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| <p>● O0 (Stroma) TAGLN, ACTA2, MGP
cardiovascular system development</p> | <p>● O5 (Endothelium) CLDN5, PECAM1, KDR
cardiovascular system development</p> | <p>● O10 (Muscle prog.) MYOG, MYOD1
muscle filament sliding</p> |
| <p>● O1 (Stroma) MAB21L2, CXCL14, PRRX1
cartilage development</p> | <p>● O6 (Cell cycle) CENPF, HMGB2, UBE2C
mitotic cell cycle processes</p> | <p>● O11 (Neural prog.) HES6, STMN2
generation of neurons</p> |
| <p>● O2 (Podocyte) PODXL, NPHS2, TCF21
renal filtration cell differentiation</p> | <p>● O7 (Stroma) COL2A1, COL9A3, CNMD
extracellular matrix organisation</p> | <p>● O12 (Endothelium) GNG11, CALM1
negative reg. of cation channel activity</p> |
| <p>● O3 (Stroma) DLK1, GATA3, IGFBP5
wound healing</p> | <p>● O8 (Neural prog.) FABP7, TTHY1, SOX2
brain development</p> | |
| <p>● O4 (Cell cycle) HIST1H4C, PCLAF, TYMS
DNA conformation change</p> | <p>● O9 (Epithelium) PAX2, PAX8, KRT19
reg. of nephron tubule differentiation</p> | |