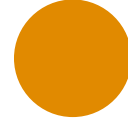
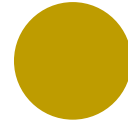


O0 (Stroma)
TAGLN, ACTA2, MGP
cardiovascular system development



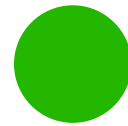
O1 (Stroma)
MAB21L2, CXCL14, PRRX1
cartilage development



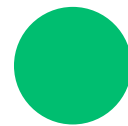
O2 (Podocyte)
PODXL, NPHS2, TCF21
renal filtration cell differentiation



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