Uncertainty Quantification course Homework assignment 7

November 2, 2023

We discuss this exercise in the class meeting on 8 November 2023.

We have a very simple epidemiological model, a so-called SEIR model (for Susceptible, Exposed, Infected, Recovered). It consists of 4 coupled ODEs for S(t), E(t), I(t) and R(t), the number of individuals in each of the four categories (also called "compartments"). The total population consists of N = S(t) + E(t) + I(t) + R(t) individuals; we assume that N is constant in time. In this assignment you use stochastic collocation to explore the uncertainty in the model output due to uncertainty in **three** input parameters: the reproduction number R_0 , the incubation period T and the infectious period τ .

On Canvas you will find two Matlab codes SEIRmodel.m and SEIRscript.m for numerical integration of a SEIR model. As mentioned, the model has three parameters: the reproduction number R_0 , the incubation period T (in days) and the infectious period τ (also in days). The output consists of timeseries for S, E, I and R. The population size is set at $N=10^5$, the initial condition is set at E(0)=0, I(0)=10, R(0)=0 and S(0)=N-E(0)-I(0)-R(0). The code is based on a model from https://cs.uwaterloo.ca/~paforsyt/SEIR.html. The time integration runs from t=0 to t=500 with timestep dt=0.01.

We assume R_0 has uniform distribution on the interval [1.5, 3.0]. Furthermore, T has uniform distribution on the interval [5, 10], and τ has a beta distribution on the interval [1, 14], with probability density function $\rho(\tau) \propto (\tau - 1)(14 - \tau)$.

Let Q be the Quantity of Interest, defined as the maximum of I(t) over time, i.e. $Q = \max_t I(t)$. Thus, Q is the peak number of infected individuals (see also the code SEIRscript.m). It is dependent on the parameters R_0, T, τ .

Use stochastic collocation to construct an approximation $\tilde{Q}(R_0,T,\tau)$ of the exact $Q(R_0,T,\tau)$. To construct the approximation, don't use a built-in / library code for interpolation, instead write your own code using Lagrange interpolating functions. For the nodes, you can use the Clenshaw-Curtis nodes. Furthermore, you can use full tensor grids for your approximation, you don't have to go to sparse grids.

Once you have built your approximation $\tilde{Q}(R_0, T, \tau)$, use Monte Carlo sampling to assess the probability distribution of \tilde{Q} . Make a histogram plot to show the distribution of the MC samples. What are its mean and standard deviation? Can you give an estimate of the probability that \tilde{Q} will be higher than 15000?