



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