GObp in module 5.1 pyrimidine ribonucleotide metabolic processlymph vessel morphogenesisretinal ganglion cell axon guidance Term negative regulation of fat cell differentiationmorphogenesis of an epithelial sheetaorta developmentbranching involved in ureteric bud morphogenesispositive regulation of phospholipase C activity-2.0 0.0 1.0 1.5 log2Enrichment GOcc in module 5.1 cullin family protein bindingcatalytic activity, acting on a glycoprotein-Term retinoic acid receptor binding chemokine binding SNAP receptor activitytau protein binding 0.0 0.5 1.5 2.0 1.0 log2Enrichment GOmf in module 5.1 cleavage furrowendoplasmic reticulum-Golgi intermediate compartment membrane-1.5 0.0 0.5 1.0 log2Enrichment