# Uncertainty management in multidisciplinary design of 

# critical safety systems 

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Managing the uncertainty in multidisciplinary design of safety critical systems requires not only the availability of a single approach or methodology to deal with uncertainty but a set of different strategies and scalable computational tools (i.e. by making use of the computational power of a cluster and grid computing). The availability of multiple tools and approaches for dealing with uncertainties allows to cross-validate the results and increase the confidence in the performed analysis.

This paper presents a unified theory and an integrated and open general purpose computational framework to deal with scarce data, aleatory and epistemic uncertainties. It allows to solve different tasks necessary to manage the uncertainty, such as: uncertainty characterization, sensitivity analysis, uncertainty quantification and robust design. The proposed computational framework is generally applicable to solve different problems in different fields and numerically efficient and scalable allowing for a significant reduction of the computational time required for uncertainty management and robust design.

The applicability of the proposed approach is demonstrated by solving a multidisciplinary design of a critical system proposed by NASA Langley in the multidisciplinary uncertainty quantification challenge problem.

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## I. Introduction

In order to design safe components and systems, the explicit inclusion of uncertainties from different sources is an indispensable step. In fact, under realistic conditions, these products are affected by uncertainties, caused by the lack of sufficient knowledge and/or by natural unpredictable external events. Uncertainty analysis is essential for modellers to obtain a robust representation of model predictions consistent with the state-of-knowledge. If the effects of the uncertainties in the "optimized" products are ignored, these products may perform unsatisfactorily in realistic conditions; for instance, they can show a very low reliability, high reparation and maintenance costs. On the other hand, in a robust design a product or system is less sensitive to the uncertainties and hence, it reaches low variability of the overall performances that can allow significant reductions in terms of e.g. the manufacturing and operating costs).

The design of safety critical systems faces very complex problems due to the presence of varying levels of aleatory and epistemic uncertainty. Aleatory uncertainty is inherent in many natural systems, and therefore cannot be reduced, but can be described and its effect quantified. Epistemic uncertainty is not completely avoidable, since it is not possible to perfectly model or predict real world situations, although epistemic uncertainty can be reduced, better characterized and quantified by using available knowledge. Despite the different levels of uncertainty, decision makers still need to make clear choices based on the available information. They need to trust the methodology adopted to propagate uncertainties through multi-disciplinary analysis, in order to quantify the risk with the current level of information and avoid wrong decisions due to artificial restrictions introduced by the modelling.

Multiple mathematical concepts can be used to characterize variability and uncertainty. Probability distributions can be used to represent the relative frequency of a given state of the system, or they can represent the degree of belief or confidence that a given state of the system exists. ${ }^{1}$ Often, very limited information is available, and collecting more data or samples might not be possible or too expensive. Given the limitations of amount of data, quantification methods often rely on subjective judgment and assumptions and it may not always seem reasonable to characterize the uncertainties in a classical probabilistic way. To avoid the inclusion of subjective and often unjustified hypothesis, the imprecision and vagueness of the data can be treated by using concepts of imprecise probabilities. Imprecise probability combines probabilistic and set theoretical components in a unified theory allowing the identification of bounds on probabilities for the
events of interest. ${ }^{2}$

Random set theory is specially suited to model under the same framework uncertainty represented as cumulative distribution functions (CDFs), intervals, ${ }^{3}$ probability boxes, ${ }^{4}$ possibility distributions ${ }^{5}$ (they are closely related to normalized fuzzy sets) and Dempster-Shafer ${ }^{6,7}$ structures without making any implicit or explicit assumptions. In other words, random set theory is a technique that permits to model naturally the aforementioned representations of uncertainty.

In this work, novel and efficient strategies are proposed to deal with aleatory and epistemic uncertainty. Random set theory is used as a unifying theoretical framework, to model different representations of the uncertainty. Additionally, the developed procedures have been implemented in an integrated computational framework allowing to solve realistic problems using a number of different approaches and methodologies. This provides an excellent tool for cross-validating the results obtained at each stage of the analysis and hence to increase the confidence in the adopted methodology and in the results. The applicability of approach is demonstrated by solving the NASA Langley multidisciplinary uncertainty quantification (UQ) challenge problem. ${ }^{8}$

## Motivation of the study

The development of safety-critical systems that must be designed to operate in harsh environments with a wide array of operating conditions (e.g. new vehicles, aircraft, nuclear power plants etc.) is a challenging problem. Furthermore, the failure of such systems might have high consequences for which quantitative data is either very sparse or prohibitively expensive to collect. Hence, uncertainty management is necessary to provide support to the decision makers through a series of different and interconnected analyses. For instance, estimating the importance of collecting additional information allows to characterize and reduce uncertainty; by performing sensitivity analysis, it is possible to identify the parameters that contribute the most to the variability of the output; uncertainty propagation allows to study the effects of uncertainty on the performance of the system and to identify extreme-case scenarios; finally, optimizing the design explicitly taking into account the effect of uncertainties allows to design a robust system.

Recent reports have clearly shown that the risk assumed by the decision maker is often wrongly estimated due to inadequate assessment of uncertainty. ${ }^{8}$ Modelling and simulation standards require estimates of
uncertainty (and descriptions of any processes used to obtain these estimates) in order to increase confidence and consistency in safety predictions and encourage the development of improved methods for quantifying and managing uncertainty. In this context, the NASA Langley multidisciplinary uncertainty quantification (UQ) challenge problem has been addressed in order to determine limitations and ranges of applicability of existing UQ methodology and to advance the state of the practice in UQ problem. ${ }^{9}$

The NASA challenge problem has represented a unique opportunity to test, validate and advance the capability of the computational framework, namely OpenCossan. ${ }^{10}$ This computational framework is able to deal with different representations of uncertainty and has been adopted to solve all the tasks proposed by the challengers.

Since many of the employed methods are rooted on random set theory, a brief introduction of the theory for the representation of the joint existence of aleatory and epistemic uncertainty is presented in Section II. The developed approaches for uncertainty quantification and management are presented in Section III. Section IV presents the computational framework and some details on computational complexity. Section V summarizes the main aspects, goals and difficulties of the NASA Langley multidisciplinary UQ challenge problem as well as the results of the various tasks of the challenge problem. Finally, the potentiality and applicability of the developed computational framework and the proposed approaches are discussed.

## II. Theoretical background

Random set theory is specially suited to model, under the same framework, uncertainty represented as cumulative distribution functions (CDFs), intervals, distribution-free probability boxes, possibility distributions and Dempster-Shafer structures ${ }^{6,7}$ without making any implicit or explicit assumption at all. ${ }^{11,12}$ In other words, random set theory is a technique that permits to model the aforementioned representations of uncertainty. Random sets can be understood as random variables that sample, instead of points, sets (called focal elements) as realizations.

In this context, many of the proposed solutions to the challenge problem make strong use of this kind of representation. In consequence, in the following, a brief review of the main concepts of random set theory that will be required in the subsequent discussion is provided. Also some new concepts developed in order to solve the challenge problem will be introduced.

## A. A succinct review of random set theory

Let us consider a universal set $\mathscr{X} \neq \emptyset$ and its power set $\mathscr{P}(\mathscr{X})$. Let $\left(\boldsymbol{\Omega}^{\prime}, \sigma_{\boldsymbol{\Omega}^{\prime}}, P_{\boldsymbol{\Omega}^{\prime}}\right)$ be a probability space and $\left(\mathscr{F}, \sigma_{\mathscr{F}}\right)$ be a measurable space where $\mathscr{F} \subseteq \mathscr{P}(\mathscr{X})$. A random set $\Gamma$ is a $\left(\sigma_{\Omega^{\prime}}-\sigma_{\mathscr{F}}\right)$-measurable mapping $\Gamma: \boldsymbol{\Omega}^{\prime} \rightarrow \mathscr{F}, \alpha \mapsto \Gamma(\alpha)$. We will say that every $\gamma:=\Gamma(\alpha) \in \mathscr{F}$ is a focal element while $\mathscr{F}$ is a focal set.

Analogously to the definition of a random variable, this mapping can be used to define a probability measure on $\left(\mathscr{F}, \sigma_{\mathscr{F}}\right)$ given by $P_{\Gamma}:=P_{\Omega^{\prime}} \circ \Gamma^{-1}$. That is, an event $\mathcal{R} \in \sigma_{\mathscr{F}}$ has the probability

$$
\begin{equation*}
P_{\Gamma}(\mathcal{R})=P_{\boldsymbol{\Omega}^{\prime}}\left\{\alpha \in \boldsymbol{\Omega}^{\prime}: \Gamma(\alpha) \in \mathcal{R}\right\} \tag{1}
\end{equation*}
$$

The random set $\Gamma$ will be also referred to as $\left(\mathscr{F}, P_{\Gamma}\right)$. When all the focal elements of $\mathscr{F}$ are singletons, then $\Gamma$ becomes a random variable $X$; hence, $\Gamma(\alpha)=X(\alpha)$ and the probability of occurrence of the event $\mathcal{F}$, is $P_{X}(\mathcal{F}):=\left(P_{\boldsymbol{\Omega}^{\prime}} \circ X^{-1}\right)(\mathcal{F})=P_{\boldsymbol{\Omega}^{\prime}}\{\alpha: X(\alpha) \in \mathcal{F}\}$ for every $\mathcal{F} \in \sigma_{X}$. In the case of random sets, it is not possible to compute exactly $P_{X}(\mathcal{F})$ but its upper and lower probability bounds. Dempster ${ }^{6}$ defined those upper and lower probabilities by,

$$
\begin{align*}
& \operatorname{LP}_{\left(\mathscr{F}, P_{\Gamma}\right)}(\mathcal{F}):=P_{\Omega^{\prime}}\{\alpha: \Gamma(\alpha) \subseteq \mathcal{F}, \Gamma(\alpha) \neq \emptyset\}  \tag{2a}\\
& \mathrm{UP}_{\left(\mathscr{F}, P_{\Gamma}\right)}(\mathcal{F}):=P_{\Omega^{\prime}}\{\alpha: \Gamma(\alpha) \cap \mathcal{F} \neq \emptyset\} \tag{2b}
\end{align*}
$$

where $\operatorname{LP}_{\left(\mathscr{F}, P_{\Gamma}\right)}(\mathcal{F}) \leq P_{X}(\mathcal{F}) \leq \mathrm{UP}_{\left(\mathscr{F}, P_{\Gamma}\right)}(\mathcal{F})$.

Copulas A copula is a function $C:[0,1]^{d} \rightarrow[0,1]$ that relates a joint cumulative density functions (CDFs) with its marginals, carrying in this way the dependence information in the joint CDF such that each of its marginal CDFs is uniform on the interval $[0,1]$. According to Sklar's theorem (see Refs. 13, 14), a multivariate $\operatorname{CDF} F_{X_{1}, X_{2}, \ldots, X_{d}}\left(x_{1}, \ldots, x_{d}\right)=P\left[X_{1} \leq x_{1}, \ldots, X_{d} \leq x_{d}\right]$ of a random vector $\left(X_{1}, X_{2}, \ldots, X_{d}\right)$ with marginals $F_{X_{i}}\left(x_{i}\right)=P\left[X_{i} \leq x_{i}\right]$ can be written as $F_{X_{1}, X_{2}, \ldots, X_{d}}\left(x_{1}, \ldots, x_{d}\right)=C\left(F_{X_{1}}\left(x_{1}\right), \ldots, F_{X_{d}}\left(x_{d}\right)\right)$, where $C$ is a copula. The copula $C$ is itself a CDF and it contains all information on the dependence structure between the components of $\left(X_{1}, X_{2}, \ldots, X_{d}\right)$ whereas the marginal cumulative distribution functions $F_{X_{i}}$ contain all information on the marginal distributions.

The reader is referred to Ref. 15 for an excellent introduction to copulas.

## B. Random sets, CDFs, distribution-free probability boxes and intervals

The original definition of random sets is very general; Alvarez ${ }^{11,12}$ showed that making the particularizations $\boldsymbol{\Omega}^{\prime}:=(0,1]^{d}, \sigma_{\boldsymbol{\Omega}^{\prime}}:=(0,1]^{d} \cap \mathscr{B}^{d}$, where $\mathscr{B}$ stands for the Borel $\sigma$-algebra on $\mathbb{R}$, and $P_{\Gamma} \equiv \mu_{C}$ for some copula $C$ that contains the dependence information within the joint random set, and using intervals and $d$-dimensional boxes as elements of $\mathscr{F}$, it is enough to model possibility distributions, distribution-free probability boxes, intervals, CDFs and Dempster-Shafer structures or their joint combinations (for a definition of joint Dempster-Shafer structure and joint random set the reader is referred to Ref. 12). Here, $P_{\Gamma} \equiv \mu_{C}$ denotes the fact that $P_{\Gamma}$ is the probability measure generated by $P_{\boldsymbol{\Omega}^{\prime}}$ which is defined by the Lebesgue-Stieltjes measure corresponding to the copula $C$, i.e. $\mu_{C}$. In other words, $P_{\Gamma}(\Gamma(G))=\mu_{C}(G)$ for $G \in \sigma_{\Omega^{\prime}}$.

In the rest of this subsection, $\left(\boldsymbol{\Omega}^{\prime}, \sigma_{\boldsymbol{\Omega}^{\prime}}, P_{\boldsymbol{\Omega}^{\prime}}\right)$ will stand for a probability space with $\boldsymbol{\Omega}^{\prime}:=(0,1], \sigma_{\boldsymbol{\Omega}^{\prime}}:=$ $(0,1] \cap \mathscr{B}:=\cup_{\theta \in \mathscr{B}}\{(0,1] \cap \theta\}$ and $P_{\Omega^{\prime}}$ will be a probability measure corresponding to the CDF of a random variable $\tilde{\alpha}$ uniformly distributed on $(0,1]$, i.e. $F_{\tilde{\alpha}}(\alpha):=P_{\boldsymbol{\Omega}^{\prime}}[\tilde{\alpha} \leq \alpha]=\alpha$ for $\alpha \in(0,1]$; that is, $P_{\boldsymbol{\Omega}^{\prime}}$ is a Lebesgue measure on $(0,1]$.

## 1. Cumulative distribution functions

When a variable is expressed as a random variable on $X \subseteq \mathbb{R}$, the probability law of the random variable can be expressed using a CDF $F_{X}$ (recall $F_{X}(x)=P_{\Gamma}(X \leq x)$ for $\left.x \in X\right)$. That CDF can be represented as the random set $\Gamma: \boldsymbol{\Omega}^{\prime} \rightarrow \mathscr{F}, \alpha \mapsto \Gamma(\alpha)$ where $\mathscr{F}$ is the system of focal elements $\Gamma(\alpha):=F_{X}^{-1}(\alpha)$ for $\alpha \in \boldsymbol{\Omega}^{\prime}$ (the inverse of the CDF $F_{X}$ is defined by $F_{X}^{-1}(\alpha):=\inf \left\{x: F_{X}(x) \geq \alpha, \alpha \in(0,1]\right\}$; take into account that this definition uses the infimum since CDFs are weakly monotonic and right-continuous). Note that the representation of the CDF as a random set only contains an aleatory component, which is given either by $\alpha$, or by its corresponding sample $x=F_{X}^{-1}(\alpha)$; there is not an epistemic component in this representation.

## 2. Intervals

An interval $I=[l, u]$ can be represented as the random set $\Gamma: \Omega^{\prime} \rightarrow \mathscr{F}, \alpha \mapsto \Gamma(\alpha)$ (i.e. $\left.\left(\mathscr{F}, P_{\Gamma}\right)\right)$ defined on $\mathbb{R}$ where the focal set contains the unique focal element $[l, u]$, that is, $\mathscr{F}=I$ and $\alpha \in(0,1] \equiv \boldsymbol{\Omega}^{\prime} ;$ in this case, $P_{\Gamma}$ is specified by Eq. (1). In other words, all the samplings of $\alpha \in \Omega^{\prime}$ draw the interval $[l, u]$. Note that the representation of intervals as a random set does not contain an aleatory component, inasmuch as
it does not matter which value $\alpha$ takes, because all $\alpha$-s map to the same focal element $I$. In this case, the epistemic component is given by the interval itself, $I$.

## 3. Probability boxes

A probability box or $p$-box (term coined by Ferson et $\mathrm{al.}^{4}$ ) $\langle\underline{F}, \bar{F}\rangle$ is a set of CDFs $\{F: \underline{F} \leq F \leq$ $\bar{F}, F$ is a CDF $\}$, delimited by upper and lower CDF bounds $\underline{F}$ and $\bar{F}: \mathbb{R} \rightarrow[0,1]$, which collectively represent the epistemic uncertainty about the CDF of a random variable. This class of functions may not have additional restrictions or may belong, as well, to a reduced class of CDFs; using that discrimination, probability boxes can be naturally grouped into two disjoint groups: free and distributional.

DISTRIBUTION-FREE P-BOXES Distribution-free p-boxes (also known as non-parametric p-boxes) appear when the CDF of a random variable cannot be specified precisely, given that the CDF family is unknown; in this case only the upper and lower CDF bounds $\underline{F}$ and $\bar{F}$ bounds of the probability box are specified. These bounds can either be defined in advance or can be estimated using for example the methods listed in Zhang et. al. ${ }^{16}$ and references therein. Note that distribution-free p-boxes do not make any assumption about the family or shape of the uncertain CDFs that belong to the p-box.

There are two alternatives but equivalent methods to represent distribution-free p-boxes using random set theory.

The first method was proposed in Refs. 11,12. Using this method, a distribution-free probability box delimited by lower and upper CDF bounds $\underline{F}$ and $\bar{F}$ can be represented as the random set $\Gamma: \boldsymbol{\Omega}^{\prime} \rightarrow$ $\mathscr{F}, \alpha \mapsto \Gamma(\alpha)$ (i.e. $\left.\left(\mathscr{F}, P_{\Gamma}\right)\right)$ defined on $\mathbb{R}$ where $\mathscr{F}$ is the class of focal elements $\Gamma(\alpha):=\langle\underline{F}, \bar{F}\rangle^{-1}(\alpha):=$ $\left[\bar{F}^{-1}(\alpha), \underline{F}^{-1}(\alpha)\right]$ for $\alpha \in(0,1] \equiv \boldsymbol{\Omega}^{\prime}$ with $\underline{F}^{-1}(\alpha)$ and $\bar{F}^{-1}(\alpha)$ denoting the inverses of $\underline{F}$ and $\bar{F}$ and $P_{\Gamma}$ is specified by Eq. (1).

The second alternative method, proposed here, considers a random variable which follows a CDF $F$ with parameters $\theta_{i}$ that belong to the interval $I_{i}$ for $i=1,2, \ldots, m$ (i.e. $F\left(\cdot ; \theta_{1}, \theta_{2}, \ldots, \theta_{m}\right)$ ). This representation is in comparison to the first method, which models a p-box using only its lower and upper CDF bounds $\underline{F}$ and $\bar{F}$. Using the random set representation, a focal element of the probability box $\langle\underline{F}, \bar{F}\rangle$ can be represented as the image through the function $F^{-1}$ of the input intervals $\left\{I_{i}: i=1,2, \ldots, m\right\}$ together with the sample of $\alpha$ which is a uniform random variable on $(0,1] \equiv \boldsymbol{\Omega}^{\prime}$. In consequence, it can be represented as the
random set $\Gamma: \boldsymbol{\Omega}^{\prime} \rightarrow \mathscr{F}, \alpha \mapsto \Gamma(\alpha)$ (i.e. $\left.\left(\mathscr{F}, P_{\Gamma}\right)\right)$ defined on $\mathbb{R}$ where $\mathscr{F}$ is the system of focal elements $\left\{F^{-1}\left(\alpha ; I_{1}, I_{2}, \cdots, I_{m}\right): \alpha \in \boldsymbol{\Omega}^{\prime}\right\}$ and $P_{\Gamma}$ is specified by Eq. (1). Observe that each focal element has an aleatory component $\alpha$ and an epistemic component in the Cartesian product $\times_{i=1}^{m} I_{i} \triangleq I_{1} \times I_{2} \times \cdots \times I_{m}$.

This representation of distribution-free p-boxes shows that for a single realization of the aleatory component $\alpha$, a focal element contains the image through $F^{-1}$ of all the possible combinations of values of the intervals for the parameters of the parental CDF $F$. It derives from the fact that a focal element is defined as: $\langle\underline{F}, \bar{F}\rangle^{-1}(\alpha)=\{x: F(x)=\alpha, F \in\langle\underline{F}, \bar{F}\rangle\}, \bar{F}^{-1}(\alpha)=\inf _{\boldsymbol{\theta} \in \times_{i=1}^{m} I_{i}} F^{-1}\left(\alpha ; \theta_{1}, \ldots, \theta_{m}\right)$ and $\underline{F}^{-1}(\alpha)=\sup _{\boldsymbol{\theta} \in X_{i=1}^{m} I_{i}} F^{-1}\left(\alpha ; \theta_{1}, \ldots, \theta_{m}\right)$.

Note that only distribution-free probability boxes can be represented using random set theory. However, in the analysis of the challenge problem a different approach has been used to represent distributional probability boxes as will be explained in the following lines.

Distributional p-boxes and the double loop Monte Carlo strategy Distributional p-boxes (also known as parametric p-boxes) appear when there is uncertainty in the representation of the parameters of a given CDF (hereafter called the parental $C D F$.) These parameters are imprecisely specified as intervals. For instance, consider a quantity that is known to be Gaussian with mean within the interval $[1,2]$ and standard deviation somewhere in $[3,4]$; Ferson et. al. ${ }^{4}$ describes how to obtain such probability boxes. All CDFs that are normal and have means and standard deviations inside these respective intervals will belong to this probability box. The upper and lower CDF bounds $\underline{F}$ and $\bar{F}$ of the p-box enclose many non-normal distributions, but these would be excluded from the p-box by specifying the normal CDF as the parental distribution family.

According to the second representation of distribution-free p-boxes, the focal element corresponding to a realization $\alpha$ of the aleatory component contains the image through $F^{-1}$ of all possible $\boldsymbol{\theta} \in X_{i=1}^{m} I_{i}$. As consequence, a set of focal elements of the probability box would be a family of intervals each of them being a mapping of $X_{i=1}^{m} I_{i}$ through $F^{-1}$. Hence, for a fixed value of $\boldsymbol{\theta}$ it is not possible to identify in that set of intervals the points that would belong to some CDF. For this reason, random set theory can not be used to model distributional p-boxes.

Distributional p-boxes can be dealt with using a double loop Monte Carlo strategy, in which the outer loop draws $\boldsymbol{\theta}$-s from $X_{i=1}^{m} I_{i}$ and the inner loop samples $\alpha$-s from a uniform distribution in ( 0,1$]$. In this case,
using the principle of maximum entropy, we will assume a uniform distribution in $\times_{i=1}^{m} I_{i}$. This approach has been used to solve some of the tasks of the challenge problem. Please note that the outer loop can be used to drive an optimization/search process in $\times_{i=1}^{m} I_{i}$ to identify the lower and upper bounds. In this case, it is not necessary to assume a uniform distribution in $\times_{i=1}^{m} I_{i}$.

## C. Sampling from a random set

A sample from a random set is simply obtained by generating an $\alpha$ from a uniform distribution on $(0,1]$ and then, retrieving the corresponding focal element $\Gamma(\alpha)$; for example, for sampling from a distributionalfree probability box an $\alpha$ uniformly distributed in ( 0,1 ] is drawn and then its corresponding " $\alpha$-cut" $\left[\bar{F}^{-1}(\alpha), \underline{F}^{-1}(\alpha)\right]$ is obtained. In the case of multivariate random sets, a sample $\boldsymbol{\alpha} \in \boldsymbol{\Omega}^{\prime}$ is drawn from the copula $C$ that models the dependence between the input variables. Then, the corresponding marginal focal elements are obtained and combined as explained in the next subsection. Take into account that $n$ samples of a random set form the Dempster-Shafer structure $\left(\mathscr{F}_{n}, m\right)$; here $\mathscr{F}_{n}$ denotes the set of all sampled focal elements; the basic mass assignment $m$ associated to each focal element is equal to $1 / n$; note that a Dempster-Shafer structure is itself a finite random set. ${ }^{11,12}$

Samples from distributional p-boxes can be obtained resorting to a double Monte Carlo loop as explained in the previous Section.

## D. Combination of focal elements

After sampling each input variable, a combination of the sampled focal elements is carried out. Usually, the joint focal elements are given by the Cartesian product $X_{i=1}^{d} \gamma_{i} \subseteq \mathscr{X}$ where $d$ is the number of input variables, $\gamma_{i}:=\Gamma^{i}\left(\alpha_{i}\right)$ are the sampled focal elements from every input variable (that is, $\gamma_{i}$ represents a sampled marginal focal element). Some of these $\gamma_{i}$ are intervals, some other, points. Inasmuch as every sample of a input variable can be represented by $\gamma_{i}$ or by the corresponding $\alpha_{i}$, the joint focal element can be represented either by the $d$-dimensional box $\boldsymbol{\gamma}:=X_{i=1}^{d} \gamma_{i} \subseteq \mathscr{X}$ or by the point $\boldsymbol{\alpha}:=\left[\alpha_{1}, \alpha_{2}, \ldots, \alpha_{d}\right] \in \boldsymbol{\Omega}^{\prime}$ (see Figure 1).


Figure 1: Focal elements in the $\mathscr{X}$ (Panel a) and in the $\boldsymbol{\Omega}^{\prime}$-space (Panel b), respectively. The focal elements are the realizations of input variables which are depicted either as the points $\boldsymbol{\alpha}$ in the $\boldsymbol{\Omega}^{\prime}$-space or as (multidimensional) boxes, corresponding to the focal element $\Gamma(\boldsymbol{\alpha})$, in the $\mathscr{X}$ space. The figure (a) shows also the failure surface, $g(\boldsymbol{x})=0$, that defines the safe $\mathcal{S}$ and failure $\mathcal{F}$ domains. In the $\boldsymbol{\Omega}^{\prime}$-space (b) are defined the regions $F_{\mathrm{LP}}$ and $F_{\mathrm{UP}}$ together with the failure surfaces $\underline{g}(\boldsymbol{\alpha})=0$ and $\bar{g}(\boldsymbol{\alpha})=0$, where $\underline{g}(\boldsymbol{\alpha}):=\min _{\boldsymbol{x} \in \Gamma(\boldsymbol{\alpha})} g(\boldsymbol{x})$ and $\bar{g}(\boldsymbol{\alpha}):=\max _{\boldsymbol{x} \in \Gamma(\boldsymbol{\alpha})} g(\boldsymbol{x})$. Those boxes in $\mathscr{X}$ which contain at least one point of the failure region $\mathcal{F}$ have a corresponding $\boldsymbol{\alpha}$ point in the region $F_{\mathrm{UP}}$; while those boxes in $\mathscr{X}$ which are completely contained in the region $\mathcal{F}$ have a corresponding $\boldsymbol{\alpha}$ point in the region $F_{\mathrm{LP}}$.

## E. The epistemic and the aleatory spaces

Along this paper, two spaces are defined for modelling the aleatory and the epistemic uncertainties and which are called the aleatory space $\boldsymbol{\Omega}$ and the epistemic space $\boldsymbol{\Theta}$, respectively (see Figure 2 ).

The aleatory space A sample from a random set is obtained by drawing an $\boldsymbol{\alpha} \in \boldsymbol{\Omega}^{\prime}$ from the copula $C$. Since a sample from an interval does not contain an aleatory component, if we strip from space $\boldsymbol{\Omega}^{\prime}$ all those components which belong to intervals, then a subspace $\boldsymbol{\Omega}$ of $\boldsymbol{\Omega}^{\prime}$ is obtained. The subspace $\boldsymbol{\Omega}$ contains only probabilistic information without spurious random variables. This set is called from now on the aleatory space $\boldsymbol{\Omega}$. Without loss of generality all copulas in our discussion will be defined on $\boldsymbol{\Omega}$, and all subsequent discussion will be performed with respect to the set $\boldsymbol{\Omega}$.

The epistemic space The epistemic space $\boldsymbol{\Theta}$ is formed by the Cartesian product of all intervals $\left\{I_{i}, i=\right.$ $1,2, \ldots, q\}$ that contain epistemic uncertainty, that is $\boldsymbol{\Theta}=X_{i=1}^{q} I_{i}$. Since the epistemic uncertainty can be reduced when additional information is available, we will assume that a point $\boldsymbol{\theta}^{*} \in \boldsymbol{\Theta}$ in the epistemic space will represent the "true uncertainty model", which will result once all epistemic uncertainty is removed from $\Theta$. When new information is available, the epistemic space will shrink to a subset of it called the reduced epistemic space.

For example, let's consider a problem with four input variables: two correlated random variables $X$ and $Y$ modelled as a bivariate normal distribution and two independent variables $W$ and $Z$ which are modelled by the intervals $I_{W}$ and $I_{Z}$, correspondingly. The joint CDF of $X$ and $Y$ is defined by the mean vector $\left[\mu_{X}, \mu_{Y}\right]^{T}$, variances $\left[\sigma_{X}^{2}, \sigma_{Y}^{2}\right]^{T}$ and a Pearson correlation coefficient $\rho_{X Y}$. If we assume that all those five parameters are also unknown and represented by intervals, namely, $I_{\mu_{X}}, I_{\mu_{Y}}, I_{\sigma_{X}^{2}}, I_{\sigma_{Y}^{2}}$ and $I_{\rho_{X Y}}$, respectively, then, the aleatory space $\boldsymbol{\Omega}$ is $(0,1]^{2}$ while $\boldsymbol{\Omega}^{\prime}$ is $(0,1]^{4}$; in addition, a Gaussian copula is defined on the aleatory space $\boldsymbol{\Omega}$. Finally, the epistemic space $\boldsymbol{\Theta}$ is a seven-dimensional space formed by the Cartesian product $I_{\mu_{X}} \times I_{\mu_{Y}} \times I_{\sigma_{X}^{2}} \times I_{\sigma_{Y}^{2}} \times I_{\rho_{X Y}} \times I_{W} \times I_{Z}$. Notice that the point $\boldsymbol{\theta}^{*}$ belongs to that space.

## F. The system representation as a function of the aleatory and the epistemic uncertainty

Let us denote by $\mathcal{G}: \mathscr{X} \rightarrow \mathbb{R}$ a function that represents the system; this function maps from the input space $\mathscr{X}$ of input variables to the real line and let $\mathcal{W}: \Omega \times \boldsymbol{\Theta} \rightarrow \mathscr{X}$ be a function which returns the point in $\Gamma(\boldsymbol{\alpha})$ after reducing the epistemic uncertainty in $\boldsymbol{\Theta}$ to $\boldsymbol{\theta}$. The function $\mathcal{W}$ exists only if the random set $\Gamma$ models intervals, CDFs, p-boxes or their joint combination. This function does not exist if $\Gamma$ models Dempster-Shafer structures or possibility distributions, but this is not the case in this paper. Note on the one hand, that the image of $\boldsymbol{\Theta}$ through $\mathcal{W}(\boldsymbol{\alpha} ; \cdot)$ is the focal element $\Gamma(\boldsymbol{\alpha})$; on the other hand, the image of $\boldsymbol{\Omega}$ through $\mathcal{W}(\cdot ; \boldsymbol{\theta})$ can be modelled as a CDF with parameter vector $\boldsymbol{\theta}$, that is, $F(\cdot ; \boldsymbol{\theta})$. Take into account that the definition of function $\mathcal{W}$ uses, in the case of CDFs and p-boxes, the inverse CDF of the input variable in consideration.

We will define the function $\mathcal{H}: \boldsymbol{\Omega} \times \boldsymbol{\Theta} \rightarrow \mathbb{R}$ as $\mathcal{H}=\mathcal{G} \circ \mathcal{W}$, that is, $\mathcal{H}$ represents the system as well, but its domain is the Cartesian product of the aleatory and epistemic spaces (see Figure 2).


Figure 2: Representation of the aleatory and epistemic spaces and their propagation through the model. Here $\boldsymbol{\theta}^{*}$ represents the true uncertainty model after all epistemic uncertainty has been removed. The subset of the epistemic space that appears after new information is available is the reduced epistemic space. The function $\mathcal{W}$ produces a realization for a given $\boldsymbol{\alpha}$ and $\boldsymbol{\theta}$; this output becomes the input for the system $\mathcal{G}$. The composition of both functions forms the function $\mathcal{H}$.

## G. Mapping of a focal element through a system: the extension principle of random sets

The capability to propagate intervals, CDFs, p-boxes, and their combination through a system represents the core of the developed computational framework. In order to find the image of a focal element, $\gamma_{i} \subseteq \mathscr{X}$, through a function $\mathcal{G}: \mathscr{X} \rightarrow \mathbb{R}$, the extension principle of random sets is used (this principle states how to propagate a random set through a function - see Ref. 17). This can be done by means of optimization methods,,${ }^{18}$ sampling methods, ${ }^{19}$ a vertex method, ${ }^{20}$ or the interval arithmetic method. ${ }^{21,3}$ In the following, the optimization and the sampling methods will be explained in detail, since both methods have been employed to solve the NASA UQ challenge problem.

The optimization method If a focal element $\gamma_{i}:=\Gamma\left(\boldsymbol{\alpha}_{i}\right)$ is connected and compact and $\mathcal{G}$ is continuous, then the image of the set $\gamma_{i}$ through $\mathcal{G}$, can be calculated as

$$
\begin{equation*}
\mathcal{G}\left(\Gamma\left(\boldsymbol{\alpha}_{i}\right)\right)=\left[\underline{\mathcal{G}}\left(\boldsymbol{\alpha}_{i}\right), \overline{\mathcal{G}}\left(\boldsymbol{\alpha}_{i}\right)\right] \tag{3}
\end{equation*}
$$

where,

$$
\begin{equation*}
\underline{\mathcal{G}}\left(\boldsymbol{\alpha}_{i}\right):=\min _{\boldsymbol{x}_{i} \in \Gamma\left(\boldsymbol{\alpha}_{i}\right)} \mathcal{G}\left(\boldsymbol{x}_{i}\right) \quad \overline{\mathcal{G}}\left(\boldsymbol{\alpha}_{i}\right):=\max _{\boldsymbol{x}_{i} \in \Gamma\left(\boldsymbol{\alpha}_{i}\right)} \mathcal{G}\left(\boldsymbol{x}_{i}\right) ; \tag{4}
\end{equation*}
$$

are limit state functions defined in $\boldsymbol{\Omega}$. Using the function $\mathcal{H}$ defined in Section F , Eqs. (4) can be written as an optimization over the epistemic space:

$$
\begin{equation*}
\underline{\mathcal{G}}\left(\boldsymbol{\alpha}_{i}\right):=\min _{\boldsymbol{\theta} \in \boldsymbol{\Theta}} \mathcal{H}\left(\boldsymbol{\alpha}_{i}, \boldsymbol{\theta}\right) \quad \overline{\mathcal{G}}\left(\boldsymbol{\alpha}_{i}\right):=\max _{\boldsymbol{\theta} \in \boldsymbol{\Theta}} \mathcal{H}\left(\boldsymbol{\alpha}_{i}, \boldsymbol{\theta}\right) \tag{5}
\end{equation*}
$$

This approach is usually employed when $\mathcal{G}$ is a nonlinear function of the system parameters. The main drawback of this method is that it requires a high computational effort in a complex and large scale system.

The sampling method (random search) The image of the focal element $\gamma_{i}$ through $\mathcal{G}$ can also be estimated using a sampling technique (this should not be confused with the double loop simulation used to deal with distributional p-boxes). Remembering that the focal element $\gamma_{i}$ is a multi-dimensional box; random samples can be generated inside that box and then they are mapped through $\mathcal{G}$; then Eqs. (4) are approximated by the smallest and largest values of the images of those samples. This method is easy to implement but it requires huge number of samples (due to the curse of dimensionality). The sampling method gives does not guarantee that the true minimum and maximum are identified even using a very large number of samples.

## III. Proposed approach for uncertainty management and quantification

The robust design of safety-critical systems requires not only the explicit treatment of different forms and representations of uncertainty but also, performing a number of different tasks. Generally, the design of such systems requires inputs and criteria of different disciplines and one of the main challenges in uncertainty management is how to propagate the uncertainty and understand how the uncertainty in one field affects other disciplines. More specifically,

- the first task required is to refine the current uncertainty model using new available information. This task is often called model updating (see e.g. Refs. 22, 23);
- usually sensitivity analysis is performed for the identification of those parameters whose uncertainty is the most/least consequential. This allows to drive the collection of new data and information focusing on those parameters that affect mostly the variability of the outputs;
- the propagation of mixed aleatory and epistemic uncertainties of the refined/improved model and the extreme-case system performance assessment are performed in order to identify the combinations of parameters that lead to the worst performance;
- finally the design in the presence of uncertainty is achieved. This task is computationally demanding since it requires the propagation of the uncertainty through the system for each candidate solution.

Different tools and approaches exist for uncertainty quantification and characterization that can be potentially used in the design of safety critical systems. Each method is based on some assumptions that often cannot be verified a priori. Moreover, the simulation strategies are able to produce accurate results only if the right set of parameters is selected and this often cannot be verified. Finally, the numerical implementation might contain errors.

For these reasons, it is necessary to perform the analysis using different strategies and hypotheses in order to be able to cross-validate the results. Hence, different strategies implemented in a flexible and open computational framework are briefly summarized in the next sections.

## A. Model updating

The aim of model updating is to reduce the epistemic uncertainty on the output of the model $x=\mathcal{H}(\boldsymbol{\alpha} ; \boldsymbol{\theta})$ based on the availability of a limited set of data (observations) $\mathcal{D}_{e}:=\left\{x_{k}^{e}: k=1,2, \ldots, n_{e}\right\}$. These observations of the "true uncertainty model" $\boldsymbol{\theta}^{*} \in \boldsymbol{\Theta}$ can be used to improve the uncertainty model, i.e. to reduce the original intervals of the epistemic uncertainties by excluding those combinations of parameters that fail to describe the observations as shown in Figure 2. Two different approaches will be used for model updating: a non-parametric model based on some statistical tests and a Bayesian method.

Along the rest of this paper hats $(\hat{F})$ and tildes $(\tilde{F})$ will be used for referring to empirical CDFs and a kernel density estimations of CDFs, respectively.

## 1. Non-parametric statistic method based on the Kolmogorov-Smirnov test

A simple and fast approach to improve the uncertainty model is based on the comparison of the CDFs of the observations of the true uncertainty model and those obtained by means of random combinations of the input parameters in order to identify tighter intervals which form a reduced epistemic space and which are in agreement with the observations.

Let us consider the epistemic space $\boldsymbol{\Theta}$ of the involved variables. Random realizations $\boldsymbol{\theta}_{i}$ in the epistemic space $\boldsymbol{\Theta}$ are generated assuming, for example, a uniform PDF on $\boldsymbol{\Theta}$ (in agreement with the Laplace's principle of indifference). Thereafter the points $\left\{\boldsymbol{\alpha}_{j}, j=1,2, \ldots, n\right\}$ are sampled from the aleatory space $\boldsymbol{\Omega}$ according to the copula $C$ (Nelsen ${ }^{15}$ provides methods to do it), in order to simulate $n$ observations from the system $\mathcal{H}$ as $x_{j}^{i}=\mathcal{H}\left(\boldsymbol{\alpha}_{j}, \boldsymbol{\theta}_{i}\right)$. For a single realization $\boldsymbol{\theta}_{i}$, the Kolmogorov-Smirnov statistic, which is defined as

$$
\begin{equation*}
D_{i}=\sup _{x}\left|\hat{F}\left(x \mid \boldsymbol{\theta}_{i}\right)-\hat{F}_{e}(x)\right| \tag{6}
\end{equation*}
$$

is used to measure the similarity between the CDFs obtained with the sampled set $\left\{x_{j}^{i}, j=1,2, \ldots, n\right\}$ and the set of observations $\mathcal{D}_{e}$. Here $\hat{F}\left(\cdot \mid \boldsymbol{\theta}_{i}\right)$ and $\hat{F}_{e}$ are the empirical CDFs obtained using the random samples drawn according to the epistemic parameters $\boldsymbol{\theta}_{i}$ and the provided experimental data, respectively.

The Kolmogorov-Smirnov test is used to obtain confidence limits on $\hat{F}\left(\cdot \mid \boldsymbol{\theta}_{i}\right)$ by choosing different critical values of the test statistic $D$. This implies that a band of width $\pm D$ around $\hat{F}_{e}(x)$ will entirely contain $\hat{F}\left(\cdot \mid \boldsymbol{\theta}_{i}\right)$ with probability $1-c$. This allows to identify those combinations of epistemic parameters such that $P\left(D_{i}>D\right)=c . \quad c=0$ means that all the CDFs $\hat{F}\left(\cdot \mid \boldsymbol{\theta}_{i}\right)$ are accepted and the refinement of the input intervals is not possible, whereas $c=1$ implies that $\hat{F}\left(\cdot \mid \boldsymbol{\theta}_{i}\right)$ comes exactly from the same model that has generated the target distribution $\tilde{F}_{e}(x)$, i.e. no epistemic uncertainty is present.

The selection of $D$ is a critical task and generally depends of the amount of available information (i.e. number of observations). A practical approach is to use two different data sets that come from the same process to estimate the critical level of the measure of similarity $D_{\hat{v}}$ (using Eq. (6)). The computed validation distance $D_{\hat{v}}$ can be used to set the required confidence level, accepting all the combinations of epistemic parameters with $D_{i}<D_{\hat{v}}$. When an independent validation data set is not available, a cross validation data
set can be constructed to test the model in order to limit problems such as overfitting. This cross validation data set can be obtained by means of re-sampling techniques. ${ }^{24}$ Cross-validation is important to protect against hypotheses suggested by the data ${ }^{25}$ specially where further samples are costly or simply impossible to collect.

The non-parametric approach based on the Kolmogorov-Smirnov test is a simple and fast method for performing uncertainty characterization (and model updating). However, it is important to keep in mind the limitations of the approach. In fact, the method assumes that the measure of similarity $D_{i}$ is distributed according to the Kolmogorov distribution, ${ }^{26}$ which is strictly true only for large sample sets. It is possible to use some smoother techniques such as the Gaussian kernel density estimation to overcome this limitation. Gaussian kernel density estimates for $\mathcal{D}_{e}$ are given by

$$
\begin{equation*}
\tilde{F}_{e}(x)=\frac{1}{n \sigma \sqrt{2 \pi}} \int_{-\infty}^{x} \sum_{j=1}^{n_{e}} \exp \left(-\frac{1}{2}\left(\frac{x^{\prime}-x_{j}^{e}}{\sigma}\right)^{2}\right) \mathrm{d} x^{\prime} ; \tag{7}
\end{equation*}
$$

here $\sigma$ stands for the standard deviation of the Gaussian kernels that represents the smoothing parameter, proportional to the so-called bandwidth. Assuming $x$ is a continuous random variable, for $n_{e} \rightarrow \infty$ the Gaussian kernel density estimate converges to the true underlying density. The support of the associated PDFs $\tilde{f}_{e}(x)$ (i.e. $\left\{x: \tilde{f}_{e}(x)>0\right\}$ ) and the bandwidth of the kernel have strong influence on the resulting estimate. We suggest to use the approach in Ref. 27 to estimate the support of the PDF and Silverman's rule of thumb ${ }^{28}$ to estimate the bandwidth of the kernels. Using realizations from Eq. (7) the measure of similarity can be calculated via Eq. (6) where $\hat{F}_{e}(x)$ is replaced by $\tilde{F}_{e}(x)$. Please note that the Gaussian kernels can be used to define a new critical measure level indicated with $D_{\tilde{v}}$.

To summarize, the following pseudo-algorithm is used:

1. Estimate the parameters $\sigma$ and the Gaussian kernel $\operatorname{CDF} \tilde{F}_{e}$ using Eq. (7);
2. Estimate $D_{\hat{v}}$ and $D_{\hat{v}}$;
3. Generate realizations on the epistemic space, $\boldsymbol{\theta}_{i}$;
4. Draw $n$ points from the aleatory space $\boldsymbol{\Omega}$, using copula $C$; we will call these samples $\left\{\boldsymbol{\alpha}_{j}: j=1, \ldots, n\right\}$;
5. Evaluate the model $x_{j}^{i}:=\mathcal{H}\left(\boldsymbol{\alpha}_{j} ; \boldsymbol{\theta}_{i}\right)$ for $j=1, \ldots, n$;
6. Estimate the empirical CDF $\hat{F}\left(\cdot \mid \boldsymbol{\theta}_{i}\right)$ of the set of samples $\left\{x_{j}^{i}, j=1,2, \ldots, n\right\}$;
7. Using Eq. (6), compute the measure of similarity $D_{i}$;
8. If $D_{i}<D_{\tilde{v}}$ (or $D_{i}<D_{\hat{v}}$ ) collect $\boldsymbol{\theta}_{i}$. The set of collected points identify a reduced space in the original epistemic space.

## 2. Bayesian updating on the epistemic space

Bayesian inference is a statistical method in which the Bayes' rule is used to update the probability estimate for a hypothesis as additional information is available.

Suppose we are given a set of observed data points $\mathcal{D}_{e}:=\left\{x_{k}^{e}: k=1,2, \ldots, n_{e}\right\}$ called the evidence, and which are sampled from a PDF $p\left(\cdot ; \boldsymbol{\theta}^{*}\right)$ which belongs to a certain family of PDFs $\{p(\cdot ; \boldsymbol{\theta}): \boldsymbol{\theta} \in \boldsymbol{\Theta}\}$ called the parametric model. The idea of Bayesian inference is to update our belief about the vector of parameters $\boldsymbol{\theta}$ provided that $\boldsymbol{\theta}^{*}$, the true set of parameters of the PDF, is unknown. Bayes' theorem updates that belief using two antecedents:

- a prior PDF $p(\boldsymbol{\theta})$, which indicates all available knowledge about $\boldsymbol{\theta}^{*}$ before the evidence $\mathcal{D}_{e}$ is observed;
- and the likelihood function $P\left(\mathcal{D}_{e} \mid \boldsymbol{\theta}\right)$, which is a function related to the probability of observing the samples $\mathcal{D}_{e}$ assuming that the true parameter underlying the model PDF $p(x ; \boldsymbol{\theta})$ is $\boldsymbol{\theta}$; it is defined as

$$
\begin{equation*}
P\left(\mathcal{D}_{e} \mid \boldsymbol{\theta}\right)=\prod_{k=1}^{n_{e}} p\left(x_{k}^{e} ; \boldsymbol{\theta}\right) \tag{8}
\end{equation*}
$$

when a set of independent and identically distributed observations $\mathcal{D}_{e}$ is available. Please note that in practice (i.e. for the numerical implementation) the log-likelihood is used instead of the likelihood.

The updated belief about the vector of parameters $\boldsymbol{\theta}$ after observing the evidence $\mathcal{D}_{e}$, is modelled by the so-called posterior PDF $p\left(\boldsymbol{\theta} \mid \mathcal{D}_{e}\right)$ which is calculated by:

$$
\begin{equation*}
p\left(\boldsymbol{\theta} \mid \mathcal{D}_{e}\right)=\frac{P\left(\mathcal{D}_{e} \mid \boldsymbol{\theta}\right) p(\boldsymbol{\theta})}{P\left(\mathcal{D}_{e}\right)} \tag{9}
\end{equation*}
$$

where the probability of the evidence,

$$
\begin{equation*}
P\left(\mathcal{D}_{e}\right)=\int_{\boldsymbol{\Theta}} P\left(\mathcal{D}_{e} \mid \boldsymbol{\theta}\right) p(\boldsymbol{\theta}) \mathrm{d} \boldsymbol{\theta} \tag{10}
\end{equation*}
$$

can be understood as a normalizing constant. Bayesian updating hopes that after using the evidence $\mathcal{D}_{e}$ the posterior PDF $p\left(\boldsymbol{\theta} \mid \mathcal{D}_{e}\right)$ is sharply peaked about the true value of $\boldsymbol{\theta}^{*}$. We will update our belief about the true set of parameters $\boldsymbol{\theta}^{*} \in \boldsymbol{\Theta}$ propagating the evidence through the Bayes' equation numerically. Samples of the posterior PDF can be generated without the necessity to evaluate $p\left(\boldsymbol{\theta} \mid \mathcal{D}_{e}\right)$, using an algorithm called Transitional Markov Chain Monte Carlo (TMCMC). ${ }^{29}$

As the prior PDF, we will use a uniform distribution on the epistemic space $\boldsymbol{\Theta}$, that is $\boldsymbol{\theta} \sim \operatorname{Unif}(\boldsymbol{\Theta})$, in accordance to the Laplace's principle of indifference (or more generally, the principle of maximum entropy).

Different likelihood functions can be used, based on different mathematical assumptions; in the following two methods will be proposed: a method that used a kernel density estimator to approximate $p\left(\cdot \mid \boldsymbol{\theta}_{i}\right)$ and a approximate Bayesian computational method.

Bayesian computational method In this case, the likelihood is estimated through kernel density. Assuming that the samples $\mathcal{D}_{e}$ were drawn from $p\left(x ; \boldsymbol{\theta}_{i}\right)$, the likelihood $P\left(\mathcal{D}_{e} \mid \boldsymbol{\theta}_{i}\right)$ is defined in the following way:

1. Draw $n$ points, $\left\{\boldsymbol{\alpha}_{j}: j=1, \ldots, n\right\}$, from the aleatory space $\boldsymbol{\Omega}$, using copula $C$;
2. Calculate $x_{j}^{i}:=\mathcal{H}\left(\boldsymbol{\alpha}_{j} ; \boldsymbol{\theta}_{i}\right)$ for $j=1, \ldots, n$;
3. Using kernel density estimation and the samples $\left\{x_{j}^{i}: j=1, \ldots, n\right\}$, estimate the $\operatorname{CDF} \tilde{F}\left(\cdot \mid \boldsymbol{\theta}_{i}\right)$ and its associated PDF $\tilde{p}\left(x \mid \boldsymbol{\theta}_{i}\right) \equiv p\left(x ; \boldsymbol{\theta}_{i}\right)$. This step is required because $\tilde{p}\left(x \mid \boldsymbol{\theta}_{i}\right)$ cannot be obtained analytically;
4. Calculate the likelihood function $P\left(\mathcal{D}_{e} \mid \boldsymbol{\theta}_{i}\right)$ as in Eq. (8).

Approximate Bayesian computational method The likelihood calculated by means of the "Bayesian computational method" applies Bayes' theorem directly and without strong assumptions. However it requires a large number of model evaluations and a relative larger data set to converge. ${ }^{30}$ Recently, approximate Bayesian computational methods have been proposed to reduce the computational costs of the expensive or intractable likelihood function. ${ }^{31,32}$ The likelihood can be for instance approximated with the following expression:

$$
\begin{equation*}
P\left(\mathcal{D}_{e} \mid \boldsymbol{\theta}_{i}\right)=\prod_{k=1}^{n_{e}} \frac{1}{\sqrt{2 \pi} \sigma} \exp \left(-\frac{1}{2}\left(\frac{\delta_{k}}{\sigma}\right)^{2}\right) \tag{11}
\end{equation*}
$$

where $\delta_{k}$ is the absolute value of the difference between the empirical $\operatorname{CDF} \hat{F}\left(\cdot \mid \boldsymbol{\theta}_{i}\right)$ obtained for an individual realization $\boldsymbol{\theta}_{i}$ of the epistemic space $\boldsymbol{\Theta}$, evaluated at each point $\left\{x_{k}^{e}, k=1,2, \ldots, n_{e}\right\}$ and the empirical CDFs of the experimental dataset $\mathcal{D}_{e}$, that is:

$$
\begin{equation*}
\delta_{k}=\left|\hat{F}\left(x_{k} \mid \boldsymbol{\theta}_{i}\right)-\hat{F}_{e}\left(x_{k}^{e}\right)\right| \tag{12}
\end{equation*}
$$

for $k=1,2, \ldots, n_{e}$. Please note that the Bayesian updating approach is generally applied to identify a fixed estimate of $\boldsymbol{\theta}$ as close as possible to $\boldsymbol{\theta}^{*}$. Here, the approach has been used to identify a reduced epistemic space containing the true values of the unknown parameters. If a constant $\sigma$ is used, the Bayesian updating formulation, here introduced, is equivalent to a minimization in the least square sense of the distance between the CDFs $\hat{F}\left(\cdot \mid \boldsymbol{\theta}_{i}\right)$ and $\hat{F}_{e}$. However the value of $\sigma$ is unknown and hence it represents an additional parameter that needs to be estimated. ${ }^{33}$

This last approach is indeed based more on practical considerations than on a sound mathematical basis, and is open to criticisms since the differences $\delta_{k}$ are assumed to be independent and normally distributed with zero mean and unit variance, and that even though $\delta_{k}$ is normally distributed, it will only take values in the interval $[0,1]$ since the CDF ranges between 0 and 1.

Using the above defined prior PDF and likelihood functions, the TMCMC algorithm ${ }^{29}$ is employed in order to find samples of the posterior $p\left(\boldsymbol{\theta} \mid \mathcal{D}_{e}\right)$. The likelihood $P\left(\mathcal{D}_{e} \mid \boldsymbol{\theta}_{i}\right)$ is calculated, using the approximate Bayesian computational method, by the following procedure:

1. Draw $n$ points $\left(\left\{\boldsymbol{\alpha}_{j}: j=1, \ldots, n\right\}\right)$ from the aleatory space $\boldsymbol{\Omega}$, using copula $C$;
2. Calculate $x_{j}^{i}:=\mathcal{H}\left(\boldsymbol{\alpha}_{j} ; \boldsymbol{\theta}_{i}\right)$ for $j=1, \ldots, n$;
3. Using the samples $\left\{x_{j}^{i}: j=1, \ldots, n\right\}$, estimate the empirical CDF $\hat{F}\left(\cdot \mid \boldsymbol{\theta}_{i}\right)$;
4. Compute $\delta_{k}=\left|\hat{F}\left(x_{k}^{e} \mid \boldsymbol{\theta}_{i}\right)-\hat{F}_{e}\left(x_{k}^{e}\right)\right|$ at each point $x_{k}^{e} \in \mathcal{D}_{e}$;
5. Calculate the likelihood function $P\left(\mathcal{D}_{e} \mid \boldsymbol{\theta}_{i}\right)$ as in Eq. (11).

## B. Sensitivity analysis

The aim of sensitivity analysis is to identify and rank the parameters that contribute mostly to the variability of the output of a system $\mathcal{H}$. Two approaches can be used: the Hartley-like measure of nonspecificity and
the global sensitivity analysis based on Sobol' and total sensitivity measures. Both approaches can be used to perform global sensitivity analysis in presence of epistemic uncertainty.

## 1. Nonspecificity technique

Before delving into this method of sensitivity analysis, a small introduction to the nonspecificity measure is presented.

The nonspecificity, proposed by Klir and coworkers, ${ }^{34,35}$ is a measure of the amount of information required to remove the epistemic uncertainty; it is used in cases when we have to select a unique element from a set, but we are totally indifferent about which element of the provided ones to choose.

The nonspecificity is based on the so called Hartley-like measure, which for a d-dimensional box (or focal element) $A=X_{i=1}^{d}\left[l_{i}, u_{i}\right]$, like the ones that we are considering in this paper is given by:

$$
\begin{equation*}
\operatorname{HL}(A)=\log _{2}\left(\prod_{i=1}^{d}\left(1+u_{i}-l_{i}\right)\right) . \tag{13}
\end{equation*}
$$

The nonspecificity of a random set with an infinite number of focal elements is given by (see Ref. 36):

$$
\begin{equation*}
\mathrm{HL}\left(\left(\mathscr{F}, P_{\Gamma}\right)\right)=\int_{\boldsymbol{\Omega}} \mathrm{HL}(\Gamma(\boldsymbol{\alpha})) \mathrm{d} C(\boldsymbol{\alpha}) ; \tag{14}
\end{equation*}
$$

two special cases of Eq. (14) are:

- the nonspecificity of a Dempster-Shafer structure $\left(\mathscr{F}_{n}, m\right)$ with focal set $\mathscr{F}_{n}=\left\{A_{1}, \ldots, A_{n}\right\}$ and basic mass assignment $m$ :

$$
\begin{equation*}
\operatorname{HL}\left(\left(\mathscr{F}_{n}, m\right)\right)=\sum_{i=1}^{n} \operatorname{HL}\left(A_{i}\right) m\left(A_{i}\right) ; \tag{15}
\end{equation*}
$$

- the nonspecificity of a distribution-free probability box $\langle\underline{F}, \bar{F}\rangle$ :

$$
\begin{equation*}
\mathrm{HL}(\langle\underline{F}, \bar{F}\rangle)=\int_{0}^{1} \log _{2}\left(1+\underline{F}^{-1}(\alpha)-\bar{F}^{-1}(\alpha)\right) \mathrm{d} \alpha \tag{16}
\end{equation*}
$$

The nonspecificity is a measure of epistemic uncertainty, and in consequence, it is useful for assessing the variability of the output due to the epistemic uncertainty in the input of the model.

The method, which is detailed in Ref. 37, calculates a Dempster-Shafer structure that is the result of propagating the epistemic uncertainty through the system $\mathcal{H}$ (using the extension principle for random sets). Then, the Hartley-like measure of nonspecificity of that output Dempster-Shafer structure is evaluated. More specifically:

1. $n$ samples $\boldsymbol{\alpha}_{i} \in \boldsymbol{\Omega}$ are drawn from copula $C$. Thereafter, the image of the focal element $\boldsymbol{\alpha}_{i}$ through $\mathcal{H}$ is calculated by means of Eq. (5) as $\left[\min _{\boldsymbol{\theta} \in \boldsymbol{\Theta}} \mathcal{H}\left(\boldsymbol{\alpha}_{i}, \boldsymbol{\theta}\right), \max _{\boldsymbol{\theta} \in \boldsymbol{\Theta}} \mathcal{H}\left(\boldsymbol{\alpha}_{i}, \boldsymbol{\theta}\right)\right]$. This set of $n$ focal elements is used to construct a Dempster-Shafer structure $\left(\mathscr{F}_{n}, m\right)$ (as explained in Section II-C) which represents the propagation of the aleatory and epistemic uncertainty through the system. The basic mass assignment $m$ of each focal element is $1 / n$.
2. Compute $\mathrm{HL}_{\text {unpinched }}=\operatorname{HL}\left(\left(\mathscr{F}_{n}, m\right)\right)$ according to Eq. (15).
3. Consider a set of points $\left\{p_{r}: 0 \leq p_{1}<p_{2}<\ldots<p_{r}<\ldots<p_{z} \leq 1\right\}$ which are evenly distributed in the interval $[0,1]$.
4. For each point $p_{r}$, do the following:
(a) Each interval, $I_{j}=\left[l_{j}, u_{j}\right]$ that forms the epistemic space $\boldsymbol{\Theta}$, is pinched (or reduced) to the value given by $l_{j}+p_{r} \cdot\left(u_{j}-l_{j}\right)$, while leaving all other intervals unchanged (take into account that pinching of groups of input variables can be performed as well). After pinching the $j$-th input variable, a subset of $\boldsymbol{\Theta}$, namely $\boldsymbol{\Theta}_{r j}$, which includes the pinched inputs is formed.
(b) $n$ samples $\boldsymbol{\alpha}_{i} \in \boldsymbol{\Omega}$ are drawn from copula $C$; thereafter, the image of the focal element $\boldsymbol{\alpha}_{i}$ through $\mathcal{H}$ is calculated by means of Eq. (5) as $\left[\min _{\boldsymbol{\theta} \in \boldsymbol{\Theta}_{r j}} \mathcal{H}\left(\boldsymbol{\alpha}_{i}, \boldsymbol{\theta}\right), \max _{\boldsymbol{\theta} \in \boldsymbol{\Theta}_{r j}} \mathcal{H}\left(\boldsymbol{\alpha}_{i}, \boldsymbol{\theta}\right)\right]$. This set of $n$ focal elements is used to construct a Dempster-Shafer structure $\left(\mathscr{F}_{n}^{r j}, m\right)$ for each pinching (as explained in Section II). The basic mass assignment $m$ of each focal element is $1 / n$.
(c) The nonspecificity $\mathrm{HL}_{r j}:=\mathrm{HL}\left(\left(\mathscr{F}_{n}^{r j}, m\right)\right)$ of each of those output Dempster-Shafer structures is computed, as in Eq. (15).
(d) The nonspecificity measure of the output Dempster-Shafer structure is normalized against the nonspecificity measure computed before pinching. In this way, the index $I_{r j}=\mathrm{HL}_{r j} / \mathrm{HL}_{\text {unpinched }}$ is calculated.
5. The index $I_{j}$ is calculated as the mean square of all indexes $I_{r j}$ (remember that $\left.\mathrm{E}\left[I_{j}^{2}\right]=\mathrm{E}\left[I_{j}\right]^{2}+\operatorname{Var}\left[I_{j}\right]\right)$; those indexes are ranked according to their mean square.
$I_{j}$ is used as a measure of the propagation of the epistemic uncertainty to the output of the system $\mathcal{H}$. The smaller $I_{j}$ is, the larger is the sensitivity of the system to the epistemic uncertainty in the input variable
$j$. The mean squared has been chosen as a ranking criterion in order to account for not only the bias but also the variance of the estimator. Note that a precise estimation of $I_{j}$ is not necessary, since only the ranking of the variables $I_{j}$ is required; therefore, $n$ is usually a small number.

## 2. Global Sensitivity analysis

The second approach is based on global sensitivity analysis to estimate the Sobol' and the total indices. ${ }^{38}$ The global sensitivity approach cannot be applied directly to solve the problems where the uncertainty is described as a distributional/free p-boxes and intervals. In fact, this method requires the exact knowledge of the PDF of the input variables and the variance of a measurable model output. In consequence, an alternative mathematical model to $\mathcal{H}$ (as defined in Section II-F) has to be defined in the next.

Consider a model $\mathcal{H}^{*}: \boldsymbol{\Theta} \rightarrow \mathbb{R}$, that is, $Y=\mathcal{H}^{*}(\boldsymbol{\theta})$, where $\boldsymbol{\theta}=\left[\theta_{1}, \ldots, \theta_{q}\right]$ is a vector of random variables and $Y$ is a chosen univariate model output.

Let us associate $\boldsymbol{\Theta}$ with the epistemic space; for a given value of $\boldsymbol{\theta}_{j} \in \boldsymbol{\Theta}$, the function $\mathcal{H}^{*}$ returns the area between a CDF $F\left(\cdot \mid \boldsymbol{\theta}_{j}\right)$ and a reference $\operatorname{CDF} F(\cdot \mid \overline{\boldsymbol{\theta}})$ :

$$
\begin{equation*}
y_{i}:=\mathcal{H}^{*}\left(\boldsymbol{\theta}_{i}\right):=\int_{-\infty}^{+\infty}\left|F\left(x \mid \boldsymbol{\theta}_{i}\right)-F(x \mid \overline{\boldsymbol{\theta}})\right| \mathrm{d} x \tag{17}
\end{equation*}
$$

here $\overline{\boldsymbol{\theta}}$ denotes the center of gravity of $\boldsymbol{\Theta}$ (in other words, $\overline{\boldsymbol{\theta}}$ is a vector formed by the mean value of each input epistemic parameter), and $F(\cdot \mid \boldsymbol{\theta})$ represents the CDF obtained after mapping all aleatory uncertainty through the system $\mathcal{H}$, for a given set of epistemic parameters $\boldsymbol{\theta}$. Since the global sensitivity analysis is based on the variance decomposition, any reference CDF can be used in the model $\mathcal{H}^{*}$.

The procedure to estimate the empirical $\operatorname{CDF} \hat{F}(\cdot \mid \boldsymbol{\theta})$ as an approximation to $F(\cdot \mid \boldsymbol{\theta})$ is as follows:

1. Draw $n$ points $\left(\left\{\boldsymbol{\alpha}_{j}: j=1, \ldots, n\right\}\right)$ from the aleatory space $\boldsymbol{\Omega}$, using copula $C$;
2. Evaluate the model $x_{j}^{i}:=\mathcal{H}\left(\boldsymbol{\alpha}_{j} ; \boldsymbol{\theta}\right)$ for $j=1, \ldots, n$;
3. Estimate the empirical CDF $\hat{F}(\cdot \mid \boldsymbol{\theta})$ of the set of samples $\left\{x_{j}^{i}, j=1,2, \ldots, n\right\}$.

Using the above procedure, a sample from the random variable $Y$, namely $y_{i}$ can be estimated by means of Eq. (17) for each realization of input $\boldsymbol{\theta}_{\boldsymbol{i}}$ by using the empirical CDFs $\hat{F}(\cdot \mid \boldsymbol{\theta})$ and $\hat{F}(\cdot \mid \overline{\boldsymbol{\theta}})$. Please note that when the model produces a scalar value for each realization of the input $\boldsymbol{\Theta}_{i}$, e.g. it returns the expected
value $\left(y_{i}=E\left(x \mid \boldsymbol{\theta}_{\boldsymbol{i}}\right)\right)$ or a quantile of a distribution, it is not necessary to evaluate Eq. (17) but the model output can be used directly (see Section B-2).

Finally, the first order Sobol' indices are calculated as follows ${ }^{39}$

$$
\begin{equation*}
S_{i}=\frac{\operatorname{Var}_{\theta_{i}}\left[\mathrm{E}_{\boldsymbol{\theta} \sim i}\left(Y \mid \theta_{i}\right)\right]}{\operatorname{Var}[Y]} \tag{18}
\end{equation*}
$$

where $\operatorname{Var}[Y]$ represents the unconditional variance of the quantity of interest and $\operatorname{Var}_{\theta_{i}}\left[\mathrm{E}_{\boldsymbol{\theta} \sim i}\left(Y \mid \theta_{i}\right)\right]$ the variance of conditional expectation. The total sensitivity index, $T_{i}$, measures the contribution to the output variance of $\theta_{i}$ of the input factors including all interactions with any other input variables,

$$
\begin{equation*}
T_{i}=1-\frac{\operatorname{Var}_{\boldsymbol{\theta}_{\sim i}}\left(\mathrm{E}_{\theta_{i}}\left(Y \mid \boldsymbol{\theta}_{\sim i}\right)\right)}{\operatorname{Var}(Y)} \tag{19}
\end{equation*}
$$

Note that unlike the first order indices, the sum of total indices can exceed one.

The proposed approach allows to decompose the variance of the output $Y$ into parts attributable to the variance of the input variables $\boldsymbol{\theta}$; in other words, it allows to identify and rank the contribution of the epistemic uncertainty, i.e. interval of the parameters, on the p-boxes of quantity of interest. The magnitude of the sensitivity indices are proportional to the contribution to the output variance, i.e. input factor associated with a large sensitivity index contributes most to the variance of the output. Hence, adopting the approach proposed here, the global sensitivity analysis allows us to identify the contribution of the epistemic uncertainty of input factors on the variance of the model.

Different techniques exist to compute the sensitivity indices such as the extended-"Fourier Amplitude Sensitivity Test" (FAST) ${ }^{40,41}$ and the Saltelli's method. ${ }^{38}$ The FAST method allows to estimate first order Sobol' indices, whereas Saltelli's method computes also the total indices.

## C. Uncertainty Propagation

The focus of the uncertainty propagation analysis is to quantify the effect of the uncertain model parameters on quantities of interest such as the mean, variance and quantiles of the system's response or its failure probability. The generalized probabilistic model makes the UQ rather challenging task in terms of computational cost. The challenge is to compute the lower and upper bounds of the quantities of interest. Monte Carlo method remains the most versatile and simple tool to propagate epistemic and aleatory uncertainty.

1. Optimization in the epistemic space (standard approach)

In this approach, the quantity of interest (e.g. mean or failure probability estimation) defines the objective function; and the bounds on that objective function are calculated by means of a global search in the epistemic space $\boldsymbol{\Theta}$. On one hand, the lower and upper bounds of the mean are obtained as:

$$
\begin{equation*}
\underline{\mu}=\min _{\boldsymbol{\theta} \in \boldsymbol{\Theta}} \mu(\boldsymbol{\theta}) \quad \bar{\mu}=\max _{\boldsymbol{\theta} \in \boldsymbol{\Theta}} \mu(\boldsymbol{\theta}) \tag{20}
\end{equation*}
$$

where the mean of the response model is given by:

$$
\begin{equation*}
\mu(\boldsymbol{\theta})=\int_{\boldsymbol{\Omega}} \mathcal{H}(\boldsymbol{\alpha} ; \boldsymbol{\theta}) \mathrm{d} C(\boldsymbol{\alpha}) \tag{21}
\end{equation*}
$$

On the other hand, the lower and upper bound of the failure probability, defined as the excedance of a critical threshold level $\mathcal{H}^{\text {crit }}$ of the model response, are obtained as

$$
\begin{equation*}
\underline{P_{f}}=\min _{\boldsymbol{\theta} \in \boldsymbol{\Theta}} P_{f}(\boldsymbol{\theta}) \quad \overline{P_{f}}=\max _{\boldsymbol{\theta} \in \boldsymbol{\Theta}} P_{f}(\boldsymbol{\theta}) \tag{22}
\end{equation*}
$$

here $P_{f}(\boldsymbol{\theta})$ stands for the failure probability, that is,

$$
\begin{equation*}
P_{f}(\boldsymbol{\theta}):=\int_{\Omega} \mathcal{I}\left[\mathcal{H}(\boldsymbol{\alpha} ; \boldsymbol{\theta})>\mathcal{H}^{c r i t}\right] \mathrm{d} C(\boldsymbol{\alpha}) \tag{23}
\end{equation*}
$$

Monte Carlo method is used to calculate the bounds Eq. (20) and Eq. (22), by means of a double loop simulation:

- The outer loop drives an optimization/search process in the epistemic space $\boldsymbol{\Theta}$ to identify the lower and upper bounds Eq. (20) and Eq. (22). This search is performed by Monte Carlo sampling taking into account that this optimization method is very inefficient in high dimensional spaces since the search space grows exponentially with the number of variables. Better optimization strategies such as Genetic Algorithms can also be adopted as shown in Section V-C.
- The inner loop propagates the aleatory uncertainty and estimates the statistical quantities of interest (e.g. expected value, failure of probability, CDF, etc). In this way, several $\boldsymbol{\alpha}_{j}$ are sampled from copula $C$ in order to estimate integrals of Eq. (21) and Eq. (23). Take into account that this Monte Carlo integration in the aleatory space $\boldsymbol{\Omega}$ is insensitive to the dimensionality of the problem although it can
be inefficient in case of the calculation of integral of Eq. (23), when the probability of failure is very small. The estimation of the integrals can be speed up by adopting the so called Advanced Monte Carlo methods such as Importance Sampling, Subset Simulation and Line Sampling. ${ }^{19}$


## 2. Propagation of focal sets (counter approach)

The second approach for uncertainty propagation, which is proposed in Refs. 11, 12, 42, is based on the propagation of focal sets through a function. Using random set theory, as explained in Section II, it can be seen that the aleatory space $\boldsymbol{\Omega}$ contains the regions $F_{\mathrm{LP}}:=\{\boldsymbol{\alpha} \in \boldsymbol{\Omega}: \Gamma(\boldsymbol{\alpha}) \subseteq F, \Gamma(\boldsymbol{\alpha}) \neq \emptyset\}$ and $F_{\mathrm{UP}}:=\{\boldsymbol{\alpha} \in \boldsymbol{\Omega}: \Gamma(\boldsymbol{\alpha}) \cap F \neq \emptyset\}$ which are correspondingly formed by all those points whose respective focal elements are completely contained in the failure set $F=\left\{\boldsymbol{x} \in \mathscr{X}: g(\boldsymbol{x})>\mathcal{H}^{\text {crit }}\right\}$ or have in common at least one point with $F$ correspondingly (see Figure 1b). Notice that the set $F$ is defined in the space of input variables $\mathscr{X}$; in this case, the lower Eq. (2a) and upper Eq. (2b) probability measures of $F$ can be calculated by:

$$
\begin{equation*}
\underline{\underline{P_{f}}}=\mathrm{LP}_{\left(\mathscr{F}, P_{\Gamma}\right)}(F)=\int_{\boldsymbol{\Omega}} \mathcal{I}\left[\boldsymbol{\alpha} \in F_{\mathrm{LP}}\right] \mathrm{d} C(\boldsymbol{\alpha}) \quad \overline{\overline{P_{f}}}=\mathrm{UP}_{\left(\mathscr{F}, P_{\Gamma}\right)}(F)=\int_{\boldsymbol{\Omega}} \mathcal{I}\left[\boldsymbol{\alpha} \in F_{\mathrm{UP}}\right] \mathrm{d} C(\boldsymbol{\alpha}) \tag{24}
\end{equation*}
$$

provided that $F_{\mathrm{LP}}$ and $F_{\mathrm{UP}}$ are $\mu_{C}$-measurable sets; here $\mathcal{I}$ stands for the indicator function.
Eq. (24) can be evaluated by means of simple Monte Carlo method sampling $n$ points from the copula $C$, namely $\boldsymbol{\alpha}_{1}, \boldsymbol{\alpha}_{2}, \ldots, \boldsymbol{\alpha}_{n} \in \boldsymbol{\Omega}$, and then retrieving the corresponding focal elements $\gamma_{j}:=\Gamma\left(\boldsymbol{\alpha}_{j}\right), j=1, \ldots, n$ from $\mathscr{F}$. Afterwards, integrals Eq. (24) are computed by the unbiased estimators $\underline{\hat{P}_{f}}$ and $\hat{\overline{P_{f}}}$, which are given by:

$$
\begin{equation*}
\underline{\underline{\hat{P}_{f}}}=\frac{1}{n} \sum_{j=1}^{n} \mathcal{I}\left[\boldsymbol{\alpha}_{j} \in F_{\mathrm{LP}}\right] \quad \hat{\overline{\overline{P_{f}}}}=\frac{1}{n} \sum_{j=1}^{n} \mathcal{I}\left[\boldsymbol{\alpha}_{j} \in F_{\mathrm{UP}}\right] \tag{25}
\end{equation*}
$$

The image of $\Gamma\left(\boldsymbol{\alpha}_{i}\right)$ through the function $\mathcal{G}$ can be computed using the optimization method, as described by equations (3) and (4). Since, $\mathcal{I}\left[\mathcal{G}\left(\Gamma\left(\boldsymbol{\alpha}_{i}\right)\right) \subseteq F\right]=\mathcal{I}\left[\overline{\mathcal{G}}\left(\boldsymbol{\alpha}_{i}\right)>\mathcal{H}^{c r i t}\right]=\mathcal{I}\left[\boldsymbol{\alpha}_{i} \in F_{\mathrm{LP}}\right]$ and $\mathcal{I}\left[\mathcal{G}\left(\Gamma\left(\boldsymbol{\alpha}_{i}\right)\right) \cap F \neq\right.$ $\emptyset]=\mathcal{I}\left[\underline{\mathcal{G}}\left(\boldsymbol{\alpha}_{i}\right)>\mathcal{H}^{c r i t}\right]=\mathcal{I}\left[\boldsymbol{\alpha}_{i} \in F_{\mathrm{UP}}\right]$ it follows that Eqs. (25) can be written as:

$$
\begin{equation*}
\underline{\underline{\hat{P}_{f}}}=\frac{1}{n} \sum_{i=1}^{n} \mathcal{I}\left[\overline{\mathcal{G}}\left(\boldsymbol{\alpha}_{i}\right)>\mathcal{H}^{c r i t}\right] \quad \hat{\overline{\overline{P_{f}}}}=\frac{1}{n} \sum_{i=1}^{n} \mathcal{I}\left[\underline{\mathcal{G}}\left(\boldsymbol{\alpha}_{i}\right)>\mathcal{H}^{c r i t}\right] \tag{26}
\end{equation*}
$$

Observe that this approach operates by inverting the order of execution of the loops in the double loop described above:

- the outer loop propagates the aleatory uncertainty by sampling the points $\boldsymbol{\alpha}_{1}, \boldsymbol{\alpha}_{2}, \ldots, \boldsymbol{\alpha}_{n} \in \boldsymbol{\Omega}$ using copula $C$.
- the inner loop drives an optimization/search process in $\Gamma\left(\boldsymbol{\alpha}_{i}\right)$ in order to find the image of the input focal element through the system $\mathcal{G}$; this step is performed when evaluating Eqs. (3) and (4).

One of the main advantages of the random set theory is that, for a problem where inputs are defined using any possible imprecise probability framework (CDFs, intervals, distribution-free probability boxes, possibility distributions, Dempster-Shafer structures, etc.), it allows to employ the methods developed by the community of stochastic mechanics for estimating the failure probabilities of the two limit state functions $\underline{\mathcal{G}}$ and $\overline{\mathcal{G}}$, i.e. calculating of bounds on probability [ $\left[\underline{\underline{\hat{P}_{f}}}, \overline{\overline{\overline{P_{f}}}}\right]$. In case that the calculation of very small probability bounds is requested, the plain Monte Carlo simulation described here is not efficient. Advanced Monte Carlo methods can be used to estimate small probabilities of failure as described in e.g. Ref. 43,19.

It is worth noting that although the random set theory is in general not applicable in the case of distributional p-boxes, the method presented in Section C-1 can still be used as far as the bounding CDFs of the input p-boxes can be identified. However, applying this approach to distributional p-boxes treats those p-boxes as distribution-free ones. This inevitably leads to loss of information which results in the underestimation and overestimation of the lower and upper bounds respectively, when compared to the method of optimization in the epistemic space (standard approach).

## 3. Numerical considerations

Two degrees of error can be identified using both approaches for UQ. The first error concerns the estimation of the statistics and failure probability, which can be reduced by increasing the number of samples or by implementing an efficient sampling technique. ${ }^{43,19}$ In reliability analysis, the limited set of samples may lead to both an underestimation and to an overestimation. The confidence of the estimator can always be improved adopting a larger set of samples but at the cost of increasing the computational demand.

The second error concerns the global search. In general it is not possible to guarantee the identification of global optima. Only when the feasible (search) domain of the input variables is small ( $\approx 5$ variables), a
thorough search can lead to a good approximation of the global optima. The search error can only affect the results in one direction. For example, if a global minimum is searched, the identified minimum can only be greater than the global one; in the same way, the identified maximum can only be smaller than the global maximum.

Under the assumption that the sampling error for estimating the failure probability is very small, the "optimization in the epistemic space" approach (Section C-1) always results in an overestimation of the lower bound and an underestimation of the upper bound, which may lead to an optimistic decision.

## D. Extreme case analysis

The extreme case analysis consists in identifying the the combinations of epistemic realizations $\boldsymbol{\theta}$ that leads to the worst/best behaviour of the system. This analysis can be seen as an inverse problem of the uncertainty propagation, the forward problem, described in Section C.

This problem is a by-product of the uncertainty propagation but the ability to solve it depends on the approach used to perform the forward problem. The extreme case analysis can not be performed using the approach "propagation of focal sets" presented in Section C. This is because distributional p-boxes are treated as distribution-free p-boxes. Hence, extreme cases might result associated with distributions that lay inside the p-boxes but that do not comply with the associated parental distributions. Only the approach "optimization of the epistemic space" can be used because the approach holds a bijective mapping between the inputs in the epistemic domain and the quantity of interest.

Solving Eqs. (20) and (23)) it is possible to identify directly realizations of the epistemic space $\boldsymbol{\theta}$ that produce the bounds of quantity of interest, as

$$
\begin{align*}
& \boldsymbol{\theta}_{\underline{\mu}}=\arg \min _{\boldsymbol{\theta} \in \boldsymbol{\Theta}} \mu(\boldsymbol{\theta})  \tag{27}\\
& \boldsymbol{\theta}_{\bar{\mu}}=\arg \max _{\boldsymbol{\theta} \in \boldsymbol{\Theta}} \mu(\boldsymbol{\theta})  \tag{28}\\
& \boldsymbol{\theta}_{\underline{P_{f}}}=\arg \min _{\boldsymbol{\theta} \in \boldsymbol{\Theta}} P_{f}(\boldsymbol{\theta})
\end{align*} \boldsymbol{\theta}_{\overline{P_{f}}}=\arg \max _{\boldsymbol{\theta} \in \boldsymbol{\Theta}} P_{f}(\boldsymbol{\theta}) .
$$

Unfortunately, the uncertainty makes the inverse problem an ill-posed and difficult to solve problem. For instance, the objective of the optimization can involve the calculation of some statistics. These are generally estimated by means of samples and those statistics are not exact but approximate. Stochastic optimization methods ${ }^{44}$ are specially suited to make optimization with random objective functions.

In addition, the necessity to separate epistemic and aleatory uncertainty makes the extreme case analysis even more difficult. This is because an extreme case can derive from different combinations of epistemic/aleatory uncertainty ( $\boldsymbol{\alpha}, \boldsymbol{\theta})$.

## E. Robust Design

The final task in the design of a safety critical system is to perform a robust design optimization. The main aim of the robust design is to consider explicitly the effects of the uncertainties in the optimization problem. A solution of this problem can be obtained by performing an optimization analysis able to identify the design point with improved robustness and reliability characteristics. ${ }^{9}$

This requires to repeatedly evaluate the performance of the system that can be defined as e.g. expected values, probability of failure. The approach described in Section C can be adopted for the estimation of these quantities (inner loop) and it generally requires considerable numerical efforts. In addition, it has to be performed for each candidate solutions of the optimization procedure (the outer loop).

Generally in robust design only one bound is of interest. For instance we would like to reduce the probability of failure. In this sense, the optimal design point $\boldsymbol{d}^{\text {opt }}$, would be given for example by:

$$
\begin{equation*}
\boldsymbol{d}^{o p t}=\arg \min _{\boldsymbol{d} \in \boldsymbol{D}} \overline{P_{f}}(\boldsymbol{d})=\arg \min _{\boldsymbol{d} \in \boldsymbol{D}} \max _{\boldsymbol{\theta} \in \boldsymbol{\Theta}} \int_{\boldsymbol{\Omega}} \mathcal{I}\left[\mathcal{H}(\boldsymbol{\alpha} ; \boldsymbol{\theta} ; \boldsymbol{d})>\mathcal{H}^{c r i t}\right] \mathrm{d} C(\boldsymbol{\alpha}) \tag{29}
\end{equation*}
$$

or by

$$
\begin{equation*}
\boldsymbol{d}^{o p t}=\arg \min _{\boldsymbol{d} \in \boldsymbol{D}} \bar{\mu}(\boldsymbol{d})=\arg \min _{\boldsymbol{d} \in \boldsymbol{D}} \max _{\boldsymbol{\theta} \in \boldsymbol{\Theta}} \int_{\boldsymbol{\Omega}} \mathcal{H}(\boldsymbol{\alpha} ; \boldsymbol{\theta} ; \boldsymbol{d}) \mathrm{d} C(\boldsymbol{\alpha}) \tag{30}
\end{equation*}
$$

where $\boldsymbol{D}$ is the design space and $\mathcal{H}$ becomes a function $\boldsymbol{\Omega} \times \boldsymbol{\Theta} \times \boldsymbol{D} \rightarrow \mathbb{R}$.

Nevertheless, the estimation of bounds of the system performance for this subproblem, remains a computational challenge. Thus, due to the tremendous numerical cost involved, caused by the repeated assessment of the system response for different candidate solutions, the direct solution of this subproblem may render the computational task unfeasible, even for academic problems. Then, it is necessary to resort to specific techniques such as the use of surrogate models in order to decrease the computational costs. ${ }^{45,46}$ Surrogatemodels mimic the behaviour of the original model, by means of an analytical expression with negligible computational cost. The approximation is constructed by selecting some predefined interpolation points in the design space, at which the maximal failure probability is estimated; then, a surrogate model is adjusted
to the data collected in a least squares sense. As the construction of this approximation over the entire domain can be demanding, it may be easier to generate an approximation of the failure probabilities over a sub-domain, ${ }^{47}$ i.e. to generate a local surrogate model. Local surrogate model might require generally less evaluation points to be constructed although they have to be continuously updated in order to follow the current values of the design variables. Artificial neural networks are very versatile surrogate models; other methods such as kriging can be used as well. ${ }^{45,46}$ Surrogate models should not introduce unnecessary approximations and errors. Hence, only the most computational expensive part of the model should be replaced keeping the original models for the less demanding parts.

## IV. Numerical Implementation

The uncertainty quantification and management require the availability of flexible numerical tools able to deal with the different representations of uncertainty. Furthermore, since the non-deterministic analysis is computationally quite demanding, such numerical tools need to be very efficient and scalable. In fact, since such analyses need to be repeated a large number of times, the computational cost could be excessive even when the solver is reasonably fast (e.g. the computation of $g$ in the challenge problem requires 2 seconds on a common desktop computer). For these reasons, the proposed approach has been developed and integrated into the OpenCossan framework. ${ }^{10}$

## A. OpenCossan

OpenCossan is a collection of open source algorithms, methods and tools released under the LGPL licence, ${ }^{48}$ and under continuous development at the Institute for Risk and Uncertainty at the University of Liverpool, UK. The source code is available upon request at the web address http://www.cossan.co.uk.

OpenCossan is also the computational core of a general purpose software, namely COSSAN-X, originally developed by the research group of Prof. G.I. Schuëller at the University of Innsbruck, Austria. ${ }^{49,50}$ As a general purpose software, it means that a reasonably wide range of engineering and scientific problems can be treated by the software.

This computational core, developed in MATLAB ${ }^{\circledR}$ using an object-oriented programming paradigm, includes several predefined solution sequences to solve a number of different problems. The framework is
organized in classes, i.e. data structures consisting of data fields and methods together with their interactions and interfaces. Thanks to the modular nature of OpenCossan, it is possible to define specialized solution sequences including reliability methods, optimization strategies and surrogate modelling or parallel computing strategies to reduce the overall cost of the computation.

OpenCossan provides intuitive, clear, well documented and human readable interfaces to the classes. Furthermore, the developed numerical methods are highly scalable and parallelizable, thanks to its integration with distributed resource management, such as openlava and GridEngine. These job management tools allow to take advantages of high performance computing, as shown in the next sub-session.

## B. High performance computing

The proposed strategies for solving the challenge problem are generally very demanding in terms of computational resources. For instance, sensitivity analysis and uncertainty quantification might require more than $10^{6}$ up to $10^{9}$ model evaluations (see e.g. Section V-B)).

Even though the computational cost to evaluate the model might be low, the huge number of model evaluations required by the analyses represents a computational challenge. A possible way to reduce the execution time of the analyses is to employ surrogate models to approximate the input/output relations with faster analytical approximations. This, however, introduces loss of accuracy in the analysis, and such surrogate models have to be accurately calibrated before being employed in the analysis.

Alternatively, multiple independent instances of the solver can be executed simultaneously for different values of the input to the system, allowing for a reduction of the analyses time without any loss of accuracy.

Hence, in order to reduce the computational wall-clock time required by the analyses two types of parallelization can be used. The first type of parallelization is used to speed-up the analysis of most internal loop required by the simulations. In this case, a special job on a pool of MATLAB workers is created on each multi-core machine, connecting the MATLAB client to the parallel pool (e.g. using the command parpool). Features from the MATLAB parallel toolbox e.g., parfor, can be used to distribute the tasks on the MATLAB clients. This type of parallelism can be implemented on each single computational node. Clearly such kind of parallelization can only be used if the model is evaluated in MATLAB. In case the analysis of the inner loop requires the call of an external solver (such as a FE/CFD analysis) the multi-thread, shared
memory parallelism capabilities of the external software need to be adopted in order to enable the first level of parallelization. The second level of parallelization exploits cluster and grid computing, i.e. the availability of machines connected in an heterogeneous network. In this case, the total number of simulations is slitted in a multiple number of independent batch jobs. The jobs are then submitted to the job scheduler/manager and distributed efficiently on the available machines of the grid/cluster.

As a final consideration, these two types of parallelization can be combined together. As an example, the model evaluation required by global sensitivity analysis can be spread using batch jobs along multiple computational nodes. Then, for each batch processed on each node of the cluster, a subset of analyses is performed in parallel on the cores of the node in order to compute the quantity of interest, e.g., Monte Carlo simulation can be performed to evaluate a stochastic model and to compute the empirical CDFs of the quantity of interest.

Although, using OpenCossan framework, the parallelization of the analysis is straightforward, the parallelization of a generic model might be quite challenging. In fact, independent multiple stream and substream should be generated by the master node and distributed to the workers. In MATLAB, a combined multiple recursive generator (mrg32k3a) can be used to generate such independent sub-streams. When usersupplied code is involved, the standard approach, taken by OPENCossan to parallelize MATLAB functions with independent jobs, is to compile such functions using mcc and then distribute the compiled code to the node of the cluster (workers). Hence, it is possible to execute in parallel MATLAB code without the necessity to install MATLAB on each computational node of the cluster, but only accessing the MATLAB runtime libraries. When this approach is not possible, for instance due to license limitations to deploy code, multiple headless instances of MATLAB are executed (available MATLAB licenses on each cluster node are necessary).

## V. Numerical application

## NASA Langley multidisciplinary uncertainty quantification challenge

The necessity to determine limitation and range of applicability of existing uncertainty quantification (UQ) methodologies and to advance the state of the practice in UQ problem of direct interest of NASA has lead to
the development of a challenge problem. The reader is referred to Ref. 9 for a full description of the NASA UQ challenge problem.

A mathematical model that describes the dynamic of a remotely operated twin-jet aircraft developed by NASA Langley Research Center is analyzed (see Figure 3). The model, provided as a "Black Box", contains 21 parameters, $\boldsymbol{p}, 16$ design variables, $\boldsymbol{d}$ and 8 outputs, $\boldsymbol{g}$. Furthermore, a set of intermediate variables, $\boldsymbol{x}$, that can be interpreted as outputs of the so-called fixed discipline analysis, $\boldsymbol{x}=h(\boldsymbol{p})$, are the inputs of the cross discipline analysis $\boldsymbol{g}=f(\boldsymbol{x}, \boldsymbol{d})$. One of the main objectives of the proposed problem is to identify the design parameters, $\boldsymbol{d}$, that provide optimal worst case probabilistic performance in presence of the model parameters uncertainty, $\boldsymbol{p}$ i.e. perform a robust optimization. This requires to solve a series of subproblems, such as uncertainty characterization, sensitivity analysis, among others, in order to improve the model.

In the following, the term "original model" is used to describe the uncertainty model as provided in the challenge problem; "reduced model" refers to the model with reduced uncertainty after the solution of the subproblem A and "improved model" refers to the reduced model with four parameters with the smallest ranges of uncertainty obtained from NASA. Only the main findings are reported and the reader is referred to Ref. 30 for detailed results of the challenge problem.


Figure 3: Relationship between the variables and functions of the NASA Langley multidisciplinary uncertainty quantification challenge problem. ${ }^{8}$

Decomposition of variables $p$ into its aleatory and epistemic components Table 1 lists all variables of vector $\boldsymbol{p}$ decomposed into an aleatory component and an epistemic component. Note that on the one hand, the aleatory component of a random variable or distributional p-box can be represented as a
uniform random variable in $(0,1]$; on the other hand, the epistemic component of a distributional p -box is given by the intervals that describe the parameters of the parental CDF; in this way, the aleatory $\boldsymbol{\Omega}$ and the epistemic $\boldsymbol{\Theta}$ spaces have respectively 17 and 31 dimensions.

Representation of variable $p_{1}$ In Section II it has been shown how to represent p-boxes. However, variable $p_{1}$ requires special considerations in its representation. These are discussed in detail in the following: The input variable $p_{1}$ is represented as a unimodal beta distribution whose mean $\mu$ and variance $\sigma^{2}$ are uncertain, but are known to lie in the intervals [3/5, 4/5] and [1/50, 1/25] respectively. Instead beta distributions are characterized by shape parameters $a$ and $b$ which are related to $\mu$ and $\sigma^{2}$ by:

$$
\begin{equation*}
\mu=\frac{a}{a+b} \quad \sigma^{2}=\frac{a b}{(a+b+1)(a+b)^{2}} \tag{31}
\end{equation*}
$$

that is,

$$
\begin{equation*}
a=-\frac{\mu\left(\sigma^{2}+\mu^{2}-\mu\right)}{\sigma^{2}} \quad b=\frac{(\mu-1)\left(\sigma^{2}+\mu^{2}-\mu\right)}{\sigma^{2}} \tag{32}
\end{equation*}
$$

The required unimodality implies that $a$ and $b$ are greater than 1 . For shape parameters lower than 1 the beta distribution assume the U-shaped bimodal distributions.

Representation of variables $p_{4}$ and $p_{5}$ One drawback of the proposed approach is that the copula must be perfectly modelled, without any epistemic uncertainty in its parameters. The copula that relates variables $p_{4}$ and $p_{5}$ has an interval parameter, namely $I_{8}$, which models the correlation $\rho\left(p_{4}, p_{5}\right)$. Variables $p_{4}$ and $p_{5}$ are modelled using the following formulation, which permits to split uncertainty into the aleatory and the epistemic spaces while representing the dependence with an independent copula, which does not have any epistemic component at all:

- The aleatory part of the joint probability box is given by $\alpha_{3}$ and $\alpha_{4}$ which are independent and uniform random variables on $(0,1]$. Note that $z_{3}=\Phi^{-1}\left(\alpha_{3}\right)$ and $z_{4}=\Phi^{-1}\left(\alpha_{4}\right)$ where $\Phi$ represents the standard normal CDF.
- The epistemic part of the joint distribution is given by the 5 -dimensional box $X_{i=4}^{8} I_{i}$.

A simulation from variables $p_{4}$ and $p_{5}$ can be performed by using the vector $\boldsymbol{z}=\left[z_{3}, z_{4}\right]^{T}$ and a parameter vector $\boldsymbol{\theta} \in X_{i=4}^{8} I_{i}$; the simulation uses the standard procedure for sampling from a multivariate normal PDF. This method employs the Cholesky decomposition of the covariance matrix.

Table 1: Aleatory and epistemic components of the input variables $p_{i}$, The first column provides the parameter's symbol, the second one its category (see above for a description of the categories), the third and fourth one describe its aleatory and epistemic uncertainty model. Here $\rho(),, \mathrm{E}[\cdot]$ and $\operatorname{Var}[\cdot]$, denote the correlation, expected value, and variance operators respectively.

| Variable | Category | Aleatory <br> component | Epistemic component | Description |
| :---: | :---: | :---: | :---: | :---: |
| $p_{1}$ | III | $\alpha_{1} \sim \operatorname{Unif}(0,1]$ | $I_{1}=[3 / 5,4 / 5]$ | Interval of $\mathrm{E}\left[p_{1}\right]$ |
|  |  | (distribution type: | $I_{2}=[1 / 50,1 / 25]$ | Interval of $\operatorname{Var}\left[p_{1}\right]$ |
|  |  | unimodal Beta) |  |  |
| $p_{2}$ | II |  | $I_{3}=[0,1]$ | Interval |
| $p_{3}$ | I | $\alpha_{2} \sim \operatorname{Unif}(0,1]$ |  | Random variable |
| $p_{4}, p_{5}$ | III | $\alpha_{3} \sim \operatorname{Unif}(0,1]$ | $I_{4}=[-5,5]$ | Interval of $\mathrm{E}\left[p_{4}\right]$ |
|  |  | $\alpha_{4} \sim \operatorname{Unif}(0,1]$ | $I_{5}=[1 / 400,4]$ | Interval of $\operatorname{Var}\left[p_{4}\right]$ |
|  |  | (distribution type: | $I_{6}=[-5,5]$ | Interval of $\mathrm{E}\left[p_{5}\right]$ |
|  |  | multivariate gaussian) | $I_{7}=[1 / 400,4]$ | Interval of $\operatorname{Var}\left[p_{4}\right]$ |
|  |  |  | $I_{8}=[-1,1]$ | Interval of $\rho\left(p_{4}, p_{5}\right)$ |
| $p_{6}$ | II |  | $I_{9}=[0,1]$ | Interval |
| $p_{7}$ | III | $\alpha_{5} \sim \operatorname{Unif}(0,1]$ | $I_{10}=[0.982,3.537]$ | Interval of $a$ |
|  |  | (distribution type: Beta) | $I_{11}=[0.619,1.080]$ | Interval of $b$ |
| $p_{8}$ | III | $\alpha_{6} \sim \operatorname{Unif}(0,1]$ | $I_{12}=[7.450,14.093]$ | Interval of $a$ |
|  |  | (distribution type: Beta) |  | Interval of $b$ |
| $p_{9}$ | I | $\alpha_{7} \sim \operatorname{Unif}(0,1]$ |  | Random variable |
| $p_{10}$ | III | $\alpha_{8} \sim \operatorname{Unif}(0,1]$ | $I_{14}=[1.520,4.513]$ | Interval of $a$ |
|  |  | (distribution type: Beta) | $I_{15}=[1.536,4.750]$ | Interval of $b$ |
| $p_{11}$ | I | $\alpha_{9} \sim \operatorname{Unif}(0,1]$ |  | Random variable |
| $p_{12}$ | II | $34 \text { of } 73$ | $I_{16}=[0,1]$ | Interval |
| $p_{13}$ | III | Qmoericdunifstatulu of Aeronaut |  | Interval of $a$ |

Consequently, the joint distribution-free probability box formed by variables $p_{4}$ and $p_{5}$ can be represented as the random set $\Gamma:(0,1]^{2} \rightarrow \mathscr{F}, \boldsymbol{\alpha} \mapsto \Gamma(\boldsymbol{\alpha})$ where $\boldsymbol{\alpha}=\left(\alpha_{3}, \alpha_{4}\right), \mathscr{F}$ is the system of focal elements given by the preimages of $\left\{\alpha_{3} \times \alpha_{4} \times I_{4} \times I_{5} \times \cdots \times I_{8}:\left(\alpha_{3}, \alpha_{4}\right) \in(0,1]^{2}\right\}$ through $F_{p_{4} p_{5}}$. Since $\alpha_{3}$ and $\alpha_{4}$ are independent uniform random variables in $(0,1]$, they can be considered as the realization of a bidimensional product copula, defined on $(0,1]^{2}$. For the interpretation of $\alpha_{3}, \alpha_{4}, I_{4}, \ldots, I_{8}$ the reader is referred to Table 1.

## A. Subproblem A

The aim of the uncertainty characterization or subproblem A is to reduce the epistemic uncertainty components of the category II ( $p_{2}$ ) and III parameters ( $p_{1}, p_{4}, p_{5}$ ) that are inputs of a subsystem $h_{1}$. The subsystem provides a scalar output $x_{1}$ as a function of those five uncertain parameters, that is,

$$
\begin{equation*}
x_{1}=h_{1}\left(p_{1}, p_{2}, p_{3}, p_{4}, p_{5}\right) \tag{33}
\end{equation*}
$$

In this subproblem, the vector $\left[p_{1}, \ldots p_{5}\right]$ is the output of the system $\mathcal{W}$, the system $h_{1}$ is equivalent to the function $\mathcal{G}$ defined on Section II-F, the epistemic space is the Cartesian product $\boldsymbol{\Theta}:=X_{i=j}^{8} I_{j}$ and the aleatory space, which models variables $\alpha_{1}$ to $\alpha_{4}$, is defined by $\boldsymbol{\Omega}:=(0,1]^{4}$ (see Table 1 ).

Two sets of 25 observations of the "true uncertainty model" $\boldsymbol{\theta}^{*} \in \boldsymbol{\Theta}$ are available to reduce the uncertainty in $\boldsymbol{\Omega}$. The approaches described in Section III-A are here adopted.

One of the main challenges of this subproblem is provided by the limited available information ( 25 observation points for each dataset) and the relatively large dissimilarity of the empirical CDFs associated with those datasets as shown in Figure 4.

## 1. Non-parametric statistic method based on the Kolmogorov-Smirnov test

The procedure presented in Session III-A has been used to solve the subproblem A. First, the validation similarity level has been calculated after using a Gaussian KDE to compute the CDF $\tilde{F}_{e}$ for the observation sets. A validation similarity level $D_{\tilde{v}}=0.18$ has been obtained calculating the maximum distance between the two KDEs adjusted to the two datasets respectively $\tilde{F}_{e}$ (i.e. using Eq. (6)). The measure of similarity


Figure 4: Empirical CDF, $\hat{F}$, of the two set of observation points and CDF obtained adopting the Gaussian kernel density of Eq. (7), $\tilde{F}$. The dots and squares show the two datasets $\mathcal{D}_{e}$, respectively.
obtained comparing the two empirical CDFs, $\hat{F}_{e}$, of the datasets is $D_{\hat{v}}=0.24$ as shown in Figure 5. This allows to identify those points $\boldsymbol{\theta}_{i} \in \boldsymbol{\Theta}$ that conform with the observations such that $D_{i}<D_{v}$.

Assuming a uniform distribution on $\boldsymbol{\Theta}, 10000$ samples $\boldsymbol{\theta}_{i}$ are drawn and for each $\boldsymbol{\theta}_{i}, n=5000$ samples from the aleatory space $\boldsymbol{\Omega}$ are used to propagate the aleatory uncertainty through the model (using the function p_to_x1). Finally, using the empirical CDF of $x_{1}\left(\hat{F}\left(x_{1} \mid \boldsymbol{\theta}_{i}\right)\right)$, the measure of similarity $D_{i}$ is calculated against $\tilde{F}_{e}$ according to Eq. (6) (i.e. $\left.D_{i}=\sup _{x}\left|\hat{F}\left(x \mid \boldsymbol{\theta}_{i}\right)-\tilde{F}_{e}(x)\right|\right)$. Please note that due to the large number of samples used $\hat{F}_{i}\left(x_{1} \mid \boldsymbol{\theta}_{i}\right) \approx \tilde{F}_{i}\left(x_{1} \mid \boldsymbol{\theta}_{i}\right)$. The histograms of the measure of similarity $D_{i}$ are shown in Figure 5 computed for the dataset of 25 and 50 observations, respectively. It is possible to observe that $D_{i}$ is smaller when the $\operatorname{KDE} \tilde{F}_{i}\left(x_{1} \mid \boldsymbol{\theta}_{i}\right)$ and all 50 observations are used.

The measure of similarity $D_{\tilde{v}}=0.18$ identifies model outputs, $x_{1}$ obtained from the realizations in the epistemic space, $\boldsymbol{\theta}_{i}$, that are in agreement with the observations (represented in Figure 5 by the bars on the left of $\left.D_{\tilde{v}}\right)$. Calculating $P\left(D_{i}>D_{\tilde{v}}\right)=c$, two confidence levels have been obtained: $c_{\tilde{v}(25)}=0.8031$ and $c_{\tilde{v}(50)}=0.547$ when $D_{i}$ is calculated against the $\tilde{F}_{e}$ obtained using 25 and 50 observations, respectively.

Figure 6 shows the parallel coordinate plot of the epistemic realizations. Please note that for readability purposes, only 1000 realizations are shown. In a parallel plot a multi-dimensional quantity is shown graphically and represented as a polyline with vertices on the parallel axes. The vertex on the $m$-axis corresponds to the $i$-th realization of the $m$-coordinate (i.e. $\theta_{m}^{(i)}$ ). The axes of the plot have been normalized, between 0 and 1. The top panel of Figure 6 shows combination of epistemic realizations for different level of similarity measure computed against $\tilde{F}_{e}$ constructed from 25 observations. The Figure shows all the combinations of all epistemic realizations $(c=0)$, those with a similarity measure $D_{i}<D_{\hat{v}}$ (i.e. $\left.c=0.547\right)$ and $D_{i}<D_{\tilde{v}}$ (i.e. $c=8031$ ), respectively. The top panel of Figure 6 shows the parallel plot with measures of similarity calculated using all the 50 observations. $c=0.0547$ correspond to a similarity measure $D_{i}<D_{\tilde{v}}$ while $c=0.0547$ correspond an arbitrary level $D_{i}<0.1$.

The parallel coordinate plot allows to identify the epistemic uncertainty that can be reduced. For instance, all the realizations of $\mathrm{E}\left[p_{5}\right]$ with similarity level lower $D_{v}$ are in the normalized interval $[0,0.6]$ while $\mathrm{E}\left[p_{1}\right]$ is in the normalized interval $[0,0.7]$. On the contrary, the intervals of $\operatorname{Var}\left[p_{1}\right], p_{2}, \mathrm{E}\left[p_{4}\right], \operatorname{Var}\left[p_{4}\right], \operatorname{Var}\left[p_{5}\right]$ and $\rho\left(p_{4}, p_{5}\right)$ cannot be improved based on the current available data. Although the resulting model for $\theta$ obtained are collection of points, the identified realizations cover connected ranges (remember that only

1000 over 10000 realizations are shown in Figure 6). The results are summarized in Table 2.

## 2. Bayesian updating on the epistemic space

The Bayesian inference is the second approach used to reduce the epistemic uncertainty as explained in Section III-A-2. In this method, Transitional Monte Carlo Markov Chains have been used to sample 1000 realizations from the posterior $\operatorname{PDF} p\left(\boldsymbol{\theta} \mid \mathcal{D}_{n}\right)$. Two strategies have been employed to estimate the likelihood $P\left(\mathcal{D}_{e} \mid \boldsymbol{\theta}_{i}\right)$ : the standard Bayesian and an approximate Bayesian computational method.

Bayesian computational method (BC) In this case, the likelihood is computed using Eq. (8) and $p\left(x \mid \boldsymbol{\theta}_{i}\right)$ is estimated by means of a KDE, computed with $n=1000$ points from the aleatory space. Figures 7 shows the posterior distributions sampled using TMCMC with 25 and 50 observation points as evidence, respectively. Histograms of the posterior samples are normalized, assigning a value of 1 to the number of counts in the bin containing the majority of samples. After normalizing the histograms, it is possible to set a general limit of normalized counts used to exclude outliers of the TMCMC algorithm and indicated by the horizontal red lines in Figures 7.

Approximate Bayesian computational method (ABC) In this case, 200 samples are used to evaluate $\hat{F}\left(\cdot \mid \boldsymbol{\theta}_{i}\right)$ and the quantities $\delta_{k}$ by means of Eq. (12). Thereafter, the likelihood (11) is computed. Figures 8 shows the normalized posterior distributions sampled using TMCMC with 25 and 50 observation points as evidence, respectively.

The results The proposed method has been able to identify a reduced epistemic space associated to $\mathrm{E}\left[p_{1}\right]$ and $\mathrm{E}\left[p_{5}\right]$ but no conclusions can be drawn for the other input parameters. The updated ranges of the epistemic uncertainties are summarized in Table 2.

The Bayesian updating procedure successfully managed to reduce the uncertainty associated to the output $x_{1}$ as shown for example in Figure 9, for the approximate Bayesian computational method. Figure 9 shows different p-boxes of $x_{1}$ obtained with the updated epistemic uncertainty parameters, using the first set of 25 observations and the full set of 50 observations, respectively. The approximated p-boxes have been obtained using the following procedure. First, 10000 samples $\boldsymbol{\theta}_{i}$ of the epistemic variable are drawn from uniform distributions defined by the full range of the updated bounds (light gray) and by the updated


Figure 5: Histogram of the measure of similarity, $D_{i}$, between the CDF calculated sampling randomly in the epistemic space and the observations, for 25 (top panel) and 50 (bottom panel) observations $\left(\mathcal{D}_{e}\right) . D_{i}$ has been computed using the empirical CDF of the experimental data (blue bars) and the CDF obtained using Gaussian kernel smoother functions (yellow bars). The figure also shows the values of the measure of similarity between the two set of observation data computed using Gaussian kernel smoother techniques, $D_{\tilde{v}}$, and empirical CDF, $D_{\hat{v}}$, respectively.


Figure 6: Parallel coordinates plot of the 8 category II and III parameters of the input factors of $h_{1}$ (i.e. $\left.p_{i}, i=1, \cdots, 5\right)$ for 25 (top panel) and 50 (bottom panel) observations $\left(\mathcal{D}_{e}\right)$. The figure shows only 1000 realizations (over a total sample of 10000) of the epistemic space for different significant levels $c$ of the Kolmogorov-Smirnov test. $c=0$ represents of all the realizations. $c_{\tilde{v}}$ represents realizations of $\theta$ with a measure of similarity $D_{i}<D_{\tilde{v}}$. $c_{\hat{v}}$ represents realizations of $\theta$ with a measure of similarity $D_{i}<D_{\hat{v}}$ and $c_{D=0.1}$ realizations with a measure of similarity $D_{i}<0.1$.


Figure 7: Normalized histogram of $p\left(\boldsymbol{\theta} \mid \mathcal{D}_{e}\right)$ obtained using Bayesian Computational method with (a) 25 experimental observations and (b) 50 experimental observations (b) of $x_{1}$, respectively. The normalization assigns a value of 1 to the bin with the highest number of counts. The red line represent the cut-off value to determine the updated range.


Figure 8: Normalized histogram of $p\left(\boldsymbol{\theta} \mid \mathcal{D}_{e}\right)$ obtained using Approximate Bayesian Computational method with (a) 25 experimental observations and (b) 50 experimental observations (b) of $x_{1}$, respectively. The normalization assigns a value of 1 to the bin with the highest number of counts. The red line represent the cut-off value to determine the updated range.


Figure 9: P-boxes of $x_{1}$ and the empirical CDFs of the experimental data. The p-boxes have been obtained using the full range of the posterior parameters and using the range that excludes the outliers, respectively.
bounds obtained excluding the outliers (dark gray). Then, the $\operatorname{CDF} \hat{F}\left(\cdot \mid \boldsymbol{\theta}_{i}\right)$ is computed for each epistemic realization. Finally the curves enveloping all the CDFs are obtained and shown in Figure 9. It is possible to notice that the updated p-box of $x_{1}$ is tighter when all the 50 experimental observations are used. Additionally, the experimental CDFs of the calibration data set are fully contained in the light gray area (i.e. the p-boxes obtained excluding the outliers). However, the validation data lay inside the updated p-box only when the full intervals of updated parameters are considered.

The reduced uncertainty model identified by the non-parametric approach and by the Bayesian inference approach are summarized in Table 2, respectively. Although only the uncertainty of two parameters can be significantly reduced, the results provided by the proposed approaches are in agreement providing a cross validation of the developed procedures used to solve the subproblem A.

## B. Subproblem B

The aim of this subproblem is to identify and rank the input parameters of category II and III (i.e. intervals and distributional p-boxes) according to degree of refinement in the output p-boxes which one could hope to obtain by refining their uncertainty models. More specifically, in problem B1 the focus is to rank the 4 input factors that affect the variability the output $x_{i}$ of each model $h_{i}(\cdot), i=1, \ldots, 4$, respectively. In tasks B2-B3, 17 parameters need to be ranked according to the reduction in the range of $J_{1}=E\left[w\left(\boldsymbol{p}, \boldsymbol{d}_{\text {baseline }}\right)\right]$ (task B2) and $J_{2}=1-P\left[w\left(\boldsymbol{p}, \boldsymbol{d}_{\text {baseline }}\right)<0\right]$ (task B3), respectively. In those expressions, the worst-case requirement metric $w$ is defined by $w(\boldsymbol{p}, \boldsymbol{d})=\max _{1 \leq i \leq 8} g_{i}(\boldsymbol{p}, \boldsymbol{d})$. The strategy presented in Section III-B will be used.

## 1. Problem B1

Nonspecificity technique By means of the nonspecificity measure, each interval $\left[\underline{I_{i}}, \overline{I_{i}}\right]$ is reduced to the value given by $\underline{I_{i}}+p_{r} \cdot\left(\overline{I_{i}}-\underline{I_{i}}\right)$, where $p_{r} \in\{0.1,0.3,0.5,0.7,0.9\}$. For instance, interval $I_{1}=\left[\underline{\mathrm{E}\left[p_{1}\right]}, \overline{\mathrm{E}\left[p_{1}\right]}\right]=[3 / 5,4 / 5]$ is reduced to the constants $0.62,0.66,0.70,0.74$ and 0.78 and $n=50$ samples from the product copula that links aleatory variables, $\left(\alpha_{1}, \alpha_{2}, \alpha_{3}, \alpha_{4}\right)$, are employed to construct the output Dempster-Shafer structure for each reduction. Note that $\alpha_{3}$ and $\alpha_{4}$ are used to model the variables $p_{4}$ and $p_{5}$, according to the transformation explained at the beginning of Section $V$. Then, the

Table 2: Reduced uncertainty model using the non-parametric approach ( $c=0.547$ ) or 25 observations and $c=8031$ for 50 observations) and the Bayesian inference, respectively. $\mathrm{A}-$ means that the method could not reduce the epistemic uncertainty for the referred variable.

| Variable | Original <br> interval | Nonparametric <br> method | Bayesian methods |  |
| :---: | :---: | :---: | :---: | :---: |
|  |  |  | BC | ABC |
|  | 25 observations |  |  |  |
| $\mathrm{E}\left[p_{1}\right]$ | [ 0.6000, 0.80] | [ 0.6000, 0.72] | [ $0.6000,0.73]$ | [0.6030, 0.755 ] |
| $\operatorname{Var}\left[p_{1}\right]$ | [ 0.0200, 0.04] | - | - | - |
| $p_{2}$ | [ 0.0000, 1.00] | - | - | - |
| $\mathrm{E}\left[p_{4}\right]$ | [-5.0000, 5.00] | - | - | - |
| $\operatorname{Var}\left[p_{4}\right]$ | [ 0.0025, 4.00] | - | - | - |
| $\mathrm{E}\left[p_{5}\right]$ | [-5.0000, 5.00] | [-5.0000, 0.78] | - | [-5.0000, 4.50] |
| $\operatorname{Var}\left[p_{5}\right]$ | [ 0.0025, 4.00] | - | - | - |
| $\rho\left(p_{4}, p_{5}\right)$ | [-1.0000, 1.00] | - | - | - |
|  | 50 observations |  |  |  |
| $\mathrm{E}\left[p_{1}\right]$ | [ 0.6000, 0.80] | [0.63, 0.76] | [0.60, 0.75] | [0.618, 0.791] |
| $\operatorname{Var}\left[p_{1}\right]$ | [ 0.0200, 0.04] | [0.0260, 0.04] | - | - |
| $p_{2}$ | [ 0.0000, 1.00] | - | - | - |
| $\mathrm{E}\left[p_{4}\right]$ | [-5.0000, 5.00] | [-4.50, 4.80] | - | - |
| $\operatorname{Var}\left[p_{4}\right]$ | [ 0.0025, 4.00] | - | - | [0.097, 3.943] |
| $\mathrm{E}\left[p_{5}\right]$ | [-5.0000, 5.00] | [-4.90, 0.30] | - | [-5.00, 4.45 ] |
| $\operatorname{Var}\left[p_{5}\right]$ | [ 0.0025, 4.00] | - | - | - |
| $\rho\left(p_{4}, p_{5}\right)$ | [-1.0000, 1.00] | - | - | - |

nonspecificity which is a measure of epistemic uncertainty, of each of those Dempster-Shafer structures is calculated. Following a similar procedure, the rankings of input variables have been calculated (see Table 3) according to the output nonspecificity for the systems $h_{2}, h_{3}$ and $h_{4}$, respectively. In all cases, the evaluation of equation (5) was performed for each focal element using a genetic algorithm with a population of 30000 individuals and 10 generations.

Global sensitivity analysis technique The global sensitivity analysis has been performed on a redefined mathematical model $\boldsymbol{h}^{*}$ of the original $\boldsymbol{h}$ as detailed in Section III-B-2. $\boldsymbol{h}^{*}$ takes as inputs only uniform distributions (that represents the epistemic space $\mathbf{\Theta}$ ) and returns a scalar output $y_{i}$ (the area of distribution-free p-boxes) as shown in Figure 10.

For each combination $\boldsymbol{\theta}_{i} \in \boldsymbol{\Theta}$ of the input parameters, the model $h^{*}$ performs an internal Monte Carlo simulation using $n=500$ samples $\boldsymbol{\alpha}_{j}$ to calculate an empirical CDF of $x_{j}^{i}, \hat{F}\left(\cdot \mid \boldsymbol{\theta}_{i}\right)$. A sample size of 500 is sufficient to rank unequivocally the most important parameters with respect to the outputs $x_{k}, k=1, \ldots, 5$, as shown in Figure 11. Then, $\hat{F}\left(\cdot \mid \boldsymbol{\theta}_{i}\right)$ is compared with a "reference CDF", $F(\cdot \mid \overline{\boldsymbol{\theta}})$ and the final output $y_{i}$ is returned (see Eq. (17)).


Figure 10: Redefined model $h_{1}^{*}$ used for performing the Global sensitivity analysis with aleatory and epistemic uncertainty.

In order to reduce the computational noise of $h^{*}$ (i.e. the variance of the output), the common random number technique ${ }^{51}$ has been used to propagate the aleatory uncertainty (i.e. performing the internal Monte Carlo simulation for the model $h^{*}$ ). The extended-FAST method has been used with 2048 samples of $\boldsymbol{\theta}_{i}$ for each of the 8 input factors of the refined model, and in consequence, 16384 simulations are required for each
measure of $x_{k}$ while the Saltelli's method has been run with 16384 samples for a total cost of 540672 model evaluations for each $x_{k}$.

First order Sobol' indices for $x_{1}$


Figure 11: Effect of different samples size for the internal Monte Carlo simulation on the estimation of the Sobol' indices with respect to $x_{1}$ by means of extended-FAST method method.

Since the global sensitivity procedure computes the sensitivity measure of the individual components for the category III parameters (e.g. $\mathrm{E}\left[p_{1}\right], \operatorname{Var}\left[p_{1}\right]$, the numerical values for the input parameters have been calculated as: $S\left(p_{1}\right)=S\left(\mathrm{E}\left[p_{1}\right]\right)+S\left(\operatorname{Var}\left[p_{1}\right]\right), S\left(p_{4}\right)=S\left(\mathrm{E}\left[p_{4}\right]\right)+S\left(\operatorname{Var}\left[p_{4}\right]\right)+S\left(\rho\left(p_{4}, p_{4}\right)\right)$, and $S\left(p_{5}\right)=$ $S\left(\mathrm{E}\left[p_{5}\right]\right)+S\left(\operatorname{Var}\left[p_{5}\right]\right)+S\left(\rho\left(p_{4}, p_{5}\right)\right)$.

From the results summarized in Table 3, it is possible to see that the results obtained applying the two approaches are in agreement.

## 2. Problems B2 and B3

Similar strategies applied in the solution of of task B1 have been here applied. In this case, for the nonspecificity technique, the variables are mapped through the system $w\left(\boldsymbol{p}, \boldsymbol{d}_{\text {baseline }}\right)$ with only 10 focal elements and the range of the interval was measured instead of the nonspecificity of each focal element.

The redefined model $h^{*}$ has also been adopted for performing global sensitivity analyses. Here, $h^{*}$ takes as input uniform distributions representing the epistemic uncertainties and returns the output $J_{1}$ and $J_{2}$. For each realization of the epistemic uncertainty, a Monte Carlo simulation with 500 samples is performed

Table 3: Ranking of the 4 category II-III parameters according to the nonspecificity technique (NST) and global sensitivity analysis (GSA) for $x_{i}, i=1, \ldots, 4$, respectively. Note that using the global sensitivity analysis, the larger the value of the "first Sobol' index" is, the more important the input factor is. On the other hand for nonspecificity technique parameters with the lower values are more important than parameter with larger values.

| Output | Rank\#1 | Rank\#2 | Rank\#3 | Rank\#4 | Strategy |
| :---: | :---: | :---: | :---: | :---: | :---: |
| $x_{1}$ | $p_{1}(\mu)(0.235)$ | $p_{5}(\mu)(0.757)$ | $p_{4}(\mu)(0.808)$ | $p_{2}(0.850)$ | NST |
|  | $p_{1}(0.684)$ | $p_{5}(0.145)$ | $p_{4}(0.056)$ | $p_{2}(0.02)$ | GSA |
| $x_{2}$ | $p_{6}(0.063)$ | $p_{7}(a)(0.596)$ | $p_{8}(a)(0.922)$ | $p_{10}(b)(0.993)$ | NST |
|  | $p_{6}(0.701)$ | $p_{7}(0.153)$ | $p_{8}(0.021)$ | $p_{10}(<0.001)$ | GSA |
| $x_{3}$ | $p_{12}(0.026)$ | - |  |  | NST |
|  | $p_{12}(0.835)$ | $p_{15}(0.017)$ | $p_{14}(<0.001)$ | $p_{13}(<0.001)$ | GSA |
| $x_{4}$ | $p_{16}(0.121)$ | $p_{17}(a)(0.779)$ | $p_{18}(a)(0.786)$ | $p_{20}(a)(0.938)$ | NST |
|  | $p_{16}(0.761)$ | $p_{18}(0.073)$ | $p_{17}(0.025)$ | $p_{20}(0.001)$ | GSA |

to propagate the aleatory uncertainty Although the distribution of $J_{1}$ is very sensitive to the number of aleatory samples $\boldsymbol{\alpha}_{j}$, a sample size of 500 has been demonstrated to be sufficient for ranking unequivocally the most important parameters as shown in Figure 12..

The values of $\boldsymbol{g}$ are computed via $\boldsymbol{f}$ function and the CDF of $w$ computed. Finally, $J_{1}$ and $J_{2}$ are calculated from $F(w)$. Finally, the sensitivity indices of $J_{1}$ and $J_{2}$ are calculated. The extended-FAST method and the Saltelli's method has been used to estimate the sensitivity measure. The extended-FAST method has been computed using 1000 samples $\boldsymbol{\theta}_{i}$ for each input factor of the model $h^{*}$ (i.e. the intervals of the epistemic space) for a total cost of simulations 31000 whereas the Saltelli's method has been performed with 8192 samples for a total cost of 270336 model evaluations. Figures $13-14$ show the sensitivity measures of the input factors $\boldsymbol{p}$ with respect to $J_{1}$ and $J_{2}$, respectively. The most important factor that contributes to the variance of $J_{1}$ is $p_{21}$ and in particular its variance. The total indices for $p_{4}$ and $p_{5}$ show that their


Figure 12: Effect of different samples size for the internal Monte Carlo simulation on the estimation of the Sobol' indices with respect to $J_{1}$ by means of extended-FAST method method.
interaction also contributes to the variance of $J_{1}$ but it is not possible to discriminate the single contribution of the parameters $p_{4}$ and $p_{5}$. All the other components provide similar (small) contributions to the variance of $J_{1}$. Regarding the variance of $J_{2}$, the first order and total indices indicate that the parameters $p_{12}$ and $p_{1}$ are the most important parameters. The first order index indicates also a contribution from $p_{4}$. All the other components provide similar (small) contribution to the variance of $J_{2}$.

The results of the sensitivity analysis are summarized in Table 4. The most important variables in the reduction of uncertainty on $J_{1}$ are $p_{21}$ and the one that reduces the uncertainty on $J_{2}$ is $p_{12}$. Again, the different approaches have provided consistent results.

Four variables have been selected $\left(p_{1}, p_{4}, p_{12}\right.$ and $\left.p_{21}\right)$. An improved uncertainty model for these variables has been obtained from NASA. $p_{21}$ is the most important parameter for $J_{1}, p_{12}$ and $p_{1}$ are the most important parameters for $J_{2}$. The sensitivity analyses indicated that the parameters $p_{4}$ and $p_{5}$ are also important however without been able to discriminate between the two parameters. Since the parameter $p_{5}$ has been already reduced during the Uncertainty Characterisation analysis (see Section A), it has been decided to ask for an improvement of the parameter $p_{4}$, in case $p_{4}$ and $p_{5}$ where strongly correlated. The improved uncertainty intervals cannot be disclosed, as requested by the challengers.


Figure 13: First and total sensitivity measure of the $\boldsymbol{p}$ parameters respect to $J_{1}$.

Table 4: Ranking of the category II-III parameters for $J_{1}$ and $J_{2}$ computed by means of the nonspecificity technique and global sensitivity analysis (GSA).

| Output | Rank \#1 | Rank \#2 | Rank \#3 | Rank \#4 | Strategy |
| :---: | :---: | :---: | :---: | :---: | :---: |
| $J_{1}$ | $p_{21}(b)(0.726)$ | $p_{6}(0.751)$ | $p_{1}(\mu)(0.763)$ | $p_{7}(a)(1.007)$ | NST |
| $J_{1}$ | $p_{21}(0.089)$ | $p_{5}(0.036)$ | $p_{4}(0.031)$ | - | GSA |
| $J_{2}$ | $p_{12}(0.189)$ | $p_{1}(\mu)(0.571)$ | $p_{5}(\mu)(0.945)$ | - | NST |
| $J_{2}$ | $p_{12}(0.666)$ | $p_{1}(0.393)$ | $p_{4}(0.201)$ | $p_{5}(0.179)$ | GSA |



Figure 14: First and total sensitivity measure of the $\boldsymbol{p}$ parameters respect to $J_{2}$.

## C. Subproblem C

For this subproblem, we were asked to find the range of the metrics $J_{1}=E\left[w\left(\boldsymbol{p}, \boldsymbol{d}_{\text {baseline }}\right)\right]$ and $J_{2}=$ $1-P\left[w\left(\boldsymbol{p}, \boldsymbol{d}_{\text {baseline }}\right)<0\right]$, both with the reduced and with the improved uncertainty models. The metric $J_{1}$ is the expected value of the worst-case requirement metric $w$, while the metric $J_{2}$ represents the failure probability of the system. For solving this problem two different strategies, introduced in in Section III-C, have been employed.

Optimization in the epistemic space (Standard approach) A global optimization is performed in the epistemic space $\boldsymbol{\Theta} \equiv X_{i=1}^{31} I_{i}$, in order to find those points in $\boldsymbol{\Theta}$ that produce the upper and lower bounds on $J_{1}$ and $J_{2}$. For any candidate solution provided by the optimization algorithm, i.e. $\boldsymbol{\theta}_{i} \in \boldsymbol{\Theta}$, a set of $n=1000$ random points $\left\{\boldsymbol{\alpha}_{j}, j=1,2, \ldots, n\right\}$ is drawn from the aleatory space $\boldsymbol{\Omega} \equiv(0,1]^{17}$ to estimate the metrics. The number of samples from the aleatory space has been selected after performing a convergence test. More specifically, in this test, both $J_{1}$ and $J_{2}$ are estimated with increasing values of $n$ (i.e. 100,500 , 1000,5000 and 10000) for 5 representative realizations of the epistemic space, as shown in Figure 15. From the figure, it can be seen that $n=1000$ points are sufficient for estimating $J_{1}$ and $J_{2}$, with a C.o.V. of 0.1 and 0.05 respectively. The confidence of these estimates can be improved by using a larger sample size at the expense of increasing even more computational cost of the analysis.

The search for lower and upper bounds is performed by means of Monte Carlo optimization using Latin Hypercube sampling, with approximately 50000 samples. A total of $5 \times 10^{7}$ evaluations of the function x_to_g (model $\boldsymbol{f}$ ) are thus, required to complete the analysis. Here, Monte Carlo is a convenient method to solve the optimization, as the objective functions $J_{1}$ and $J_{2}$ can be quite noisy, varying approximately between $\mp 10 \%$ of the true value. In order to reduce the effect of the estimation error introduced by using finite sample sets, the objective functions maximum and minimum of $J_{i=1,2}$, are redefined as lower $J_{i}\left(1-t_{\alpha / 2}\right.$ C. o. V.) and upper $J_{i}\left(1+t_{\alpha / 2}\right.$ C.o. V.) estimations, respectively, where $\alpha=0.14$ and $t_{\alpha / 2}=1.48$ is the 86 th t-Student percentile (see also ${ }^{52}$ ).

Note that, in order to run the analysis within a reasonable time, parallelization lies at the foundations of this approach. On a common dual-core personal computer, a single estimation of $J_{i}$ takes approximately 3.4 minutes, thus a total of $\sim 120$ days for a complete analysis. By means of a double parallelization, as
described in Section IV, it has been possible to reduce the running time by two orders of magnitude, making it possible to complete the analysis in just $\sim 80$ hours.

Propagation of focal sets (counter approach) Using the propagation of focal sets method, $n=$ 1000 random vectors $\left\{\boldsymbol{\alpha}_{j}, j=1,2, \ldots, n\right\}$ are drawn from the aleatory space $\boldsymbol{\Omega} \equiv(0,1]^{17}$. Thereafter, the procedure described in Section III-C-2 was applied. In order to evaluate equations (3) and (4), genetic algorithms with a population of 125 individuals and 50 generations are adopted requiring a total computational cost of $5 \times 10^{6}$ evaluations of $w$. Figure 16 shows the convergence of the genetic algorithms for two representative focal elements. The convergence is achieved using 30 generations for the identification of the minimum/maximum of the Eq. (4).

For this approach, parallelization is also essential. In fact, approximately $5 \times 10^{6}$ evaluations of the function $x_{\_}$to_g are required to complete a full analysis. Although, in this case, the use of GA makes the parallelization a little more articulated (jobs need to be sent at any iteration of the algorithm), it is still possible to significantly reduce the running time up to two orders of magnitude (as in the standard approach). It is worth noting that the overall number of function evaluations makes this approach about 10 times more efficient than the standard approach.

RESULTS The results of the reduced uncertainty model and the improved model are summarized in Table 5. Using the proposed methods, it has been possible to bound the actual solution for the targeted metrics. As expected, the improved uncertainty model is far more informative than the reduced model, which is shown by a sensible reduction in the upper bound of $J_{1}$. An even more significant difference is documented for the range of $J_{2}$ (see Table 5), where the model of uncertainty from being totally uninformative, $J_{2} \in[0,1]$, is reduced to $J_{2} \in[0.20,0.41]$. Note also that the optimization in the epistemic space (standard approach) provided tighter bounds than the propagation of focal sets (counter approach). This result was expected inasmuch as, the random set methodology cannot cope with distributional probability boxes and has to treat them as distribution-free p-boxes, as discussed in Section III-C-2.

The computational costs using the optimization approach in the epistemic space is less intensive than the propagation of focal sets inasmuch as only four optimization tasks are required to find the lower and upper bounds of $J_{1}$ and $J_{2}$ while the counter approach requires a pair of optimization tasks for each focal

Table 5: Bounds of the variable $J_{1}$ and $J_{2}$ for the reduced and improved uncertainty model obtained by means of the two proposed approaches (Optimization in the epistemic space and Propagation of focal sets), respectively.

| Reduced Uncertainty model | Improved Uncertainty model | Strategy |
| :--- | :--- | :--- |
| $J_{1}=\left[1.37 \times 10^{-2}, 4.97\right]$ | $J_{1}=\left[2.88 \times 10^{-2}, 1.11\right]$ | Optimization in the epistemic space |
| $J_{2}=\left[6.4 \times 10^{-2}, 0.82\right]$ | $J_{2}=[0.24,0.38]$ |  |
| $J_{1}=\left[-1.57 \times 10^{-4}, 54.05\right]$ | $J_{1}=\left[-1.10 \times 10^{-4}, 3.05\right]$ | Propagation of focal sets |
| $J_{2}=[0,1]$ | $J_{2}=[0.20,0.41]$ |  |

element and for each quantity of interest (i.e. $J_{1}$ and $J_{2}$ ). Both approaches are based on global optimization strategies and hence, they both suffer from the curse of dimensionality. The approaches proposed require an increasingly larger sample size (number of individuals and generations) in order to explore properly the optimization domain. In consequence, it is no longer guaranteed that the calculated optima are actually the global ones. In uncertainty propagation problems, missing the global optima means computing ranges of the targeted variables that are narrower than the sought ones. In this case, the methods result in an under(inner)-estimation of the actual solution, which may lead to an under-prediction of e.g. the failure probability of the system.

## D. Subproblem D

Subproblem D aims at identifying the epistemic realizations that lead to the smallest and largest values of $J_{1}$ (task D1) and $J_{2}$ (task D2). The extreme case analysis has been performed both for the reduced uncertainty model and the improved uncertainty model, as requested. However, for conciseness, only results from the improved model will be herein presented.

The extreme case analysis in presence of uncertainty is an ill posed inverse problem. The direct identification of the epistemic realizations, $\boldsymbol{\theta}$, leading to the maximum/minimum of $J_{1}$ and $J_{2}$ from the forward simulation has not been possible. Further, due to the complexity of the problem (in terms of nonlinearity and computational costs), a specific strategies has been developed as explained in the following section.


Figure 15: Effect of the number of samples sampled from the aleatory space in the inner loop on the estimation of $J_{1}$ and $J_{2}$, respectively, in the optimization in the epistemic space approach applied for the solution of subproblem C.


Figure 16: Convergence of the objective function $w$ to the minimum and maximum for a representative focal element. Genetic Algorithms have been used with a population of 1000 individuals to identify the realizations in the epistemic space that minimize and maximize the objective function $w$.

## 1. Extreme values of $J_{1}$ (task D1)

In this task we are focusing on $J_{1}=E[w]$ that is the expectation (mean) of the worst-case requirement metric: $w=\max _{i=1: 8}\left(g_{i}\right)$. In order to be able to identify the realizations of the inputs $\boldsymbol{p}$ that produce the extreme values of $J_{1}$, the relationships among intermediate variables, $\boldsymbol{g}, \boldsymbol{x}$ and $\boldsymbol{p}$ are analyzed.


Figure 17: Analysis of the performance function $\boldsymbol{g}$ with respect to the output of the subdisciplines, $\boldsymbol{x}$. In the plot the ranges of $x_{i=1: 5}$ leading to large positive values of $g_{i=1: 8}$ are shown using coloured bars. Grey bars (and dashed lines) indicate variables that are not important for the maximum of the corresponding performance $g_{i}$ ).

Dependence of $J_{1}$ on $w$ The extreme values of $J_{1}$ depend on the presence of very large (but rare) values of $w$ (hereafter indicated as outliers of $w$ ). The outliers of $w$ can assume values $w>1000$, while the most probable values of $w$ are limited to values around 0 . Two well distinct classes for $w$ have been identified. A first class identifies values where $w<3$, and a second class identifies the outliers, where $w>100$ and have values as high as 1000 . Hence, $J_{1}$ may assume its smallest value only if no outliers are present. On the other hand, the more outliers are present, the larger the value of $J_{1}$.

Dependence between $\boldsymbol{g}$ and $\boldsymbol{x}$ Next, the dependence between the performance functions of the system $\boldsymbol{g}$ and the output of subdisciplines $\boldsymbol{x}$ is analysed. The interest is to identify values (and ranges) of $\boldsymbol{x}$ that produce the maxima of the performance functions $\boldsymbol{g}$.

This study is performed by means of an optimization procedure where $g_{i=1: 8}$ are the objective functions
to be maximized and $\boldsymbol{x}$ are the search variables. Genetic Algorithm with 243 individuals and 50 generations is used for analysing each performance function $g_{i}$. The results are shown in Figure 17. The analysis of the function $x_{-} t o_{-} g$ (i.e. the model $f$ ) has revealed that only the performance functions $g_{i=3: 8}$ yield values $w>100$, while $g_{1}$ and $g_{2}$ are always lower than 1 and 2.8 , respectively.

Then, the individuals that produce $g_{1}>0.1, g_{2}>0.1$ and $g_{i=3: 8}>100$ are collected and shown in Figure 17 using coloured bars.

Some variables, shown in the Figure using grey color and dashed line, do not influence the maximum of the performance functions (i.e. they can assume any value within their bounds).

From Figure 17 critical sets (or regions) for each variable $x_{i}$ can be identified. For instance, there are three sets of $x_{1}$ able to produce values of $g_{4}>100$, namely $x 1 \in[0,0.05] \cup[0.82,0.91] \cup[1.11,1.17]$. However, these sets have been found without taking into account the probability distributions associated to the inputs $p_{i=1: 5}$. The most probable regions of $\boldsymbol{x}$ has been identified by means of the double loop Monte Carlo simulation used in Section C.

Interestingly, the most probable realizations of $\boldsymbol{x}$ that produce outliers of $w$ belong to a very clear pattern of coordinates, as shown in Figure 18.


Figure 18: Parallel coordinates of $x_{i=1: 5}$ leading to the outliers of $w$. The plot shows also the bounds of the variables $x_{i=1: 5}$ identified for the improved uncertainty model.

Dependence between $\boldsymbol{x}$ and $\boldsymbol{p}$ Once the regions of $\boldsymbol{x}$ that produce the outliers of $w$ have been identified, it is necessary to establish if such critical sets can be produced by any feasible realizations of inputs $\boldsymbol{p}$. This analysis has been performed by studying the functions $p_{-}$to_x (i.e. the model $h$ ) by using a double loop Monte Carlo approach, with an outer loop of 10000 Latin Hypercube samples (for the epistemic uncertainty,
$\boldsymbol{\theta}$ ) and an internal loop (for the aleatory uncertainty, $\boldsymbol{\alpha}$ ) of 1000 samples.


Figure 19: Parallel coordinates of the inputs $p_{i=1: 21}$ leading to values of $w>1000$ and $J_{1}>1.0$. The y-axis has been normalized between the lower and upper bound of the inputs $p_{i=1: 3,6: 21} . p_{4}$ and $p_{5}$ have been normalized between $*[-5,5]$.

Epistemic realization that produce maximum of $J_{1}$ Figure 19 shows the identified realizations of $\boldsymbol{p}$ that produce critical values of $\boldsymbol{x}$ (as shown in Figure 17). Only some inputs can lead unequivocally to the critical values of $\boldsymbol{x}$, namely $p_{1}, p_{4}, p_{5}, p_{14}, p_{15}$ and $p_{21}$. In the matter of $p_{4}$ and $p_{5}$, only values in the region where $3.72<p_{4}<4.70$ and $-3.46<p_{5}<-2.70$ can produce $x_{1}$ in the critical set and hence leading to large values of $w$. Since $p_{4}$ and $p_{5}$ are normally distributed, it is possible to select distributions peaked around the identified region as shown in Table 6. The epistemic realizations of Table 6 are calculated by maximizing the joint probability $\pi_{p_{4} p_{5}}=P\left[3.72<p_{4}<4.70,-3.46<p_{5}<-2.70\right]$. Using the distribution parameters reported in Table 6, such target maximum probability is $\max _{\Theta}\left(\pi_{p_{4} p_{5}}\right)=0.9912$.

Table 6: Epistemic realizations of $p_{4}$ and $p_{5}$ leading to the maximum of $J_{1}$. The parameters of the multivariate distribution are calculated maximizing the probability $\pi_{p_{i}}$ of being inside the specified ranges (i.e. Critical range $R_{c}$ ).

|  | $\pi_{p_{i}}^{\max }$ | Critical range $\left(R_{c}\right)$ | Epistemic real. |  |  |
| :---: | :---: | :---: | :---: | :---: | :---: |
| $p_{4}$ | 0.9912 | $3.72<p_{4}<4.70$ | $E\left(p_{4}\right)=4.21$ | $V\left(p_{5}\right)=\underline{V}\left(p_{4}\right)$ | $\rho=0$ |
| $p_{5}$ | 0.9912 | $-3.46<p_{5}<-2.70$ | $E\left(p_{5}\right)=-3.04$ | $V\left(p_{5}\right)=\underline{V}\left(p_{5}\right)$ | $\rho=0$ |

Epistemic realizations corresponding to parameters $p_{1}, p_{14}, p_{15}$ and $p_{21}$ are also calculated in a similar way. Table 7 show the epistemic realizations of these inputs corresponding to the critical values, and the
second column shows the corresponding values of the maximum probabilities $\pi_{p_{i}}$.

Table 7: Epistemic realizations of $p_{1}, p_{14}, p_{15}$ and $p_{21}$ leading to the maximum of $J_{1}$. These realizations maximize the probability of the input parameter $\pi_{p_{i}}$ of being inside the specified ranges $\left(R_{c}\right)$.

|  | $\pi_{p_{i}}^{\max }$ | Critical range $\left(R_{c}\right)$ | Epistemic real. |  |
| :---: | :---: | :---: | :---: | :---: |
| $p_{1}$ | 0.141 | $0.81<p_{1}<0.9$ | $E\left(p_{1}\right)=\bar{E}\left(p_{1}\right)$ | $V\left(p_{1}\right)=\underline{V}\left(p_{1}\right)$ |
| $p_{14}$ | 0.854 | $0.00<p_{14}<0.54$ | $a\left(p_{14}\right)=\underline{a}\left(p_{14}\right)$ | $b\left(p_{14}\right)=\bar{b}\left(p_{14}\right)$ |
| $p_{15}$ | 0.940 | $0.29<p_{15}<0.78$ | $a\left(p_{15}\right)=\bar{a}\left(p_{15}\right)$ | $b\left(p_{15}\right)=6.498$ |
| $p_{21}$ | 0.077 | $0.27<p_{21}<0.45$ | $a\left(p_{21}\right)=\bar{a}\left(p_{21}\right)$ | $b\left(p_{21}\right)=\underline{b}\left(p_{21}\right)$ |

$p_{1}$ and $p_{21}$ are somehow problematic inputs in the determination of the epistemic realization. By analysing the realizations from the input parameters $p_{1}$ and $p_{21}$, it can be seen that critical values of $\boldsymbol{x}$ are obtained when $0.805<p_{1}<0.902$ and $0.27<p_{21}<0.45$, respectively. However, from the p-boxes associated to these inputs (see Figure 20), it is not possible to select any CDF within in the p-box of $p_{1}$ and $p_{21}$ that permits to exclude (or include) completely the critical realizations (shown as round dots in Figure 20).


Figure 20: P-box representation of parameter $p_{1}$ and $p_{21}$, respectively. The figures show the ranges of values that produce critical values of $\boldsymbol{x}$ (and in turn large values of $w$ ).

Epistemic realization of the remaining parameters $\boldsymbol{p}$, which do not appear to have influence in the
generation of the critical values of $w$ (see Figure 19), have been obtained by maximizing the probability $\nu_{c}=P\left[w>1000 \mid p_{i} \in R_{c}\left(p_{i}\right)\right]$ for $i=1,4,5,14,15,21$. A random search for the maximum values of the mean of $p_{i}$ has been performed. 1000 aleatory samples have been used to calculate the above conditional probability. The results are reported in Table 8. The maximum identified frequencies is $\nu_{c}^{\max }=0.572$ and minimum $\nu_{c}^{\min }=0.261$. These values are quite close meaning that the epistemic uncertainty may play a secondary role for the extreme value of $J_{1}$.

The parameters of the p-boxes have been calculated using the identified values of $E\left[p_{i}\right]$ and the maximum admissible value for $V\left[p_{i}\right]$.

Table 8: Epistemic realization that are very likely to produce the maximum of $J_{1}$. The realization has been identified maximizing the probability $\nu_{c}$.

| Parameter | Epistemic real. | Parameter | Epistemic real. |
| :---: | :---: | :---: | :---: |
| $p_{2}$ | 0.719 | $p_{12}$ | $\underline{p_{12}}$ |
| $p_{6}$ | 0.760 | $p_{13}$ | $a=0.45, b=\bar{b}$ |
| $p_{7}$ | $a=\underline{a}, b=0.73$ | $p_{16}$ | 0.590 |
| $p_{8}$ | $a=\bar{a}, b=\underline{b}$ | $p_{17}$ | $a=\underline{a}, b=1.32$ |
| $p_{10}$ | $a=3.55, b=\bar{b}$ | $p_{18}$ | $a=3.26, b=\underline{b}$ |
| $p_{12}$ | $\underline{p_{12}}$ | $p_{20}$ | $a=10.68, b=\bar{b}$ |

The realization leading to the minimum of $J_{1}$ can be directly identified from results of task C 1 (see Section C). The results are summarized in Figure 21.

## 2. Extreme values of $J_{2}$

The task D2 asks to identify the extreme case for metric $J_{2}$, where $J_{2}=P[w \geq 0]$ is the failure probability of the worst-case requirement metric $w=\max _{i=1: 8}\left(g_{i}\right)$. Differently form $J_{1}$, this metric is not sensitive to the largest values of $w$. A double loop Monte Carlo approach has been adopted to solve this problem. 1000 aleatory samples have been used to compute the failure probability $J_{2}$. It is known from Section C that both lower and upper bounds of $J_{2}$ are greater than $10^{-1}$, hence 1000 samples are enough for a sufficiently


Figure 21: Extreme case analysis of $J_{1}$ : parallel plot of the epistemic parameters. The y-axis represents normalized values of the epistemic variables.
robust estimation of $J_{2}$ in the analysis.
The realizations of the input parameters $\boldsymbol{p}$ that produce the extreme values of $J_{2}$ are shown in Figure 22. Results from this analysis show, as expected, that realizations leading to the maximum (minimum) of $J_{2}$ are generally different from those leading to the maximum (minimum) of $J_{1}$. It is also noted that many realizations are very close to the bounds of the epistemic domain.


Figure 22: Extreme case analysis of $J_{2}$ : parallel plot of the epistemic parameters. The y-axis represents normalized values of the epistemic variables.

## 3. Solution of task D3

In task D3, it is asked to identify some representative realizations of $\boldsymbol{x}$ that typify different failure scenarios. The results of this task have already been discussed in Section D-1 and visualised in Figure 17. Overall, the following failure scenarios have been identified:

- Values of $x_{i=1: 5}$ close to their upper bounds lead to large values of $g_{i=1: 8}$;
- Small values of $x_{1}$ combined with large values of $x_{i=2,3,4,5}$, lead to values of $g_{i=3,4,5,6,7,8}>1000$;
- Values of $x_{1} \in[0.84,0.9]$ combined with large values of $x_{i=2,3,4,5}$, lead to values of $g_{i=4,6,7,8}>1000$
- Values of $x_{1} \in[0.4,0.425]$, combined with large values of $x_{2,3,4,5}$, lead to values of $g_{1}>0.1$ and

$$
g_{i=3,5}>1000 .
$$

Analysing the results of the simulations used in Section C ( Genetic Algorithm with 125 individuals and 45 generations), it is also possible to study the relationship between $\boldsymbol{x}$ and $\boldsymbol{g}$. For example, large positive values of $g_{5}$, whose maximum is $g_{5}^{\max }=1021$, are insensitive to $x_{3}$ and $x_{4}$. This can be appreciated in Figure 23, where the evolution of objective function $g_{5}$ and search variables $x_{i}$ are represented. During the optimization the values of variables $x_{3}$ and $x_{4}$ change frequently, despite that the value of the objective remains the same. Analogously, for the other performances, it is found that large positive values of $g_{4}, g_{6}$, $g_{7}$ and $g_{8}$ are totally insensitive to $x_{2}$ and slightly insensitive to $x_{4}$.


Figure 23: Evolution of the objective function $g_{5}$ and search variables $x_{i}$. Note that the values of variables $x_{3}$ and $x_{4}$ change frequently during the optimization, despite the values of the objective remain the same.

## E. Subproblem E

The last task of the challenge problem is to perform a robust design of the multidisciplinary system: performing an optimization able to identify the design point $\boldsymbol{d}$ with improved robustness and reliability characteristics. This requires to perform UQ for each candidate solutions leading to unmanageable computational costs. Hence, it is necessary to adopt surrogate models. Here, it has been decided to replace with surrogate models only the computational costly part of the model and keeping the original functions for the less
demanding parts. In fact, training a surrogate model to approximate the non-linear, an noisy, functions $J_{1}$ and $J_{2}: \mathbb{R}^{21} \times \mathbb{R}^{14} \rightarrow \mathbb{R}$ would have required a huge number of training samples with no warranties on the quality of the approximation.

The subproblems E1 and E2 require the calculation of $w(\boldsymbol{p}, \boldsymbol{d})=\max _{i=1, \ldots, 8} g_{i}(\boldsymbol{x})$, where $\boldsymbol{g}(\boldsymbol{x})=\boldsymbol{f}(\boldsymbol{x}, \boldsymbol{d})$ and $\boldsymbol{x}=\boldsymbol{f}(\boldsymbol{p})$ where the most computationally expensive part is the evaluation of $\boldsymbol{g}(\boldsymbol{x})=\boldsymbol{f}(\boldsymbol{x}, \boldsymbol{d})$. Multilayer perceptron artificial neural networks ${ }^{53}$ are suggested to speed up the calculation of function $\boldsymbol{g}=\hat{\boldsymbol{f}}(\boldsymbol{x}, \boldsymbol{d})$ : $\mathbb{R}^{5} \times \mathbb{R}^{14} \rightarrow \mathbb{R}^{8}$; in other words, the artificial neural networks act here as nonlinear response surfaces.

To train the artificial neural networks, training examples of $\{\boldsymbol{x}, \boldsymbol{d}, \boldsymbol{g}\}$ are passed to an error backpropagation algorithm. A set $\boldsymbol{d}^{(i)}: i=1,2, \ldots, 2000$ of Latin Hypercube quasi-random points were generated in the 14-dimensional space of the design variables.The design variables $\boldsymbol{d}$ can theoretically assume any real valued quantity, but they have been actually generated in a bounded space to generate a local surrogate model. For the first local meta-model, the following bounds has been assigned to each design variable, $d_{i}:\left[\min \left(0.5 * d_{i, \text { baseline }}, 1.2 * d_{i, \text { baseline }}\right), \max \left(0.5 * d_{i, \text { baseline }}, 1.2 * d_{i, \text { baseline }}\right)\right]$. Please note that the baseline $d_{i}$ can also be negative, and this definitions guarantees that the baseline is included in the bounds.In case the optimization procedure would have found a optimum design laying on one of the bounds of the training region, a new local surrogate model would have needed to be trained, around the identify optimum. Then, the optimization procedure is restarted.

The generation of samples of $\boldsymbol{x}$ is more involved. One possible approach is to determine the bounding box of $\boldsymbol{x}$ using an optimization procedure over the function $\boldsymbol{x}=\boldsymbol{h}(\boldsymbol{p})$ and then draw samples from this box. However, with this approach samples will be drawn from regions where it is less probable to obtain values of $\boldsymbol{x}$, and where the neural network does not need to give an accurate prediction.

To concentrate the generation of training samples only in the region of space of higher probability of $\boldsymbol{x}$, a set $\boldsymbol{\theta}^{(i)}: i=1,2, \ldots, 2000$ of Latin Hypercube quasi-random points was generated in the 31-dimensional box of the epistemic space. For each set $\{\boldsymbol{\theta}, \boldsymbol{d}\}^{(i)}, 200$ Monte Carlo samples are generated in the aleatory uncertainty space, obtaining $\boldsymbol{x}^{(j)}: j=1,2, \ldots, 200$ realizations of the function $h(\boldsymbol{p})$. The main draw back of this procedure is that few samples will be generated in the tails of the distributions, thus the neural network will perform badly in the prediction of the extreme values.

In the end, 400000 points $\boldsymbol{x}, \boldsymbol{d}$ are available to compute the model outputs $\boldsymbol{f}(\boldsymbol{x}, \boldsymbol{d})$ and then train the
surrogate model $\hat{\boldsymbol{f}}(\boldsymbol{x}, \boldsymbol{d})$. Since the minimum values $g_{j}$, for variables $j=3$ to 8 are very close to zero and very small compared to the respective maximum value of variable $g_{j}$, the following nonlinear transformation of variables $g_{j}$ for $j=3,4, \ldots, 8$ is employed:

$$
\begin{equation*}
z\left(g_{j}\right):=\frac{1}{200\left|\min \left(g_{j}\right)\right|}-\frac{1}{100\left(g_{j}+2\left|\min \left(g_{j}\right)\right|\right)} \tag{34}
\end{equation*}
$$

This nonlinear transformation of variables $g_{j}$ stretches the interval of $g_{j}$ for those values that are close to zero but preserving the sign of $g_{j}$. This is a very important characteristic since zero represents the limit value between the failure and the safe region.

Using nonlinear transformation Eq. (34), we set map $g_{j}$ to:

$$
y_{j}= \begin{cases}g_{j} & \text { for } j=1,2  \tag{35}\\ 100 z\left(g_{j}\right) & \text { for } j=3, \ldots, 8\end{cases}
$$

in this way, $\mathcal{T}=\left\{\left(\boldsymbol{x}^{(s)}, \boldsymbol{y}^{(s)}\right): s=1,2, \ldots, 400000\right\}$ served as the set of samples that were used for training, validating and testing the artificial neural networks.

Given different levels of non-linearities in the relations between the inputs and each $g_{j}$, one multi layer perceptron has been trained for each $g_{j}$, and the optimal network architecture, i.e. characterized by the smallest regression error, has been identified for each output quantity. The first 300000 samples of $\mathcal{T}$ have been used to train each multilayer perceptron using the Levenberg-Marquardt learning algorithm, a least squares curve fitting algorithm. The rest of the samples were used for validating and testing the artificial neural networks. The LGPL library FANN (Fast Artificial Neural Network), ${ }^{53}$ integrated in OpenCossan, has been used. Finally, the surrogate model will approximate $\boldsymbol{g}(\boldsymbol{x})$ by applying the invers of the non-linear transformation of Equation 34

$$
g_{j}= \begin{cases}y_{j} & \text { for } j=1,2  \tag{36}\\ \left(\frac{1}{2\left|\min \left(g_{j}\right)\right|}-y_{j}\right)^{-1}-2\left|\min \left(g_{j}\right)\right| & \text { for } j=3, \ldots, 8\end{cases}
$$

where $y_{j}$ is the output of the artificial neural network. Tanks to the non-linear transformation, the neural network will provide a very accurate response for very small values of $g$, e.g., centered around 0 , at the expenses of a less accurate prediction for values of bigger magnitude.

The robust design requires to minimize the upper bound of $J_{1}$ and $J_{2}$ and those values need to be estimated for each candidate design $\boldsymbol{d}$.

Genetic algorithms have been used to identify the optimal $\boldsymbol{d}$ that minimize the largest value of $J_{1}$ using a population size of 50 individuals. $100 \alpha$-cuts are randomly generated for the input parameter $\boldsymbol{p}$ and 2500-sample internal Monte Carlo simulations are used to identify the upper and lower bounds of $w_{\alpha}$. This allows to estimate $\overline{J_{1}}$ for each candidate design in approximately 25 s , and hence leading to approximately 10 minutes of computational time for each generation using a local parallelization strategy (as explained in Section IV-B) on a Intel Xeon Processor E5-2450-v2 (8 cores at 2.5 GHz ). The identified optimum is:

$$
\begin{align*}
& \boldsymbol{d}_{E 1}=[0.0140,-0.2568,-0.0944,-0.4405,-0.1508,-0.1029,-0.0713, \ldots \\
&0.2002,-0.4431,0.2579,0.0044,-0.2086,0.6330,-0.0166] \tag{37}
\end{align*}
$$

corresponding to an optimum value of $\bar{J}_{1, \text { opt }}=0.0044$. Subsequent run of the optimization algorithm demonstrated that the optimum found is robust.

The optimal design identified is better than the baseline in respect to the range of $J_{1}$. In fact, the range of $J_{1}$ at the optimum design is $\left[\underline{J_{1}}, \overline{J_{1}}\right]=\left[1.798 \cdot 10^{-4}, 0.0044\right]$ and it is narrower than the range identified in Table 5. A optimum design point $\boldsymbol{d}$ that minimize the largest value of $J_{2}$ has also been identified. MonteCarlo simulation has been used to compute the upper value of the probability of failure, $\max \left(J_{2}\right)$. In order to reduce the coefficient of variation of the probability of failure estimator, $1500 \alpha$-cuts have been used. In order to asses the robustness of the identified optimum, the optimization have been performed 3 times using different initial populations and the results are shown in Table 9.

Although, the maximum of $J_{2}$ is very close, the identified design variables shows a large variability, in particular variables $d_{1}, d_{5}$ and $d_{14}$. In order to asses the importance of these design variables with respect to the computation of the upper bound of $J_{2}$, a global sensitivity analysis of the model with respect the design variables has been carried out. The first order indices has been computed by means of the extended-FAST method with 1000 samples. The most important (design) variables are $d_{9}, d_{4}, d_{12}$ and $d_{14}$, and the remaining variables has a lower, similar importance. It can be noticed that the range of $J_{2}$ at the optimum is larger than the range of $J_{2}$ using the base design (Table 5). However, it is important to keep in mind that the aim of the robust design is to reduce the upper bound of $J_{2}$ and not its range.

| run | $d_{1}$ | $d_{2}$ | $d_{3}$ | $d_{4}$ | $d_{5}$ | $d_{6}$ | $d_{7}$ | $d_{8}$ | $d_{9}$ |
| :---: | :---: | :---: | :---: | :---: | :---: | :---: | :---: | :---: | :---: |
| 1 | -0.0013 | -0.2322 | -0.0993 | -2.0426 | -0.2417 | -0.1681 | -0.0979 | -0.4362 | -0.5958 |
| 2 | 0.0052 | -0.2658 | -0.0874 | -1.0996 | 0.2852 | -0.1798 | -0.0981 | -0.4362 | -0.5958 |
| 3 | -0.0001 | -0.2722 | -0.1003 | -1.6712 | 0.3191 | -0.1640 | -0.0981 | -0.4362 | -0.5958 |
|  | run | $d_{10}$ | $d_{11}$ | $d_{12}$ | $d_{13}$ | $d_{14}$ | $J_{2}$ |  |  |
|  | 1 | 0.0730 | 0.0053 | -0.2012 | 0.5144 | -0.0083 | [0.0053 | .2973] |  |
|  | 2 | 0.3337 | 0.0053 | -0.2014 | 0.5875 | 0.0054 | [0.0047 | .2993] |  |
|  | 3 | 0.3230 | 0.0053 | -0.2127 | 0.5641 | 0.0187 | [0.0033-1 | . 3073 ] |  |

Table 9: Robust desing $\boldsymbol{d}_{E 2}$ with respect the upper bound of $J_{2}$

Finally, the sensitivity analysis for the identified design points that minimize the $\overline{J_{1}}$ and $\overline{J_{2}}$, have been rerun. The extended-FAST method have been computed using 16384 samples for each input factor for a total cost of 507904 simulations. Each model evaluation requires the propagation of the aleatory uncertainty and 500 Monte Carlo samples have been used. The sensitivity analysis has been performed using the real model $\boldsymbol{f}(\boldsymbol{x}, \boldsymbol{d})$ and not the surrogate model.

Figure 24 shows the sensitivity analysis of the epistemic input factors respect the performances $J_{1}$ and $J_{2}$ evaluated at the design points $\boldsymbol{d}_{E 1}$ and $\boldsymbol{d}_{E 2}$, respectively. The design point $\boldsymbol{d}_{E 1}$ seems to be very robust since all the sensitivity measures are all very small ( 1 order of magnitude smaller compared to the sensitivity of Section B. The most important factor that contribute the the variance of $J_{1}$ is the expected value of $p_{5}$. The most important factor that contribute the the variance of $J_{2}$ is the parameter $p_{14}$ followed by parameters $p_{2}, p_{1}$ and $p_{16}$. It is interesting to notice that the sensitivity analysis using the baseline for $\boldsymbol{d}$ does not show the importance of $p_{14}$ and $p_{2}$.

## VI. Conclusions

The development and design of robust safety-critical systems is a challenging problem since in general quantitative data is either very sparse or prohibitively expensive to collect. Moreover, the failure of such


Figure 24: First order sensitivity measure of the category II and III input factors evaluated. The first order sensitivity computed using extended-FAST method.
systems might have severe consequences. In order to increase confidence and consistency in safety predictions, modelling and simulation standards require estimates of uncertainty and descriptions of any processes used to obtain these estimates.

In this paper, a unified theory and an integrated and open general purpose computational framework to deal with scarce data, aleatory and epistemic uncertainties has been presented. The proposed computational tools are generally applicable to solve a reasonable large number of different problems and numerically efficient and scalable. The applicability of the proposed strategy has been shown solving addressing the NASA Langley UQ challenge problem.

The presented results of this challenge problem clearly show that there are many ways of performing analysis when different types of uncertainties, namely epistemic and aleatory, are present. All of these methods have at some point made some weak or strong assumptions in order to find an answer. This forms a sound basis for future improvements and developments. In fact, it is envisaged that this quantitative comparison of the approaches will be most instrumental and useful for the engineering community, since it will highlight the advantages and disadvantages of existing methods for the handling joint existence of epistemic and aleatory uncertainty. As a general remark, the suggested procedures reveal the capability of random set theory to represent without any assumption epistemic and aleatory uncertainty. A major drawback of the proposed techniques was that many of them were based, up to some point in global optimization algorithms, which is known to be difficult for noisy and high dimensional objective functions, and will lead to spurious results when convergence to non-global optima occurs. It is left as an open problem how to circumvent that optimization step when mapping focal elements through a function. Furthermore, the proposed techniques are still very computational demanding requiring up to millions of model evaluations. Clearly this can only be archived resorting to some sort of parallelizations strategies and to the computational power of cloud and cluster computing.

Considering different approaches to solve the same engineering problem might be seen a waste of resources and time. However, all the existing approaches for dealing with epistemic and aleatory uncertainty require fine tuning of their parameters in order to be efficient and accurate. Hence, it is of paramount importance to be able to verify and cross-validate the results against different procedures. In this respect, the availability of an open, flexible and modular computational framework implementing a number of different numerical
strategies is essential.

Apart from assessing existing procedures in model updating, sensitivity analysis, quantification of bounds on statistics, and optimal design, the challenge results are expected to serve as a reference for the engineering community in order to test new algorithms and computational procedures.

## Acknowledgments

Authors are particularly grateful to anonymous reviewers who considerably helped in improving the manuscript.

Matteo Broggi is a member of the Virtual Engineering Centre, which his located at the Daresbury Laboratory of the Science and Technology Facilities Council and partially funded by the North West Development Agency, the European Regional Development Fund and the Hartree Centre. These organizations are greatly acknowledged for their support. Part of this research was supported by the Hartree Centre, which is acknowledged for its support.

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