

Computing Upper Probabilities of Failure Using Optimization Algorithms Together with Reweighting and Importance Sampling

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The presentation addresses efficient computation of the upper probability of failure of engineering structures, when the uncertainty is modelled by a family of probability densities. We develop an algorithm significantly reducing the sample sizes required in the optimization algorithm by adopting a recursive importance sampling scheme.

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1. Parametrized and Upper Probabilities

Let $\{X_t\}_{t \in \mathcal{T}}$ be a family of multivariate random variables $X_t : \Omega \rightarrow D \subseteq \mathbb{R}^{d_1}$ with corresponding density functions $f_t : D \rightarrow \mathbb{R}$ which are parametrized by $t = (\tau_1, \dots, \tau_{d_2}) \in \mathcal{T}$. Further let $h : D \rightarrow \{0, 1\}$ be an indicator function on D where $h(x) = 1$ means failure and 0 no failure. Then the probability of failure depending on t is $p(t) = P(h(X_t) = 1) = \int_D h(x)f_t(x) dx$ and the upper probability of failure \bar{p} is obtained by solving the global optimization problem $\bar{p} = \max_{t \in \mathcal{T}} p(t)$. To explain the situation we start with a numerical example.

2. Numerical Example

We consider the re-calculation of the reliability of a part of an airplane. The corresponding FE-model has $4.7 \cdot 10^6$ degrees of freedom and needs 48 different input parameters. Only the two most decisive input parameters $x = (x_1, x_2)$ are taken into account which are the Young's moduli of two metal components. The output of the FE-computations is the value of the above indicator function h in some complementary components.

The uncertainty about the values x_1 and x_2 is modelled by a family of two-dimensional random variables $\{X_t\}_{t \in \mathcal{T}}$ with $X_t \sim \mathcal{N}(\mu(t), \Sigma(t))$ parametrized by $t = (\varphi, \rho)$. The means $\mu(t)$ may vary on a disc of radius $r = 2$ GPa and midpoint

(m, m) with $m = 70$ GPa. Since for upper probabilities the boundary of the disc is relevant we consider only the circle

$$\mu(t) = \mu(\varphi, \rho) = 2 \cdot \begin{bmatrix} \cos(\varphi) \\ \sin(\varphi) \end{bmatrix} + \begin{bmatrix} m \\ m \end{bmatrix}, \varphi \in [0, 2\pi].$$

The coefficient of correlation varies in an interval, $t_2 = \rho \in [0, 0.8]$, and the coefficient of variation $\nu = 6.5\%$ is assumed to be deterministic. This leads to a parametrization of the covariance matrix

$$\Sigma(t) = \Sigma(\varphi, \rho) = \begin{bmatrix} \sigma_1(\varphi)^2 & \sigma_1(\varphi)\sigma_2(\varphi)\rho \\ \sigma_1(\varphi)\sigma_2(\varphi)\rho & \sigma_2(\varphi)^2 \end{bmatrix}$$

with $\sigma_i(\varphi) = \nu\mu_i(\varphi)$.

3. Estimating $p(t)$ and Their Derivatives

An optimization algorithm for obtaining upper probabilities needs a sequence of parameter values t_1, \dots, t_m and their probabilities (function values) $p(t_1), \dots, p(t_m)$. To estimate these probabilities $p(t_i)$ by Monte Carlo simulations it is crucial that for all t_k the simulations are based on the same single set $\omega = \{\omega_1, \dots, \omega_N\}$ of N random numbers, see Troffaes et al. (2018). These random numbers are then transformed to sets $R_{\omega, t_k} = \{r_1, \dots, r_N\}$ of sample points distributed according to densities f_{t_k} as for the estimates needed. For parametrized Gaussian distributions it means starting from random numbers $\omega_i \sim \mathcal{N}(0, I)$ and transforming to sample points

$r_i(t) \sim \mathcal{N}(\mu(t), \Sigma(t))$ by $r_i(t) = \mu(t) + C(t)\omega_k$ with Cholesky factor $C(t)$ of $\Sigma(t)$.

Estimating $p(t_k) \approx \frac{1}{|R_{\omega,t_k}|} \sum_{r \in R_{\omega,t_k}} h(r)$ by Monte Carlo simulation independently for all $k = 1, \dots, m$ would lead to mN evaluations of function h . For reducing this high computational effort it is important for an estimate of $p(t_n)$ in step n of optimization to re-use $h(r)$ and sample points r in sets R_{ω,t_k} from previous steps $k < n$. We use reweighting or importance sampling as in Fetz (2017) and Owen (2018), but here on each set of a partition $D = D_1^n \cap \dots \cap D_m^n$:

$$p(t_n) = \sum_{k=1}^n \int_{D_k^n} h(x) \frac{f_{t_n}(x)}{f_{t_k}(x)} f_{t_k}(x) dx$$

$$\approx \sum_{k=1}^n \frac{1}{|R_{\omega,t_k}^n|} \sum_{r \in R_{\omega,t_k}^n} h(r) \frac{f_{t_n}(r)}{f_{t_k}(r)} =: p_{\mathcal{R}_\omega^n}(t_n)$$

where $R_{\omega,t_k}^n = R_{\omega,t_k} \cap D_k^n$ is the set R_{ω,t_k} of sample points restricted to the set D_k^n of the current partition and \mathcal{R}_ω^n the set $\{R_{\omega,t_1}^n, \dots, R_{\omega,t_n}^n\}$ of all samples considered in step n . The expensive $h(r)$ is evaluated for $r \in R_{\omega,t_n}^n$ only, all other $h(r)$ are already known from the steps before. The ratios between the original density f_{t_n} and the densities f_{t_k} used instead are the importance sampling ratios which should not be too large (Owen (2018)), because large ratios mean fewer sample points in a considered area leading to less reliable estimates. Taking this into account we define

$$D_k^n = \{x \in D : f_{t_k}(x) \geq \max_{j \leq n, j \neq k} f_{t_j}(x)\}.$$

We do not have to know D_k^n in detail. It is sufficient to know the sets R_{ω,t_k}^n of sample points r keeping only these r where $f_{t_k}(r) > f_{t_j}(r)$, $j = 1, \dots, n, j \neq k$. We generate such sets recursively for $n > 1$ setting $R_{\omega,t_1}^1 = R_{\omega,t_1}$ and

$$R_{\omega,t_n}^n = \{r \in R_{\omega,t_n} : f_{t_n}(r) > \max_{k < n} c_k^{n-1} f_{t_k}(r)\}$$

for the newest and updating all previous ($k < n$):

$$R_{\omega,t_k}^n = \{r \in R_{\omega,t_k}^{n-1} : c_k^{n-1} f_{t_k}(r) > f_{t_n}(r)\}.$$

In addition we may use factors c_k^{n-1} to be more tolerant allowing lower densities to get fewer new sample points. These factors are also recursively defined as $c_k^n = c_n c_k^{n-1}$ where the cumulative multiplications ensure that $R_{\omega,t_k}^n \subseteq \dots \subseteq R_{\omega,t_k}^{k-1}$.

Further we may go one step back in the recursion if $|R_{\omega,t_n}^n|$ is less than some percentage q of N .

For the computation of the i^{th} partial derivative at t_n needed in the optimization algorithm we use $(p_{\mathcal{R}_\omega^n}(t_n + h^{(i)}) - p_{\mathcal{R}_\omega^n}(t_n)) / h_i^{(i)}, h_j^{(i)} = 10^{-8} \delta_{ij}$.

4. Optimizing and Results

For our optimization problem we choose $q = 10\%$, a sequence $c_n = (c_{n-1} - 1)/2 + 1$ with $c_1 = 1.5$ preventing too large tolerances, and $N = 50\,000$. The MATLAB global optimization algorithm needs 2777 parameter values t_k requiring 218 160 evaluations of h compared to 2777 N without this new method which re-uses the $h(r)$ from former samples together with good importance sampling ratios on the sets D_k^n of the partition increasing the accuracy of the estimates. In Fig. 1 we show the contour lines of estimates for p (classical and with $p_{\mathcal{R}_\omega^n}$). All parameters t_k used in optimization are visualized as dots where the colour/size indicates the sample sizes used (gray: no new samples, red: new samples needed, the greater the dot the more sample points). We obtain the upper probability $\bar{p} = 0.021$ at $t = (2.438, 0)$.

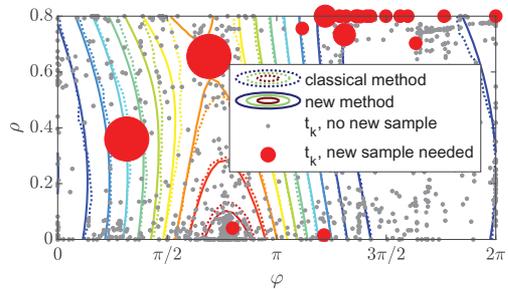


Fig. 1. Contour lines of estimates of p and points t_k .

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