

E2Factivation\_1, E2Fdegradation\_1, E2Fsynthesis\_1, \_1, \_20, acycECDK2degradation\_1, cycDCDK4degradation\_1, cycDCDK4p16complexformation\_1, cycDCDK4p27complexdeassociation\_1, cycDCDK4p27complexformation\_1, cycDCDK4synthesis\_1, cycECDK2activation\_1, cycECDK2deactivation\_1, cycECDK2degradation\_1, cycECDK2p27complexformation\_1, cycECDK2p27deassociation\_1, cycECDK2synthesis\_1, p16degradation\_1, p16synthesisinhibitedbypRB\_1, p27degradation\_1, p27degradationviacycECDK2\_1, p27synthesis\_1, pRBE2Fcomplexformation\_1, pRBdegradation\_1, pRBpdephosphorylation\_1, pRBpdephosphorylation\_1, pRBsynthesis\_1 OP: Y10\_1, Y11\_1, Y1\_1, Y2\_1, Y3\_1, Y4\_1, Y5\_1, Y6\_1, Y7\_1, Y8\_1, Y9\_1  
 species: Y10\_1, Y11\_1, Y1\_1, Y2\_1, Y3\_1, Y4\_1, Y5\_1, Y6\_1, Y7\_1, Y8\_1, Y9\_1  
 {Y10\_1, Y11\_1, Y1\_1, Y2\_1, Y3\_1, Y4\_1, Y5\_1, Y6\_1, Y7\_1, Y8\_1, Y9\_1}