

## PERSPECTIVES

# CT-Based Visualization and Quantification of Bone Microstructure *In Vivo*

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### Abstract

It is well-accepted that in addition to bone volume, trabecular bone microstructure is an important factor influencing bone strength, which in turn is the most important bone parameter indicating bone fracture risk. The objective of this article is to describe some of the computed tomography (CT)-based techniques used to measure the microarchitectural aspects of bone quality, how these can be quantified and how these quantitative endpoints can be used in the assessment of bone competence. Microarchitectural bone imaging is a nondestructive, noninvasive, and precise procedure with which both the apparent density and trabecular microstructure of intact bones and bone samples can be assessed in a single measurement. Recently introduced high-resolution *in vivo* CT imaging systems now allow repetitive measurements of bone microarchitecture so that small local trabecular changes can be monitored over time. The procedure can help improve predictions of fracture risk, clarify the pathophysiology of skeletal diseases, and define the response to therapy and interventions. Hopefully, this improved understanding will lead to more successful approaches in the prevention and treatment of age- and disease-related fractures. *IBMS BoneKEy*. 2008 November;5(11):410-425.

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### Introduction

Osteoporosis is not a disease of low bone mass *per se*, but of low bone strength. The present “gold standard” to determine bone strength is through mechanical testing. Direct mechanical testing is a straightforward procedure, but is limited by its destructiveness. Therefore, this method is not applicable *in vivo*, and although it can be used *in vitro*, a sample can only be tested once, which limits the assessment of direction-dependent failure characteristics. Furthermore, these tests are prone to errors related to boundary artifacts and to the often small size of the specimens, hampering high precision measurements (1).

Bone mineral density (BMD) measurements are frequently used as a surrogate measure of bone strength. Significant correlations between the apparent bone density and different mechanical properties have been demonstrated for large populations using power-law regressions (2;3). A meta-

analysis showed that for trabecular bone samples, around 64% of the variability in bone strength *in vitro* is determined by its apparent density (4). Bone microstructure explains a significant part of the remaining variation; by including architectural parameters in the analysis the predictive power increases to over 90% for bone stiffness (5-7). Given that bone stiffness and strength are highly correlated ( $R^2 > 0.95$ ; (8;9)) the effects of bone microarchitecture are very likely to strongly affect bone strength as well. Indeed, this was shown by recent experimental data that used strength as an outcome parameter and which demonstrated the significant contribution of bone architecture in predicting bone strength (10). Also, for whole bone strength, bone microarchitecture plays a significant role. In a study (11) on the dose-dependent effect of ibandronate treatment on bone mass and architecture in ovariectomized (OVX) monkeys, it was found that bone mineral content (BMC) as assessed by dual-energy X-ray absorptiometry was the best single

predictor of whole bone strength ( $r^2 = 67\%$ ); by including architectural indices in a multiple linear regression analysis the correlation increased to 88%.

The introduction of microstructural imaging systems such as desktop micro-computed tomography ( $\mu$ CT) has made it possible to assess both the apparent density and the three-dimensional (3D) trabecular microstructure of intact bones in a single measurement. It has given researchers a powerful tool for the exploration of age-related bone loss and metabolic diseases such as osteoporosis. In this *Perspective* we will present methodologies to quantify bone microstructure, to measure bone deformation under load and to assess bone competence *in silico*. The focus will be on recently introduced *in vivo* systems that allow us to image and quantify bone microarchitecture in small animals and at peripheral sites in humans.

### Three-Dimensional Imaging of Bone Microarchitecture

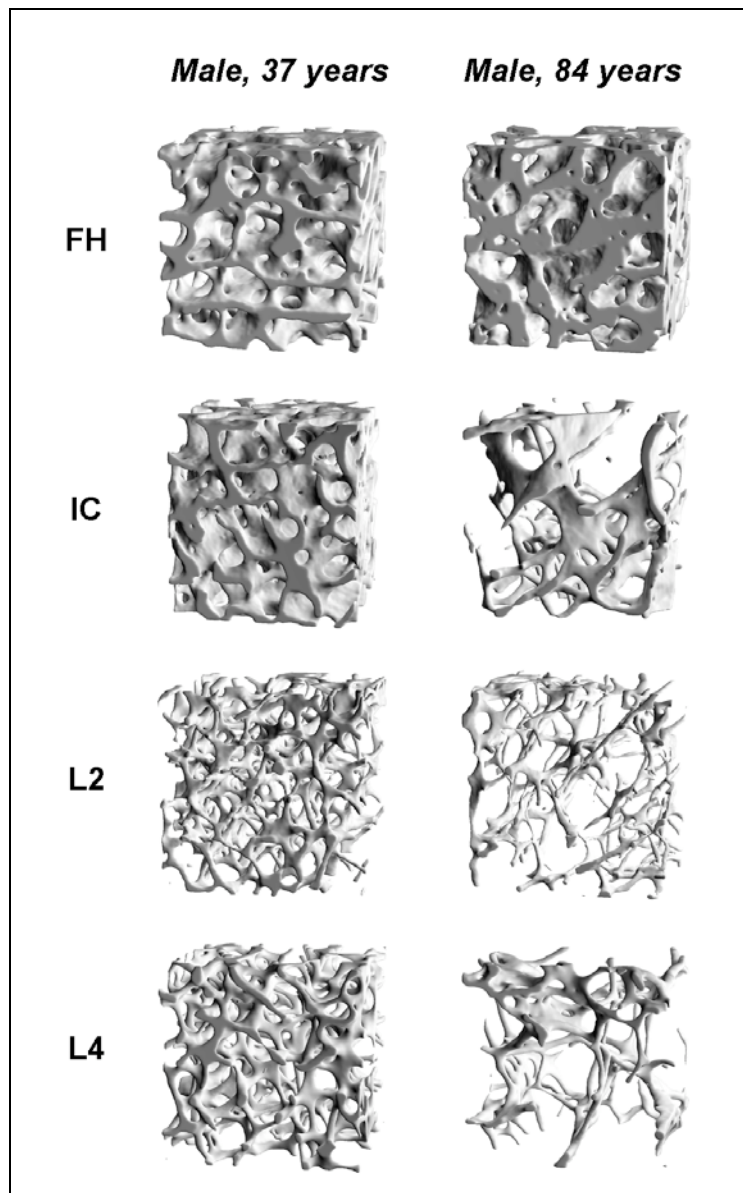
Computed tomography is an approach used to image and quantify trabecular bone in 3D, providing multi-scale biological imaging capabilities with isotropic resolutions ranging from a few millimeter (clinical CT), to few tens of micrometers ( $\mu$ CT) down to one hundred nanometer (synchrotron radiation nanoCT). An overview of the principles and methodologies is beyond the scope of this paper and can be found elsewhere (12;13).  $\mu$ CT has become the method of choice in quantifying 3D bone microstructure. Commercially available desktop  $\mu$ CT systems provide nominal resolutions ranging from roughly 1 to 100  $\mu$ m, allowing investigators to measure specimens with diameters ranging from a few millimeters to 100 mm. Desktop  $\mu$ CT is a precise and validated technique (14-16) that has been used extensively to study bone microarchitecture (16-19) and has demonstrated that trabecular bone can have very distinct microarchitectures that are remarkably heterogeneous throughout the skeleton (20-22) (Fig. 1).

### Quantification of Bone Micro-Architecture

Quantitative endpoints have become an important factor for success in basic research and the development of novel therapeutic strategies in biology and biomedicine. One method of quantitatively describing bone architecture and the changes associated with age or stage of a disease is the calculation of morphometric indices, also referred to as quantitative bone morphometry. In the past, structural properties of trabecular bone have been investigated by the examination of 2D sections of bone biopsies. 3D morphometric parameters were then derived from 2D images using stereological methods (23).

In order to take full advantage of the volumetric data as obtained from  $\mu$ CT measurements, several new 3D image processing methods have been presented, allowing direct quantification of bone microarchitecture (21;24). These techniques calculate actual distances in 3D space and, therefore, do not rely on an assumed model type and are not biased by possible deviations.

More recently, local morphometry was introduced to measure morphometric indices of single trabecular elements, such as individual plates and rods. For this purpose, the elements have to be extracted from the structure, which can be done using a method called volumetric spatial decomposition (25). With this method it is hence possible to perform more detailed analyses of three-dimensional structures and to aid in a further understanding of the individual contribution of rods and plates to the competence of trabecular bone (26;27). As an example, in a study on human vertebral bone samples, a multiple linear regression model combining mean trabecular spacing (Tb.Sp), mean slenderness of the rods ( $\langle$ Ro.SI $\rangle$ ), and the relative amount of rod volume to total bone volume (Ro.BV/BV) was able to explain 90% of the variance in bone stiffness. This model could not be improved by adding BV/TV as an independent variable (26).

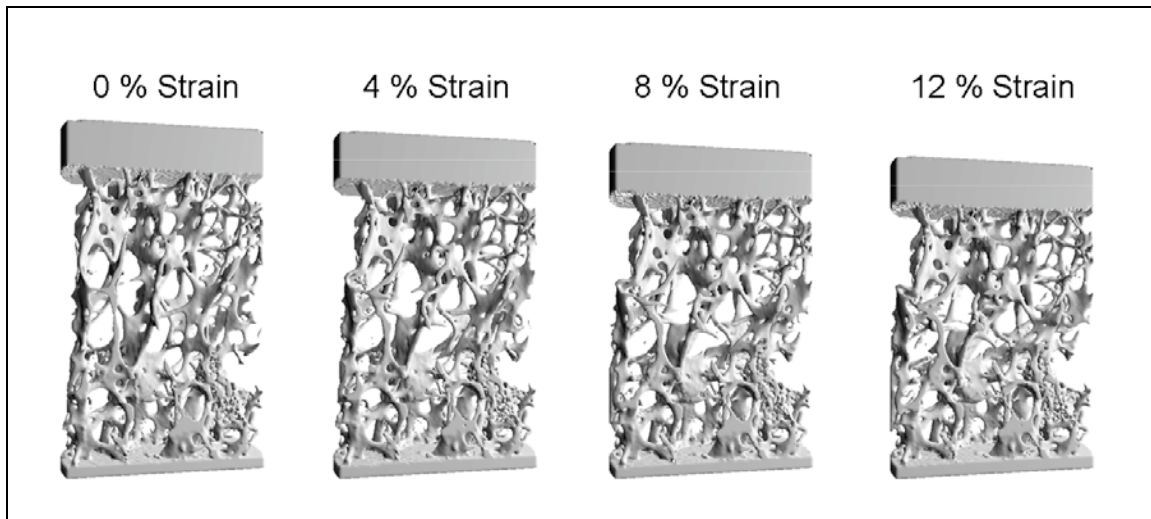


**Fig. 1.** Bone biopsies of a 37-year-old and an 84-year-old male taken at the femoral head (FH), iliac crest (IC), and lumbar spine (L2 and L4). Note the strong heterogeneity in bone architecture, not only between individuals but also between anatomical sites.

### Visualization of Bone Under Load

Basic knowledge of local trabecular failure is still lacking. In estimating the risk of spontaneous fractures, an extended understanding of the failure behavior of trabecular bone is essential. By combining experimental testing with  $\mu$ CT it has become possible to visualize and quantify bone deformations under load as well as fracture initiation and progression in a non-destructive and 3D fashion (28;29). Several

deformation modes were found in human vertebral bone; some trabeculae bend, others buckle, and some are compressed (30) (Fig. 2 and [Suppl. Material 1](#)). It is important to note that failure behavior might be quite different for bone in an *in vivo* environment where boundary conditions differ significantly from *in vitro* setups. For the purpose of extrapolating the findings on the material level to the *in vivo* situation in whole bones, we recently built a larger image-guided failure assessment (IGFA)



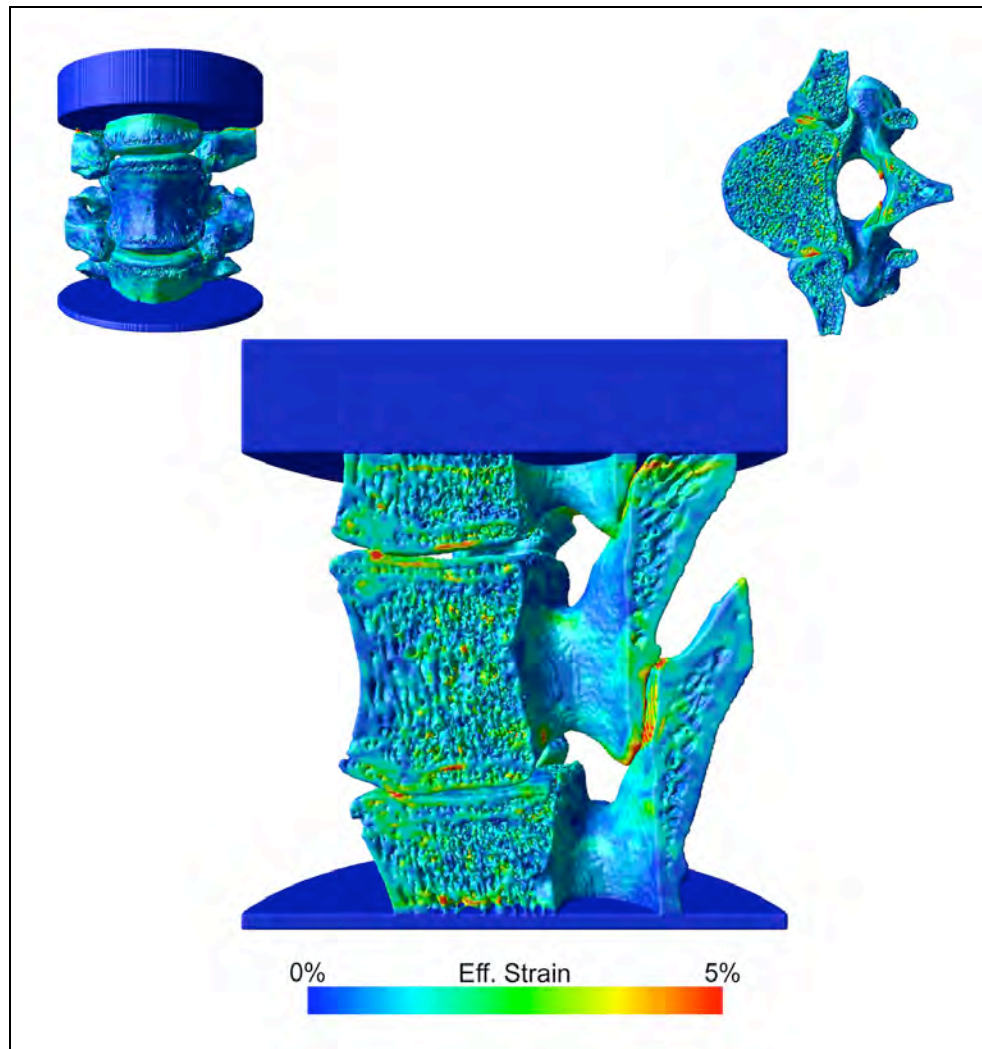
**Fig. 2.** Image-guided failure assessment in a human spine sample using time-lapsed tomographic imaging. The images show compression, imaged in steps of 4% strain. Note that only half of the specimen is depicted here, which is possible due to the non-invasive character of the IGFA technique (images courtesy of Dr. Martin Stauber, ETH Zürich, Switzerland).

device that can fit bones up to 10 cm in diameter. A pilot study on an ovine spinal unit clearly demonstrated its functionality in determining fracture location and progression under load ([Suppl. Material 2](#)). This technique of IGFA has been validated as compared to classical continuous mechanical testing (29;30). The deformation behavior has been quantified by determining the strains between local anatomical features (nodes) in the structure that were followed directly in the structure throughout all the steps (31). Although inter-node strains can identify loaded elements, they cannot determine the precise deformation of these elements. A better description of local strains is expected from experimental strain mapping approaches (32-34). Strain mapping is a mathematical technique to calculate how a bone or bone sample has deformed between two loading steps. For that purpose, the displacement field is calculated that maps one loading condition (e.g., an undeformed object) to another loading condition (the deformed object). Based on the displacement field, the local strains can then be easily determined. As an example, deformable registration was recently implemented (34) to calculate the displacement field that mapped two images of an ovine spinal unit under compression (Fig. 3; the same experiment as illustrated in [Suppl. Material 2](#)). Based on the Gaussian

filtered displacement field, the Green-Lagrange strain tensor was calculated.

### **Computational Assessment of Bone Competence**

Although the novel experimental IGFA approaches allow new and valuable insight into the bone failure process, their application is limited to *in vitro* investigations due to their destructive nature. Computational failure prediction would therefore be a valuable tool in combination with noninvasive *in vivo* imaging to assess bone competence in living subjects. Finite element (FE) analysis is the most widely used computational technique for structural analysis in engineering. It can provide insight into the load transfer through the bone architecture and with that improves our understanding of how differences in bone microarchitecture influence bone strength. Microstructural FE ( $\mu$ FE) models allow researchers to perform 'virtual experiments' on computer reconstructions of trabecular bone. These models typically represent small trabecular regions in the order of 5 to 10 mm cubes, a scale at which the bone behaves as a continuum (35). As an example of such an analysis, the principal strains in a human bone biopsy subject to a compressive load are shown (Fig. 4). Loading and deformation for each



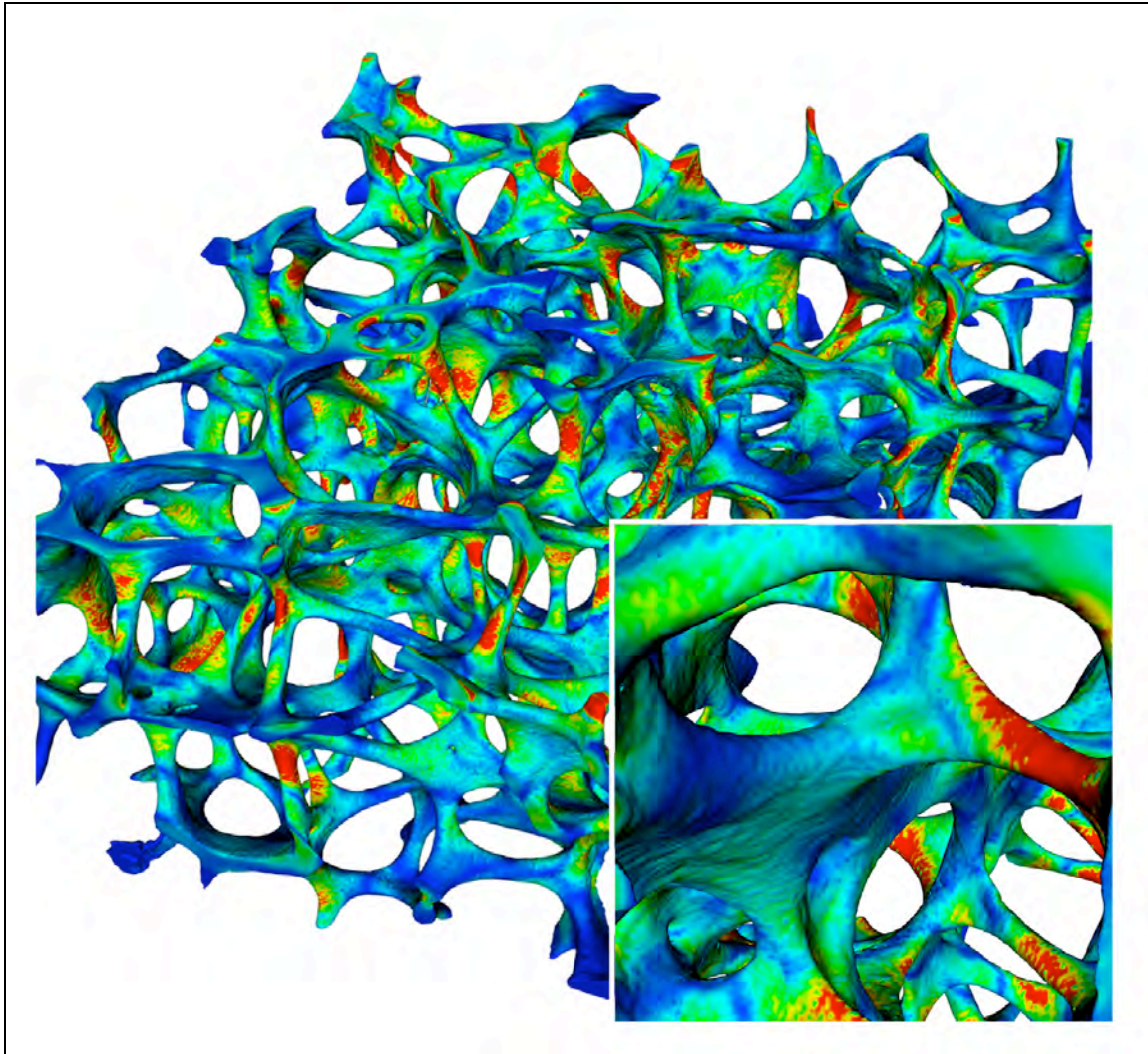
**Fig. 3.** Effective strains in an intact ovine spinal unit as determined from strain mapping based on two time-lapsed tomographic images (images courtesy of D. Christen, ETH Zürich, Switzerland).

individual trabecula can be determined and explored with high detail. Animation of the predicted deformations and associated stresses can help us to better understand the local failure behavior of the trabecular bone network ([Suppl. Material 3](#)).

For linear deformation conditions, comparison between biomechanical compression tests on excised trabecular bone samples and  $\mu$ FE show very good agreement ( $R^2=0.92$ ) when a homogeneous, isotropic tissue modulus is applied (36;37); this holds true for normal as well as osteoporotic bone (38). Also for non-linear deformation conditions, good agreement was found between  $\mu$ FE results and experimental tests. It has been shown that

at the apparent level, these  $\mu$ FE models can accurately predict trabecular bone failure for bovine (39) as well as for human trabecular bone (40); the  $\mu$ FE-predicted apparent stresses and strains at failure were equal to experimentally measured values for the same bone specimens. Whether the good agreement also holds at the level of single trabeculae and hence whether the precise location of local bone failure can be determined from  $\mu$ FE has yet to be determined. Nevertheless, the clinically most interesting parameter, apparent bone strength, can be accurately assessed, demonstrating that the quality of  $\mu$ FE analyses has reached such accuracy that they can be used as an alternative to destructive mechanical tests (39).





**Fig. 4.** Effective strains in a human iliac crest biopsy, subject to a simulated axial load, and calculated by  $\mu$ FE analysis. Blue: low strain; red: high strain. The inset shows part of the model in higher magnification (images courtesy of Andreas J. Wirth, ETH Zürich, Switzerland).

The steady increase in computational power over the last several years, using multiple processors in supercomputers, has made it possible to analyze intact human bones, or large portions thereof. Highly-detailed  $\mu$ FE models, with resolution of 60-80  $\mu$ m, of the proximal human femora (41) and of human vertebra (42;43) have been presented, with up to 100 million elements. A recent implementation even allowed the solution of a model with nearly 1.5 billion degrees of freedom within 30 minutes, to simultaneously increase the level of detail and the size of the analyzed bone volume (44;45). These models have addressed aspects of osteoporosis on the stress

distribution in trabecular bone, and assessed the importance of the thin cortical shell in human vertebrae. Recent  $\mu$ FE analyses (element size 82  $\mu$ m) on 9.1 mm thick sections at the distal radius show excellent reproducibility in stiffness and strength, with precision errors better than 1.2% CV precision for a standard analysis region (46). These models also show excellent accuracy. When using a homogeneous tissue modulus, the apparent stiffness as calculated by linear  $\mu$ FE correlated highly ( $R^2=0.972$ ) with experimentally assessed bone strength; the correlation could even be improved ( $R^2=0.983$ ) when moduli were determined

from scaled CT-attenuation values (47). Hence, these models allow the assessment of mechanical bone properties fully non-destructively and with that the prediction of bone strength in longitudinal *in vivo* studies in a time-lapsed fashion.

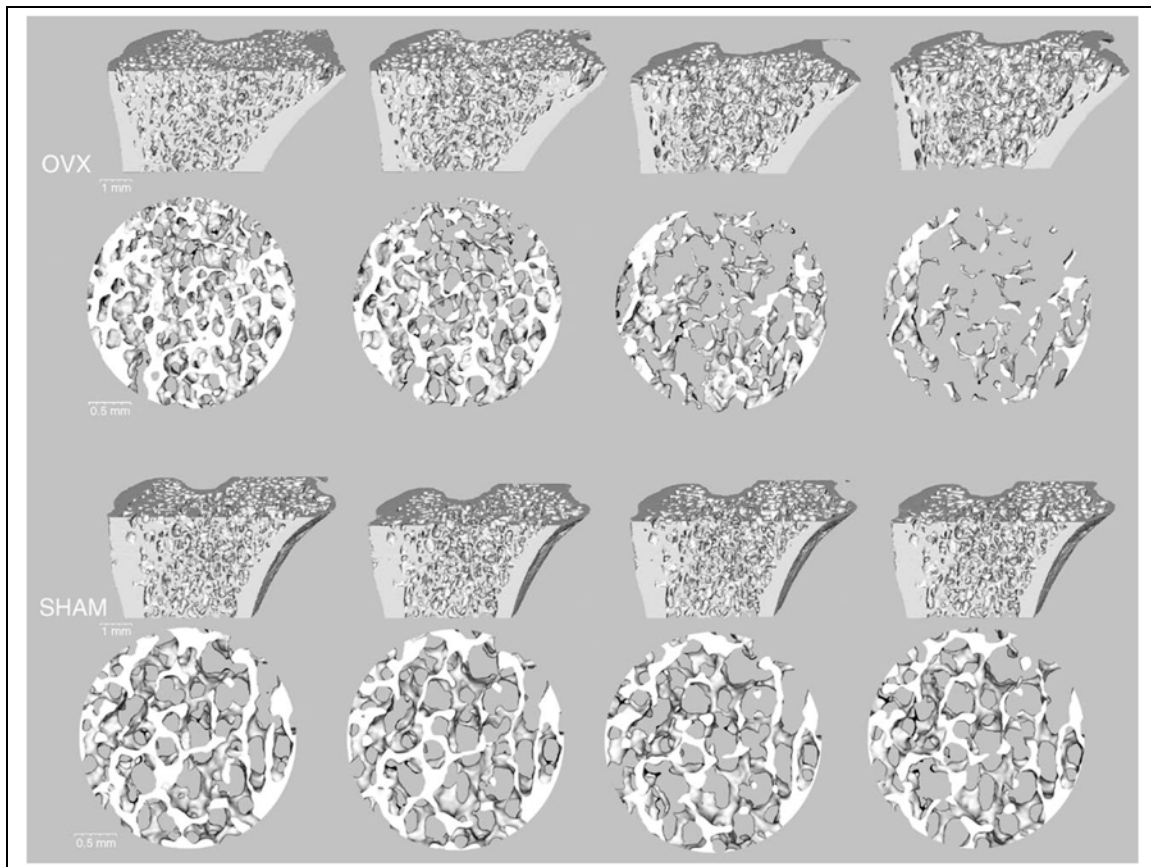
### Recent Developments in Small Animal *In Vivo* $\mu$ CT

The feasibility of performing *in vivo*  $\mu$ CT was first demonstrated in the rat proximal tibia using synchrotron radiation (48), and was used to depict bone loss after OVX (49). The recent introduction of desktop  $\mu$ CT systems, for *in vivo* scanning of small animals (50;51), with resolution better than 10  $\mu$ m has opened the way to more practical and better accessible systems. A great feature of *in vivo* scanning is that it allows for longitudinal scanning of the same animal. Hence, a time series of several images of the same bone can be obtained, which allows for the visualization and quantification of changes in bone geometry as well as changes to individual trabeculae (Fig. 5). In a way, this is similar to the time series as obtained with IGFA, but there are some important differences.

First, the time steps are in the order of weeks, such that biological adaptations causing microstructural changes can be followed over time. Hence, growth, aging, response to interventions and response to treatment can be closely monitored. Visual examinations of image-registered data show compelling evidence that small changes can be detected in the shape and orientation of single trabeculae and even parts thereof (51). However, the proper metrics to quantify these changes have yet to be introduced. Furthermore, reproducibility data for *in vivo* scanning of small animals is scarce. One study showed small errors of less than 3% in bone volume measurements and errors less than 0.5% in measurements of trabecular thickness (51). These values may differ for other types of scanners, and may also depend on the animal under investigation, its genetic background, its age, and on the specific bone location. In principle, *in vivo* reproducibility may reach that of *in vitro* studies, where excellent reproducibility has been reported for cortical (precision errors

ranging from 0.06–2.16% CV) and trabecular (precision errors ranging from 0.59–5.24%) bone parameters, respectively (52).

However – and this is the second difference – radiation is a concern in *in vivo* measurement because ionizing radiation can cause damage to bone cells and impair their function and viability. Hence, a measurement can interfere with the normal bone response. Of course, this is an unwanted side effect that may limit use of *in vivo* scanning, particularly when the detrimental effects would be large compared to the effects one would like to measure, or when the architectural features that one wants to resolve are small. The latter would require a small voxel size and hence a high radiation dose, considering that a reduction in voxel size with a factor of two requires an eight-fold increase in X-ray dose to obtain the same signal-to-noise ratio. Hence, irradiation effects have received quite some attention so as to obtain the best resolution-for-dose; nevertheless, the precise effects of current *in vivo*  $\mu$ CT devices are still subject to some debate. It is clear that the total radiation dose depends on  $\mu$ CT design, scanning protocol, and the number of repeat measurements; it also seems obvious that the time interval between measurements can play a role. Furthermore, potential effects on bone structure may depend on the animals under investigation (rats or mice), be genetically influenced, and be affected by the age of the animals. In rats, Klinck *et al.* and Brouwers *et al.* detected no effects on bone structure (53;54) nor on cell viability (53) even after eight weekly scans. However, Waarsing *et al.* noticed a small effect in rats that were scanned less often (55). This could be related to several issues such as the longer follow-up, the use of a different scanning system with different dosage or the age of the animals. In mice, multiple measurements do seem to have a small effect on trabecular bone volume (~10-20% less bone relative to non-irradiated limbs) (54); however, that same study also showed that effects of OVX were much larger and that there was no interaction between radiation and OVX treatment. In any case, more thorough studies seem warranted to investigate the optimal imaging



**Fig. 5.** Three-dimensional images of the tibial metaphysis of an OVX and sham-operated animal (median from each group) at the time of operation and follow-up measurements 1, 2 and 3 months later. The image data are transected to illustrate the internal trabecular structure, and all images were aligned using 3D image registration. Details from each time point (0.3 mm slice) are illustrated in circular cutouts (reprinted from Boyd *et al. Bone*. 2006 Oct;39(4):854-62, with permission from Elsevier).

settings providing the best resolution-for-dose and the maximal number of repetitions for a given time interval.

A third difference is that *in vivo* scanning is technically more demanding: more effort is needed to match two consecutive scans. The principle of this process, called image registration, is to reorient a  $\mu$ CT data set such that it aligns with the data set of an earlier measurement. The differences after registration identify the locations where bone has been resorbed, and where it has been added. When changes are small, this is a relatively straightforward approach; however, it becomes a non-trivial task when architectural changes are large (51). In that case, *a priori* knowledge, *e.g.*, a fixed position of an arterial opening in the cortex, can help to identify the correct reorientation. Also, in the case of longitudinal growth, one

would typically register the epiphyseal and metaphyseal bone separately to avoid mismatches due to growth by apposition at the growth plate (55). Hence, zooming in on a specific region might give a different matching than when applied on an entire bone, which implies that it is not evident how to determine the precise locations of bone loss and bone gain. Clearly, more research is needed investigating the best strategies for regional analysis in longitudinal studies where bone size and structure are changing over time in health, disease and treatment, especially in growing children or fracture healing, where large changes can be expected in a relatively short time frame.

From a methodological point of view, a great advantage of using time-lapsed *in vivo* scanning is that each animal can act as its own control, thereby increasing statistical



power. Small changes in bone structure may become detectable that would otherwise go unnoticed because of normal population variations. This was nicely demonstrated in a study on morphological changes after OVX in rats (56); the *in vivo* analysis found significant changes at an earlier time point in OVX rats and detected small changes in sham-operated rats that were not detected by the cross-sectional design. Increased statistical power also means that fewer animals are needed to detect a certain effect. Furthermore, there is no need to sacrifice animals at intermediate time points; hence, this will lead to a further reduction in the number of animals, with the reduction factor identical to the number of measurements.

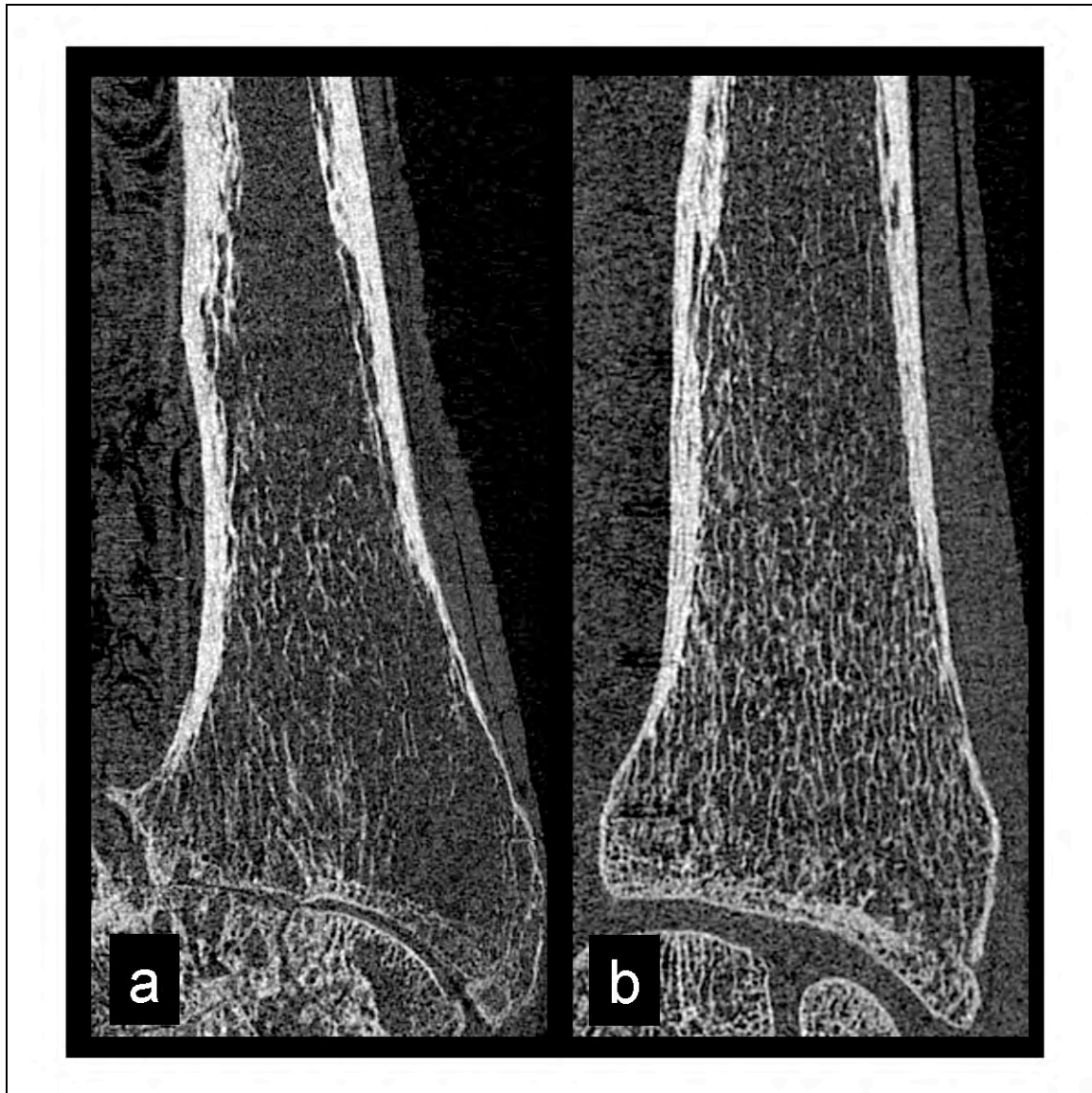
Several reports have been presented on different aspects of microarchitectural changes. Using *in vivo*  $\mu$ CT in rats, Waarsing *et al.* observed that, due to normal aging, the combination of bone loss and bone formation resulted in alignment of trabeculae across the growth plate; this process was even more pronounced in aging OVX rats (55). Brouwers *et al.* determined the effects of the time point of bisphosphonate treatment after OVX in rats (57). The rapid bone loss and structural degradation as seen after OVX could be completely inhibited by a single injection of zoledronic acid at the time of OVX, whereas an injection eight weeks after OVX improved bone mass and structure but no full recovery was seen. The effects of genetic background on OVX-induced bone loss in mice were recently assessed by Klinck and Boyd (58). They quantified the weekly loss of bone in an animal-specific manner and found that inbred strains with higher baseline bone density values showed higher rates of bone loss.

### ***In Vivo* Scanning of Human Bone Structure**

Paired iliac crest bone biopsies have been and still are being used to determine response to treatment (19;59). It is clear that these are far from ideal, first, because the procedure requires taking two invasive biopsies with associated problems, second, because the biopsies are necessarily small,

and third, because in fact two different trabecular architectures are being evaluated. The drawbacks of paired biopsies could be avoided with the introduction of *in vivo* imaging systems with a resolution better than 50  $\mu$ m. Unfortunately, such systems are not yet available, but the resolution at peripheral sites is approaching a level that allows elucidation of individual microstructural bone elements. Such high-resolution peripheral quantitative CT systems (HR-pQCT) have reached the market, allowing for *in vivo* patient measurements with an isotropic nominal resolution of 82  $\mu$ m (Fig. 6). Given the larger voxel size, trabeculae may be reconstructed with a thickness of just one or two voxels. This necessitates the use of the so-called plate model to derive trabecular thickness (Tb.Th) and trabecular separation (Tb.Sp) from bone volume density and trabecular number (Tb.N), rather than using the direct 3D approach as implemented for higher resolution  $\mu$ CT. Nevertheless, the HR-pQCT morphometric measures correlate well to those of  $\mu$ CT (60). Furthermore, they provide excellent short-term and long-term reproducibility. At the distal radius, *in vivo* reproducibility of density was better than 0.59%, whereas for morphological measures this was better than <3.9% (61). Nevertheless, as with every CT, these systems require X-rays; hence, radiation dose, how little it may be, will always be a concern. With respect to these dose considerations, novel low-dose CT procedures and high-resolution magnetic resonance imaging (MRI) may have considerable potential for future clinical applications to overcome some of the limitations with the presented X-ray-based modalities (62;63).

Using HR-pQCT, three recent cross-sectional studies have shown sex- and age-related changes in trabecular structure and in cortical thickness of the distal radius *in vivo* (64-66). The acquired structural data were able to differentiate between osteopenic women with and without history of fracture, whereas measurements of BMD did not (64). Morphometric sex-specific differences in bone microstructure were found, potentially explaining the virtual absence of an age-related rise in distal

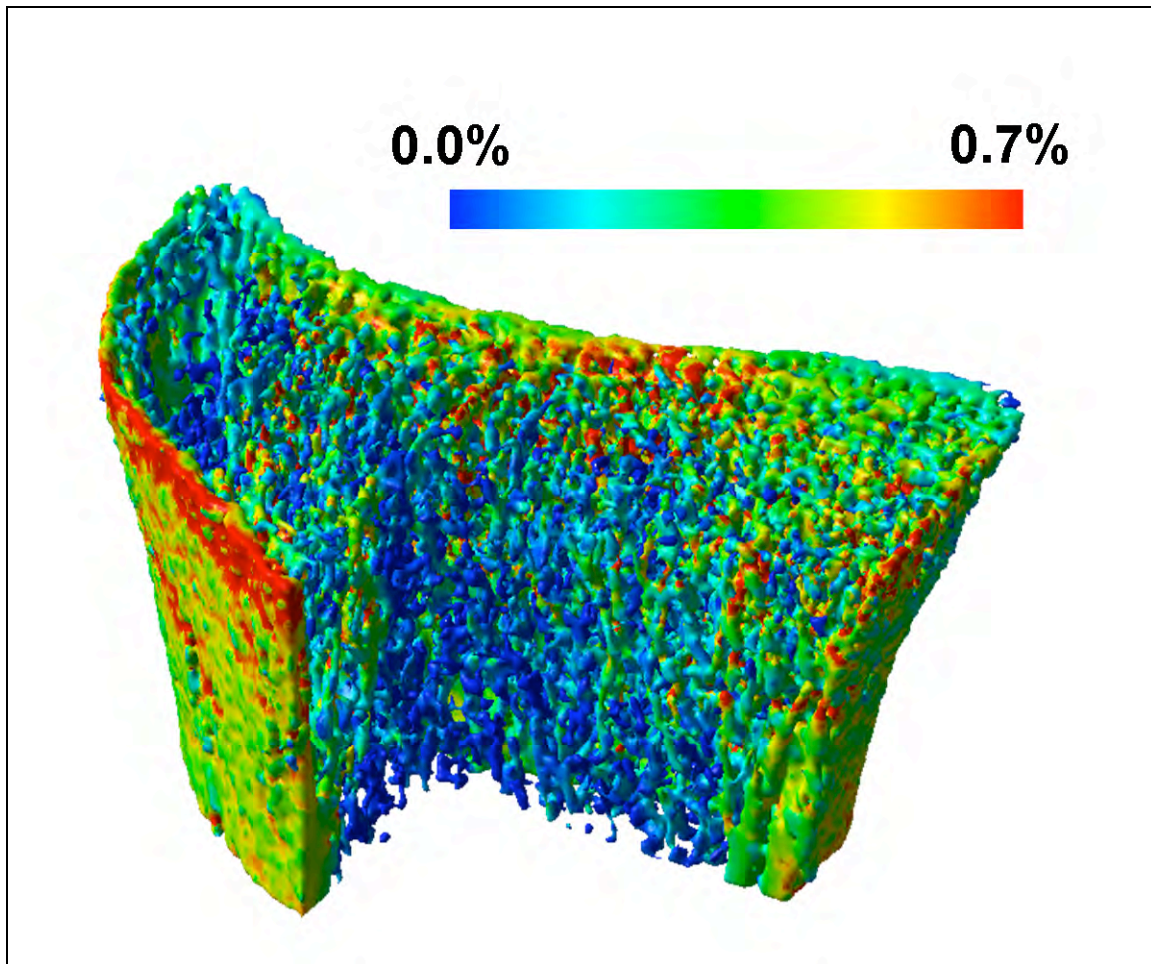


**Fig. 6.** High-resolution peripheral quantitative computed tomography (HR-pQCT) systems allow for the non-invasive measurement of human bone with resolution better than 100  $\mu\text{m}$ . The figure shows HR-pQCT images, as measured *in vitro*, of the distal radius of a 79-year-old male (a) and a 92-year-old male (b), respectively (images courtesy of Thomas L. Mueller, ETH Zürich, Switzerland).

forearm fractures in men compared to the marked increase in these fractures in older women (66). In addition, relationships between hormonal and bone turnover variables and trabecular microstructure at the distal radius were demonstrated (65).

In combination with  $\mu\text{FE}$  analyses, these HR-pQCT data allow for the determination of bone competence *in vivo*. Ulrich *et al.* created  $\mu\text{FE}$  models to analyze an *in vivo* microarchitectural model of the distal radius, measured with an earlier pQCT providing

lower resolution (67). This was the first time that tissue level stresses and strains were estimated in a living human subject. Pistoia *et al.* showed that there was good agreement between the predicted failure loads calculated from  $\mu\text{FE}$  and those measured using mechanical testing in a large sample of human cadaveric forearms. In addition, the  $\mu\text{FE}$ -derived strength parameters did substantially better in predicting experimental bone strength than classical bone densitometry (68). These data were confirmed for the newer version of



**Fig. 7.** The effective strain distribution in a 1 cm slab of bone from the distal radius as calculated from  $\mu$ FE analysis. The model was built from an *in vivo* high-resolution peripheral quantitative computed tomography (HR-pQCT) measurement; part of the bone was digitally removed to show the trabecular core.

the system that provides a nominal resolution of 82  $\mu$ m (47). The potential of HR-pQCT-based  $\mu$ FE (Fig. 7) to identify people at risk of distal radius fracture has been demonstrated (69;70), indicating that it could become a clinical tool soon.

### Conclusions

During the last decade microarchitectural bone imaging based on high-resolution CT has become the method of choice for quantifying 3D bone microstructure *in vitro*. It is a nondestructive, noninvasive, and precise procedure that allows the measurement of trabecular and compact bone, as well as the computation of microstructural and micromechanical properties. It has demonstrated its effectiveness in evaluating many

applications such as the characterization of genetic effects in different mouse models and in quantifying microarchitectural changes resulting from diseases such as osteoporosis, the effects of hormones and medications such as estrogen and bisphosphonates, and effects of bone adaptation, such as those due to mechanical loading. Recent studies have shown the great potential of *in vivo*  $\mu$ CT for a better understanding of structural changes in bone geometry and its microstructure. These structural changes can be quantified using morphometric measures derived from the 3D data sets, and the effects on bone competence can be estimated from FE analyses. The ability to noninvasively track changes over time is likely to establish *in vivo*  $\mu$ CT as a standard methodology in the coming decade. The procedure might help

improve predictions of fracture risk, clarify the pathophysiology of skeletal diseases, and define the response to therapy. We expect these findings to improve our understanding of the influence of densitometric, morphological but also loading factors in the etiology of traumatic and spontaneous fractures of the forearm, the hip and the spine. Eventually, this improved understanding may lead to more successful approaches in the prevention of such age- and disease-related fractures.

**Conflict of Interest:** Dr. Müller reports that he is a founder and board member of b-cube AG. Dr. van Lenthe reports no conflict of interest.

**Peer Review:** This article has been peer-reviewed.

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