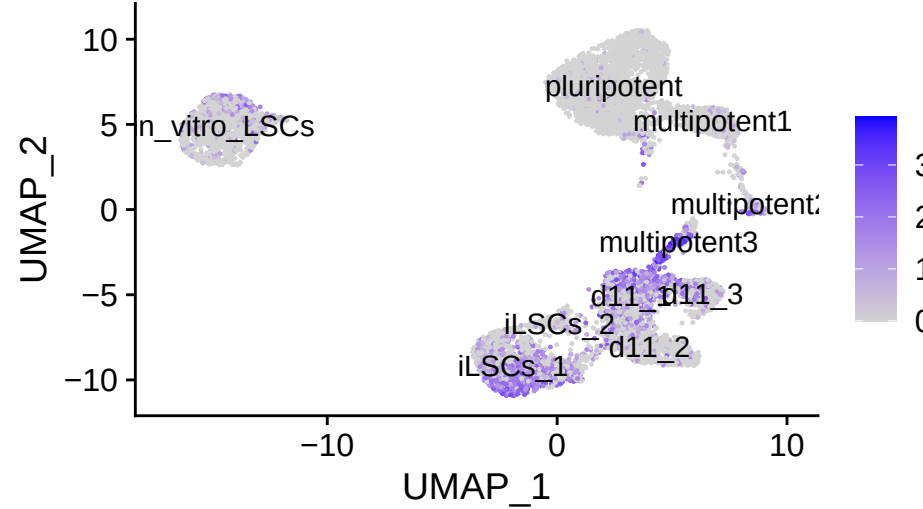


enriched go term: mesenchyme development from cluster: d_multipotent3

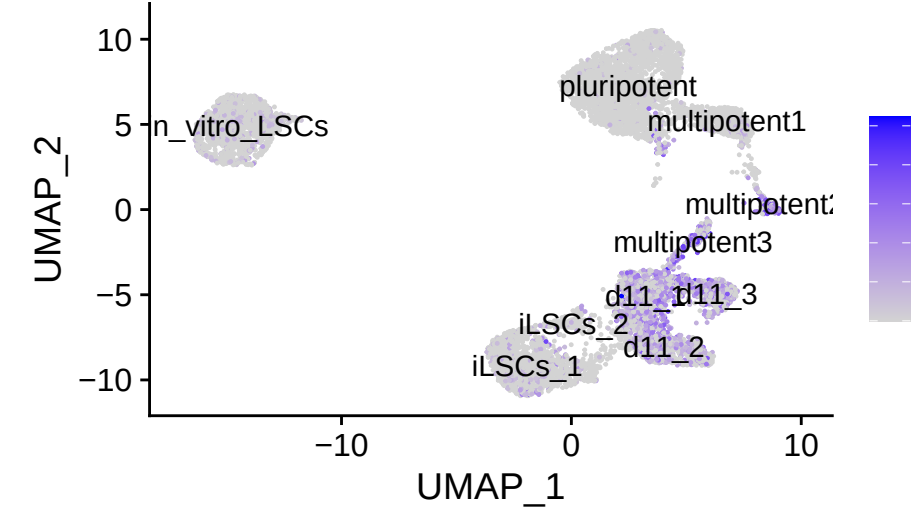
COL1A1



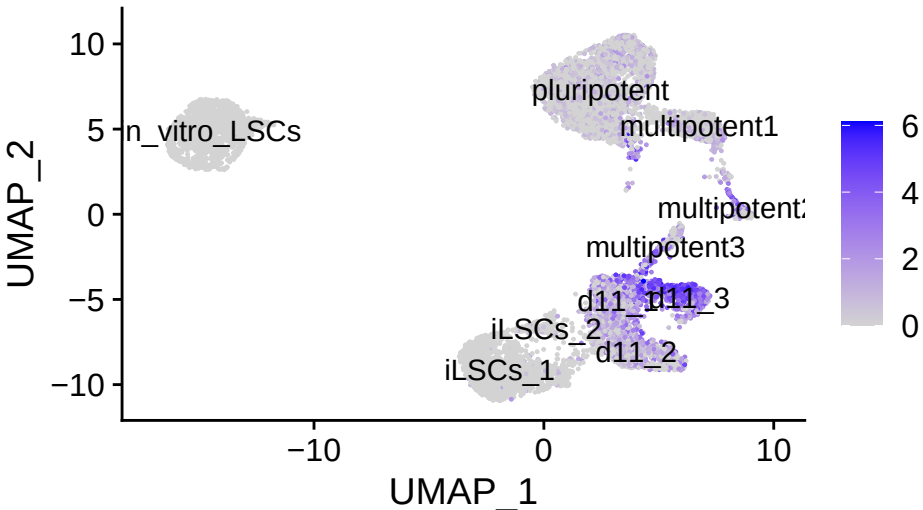
TGFB2



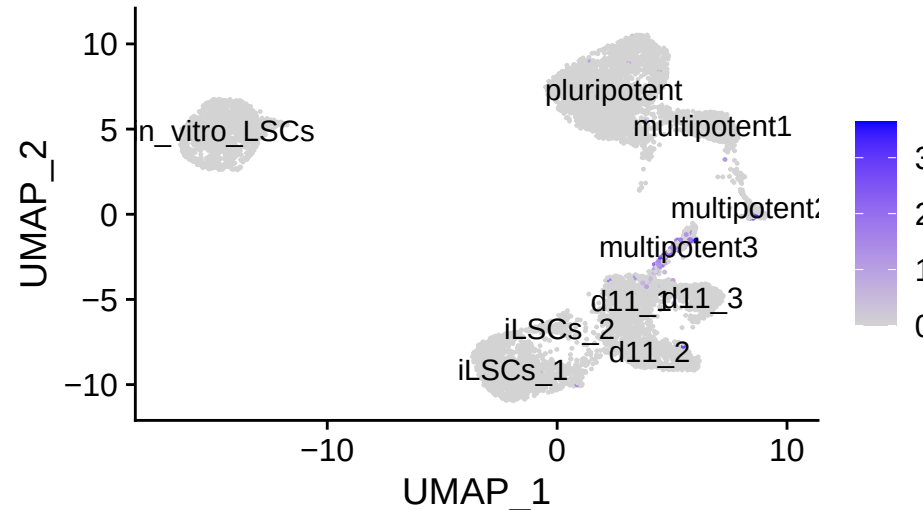
ACTA2



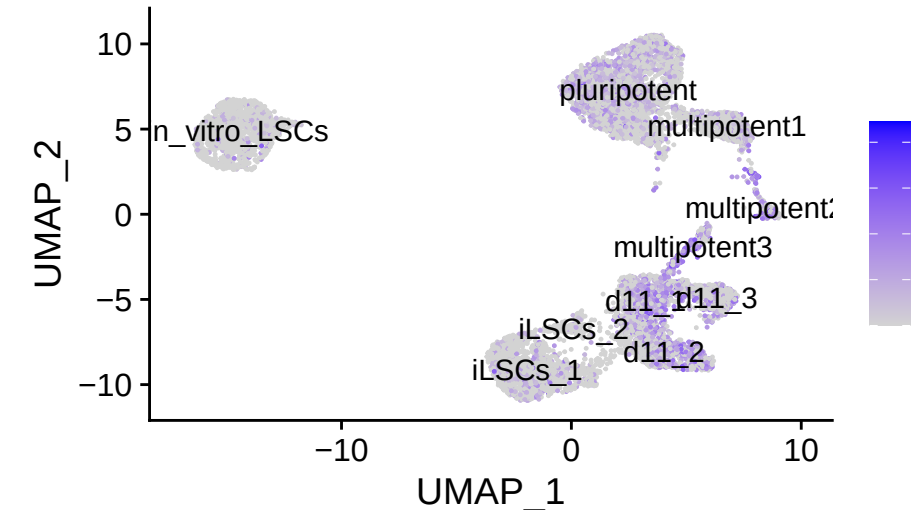
ACTC1



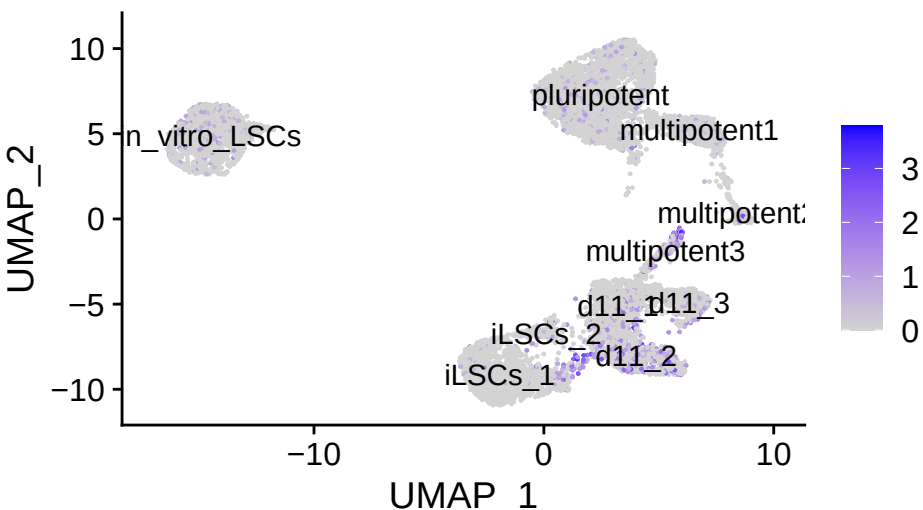
GREM1



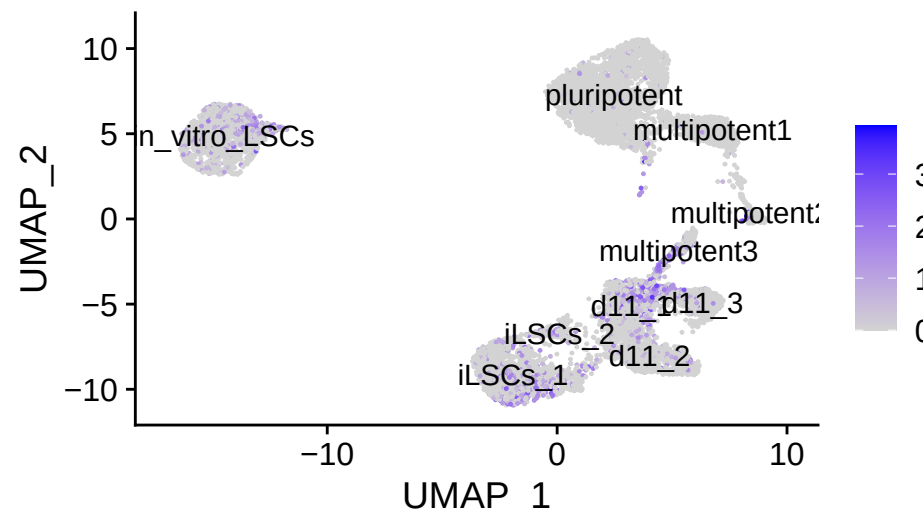
CDH2



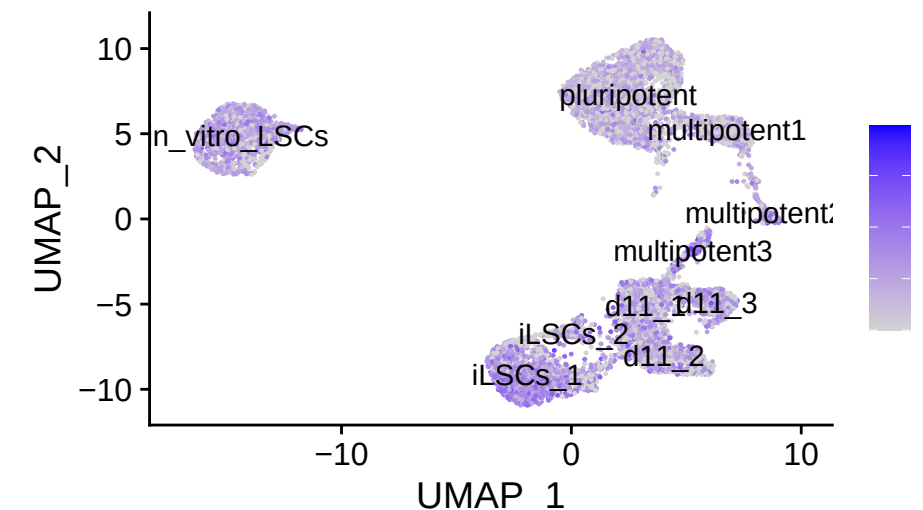
EDNRA



EDN1

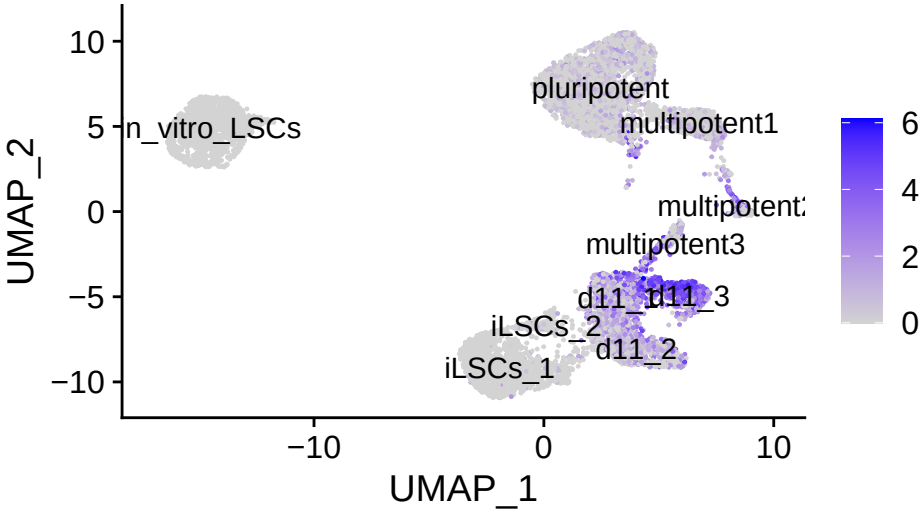


TGFBR1

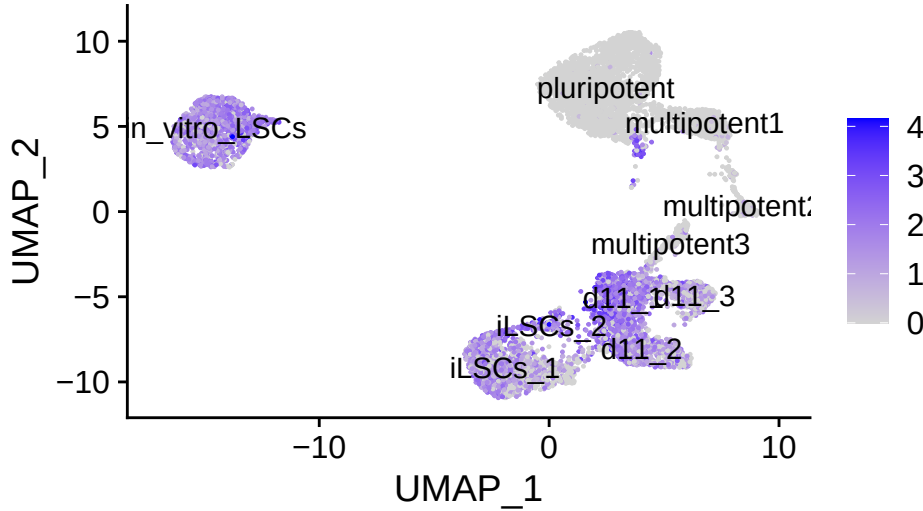


enriched go term: actin filament organization from cluster: e_d11_1

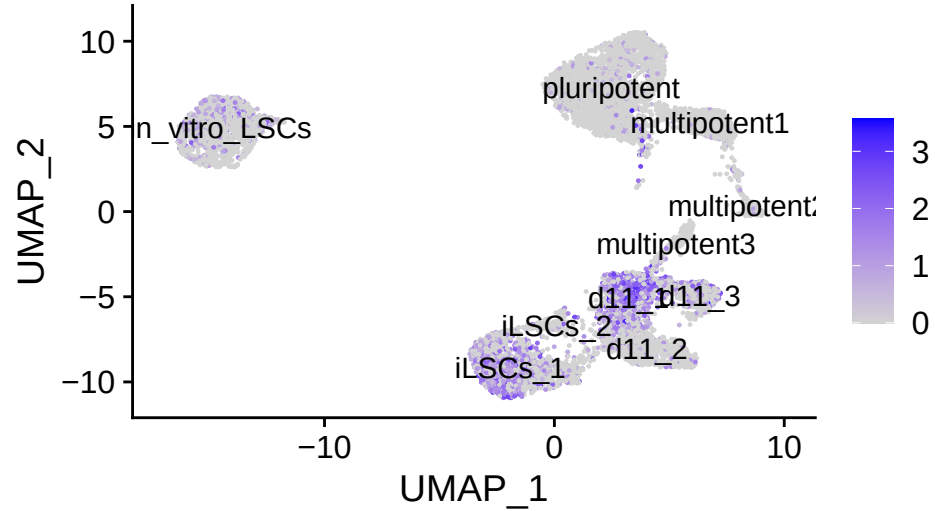
ACTC1



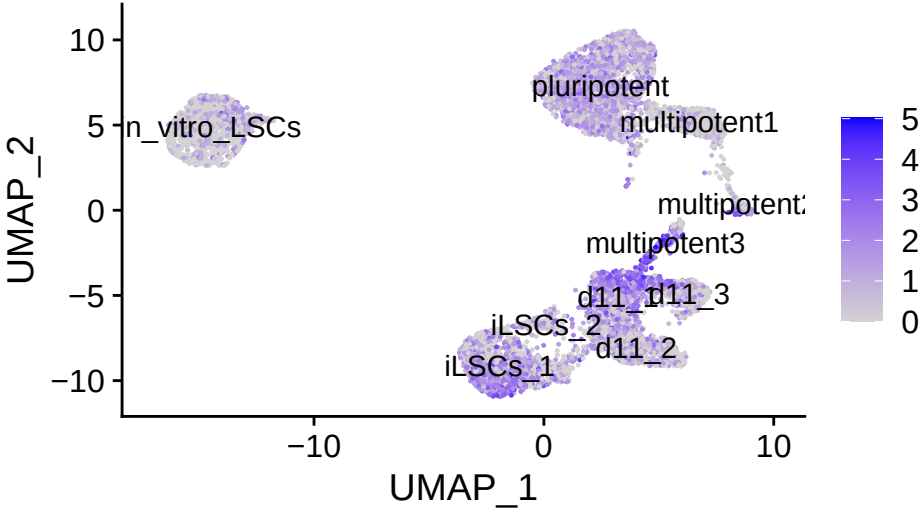
TACSTD2



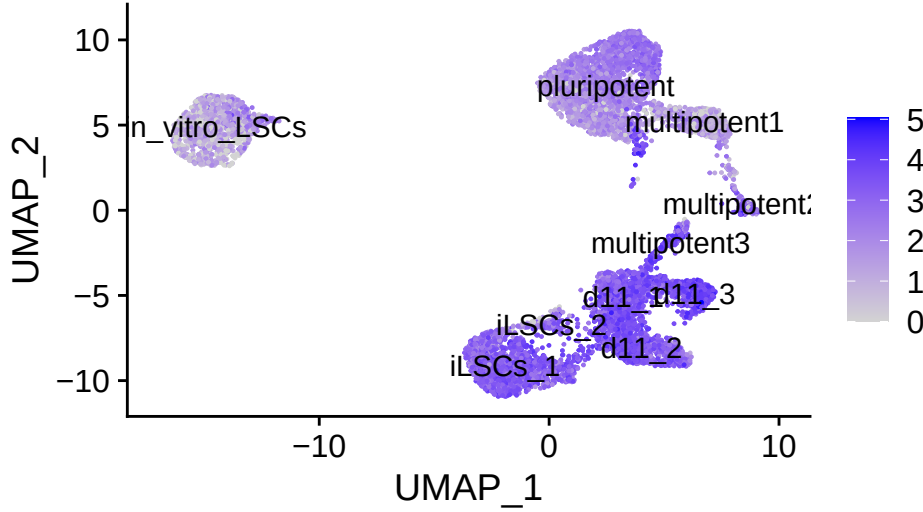
LCP1



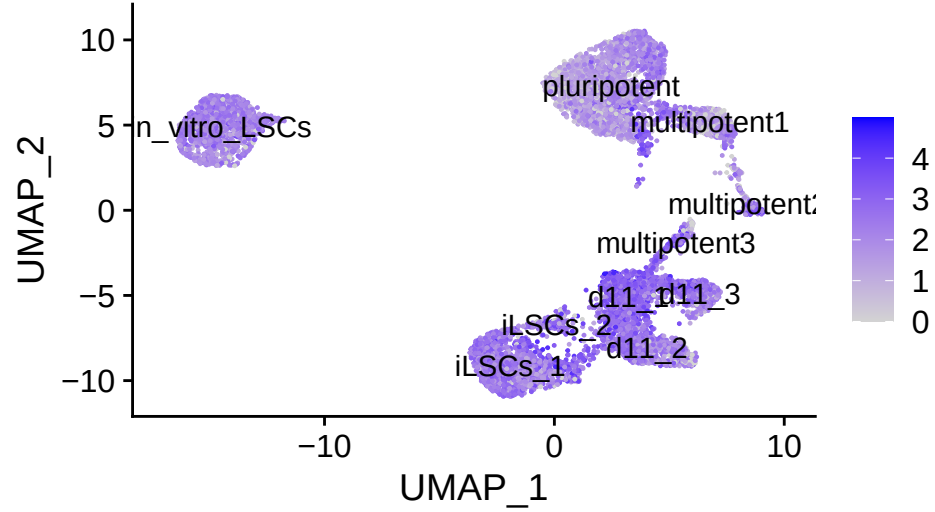
CCN2



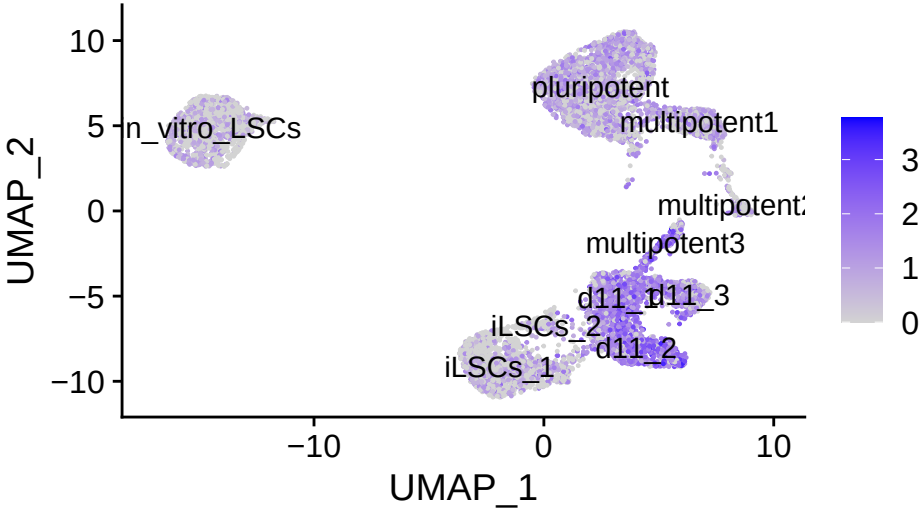
TPM1



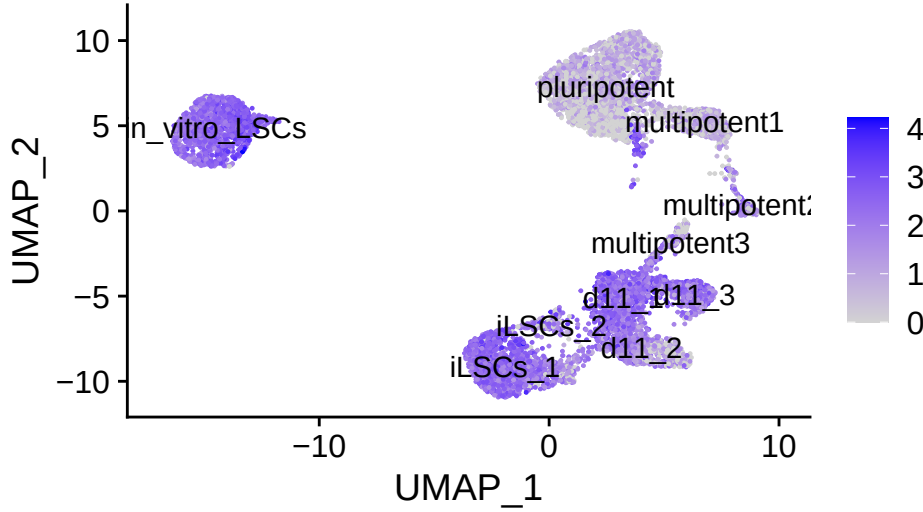
TMSB4X



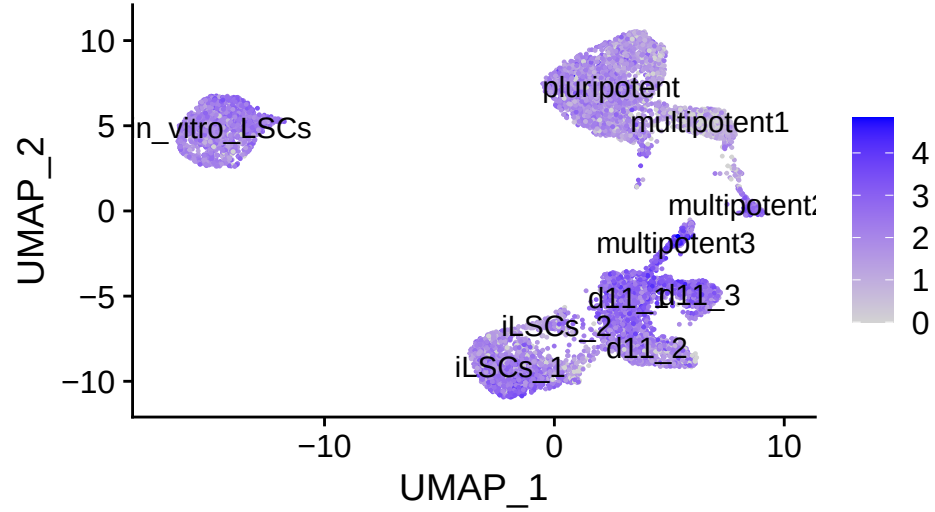
SHROOM3



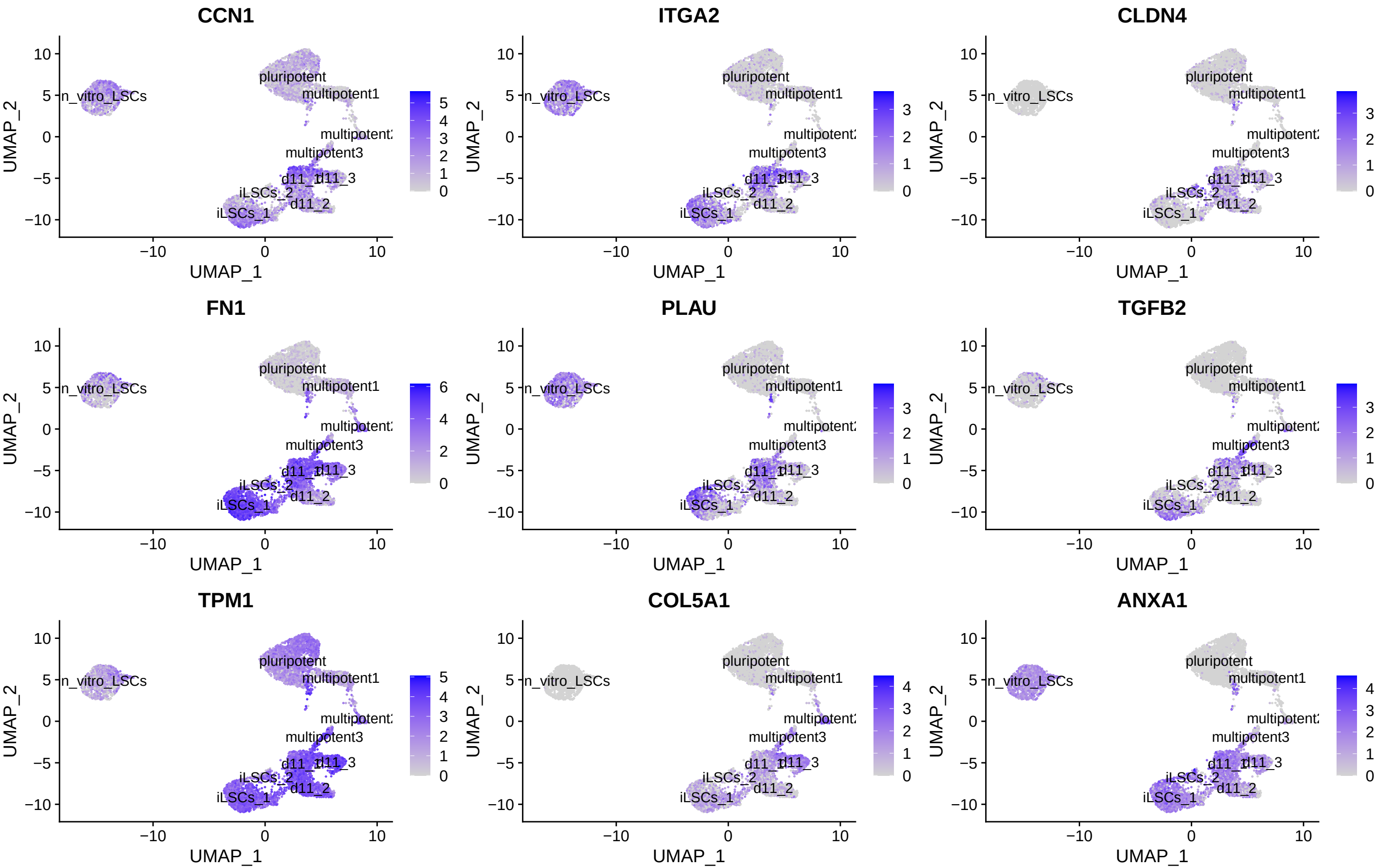
S100A10



CALD1

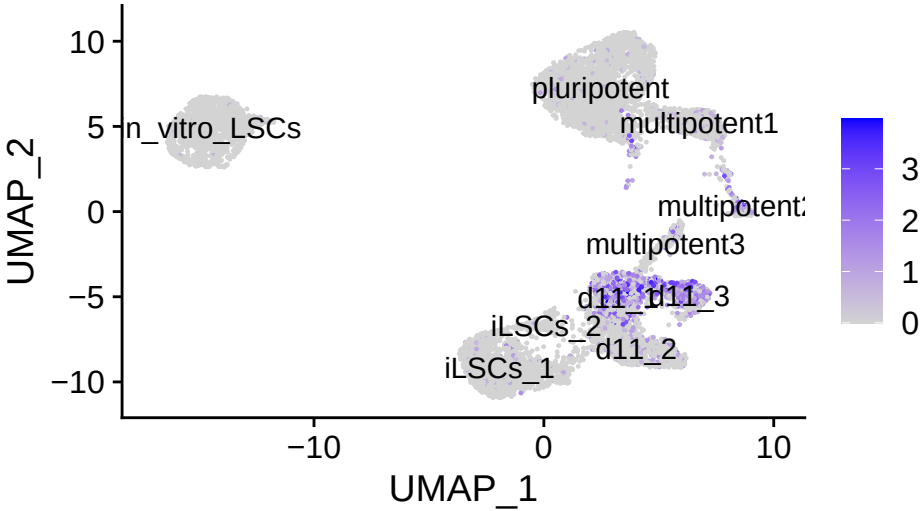


enriched go term: wound healing from cluster: e_d11_1

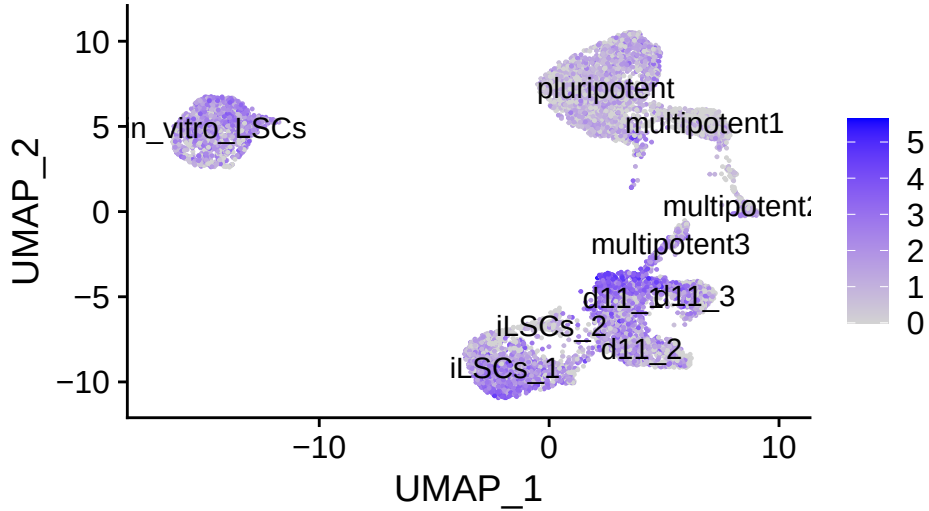


enriched go term: cell-substrate adhesion from cluster: e_d11_1

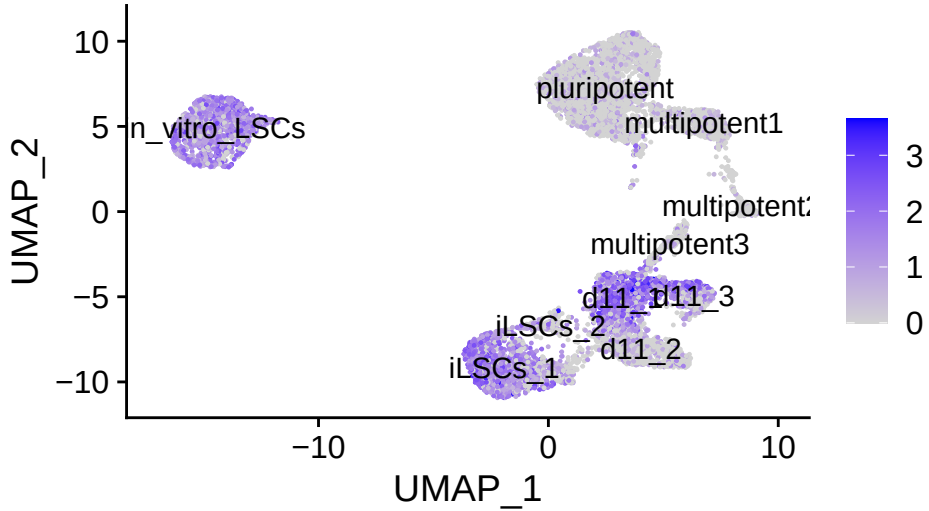
ADAMTS9



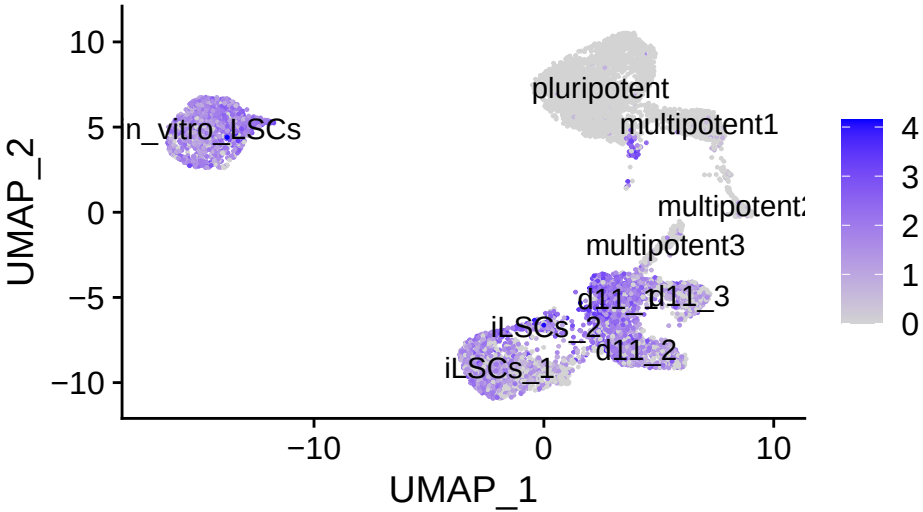
CCN1



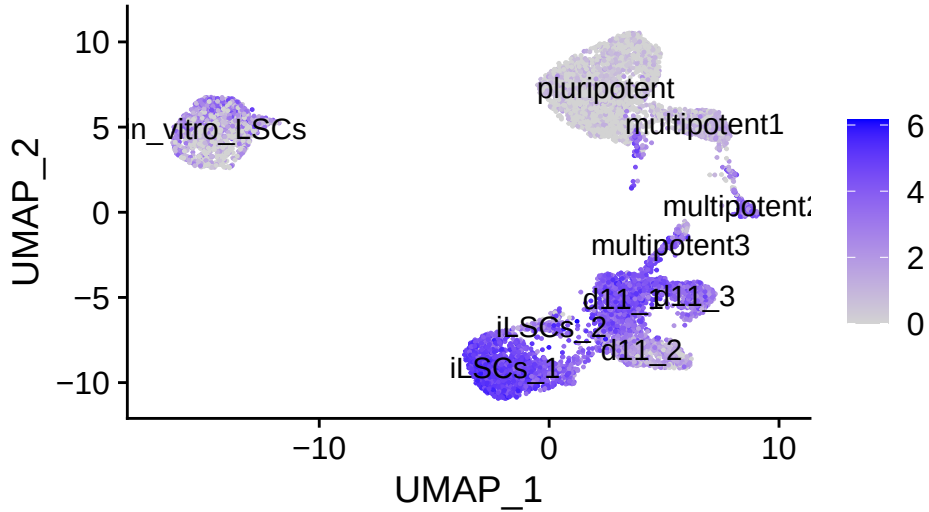
ITGA2



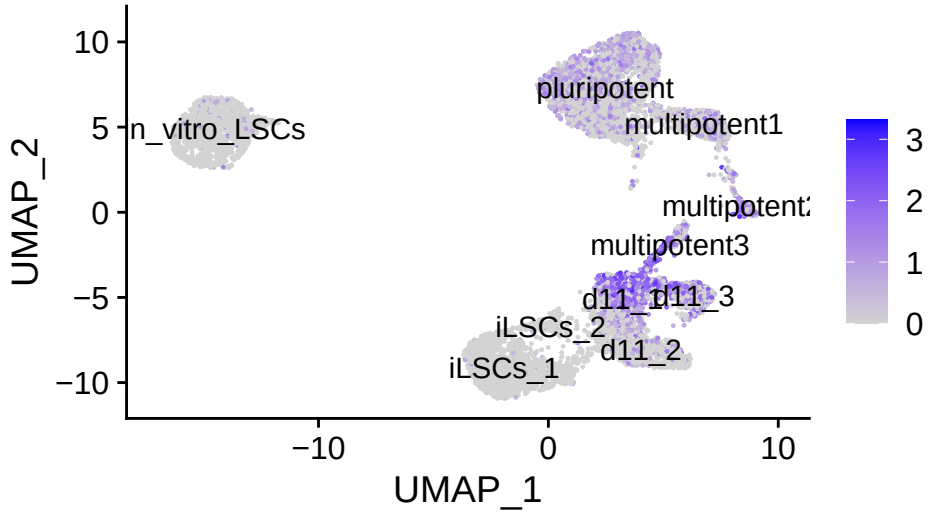
TACSTD2



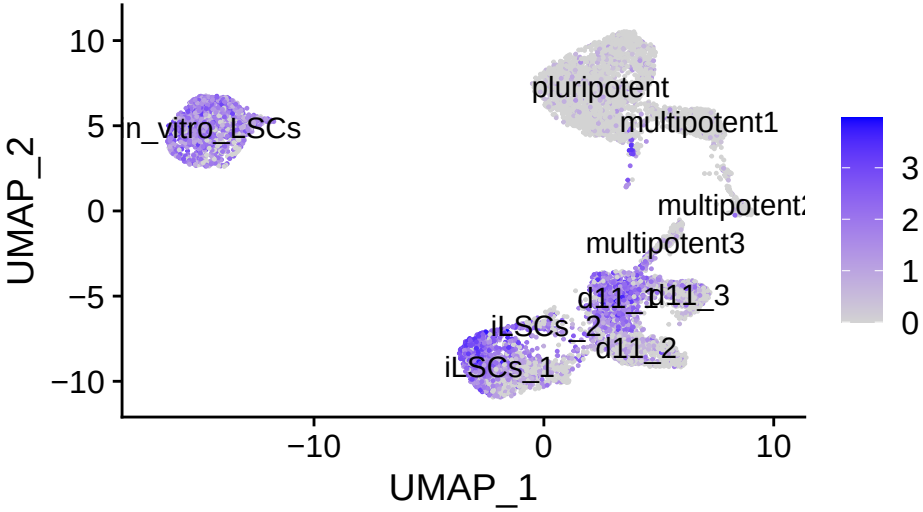
FN1



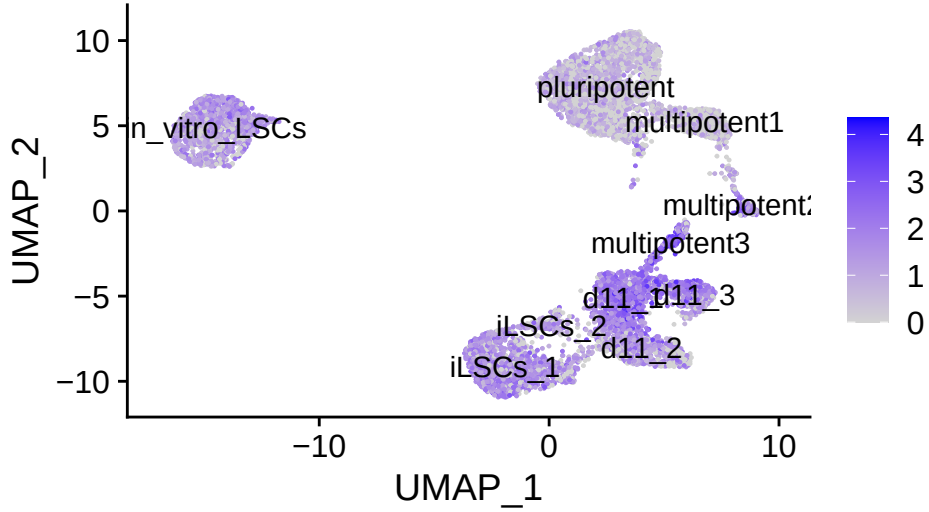
ADAMTS12



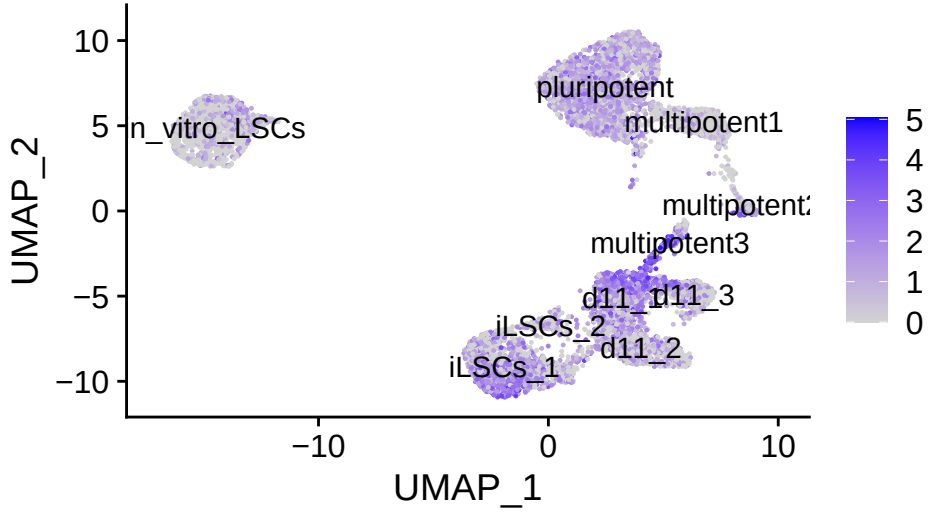
PLAU



ITGAV

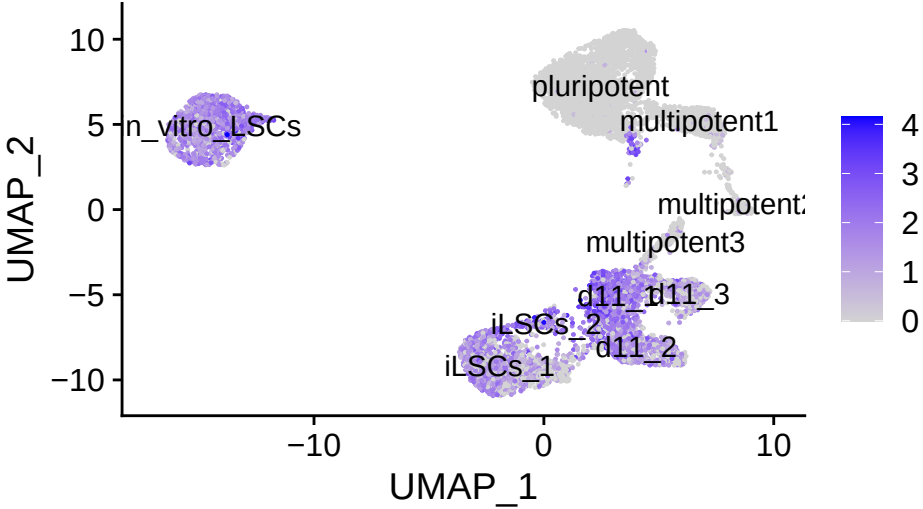


CCN2

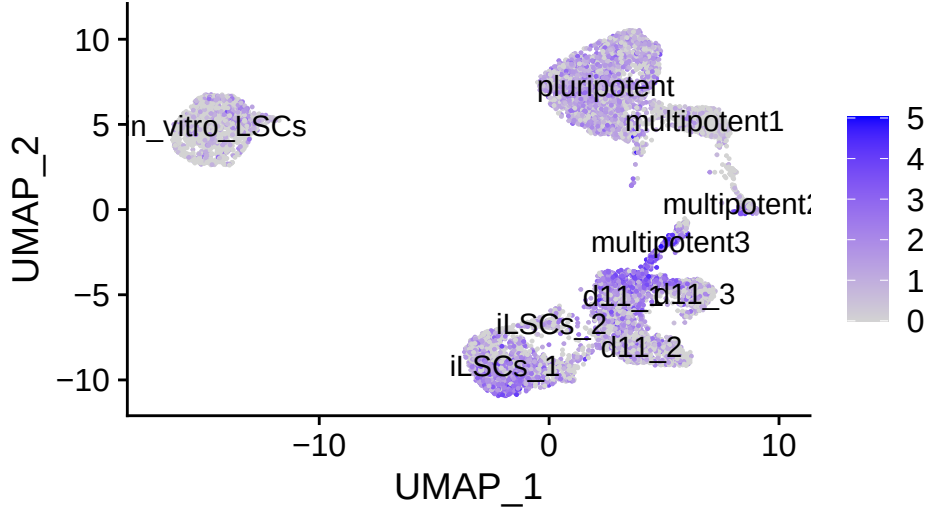


enriched go term: regulation of actin filament-based process from cluster: e_d11_1

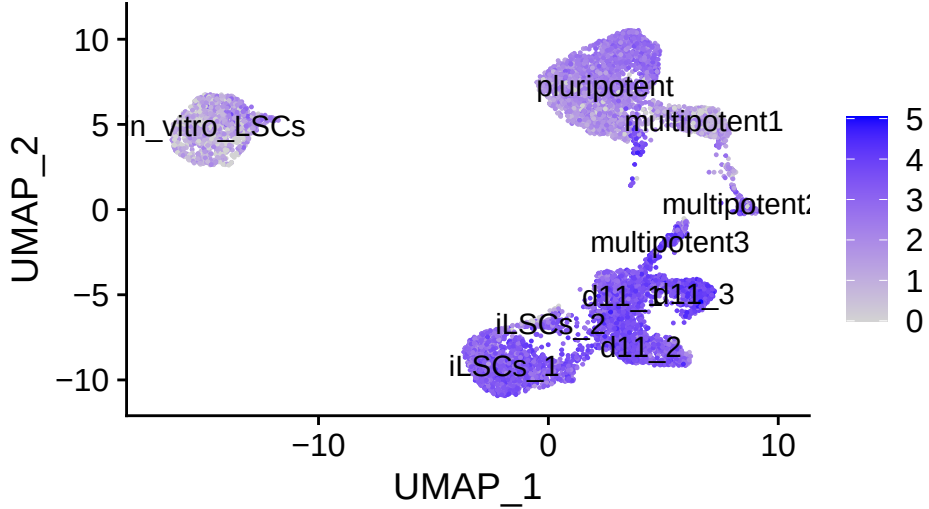
TACSTD2



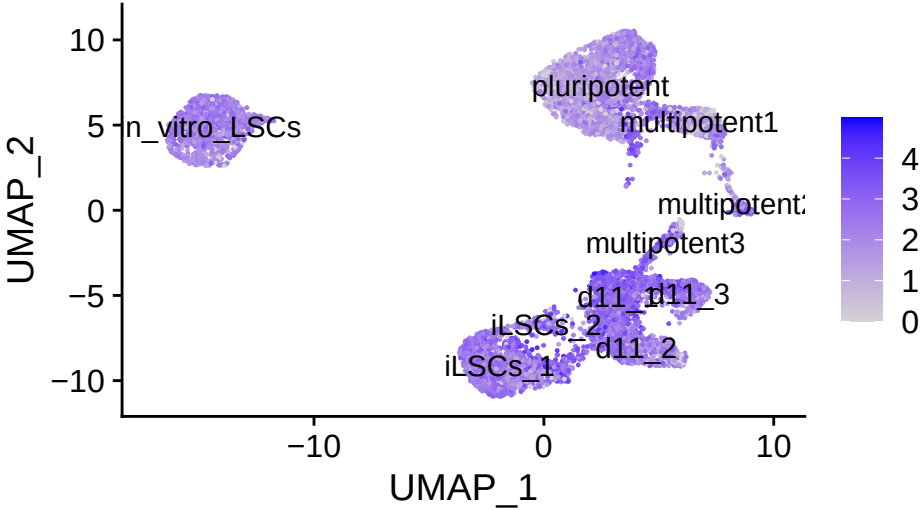
CCN2



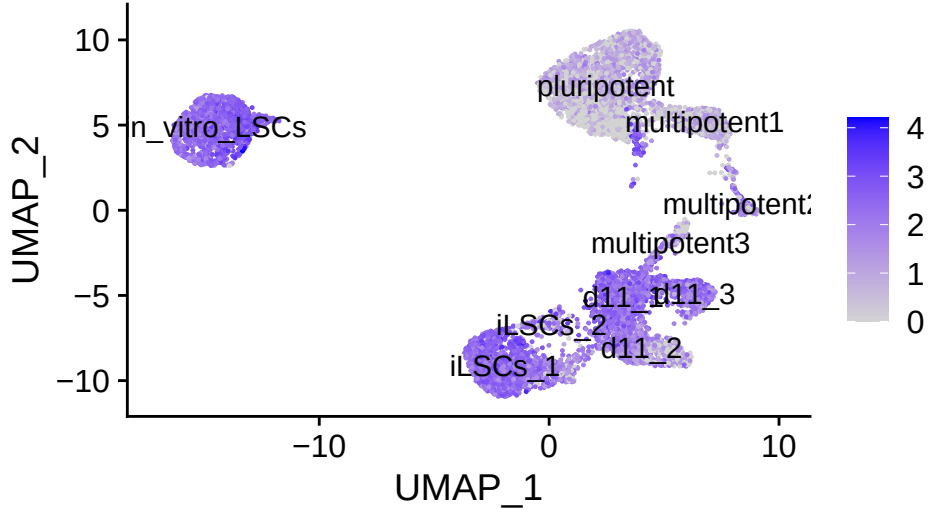
TPM1



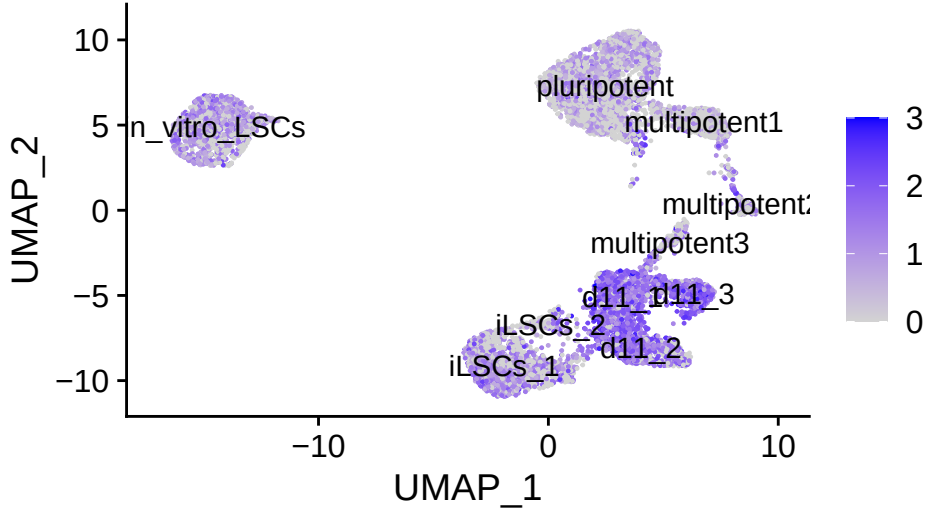
TMSB4X



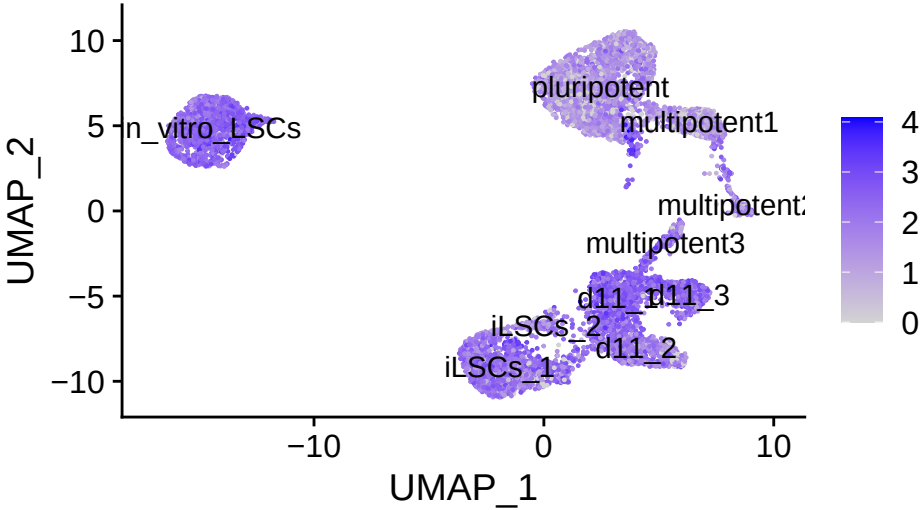
S100A10



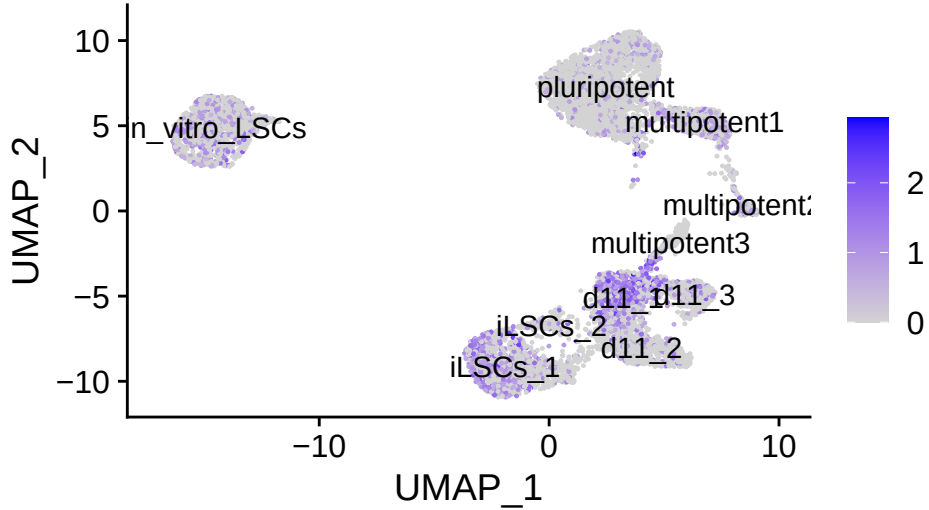
PKP2



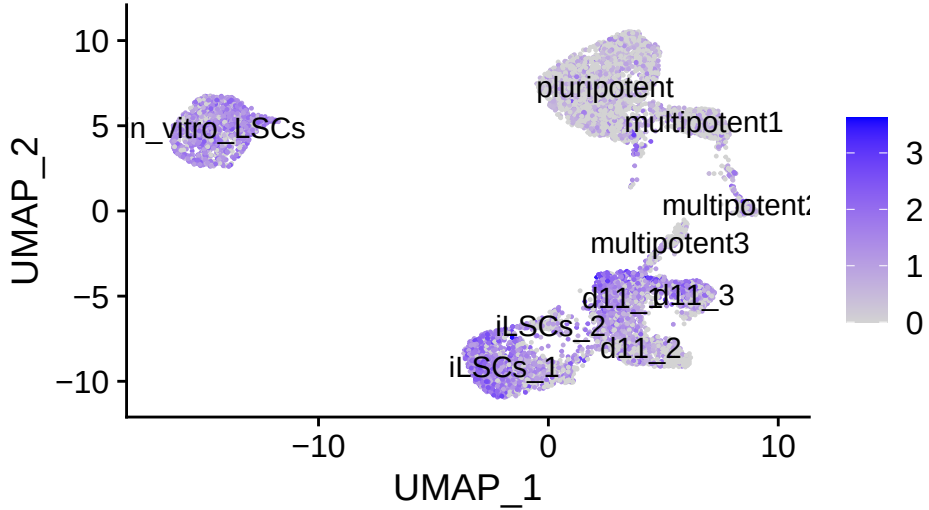
TMSB10



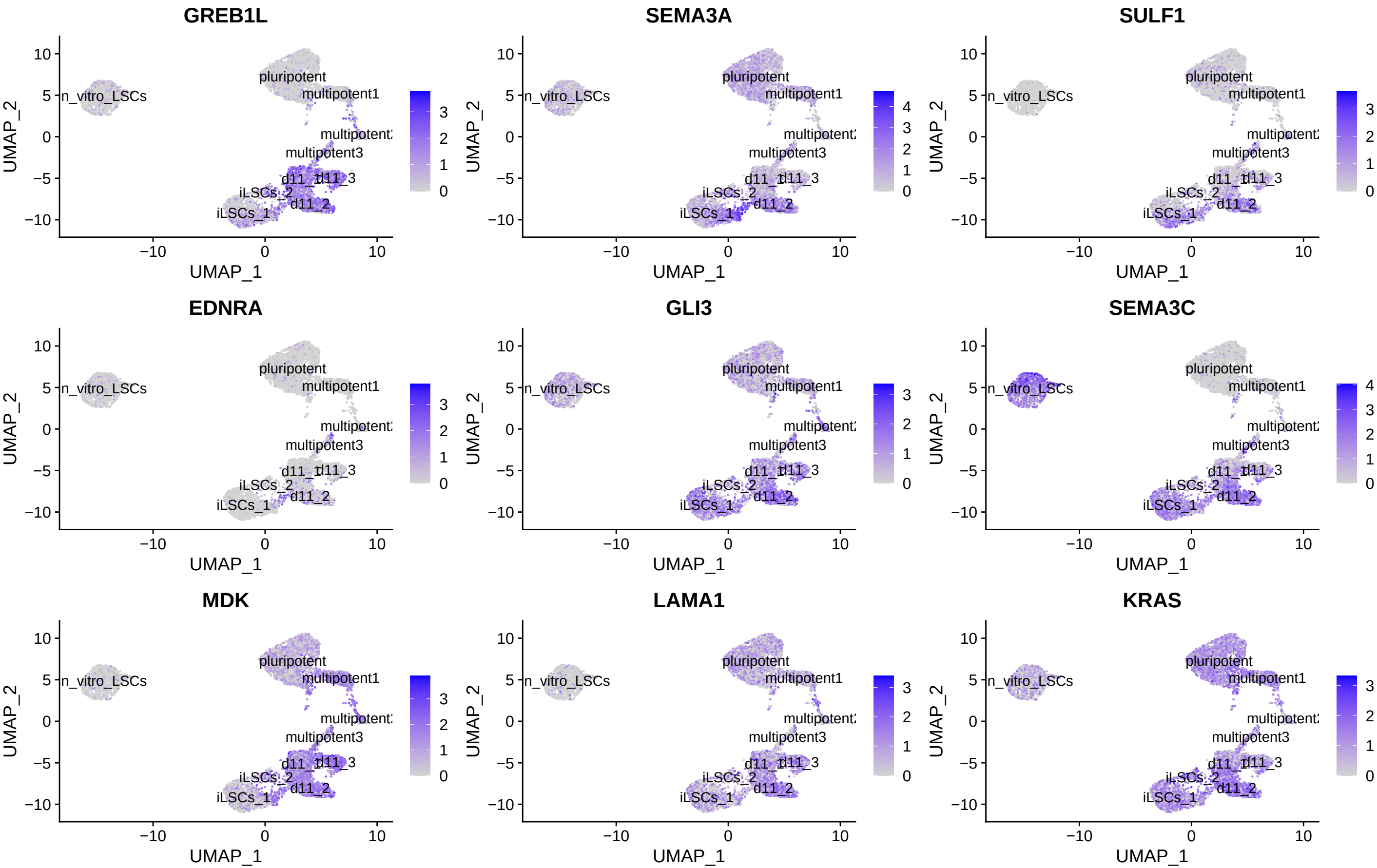
F2RL1



SDC4

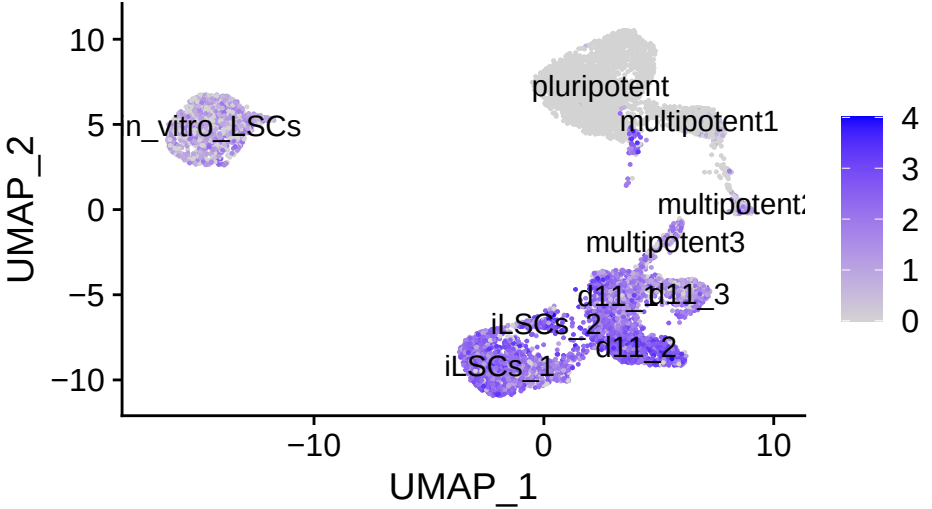


enriched go term: morphogenesis of a branching structure from cluster: f_d11_2

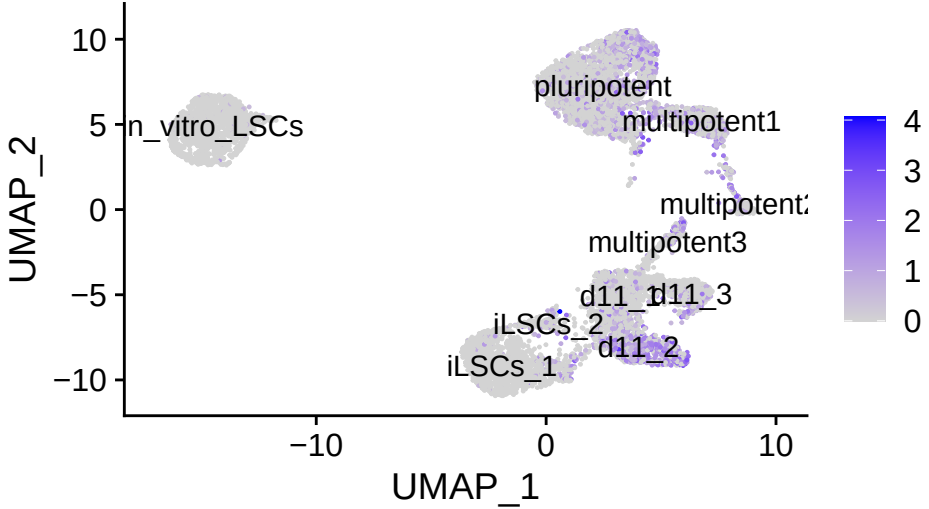


enriched go term: forebrain development from cluster: f_d11_2

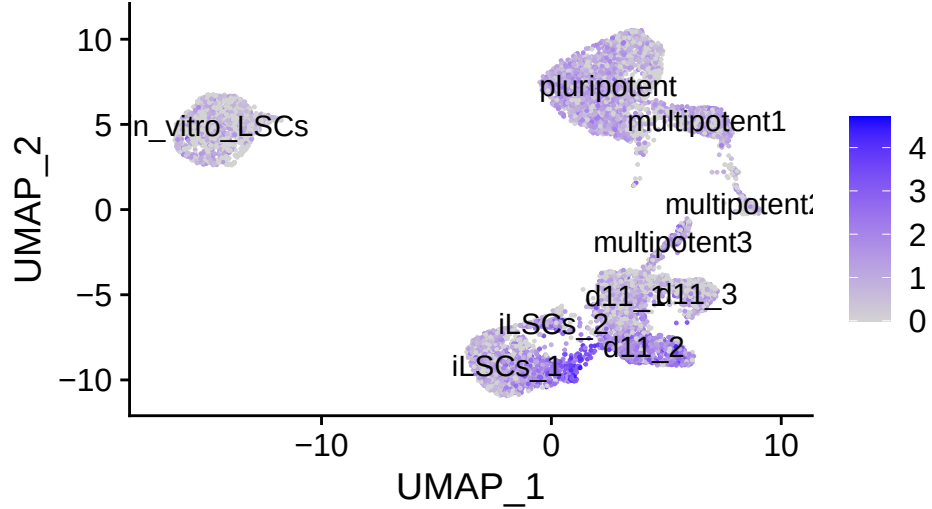
NR2F2



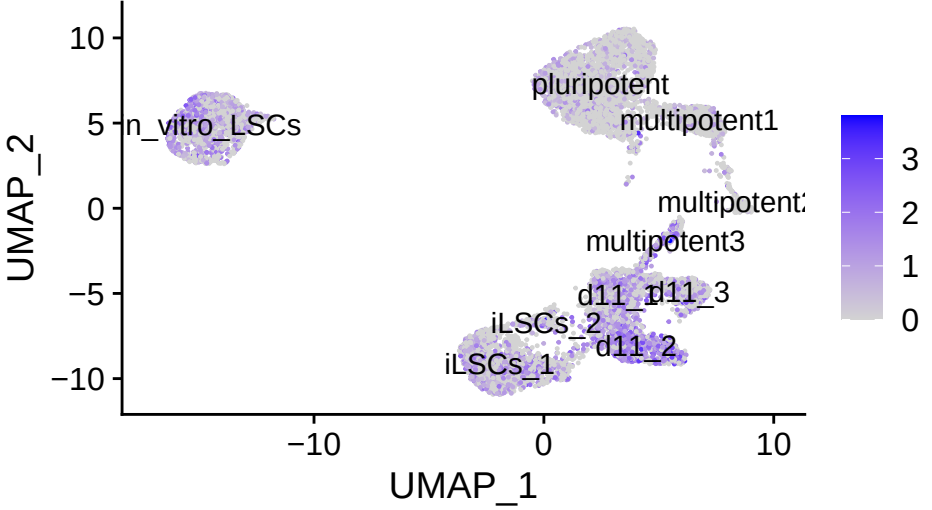
ID2



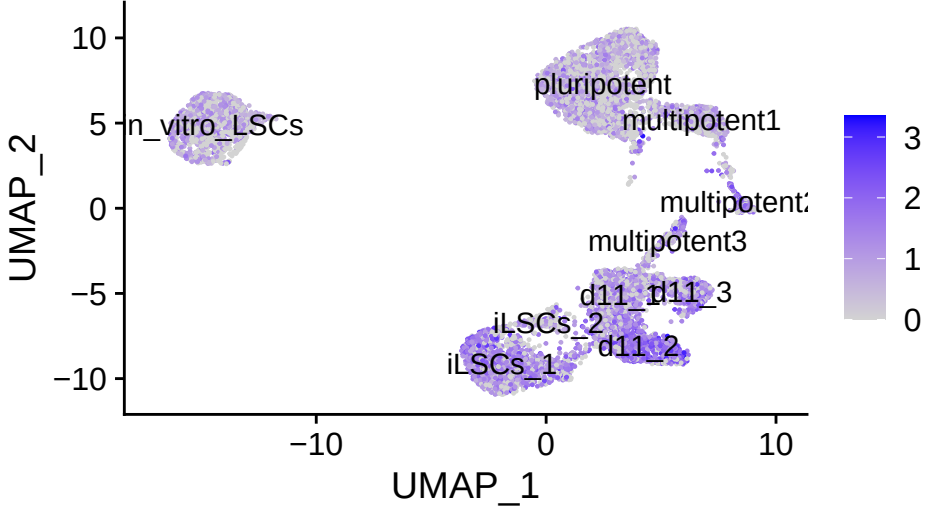
SEMA3A



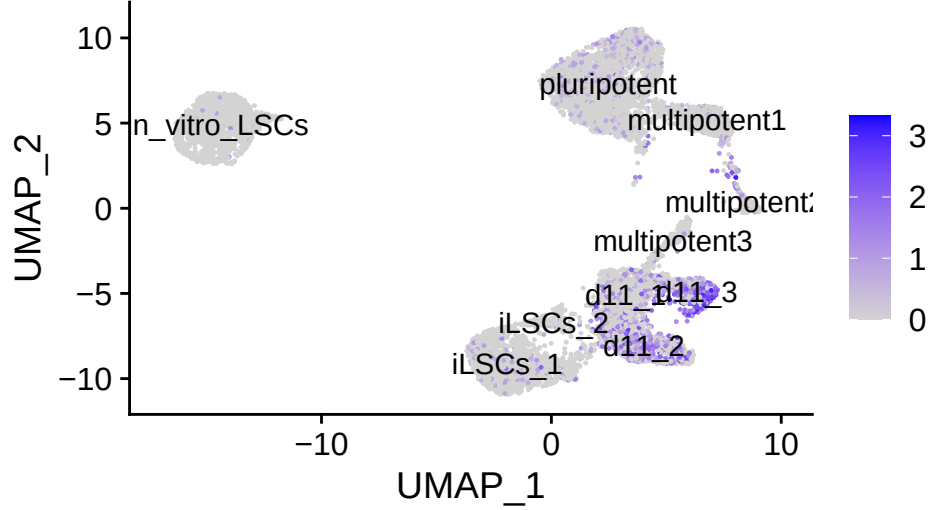
SLC8A1



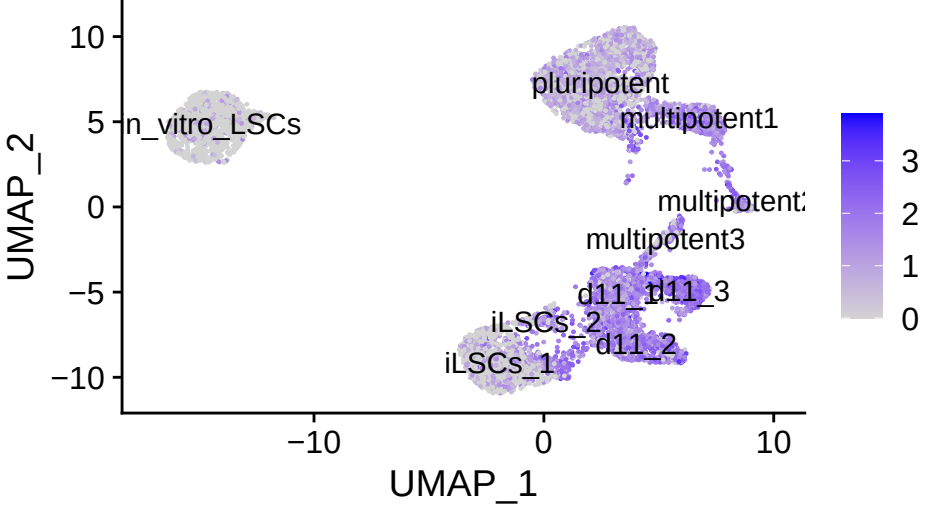
GLI3



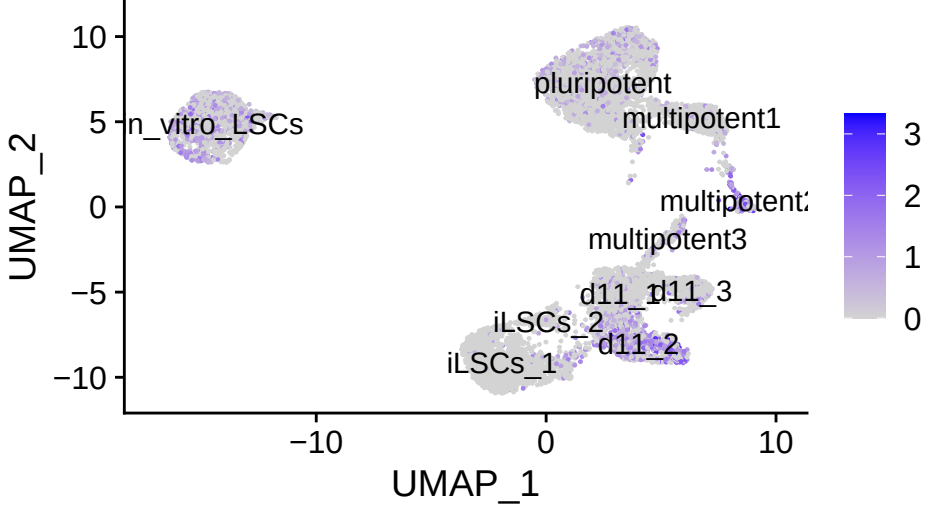
LRP2



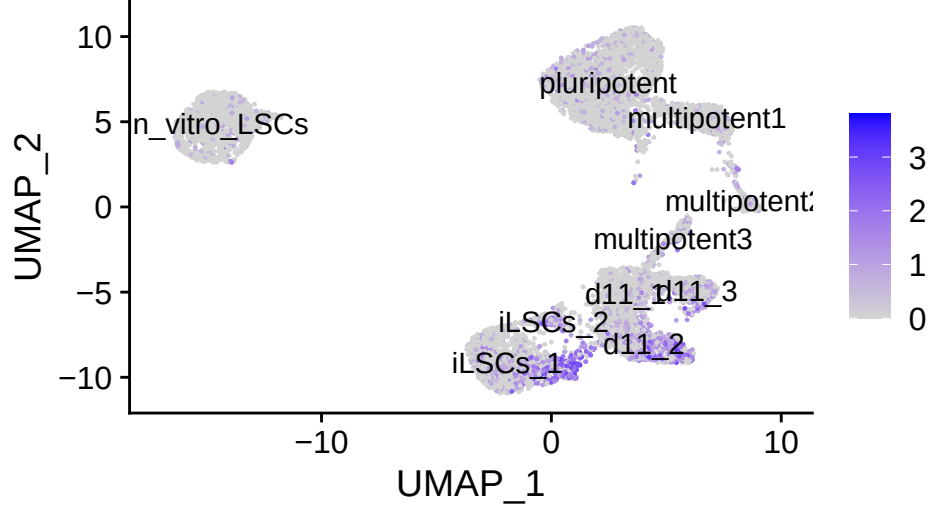
MDK



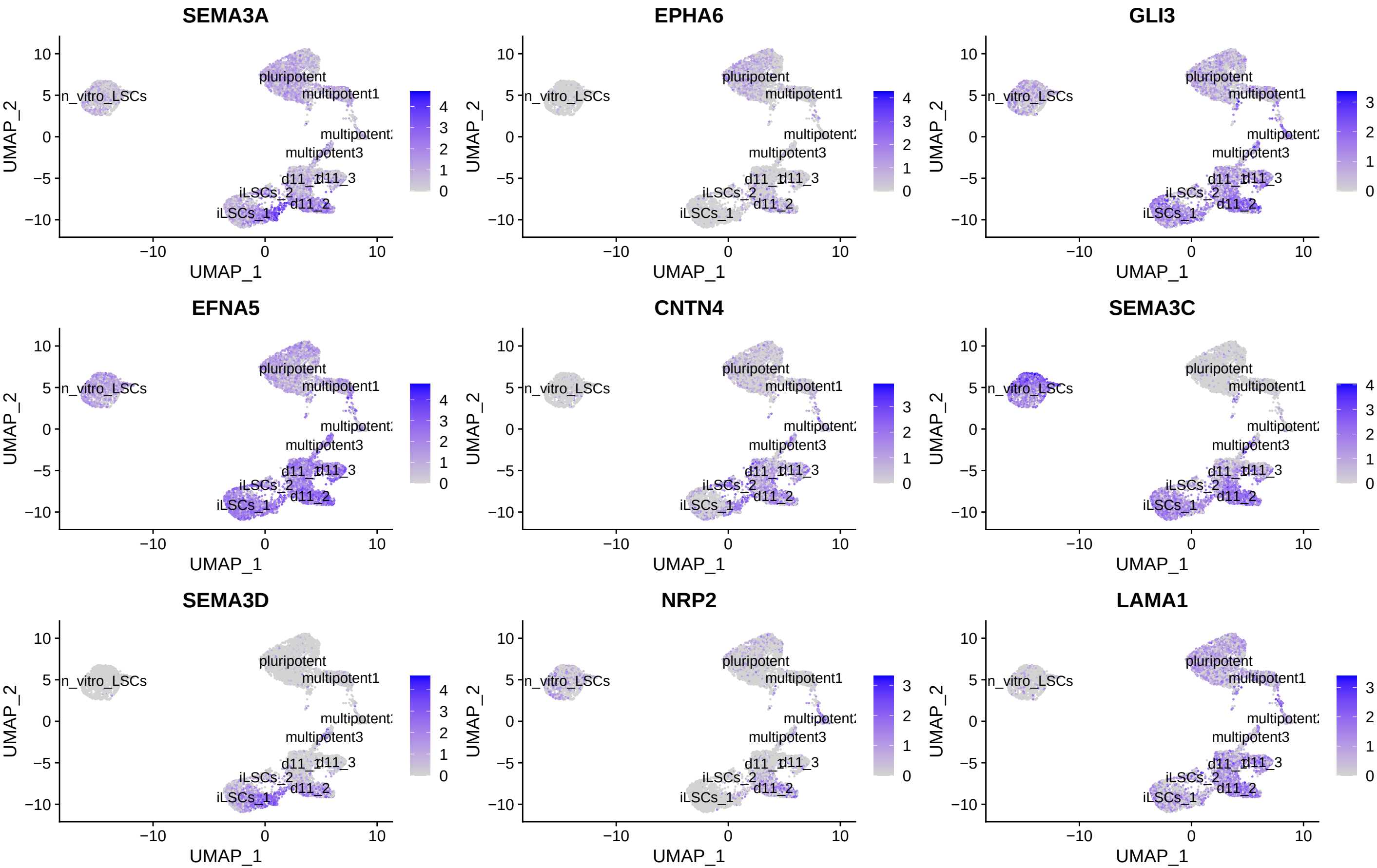
NRP2



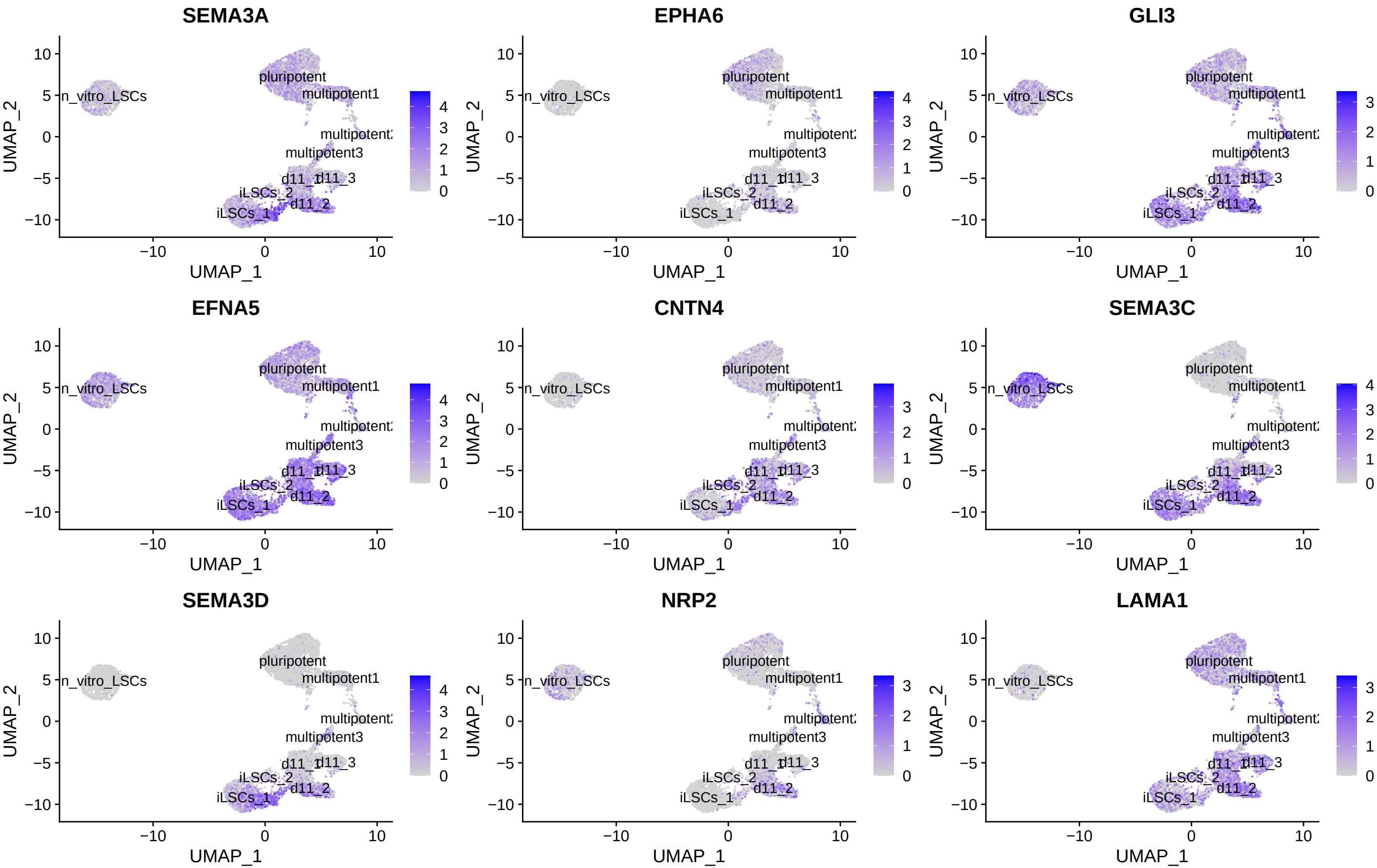
RARB



enriched go term: axon guidance from cluster: f_d11_2

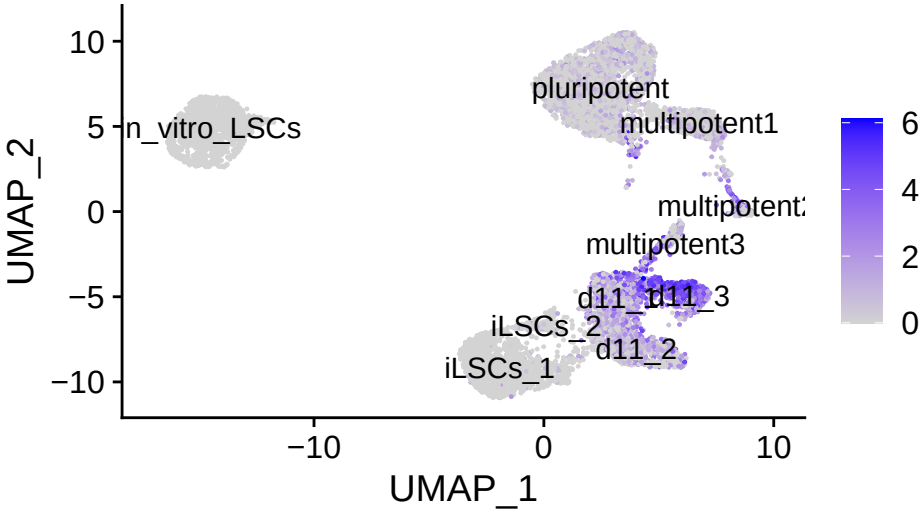


enriched go term: neuron projection guidance from cluster: f_d11_2

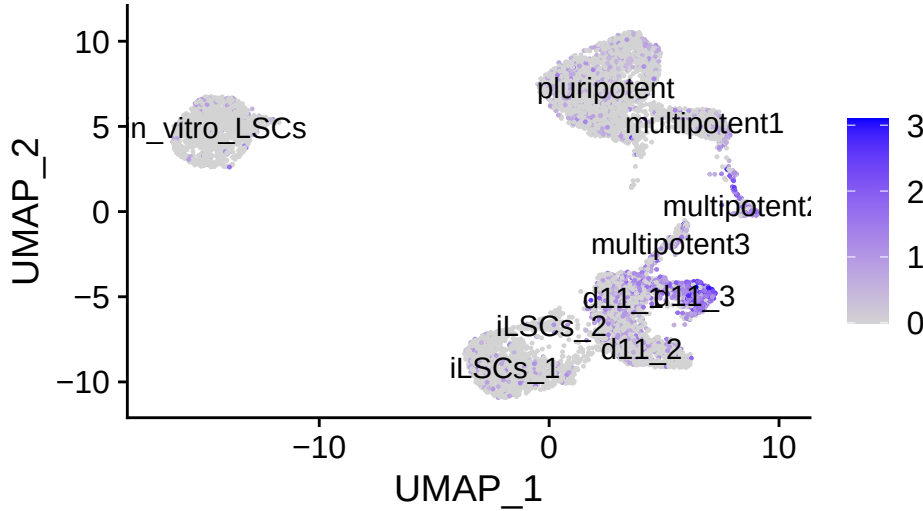


enriched go term: actin filament organization from cluster: g_d11_3

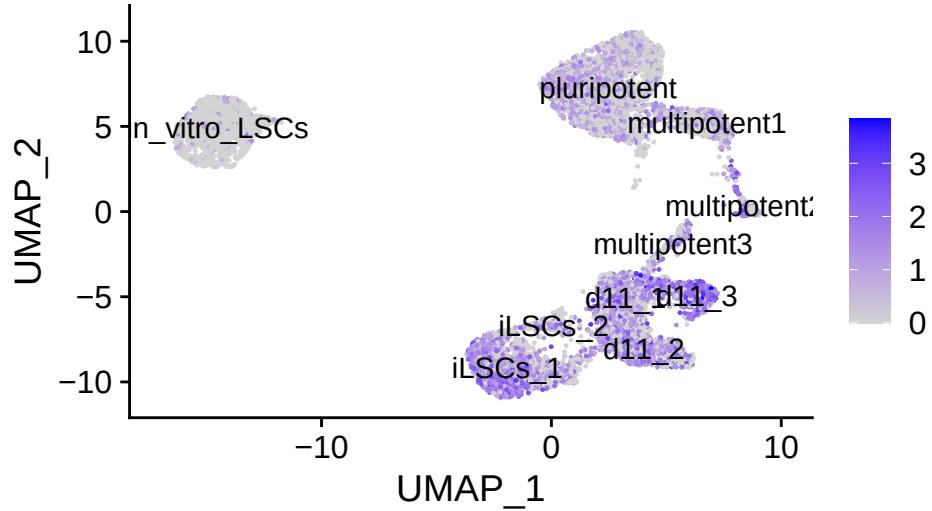
ACTC1



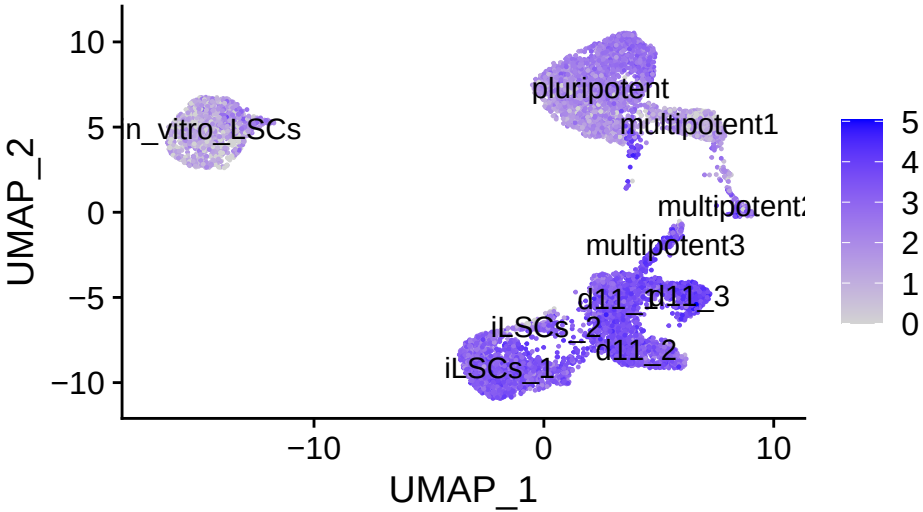
WIPF1



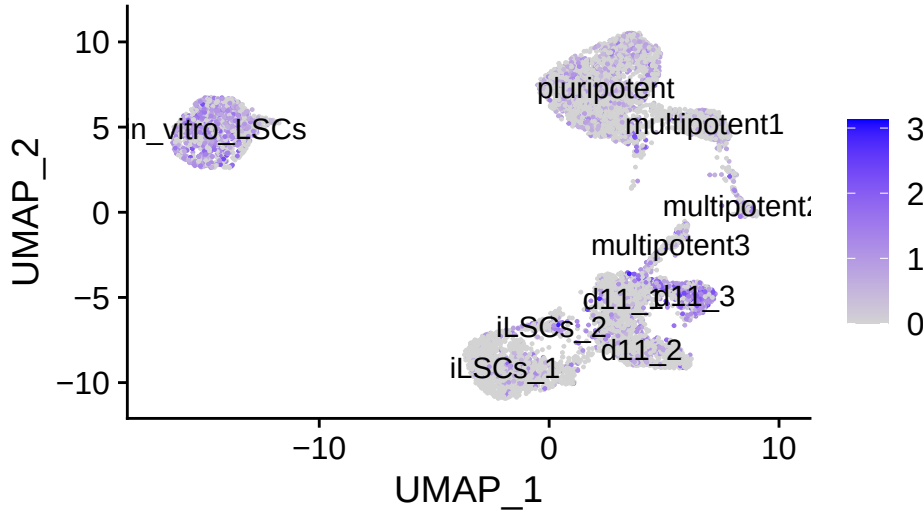
SEMA5A



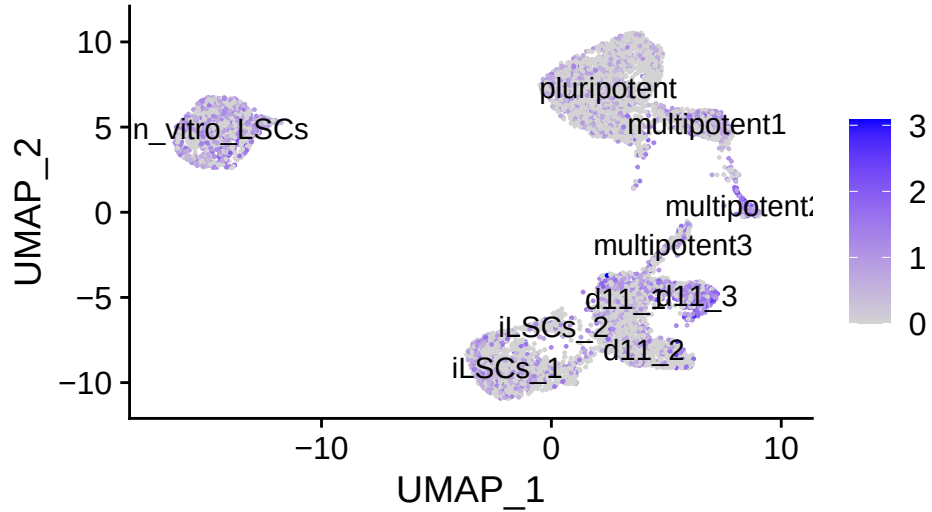
TPM1



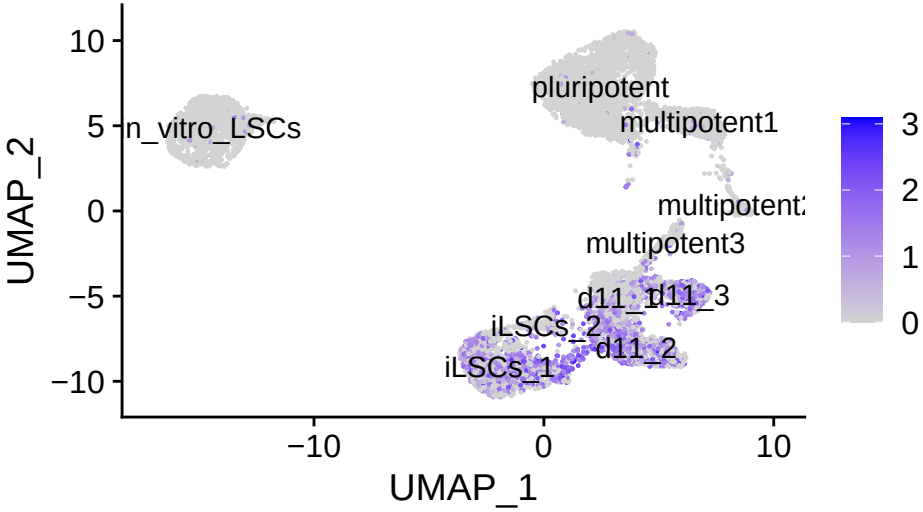
EMP2



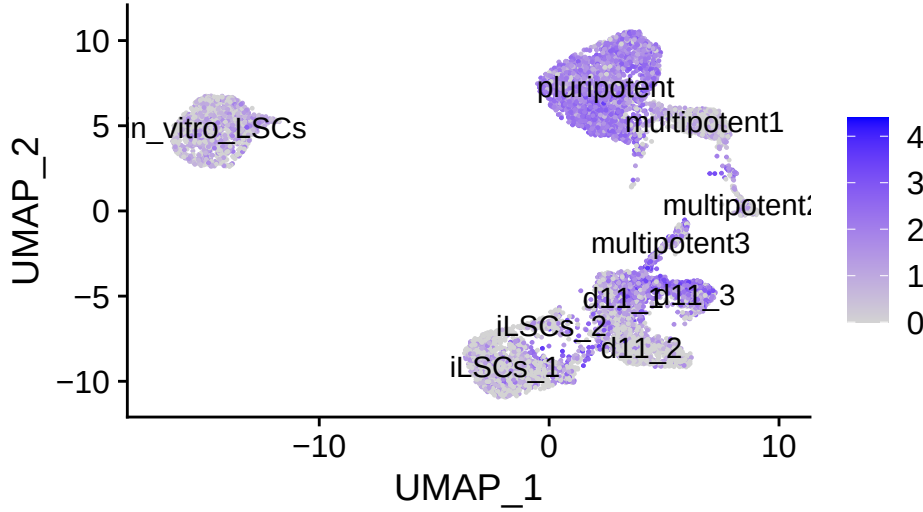
PRKCE



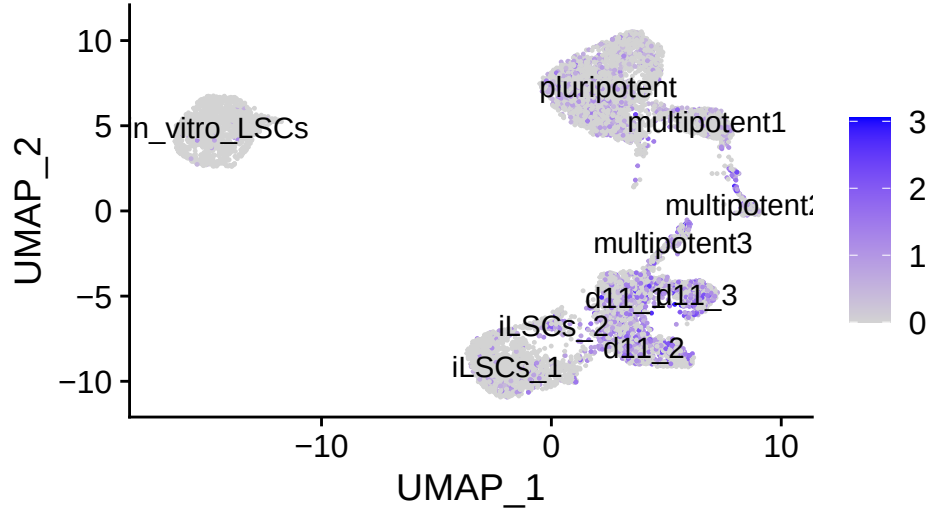
KANK4



SFRP1

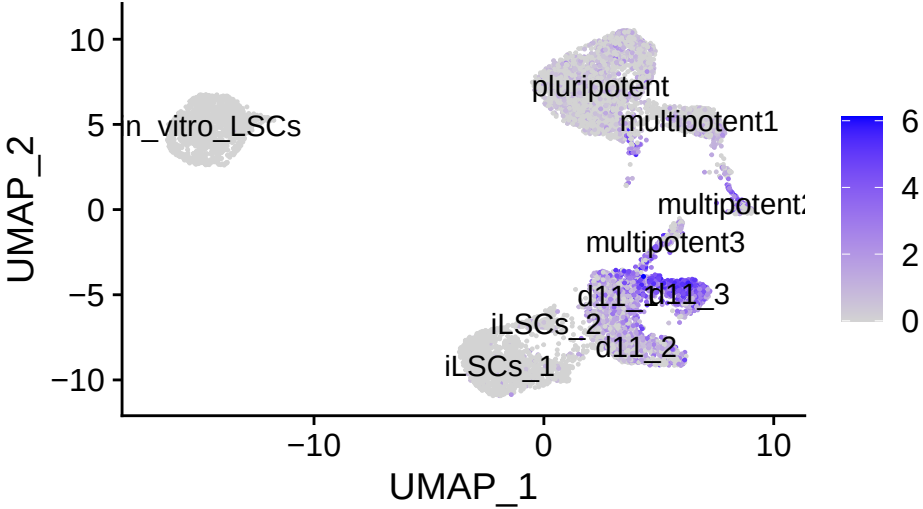


ARHGAP28

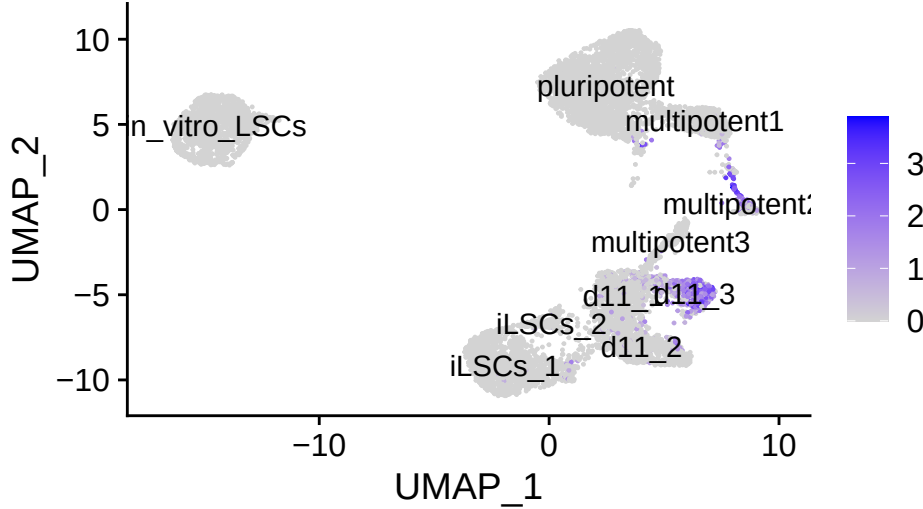


enriched go term: mesenchyme development from cluster: g_d11_3

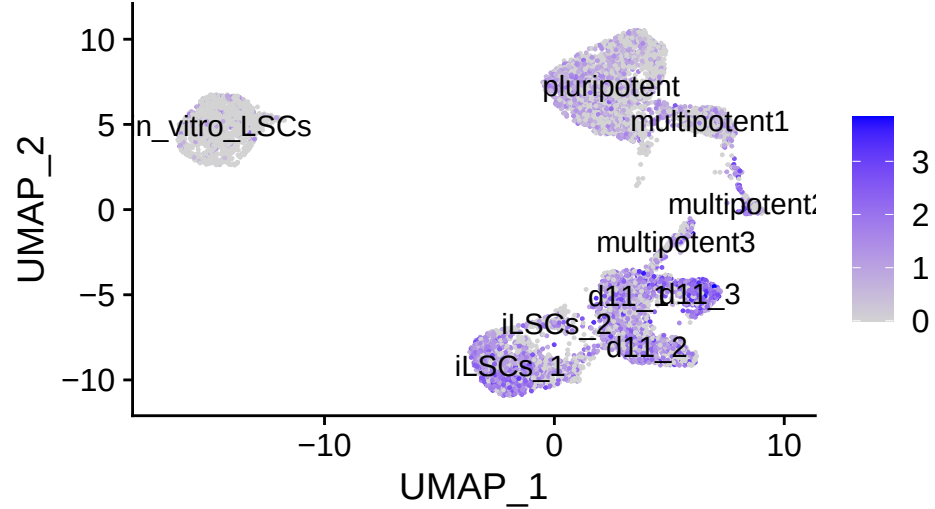
ACTC1



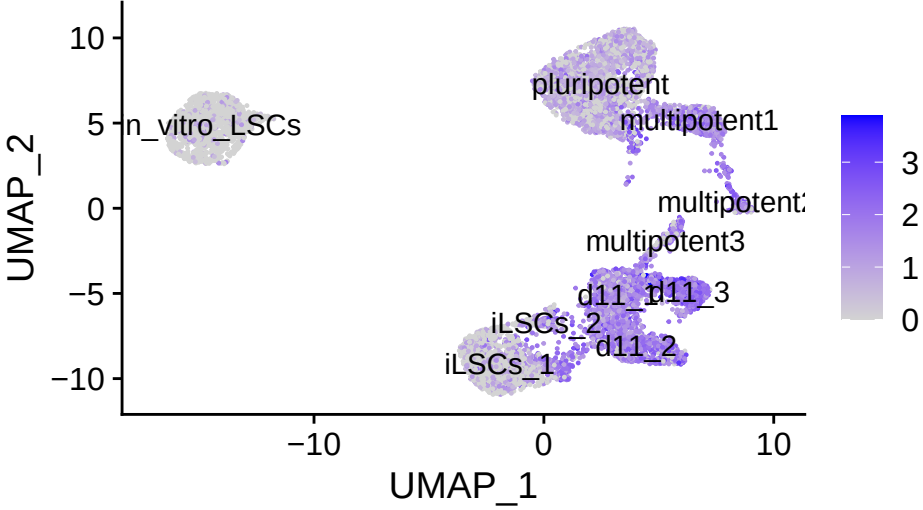
HAND1



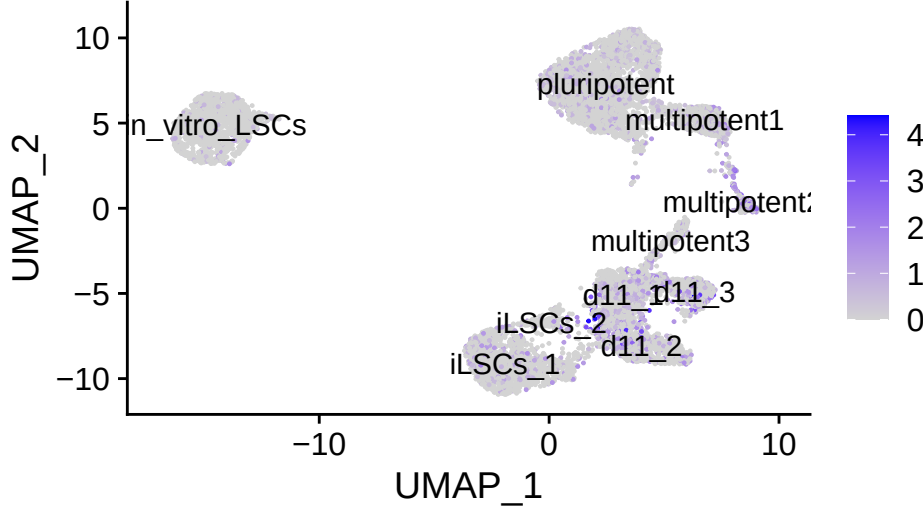
SEMA5A



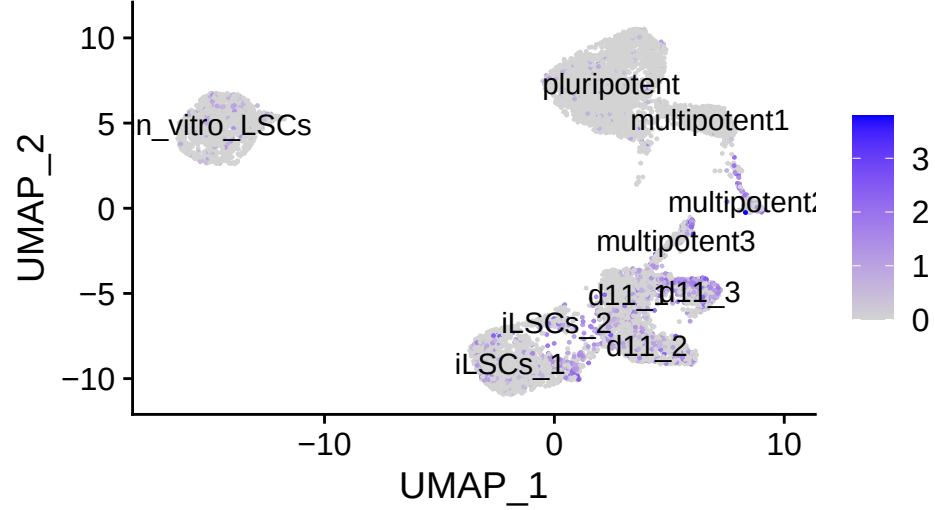
MDK



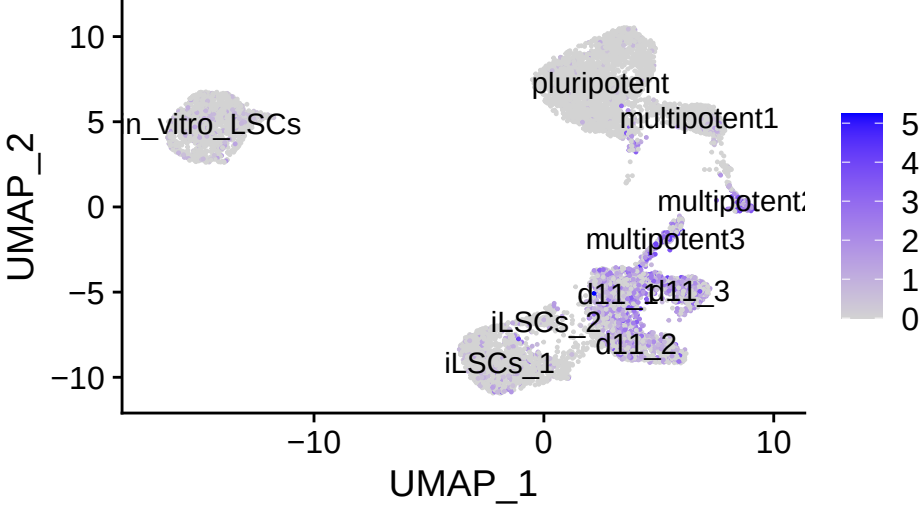
SEMA6D



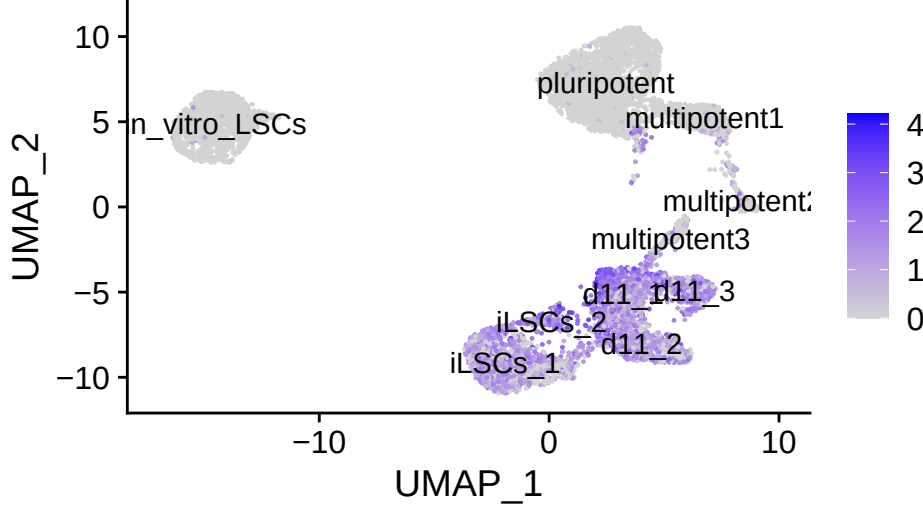
WNT5A



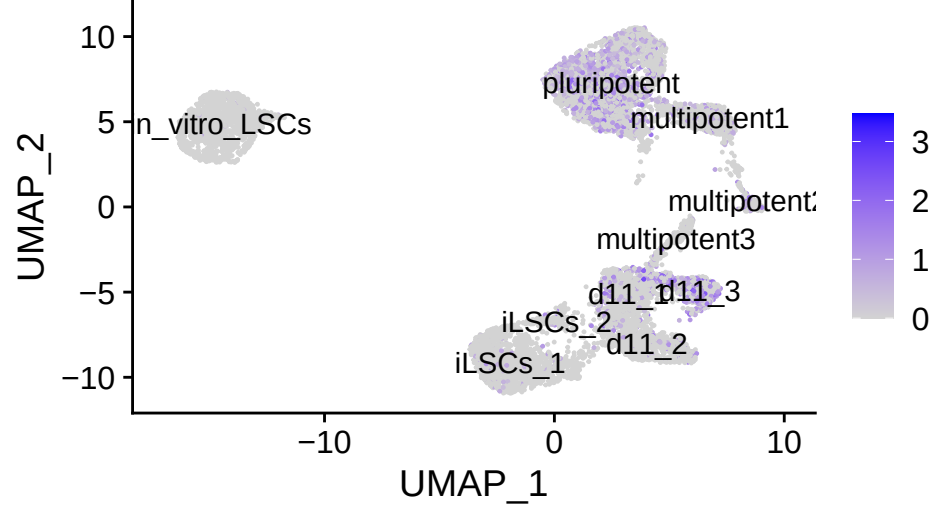
ACTA2



GATA3

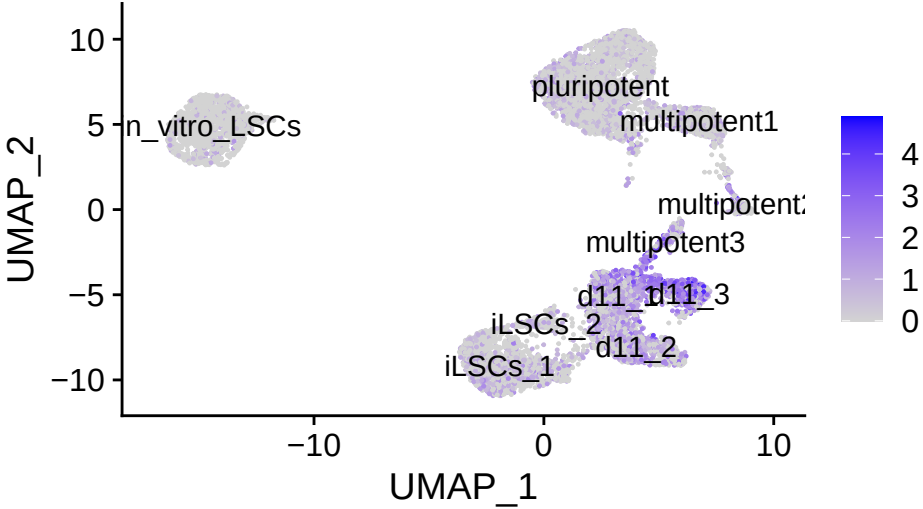


EDNRB

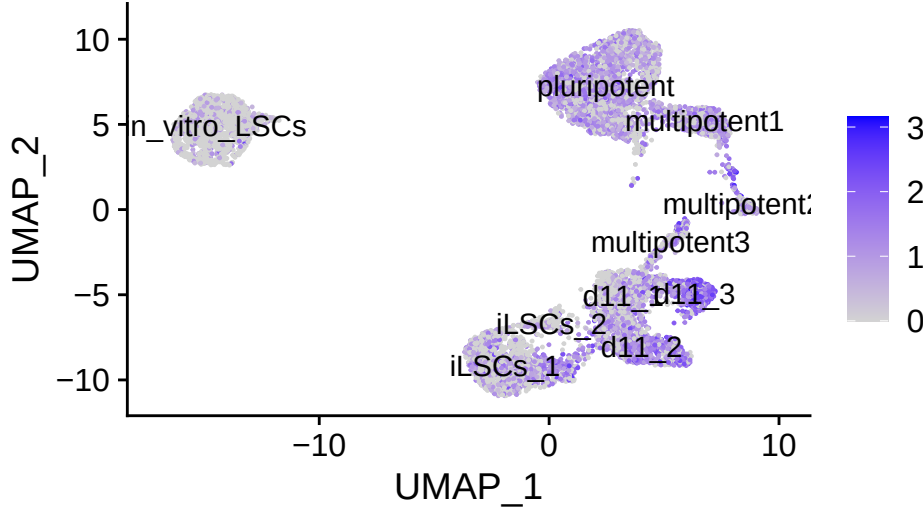


enriched go term: cell junction assembly from cluster: g_d11_3

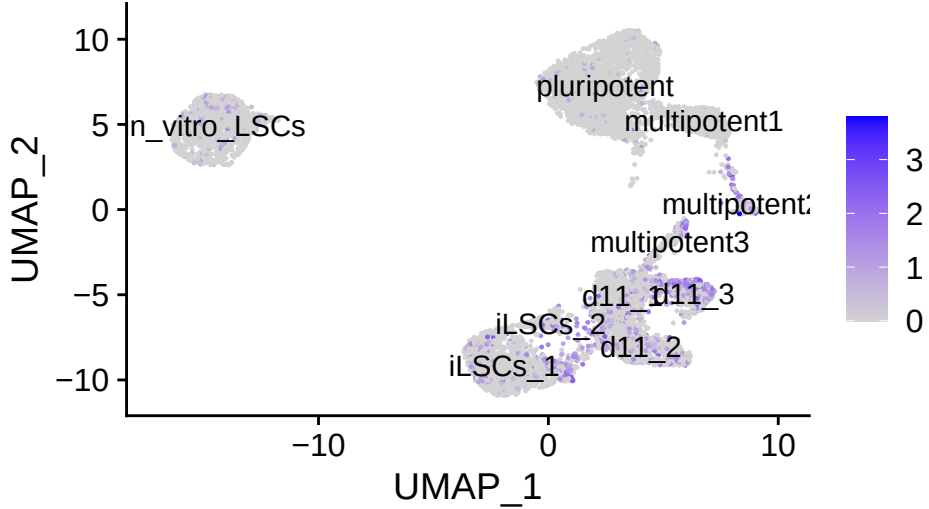
CLSTN2



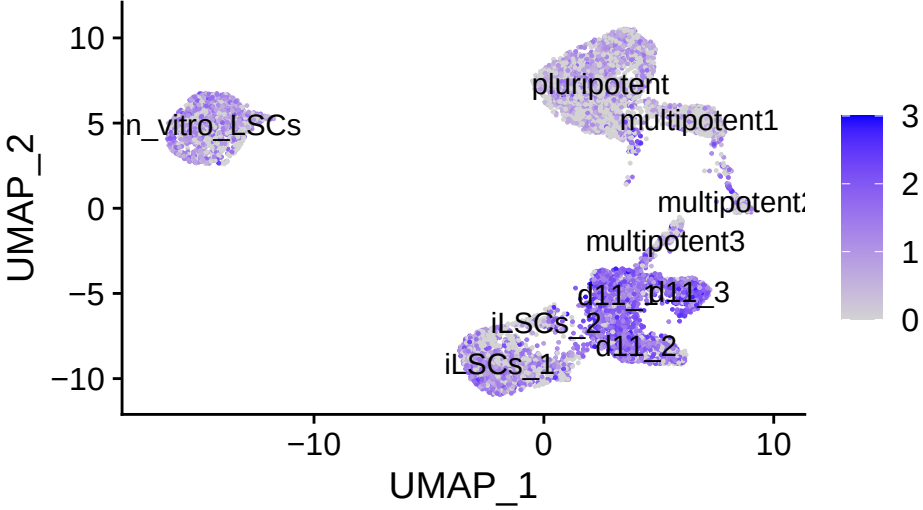
TLN2



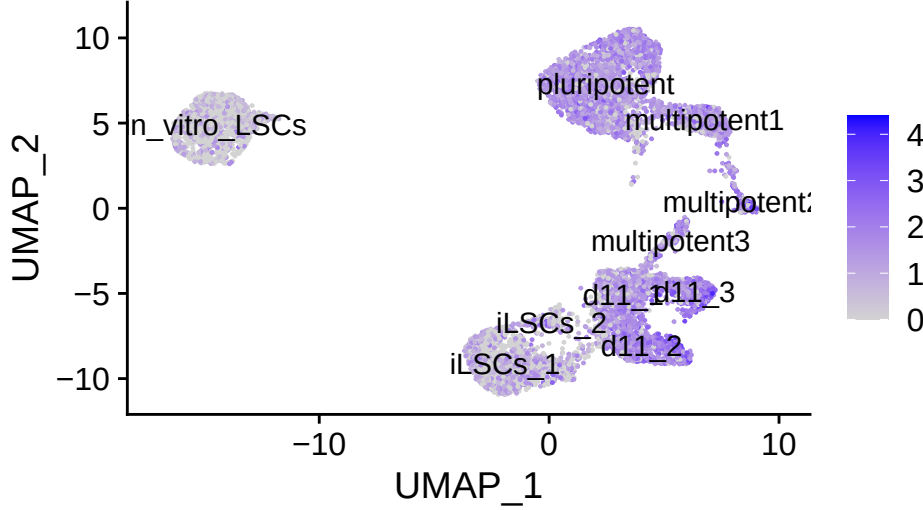
WNT5A



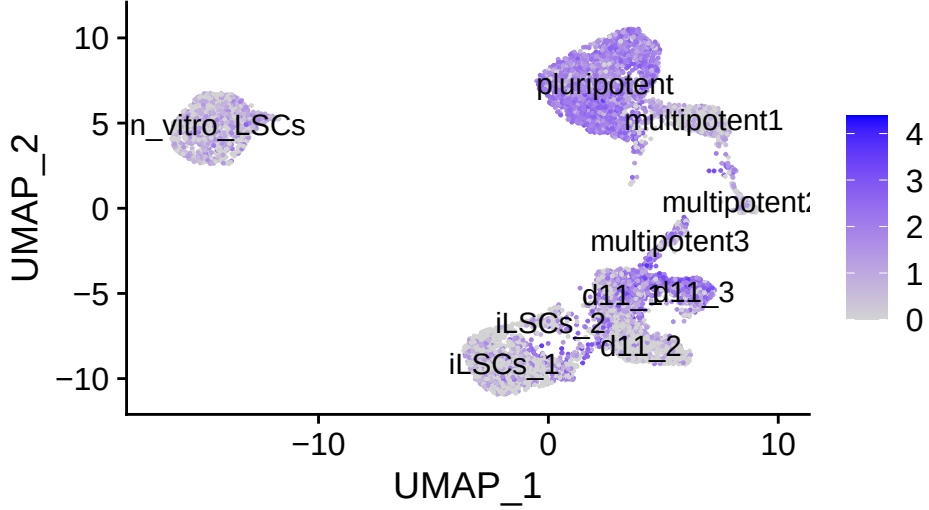
PKP2



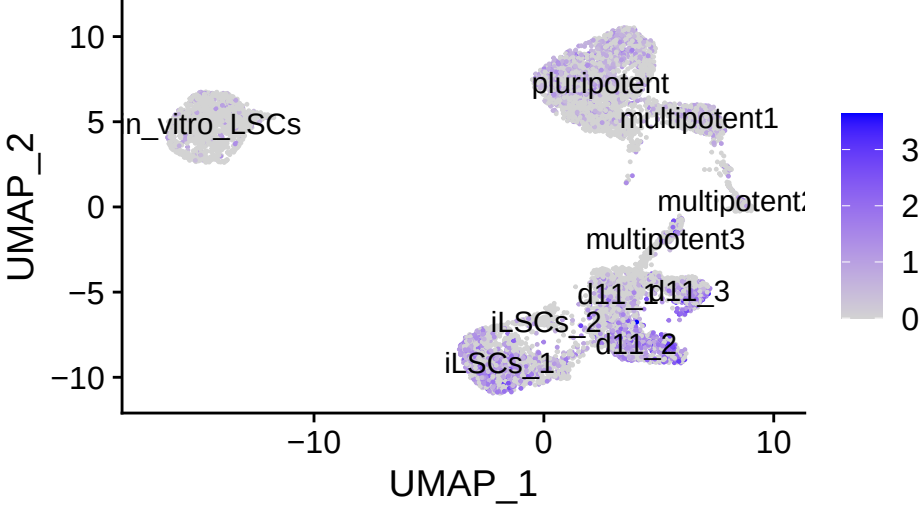
ADGRL2



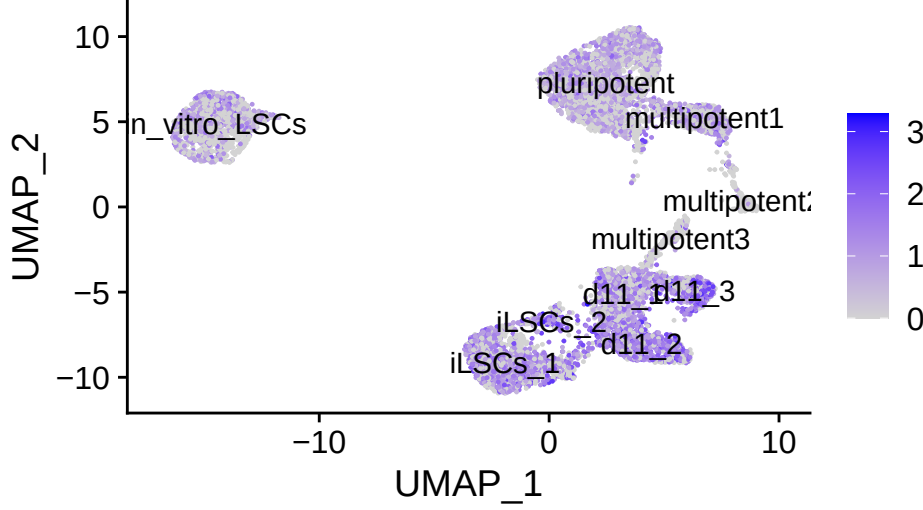
SFRP1



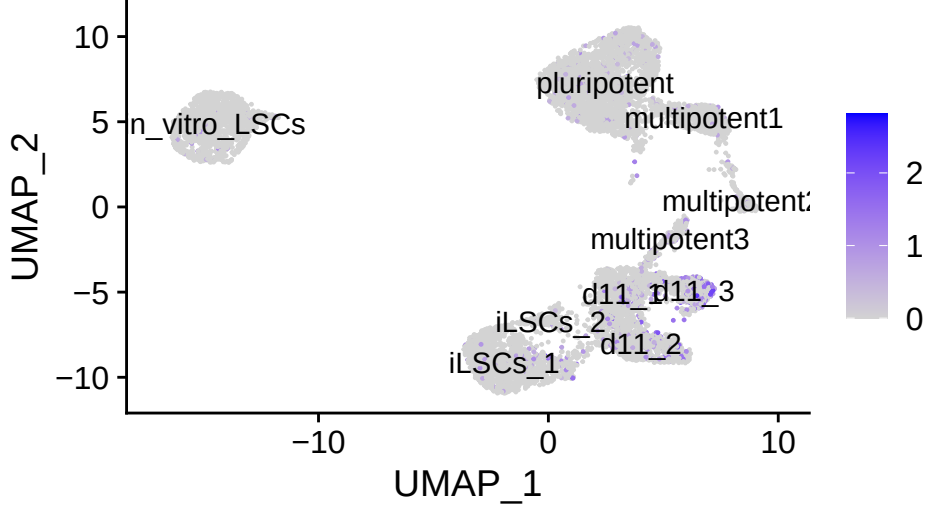
CDH8



GRHL2

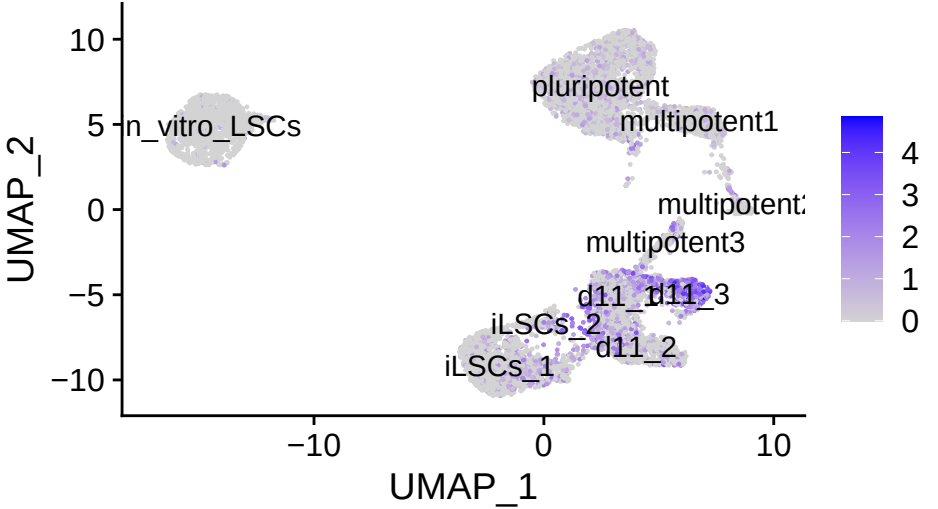


NTRK2

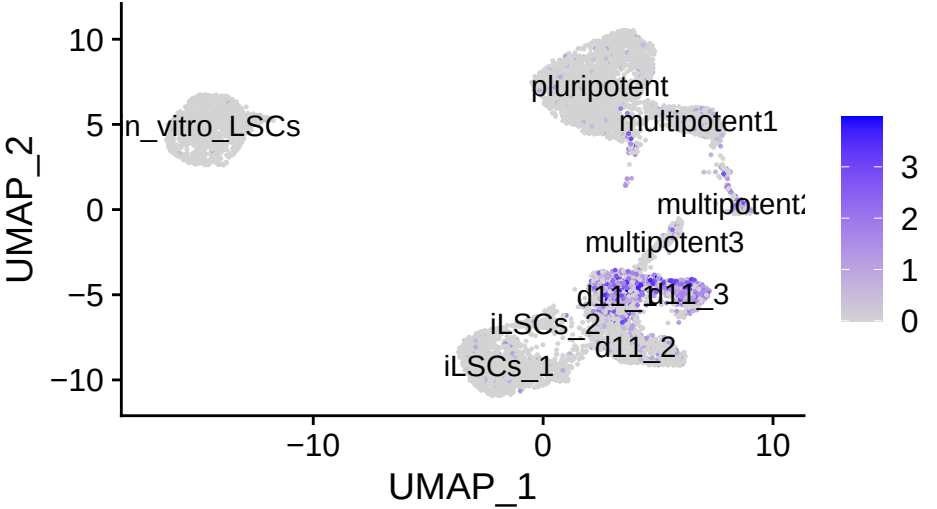


enriched go term: cell-substrate adhesion from cluster: g_d11_3

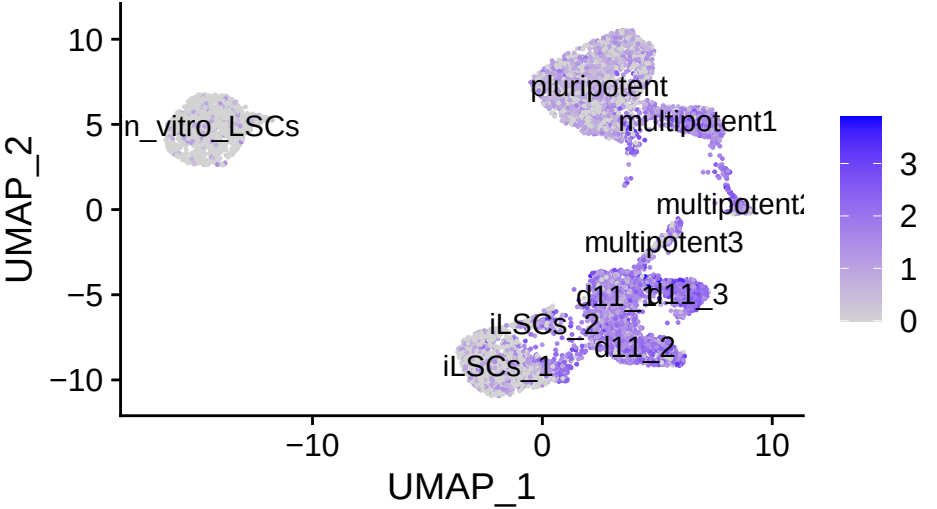
PTN



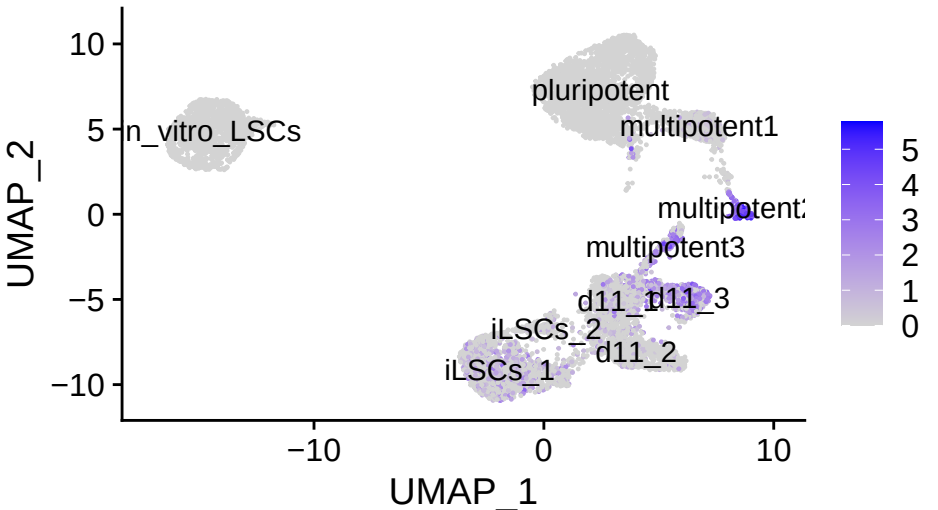
ADAMTS9



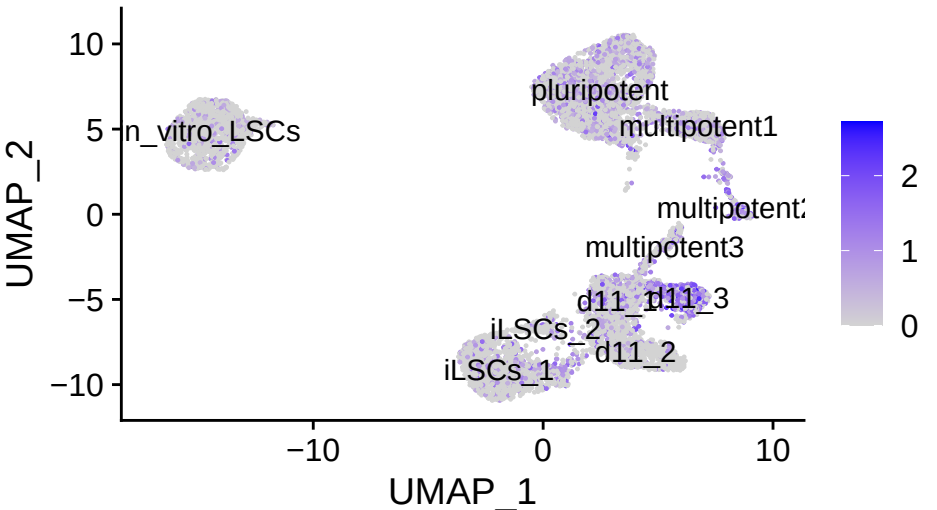
MDK



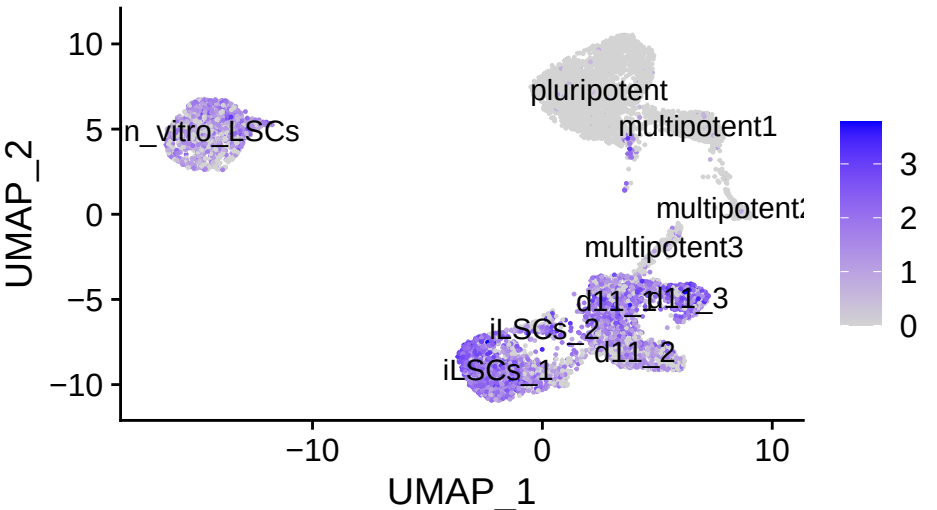
COL3A1



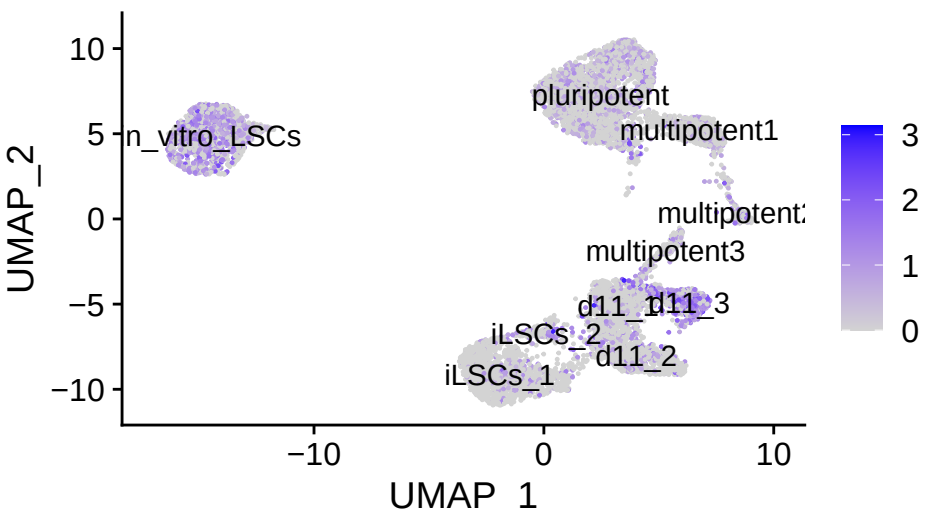
FBLN1



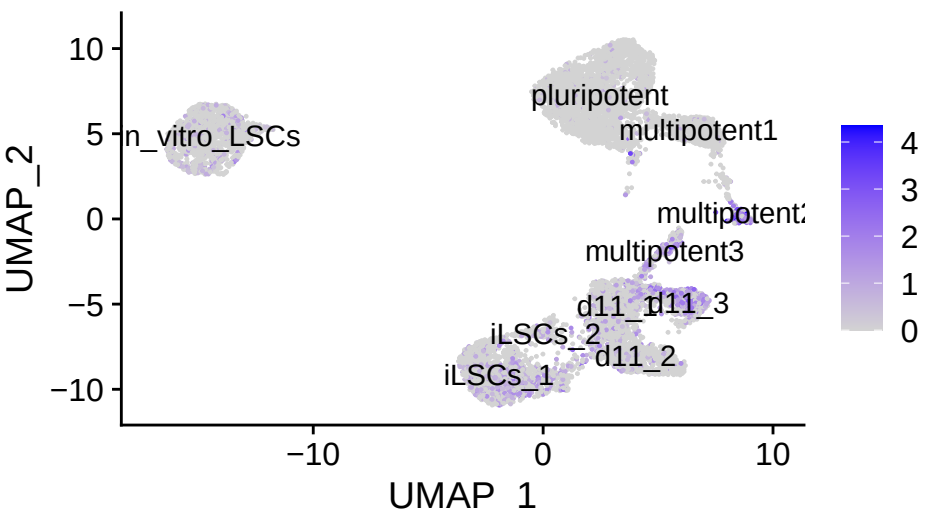
ITGB6



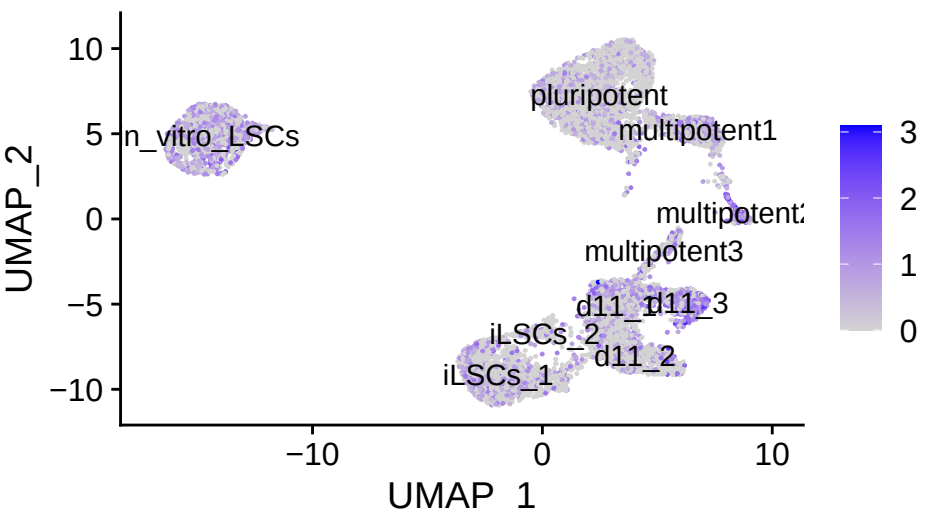
EMP2



CCDC80

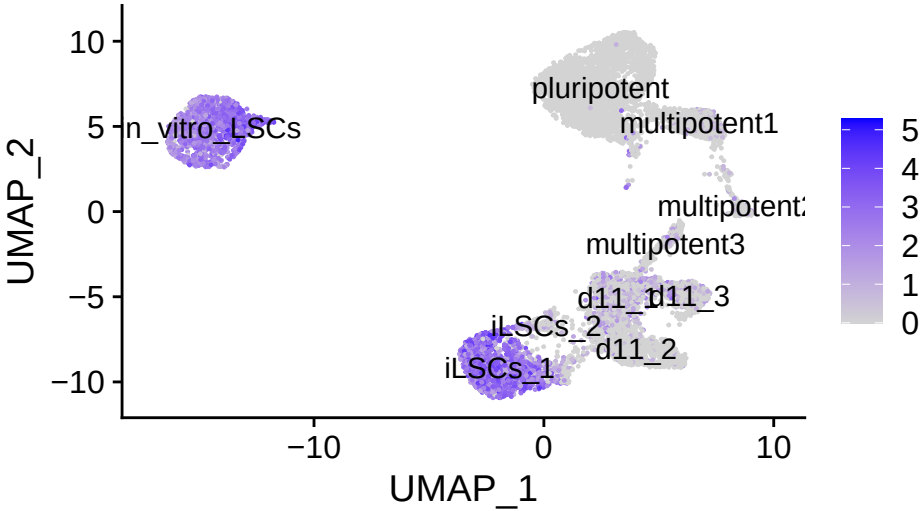


PRKCE

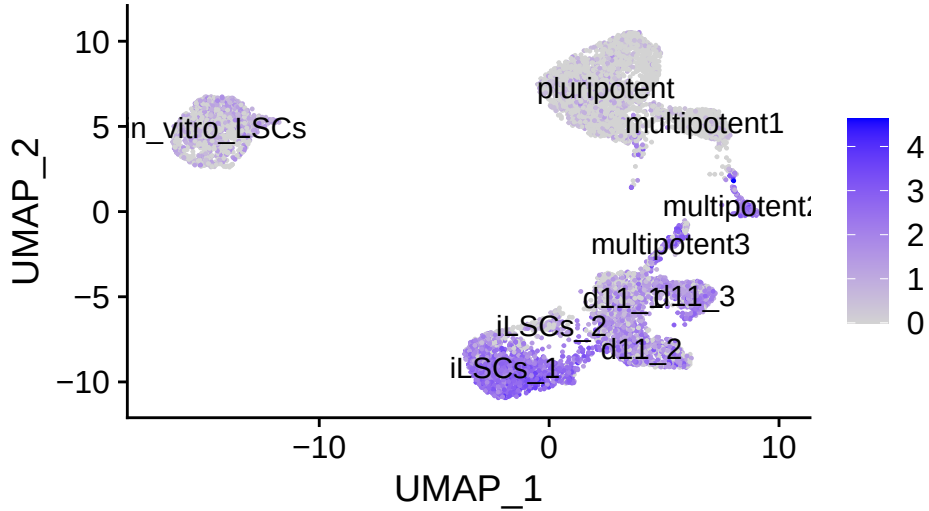


enriched go term: skin development from cluster: h_iLSCs_1

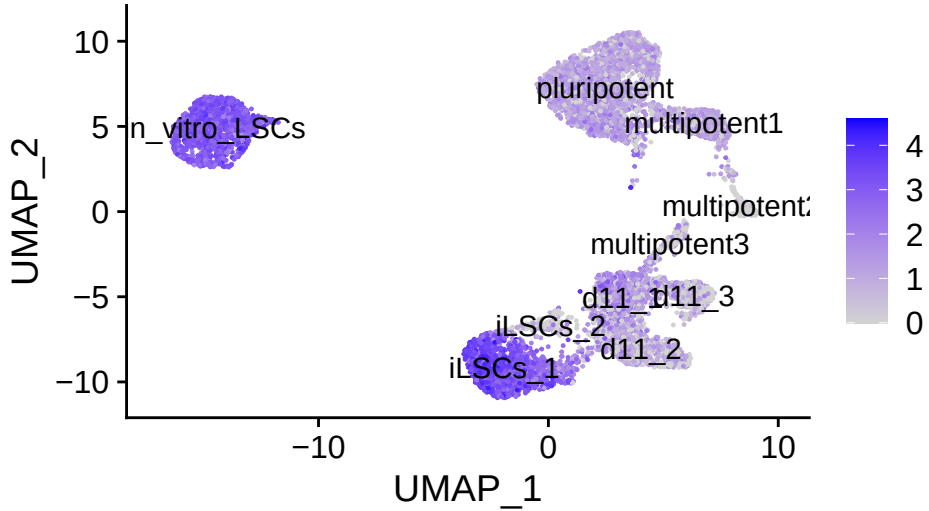
KRT17



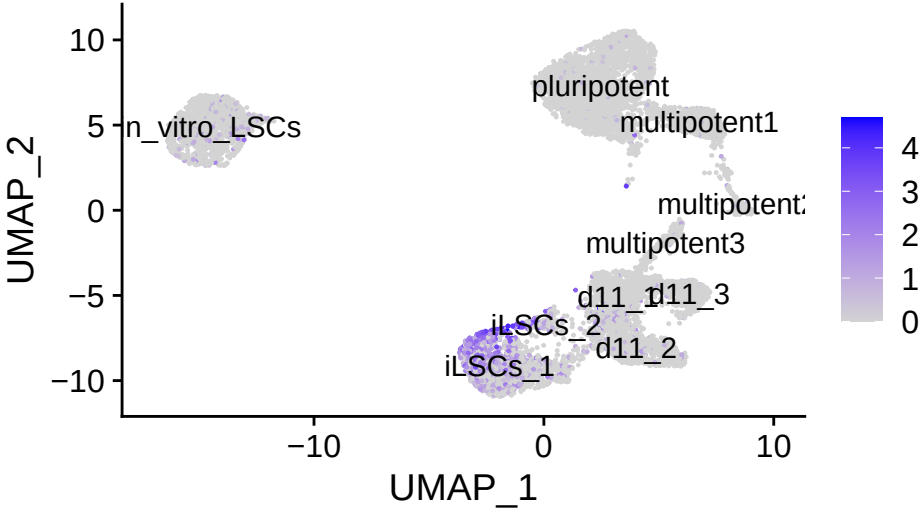
COL5A2



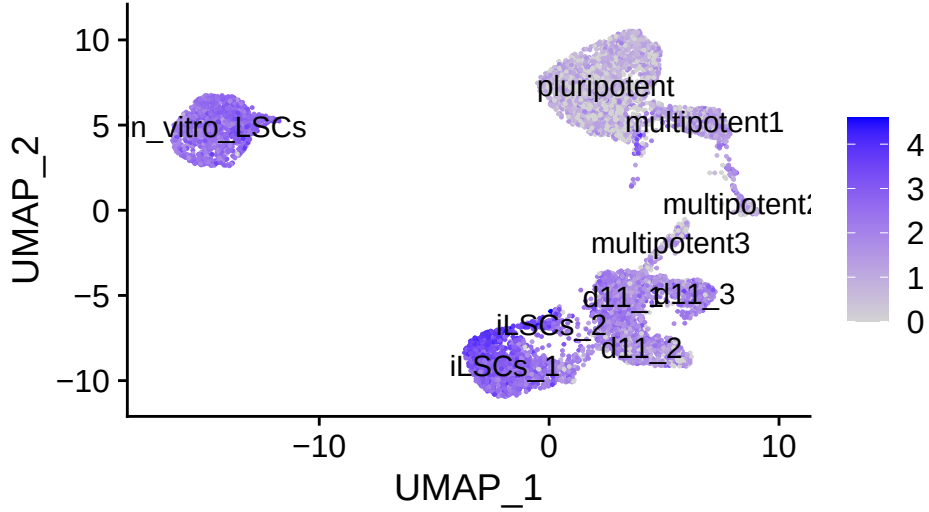
ITGA6



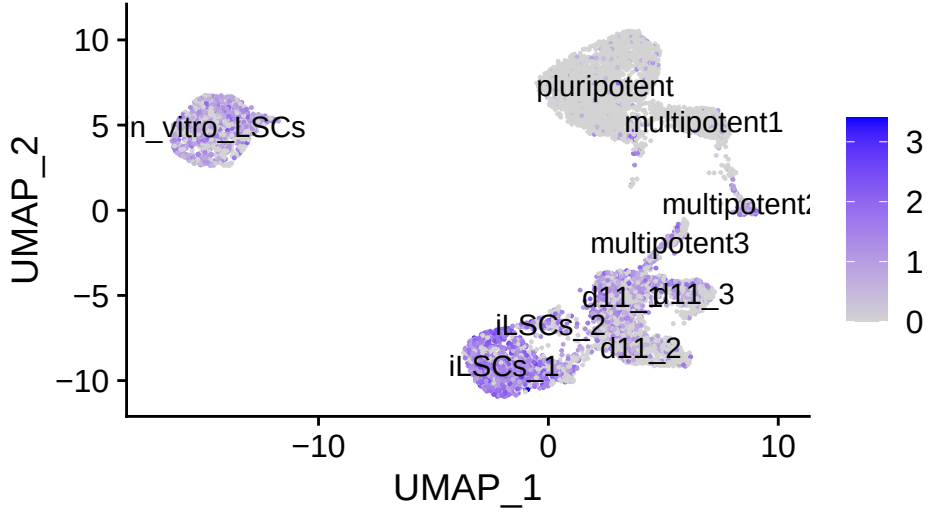
AQP3



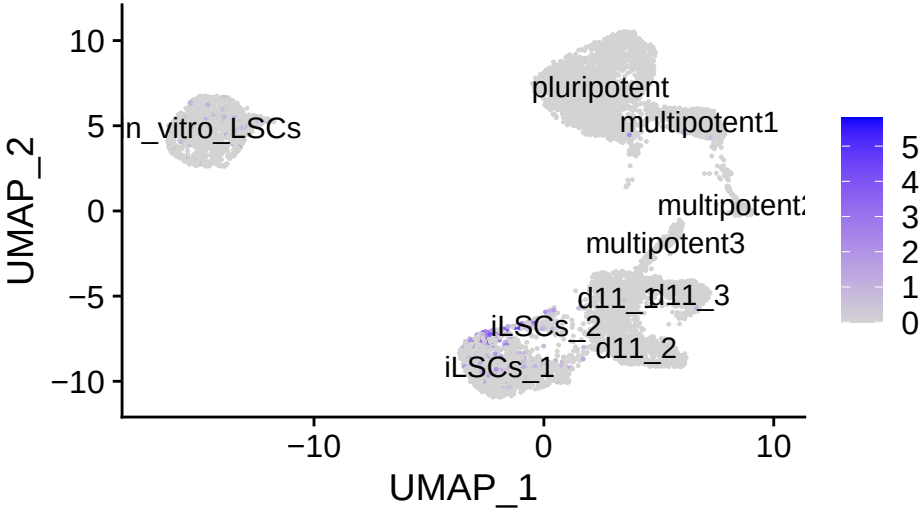
DSP



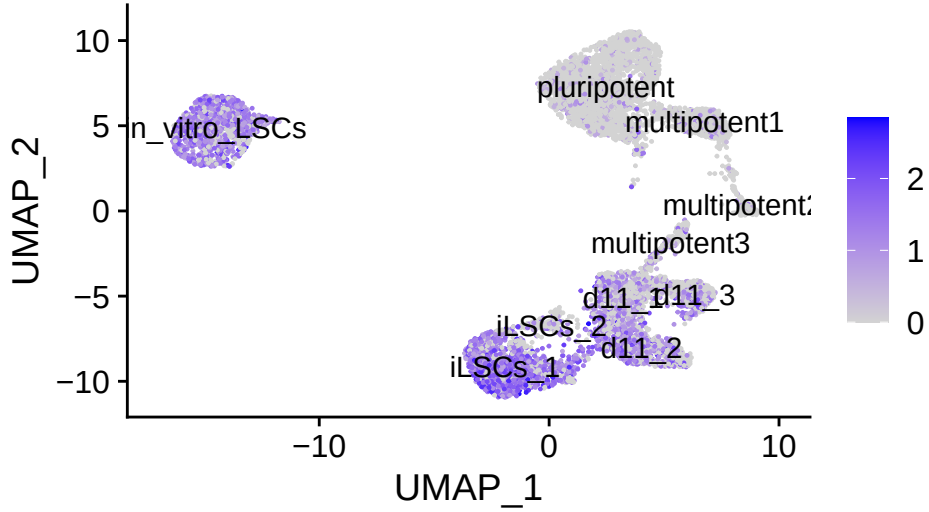
FOSL2



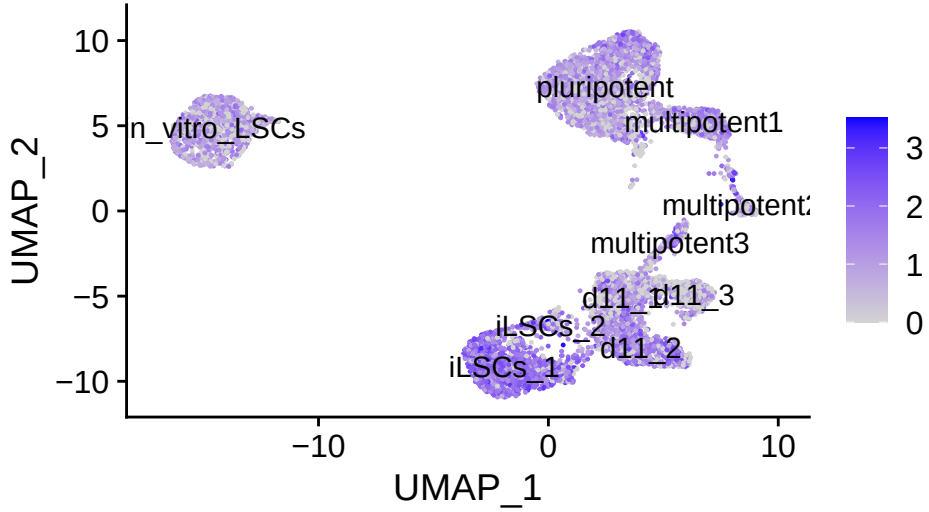
SPRR1B



MET

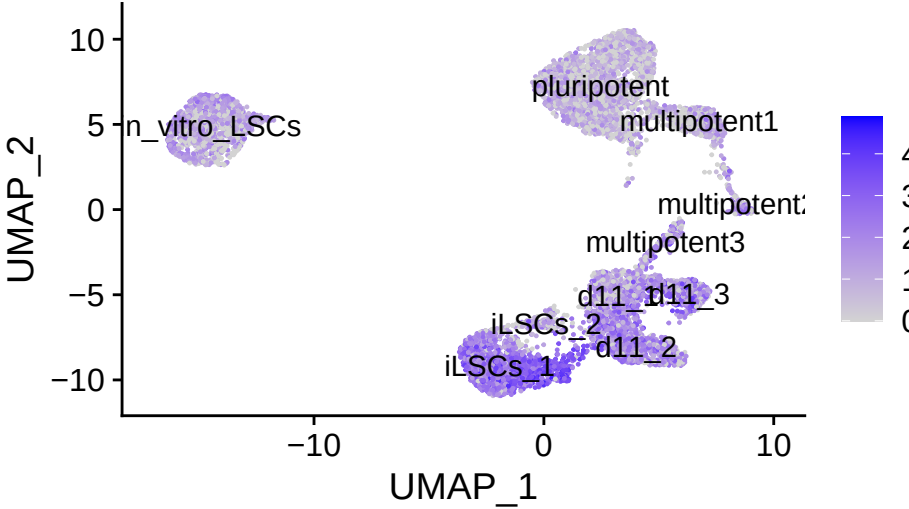


LGR4

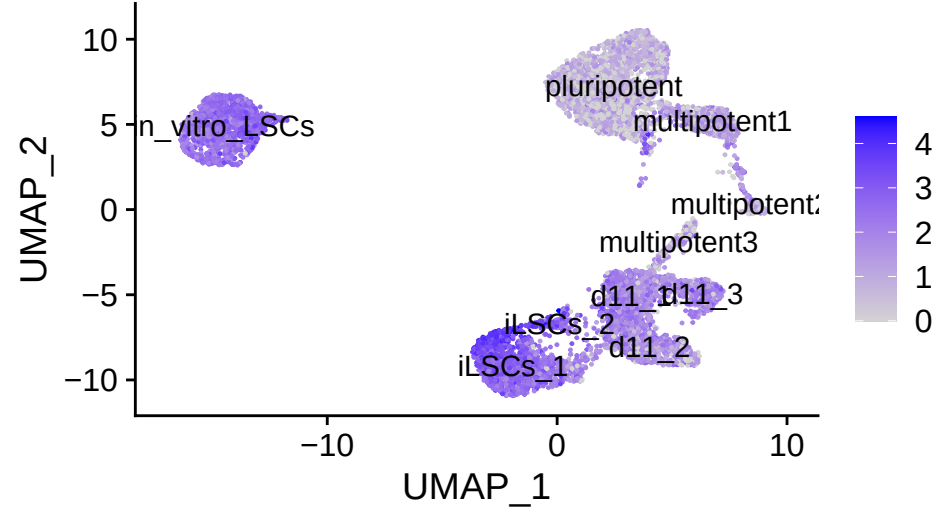


enriched go term: regulation of actin filament-based process from cluster: h_iLSCs_1

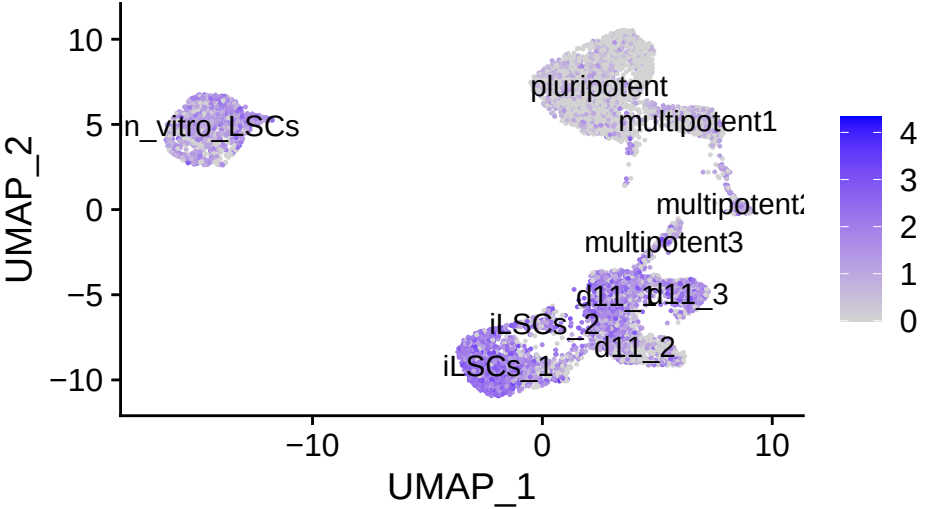
PDE4D



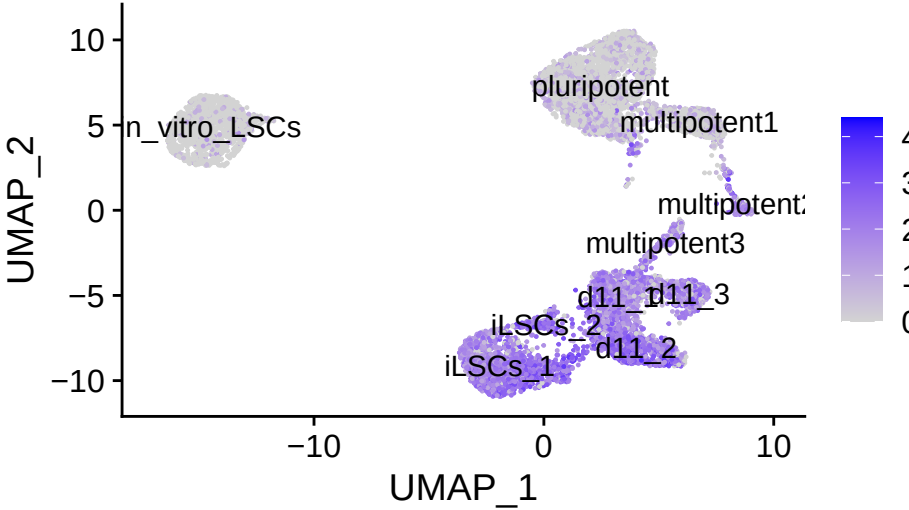
DSP



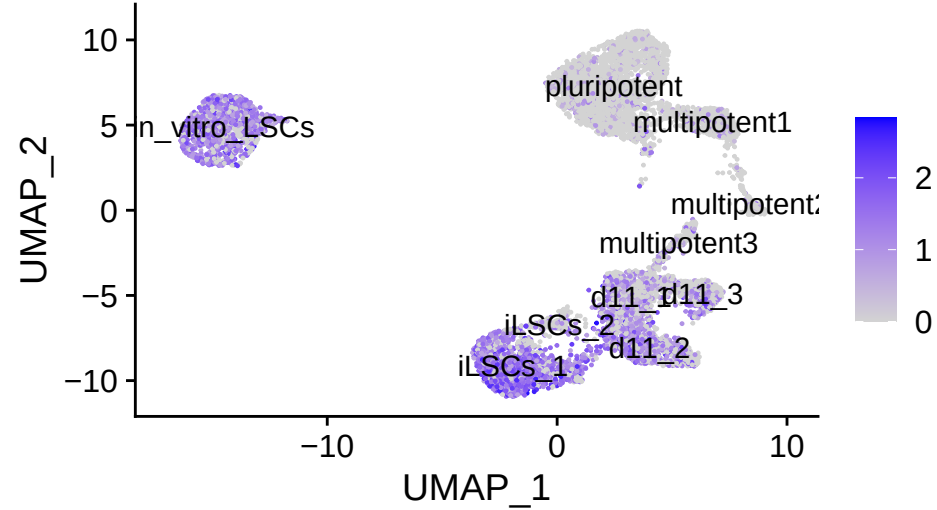
PHLDB2



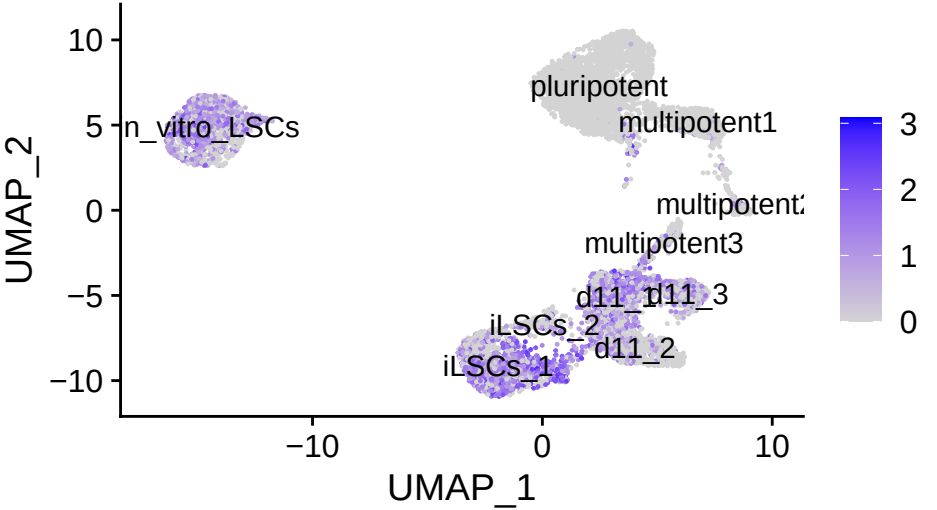
RHOBTB3



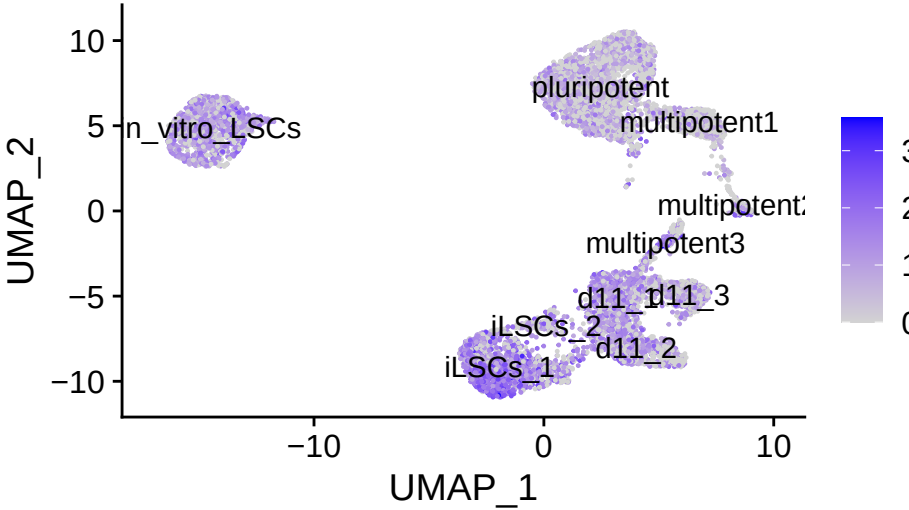
MET



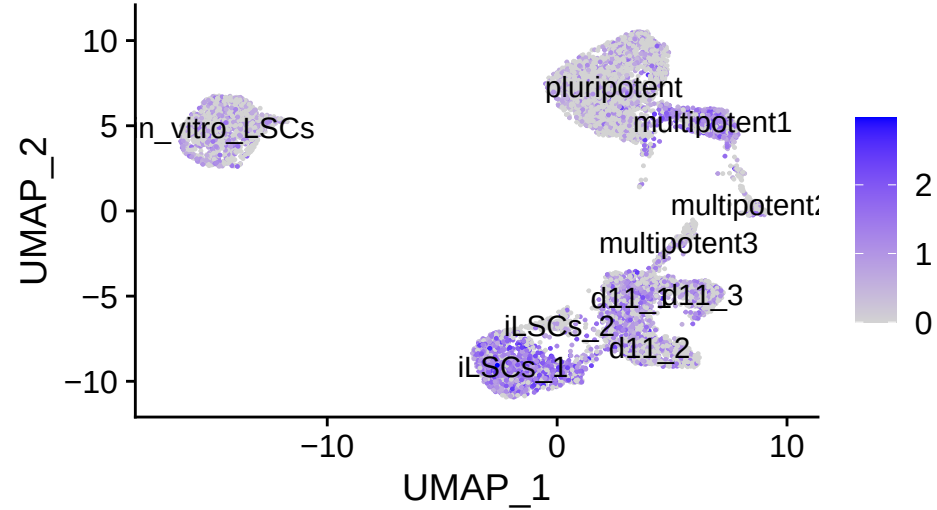
ARHGDIB



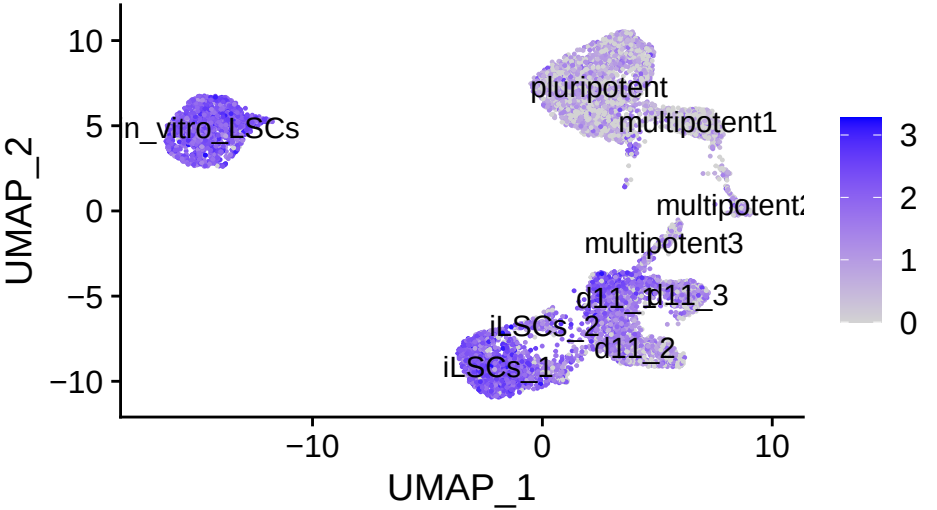
CDC42EP3



PDGFA

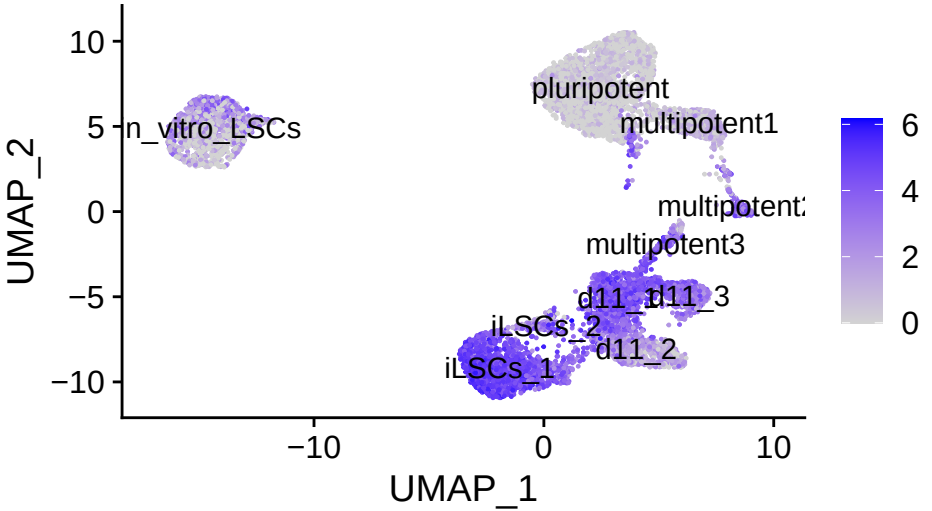


LIMA1

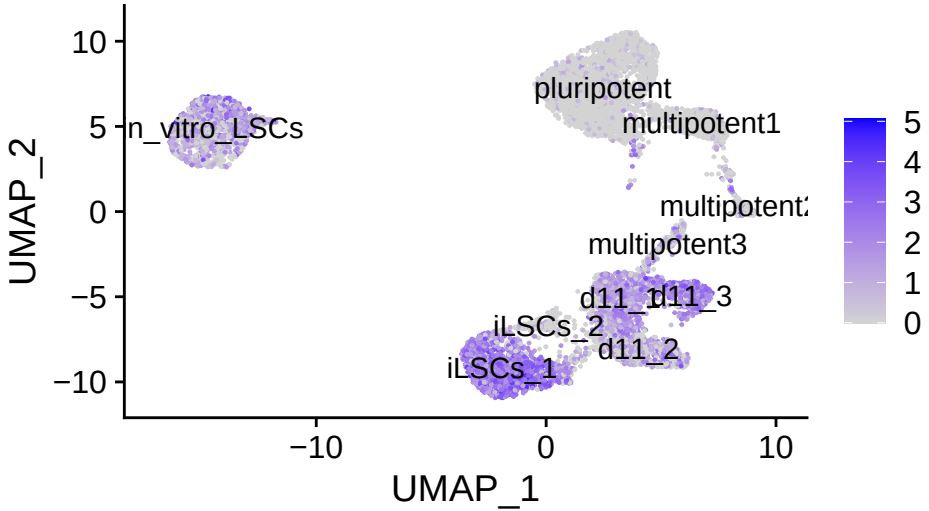


enriched go term: wound healing from cluster: h_iLSCs_1

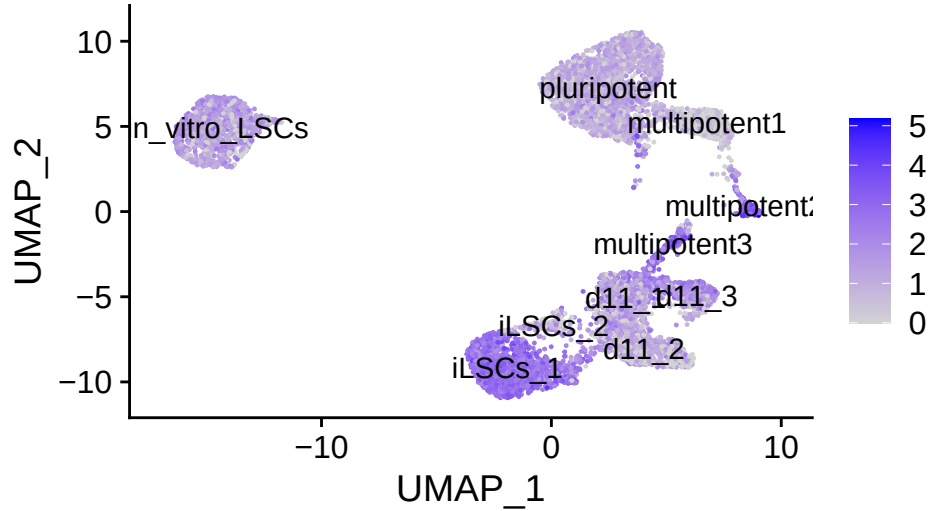
FN1



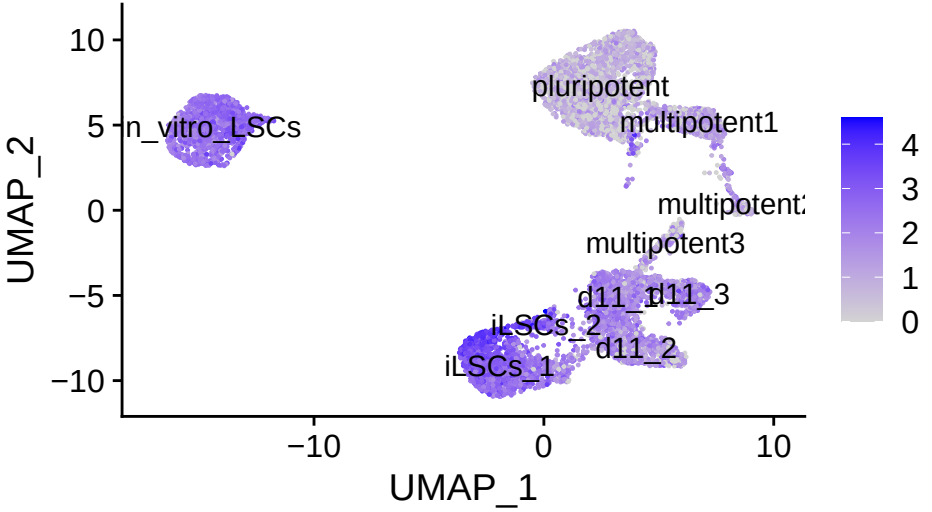
TNC



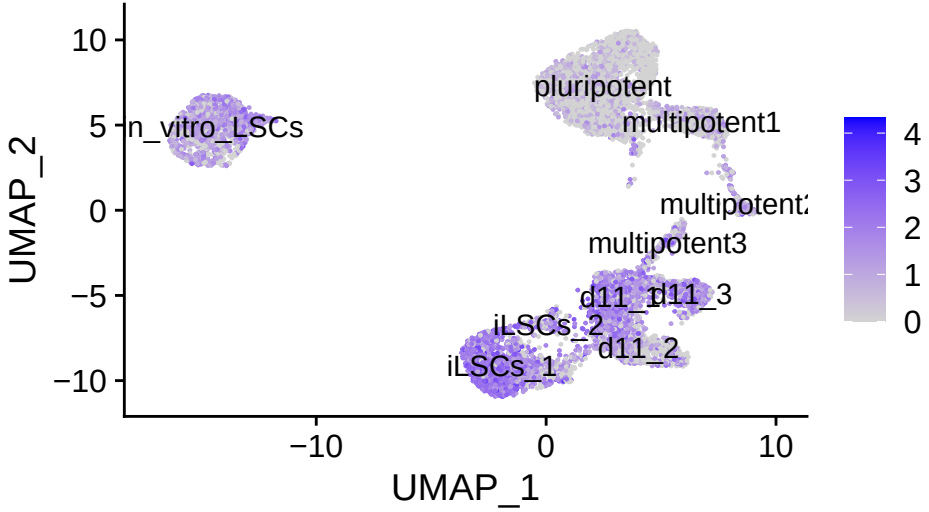
SPARC



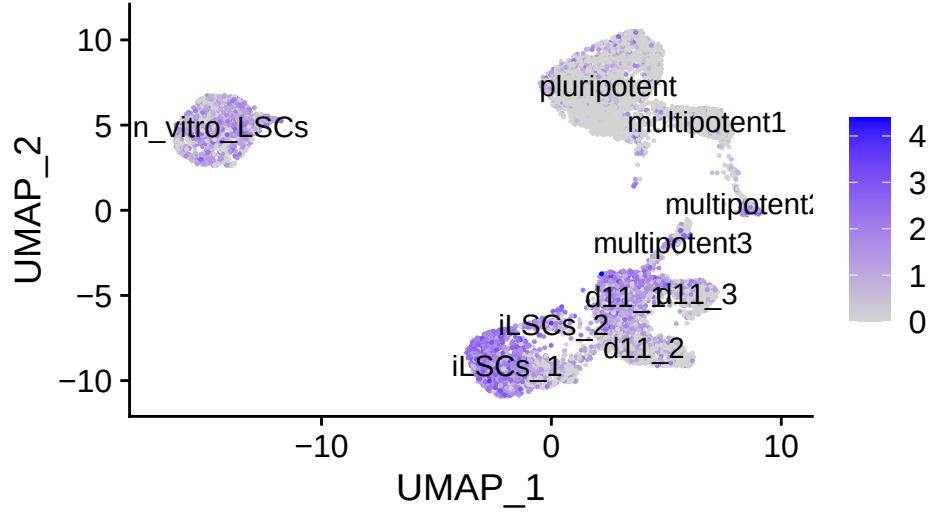
DSP



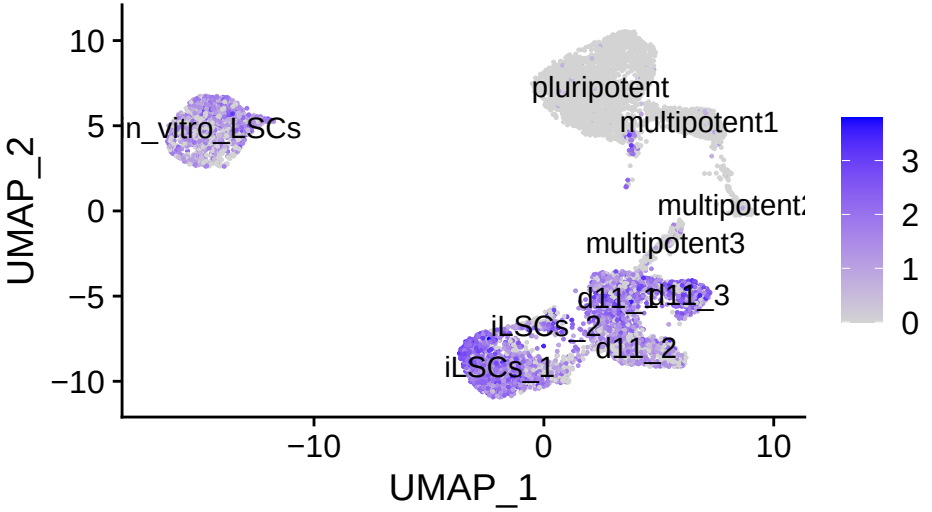
PHLDB2



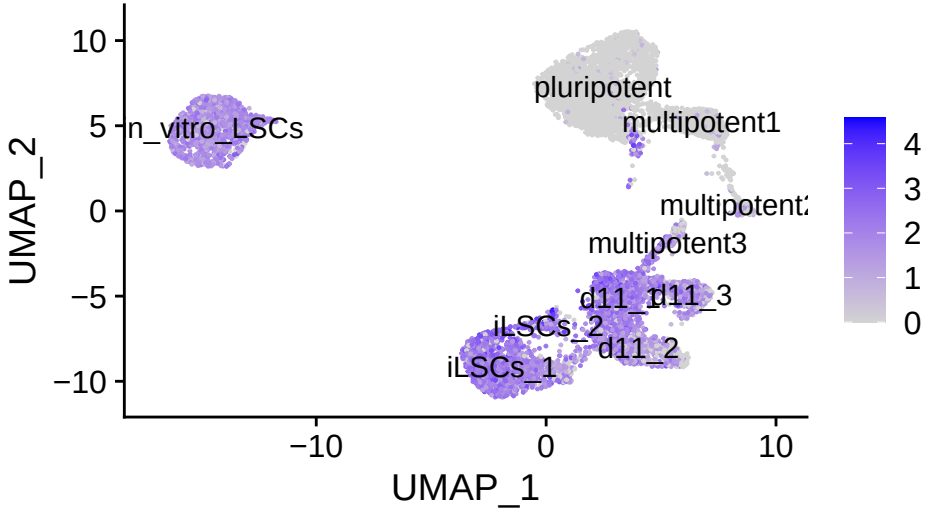
CDKN1A



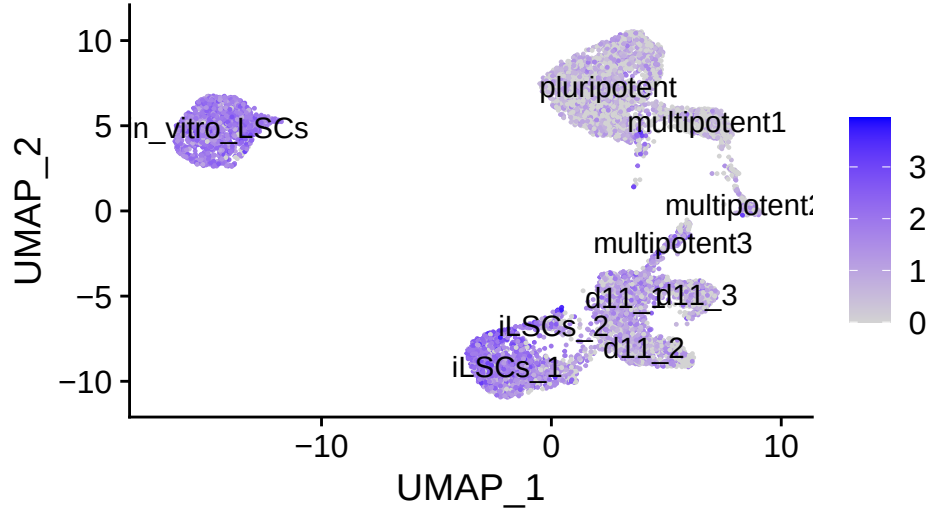
ITGB6



ANXA1

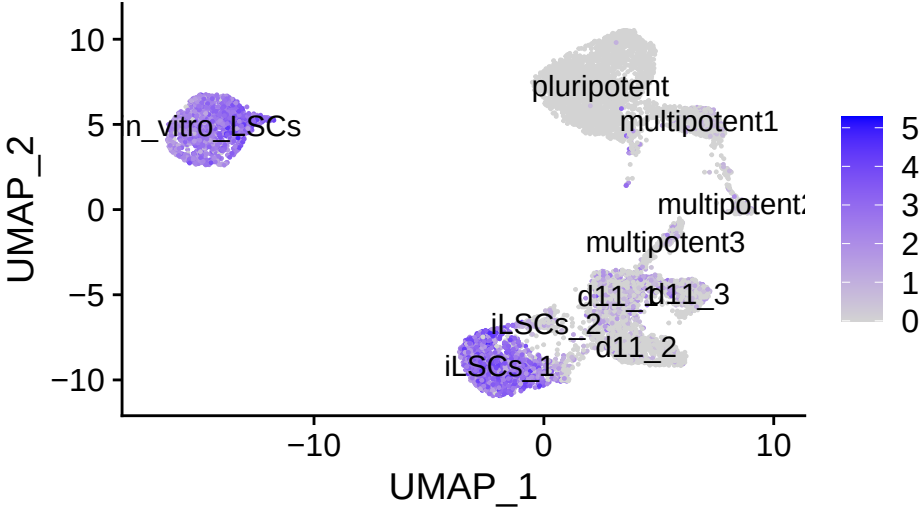


CD59

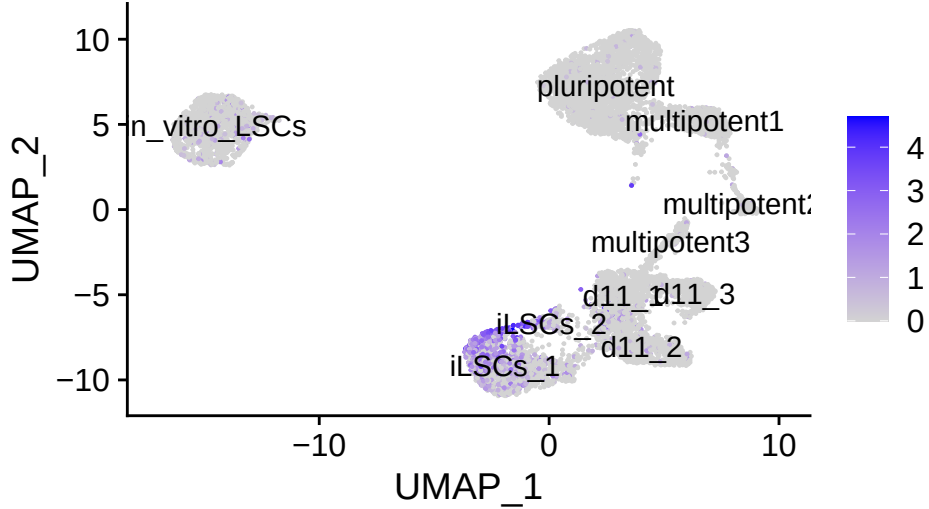


enriched go term: epidermis development from cluster: h_iLSCs_1

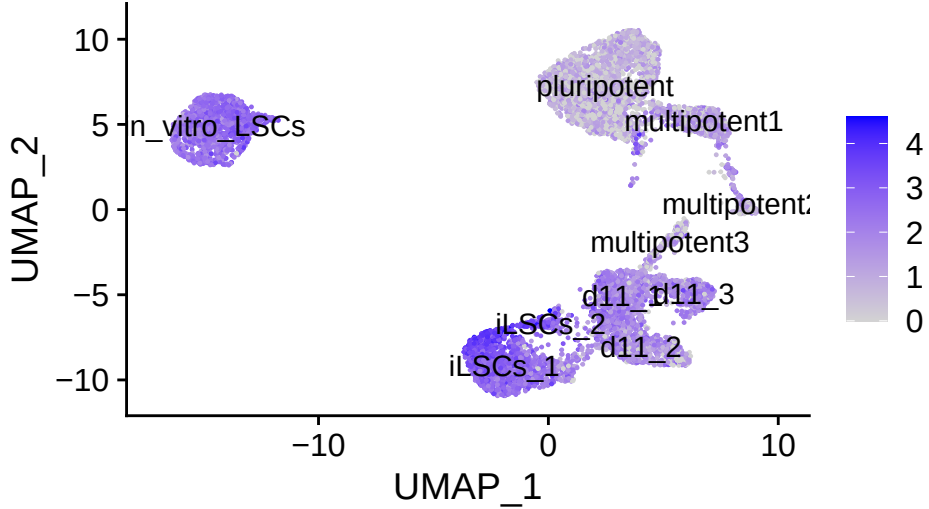
KRT17



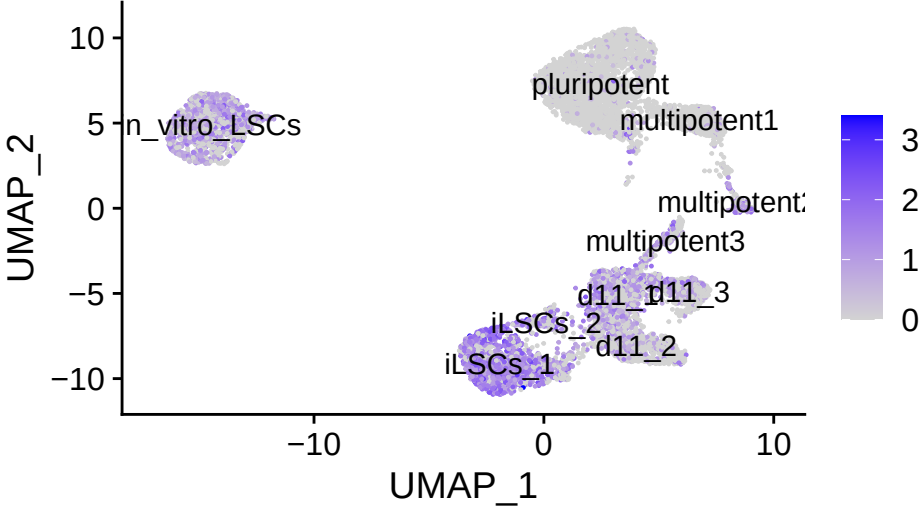
AQP3



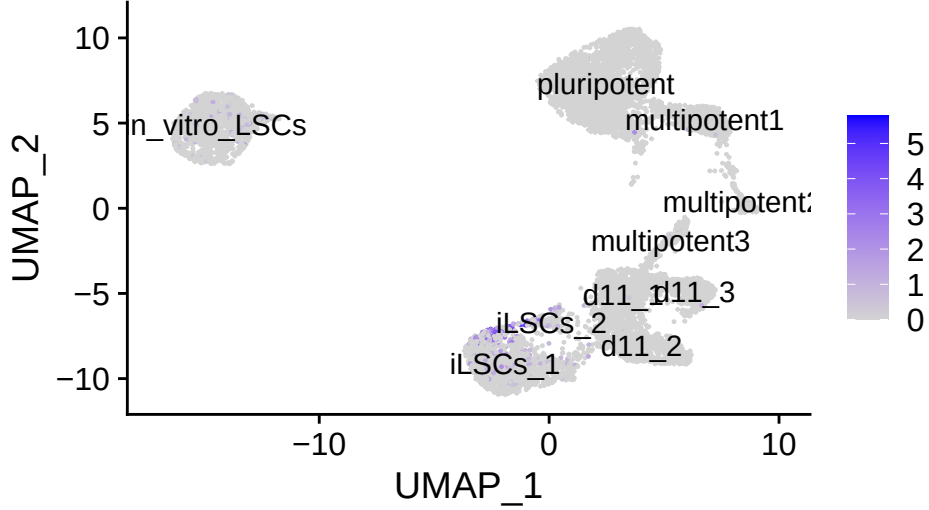
DSP



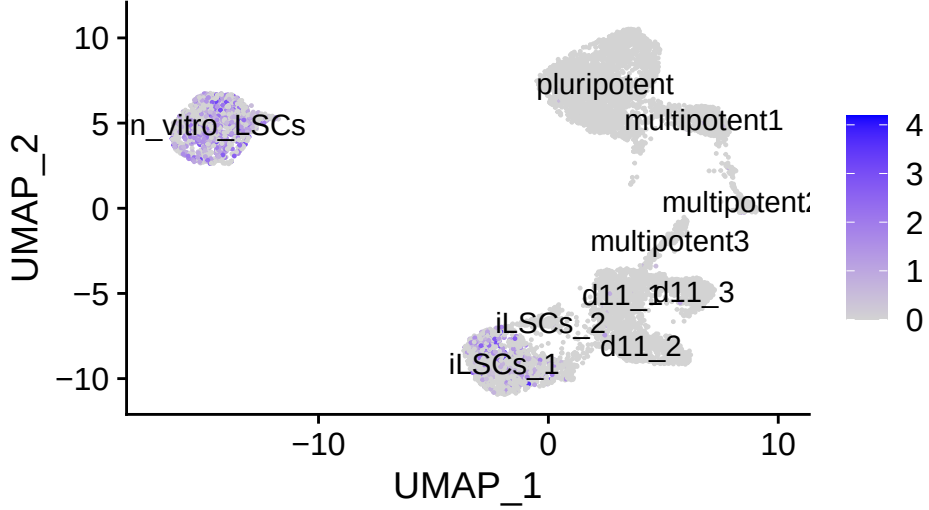
FOSL2



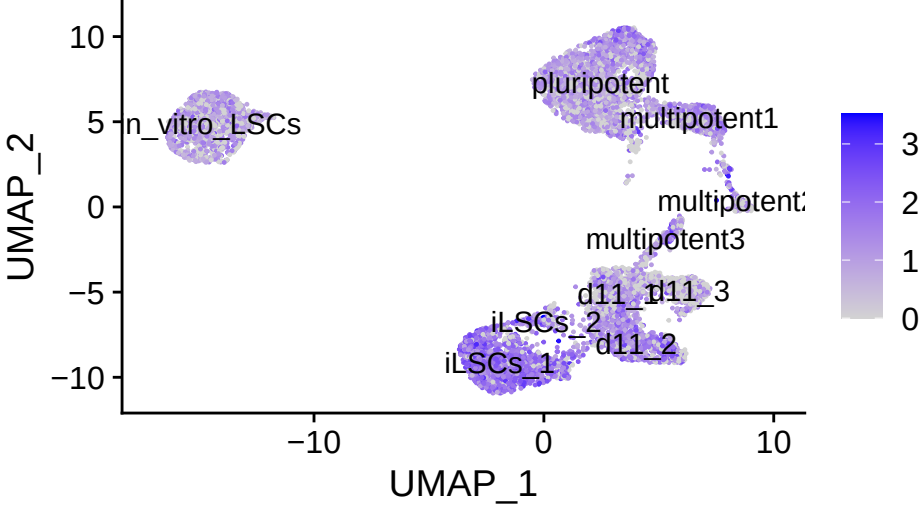
SPRR1B



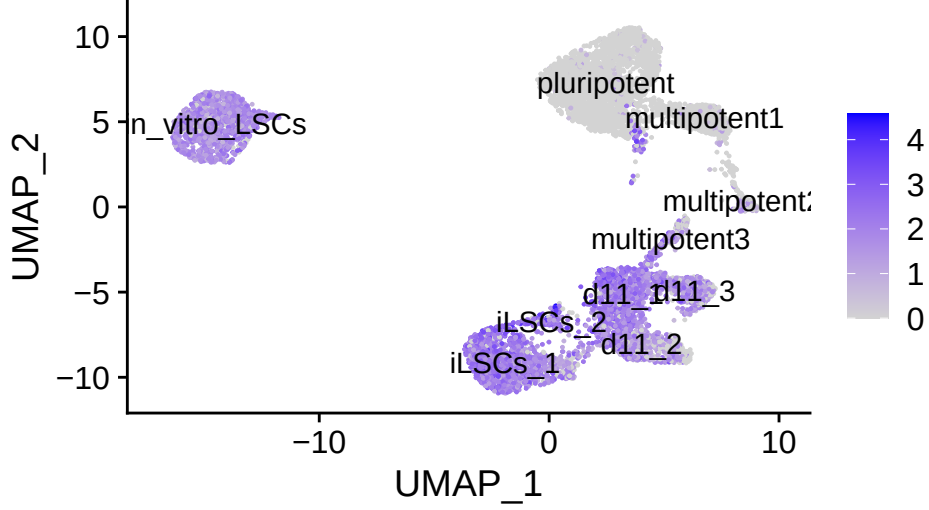
KRT15



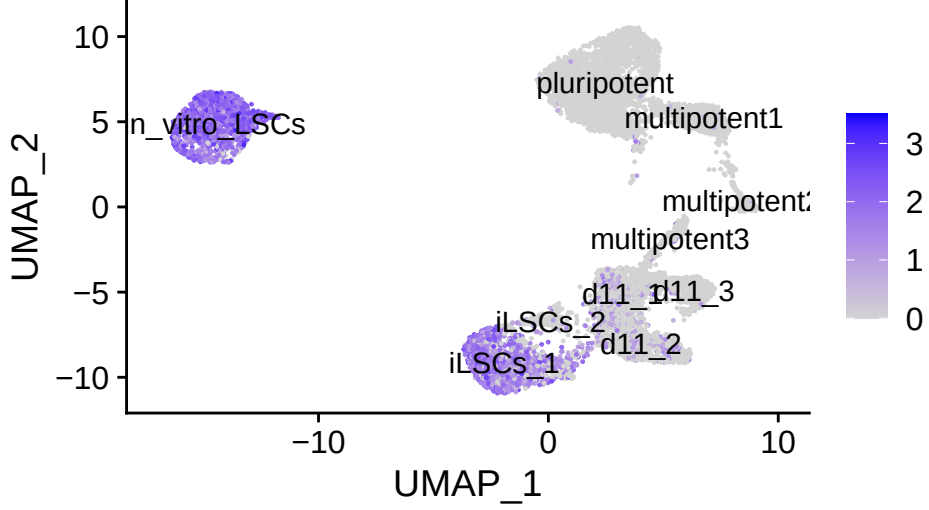
LGR4



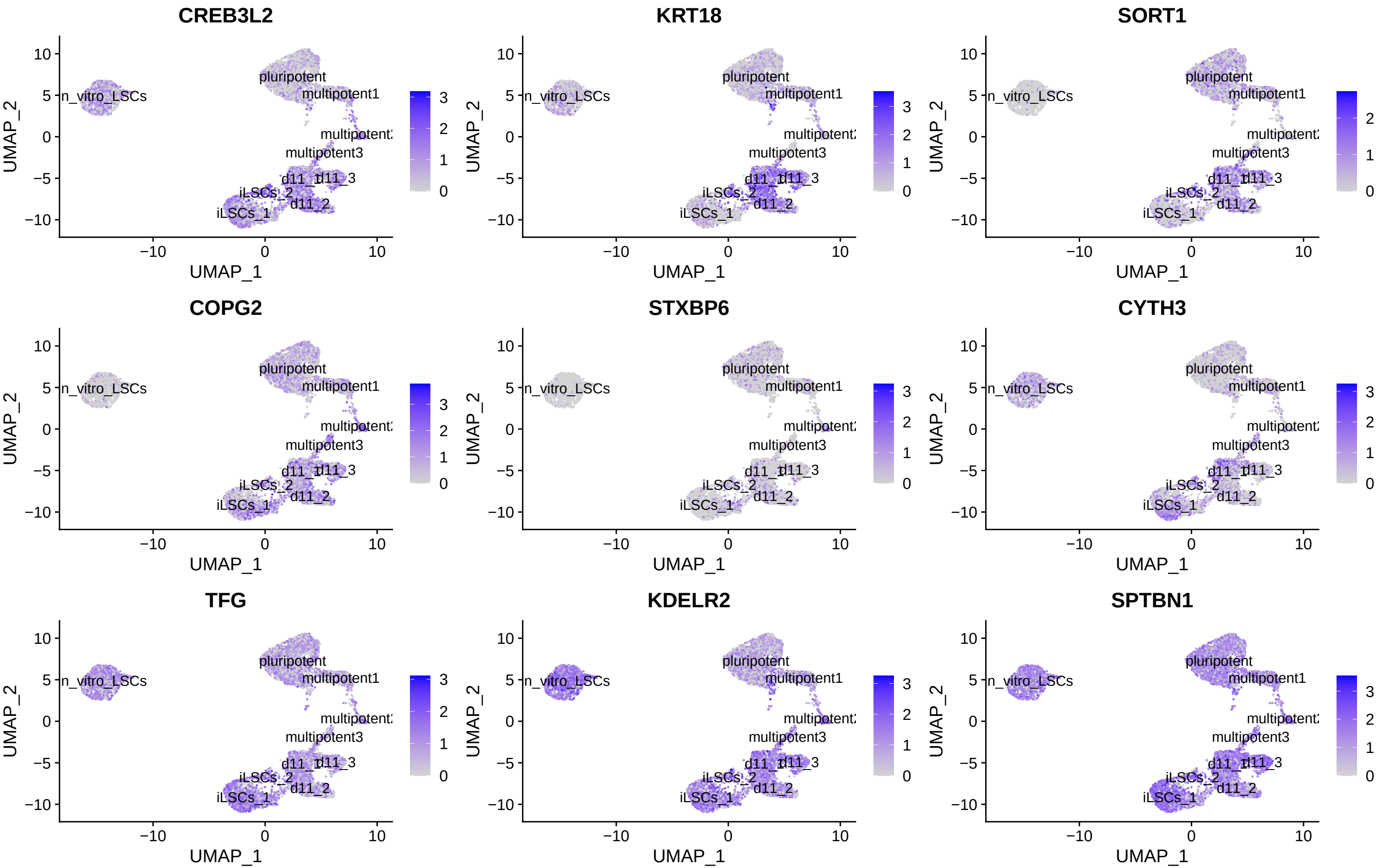
ANXA1



COL17A1

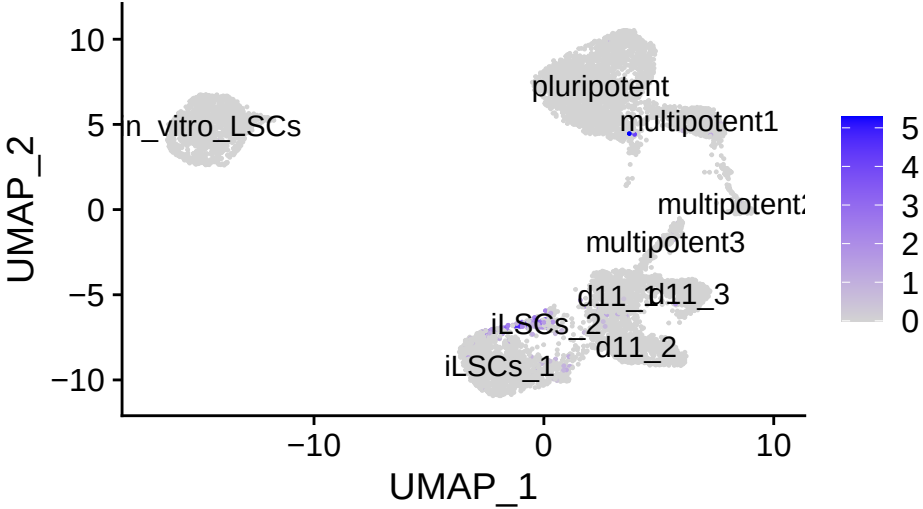


enriched go term: Golgi vesicle transport from cluster: i_iLSCs_2

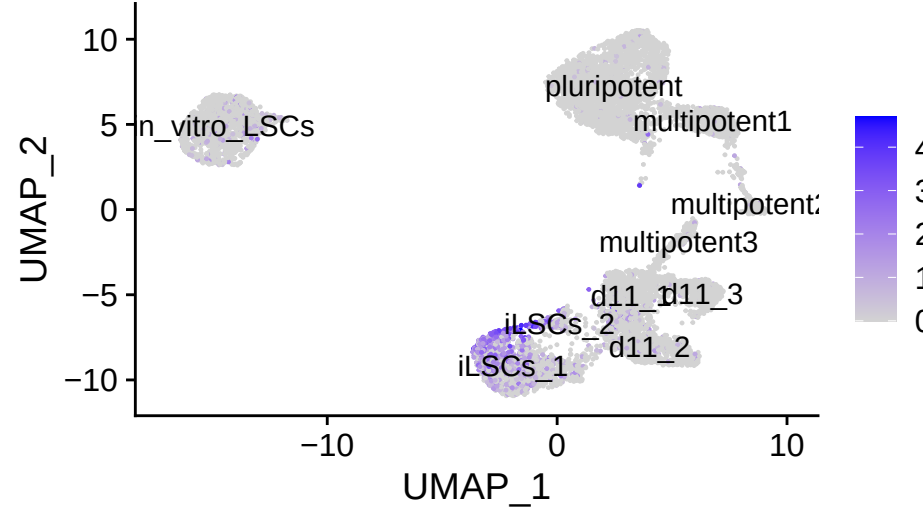


enriched go term: epidermal cell differentiation from cluster: i_iLSCs_2

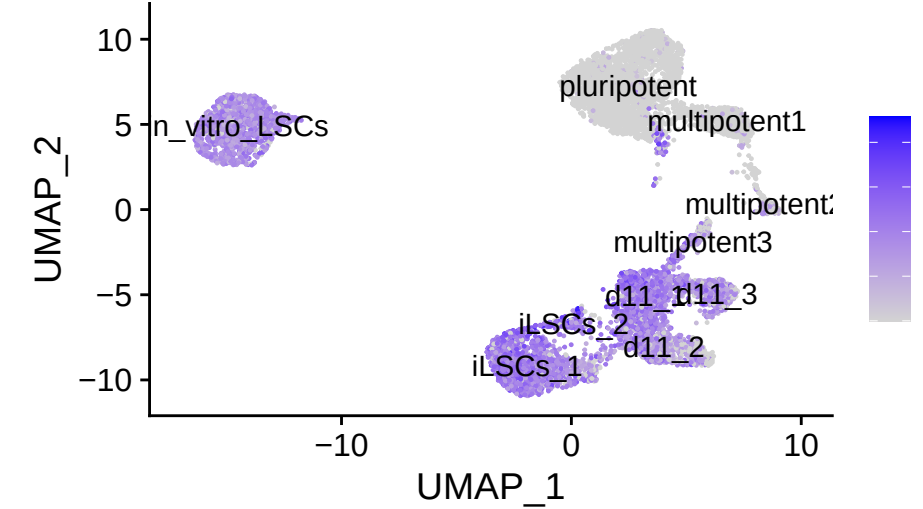
SPRR3



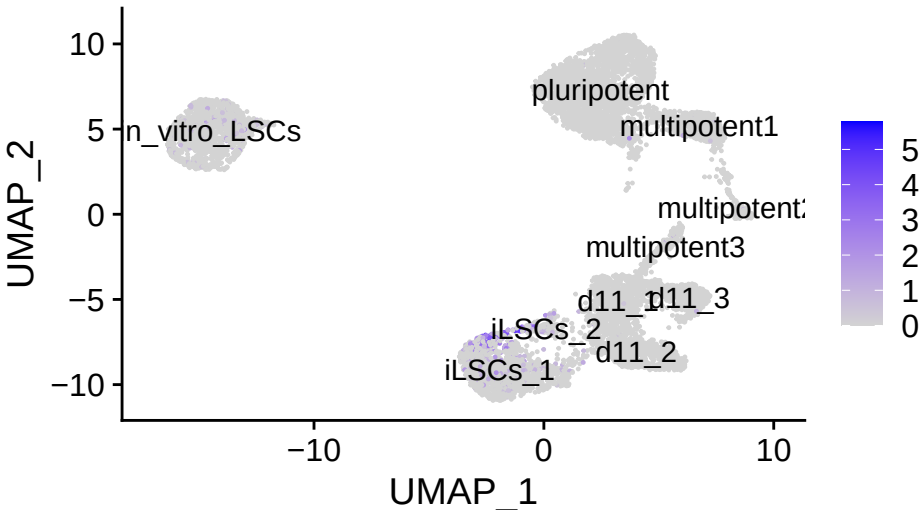
AQP3



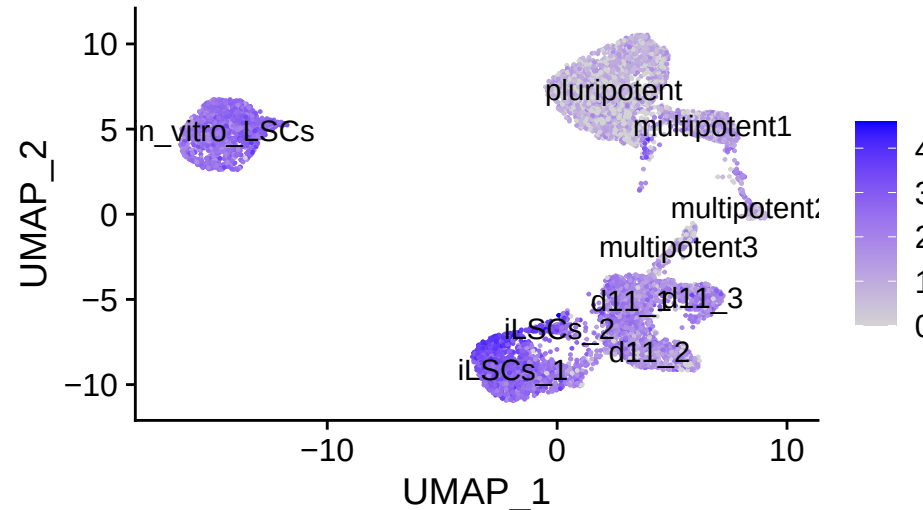
ANXA1



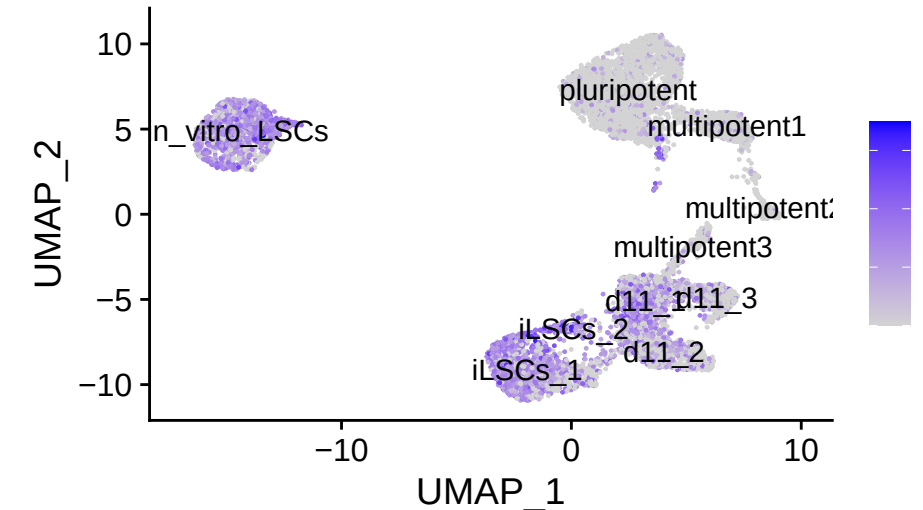
SPRR1B



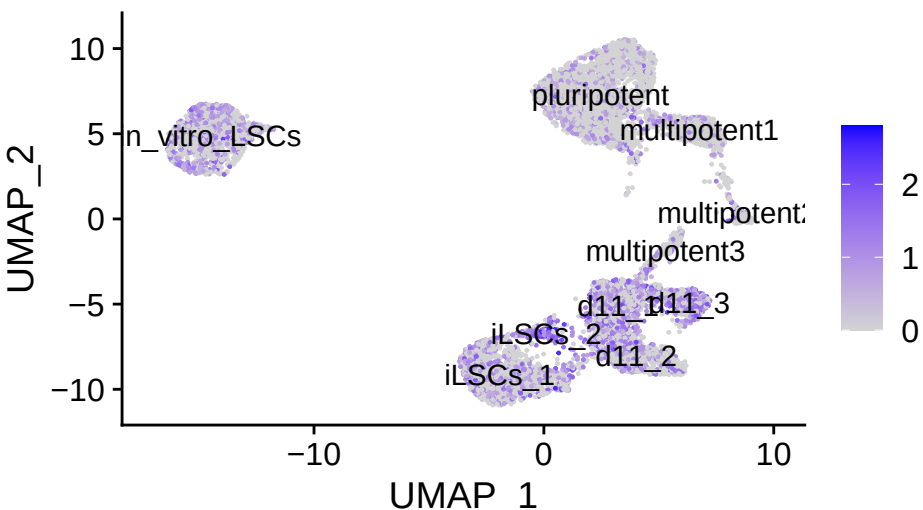
DSP



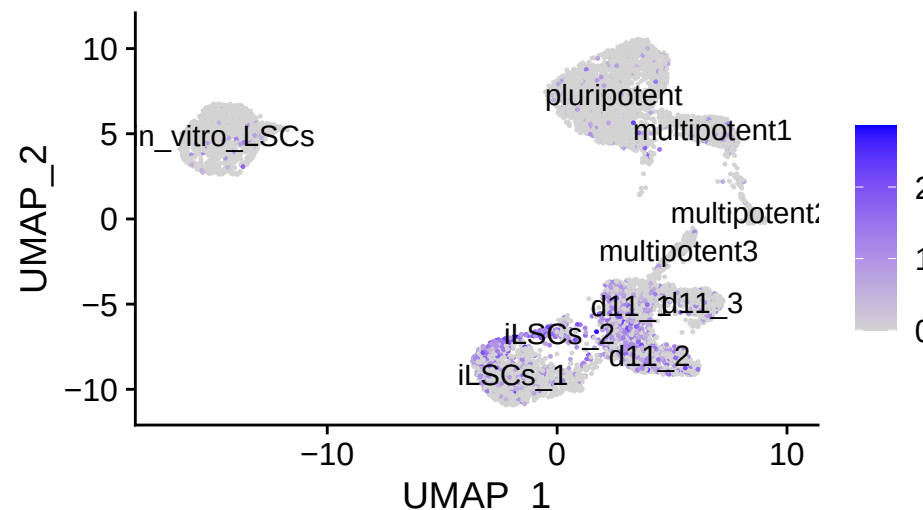
IRF6



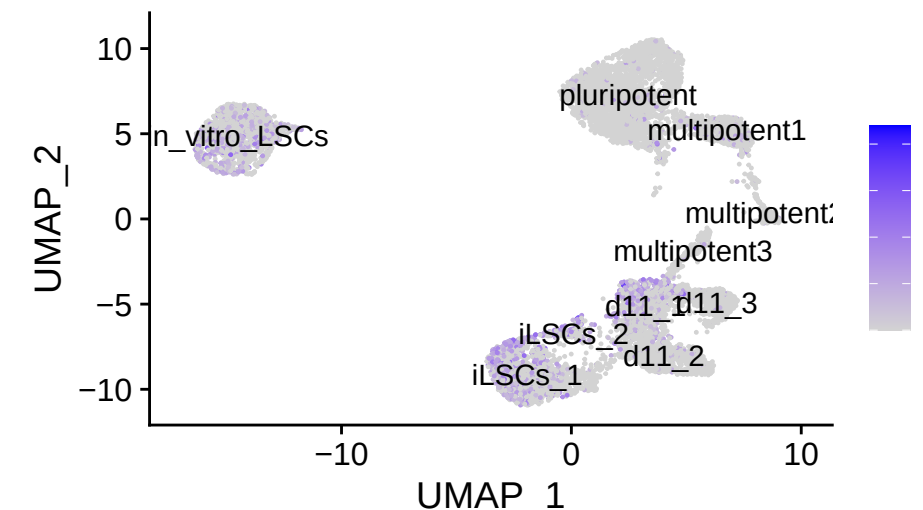
PRKCH



GRHL1

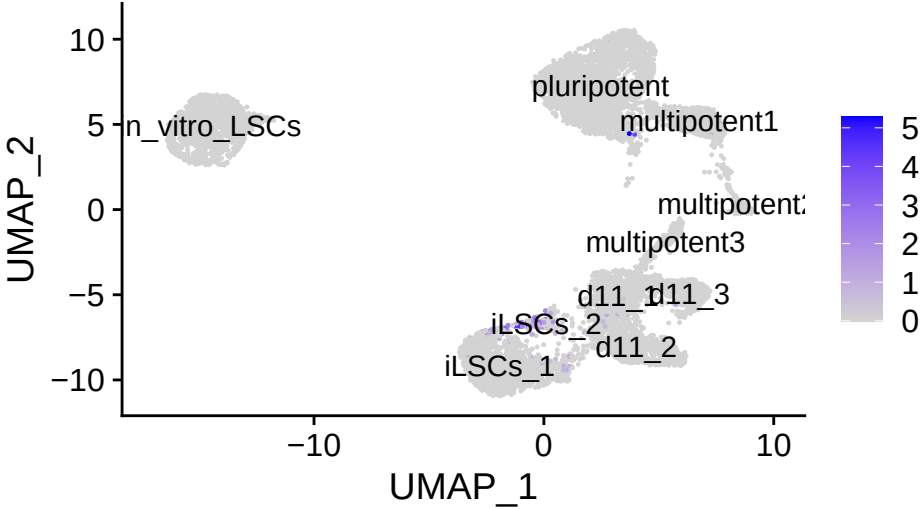


KLF4

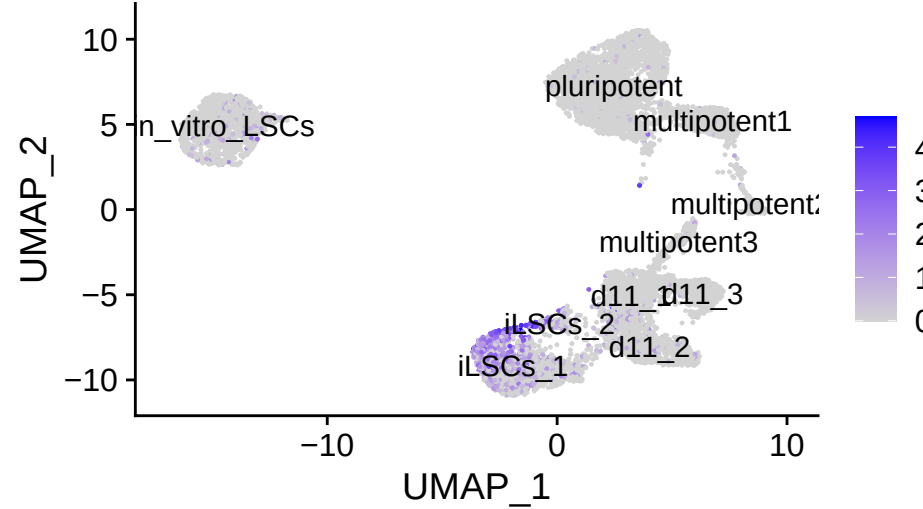


enriched go term: epidermis development from cluster: i_iLSCs_2

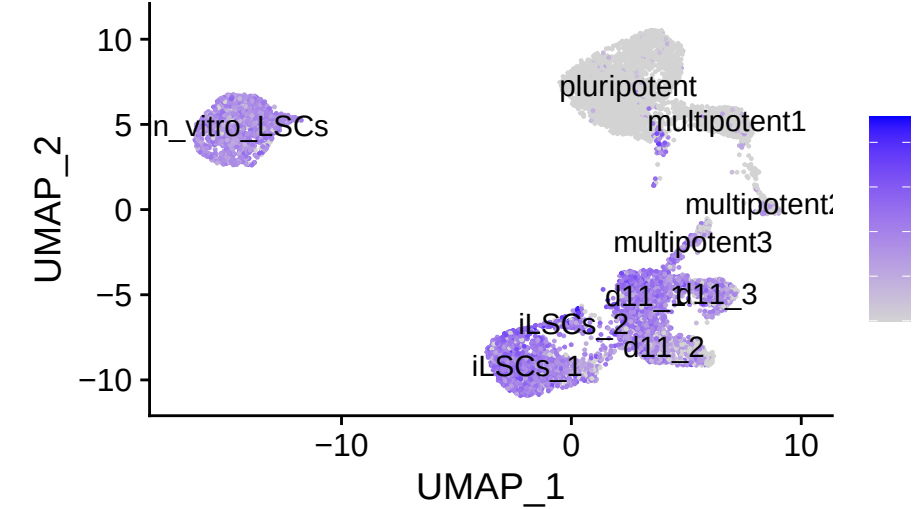
SPRR3



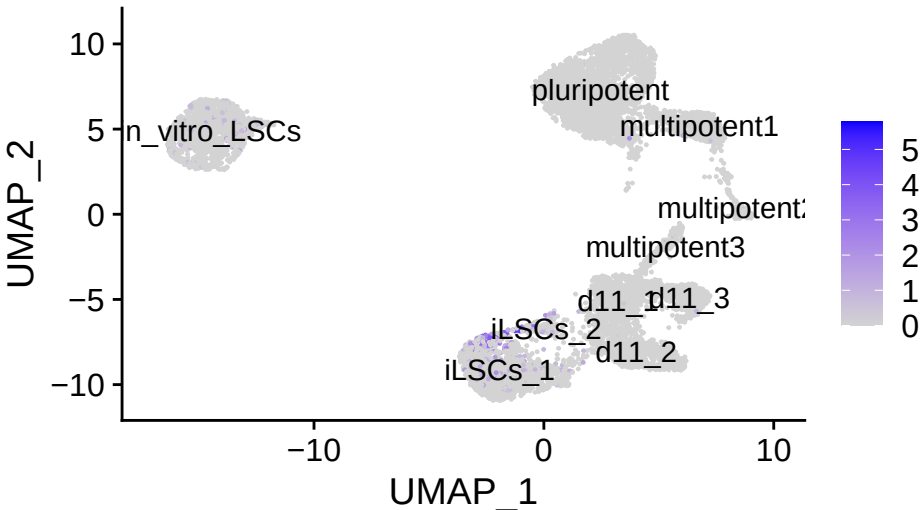
AQP3



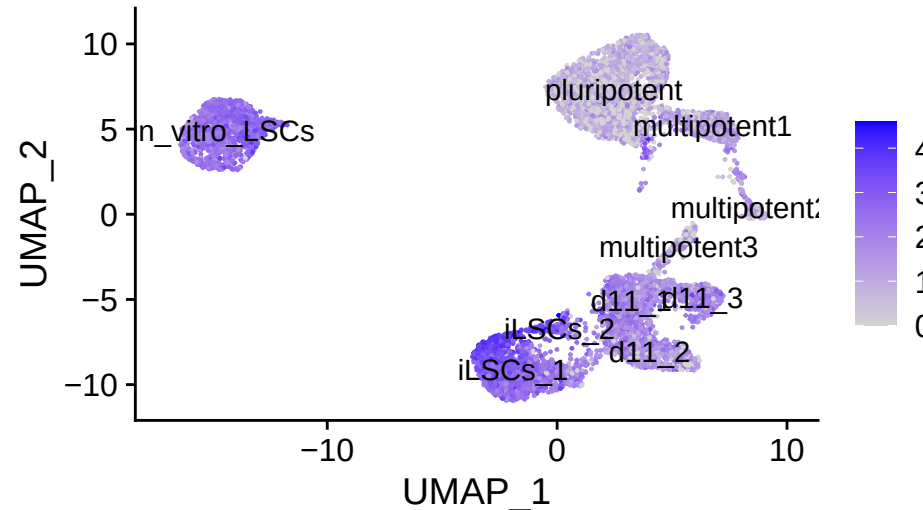
ANXA1



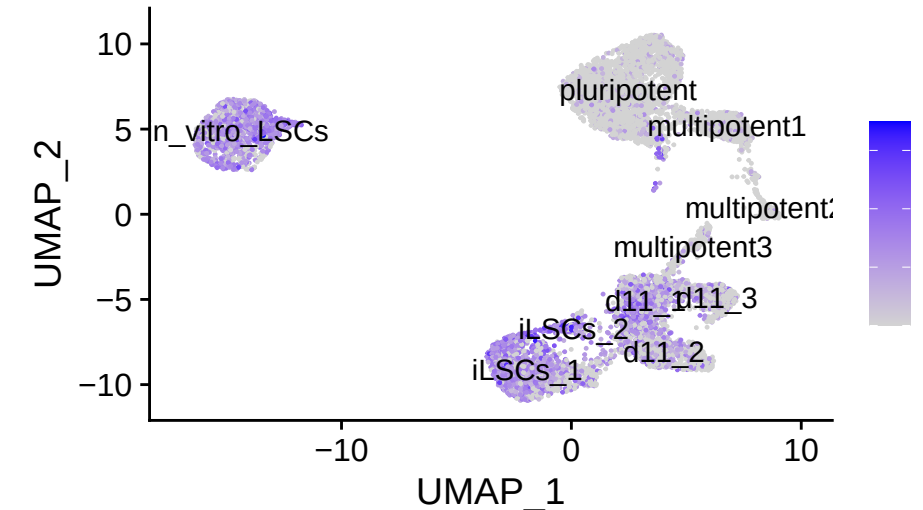
SPRR1B



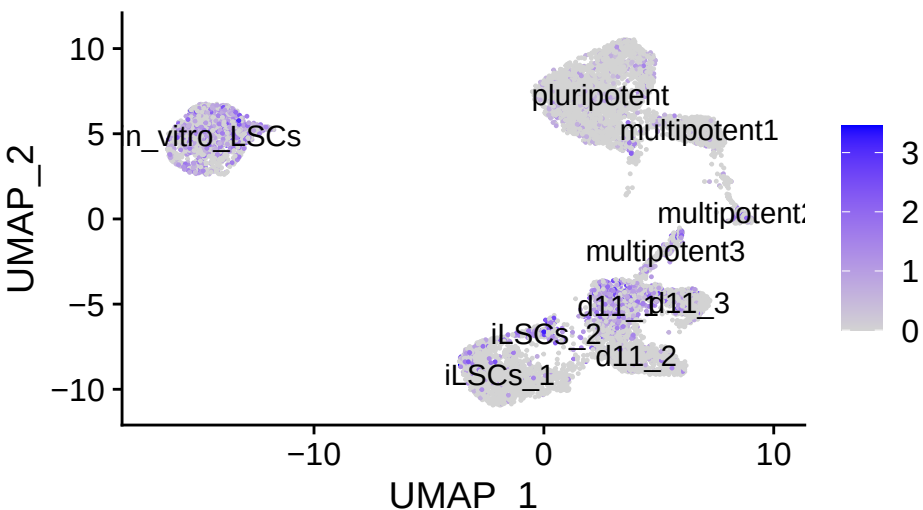
DSP



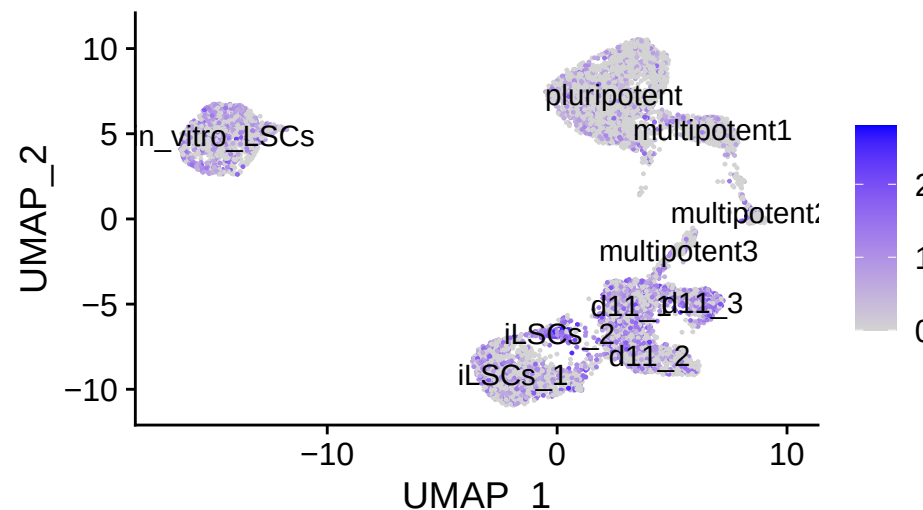
IRF6



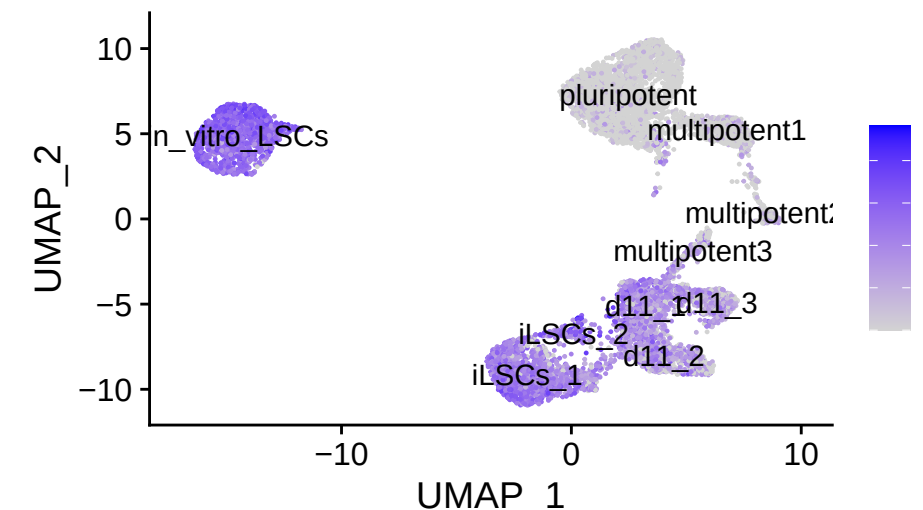
SOX9



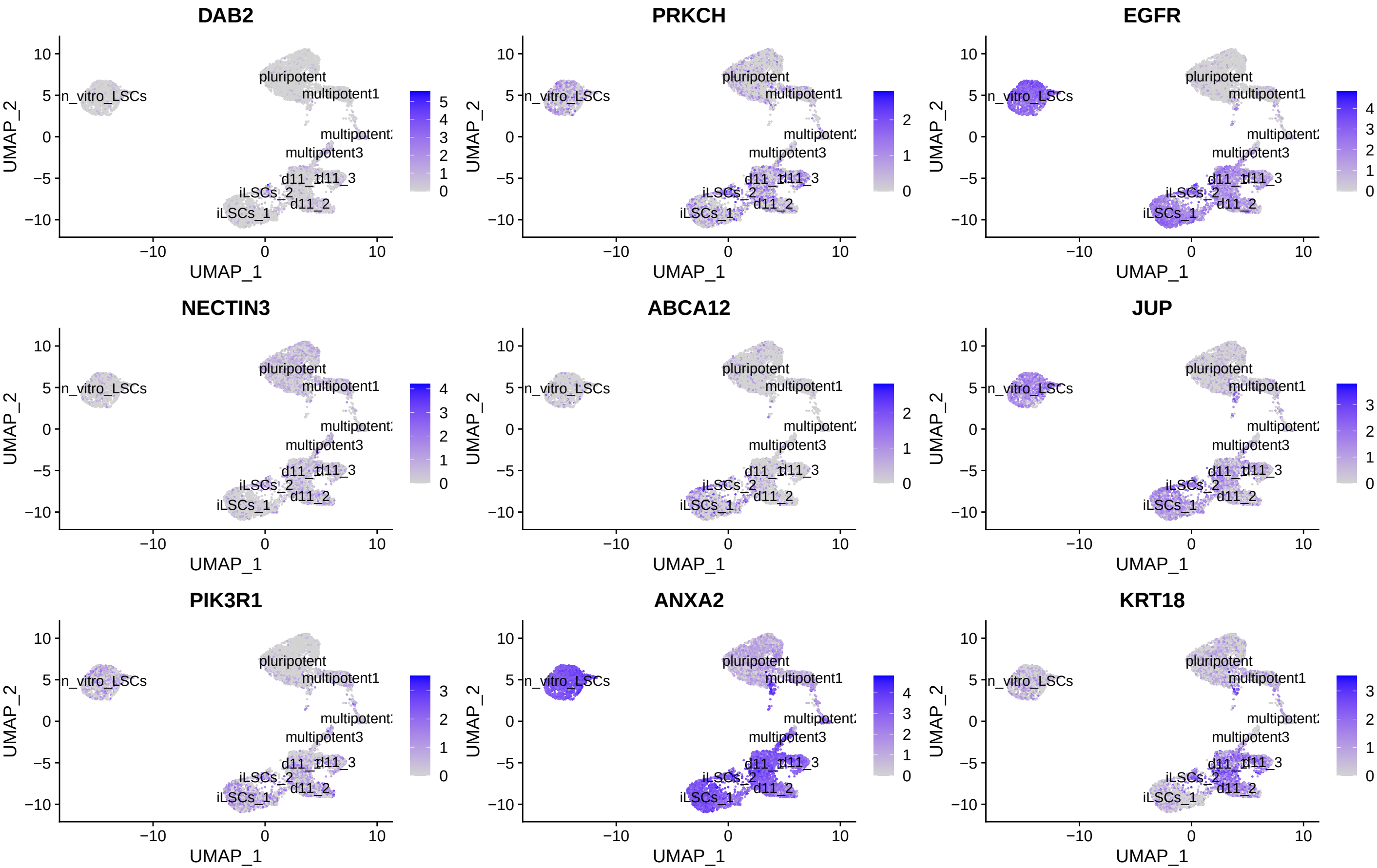
PRKCH



EGFR

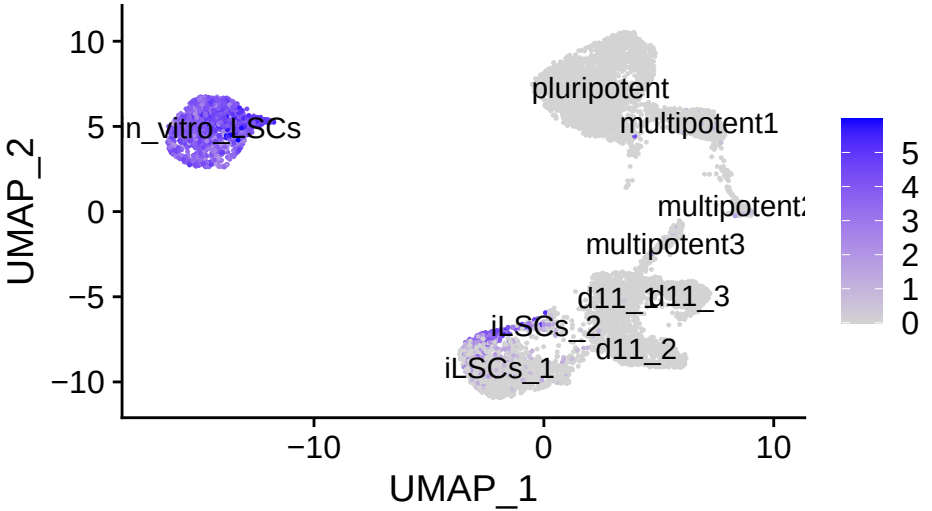


enriched go term: protein localization to cell periphery from cluster: i_iLSCs_2

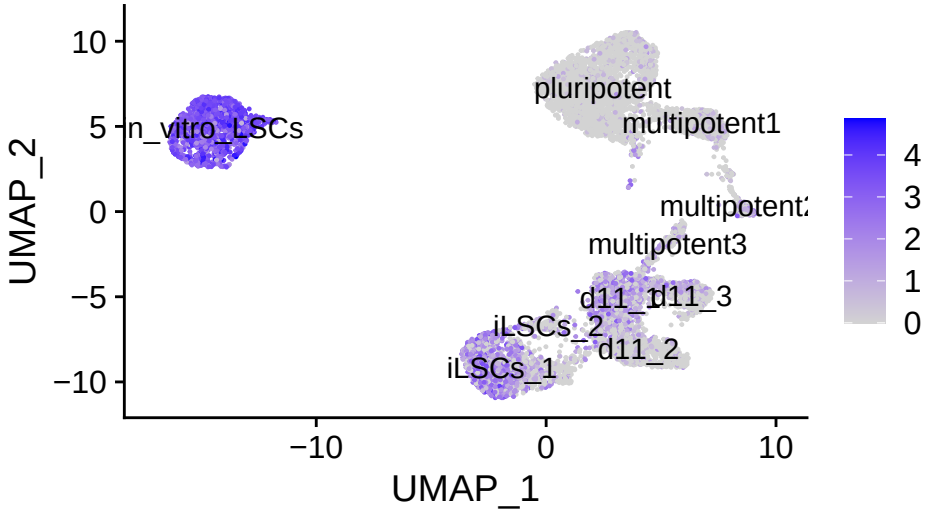


enriched go term: wound healing from cluster: j_in_vitro_LSCs

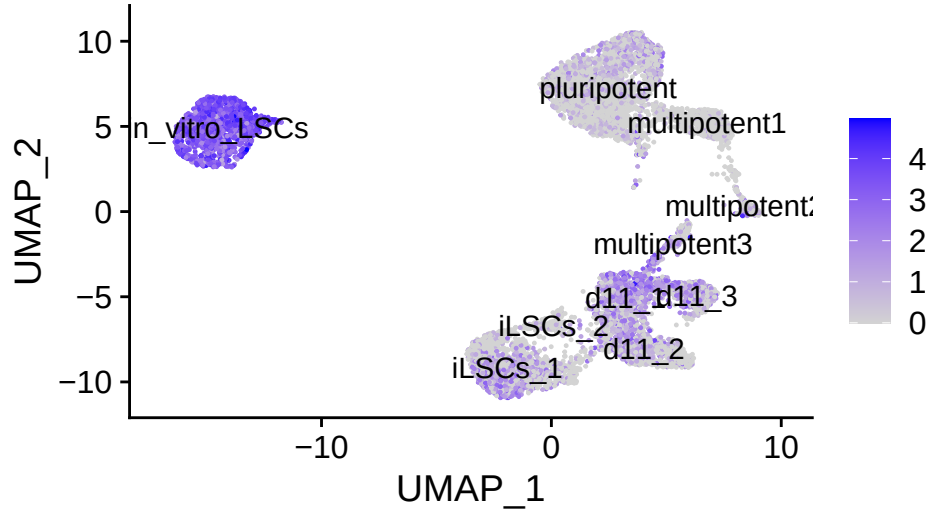
KRT6A



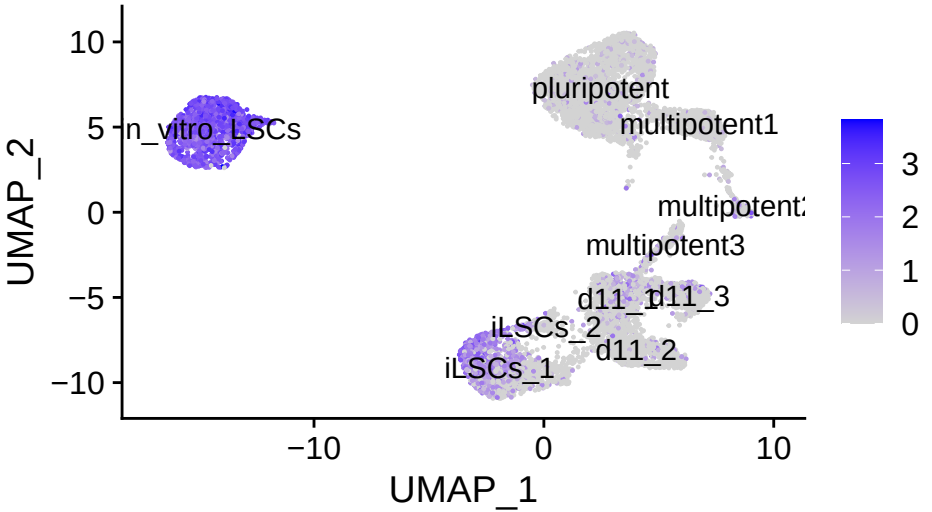
F3



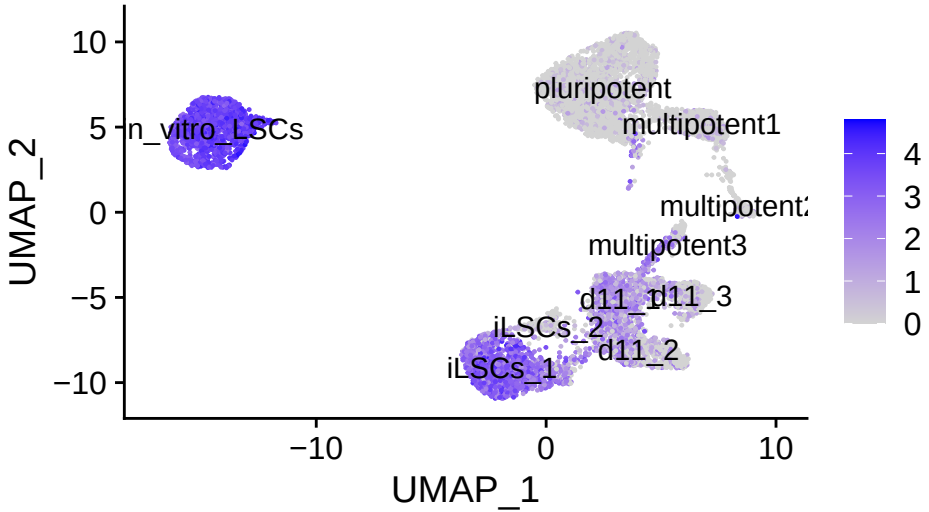
THBS1



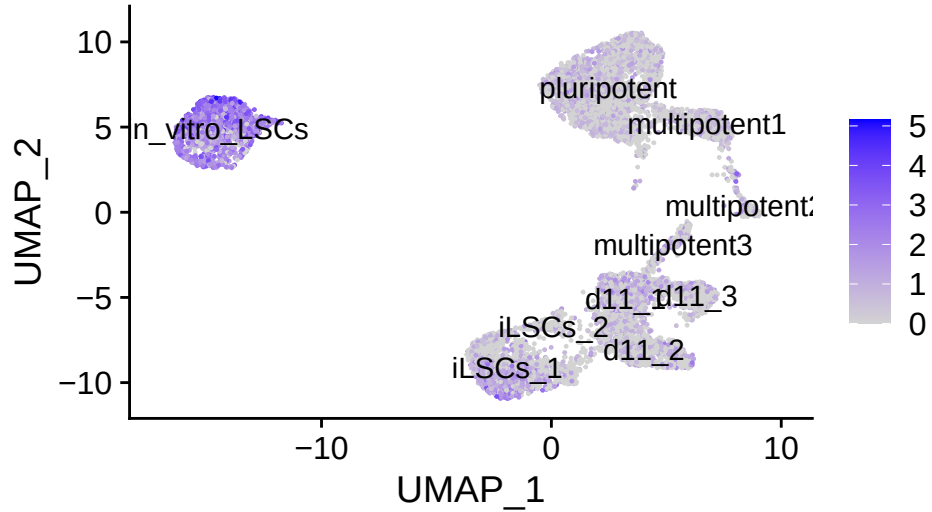
CD44



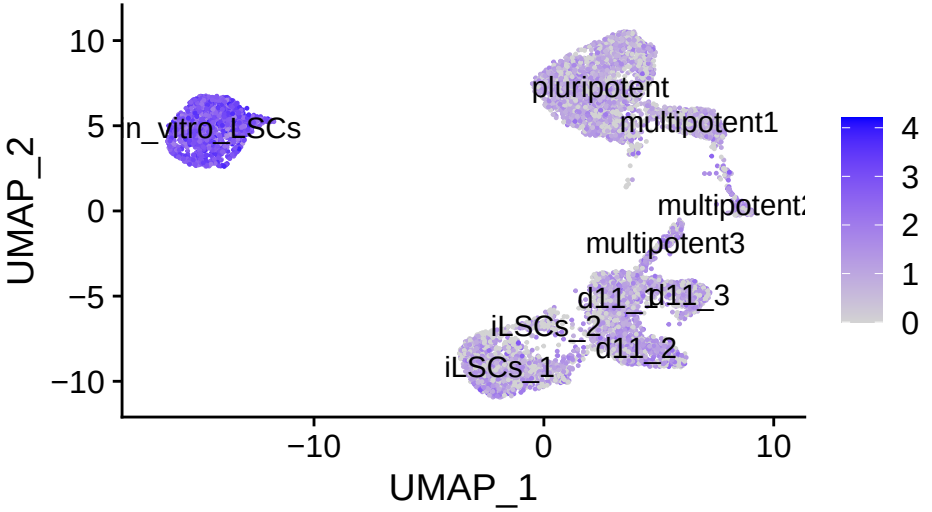
CAV1



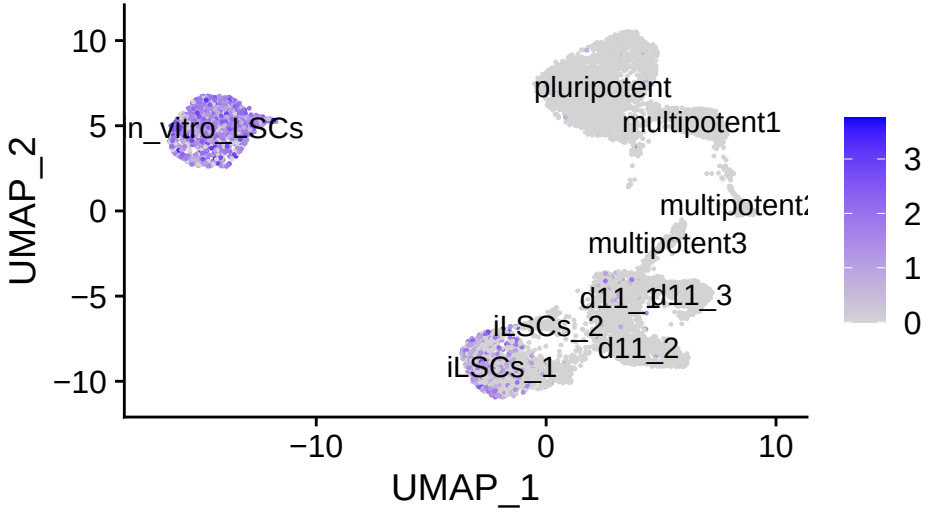
NRG1



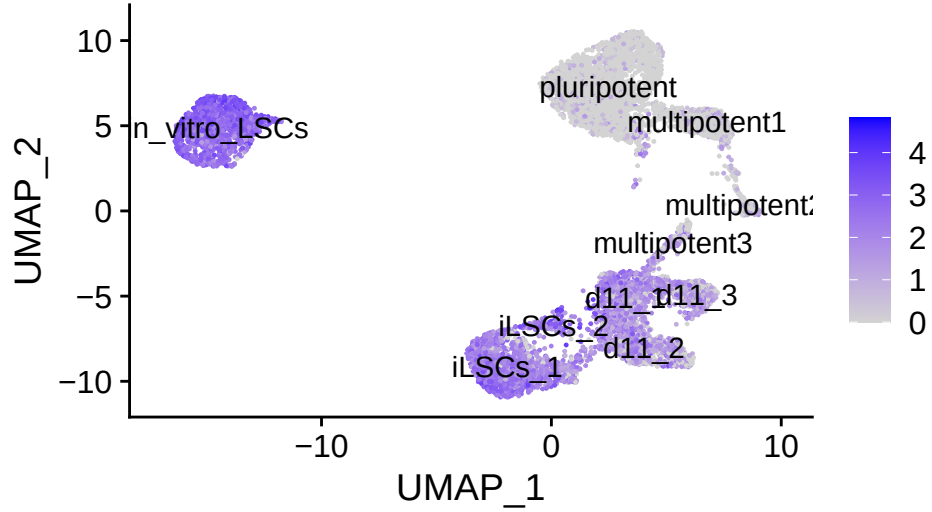
DCBLD2



IL1A

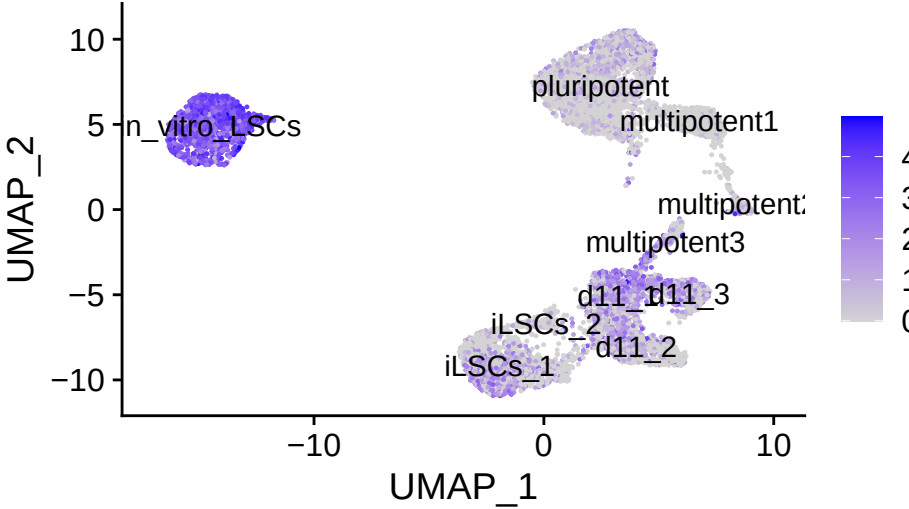


EGFR

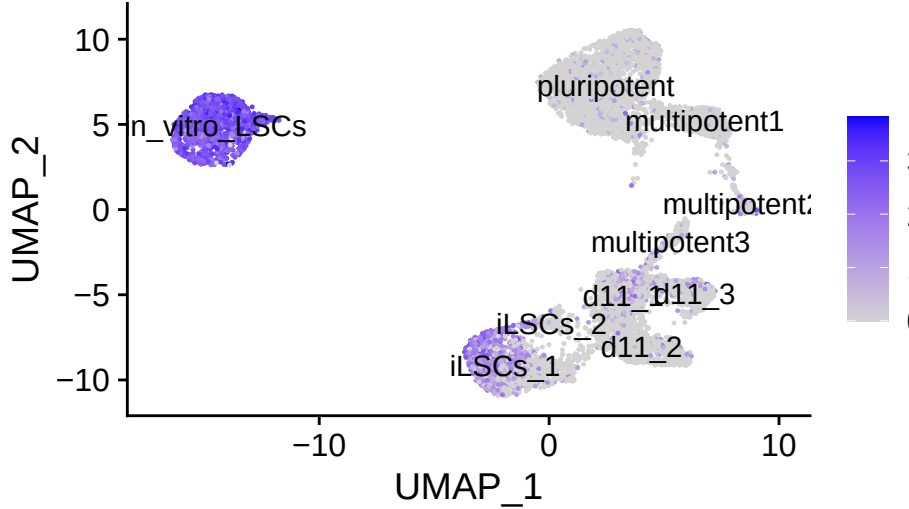


enriched go term: cell-substrate adhesion from cluster: j_in_vitro_LSCs

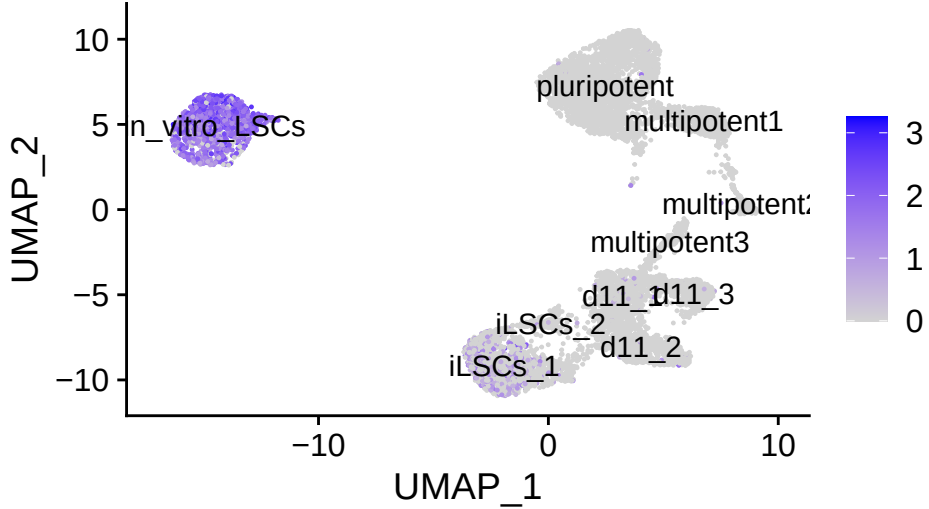
THBS1



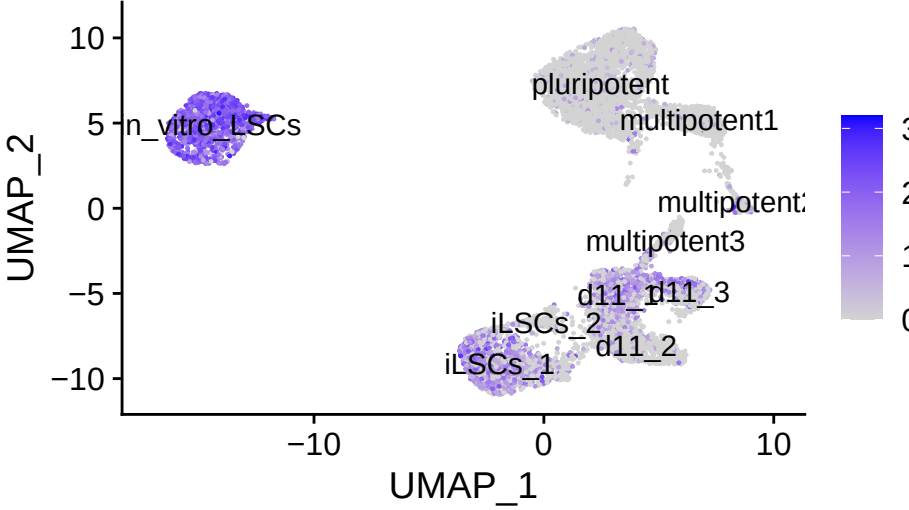
CD44



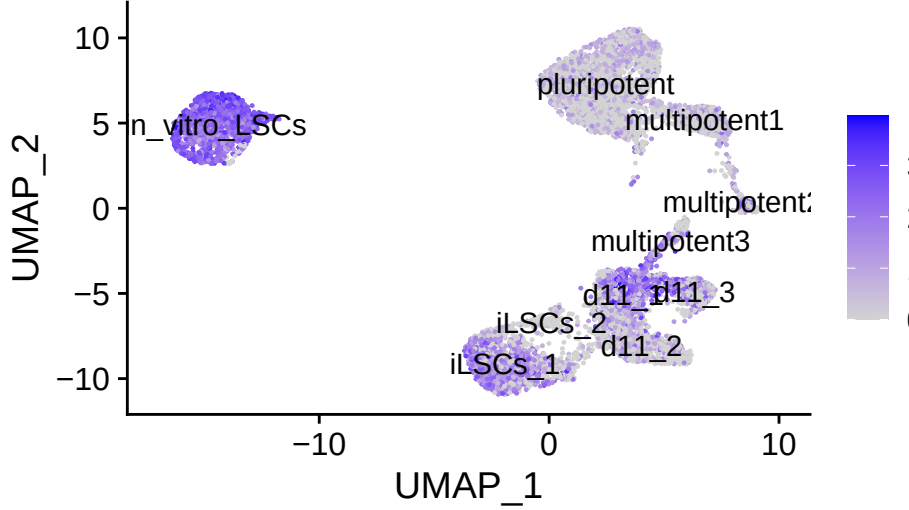
LAMB3



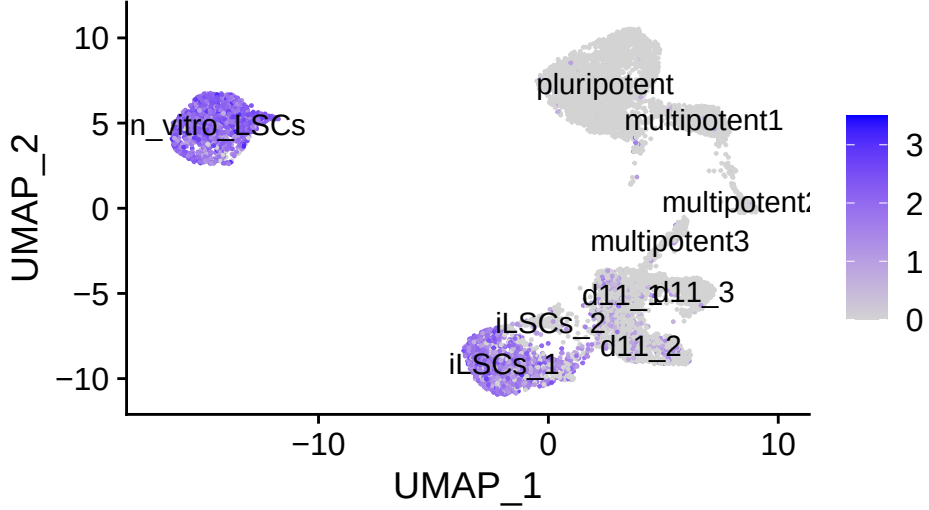
ITGA3



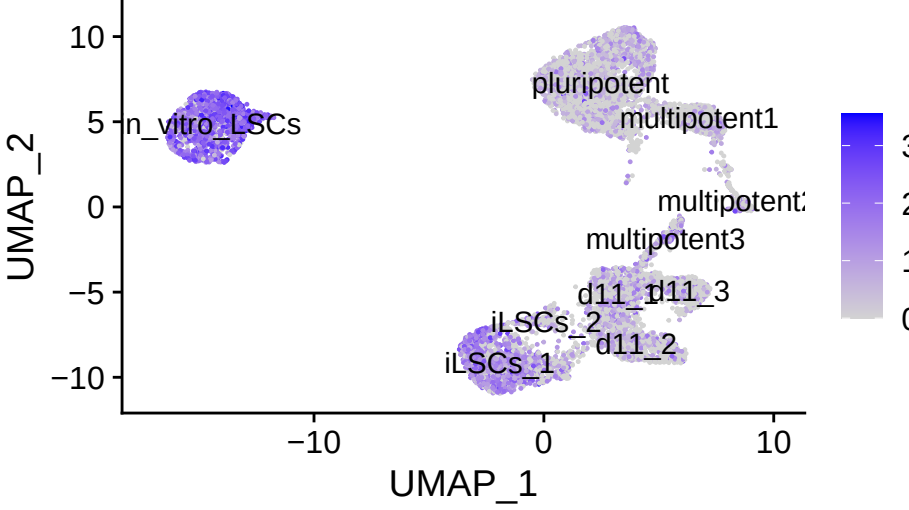
CDH13



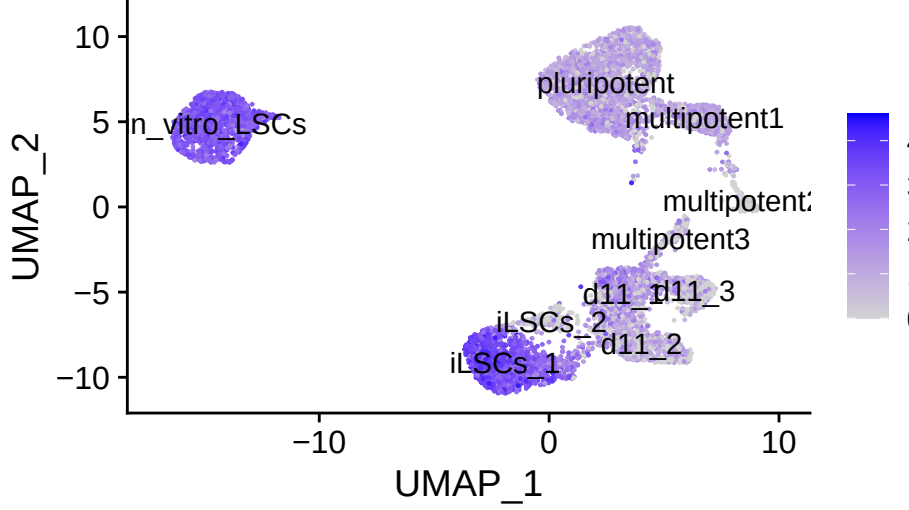
COL17A1



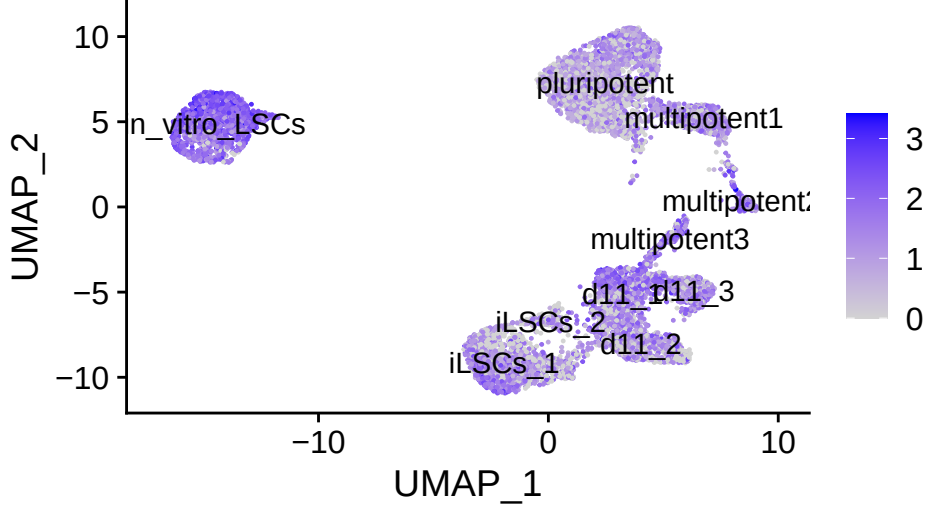
JAG1



ITGA6

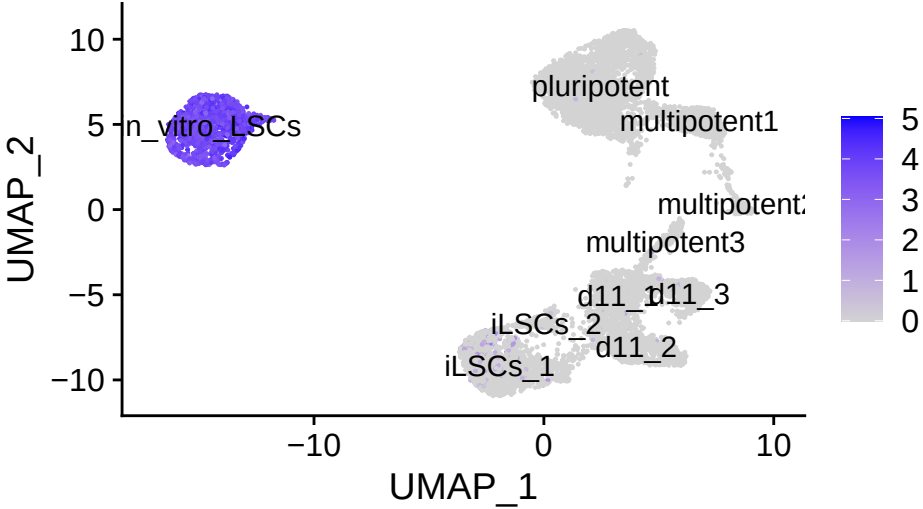


CDK6

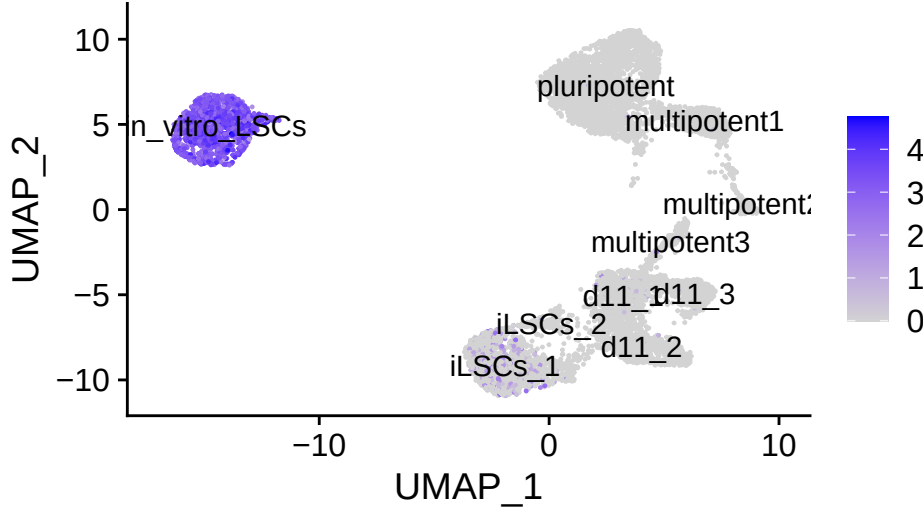


enriched go term: ameboidal-type cell migration from cluster: j_in_vitro_LSCs

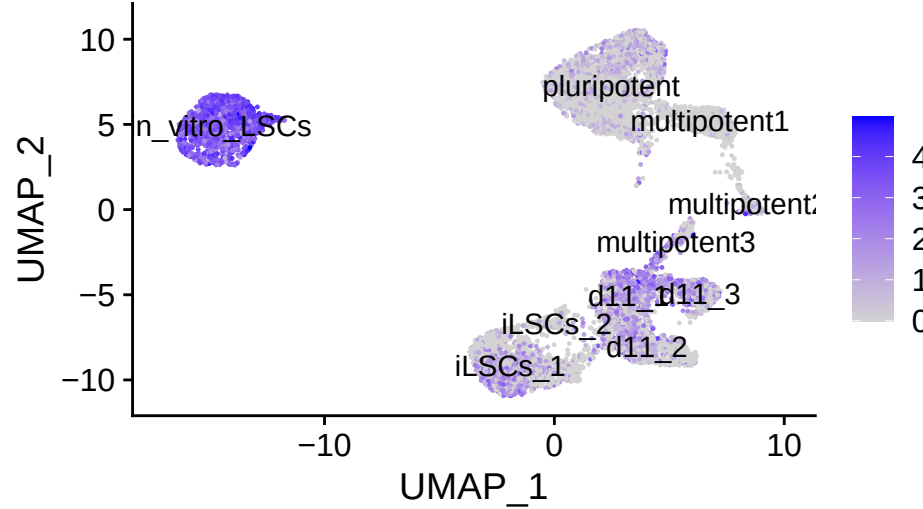
S100A2



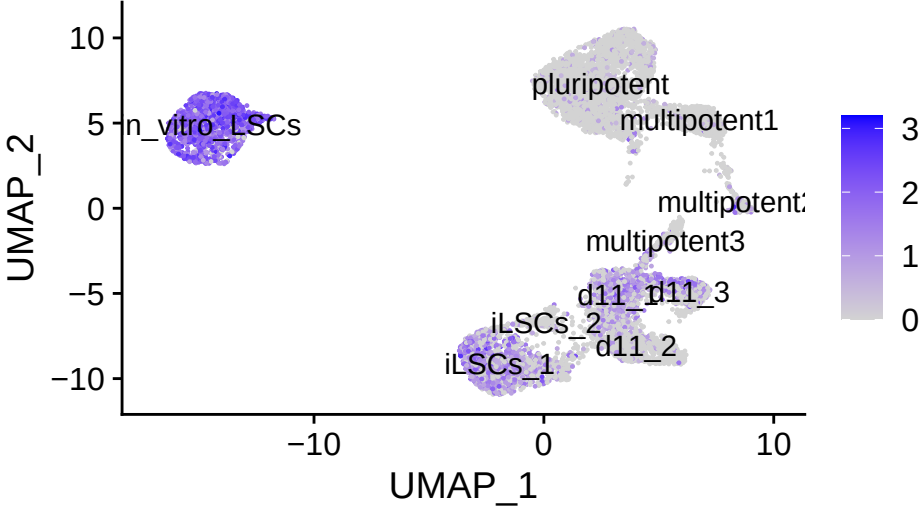
FGFBP1



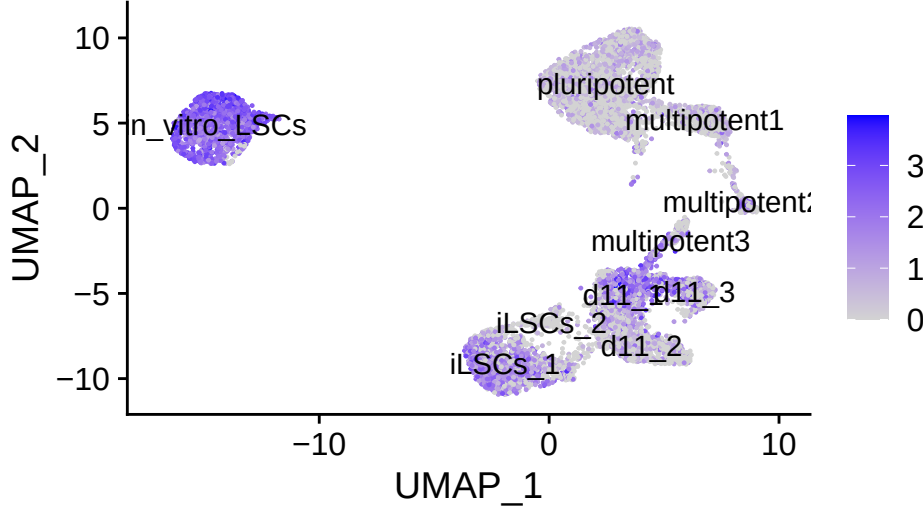
THBS1



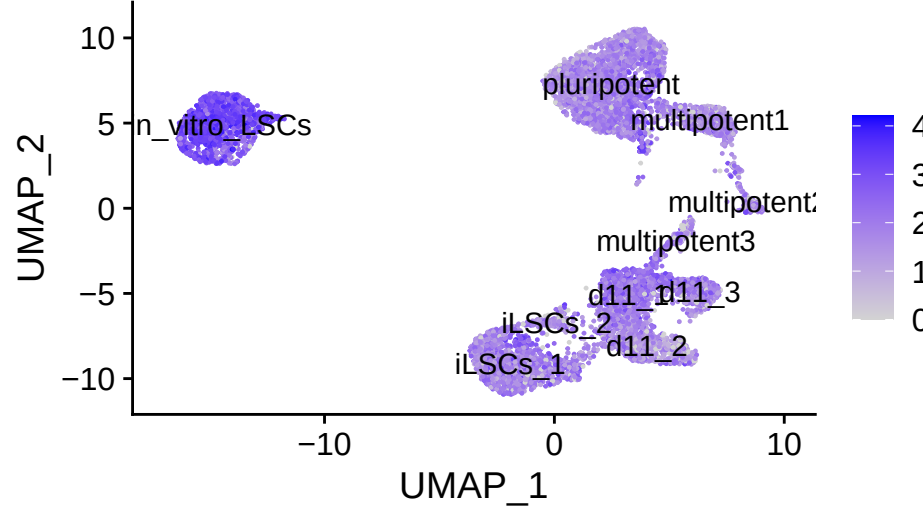
ITGA3



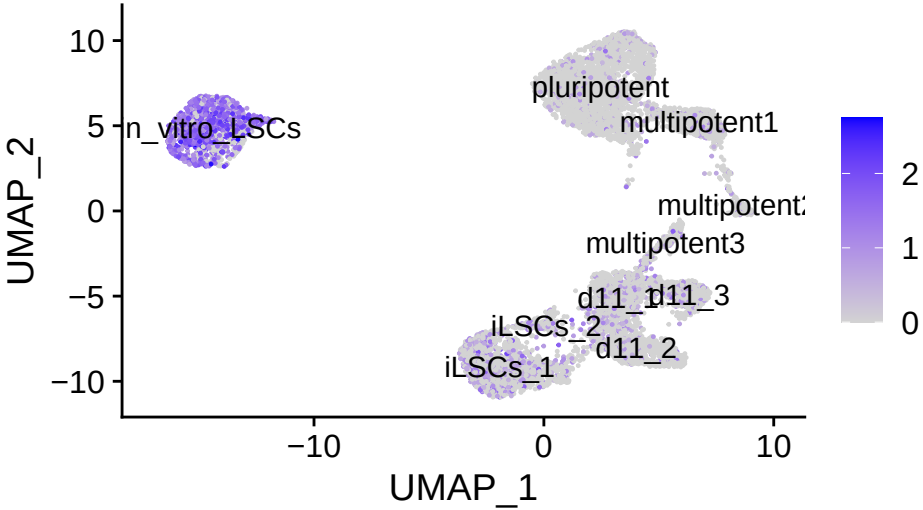
CDH13



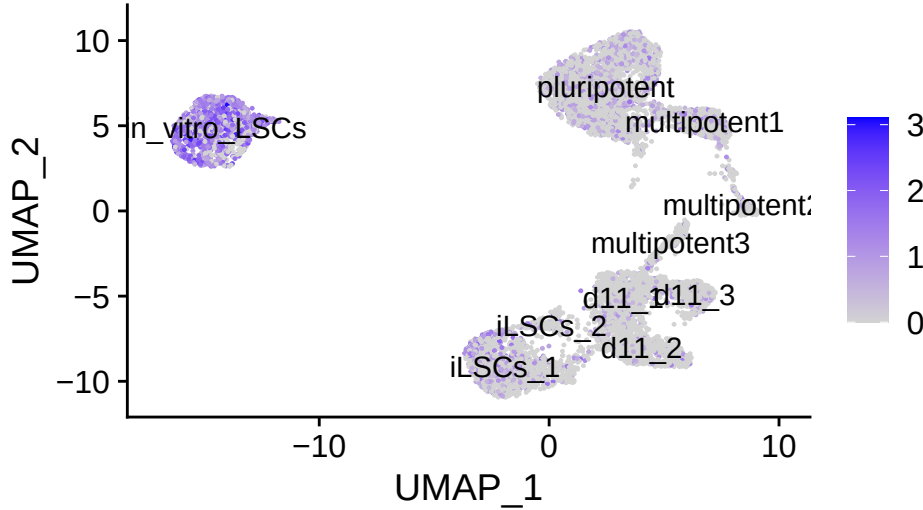
PFN1



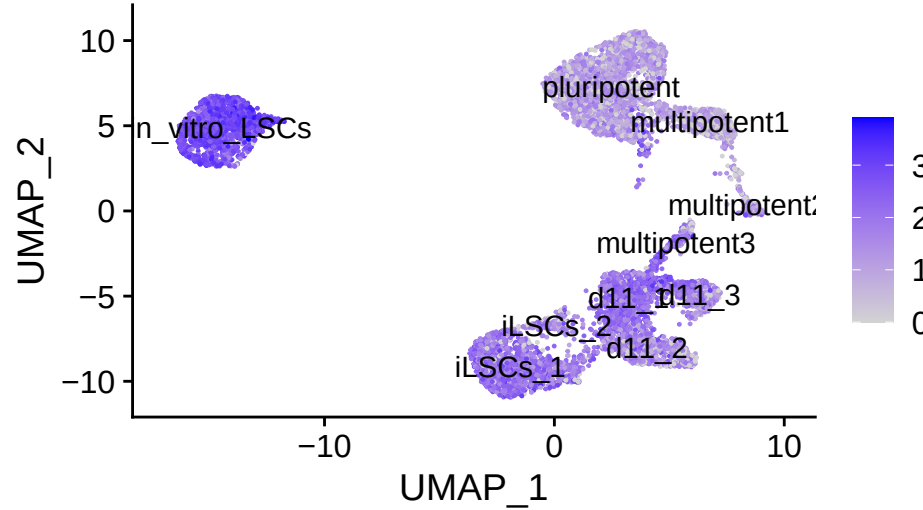
PLEC



SEMA4B



MYH9



enriched go term: actin filament organization from cluster: j_in_vitro_LSCs

