_1, _7, cycECDK2degradation_1, cyclebreak_1, cycleprogression_1, cyclinCDK4degradation_1, cyclin_1, pRBE2Fcomplexassociation_1, pRBE2FcomplexdeassociationviacycDCDK4_1, pRBpdephosphorylation_1 OP: D_1, E_1, RS_1, R_1, X_1
species: D_1, E_1, RS_1, R_1, X_1
{D_1, E_1, RS_1, R_1, X_1}