962-T1-453 Robert L. Fulton* (sharon.beam@med.va.gov), Department of Surgery, University of Louisville, Louisville, KY 40292, and Beth Bradley, Department of Mathematics, University of Louisville, Natural Science Building: Room 224, Louisville, KY 40292. Unit Step and Impulse Functions in Gene Dynamics. Preliminary report.

In a model of host response to bactera infection, (modified Lotka-Volterra system) it became apparent that Unitstep (US[t-a]) and DiracDelta(D(t-a)) functions are ideal representations of gene activity. During host response, geness are turned on, then off, promoting protein production. Gene G[t] is turned on by a promoter y[t] and off by a second protein x[t]. Then: y[t] ==conc. of promoter; x[t] ==conc. of off messenger G[t] == 1, ify[t] > critical amount and x[t] < critical amount; G[t] == 0, otherwise. If times when y[t] and x[t] are critical are plugged in, then: G'[t] == D[t-a] - D[t-b]. RNA(R[t]) is manufactured: R'[t] == KG[t] - JR[t]. Protein(P[t]), say an antibody, is built from RNA, then: P'[t] == MR[t]. K, J and M are rate coefficients. Explicit solutions were found. G'[t] is an ideal model for the unmasking of a gene. In a simple model, bacterial growth restrained by antibiotics is: B'[t] == aB[t] - B[t](cf[t]) where B[t] is the number of bacteria and c is related to the dose and sensitivity of the drug as a function of time f[t]. For the usual method of administration, tissue-blood level of antibiotic (f[t]) is (1-Cos[nt]). If mutation occurs, the solution to B'[t] == aB[t] - B[t](cf[t]]) + B[t]f[t] UnitStep[t-b] is a good model of development of antibiotic resistance. (Received September 14, 2000)