GO: Day 21 WGCNA MEmagenta.blue\_pink Cluster DNA binding GTPase activity RNA polymerase II cis-regulatory region sequence-specific DNA binding cysteine-type endopeptidase activity osmosensor activity spermidine synthase activity cysteine-type carboxypeptidase activity ferric iron binding nucleic acid binding methionine gamma-lyase activity glucosamine-6-phosphate deaminase activity ontology GTPase activator activity IMP cyclohydrolase activity BP cell morphogenesis MF positive regulation of protein phosphorylationregulation of chromatin assembly or disassembly positive regulation of receptor recycling in utero embryonic development immune system process positive regulation of T cell mediated cytotoxicity regulation of myeloid leukocyte differentiation mitotic spindle elongation -MvD88-dependent toll-like receptor signaling pathway neural tube closure 1891 MEmagenta.blue\_pink genes nuclear-transcribed mRNA poly(A) tail shortening 6 -log10(pvalue)