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Uncertainty quantification for personalized analyses of human proximal femurs

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Abstract

Computational models for the personalized analysis of human femurs contain uncertainties in bone material properties and loads, which affect the simulation results. To quantify the influence we developed a probabilistic framework based on polynomial chaos (PC) that propagates stochastic input variables through any computational model. We considered a stochastic E-ρ relationship and a stochastic hip contact force, representing realistic variability of experimental data. Their influence on the prediction of principal strains (ϵ_1 and ϵ_3) was quantified for one human proximal femur, including sensitivity and reliability analysis. Large variabilities in the principal strain predictions were found in the cortical shell of the femoral neck, with coefficients of variation of $\approx 40\%$. Between 60-80% of the variance in ϵ_1 and ϵ_3 are attributable to the uncertainty in the E-ρ relationship, while $\approx 10\%$ are caused by the load magnitude and 5-30% by the load direction. Principal strain directions were unaffected by material and loading uncertainties. The antero-superior and medial inferior sides of the neck exhibited the largest probabilities for tensile and compression failure, however all were very small ($p_f < 0.001$). In summary, uncertainty quantification with PC has been demonstrated to efficiently and accurately describe the influence of very different stochastic inputs, which increases the credibility and explanatory power of personalized analyses of human proximal femurs.

Keywords: femur, personalized medicine, uncertainty quantification, polynomial chaos, finite cell method

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1. Introduction

Computational models based on computed tomography (CT) are widely used to predict the mechanical behavior of human femurs (Bessho et al., 2007, Yosibash et al., 2013, Schileo et al., 2014, Ali et al., 2014). These have demonstrated their predictive accuracy with respect to in-vitro experiments, where a well-defined load is applied (Schileo et al., 2007, Cristofolini et al., 2010, Trabelsi et al., 2011, Ruess et al., 2012). In clinical practice, however, personalized physiological loading conditions are required. Magnitude and direction of the hip contact force are usually inferred from in-vivo measurements (Bergmann et al., 2001, 2010), but contain uncertainties because of inter- and intra-patient variations. Another major challenge is the determination of heterogeneous material properties (Taddei et al., 2007, Helgason et al., 2008, Eberle et al., 2013). The large scatter in the experiments that determine the relationship between Young's modulus E and a local densitometric measure ρ also induces uncertainties. Quantifying the influence of both uncertainties on the predicted mechanical response of the femur is mandatory when advocating computational models for clinical practice.

Uncertainty quantification is an essential part of model validation and is performed in three steps: characterizing the uncertain parameters, propagating them through the computational model, and estimating the stochastic response of interest (Oberkampf et al., 2004). Probabilistic studies that performed uncertainty quantification for computational models of human femurs are summarized in Table 1 along with their stochastic components, probabilistic methods, and aim of research.

[Table 1 about here.]

Material and hip loading uncertainties have been characterized most frequently. Some studies assumed a random but homogeneous Young's modulus for the entire femur (Bah and Browne, 2009, Mehrez and Browne, 2012) or simplified the spatial distribution of cortical and trabecular bone (Chang et al., 2001, Nicolella et al., 2006, Viceconti et al., 2006). Others considered material uncertainties within the E-ρ relationship (Chinchalkar and Taylor, 1989,

Taddei et al., 2006, Laz et al., 2007, Long et al., 2009), but each study applied a different relationship with different random variables. Regarding the loading conditions, all studies considered a quasi-static loading representing walking free or going upstairs. Uncertainty in the hip contact force was commonly described with a random variable for the magnitude while the direction was assumed deterministic. Only few studies considered also uncertainties within the force direction (Nicolella et al., 2006, Long et al., 2009, Dopico-González et al., 2010). These conceptual differences led us to consider the stochastic E-ρ relationship from Wille et al. (2012) in combination with the stochastic description of the peak hip contact force magnitude and direction from Yosibash et al. (2015), both representing realistic variabilities of experimental data.

Various probabilistic methods have been used for propagating uncertainties through a computational model, among them Monte Carlo (MC) simulation (Taddei et al., 2006, Viceconti et al., 2006, Laz et al., 2007, Dopico-González et al., 2010, Mehrez and Browne, 2012), response surface methods (Chang et al., 2001, Bah and Browne, 2009, Long et al., 2009), and the advanced mean-value method (Nicolella et al., 2006, Laz et al., 2007). These methods have in common that they are non-intrusive with respect to the computational model, i.e. they use the computational model as a black box and require no access to the solver. Accuracy and efficiency of the probabilistic methods vary, depending on the number of uncertain parameters and the stochastic response of interest. In this study we employed a different probabilistic method based on the concept of polynomial chaos (PC) (Ghanem and Spanos, 1990, Xiu and Karniadakis, 2002, Xiu, 2009), which has been shown to be superior in many engineering problems over the past two decades.

Most studies in Table 1 investigated the influence of model uncertainties within the context of a total hip replacement, which increases the model complexity considerably. Only two studies analyzed the mechanical behavior of femurs without an implant (Taddei et al., 2006, Laz et al., 2007), but limited the investigation to global performance indicators such as the maximum von Mises stress. Results typically included some descriptive statistics of the response variable (e.g. mean, standard deviation, 1st and 99th percentile), estimates of its distribution (probability density function or cumulative distribution function), and/or sensitivity parameters and probabilities of failure. None of the studies quantified uncertainties

3

₁ for the prediction of principal strains at different locations of the entire proximal femur.

₂ The objective of this research was to develop a general framework for uncertainty quan-
₃ tification, which propagates parameter uncertainties efficiently and accurately through any
₄ computational model in a non-intrusive manner. In addition to a complete stochastic de-
₅ scription of the response, the framework was required to enable global sensitivity analyses
₆ and simple reliability studies. We demonstrate uncertainty quantification for the personal-
₇ ized analysis of one human proximal femur, with focus on the prediction of principal strain
₈ magnitudes and directions.

₉ 2. Methods

₁₀ We considered in total four stochastic input variables when analyzing a patient-specific
₁₁ femur: one describing uncertainties in the E-ρ relationship and three describing uncertainties
₁₂ in the peak hip contact force magnitude and directions. The framework was used to compute
₁₃ mean and standard deviation as well as the probability density function of the stochastic
₁₄ response at various post processing locations. These stochastic results were verified with a
₁₅ Monte Carlo (MC) simulation. Additionally, a global sensitivity analysis of the stochastic
₁₆ response and a simple reliability analysis were performed.

₁₇ The framework was applied to the right proximal femur of a 56 y old male, denoted FF5
₁₈ in Yosibash et al. (2013). The donor died from a myocardial infarction, showed no skeletal
₁₉ disease, and had an approximated body weight of 800 N. Principal strain magnitudes and
₂₀ directions were computed with the Finite Cell Method (FCM) (Ruess et al., 2012), which
₂₁ performs the personalized analysis directly on the voxel data of a CT-scan. Bone FF5 had
₂₂ been scanned in water by a Philips Brilliance 64 CT Scanner (Eindhoven, Netherlands)
₂₃ resulting in a CT-scan comprised of 140 slices with 1.25 mm slice thickness and 0.26 mm in-
₂₄ plane resolution. Prior to uncertainty quantification, the accuracy of the FCM was rechecked
₂₅ by comparison to an in-vitro experiment reported in Yosibash et al. (2013).

₂₆ 2.1. Stochastic input variables

A stochastic relationship between ash density ρ_{ash} and Young's modulus E based on
pooled data from multiple experimental studies on femur tissue was presented in Wille et al.

(2012):

$$E = 12\,000 \cdot \rho_{\text{ash}}^{1.45} \cdot X_E \qquad\qquad E \text{ in [MPa]}, \ \rho_{\text{ash}} \text{ in [g/cm}^3] \qquad (1)$$

1 with $X_E \sim \ln\mathcal{N}(\mu_E = 0, \sigma_E^2 = 0.316^2)$ being a *log-normal* random variable. The two
2 parameters μ_E and σ_E^2 denote mean and variance of the associated normal distribution. X_E
3 characterizes the scatter of residual values around the regression mean (Wille et al., 2012).

4   The stochastic loading model from Yosibash et al. (2015) was considered. It describes
5 variations in the peak hip contact force during walking free and going upstairs and was
6 derived from databases *HIP98* (Bergmann, 2001) and *OrthoLoad* (Bergmann, 2008), which
7 contain *in-vivo* measurements of the hip contact force (Bergmann et al., 2001, Heller et al.,
8 2001). In total 141 data records from seven patients (3 female, 4 male, age: 55–82 y, weight:
9 49–101 kg) were considered for the stochastic loading model. Using the anatomical refer-
10 ence frame of OrthoLoad, the hip contact force is described by its magnitude F and two
11 corresponding angular directions A_x and A_y; details are given in Yosibash et al. (2015).

     We restricted this study to the stochastic loading model for going upstairs, as it has the
largest spread in data:

$$X_F \sim \mathcal{N}(0.97 \cdot BW + 1465, \ 277^2) \qquad\qquad F, BW \text{ in [N]} \qquad (2)$$

$$X_{A_x} \sim \mathcal{N}(15.25, \ 3.38^2) \qquad\qquad A_x \text{ in [deg]} \qquad (3)$$

$$X_{A_y} \sim \mathcal{N}(19.69, \ 6.49^2) \qquad\qquad A_y \text{ in [deg]} \qquad (4)$$

12 where BW denotes the body weight. The three independent *normal* random variables X_F,
13 X_{A_x}, and X_{A_y} characterize both inter- and intra-patient variability of the force magnitude
14 and direction, respectively (Yosibash et al., 2015). Inserting the assumed body weight of
15 800 N in (2) yields $X_F \sim \mathcal{N}(2241, \ 277^2)$, which completes the stochastic hip loading for the
16 personalized analysis of bone FF5.

17 *2.2. Uncertainty propagation with polynomial chaos*

18   We consider the computational model as a black box denoted by $\mathcal{M}$ into which the
19 random vector $\mathbf{X} = \left[X_E, X_F, X_{A_x}, X_{A_y}\right]^T$ is input, and denote the stochastic output of
20 interest by $Y = \mathcal{M}(\mathbf{X})$. The random variable Y can be any scalar quantity at a specific

1 location, like maximum or minimum principal strains (ϵ_1 or ϵ_3), or principal directions in

2 spherical coordinates (polar angle θ and azimuthal angle ϕ).

The main idea of PC is to represent Y by a series of orthogonal polynomials that depend on standardized random variables (Ghanem and Spanos, 1990). The distribution type of the input random variables defines the specific family of orthogonal polynomials. In our case, every input variable in $\mathbf{X}$ is related to an independent, *standard normal* random variable $U \sim \mathcal{N}(0, 1)$:

$$X_E = \exp(0.316 \cdot U_1) \tag{5}$$

$$X_F = 2241 + 277 \cdot U_2 \tag{6}$$

$$X_{A_x} = 15.25 + 3.38 \cdot U_3 \tag{7}$$

$$X_{A_y} = 19.69 + 6.49 \cdot U_4 \tag{8}$$

3 for which the Hermite polynomials form an orthogonal basis (Xiu and Karniadakis, 2002).

4 The polynomial chaos expansion (PCE) of the stochastic response reads then:

$$Y = \mathcal{M}(\mathbf{X}) = \sum_{\boldsymbol{\alpha} \in \mathcal{A}} y_{\boldsymbol{\alpha}} \Psi_{\boldsymbol{\alpha}}(\mathbf{U}) \tag{9}$$

5 where $y_{\boldsymbol{\alpha}}$ are unknown coefficients, $\boldsymbol{\alpha} = (\alpha_1, \ldots, \alpha_4)$ a set of indices, and $\mathbf{U} = [U_1, U_2, U_3, U_4]^T$

6 the vector of standard normal variables. The corresponding basis functions are $\Psi_{\boldsymbol{\alpha}}(\mathbf{U}) =$

7 $h_{\alpha_1}(U_1) \cdot h_{\alpha_2}(U_2) \cdot h_{\alpha_3}(U_3) \cdot h_{\alpha_4}(U_4)$ with h_{α_i} denoting the normalized Hermite polynomial

8 of degree α_i. In practice, the PCE (9) is truncated at a specific order p defining the highest

9 polynomial degree of $\Psi_{\boldsymbol{\alpha}}$. This limits the total number of coefficients $y_{\boldsymbol{\alpha}}$, which is given by

10 the binomial coefficient $\binom{4+p}{p}$ (Sudret, 2008). When approximating Y with PCEs of order

11 $p = 1$, 2, 3, or 4, then only 5, 15, 35, or 70 coefficients have to be determined, respectively.

12 Because the orthogonality of the basis functions is defined with respect to the expectation

13 operator $\mathbb{E}$, it holds for the Hermite polynomials that:

$$\mathbb{E}\left[\Psi_{\boldsymbol{\alpha}}(\mathbf{U})\,\Psi_{\boldsymbol{\beta}}(\mathbf{U})\right] = \int_{\mathcal{D}_{\mathbf{U}}} \Psi_{\boldsymbol{\alpha}}(\boldsymbol{u})\,\Psi_{\boldsymbol{\beta}}(\boldsymbol{u})\,f_{\mathbf{U}}(\boldsymbol{u})\,\mathrm{d}\boldsymbol{u} = \begin{cases} 1 & \boldsymbol{\alpha} = \boldsymbol{\beta} \\ 0 & \boldsymbol{\alpha} \neq \boldsymbol{\beta} \end{cases} \tag{10}$$

14 where $f_{\mathbf{U}}(\boldsymbol{u})$ is the joint probability density function of random vector $\mathbf{U}$, which defines

15 for every event $\boldsymbol{u}$ in the support space $\mathcal{D}_{\mathbf{U}}$ the respective probability measure $f_{\mathbf{U}}(\boldsymbol{u})\,\mathrm{d}\boldsymbol{u}$.

6

1   Utilizing (10) in combination with (9), the coefficients $y_{\boldsymbol{\alpha}}$ are obtained by:

$$y_{\boldsymbol{\alpha}} = \mathbb{E}\left[Y\,\Psi_{\boldsymbol{\alpha}}(\mathbf{U})\right] = \int_{\mathcal{D}_{\mathbf{U}}} \mathcal{M}\left(\boldsymbol{x}\right)\,\Psi_{\boldsymbol{\alpha}}(\boldsymbol{u})\,f_{\mathbf{U}}(\boldsymbol{u})\,\mathrm{d}\boldsymbol{u} \tag{11}$$

2   The integral (11) is approximated by a quadrature:

$$y_{\boldsymbol{\alpha}} \approx \sum_{i=1}^{Z} w_i\,\mathcal{M}\left(\boldsymbol{x}^{(i)}\right)\,\Psi_{\boldsymbol{\alpha}}(\boldsymbol{u}^{(i)}) \tag{12}$$

3   with $\boldsymbol{u}^{(i)}$ and w_i denoting the quadrature points and weights of the *Gauss-Hermite* quadra-
4   ture, respectively. The corresponding model inputs $\boldsymbol{x}^{(i)}$ are derived from $\boldsymbol{u}^{(i)}$ using (5)–(8).
5   Evaluating the multi-dimensional integral (11) by tensor products of 1-D quadrature re-
6   sults in an exponential growth of quadrature points as the number of dimensions increases.
7   This curse of dimensionality is circumvented by using Smolyak's quadrature scheme instead
8   (Smolyak, 1963); we utilized the implementation provided by Heiss and Winschel (2008).

9   When Y is approximated with a PCE of order $p = 1$, 2, 3, or 4 for instance, then the
10   computation of all 5, 15, 35, or 70 coefficients with (12) requires only 9, 41, 137, or 385
11   simulation runs with the model $\mathcal{M}$, respectively. Each simulation run evaluates then a
12   different combination of E-ρ relationship, force magnitude, and load directions, according
13   to the quadrature points. The hierarchical nature of both PCEs and Smolyak's quadrature
14   scheme allowed us to adaptively increase the number of quadrature points and to reuse lower
15   order approximations.

16   A MC simulation was conducted in order to verify the results of the PCE. MC involves
17   repeated *random* sampling of $\mathbf{X}$ and solving for each a deterministic problem, which results
18   in a sample set of Y. From this we computed the sample variance and an estimate of the
19   response distribution (histogram). Due to the slow convergence rate of the MC simulation,
20   many deterministic simulation runs are required; we performed 10000.

21   *2.3. Post-processing the stochastic response*

After all coefficients have been determined using (12), the mean μ_Y of the stochastic
response Y is described by the coefficient $y_{\boldsymbol{0}}$, and the variance σ_Y^2 of the stochastic response

7

is the sum of squares of all coefficients except y_0 (Sudret, 2008):

$$\mu_Y = \mathbb{E}[Y] = y_0 \tag{13}$$

$$\sigma_Y^2 = \mathbb{E}[(Y - \mu_Y)^2] = \sum_{\boldsymbol{\alpha} \in \mathcal{A} \backslash \mathbf{0}} y_{\boldsymbol{\alpha}}^2 \tag{14}$$

1   Please note that both parameters do not imply that Y is normally distributed, but are two
2   parameters that describe location and dispersion, respectively, of any response distribution.
3   We obtained the probability density function (PDF) of Y by kernel smoothing a large set
4   of response samples (Wand and Jones, 1995). For this purpose, the PCE was used as a
5   surrogate model to generate 10^6 samples of the response variable. These samples were also
6   used for a reliability analysis. For that we considered a simple strain based failure criterion
7   (Schileo et al., 2008, Yosibash et al., 2010, Schileo et al., 2014), which uses the yield strain for
8   femoral bone tissue in tension $\epsilon_{\lim,T} = 7\,300\,\mu\mathrm{m/m}$ and compression $\epsilon_{\lim,C} = -10\,400\,\mu\mathrm{m/m}$
9   (Bayraktar et al., 2004) as threshold values. Consequently, we defined the failure probability
10 p_f for (local) tensile or compression failure as the likelihood that the maximum principal
11 strain $\epsilon_1 \geq \epsilon_{\lim,T}$ or the minimum principal strain $\epsilon_3 \leq \epsilon_{\lim,C}$. We approximated p_f from the
12 number of response samples fulfilling the respective criteria, divided by the total number of
13 samples.

14 PCEs can be also used for a global sensitivity analysis. Sudret (2008) proved that PCEs
15 are identical to Sobol' decompositions of the model $\mathcal{M}$ (Sobol', 2001) and that the corre-
16 sponding Sobol' indices can be computed directly from the coefficients of the PCE. Sobol'
17 indices are *global* sensitivity indices, which represent the fraction of the response variance that
18 can be attributed to a specific input variable or their interactions. The first-order sensitivity
19 indices S_E, S_F, S_{A_x}, and S_{A_y} quantify in percentage the influence of each input variable taken
20 alone and were computed from the square-summed and normalized coefficients that are ex-
21 clusively associated with the respective input variable (Sudret, 2008). Also the sum of all re-
22 maining higher-order interaction indices was computed using $\sum S_{ij} = 1 - S_E - S_F - S_{A_x} - S_{A_y}$.

23 Finally, the stochastic results were visualized pointwise for an intuitive interpretation.
24 We depicted mean and variance of the maximum and minimum principal strain with a
25 sphere whose color is defined by the mean, whereas the radius depends on the standard
26 deviation. Failure probabilities were also represented by colored spheres, however all radii

1 were identical in that case. We verified the stochastic results of the PCE by comparing the

2 PDF with the respective histogram of the MC simulation. Moreover, we investigated the

3 convergence behavior of both methods approximating the variance.

4 *2.4. Deterministic simulation by FCM*

5 The FCM embeds the voxel-based geometry of bone FF5 in a simulation domain of hexa-

6 hedral cells following a regular Cartesian grid. Figure 1 illustrates this concept, which omits

7 a computational expensive segmentation and meshing procedure, as required for standard

8 finite element analyses. The FCM proved to be highly efficient for linear elastic analyses of

9 bones and was validated by in-vitro experiments in a previous study (Ruess et al., 2012).

10 [Figure 1 about here.]

The large computational cells were implemented as p-version hexahedral elements (Düster et al., 2008) and should not be mistaken for low order finite elements. Within one finite cell material properties can vary significantly, and it was shown that a coarse grid of cells is sufficient to obtain accurate results for a fine voxel resolution like the one used in the present study (Ruess et al., 2012). Inhomogeneous isotropic material properties were assumed for bone tissue (Trabelsi et al., 2011). After converting Hounsfield unit HU into equivalent mineral density ρ_{eqm} and then into ash density ρ_{ash} based on:

$$\rho_{\text{eqm}} = 10^{-3} \cdot (0.793 \cdot HU + 4.183) \qquad [\text{g/cm}^3] \qquad (15)$$

$$\rho_{\text{ash}} = 1.22 \cdot \rho_{\text{eqm}} + 0.0523 \qquad [\text{g/cm}^3] \qquad (16)$$

11 where (15) is the calibration of the CT-scan with K_2HPO_4 phantoms and (16) is from Keyak

12 and Falkinstein (2003), the heterogeneous Young's modulus was derived from (1). Because

13 the random variable in (1) is location independent, it affects the heterogeneous Young's

14 modulus everywhere the same and thus can be regarded as a global scaling factor. In all

15 computations a constant Poisson ratio $\nu = 0.3$ was used.

16 The hip contact force was modeled as a surface load, which distributes the force over

17 a locally confined contact area on the head of the femur. Since the head is approximately

18 spherical, the contact area was designed as a spherical cap (radius of sphere: 24 mm, radius

9

1. of base of the cap: 10 mm). The applied pressure load and the orientation of the contact

2. area were chosen according to the stochastic input variables X_F, X_{A_x} and X_{A_y}.

3. Within the probabilistic framework this computational model was used to assess max-

4. imum and minimum principal strains (ϵ_1, ϵ_3) and principal directions (θ, ϕ) at 884 post-

5. processing locations (554 spread along the femoral cortex, 330 uniformly filling the trabecular

6. compartment and the diaphysis).

7. *2.5. Verifying FCM's accuracy*

8. Prior to uncertainty quantification, the accuracy of the deterministic computational

9. model was rechecked by comparison to an in-vitro experiment (Yosibash et al., 2013). The

10. femur was loaded as in the experiment ($F = 1000$ N, $A_x = A_y = 0$ deg), and the median

11. of the stochastic E-ρ relationship (1) was used, since Median(X_E) = 1. Principal strains

12. were computed for the 11 surface locations (four at the superior and inferior neck, seven at

13. the medial and lateral diaphysis) at which strains had been measured (Yosibash et al., 2013,

14. Fig 2). Convergence in energy norm was investigated by increasing the polynomial order of

15. the FCM, i.e. $p_{\mathrm{FCM}} = 1, \ldots, 5$.

3. Results

17. The deterministic FCM has a high predictive accuracy: computed strains matched well

18. the ones measured in the experiment (correlation $r = 0.987$, average relative error of 18%).

19. Convergence in energy norm was achieved for $p_{\mathrm{FCM}} = 4$ (with an error of 8.4%). Thus, all

20. simulation runs were performed with $p_{\mathrm{FCM}} = 4$.

21. The stochastic results were verified by comparing the PDFs of the PCE to the respective

22. histograms of the MC simulation, as depicted exemplarily for one post-processing location

23. in Figure 2. The PCE of order $p = 4$ (385 Smolyak runs) was in excellent agreement with

24. the MC simulation (10000 runs), which is also reflected in the convergence behavior of both

25. methods (Figure 2b).

26. [Figure 2 about here.]

27. Means and standard deviations of the maximum and minimum principal strains are shown

28. in Figure 3. The vast majority of large strain values were found within the cortical shell.

10

Given that the loading had a pronounced inclination in the sagittal plane, ϵ_1 was largest at the anterior side at the distal diaphysis ($\mu = 2380$, $\sigma = 955$) and the antero-superior neck ($\mu = 1569$, $\sigma = 656$). Contrarily, the largest values for ϵ_3 were predominantly at the posterior side of the femur, with strain concentrations at the distal diaphysis ($\mu = -3230$, $\sigma = 1270$) and the medial neck above the lesser trochanter ($\mu = -2296$, $\sigma = 824$).

[Figure 3 about here.]

The largest failure probabilities within the region of the femoral neck were found to be $p_f = 13 \cdot 10^{-6}$ for tensile failure (point 351 in Figure 4a) and $p_f = 2 \cdot 10^{-6}$ for compression failure (point 231 in Figure 4b), respectively. Both locations are in the cortical shell. In general all failure probabilities were small ($p_f < 0.001$, at many locations even $p_f < 10^{-6}$).

[Figure 4 about here.]

Results of the sensitivity analysis for these two points are summarized in Table 2. ϵ_1 and ϵ_3 were most sensitive to the stochastic E-ρ relationship ($S_E \approx 60\%$ for ϵ_1 at point 351, and $S_E \approx 80\%$ for ϵ_3 at point 231). Around 10% of the variance in the principal strain magnitudes is attributed to the force magnitude; the influence of the load angles is larger for ϵ_1 ($S_{A_x} = 15.2\%$, $S_{A_y} = 12.2\%$) than for ϵ_3 ($S_{A_x} = 2.8\%$, $S_{A_y} = 2.4\%$). Different results were obtained for the principal directions: θ and ϕ of both principal strains were affected *solely* by A_x and A_y, which accounted together for more than 95% of the respective variances. The E-ρ relationship and the force magnitude had no influence on the principal directions (S_E, $S_F < 10^{-6}$). Note that the standard deviation of θ and ϕ were found to be very small ($< 5 \deg$). Although the load angles explained the vast majority of the variance, their absolute effect on the principal directions is minor.

[Table 2 about here.]

4. Discussion

A probabilistic framework was developed to quantify the influence of material and loading uncertainties on the prediction of principal strains within a personalized analysis of a

human proximal femur. The predicted strains were highly affected by the four stochastic input variables, which represented experimental variabilities in the E-ρ relationship and the hip contact force. The large standard deviations for ϵ_1 ($\sigma \leq 656$) and ϵ_3 ($\sigma \leq 824$) at the femoral neck imply broad probability distributions and thus large ranges of possible strain predictions. Even broader distributions were found at the distal diaphysis but can be attributed to the clamped boundary condition. The coefficient of variation (standard deviation divided by the mean) was $\approx 40\%$ for both ϵ_1 and ϵ_3 at all locations with sufficiently large mean values. This relative variability within the principal strain predictions is larger than the one reported in Taddei et al. (2006), which computed coefficients of variation of less than 9% for $\max \epsilon_1$ and $\max \epsilon_3$. These differences may be explained by the uncertainty in the E-ρ relationship, which is larger in the current study. For comparison purposes, assuming a bone density of $\rho_{ash} = 1.2\,\mathrm{g/cm^3}$, the 95% prediction band of the Young's modulus ranged between 8.4–29.1 GPa in this study, compared to 16.6–16.9 GPa in Taddei et al. (2006).

The global sensitivity analysis corroborated the dominating influence of the stochastic E-ρ relationship on the *magnitude* of ϵ_1 and ϵ_3. Within the femoral neck, between 60-80% of the variance of both principal strains were explained by the uncertainty in the E-ρ relationship. Because of its large influence compared to the other three input variables, a reduction of the uncertainty in the E-ρ relationship would have the largest effect on ϵ_1 and ϵ_3 and narrow the response distributions the most. This would require new experiments on bone tissue specimens with a well defined protocol that reduces the large spread in the current experimental data (Helgason et al., 2008, Wille et al., 2012).

Predictions of the principal strain *direction*, on the other hand, were completely unaffected by the uncertainties in the E-ρ relationship and the force magnitude. Only the load direction had an influence on the direction of the principal strains. However, the effect was marginal as the standard deviation of the principal strain directions was found to be very small within the femoral neck ($< 5\,\mathrm{deg}$), which is in agreement with experimental observations. Cristofolini et al. (2009) measured strains in 24 femurs for six different loading configurations and reported for all strain measurements a standard deviation of 6.7 deg.

A comparison with sensitivity results from other stochastic studies (Table 1) renders difficult. The main reason is that different sensitivity measures were used. Some studies

reported Pearson's correlation coefficient between the stochastic input variables and the response variables as sensitivity indices (Taddei et al., 2006, Viceconti et al., 2006, Dopico-González et al., 2010), which assess the strength of the linear association between them. In case the input affects non-linearly the computational model, such a correlation analysis can be very misleading. More meaningful are *local* sensitivity analyses, with sensitivity indices measuring how changes in the parameters of the stochastic input variables (e.g. mean or standard deviation) affect the probability of failure (Nicolella et al., 2006, Mehrez and Browne, 2012). In contrast, we performed a *global* sensitivity analysis, which is based on a variance decomposition for the stochastic response. Here, sensitivity indices quantify the percentage of the response variance attributable to an entire input variable or combinations of variables.

Regarding the reliability analysis, all failure probabilities were found to be very small ($p_f < 0.001$). The antero-superior neck region of this femur is more likely to suffer from local tension failure ($p_f \leq 13 \cdot 10^{-6}$) than the postero-medial neck region from compression failure ($p_f \leq 2 \cdot 10^{-6}$), when loaded with the peak hip contact force during going upstairs. However, both events are extremely improbable; an expected outcome given that the subject had no skeletal disease. Note that these local failure probabilities are not implying any global clinical failure or bone fracture. Instead they describe the likelihood of irreversible damage in the bone tissue at the respective location, which may cause fracture initiation.

The presented uncertainty quantification, including sensitivity and reliability analysis, was used largely to demonstrate the probabilistic framework based on PC. This powerful and widely accepted approach is novel in biomechanics, and to the best of our knowledge was only used for a cardiovascular simulation (Sankaran and Marsden, 2011). The PC approach was two orders of magnitude computationally more efficient than an extensive MC simulation (10000 runs) performed to verify the stochastic results. In fact, a PCE of lower order ($p = 3$) would have been sufficient to approximate the stochastic response in this study (cf. Figure 2), further reducing the number of necessary simulation runs from 385 to 137. Uncertainty quantification with a specific objective, e.g. determining percentiles or computing failure probabilities, might be possible at even lower computational costs with the advanced mean value method (Nicolella et al., 2006, Laz et al., 2007). However, a meaningful

comparison would require both methods to be applied to the same stochastic problem.

Further limitations of this study are related to the computational model and the stochastic input variables. Any personalized computational model requires assumptions on the geometry, material behavior, and boundary conditions. Uncertainties in geometry due to imprecise segmentation from CT-scans were not addressed, because they are considered negligible compared to material and loading uncertainties (Gelaude et al., 2008, Trabelsi et al., 2009). The material behavior of the bone was assumed to be linear elastic and isotropic, which is a simplification of the reality that proved to be reasonable for stance-like loading conditions (Trabelsi et al., 2011). Orthotropic or transversely isotropic material models may become necessary in case of loading conditions that introduce considerable torsional moments, e.g. during sidewise falling. Another limitation is the clamped boundary condition at the distal diaphysis. Although mean deflections were within physiological range ($< 2.5\,\mathrm{mm}$), realistic physiological boundary conditions would include kinematic constraints at the joints as well as muscle forces (Speirs et al., 2007). The absence of muscle forces in this study explains the very small strains observed in the greater trochanter area. However, we are unaware of any (stochastic) muscle force model that is suitable for the personalized analysis of human femurs. Clearly, all limitations related to the stochastic input variables pass on to the probabilistic analysis. For example, the stochastic peak hip contact force was derived from an elderly population (age 55–82 y) with total hip replacements (Yosibash et al., 2015). Here, the subject (age 56 y) falls inside the range, but had no hip implant. Moreover, the stochastic E-ρ relationship was obtained by pooling data sets of various experimental studies (Wille et al., 2012). The different experimental protocols contributed considerably to the uncertainty in the E-ρ relationship. In both cases it is important to enhance the respective data basis with further experimental results, as this will improve the description of the stochastic input parameters.

In closing, a probabilistic framework that allows to quantify the influence of parameter uncertainties on the personalized analysis of human femurs was presented. The probabilistic framework can be applied to any computational model and easily extended by additional stochastic input variables. Future studies may not only perform uncertainty quantification for the prediction of principal strains, but investigate also displacements, stresses, global

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bone stiffness, implant micro-motions etc. The global sensitivity indices can then be used to identify input parameters that have a negligible influence on the stochastic response and therefore can be safely considered as deterministic parameter. In case of clinical assessments, uncertainty quantification will significantly increase the credibility and explanatory power of computational models.

Conflict of interest statement

None of the authors have any conflict of interest to declare that could prejudice this work.

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1 Tables

Table 1: Summary of probabilistic studies performing uncertainty quantification for computational models of human femurs. The checkmark symbol (✓) indicates which components of the femur model were considered to be stochastic.

Table 2: Global sensitivity analysis for ϵ_1 at point 351 and ϵ_3 at point 231 (magnitude and θ and ϕ of the corresponding eigenvector). Shown are mean μ, standard deviation σ, related first-order sensitivity indices $S_F, S_{A_x}, S_{A_y}, S_E$, and the sum of all remaining higher-order interaction indices S_{ij}.
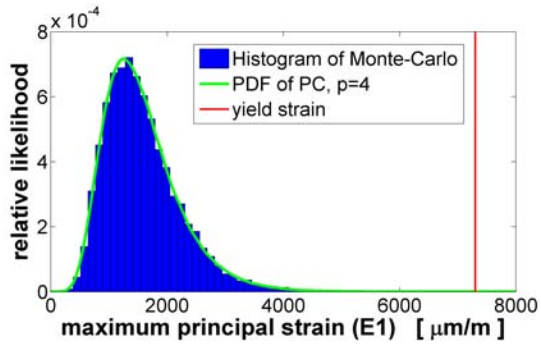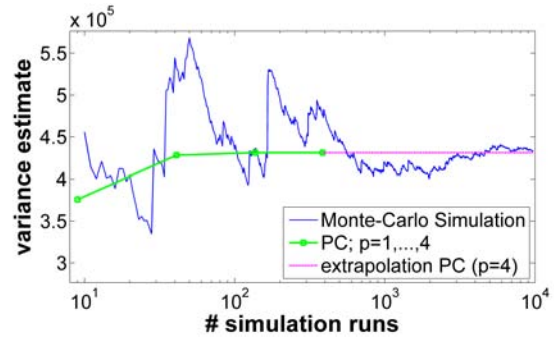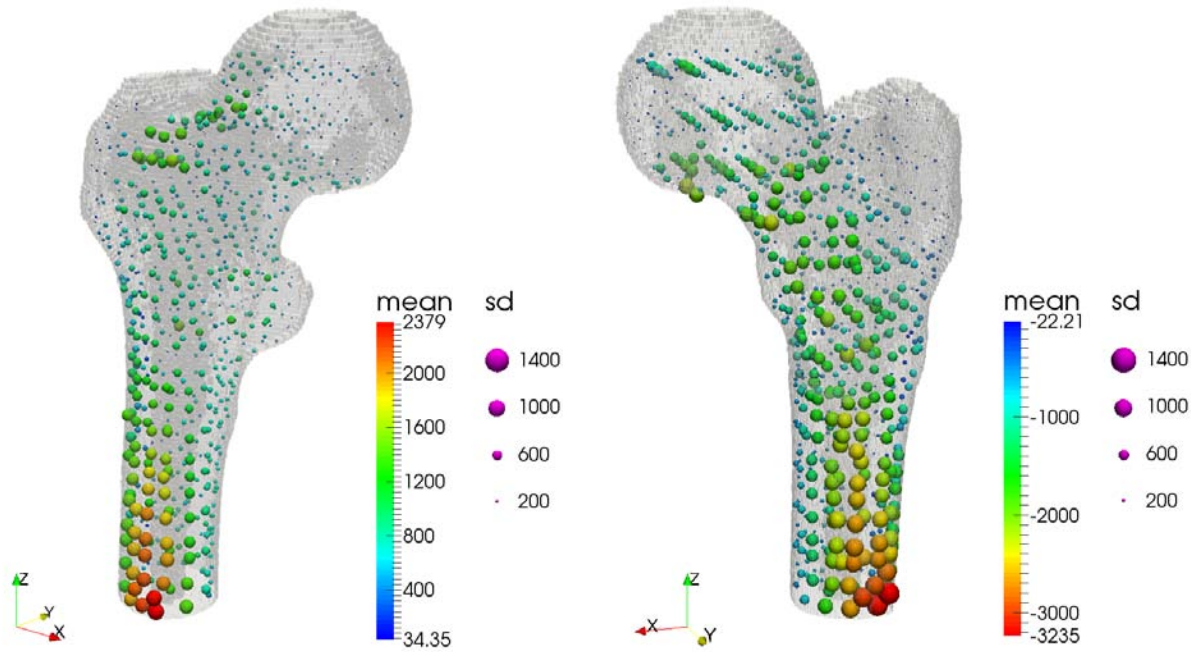
₁ **Figures**
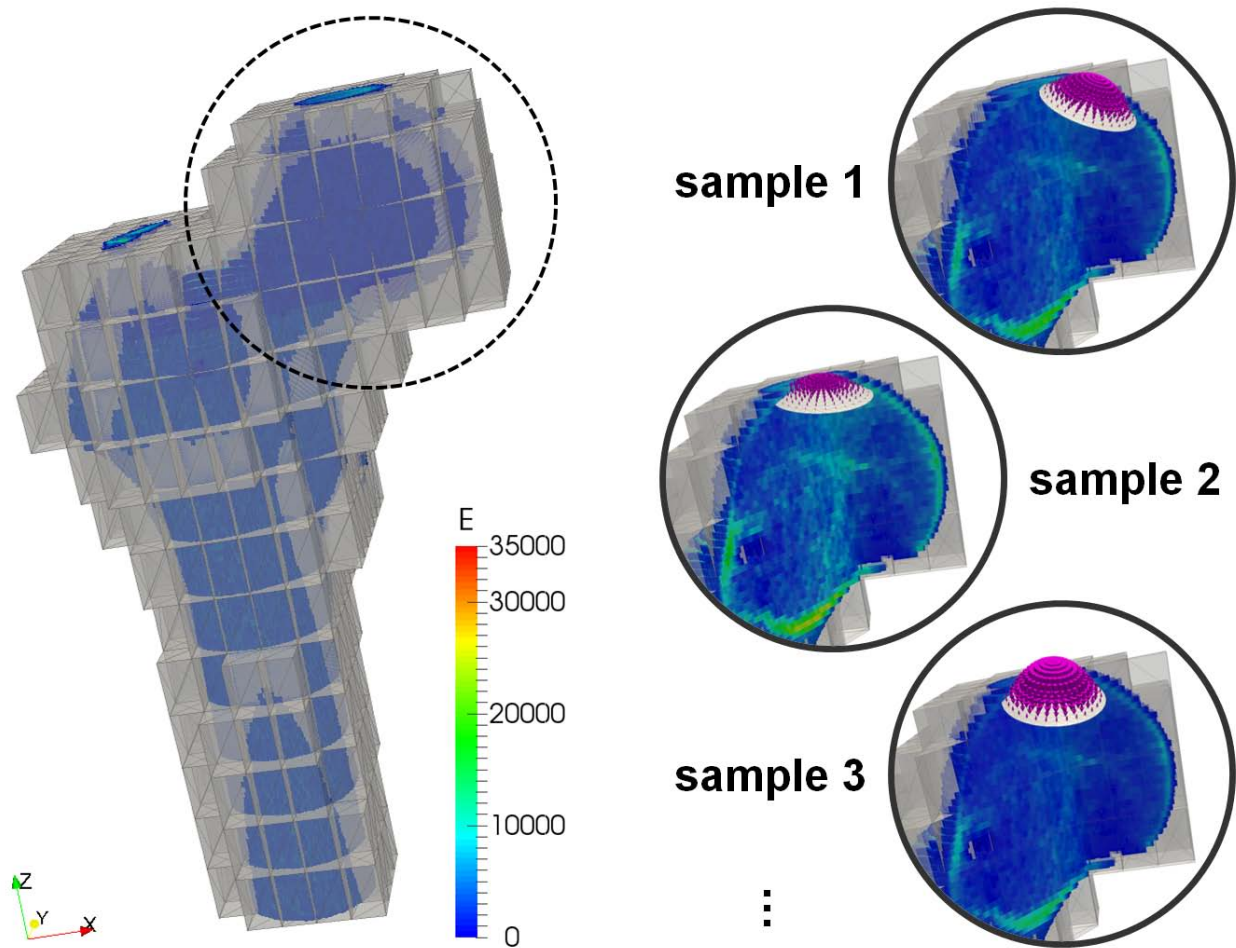
Figure 1: Deterministic finite cell model based on the CT scan of bone FF5 reported in Yosibash et al. (2013). The computational model is clamped at the distal end of the diaphysis. The hip contact force is modeled as surface load on a confined contact area on the head. During uncertainty quantification the heterogeneous Young's modulus E and the surface load (magnitude and orientation) are changed for every simulation run, as depicted by the samples that give an insight into the region of the dashed circle.

(a) Probability distribution of ϵ_1

(b) Convergence plot for estimating the variance of ϵ_1

Figure 2: Comparison of PCE with MC simulation for ϵ_1 at post-processing point 351.

(a) Maximum principal strain ϵ_1 [μm/m], antero-medial view

(b) Minimum principal strain ϵ_3 [μm/m], postero-medial view

Figure 3: Mean and standard deviation of principal strains represented by spheres (color = mean, radius of sphere = standard deviation). All 884 post-processing locations are shown at once (554 along the femoral cortex, 330 within the trabecular compartment and the diaphysis). Femur's geometry is indicated by a translucent voxel representation that was derived from the CT data. For animations of these results the reader is referred to the supplementary material.
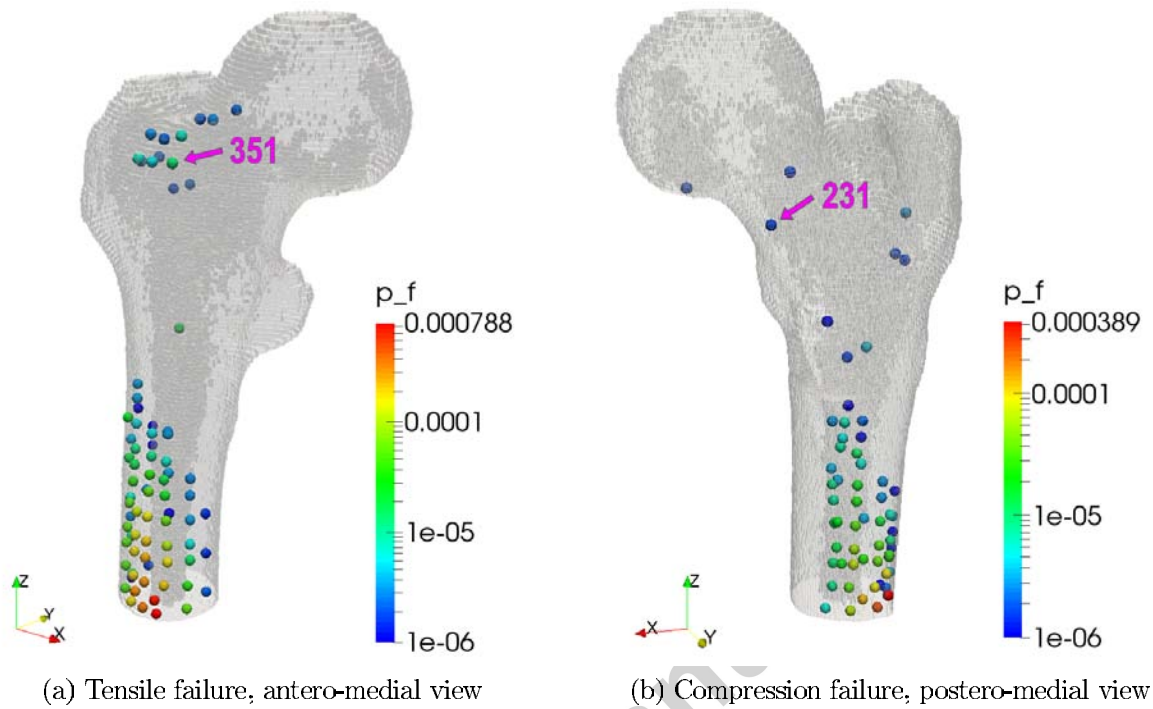
(a) Tensile failure, antero-medial view   (b) Compression failure, postero-medial view

Figure 4: Probability of failure p_f represented as colored unit sphere at locations with $p_f > 10^{-6}$. Within the femoral neck the largest probability of tensile failure is at point 351 and for compression failure at point 231. Femur's geometry is indicated by a translucent voxel representation that was derived from the CT data. For animations of these results the reader is referred to the supplementary material.

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