

**EXPLORING THE UTILITY OF SEVERAL EVALUATION METHODS
IN DISTINGUISHING CANNON BONES FROM FRACTURE-AFFLICTED
AND SKELETALLY INTACT RACEHORSES**

by
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Dedicated to my parents, Carrie and Bert, for their endless support

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LIST OF ABBREVIATIONS

| Abbreviation | Meaning |
|---------------------|------------------------------------|
| MC3 | Third Metacarpal |
| SSMD | (Proximal Forelimb) Sesamoid |
| LB | Long Bone |
| C | Control |
| RPI | Reference Point Indentation |
| BMSi | Bone Material Strength Index |
| BMD | Bone mineral density |
| BMC | Bone mineral content |
| 1st ID / ID1 | First Indentation Distance |
| TID | Total Indentation Distance |
| avgED | Average Energy Dissipated |
| IDI | Indentation Distance Increase |
| CID | Creep Indentation Distance |
| avgCID | Average Creep Indentation Distance |
| ED | Energy Dissipated |
| US | Unloading Slope |
| avgUS | Average Unloading Slope |
| LS | Loading Slope |
| TDD | Touchdown Distance |

ABSTRACT

Stress fractures are common in the limb bones of human and equine athletes alike. Repetitive skeletal loading can lead to remodeling and the accumulation of microdamage in bone, which only becomes grossly evident during catastrophic fracture of the bone due to the accumulated microdamage. Though various metrics attempting to quantify bone health exist, none have distinguished themselves as early predictors of the susceptibility of bone to fracture. In this exploratory study, we examine the ability of several evaluation methods to distinguish between third metacarpal (MC3) bones from racehorses that have experienced a limb-bone fracture and from those that have not. Third metacarpal bones were harvested from deceased Thoroughbred racehorses and categorized into four groups: MC3 bones from horses whose cause of death was not related to skeletal fracture (Control group, n = 20), MC3 bones from horses that were euthanized after fracturing proximal sesamoid bones (Sesamoid group, n = 20), MC3 bones from horses that were euthanized after fracturing a non-MC3 long bone (Long Bone group, n = 19), and MC3 bones from horses that were euthanized after fracturing an MC3 (MC3 group, n = 5). Each MC3 bone underwent testing using a variety of tools and methods at the proximal, midshaft, and distal levels of the lateral, dorsal, and medial surfaces. All tools and methods (OsteoProbe reference point indentation, BioDent reference point indentation, x-ray, micro-CT, and pQCT) exhibited some capability in differentiating between control and fracture groups. The long-term objective of this project is to create a model that will utilize data from a set of evaluations and output the susceptibility of the horse to fracture a bone, a long bone, or the MC3, specifically. Although the sample size in this study is not sufficient to create a reliably predictive logistic regression model, promising results from preliminary models provide incentive to further explore the possibility of creating one. While clinical practicality will be a vital consideration for a model in the future, establishing this basis for the capability of each evaluation at hand is a necessary first step in predicting and preventing fracture in bone.

1. INTRODUCTION

1.1 Background

The limbs of racehorses are subjected to a large number of high-magnitude loading events during their early lives. In full gallop, the forelimbs in the average Thoroughbred racehorse can experience compressive forces up to 26 N / kg, or approximately 2,870 lbf [1]. Healthy bones undergo modeling and remodeling during training to better withstand these forces; however, it's estimated that up to 2% of racing starts result in fracture [2], with between 0.3 and 1.7 in 1,000 starts resulting in fatal musculoskeletal injury [3-6]. These musculoskeletal failures contribute to the approximately 1,600 jockey injuries sustained at US racetracks each year, and the often resultant euthanasia of the horse sheds negative light on the \$100 billion dollar horseracing industry from the public perspective [69].

Wolff's law asserts that healthy bone adapts to the loading conditions under which it is placed [7]. Though some details and derivations within this axiom have since been updated [8,9], the outlined principles have remained crucial to modern skeletal biomechanics research. Numerous theories have been proposed in an attempt to explain the mechanisms behind these geometric adaptations, including skeletal microdamage stimulating osteonal remodeling [70], piezoelectricity produced by collagen fibers [10], and, perhaps most widely-accepted, shear stress caused by interstitial fluid flowing through the lacunar-canalicular network in bone [11].

The modeling and remodeling processes that allow bones to be better suited to their loading conditions, however, are not instantaneous. A bone undergoing remodeling undergoes the resorption phase in 2 to 4 weeks, while bone deposition occurs more slowly over a span of multiple months [12]. Repetitive loading on an elastic material such as bone leads to material fatigue within the bone—the stresses experienced during high-force, cyclic loading can lead to microdamage in the bone tissue. Over time, this damage can accumulate into macroscopic cracks which may then lead to catastrophic fracture of the material. This fatigue can happen in any bone subjected to cyclical loading, and bone tissue left vulnerable due to the slow timeline inherent to the remodeling process may be even more susceptible to the formation of stress fractures or progression to catastrophic failure [13].

While there are proposed training regimens that promote healthy bone remodeling and reduce the rate of bucked shins or stress fracture by allowing adequate time for microdamage to heal and osteoblasts to deposit new tissue [14,15], the incidence of limb fracture in racehorses remains common. Though current clinical metrics and methods such as bone mineral density and radiography can provide a collection of information about the state of a bone's health, the predictive power in assessing the susceptibility of fracture is still rather low.

In humans, bone mineral density measurements standard for assessing bone health have been shown to correctly identify approximately 10% of fractured bones and 91% of non-fractured bones. A different method known as statistical shape and density modeling has been found to correctly identify 55% of fractured bones and nearly 95% of non-fractured bones [71]. When multiple factors are considered together, the predictive power increases. An assessment tool that utilizes a multivariable logistic regression models called the FRACTURE Index boasts an area under a receiver operating characteristic curve of almost 77%, nearing the 90% or 95% typically sought after in medical diagnostic tools [72].

1.2 Radiography

The relationship between bone mineral content (BMC), bone mineral density (BMD), and physical activity have long been studied [16,17]. In the mid-to-late 1900s, the gold standard in quantifying BMC in human clinical settings was x-ray spectrophotometry [18,19]; since then, however, multiple methods of characterizing the inorganic components of bone have arisen. Well-collimated scintillation detectors were implemented to improve upon the traditional x-ray approach [20], and multiple energies of x-ray were utilized to parse out soft tissue absorption in a method known as dual energy x-ray absorptiometry (DXA) [21]. DXA has since become a staple in diagnosing and monitoring osteoporosis [22], assessing the effects of drugs on the bones of postmenopausal women [23], or even quantifying a patient's visceral fat to predict his or her susceptibility to diabetes or heart disease [24]. Peripheral quantitative computed tomography (pQCT) measures volumetric bone mineral density (vBMD) rather than areal bone mineral density (aBMD) as in DXA. While vBMD is able to better adjust for different bone sizes (such as those in children) and can also provide geometric information about the bone that DXA cannot [25], its design inherently limits it to use in the appendicular skeleton, which has been shown to be a poor predictor of mineral density in sites of common fracture such as the proximal femur or spine [26].

Sound-based methods such as quantifying broadband ultrasonic attenuation and ultrasonic velocity have been developed, with apparent increasing effectiveness over time, as alternatives to methods dependent on radiation [27-29].

Various studies have attempted to quantify bone mineral parameters to assess bone health in racehorses. Horses that had undergone training were found to have significantly higher distal epiphyseal subchondral sagittal groove vBMD values than their untrained counterparts [30,31]. Additionally, trabecular BMD has been shown to significantly correlate with whole-bone breaking strength in the proximal phalanx [32]. BMD in bones from horses with or without fracture, however, show conflicting results. Some studies that examined the third metacarpal and proximal phalanx have reported no significant differences in BMD between control and fracture groups [31,33], while others have found that both the BMD and stiffness are significantly higher in bones from fracture groups than from control groups [34].

In human medicine, radiography is a staple of the diagnostic imaging field. X-ray imaging is used to locate and diagnose fractures, examine lung health, and even find cavities in teeth. In equine research, x-rays are often utilized as a noninvasive way to measure geometric properties of bone, particularly those of cortical bone. Finding bone length [35], cortical bone thickness [36,37], location and severity of fracture [38,39], or even history of fracture [40] are common uses of x-rays. Studies performed using x-rays have characterized geometric values in the appendicular bones of healthy thoroughbred racehorses [41]. These radiographic studies have also found correlations between exercise speed and cortical bone modeling [36] and have explored the effects of exposing bones to exogenous growth hormones [42]. Radiography is excellent at detecting fracture, osteoarthritis, and other visually-discernable maladies. However, radiology falls short of technologies such as magnetic resonance imaging to detect early stages of osseous disease or dysfunction [43], and limitations such as superimposition can inhibit accurate anatomic imaging or density measurements.

1.3 Reference Point Indentation

Reference point indentation (RPI) is an emerging technology that creates microindentations on the surface of a sample to gather information about its material properties. Determining the hardness of a material via indentation has been used for decades [44], often in metals and other engineered materials. Indentation techniques come in a variety of forms, from spherical to conical

& pyramidal indenters [45,46] and even nanometer-scale indentations [47]. While various methods for gathering information about bone using indentation have been proposed or performed [48-50], many would be difficult to utilize *in vivo*. Nanoindentation testing requires extremely precise contact angles and microscopic evaluation, both of which may be prohibitively difficult in a clinical setting and would likely require biopsies to be collected [51]. Many researchers have lately turned to one of two microindentation systems produced by Active Life Scientific, Inc. that have a greater potential clinical relevance than previously-used nanoindentation techniques. The OsteoProbe, a handheld single-impact device that has recently been approved for clinical use in Europe [52], returns a single parameter: bone material strength index (BMSi). Studies utilizing the OsteoProbe have elucidated differences between the bones of postmenopausal women with type 2 diabetes versus those without [53,54], have found correlations between BMSi in patients with a history of fragility fracture and those without [55,56], and have even discovered a relationship between low BMSi and chronic kidney disease [57]. The BioDent, a benchtop cyclic-RPI system, measures a number of parameters related to a bone's ability to resist microfracture than BMSi (Table 1.1).

Table 1.1. The BioDent system calculates parameters based on distance, stiffness, and plasticity. Listed below are each of the parameters and what a low outputted value for each indicates about the material properties of the sample [58].

| Parameter | ↓ value means... |
|--|--|
| 1 st -cycle Indentation Distance (1 st ID) | Hard, dense, highly-mineralized |
| Total Indentation Distance (TID) | Resistant to crack propagation; tough*; high bone quality |
| Indentation Distance Increase (IDI) | Resistant to fracture; low brittleness of bone |
| Creep Indentation Distance (CID) | Tissue has low viscoelasticity, high damage susceptibility |
| Unloading Slope (US) | Low elastic modulus |
| Loading Slope (LS) | Low resistance to plastic deformation; not stiff |
| Average Energy Dissipated (ED) | Material resistant to plastic deformation |

* Note: conflicting conclusions have been drawn on the degree of correlation between TID and material toughness

Of the multiple BioDent parameters, IDI is considered to best correlate with whole-bone mechanical behavior [3]. Studies have found that IDI and TID were significantly decreased in tibiae from human patients that had experienced osteoporotic femoral fractures compared to control patients [59,68]. In horses, IDI has been found to be associated with training and fracture

history. In a study in which the medial condyles of third metacarpal bones from 31 Thoroughbred racehorses were examined, IDI was found to be higher in untrained horses compared to horses undergoing race training, and higher in horses that had died as a result of a musculoskeletal injury compared to those with other causes of death [60]. These results suggest an increased resistance to indentation on the articular surface in horses that had undergone training and in those that were skeletally intact at the time of death, potentially indicating successful bone adaptation as a response to repetitive loading in accordance with Wolff's law in these groups.

1.4 Multivariable Regression Models

Many regression analyses take multiple explanatory or response variables into consideration. While one factor may correlate well with an outcome, the statistical model may be improved by introducing more explanatory variables, especially when the explanatory variables themselves do not correlate with one another [61]. One study in human cadavers compared fracture strength of the femoral neck to other clinical measurements such as areal bone mineral density, cortical porosity, RPI, and advanced glycation end-products. Each of these parameters alone proved to correlate well with fracture strength; however, when the same data were analyzed using a multiple linear regression model, it was found that combinations of BMD with any other parameter resulted in a higher correlation to fracture strength than any one variable alone [2]. Another study examining risk factors for proximal sesamoid fractures in Thoroughbred racehorses utilized a multivariable logistic regression to predict fracture risk based on a horse's sex, number and type of workouts, and distance run prior to death. It was discovered that fracture risk was higher in sexually-intact males than females and in horses that had run greater cumulative distances prior to their deaths [62].

1.5 Study Aims and Hypotheses

The primary purpose of this study was to explore and compare against one another multiple clinical and preclinical tools used in imaging or otherwise measuring bones. We first examined whether or not any tools or methods could be used to distinguish between third metacarpal bones from horses that have experienced a skeletal fracture and those that have not. Moving forward, we aimed to see if this distinction could be detected using only clinically relevant methods that may

be suitable on a standing horse. These aims were undertaken to provide a basis for the ultimate goal of the project: to select a series of tests whose measurements can be used in a statistical model to compare a sample to known “intact” and “fractured” populations, effectively predicting the sample’s susceptibility to fracture.

Multiple hypotheses have been formulated based on previously published results. Prior studies have found that bone mineral density correlates to whole-bone breaking strength in horse limbs [32], resistance to indentation correlates with training and fracture histories [60], and geometric parameters such as cross-sectional area correlate with training history [30]. Based on these results, we hypothesized that reference point indentation, CT, and x-ray imaging may provide powerful insight into the extent of healthy adaptation to loading and general health of a bone.

While a higher resistance to indentation on the surface of bone may seem to intuitively indicate a strong bone, it was hypothesized that high resistance to indentation may indicate higher susceptibility to fracture. Because the deposition of woven bone on the periosteal surface of bone is an indicator of healthy adaptation to loading, and because woven bone would presumably resist indentation less than lamellar bone, higher indentation distances were predicted to be seen in the control group than in the fracture groups.

More intuitively, greater BMD and geometric parameters such as cross-sectional area or cortical thickness were hypothesized to be associated with bones from horses that had not experienced fracture based on general mechanical principles. As bone is subjected to repetitive compression, apposition and mineralization are natural mechanisms to better support these forces.

Because third metacarpal bones tend to undergo modeling on the dorsal surfaces during training [30], it was hypothesized that this surface would be particularly conclusive in distinguishing between bones that were adapting properly to training and those that were not. Similarly, because distal condylar fractures are common in third metacarpal bones, it was hypothesized that the distal region (75% length or, in pQCT, 90% length) may be of interest when examining bone properties.

Ultimately, it was hypothesized that a number of clinically relevant tests could be used in concomitance to achieve the aim of the study: distinguishing with considerable power between bones from horses with a history of skeletal fracture or those without.

2. METHODS

2.1 Sample Collection and Selection

Thoroughbred racehorses from Indiana racetracks, when euthanized due to skeletal injury or died of causes not related to skeletal fracture, were sent to the Animal Disease Diagnostic Laboratory at Purdue University. Third metacarpal (MC3) bones were harvested during autopsy, wrapped in saline-soaked gauze, and frozen at -20° C. Horses were often transported to Purdue a day after death, resulting in a typical 24 – 30 hour time period between death and freezing of the third metacarpal bones. If a horse was autopsied on the same day on which it died, the third metacarpal bones would be refrigerated for approximately 24 hours prior to freezing to maintain consistency between horses.

Each set of MC3s was classified into one of four fracture groups based upon the reasons for euthanasia: Control (C), from horses that had died of causes not related to skeletal fracture; third metacarpal (MC3), from horses that had been euthanized due to MC3 fracture; Long Bone (LB), from horses that had been euthanized due to a non-MC3 fracture such as a tibial fracture; and Sesamoid (SSMD), from horses that had been euthanized due to a proximal sesamoid fracture. Among these groups, sample sizes for the present study were chosen based upon the data from a previous study and from the availability of bones. For C, LB, and SSMD groups, a sample size of n=20 for each group was selected. A sample size of n=5 was acquired for the MC3 group, as fewer horses with this specific fracture type were available. We had an additional n=5 that could be considered for the MC3 group, but MC3 fracture often coincided with fracture of neighboring bones as well. These samples were excluded from this study to minimize confounding factors within experimental groups.

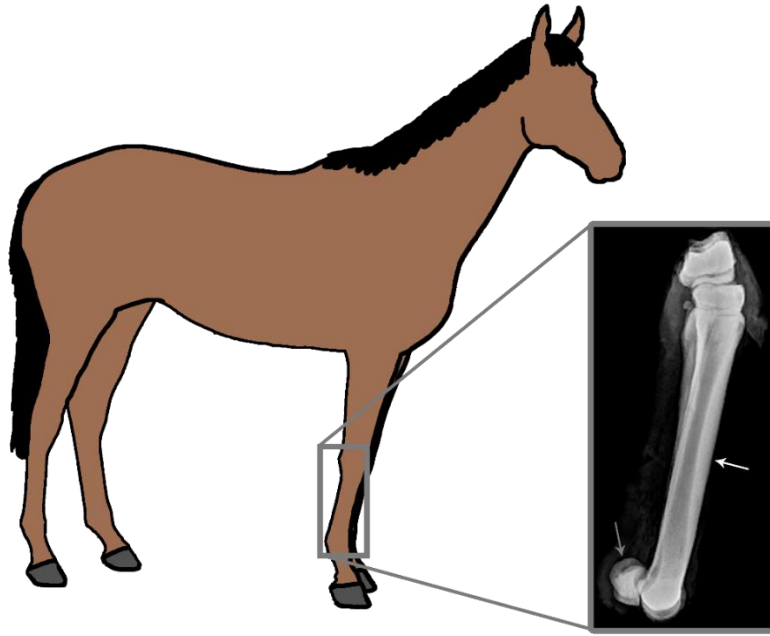


Figure 2.1. Third metacarpal bones (white arrow) were collected from deceased Thoroughbred racehorses. Proximal sesamoids (gray arrow) are a common site of fracture in racehorses.

The experimental groups were age- and sex-matched. When a one-way ANOVA test was performed, neither age ($p = 0.069$) nor mass ($p = 0.329$) returned significant differences. To accommodate for the small sample size of the MC3 group, a separate one-way ANOVA was performed between the C and SSMD groups and a combined LB / MC3 group. This resulted in a significant difference in age ($p = 0.028$), but not in mass ($p = .284$). Though the difference in age was considered statistically significant, the largest discrepancy of average age between groups was approximately 9 months—given the variation within the data and intuitive discretion, it was concluded that age would not be necessary to include as a confounding factor.

During the study, it was discovered that a pair of bones belonging to the LB group were 13% longer than the average MC3 in the study and nearly 2 cm longer than the next longest bone. Although variation is to be expected, these bones were deemed as a likely mislabeled set of third metatarsals and were disqualified from the study (LB group: $n = 19$).

2.2 X-ray

X-ray imaging was utilized to measure the cortical thickness in each bone being studied. Two dimensional digital radiographic images of the third metacarpal bones were taken by technicians in the Diagnostic Imaging Department at the Purdue Veterinary Teaching Hospital

using x-ray equipment by GE. X-rays were taken while the bones were frozen. The bones being x-rayed may have undergone 0 – 2 freeze-thaw cycles prior to imaging, though this is not expected to significantly affect cortical thickness. Two images were taken of each pair of bones: one in the dorsal / palmar view, and one in the medial / lateral view. Prior to imaging, radiopaque “left” and “right” markers were used to differentiate the bones, and a 10-mm scale ball was positioned between or beside the bones to assist in later analysis (Figure 2.2).

Keystone software (Asteris, Inc.) was used to analyze the thickness of cortical bone at 25%, 50%, and 75% of the length of the bone along the dorsal, palmar, lateral, and medial surfaces, where 25% is near the proximal end of the MC3. The software’s “Calibrate” capability was utilized with the 10-mm scale ball, followed by use of the “Length” tool to find locations at 25%, 50%, and 75% of the bone’s length (in the proximal-to-distal direction). Finally, the “Length” tool was used to measure the cortical thickness at each site. Medial and lateral cortical thicknesses were determined using images taken in the dorsal / palmar view, and dorsal and palmar cortical thicknesses were determined using the medial / lateral view.

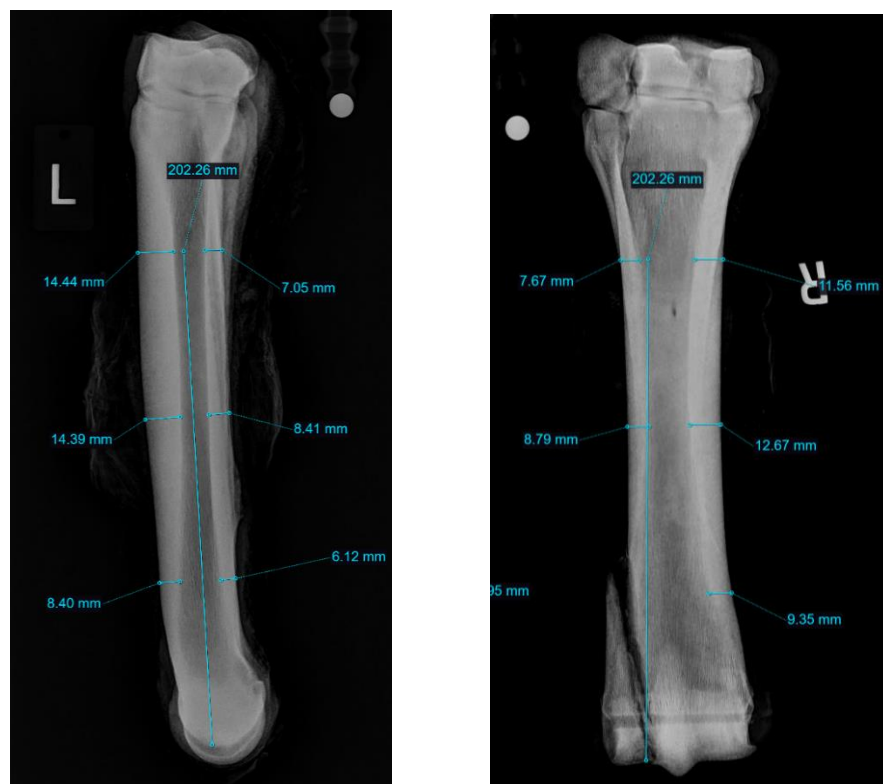


Figure 2.2. Cortical thickness measurements taken where possible at 25%, 50%, and 75% lengths from x-ray images taken from medial-lateral (left) and anterior-posterior (right) views. 10mm scale ball and left/right markers can also be seen. The annotation running down the length of the bone, shown here at the 25% position, was used to visualize where to take cortical thickness measurements.

2.3 OsteoProbe

Impact microindentation to determine the bone material strength index (BMSi) at 12 different sites along each bone was achieved using the OsteoProbe handheld microindentation tool (Active Life Scientific, Inc.).

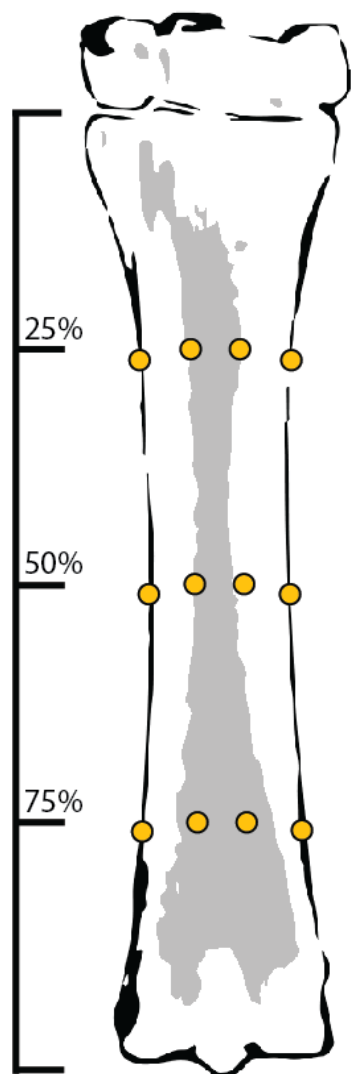


Figure 2.3. OsteoProbe indentations are made at 12 sites on each bone. The dorsal surface is avoided to simulate avoiding the digital extensor tendon *in vivo*.

When bones were harvested with skin still intact, initial OsteoProbe testing was performed through the skin to simulate clinical use. Ten percutaneous indentations, followed by 5 indentations on a block of polymethyl methacrylate (PMMA) for normalization purposes, were performed on the lateral, medial, dorsolateral, and dorsomedial surfaces at 25, 50, and 75% of the length of the bone. Indentations were taken at dorsolateral and dorsomedial surfaces, rather than on the dorsal surface, to avoid the common digital extensor tendon spanning the dorsal surface of the third metacarpal bone (Figure 2.3). OsteoProbe testing was typically performed after 1 freeze-thaw cycle, though the number of cycles at the time of testing for samples in this dataset ranges from 0 to 2.

During testing, the third metacarpal bones were held in place by a bench vice padded with paper towels, with the dorsal surface facing upright while collecting measurements on the dorsolateral and dorsomedial surfaces, the lateral surface upright while collecting measurements on the lateral surface, and the medial surface upright while collecting measurements along the medial surface. Holding the OsteoProbe in one hand, the skin at the site of indentation was held taut during testing by the operator's other hand. Ten 5 N indentations approximately 1 mm apart were made at each testing site, immediately followed by five 5 N indentations on a block of homogeneous PMMA for use in data normalization performed by OsteoProbe software.

After percutaneous OsteoProbe measurements were performed at all 12 sites of a bone, the skin and tendons were removed using a dissecting knife and scalpel. At the planes of 25, 50, and 75% of the bone length, areas approximately 1 inch wide

and spanning the dorsal surface from the lateral to the medial surface were cleared of periosteum and remaining connective tissue using a scalpel and periosteal elevator.

OsteoProbe testing was then repeated on the exposed bone surface at each site on the third metacarpal, following the previous protocol with the exception of the skin being held taut. Sites of indentations made during percutaneous testing were visually located and avoided by at least 1 – 2 mm.

2.4 BioDent

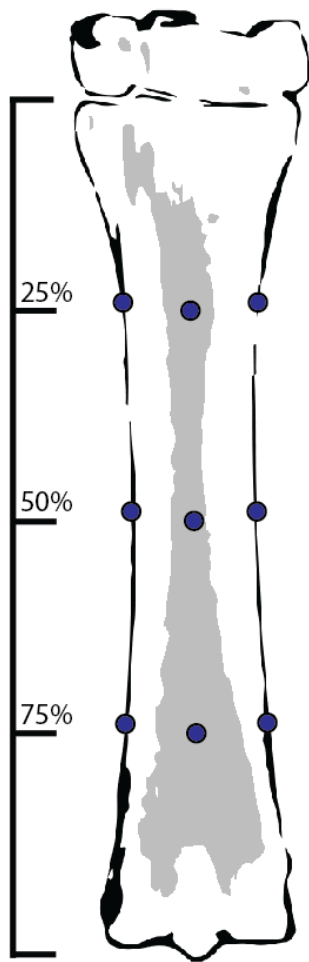


Figure 2.4. BioDent indentations are made at 9 locations on each bone. The dorsal surface is not avoided because the BioDent's benchtop setup is not considered clinically relevant.

Immediately following OsteoProbe measurements taken directly on the bone, cyclic reference point indentation testing was performed using the BioDent benchtop microindentation system (Active Life Scientific, Inc.). Where possible, BioDent testing was performed during the same freeze-thaw cycle as OsteoProbe testing. This was achieved in all but 6 samples in this study. In all bones except anomalous cases in this data set, indentation testing was performed on thawed bones after they had been frozen once. Though it has been shown that the number of freeze-thaw cycles does not have a significant effect on RPI measurements [73], consistency was striven for throughout the study. Testing was performed at the 25, 50, and 75% length sites along the dorsal, medial, and lateral surfaces of the bones. Unlike OsteoProbe testing, the dorsal surface was measured in place of the dorsolateral and dorsomedial surfaces. Because of the benchtop restrictions of the BioDent equipment, percutaneous measurements were not deemed clinically relevant and avoidance of the common digital extensor tendon was therefore not considered.

Prior to testing each bone, measurements on a homogeneous PMMA block were made in “tuning mode” until a Touchdown Distance of 150 – 200 microns was achieved. Bones were placed inside a stainless-steel pan for stability and sanitation purposes during testing, and the distal end of the bone was propped up on wetted paper towels when necessary for the testing surface to be perpendicular to

the indentation probe. When testing the medial and lateral surfaces, bones were propped up using a sand-filled zipper-lock sandwich bag and secured to the metal pan using a C-clamp.

At each site, the “BP2” reference probe was lowered onto the surface of the bone until a force between 1,300 and 1,350 grams was achieved. BP2 probes are described by the manufacturer as semi-sharp probes with blunt ends, and are recommended for testing done on excised bone. The testing protocol was then cycled, with the test probe first initiating 4 pre-load cycles of 1 N at 5 Hz to penetrate any periosteum that may have remained on the bone’s surface. 10 cycles at 40 N at a frequency of 2 Hz were then performed to collect and calculate data with parameters regarding distance, stiffness, and plasticity. Each indentation could be broken into three phases: a loading phase, a holding phase, and an unloading phase. The holding phase described when the probe maintained a constant, maximum force (40 N) for one-third of the measurement cycle (approximately 0.17 seconds). Three (or up to 6, depending on variability of data collected) sets of cycles were performed at each testing location, each at least 1 mm away from prior indentation sites.

As outlined in Table 1.1, seven types of metrics are automatically collected during indentation: 1st cycle indentation distance, total indentation distance, indentation distance increase, creep indentation distance, loading slope, unloading slope, and average energy dissipated. Total indentation distance (TID) and indentation distance increase (IDI) have been used widely in existing literature due to their potential ability to express the overall quality of bone and bone brittleness, respectively. Active Life Scientific also asserted that average energy dissipated (avg ED) may have been closely associated with the formation of microdamage in bone, which made this parameter also particularly relevant to the nature of this study. While parameters aside from TID, IDI, and avg ED were collected and examined, little emphasis was placed on their analysis due to the lack of reported results in previous studies and the consideration of what information they conveyed about the bone tissue.

2.5 Peripheral Quantitative Computed Tomography (pQCT)

pQCT measurements were taken using XCT 3000 equipment produced by Stratec, SE. Data was collected at five planes along the bone: the 25%, 50%, and 75% lengths as used in x-ray and indentation measures, as well as the 10% and 90% lengths to capture the metaphyses of the bone (Figure 2.5).

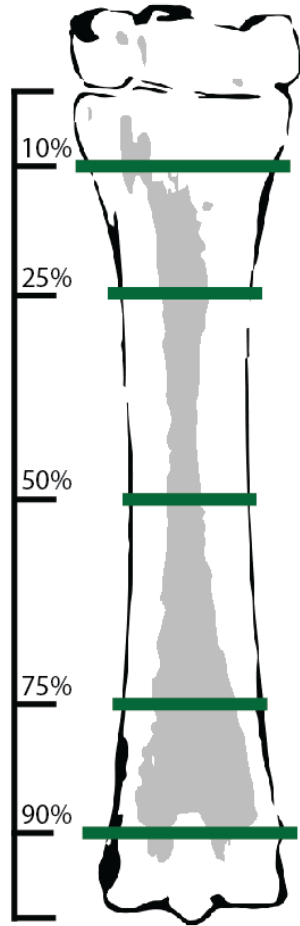


Figure 2.5. pQCT scans were taken at the planes corresponding to 10%, 25%, 50%, 75%, and 90% the length of each bone.

The locations of the planes were determined using length measurements obtained from x-rays taken previously. Voxel size was $0.1 \text{ mm} \times 0.1 \text{ mm} \times 2.2 \text{ mm}$, with the 2.2 mm side running parallel to the length of the bone. Bone was differentiated from surrounding tissue using a macro (courtesy of Dan Schiferl; Bone Diagnostics, Inc.) utilizing user-defined thresholding based on the magnitude of attenuation at each voxel. The thresholding assumes that fat has a density of 0 mg/mm^3 , water and soft tissue have a density of 60 mg/mm^3 , trabecular bone has a density of 700 mg/mm^3 , and cortical bone has a density of $1,200 \text{ mg/mm}^3$. The macro used these defined values to measure 25 parameters from each image (Table 3.1).

The degree of attenuation, translated to density for the purposes of our study, was used to distinguish between cortical bone, trabecular bone, and surrounding material or tissue. At the 10%, 25%, 50%, and 75% levels, trabeculae were automatically contoured using a threshold value of 711 mg/mm^3 . At the 90% level, a value of 169 mg/mm^3 was used. To locate the endosteal surface, a similar algorithm utilized thresholds of 900 mg/mm^3 at the 10% and 75% levels, 600 mg/mm^3 at 25% and 50% levels, and $1,200 \text{ mg/mm}^3$ at the 90% level. To ensure that no cortical bone was included in the trabecular measurements, the endosteal perimeter was contracted by 5%.

When imaging bones from the MC3 experimental group, medical tape was used to secure fractured pieces of the bone together, where possible. Values at sites affected by comminuted fracture were imputed.

2.6 Micro-Computed Tomography (μ CT)

μ CT images were taken using Quantum GX equipment produced by PerkinElmer Inc. Data was collected along the lateral, medial, and dorsal surfaces at 50% the length of each right-side bone in randomly selected samples from the C and LB experimental groups ($n_C = 10$, $n_{LB} = 6$) (Figure 2.6). The number of samples tested were determined primarily by financial restrictions, and exclusion of the MC3 and SSMD groups allowed for sample sizes adequate for statistical comparison between the C and LB groups. Factors such as difficulty in imaging fractured or fragmented bones in the MC3 group and the pathogenesis of proximal sesamoid fractures and their relevance to third metacarpal bones were also taken into consideration.

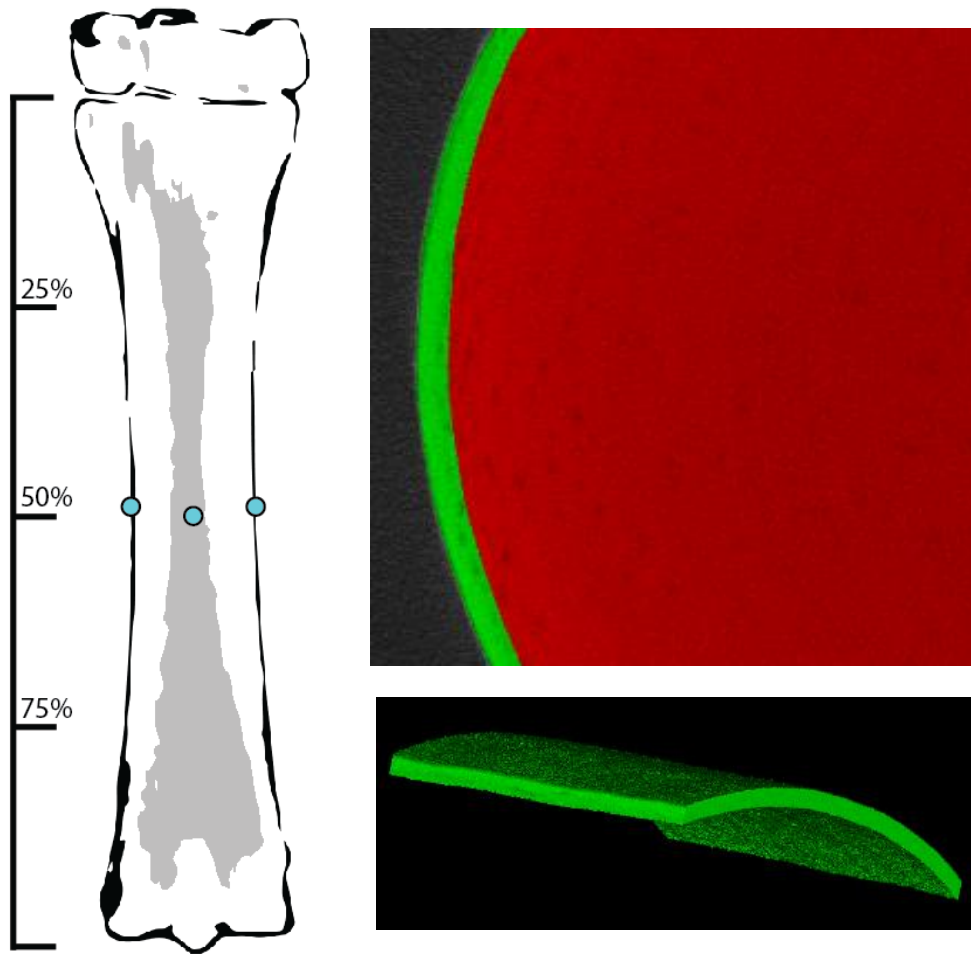


Figure 2.6. Left: μ CT images were taken at the plane corresponding to 50% of the length of each bone, and BMD measurements were made at the corresponding dorsal, medial, and lateral surfaces. Top right: the outer 550 μ m of the bone was eroded in AnalyzePro software. Bottom right: isolated example 15 mm² region of analysis.

Room-temperature bones were secured to the bed with medical tape and scanned at the 50% length plane at 90 kV and 88 mA, a process taking approximately 14 minutes per bone. A voxel resolution of 11 microns was achieved for each image. After this initial scan, 15 mm² areas were selected at the dorsal, medial, and lateral surfaces on which to take subvolume images and bone mineral density measurements.

AnalyzePro software produced by AnalyzeDirect, Inc. was used to analyze the subvolume images. The bone surface was eroded 550 µm and isolated in order to analyze only the periosteal surface of each sample (Figure 2.6). This surface was of interest because of its physiological relevance in bone modeling and because it was the same approximate tissue on which indentation testing was performed. Subvolume images were also examined visually to inspect for potential signs of bone modeling such as woven bone along the periosteal surfaces.

2.7 Statistics

Statistical tests were carried out using IBM SPSS Statistics 24, unless otherwise noted. Before inter-group analysis was performed, measurements from right and left limbs in each horse were compared against one another via paired t-test. To avoid unnecessary confounding variation, samples were grouped by experimental group and location: for example, each cortical thickness value on the dorsal surface at the 75% length location in the Control group was included in one test.

It was necessary to consider Family-wise error rates when dealing with a large volume of comparisons. The likelihood of type I errors, or false positives, increases as multiple hypothesis tests are performed at once due to the nature of the tests themselves. Controlling procedures such as Bonferroni or Šidák corrections can be implemented to account for this phenomenon. Here, Bonferroni corrections for multiple comparisons were implemented in each paired t-test by dividing the critical p-value by the number of comparisons being performed with each test. If the test found no relevant significant ($p < 0.05 / n$) differences between measurements taken on right and left limbs of a given horse, the contralateral measurements were averaged together in an attempt to provide a more complete picture of each horse without the need to manage two sets of data or arbitrarily selecting a single limb. If significant differences were discovered, logical discretion was used to dictate whether left and right values should be averaged together for analysis.

A custom syntax was created for comparing data by site and experimental group: a linear fixed effect model with site, group, and site * group interactions as fixed effects and individual horses as random effects (with default covariance type for random effects, *variance components*, selected). The model utilizes restricted maximum likelihood (REML) methods as opposed to ANOVA, as REML is able to more effectively manage unbalanced experimental designs and also allows for inferences about covariance factors in the model. Bonferroni adjustments were implemented to control for the multiple comparisons being made in each test. A critical p-value of 0.05 selected for use with the linear model. The syntax with an example parameter of interest is shown in Figure 2.7.

```
MIXED BMSi BY site group horseno
/CRITERIA=CIN(95) MXITER(100) MXSTEP(5) SCORING(1) SINGULAR(0.000000000001) HCONVERGE(0,
ABSOLUTE) LCONVERGE(0, ABSOLUTE) PCONVERGE(0.000001, ABSOLUTE)
/FIXED=site group site*group | SSTYPE(3)
/METHOD=REML
/RANDOM=horseno | COVTYPE(VC)
/EMMEANS=TABLES(site) compare(site) ADJ(BONFERRONI)
/EMMEANS=TABLES(group) compare(group) ADJ(BONFERRONI)
/EMMEANS=TABLES(site*group) compare(site) ADJ(BONFERRONI)
/EMMEANS=TABLES(site*group) compare(group) ADJ(BONFERRONI)
```

Figure 2.7. A linear mixed model was created in SPSS Statistics to analyze data (here, BMSi) with effects being test location (site), experimental group (group), and individual horse (horseno).

For each test, each of the four experimental groups were initially included in the linear mixed model. After obtaining these results, the LB and MC3 groups were combined into one group and the test was run again. This was done to account for the small sample size of the MC3 group and because third metacarpal bones are a type of long bones. Mann-Whitney tests were performed for each metric between bones in the LB and MC3 groups to determine if they were functionally equivalent for the purposes of this study.

If an individual comparison was to be made between more than two fracture groups, one-way ANOVA tests were performed in SPSS. Post-hoc Tukey and Bonferroni tests were utilized to gain further insights into the results, when appropriate.

To explore observed intra-bone differences in percutaneous BMSi between different experimental groups, a proxy variable obtained by taking the differences in BMSi between the medial and dorsolateral surfaces at each length was created. These surfaces were selected based on the medial surfaces being the apparent sites of highest BMSi and the dorsolateral surface tending to have lower values. The dorsomedial surface could have also acted as the surface to

which the medial surface was compared. The values obtained by taking the differences at the midshaft of the bones, where the strongest trend of this pattern was observed, were compared using one-way ANOVA.

2.8 Multivariable Regression Model

At the time of this study, the predictive statistical model was still in its infancy. MedCalc statistical software was used to perform logistic regression analyses on collections of data based primarily on educated discretion rather than sensitivity analyses. Due to the perceived physiological relevance of and observed intergroup differences at the 50% dorsal site on third metacarpal bones, all included data was collected from this location unless otherwise noted.

First, all variables were analyzed using a logistic regression model in isolation to determine their capability in distinguishing between bones from horses with or without fracture and whether or not they might be of use in models including multiple variables. Because regression models only allow dichotomous outputs, experimental groups were divided into Control and Fracture (SSMD, LB, and MC3 consolidated into one group). Receiver operating characteristic (ROC) curves were built from the logistic regression data, with the area under the curve (AUC) serving as an indicator of how much predictive power the model may possess.

When moving forward with selecting which variables to include in the next iteration of the model, the perceived utility, collinearity with other variables, and clinical relevance were considered. The first multivariable model was assembled prior to complete collection of pQCT data and included TID, avg ED, no-skin BMSi, and cortical thickness due to promising results seen throughout this study. A next iteration of the model, containing only clinically relevant parameters (percutaneous BMSi, cortical thickness, and mass of the horse) was then compiled. After completion of pQCT data collection, a final iteration was produced that included parameters selected for their perceived uniqueness, clinical relevance, and capability: BMD at 90% length, the difference in BMSi at the dorsolateral and medial sites at 50% length, and cortical thickness at the mid-dorsal surface. Each of these parameters could feasibly be measured in a standing horse, each examined a different property of bone, and each detected significant differences between experimental groups in this study.

3. RESULTS

3.1 Left & Right Limb Comparison

In this study, measurements were taken on both left and right third metacarpal bones in each horse when possible. However, in a clinical setting, testing would be more cost- and time-efficient if only one leg was measured. Additionally, being able to average left and right measurements together would make the data analysis in this study more concise and discernable. To validate that measurements from right and left limbs do not significantly differ from one another, paired t-tests were performed at each test location within each fracture group with Bonferroni correction of the p-value for multiple comparisons. Bonferroni corrections were performed by dividing the standard p-value of 0.05 by the number of statistical tests being performed simultaneously. If no significant differences were not discovered, the mean of the values from the left and right bones were used in analysis. If significant differences were discovered, logical discretion was used to dictate whether left and right values should be averaged together for analysis.

The MC3 fracture group was not subjected to left-right paired t-tests due to a lack of intact left / right pairs (n=1) leading to prohibitively small sample sizes. Left / right testing was also not performed with μ CT data because only right limbs were tested.

3.1.1 X-ray

No significant differences were discovered between left and right bones in any of the experimental groups after the Bonferroni correction ($p < 0.006$) for multiple comparisons was implemented. Contralateral MC3 bones were rarely found to have the exact same length, though no significant differences existed when left and right limbs were compared via paired t-test or independent sample t-test. When cortical thicknesses were normalized by the length of the bone itself, there remained no significant differences between left and right limbs in any experimental group.

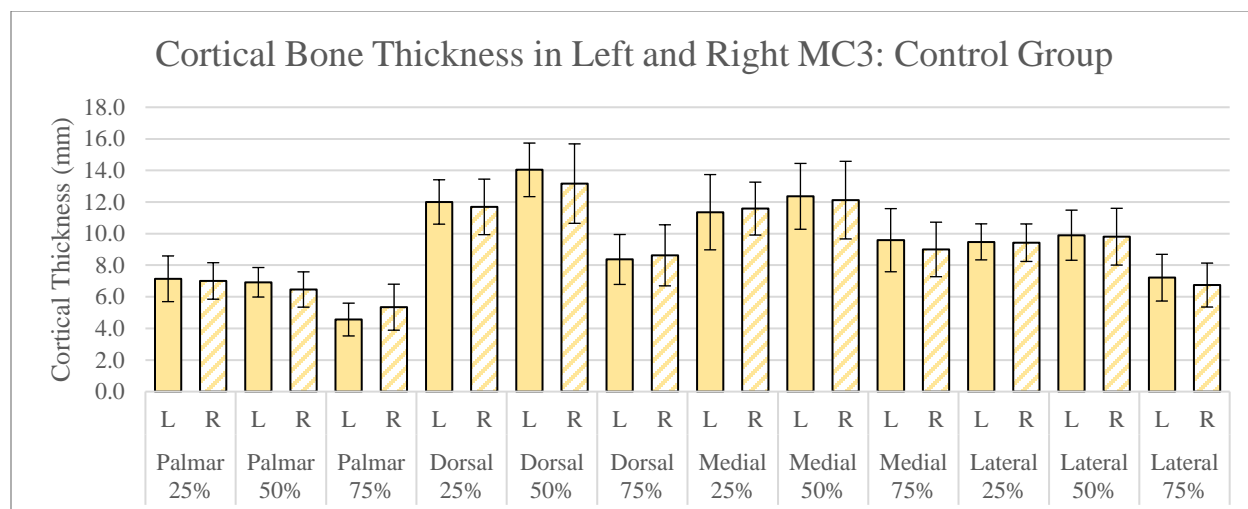


Figure 3.1. No significant ($p < 0.006$) differences between left and right MC3 cortical thickness in C group. ($n = 20$). Bar and error bars represent mean \pm 1 standard deviation. Paired t-test.

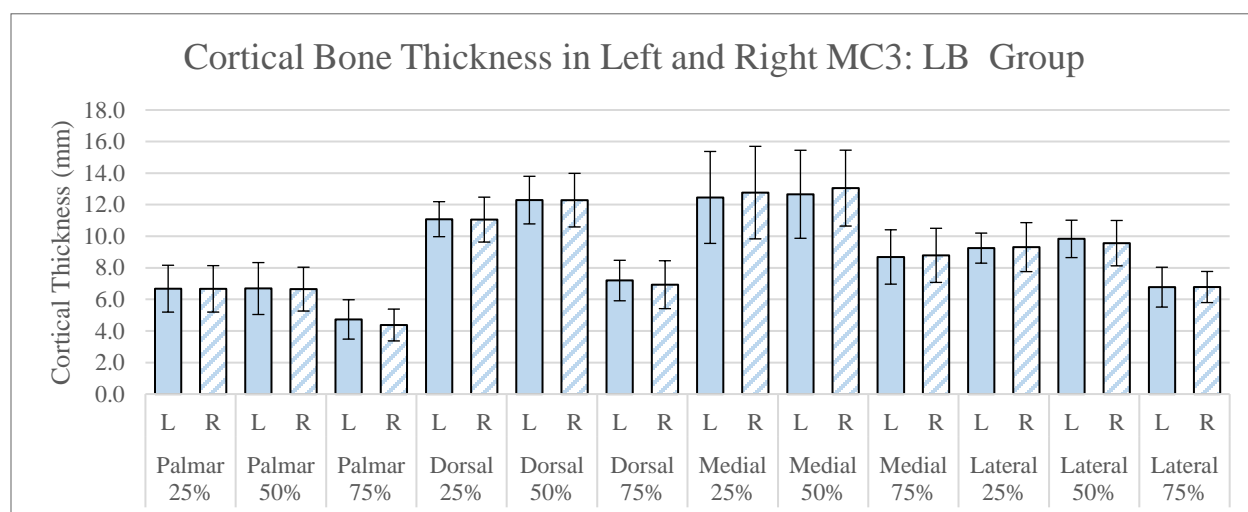


Figure 3.2. No significant ($p < 0.006$) differences between left and right MC3 cortical thickness in LB group. ($n = 19$). Bar and error bars represent mean \pm 1 standard deviation. Paired t-test.

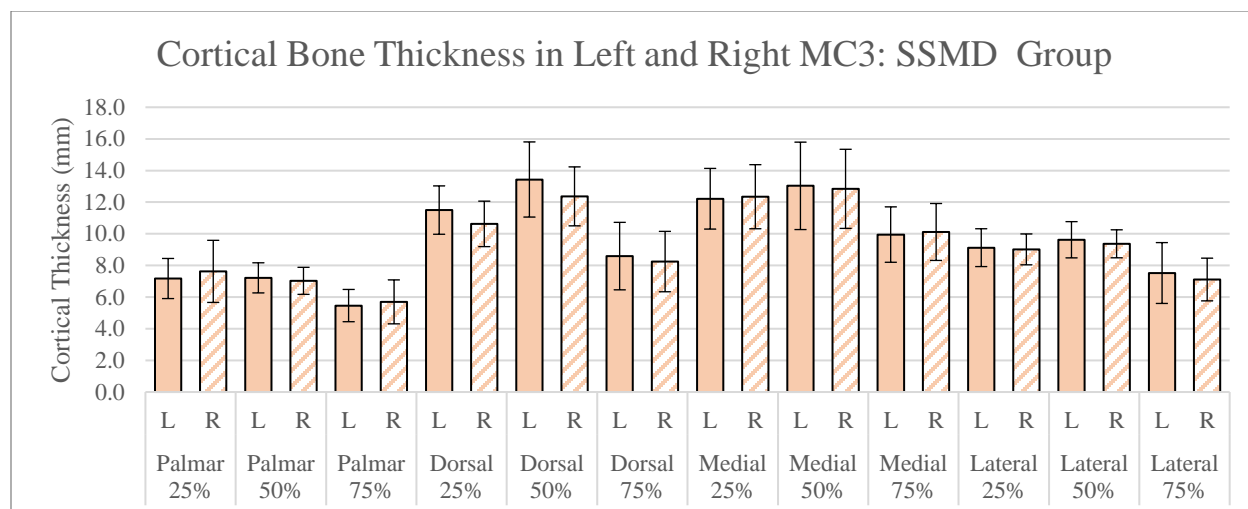


Figure 3.3. No significant ($p < 0.006$) differences between left and right MC3 cortical thickness in SSMD group.. (n = 19). Bar and error bars represent mean \pm 1 standard deviation. Paired t-test.

3.1.2 OsteoProbe

Skin-on

Paired t-tests with Bonferroni correction for multiple comparisons were performed at each site within each experimental group. No significant ($p < 0.004$) differences in percutaneous BMSi between left and right third metacarpal bones existed.

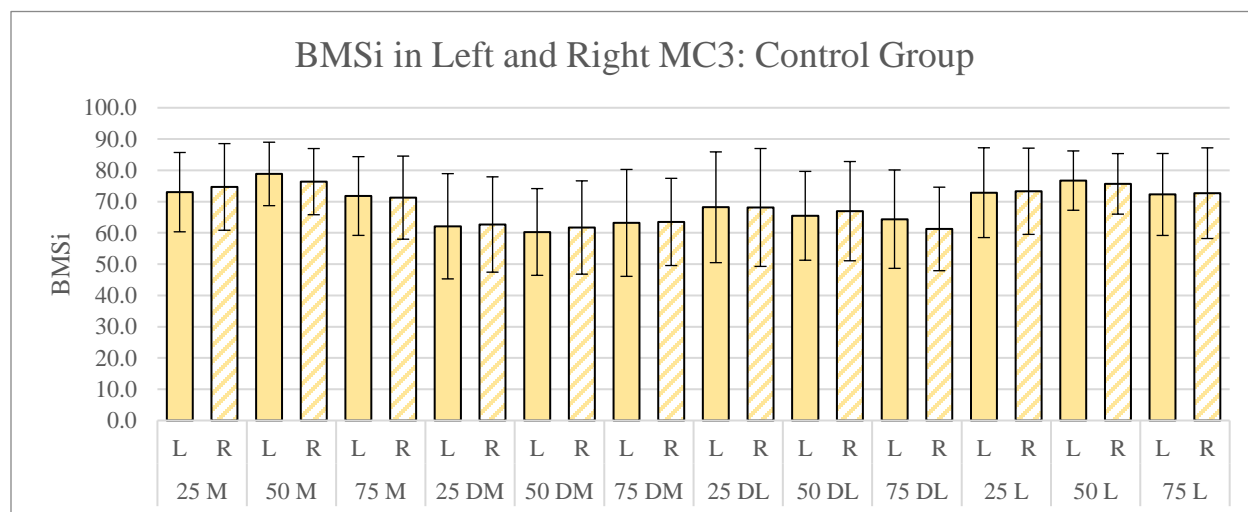


Figure 3.4. No significant ($p < 0.004$) differences between left and right MC3 BMSi in Control group. (n = 16 – 18, depending on site). Bar and error bars represent mean \pm 1 standard deviation. Paired t-test. M = medial, DM = dorsomedial, DL = dorsolateral, L = lateral.

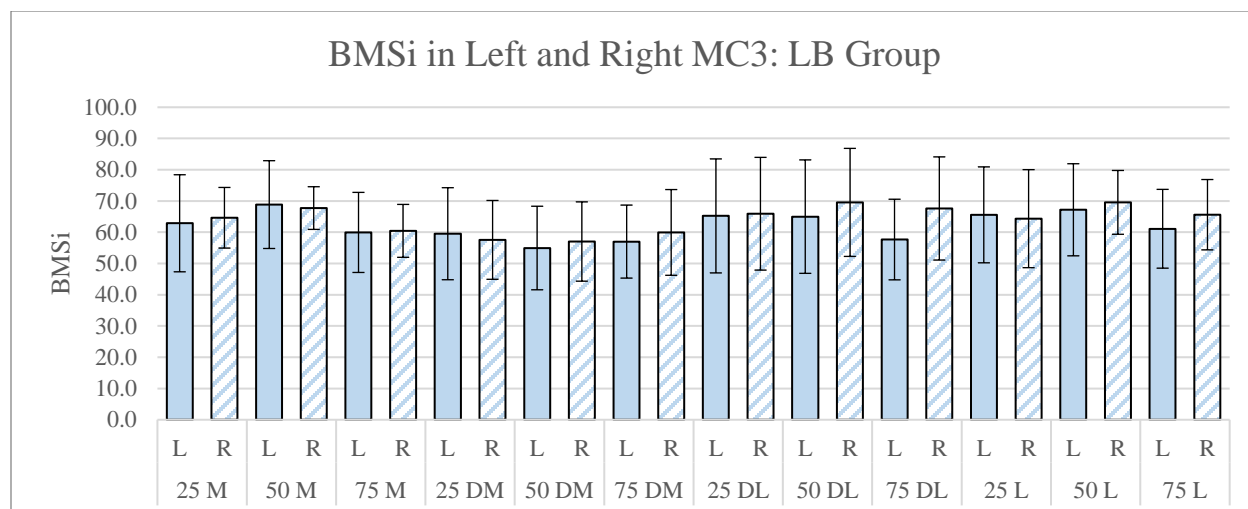


Figure 3.5. No significant ($p < 0.004$) differences between left and right MC3 BMSi in LB group. ($n = 14 - 15$, depending on site). Bar and error bars represent mean ± 1 standard deviation. Paired t-test. M = medial, DM = dorsomedial, DL = dorsolateral, L = lateral.

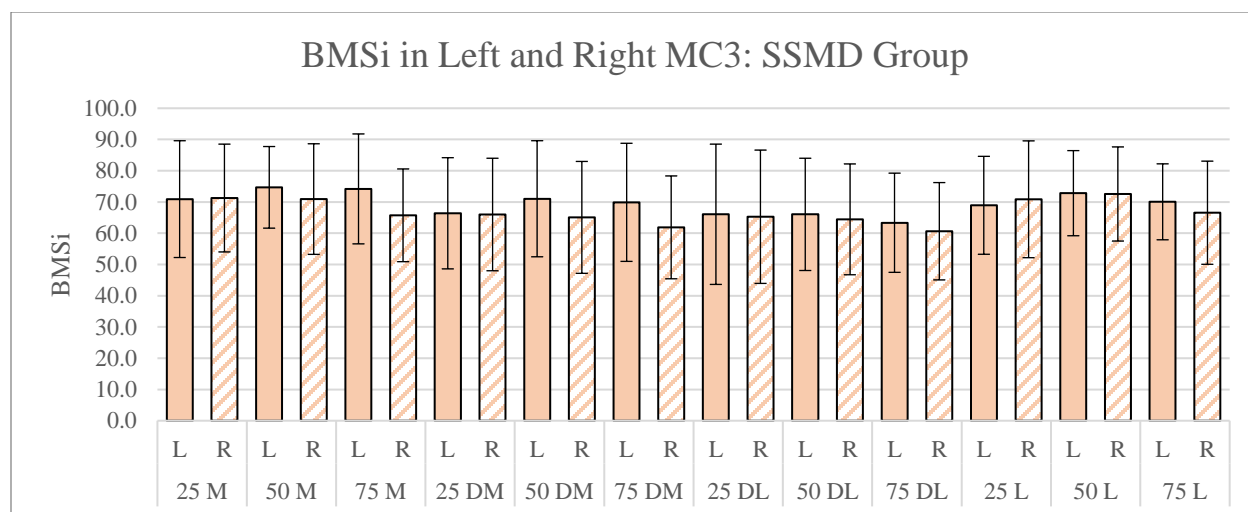


Figure 3.6. No significant ($p < 0.004$) differences between left and right MC3 BMSi in SSMD group. ($n = 15 - 16$, depending on site). Bar and error bars represent mean ± 1 standard deviation. Paired t-test. M = medial, DM = dorsomedial, DL = dorsolateral, L = lateral.

No Skin

Paired t-tests with Bonferroni correction for multiple comparisons were performed at each site within each experimental group. No significant ($p < 0.004$) differences in no-skin BMSi between left and right third metacarpal bones existed.

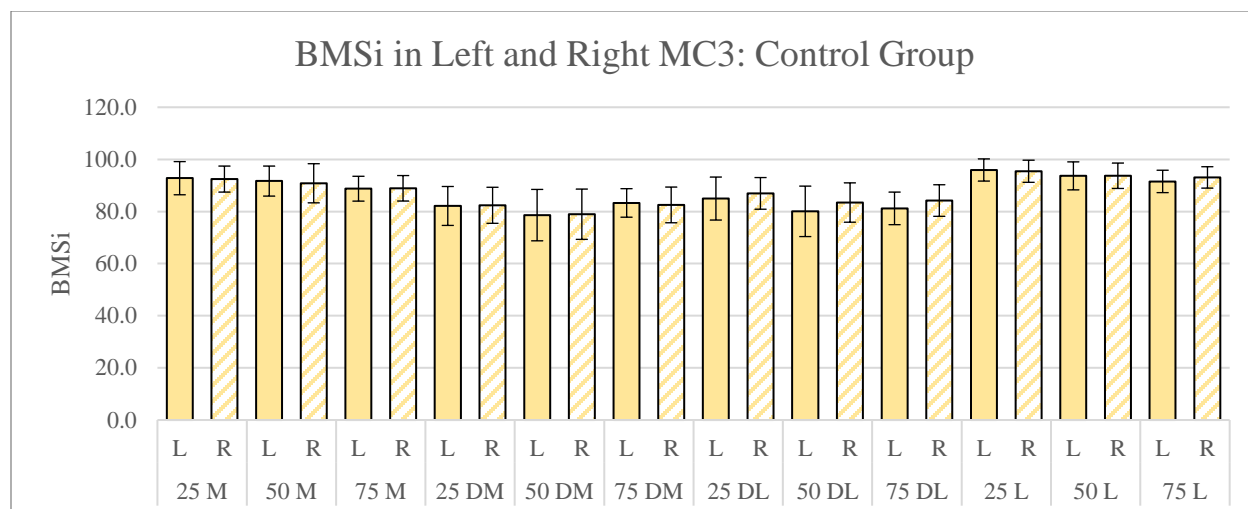


Figure 3.7. No significant ($p < 0.004$) differences between left and right MC3 BMSi in Control group. ($n = 19 - 20$, depending on site). Bar and error bars represent mean ± 1 standard deviation. Paired t-test. M = medial, DM = dorsomedial, DL = dorsolateral, L = lateral.

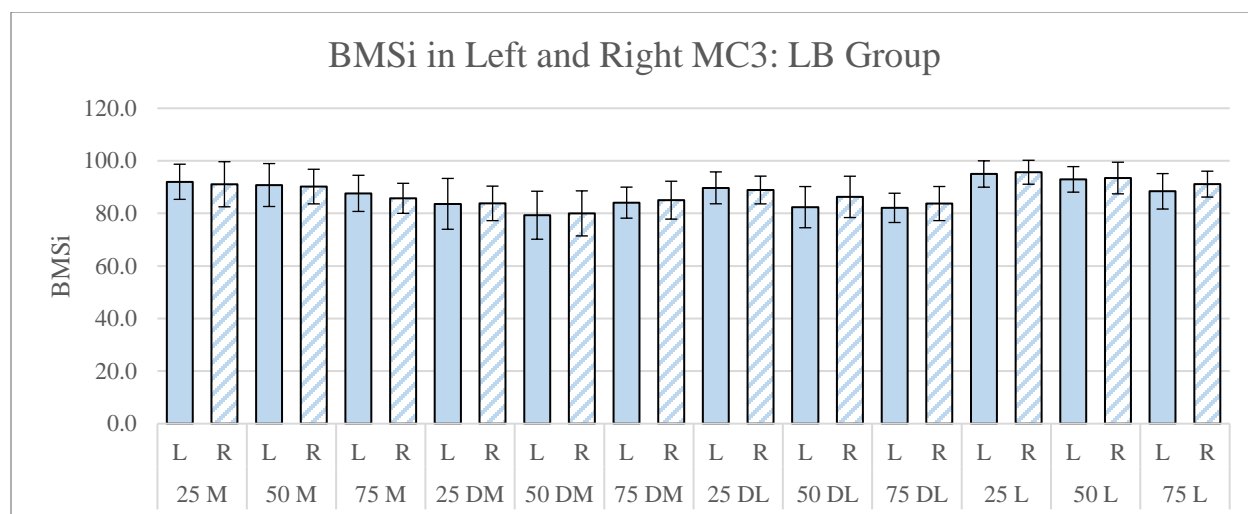


Figure 3.8. No significant ($p < 0.004$) differences between left and right MC3 BMSi in LB group. ($n = 18$). Bar and error bars represent mean ± 1 standard deviation. Paired t-test. M = medial, DM = dorsomedial, DL = dorsolateral, L = lateral.

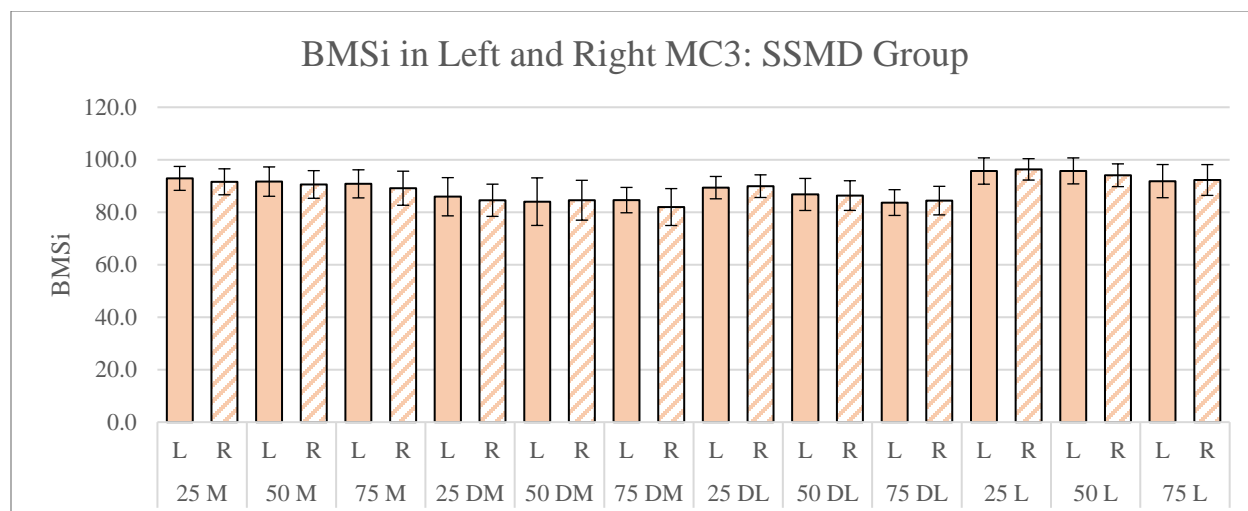


Figure 3.9. No significant ($p < 0.004$) differences between left and right MC3 BMSi in SSMD group. ($n = 19 - 20$, depending on site). Bar and error bars represent mean ± 1 standard deviation. Paired t-test. M = medial, DM = dorsomedial, DL = dorsolateral, L = lateral.

3.1.3 BioDent

When comparing BioDent parameters from right and left limbs among each of the fracture groups after Bonferroni correction, five significant ($p < 0.005$) differences were discovered. Four of these differences regarded loading or unloading slopes, indicating potential differences in stiffness or elastic moduli, and one regarded average energy dissipated. These discrepancies were not seen as a compelling argument to perform all left- and right-limb analyses separately, so left and right data was averaged together for the remaining analyses.

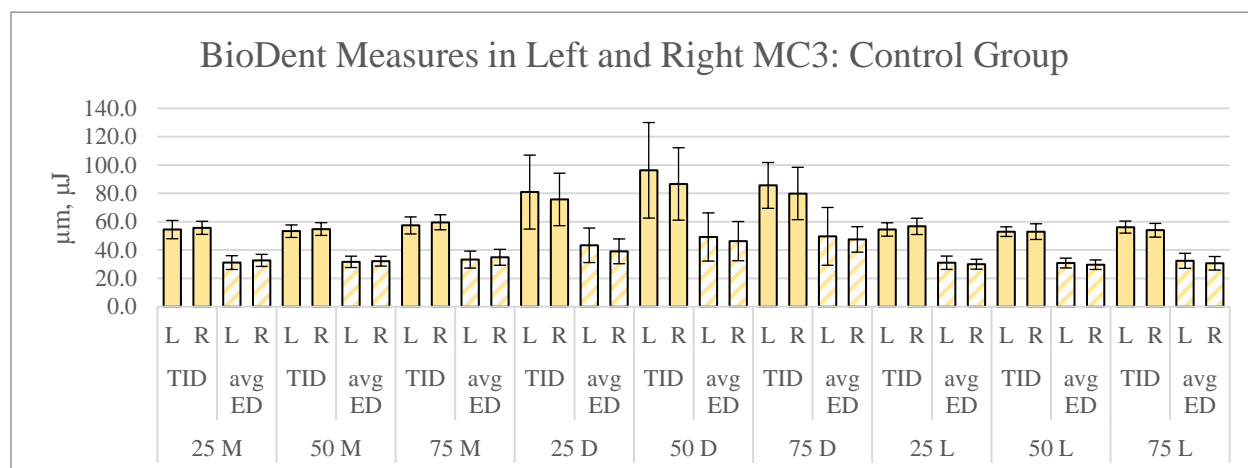


Figure 3.10. No significant ($p < 0.005$) TID or avgED differences between left and right limbs in the Control group. ($n = 16 - 20$, depending on site). Bar and error bars represent mean ± 1 standard deviation. Paired t-test. TID = total indentation distance, avg ED = average energy dissipated, M = medial, D = dorsal, L = lateral.

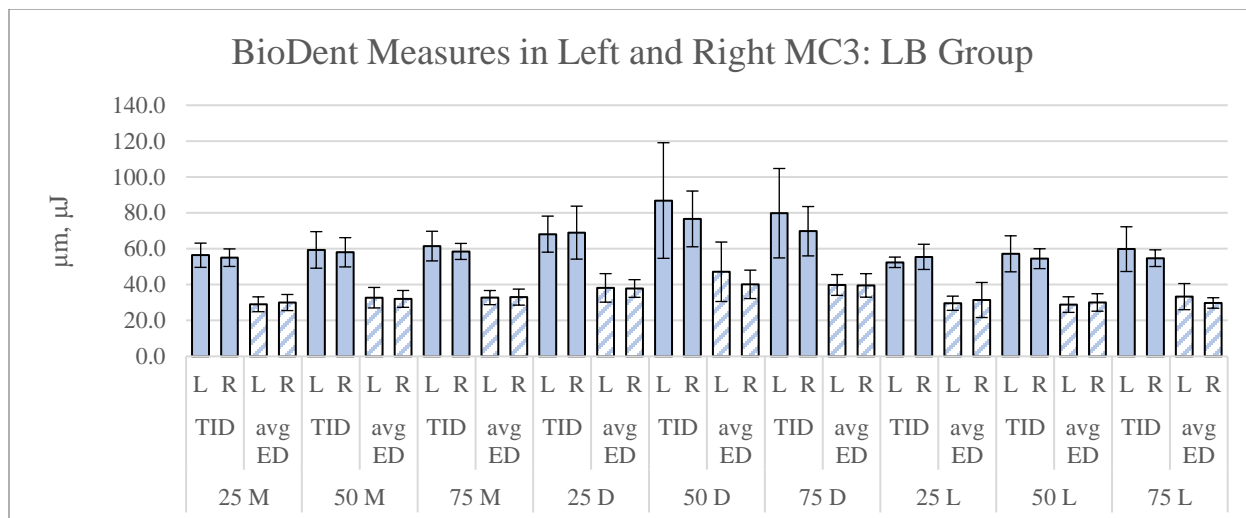


Figure 3.11. No significant ($p < 0.005$) TID or avgED differences between left and right limbs in the LB group. ($n = 17 - 18$, depending on site). Bar and error bars represent mean ± 1 standard deviation. Paired t-test. TID = total indentation distance, avg ED = average energy dissipated, M = medial, D = dorsal, L = lateral.

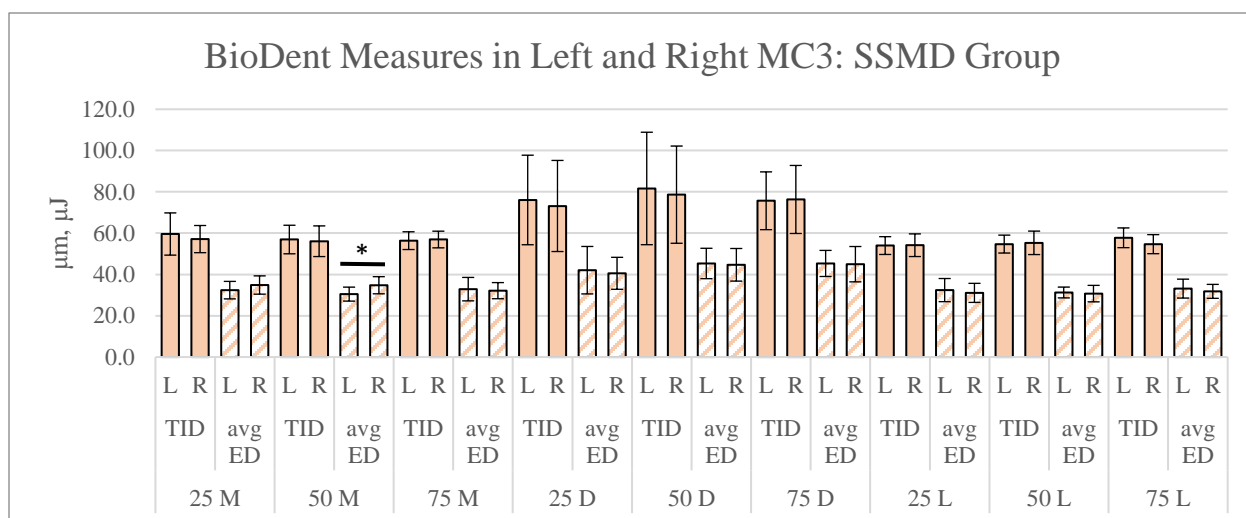


Figure 3.12. Significant ($p < 0.005$) avgED difference detected between left and right limbs at the 50% medial site in the SSMD group. ($n = 16 - 20$, depending on site). Bar and error bars represent mean ± 1 standard deviation. Paired t-test. TID = total indentation distance, avg ED = average energy dissipated, M = medial, D = dorsal, L = lateral.

Five significant ($p < 0.005$) differences existed between left and right limbs (Table 3.1), four of which belonged to parameters that would likely not be included in the final predictive model due to reasons outlined in Section 2.4.

Table 3.1. Significant differences between left and right third metacarpal bones are outlined below, with the value of the left mean or standard deviation in the top row for each instance and the value of the right in the bottom row.

| Group | Site | Parameter | Mean (Left / Right) | Standard Deviation (Left / Right) | p-value |
|-------|-------------|-----------|---------------------------|--|---------|
| C | 50% lateral | US1 | 0.71 | 0.052 | 0.003 |
| | | | 0.761 | 0.064 | |
| C | 25% lateral | US1 | 0.698 | 0.04 | 0.001 |
| | | | 0.752 | 0.059 | |
| C | 25% lateral | avg US | 0.753 | 0.036 | 0.002 |
| | | | 0.801 | 0.059 | |
| SSMD | 50% medial | avg ED | 30.48 | 3.39 | 0.003 |
| | | | 34.77 | 4.15 | |
| SSMD | 25% medial | avg LS | 0.564 | 0.036 | 0.001 |
| | | | 0.532 | 0.039 | |

Though select differences remained after Bonferroni corrections were applied to BioDent data, there were no consensus on left bones having greater values than right or vice versa. The control group was not lacking in significant left / right differences compared to the fracture groups, suggesting that significantly different values between limbs does not necessarily forecast a fracture. Additionally, because stiffness and elastic modulus at this scale are not perceived to be critical considerations for our purposes and because a discrepancy in avgED was only observed in one location among one fracture group, these differences were not seen as compelling reasons to not average all left and right data together in each pair of bones.

Therefore, data from the left and right limbs were averaged together to create one set of values for each horse during the proceeding data analysis.

3.1.4 pQCT

Among the 26 parameters measured via pQCT on the 5 sites in each of the experimental groups (excluding MC3 due to lack of intact pairs of bones), one significant difference emerged. At the 25% length plane of bones in the SSMD experimental group, periosteal circumference was found to be significantly larger in left bones ($p = 0.029$). However, the circumference is reported to be approximately twice as large in the left bones in this location as it is in the right—this difference would be visibly discernable and obvious. Upon further inspection, it appears as if

software settings may be misaligned in regards to this parameter, as many bones are reported as having a circumference of zero. Periosteal circumference will not be used in this study.

3.2 Fracture Group Comparisons

3.2.1 X-Ray

When an ANOVA was performed to compare the length of third metacarpal bones between the experimental groups, no significant differences existed.

Cortical thicknesses were analyzed using a linear mixed model with site, group, and site-group interactions as fixed effects and horse ID as a random effect. No significant differences between groups or significant site * group interactions were discovered. Results did not differ when data were normalized by mass of the horse.

Because of the small sample size of the MC3 group and the possibility that third metacarpal bones are not distinct from other long bones, data belonging to the MC3 and LB groups were then combined into one group. A Mann-Whitney U test performed between the two groups determined that there are no significant differences between their cortical thicknesses at any site on the bones, supporting the decision to combine the groups. When analyzed using the linear model, site * group interactions were significant ($p = 0.049$). Post-hoc analyses indicate that the cortical thickness of the LB-MC3 group was significantly less than that of the C group at the 50% dorsal site and significantly less than both C and SSMD groups at the 75% dorsal site. It was also found that the cortex of the LB-MC3 group is significantly less thick than the SSMD group at the 75% medial site.

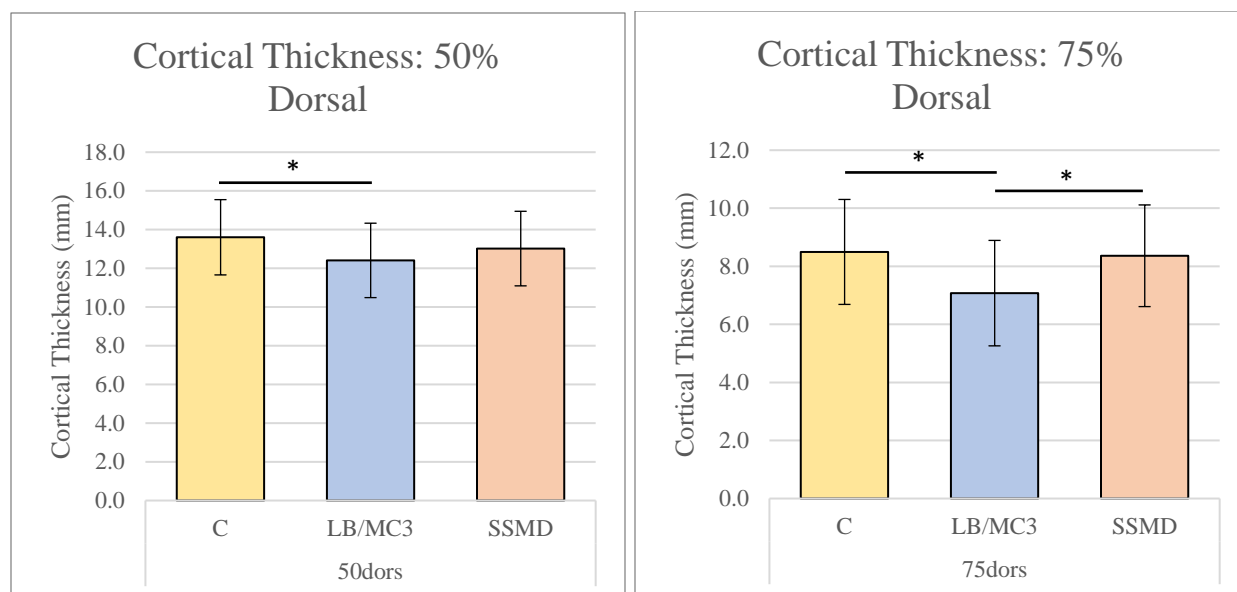


Figure 3.13. Cortical thickness significantly ($p < 0.05$) varied between fracture groups at the mid- and distal-dorsal surfaces. ($n_C = 20$, $n_{SSMD} = 20$, $n_{LB/MC3} = 24$). Bar and error bars represent mean \pm 1 standard deviation.

Further consolidating the fracture groups, the data were analyzed using the mixed model with all of the fracture groups combined into one. A significant site * group interaction was discovered, and pairwise comparisons revealed significant differences between the C group and new Fx group at the 50% and 75% dorsal sites.

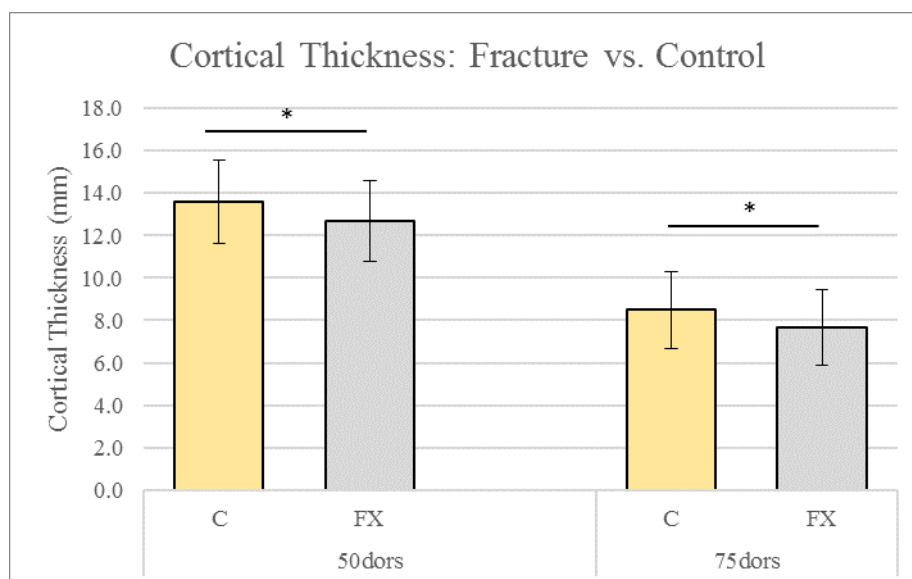


Figure 3.14. Cortical thickness was found to be significantly ($p < 0.05$) greater in the Control group than the combined Fracture group along the mid- and distal-dorsal surfaces. ($n_C = 20$, $n_{FX} = 44$). Bar and error bars represent mean \pm 1 standard deviation.

3.2.2 OsteoProbe

Skin-on

When skin-on OsteoProbe data was analyzed using the linear mixed model, it returned no significant differences between groups or significant site * group interactions. A Mann-Whitney U test determined no significant differences in BMSi at any site between the LB and MC3 groups, and when LB and MC3 data were consolidated into one group, there remained no significant differences.

When all fracture data was consolidated into one fracture group to be compared against the Control group, a significant site * group interaction was reported. Upon investigation of the pairwise comparisons, it was evident that this significance was driven by differences by site within groups, not differences by group at a given site.

No significant differences in BMSi were found at any site between C and Fx groups, though some trends did appear: bones in the C group had higher BMSi values than the Fx group along the medial and lateral surfaces ($p = 0.002$ and $p = 0.015$, respectively), whereas on the dorsolateral and dorsomedial surfaces, the opposite appeared to tend to be true ($p = 0.785$ and $p = 0.726$, respectively). When values of the difference between dorsolateral and medial BMSi values at the midshaft of each bone are treated as a distinct “proxy” variable, the values from the Control group are found to be significantly ($p < 0.05$) greater than that of the LB group and MC3 groups when tested via one-way ANOVA (Figure 3.15).

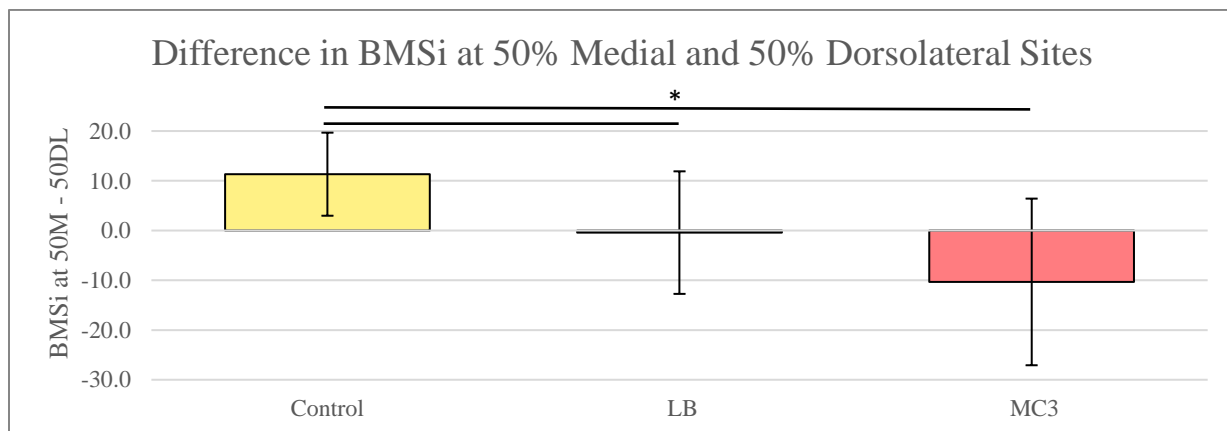


Figure 3.15. The Control group displays a pattern of higher BMSi values along the medial surface than the dorsal, while the LB and MC3 groups do not. The proxy variable obtained by taking the difference in BMSi between the 50% medial and 50% dorsolateral surfaces is significantly ($p < 0.05$) greater in the C group than in the LB or MC3 groups. Note the high standard deviation. ($n_C = 20$, $n_{LB} = 19$, $n_{MC3} = 5$). Bar and error bars represent mean \pm 1 standard deviation.

No Skin

When the same tests are performed on the no-skin OsteoProbe data, no significant group differences, site * group interactions, or medial-minus-dorsolateral differences are observed when all four fracture groups are compared or when LB and MC3 are grouped together. A Mann-Whitney U test determined a significant ($p = 0.044$) difference between the LB and MC3 groups at the 25% dorsolateral level, though this discovery bears little relevance since no positive results were found by combining these two groups.

When the control group is compared against all the combined fracture groups, no significant differences or interactions are observed. However, the same phenomenon from skin-on testing held true: on medial and lateral surfaces, the control group tends to have higher BMSi values, whereas on dorsolateral and dorsomedial surfaces, the opposite is true.

3.2.3 BioDent

When BioDent data for each parameter was analyzed using the mixed model, no significant differences between groups or site * group interactions were discovered. Trends were noticed in parameters such as Total Indentation Distance, though relatively high variation in the data or the small sample size of the MC3 group may have barred any differences from being deemed significant.

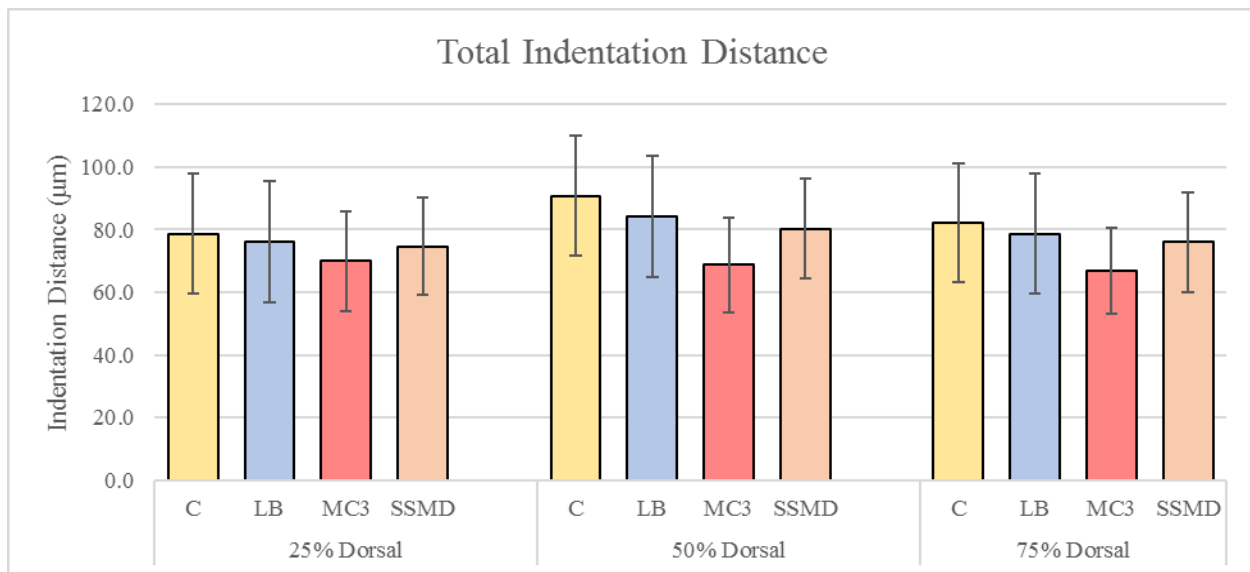


Figure 3.16. Though no significant differences arose, the MC3 group trended lower than the other experimental groups, particularly the Control group. ($n_C = 20$, $n_{LB} = 19$, $n_{MC3} = 3$, $n_{SSMD} = 20$). Bar and error bars represent mean ± 1 standard deviation.

A Mann-Whitney U test detected differences in 1st-cycle unloading slope ($p = 0.003$) and average unloading slope ($p = 0.001$) between the LB and MC3 groups at the 50% medial location, but no differences in other parameters or locations. When LB and MC3 groups are combined, no significant differences emerged. When all fracture groups are combined and compared against the Control group, a significant site * group interaction exists in Average Energy Dissipated at the 75% dorsal site. See Tables A.20 – A.22 in the Appendix for a summary of all p-values across parameters and experimental setups.

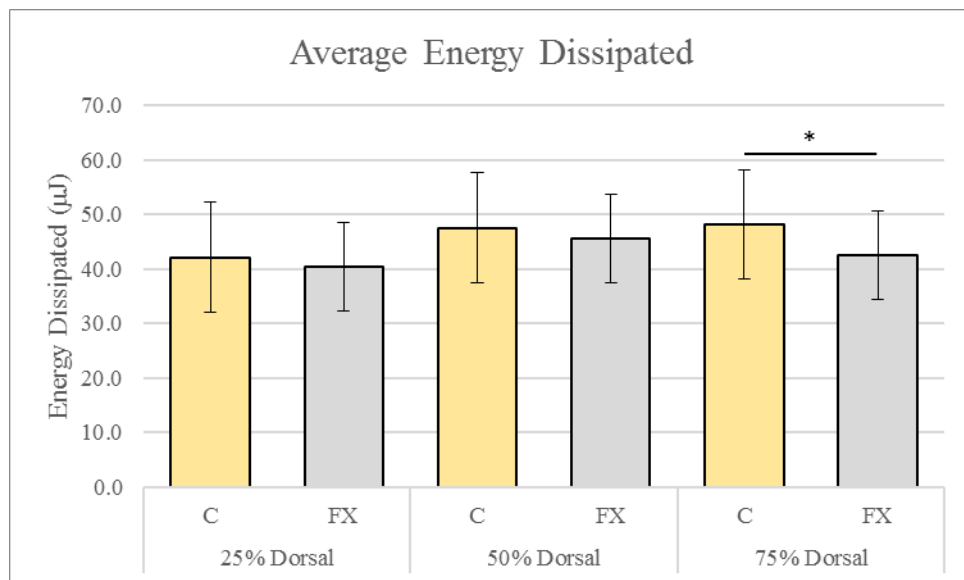


Figure 3.17. Average energy dissipated was found to be significantly ($p < 0.05$) greater in the Control group than the combined Fracture group at the 75% dorsal site. ($n_C = 20$, $n_{FX} = 42$). Bar and error bars represent mean \pm 1 standard deviation.

3.3 OsteoProbe versus BioDent Correlations

Correlations between BMSi and each of the BioDent's parameters existed in in varying degrees. TID, IDI, and avgED shared some of the strongest correlations with BMSi.

To validate the relationship between BioDent and OsteoProbe microindentation testing methods, each parameter was compared against the other modality. The 50% dorsal site was examined first due to it being perhaps the most clinically-relevant location on the third metacarpal bone. Because OsteoProbe testing was performed only on the dorsolateral and dorsomedial surfaces and not the dorsal surface itself, both were initially tested against the BioDent parameter at hand. Next, the dorsolateral and dorsomedial values were averaged together in an attempt to

interpolate what the BMSi value on the dorsal surface may be. These averaged data resulted in stronger correlations with most BioDent parameters, and were thus selected as the OsteoProbe data to be utilized in the comparison calculations.

The 50% dorsal site was found to consistently have the highest R^2 values across most BioDent parameters. Despite the medial and lateral measurements being taken on the same surface between instruments, as opposed to averaging two sites together, the distance-based parameters show nearly no correlation with one another (Table A.23).

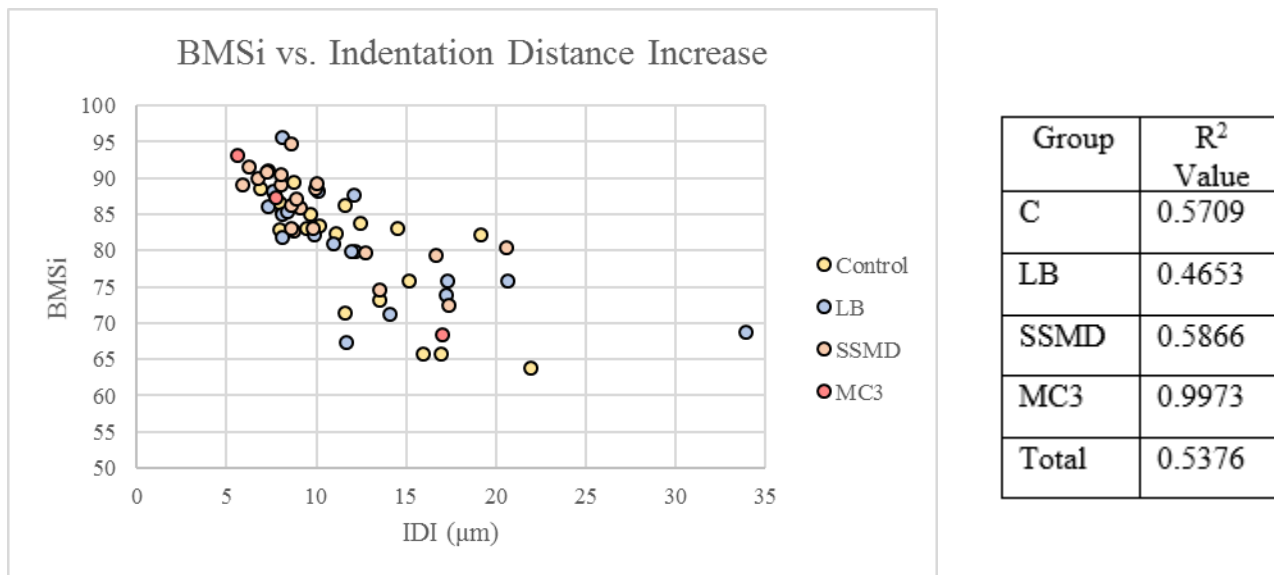


Figure 3.18. BMSi and IDI show a modest correlation in each experimental group

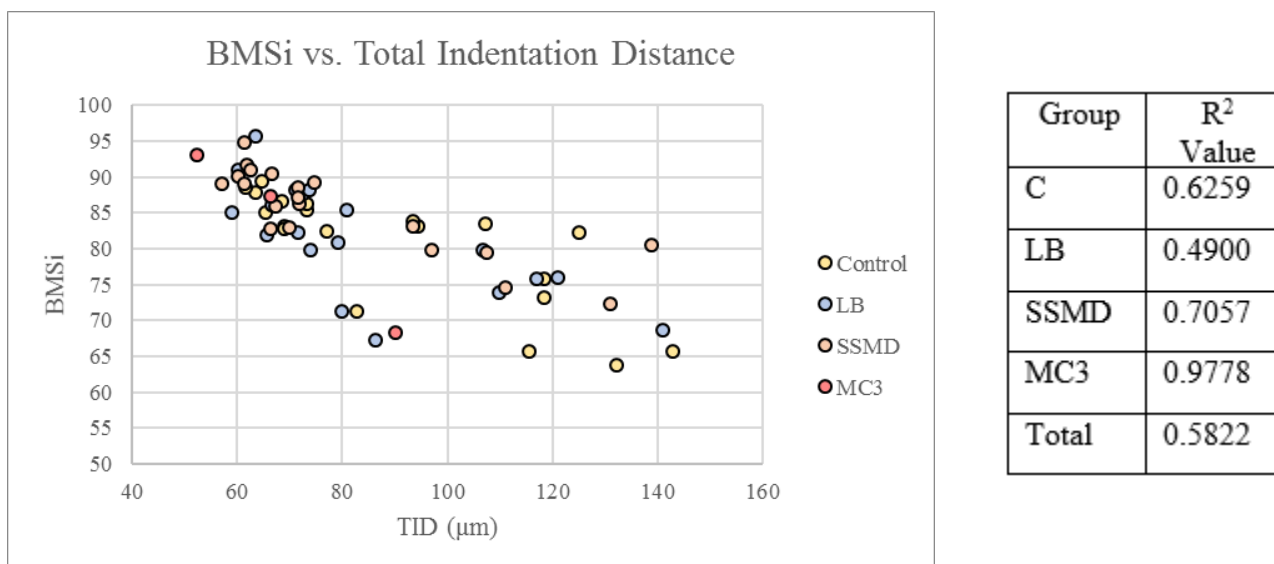


Figure 3.19. BMSi and TID show a modest correlation in each experimental group

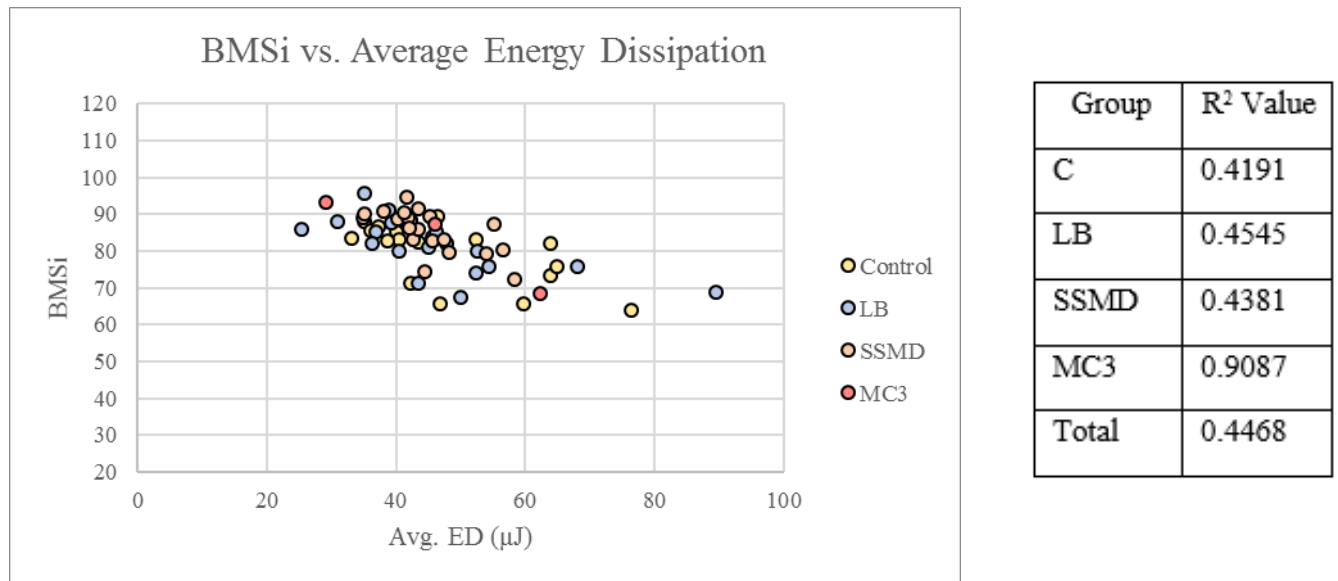


Figure 3.20. BMSi and avgED show a modest negative correlation in each experimental group.

3.4 Peripheral Quantitative Computed Tomography (pQCT)

Many parameters measured by pQCT reported significant differences, particularly significant site * group interactions. Contrary to other methodologies that were explored in this study, less differences and interactions were observed when the fracture groups are consolidated and compared against the control group. Though pQCT is a highly versatile methodology with many outputs (see Table 3.2), few will likely be considered candidates for predictor variables in the final regression model due to a lack of distinguishing ability or lack of perceived relevance.

After statistical analysis was performed, cortical thickness, cortical thickness standard deviation, endosteal circumference, and endosteal circumference (circular ring model) were found to have missing values. The results from these parameters are not accurate and should not be taken into consideration.

Table 3.2. Each parameter measured by pQCT testing. p-values for significant group differences and significant group * site interactions, when tested with all fracture groups or just Control compared against the combined Fracture group, bolded. Grayed-out values were found to have missing values after analysis and should not be considered.

| Parameter | Abbreviation | All Fx Groups | | C vs. Fx. | |
|---|--------------|---------------|--------------|-----------|--------------|
| | | Group | Site * Group | Group | Site * Group |
| Cortical bone density (mg / cm ³) | CRT_DEN | 0.000 | 0.006 | 0.068 | 0.143 |
| Cortical & subcortical content per slice (mg / mm) | CRTSUB_CNT | 0.021 | 0.012 | 0.332 | 0.143 |
| Cortical & subcortical bone density (mg / cm ³) | CRTSUB_DEN | 0.007 | 0.001 | 0.543 | 0.826 |
| Total bone density (mg / cm ³) | TOT_DEN | 0.004 | 0.005 | 0.100 | 0.235 |
| Axial area moment of inertia (via circular ring model) | I_CIRC | 0.837 | 0.012 | 0.608 | 0.378 |
| Total bone area (mm ²) | TOT_A | 0.973 | 0.009 | 0.762 | 0.331 |
| Cortical thickness (via circular ring model) (mm) | CRT_THK_C | 0.978 | 0.048 | 0.998 | 0.356 |
| Periosteal circumference (via circular ring model) (mm) | PERI_C | 0.994 | 0.009 | 0.848 | 0.309 |
| Cortical & subcortical bone area (mm ²) | CRTSUB_A | 0.100 | 0.000 | 0.391 | 0.033 |
| Axial area moment of inertia of cortical area (x-axis) | IX_CRT_A | 0.438 | 0.000 | 0.473 | 0.153 |
| Axial area moment of inertia of cortical area (y-axis) | IY_CRT_A | 0.644 | 0.000 | 0.548 | 0.468 |
| Polar area moment of inertia of cortical area | IP_CRT_A | 0.550 | 0.000 | 0.505 | 0.303 |
| Cortical moment of resistance (x-axis) | RX_CRT_A | 0.775 | 0.002 | 0.542 | 0.125 |
| Cortical moment of resistance (y-axis) | RY_CRT_A | 0.857 | 0.043 | 0.622 | 0.314 |
| Polar moment of resistance | RP_CRT_A | 0.898 | 0.004 | 0.753 | 0.197 |
| Cortical bone area (mm ²) | CRT_A | 0.893 | 0.000 | 0.680 | 0.185 |
| Cortical content per slice (mg / mm) | CRT_CNT | 0.437 | 0.189 | 0.510 | 0.647 |
| Cortical thickness standard deviation | CRT_THK_SD | 0.187 | 0.213 | 0.111 | 0.125 |
| Mean cortical thickness (mm) | CRT_THK | 0.125 | 0.169 | 0.124 | 0.111 |
| Endosteal circumference (via circular ring model) (mm) | ENDO_C | 0.867 | 0.962 | 0.606 | 0.966 |
| Endosteal circumference (mm) | ENDO | 0.315 | 0.147 | 0.403 | 0.696 |
| Total bone content per slice (mg / mm) | TOT_CNT | 0.364 | 0.422 | 0.448 | 0.704 |
| Trabecular bone area (mm ²) | TRAB_A | 0.061 | 0.088 | 0.762 | 0.331 |
| Trabecular bone content per slice (mg / mm) | TRAB_CNT | 0.384 | 0.901 | 0.448 | 0.704 |
| Trabecular bone density (mg / cm ³) | TRAB_DEN | 0.703 | 0.644 | 0.100 | 0.235 |

When considering gross mechanical properties of the bone, it is best to focus on cortical bone for its role in bearing loads and resisting fracture. Because of its ability to indicate bone health, BMD is also a factor of interest from these parameters. Therefore, taking also into

consideration that many MC3 fractures occur in the distal part of the bone, one metric that was hypothesized to be valuable in assessing a bone's susceptibility to fracture is cortical & subcortical BMD at 90% length (see Figure 3.21).

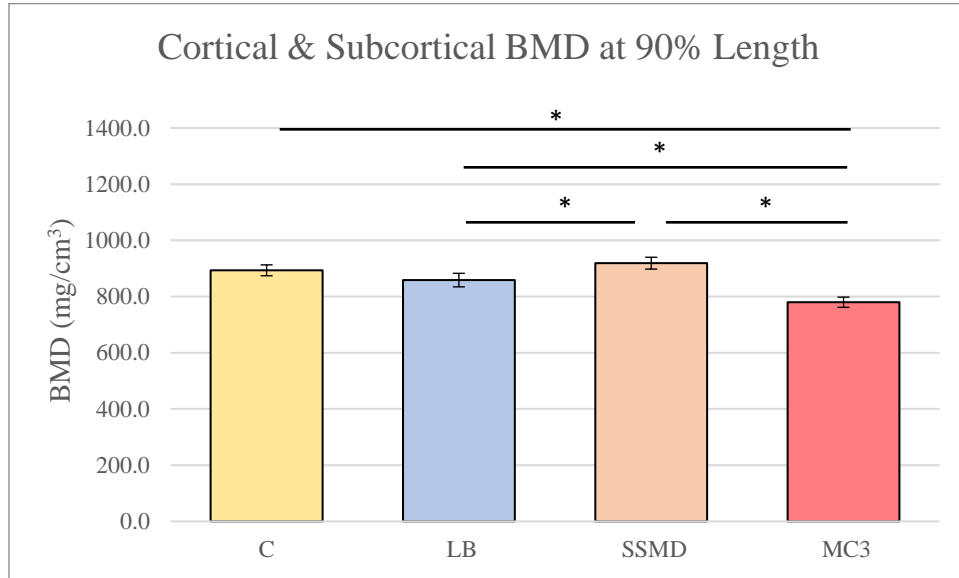


Figure 3.21. Bone mineral density in the cortical and subcortical bone significantly ($p < 0.05$) differs between the MC3 group and each other group, as well as in the SSMD group and LB group. ($n_C = 20$, $n_{LB} = 19$, $n_{SSMD} = 20$, $n_{MC3} = 5$). Bar and error bars represent mean \pm 1 standard deviation.

The MC3 group had significantly lower BMD than each other group, and the SSMD group also had higher BMD than the LB group. The Control group did not distinguish itself from any fracture group beside MC3.

3.5 Micro-Computed Tomography (μ CT)

Data from μ CT testing performed on bones from the Control and LB groups were compared via independent sample t-tests. With Control $n=10$ and LB $n=6$, a significant difference in bone mineral density was discovered at the dorsal surface ($p = 0.001$), but not at the lateral and medial surfaces ($p = 0.546$ and $p = 0.807$, respectively). See Figure 3.22.

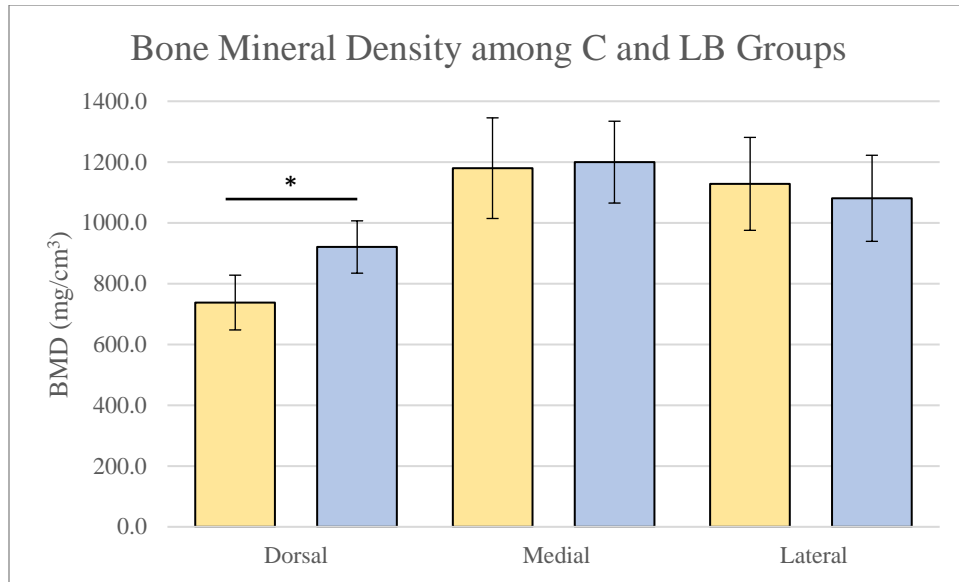


Figure 3.22. Only on the dorsal surface, the Control group (yellow) had a significantly ($p < 0.05$) lower BMD than the LB group (blue). ($n_C = 10$, $n_{LB} = 6$). Bar and error bars represent mean \pm 1 standard deviation. Independent samples t-test.

While evaluating μ CT images qualitatively, two irregularities were noted in select bones. In 5 out of 10 (50%) Control bones and 1 out of 6 (17%) of the LB group, a layer of low attenuation (i.e. darker coloring) is observed immediately below the surface of the bone (see Figure 3.23, left vs. center). In one bone from the LB group, the image has numerous dark spots within the cortex of the bone, indicating perhaps that the cortical bone in this region is porous (see Figure 3.23, right).

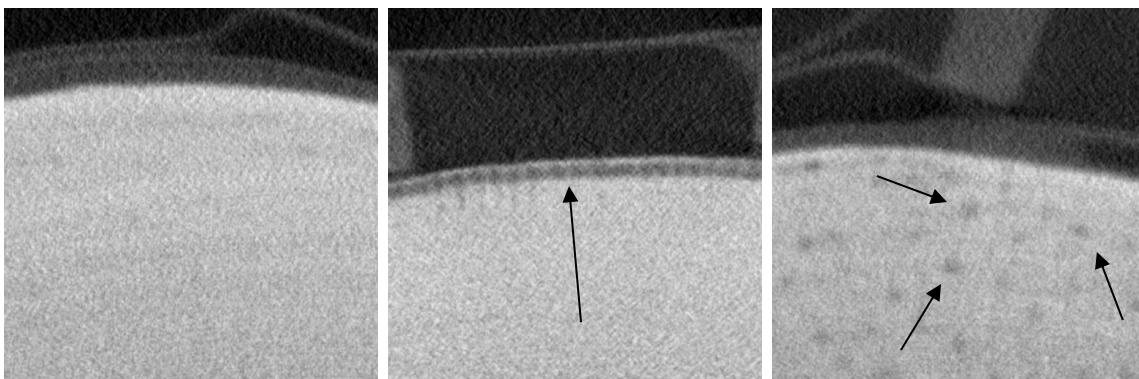


Figure 3.23. Left: a typical periosteal surface, seen here in a bone from the LB group. Center: a low-density strip is seen right below the bone surface, here in a bone from the Control group. Right: a bone apparently displaying abnormal porosity

3.6 Multivariable Model

3.6.1 Individual Analyses

In order to provide a different perspective than previous analyses that focused on difference in means, receiver operating characteristic (ROC) curves were created in MedCalc Statistical Software and their respective area under the curve (AUC) values were calculated to describe each parameter's ability to distinguish between bones from the Control group and those from the Fracture group. The ROC curves from four parameters that were deemed relevant at the 50% dorsal site are pictured in Figure 3.24.

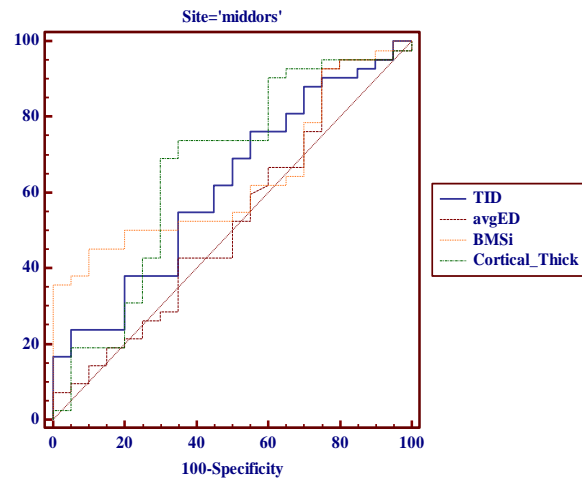


Figure 3.24. Individual parameters have low AUC values when modeled alone

Each individual parameter achieved an AUC above the minimum possible value of 0.50, though none achieved a value above 0.70. The corresponding AUC, 95% confidence interval, and standard error values are found in Table 3.3.

Table 3.3. The AUC, confidence interval, and SE values laid out for each individual parameter under consideration for the model. The low AUC values indicate that these variables don't have high capability to distinguish between healthy and at-risk bones. *Note: standard error not calculated for Average Loading Slope.*

| Parameter | AUC | 95% Confidence Interval | Standard Error |
|--------------------|-------|-------------------------|----------------|
| TID | 0.617 | 0.484 – 0.737 | 0.0768 |
| avgED | 0.527 | 0.396 – 0.655 | 0.0826 |
| BMSi | 0.637 | 0.505 – 0.755 | 0.0718 |
| Cortical Thickness | 0.663 | 0.532 – 0.778 | 0.0803 |
| ID1 | 0.621 | 0.489 – 0.742 | 0.0764 |
| US1 | 0.563 | 0.431 – 0.688 | 0.0830 |
| CID1 | 0.626 | 0.494 – 0.746 | 0.0731 |
| IDI | 0.581 | 0.449 – 0.705 | 0.0770 |
| avgCID | 0.577 | 0.445 – 0.702 | 0.0775 |
| avgUS | 0.555 | 0.423 – 0.681 | 0.0811 |
| avgLS | 0.526 | 0.395 – 0.654 | - |

When the same procedures were applied to a multivariable model containing parameters obtained data available at the time from x-rays, BioDent, and OsteoProbe at the 50% dorsal location, a more promising ROC curve was returned (Figure 3.25).

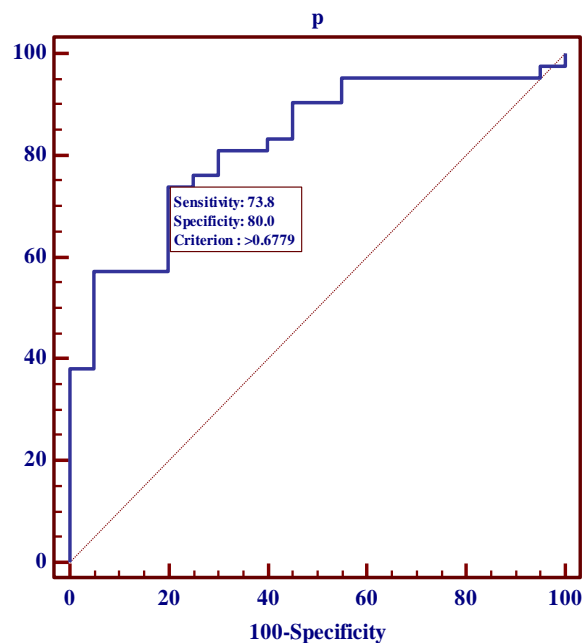


Figure 3.25. A model including TID, BMSi, avgED, and cortical thickness returns a fairly large AUC value of 0.82.

The AUC was found to be 0.82 with a 95% confidence interval of 0.71 – 0.91. The statistical optimal cutoff per the Youden index is at a specificity of 80.0 and sensitivity of 73.8, suggesting that the model has optimal distinguishing capacity at these levels.

When clinical practicality was taken into consideration, another model was created using only data from tests that can routinely be performed on live horses: BMSi, cortical thickness, and mass of the horse (Figure 3.26).

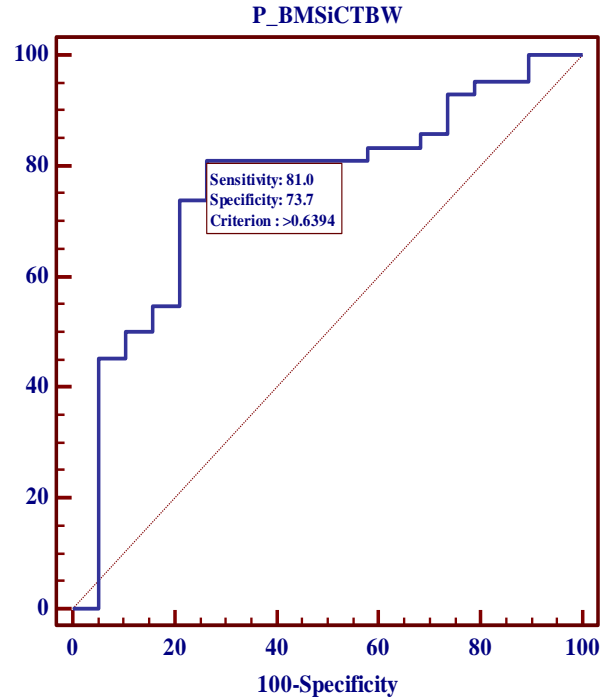


Figure 3.26. When only clinically relevant parameters (BMSi, cortical thickness, and mass) are included in the model, a returned AUC of 0.76 is still promising but not ideal

The AUC (0.761 with 95% confidence interval 0.634 – 0.861) was not as large as the previous model; however, it still suggests considerable distinguishing ability and reducing the number of predictor variables may help reduce the error inherent to the model.

Another model was created using three clinically-relevant parameters that were selected upon two primary criteria: uniqueness and utility. Cortical thickness at the 50% dorsal surface, bone mineral density of the cortex and subcortex at the 90% level, and the difference in percutaneous BMSi at the dorsolateral and medial surfaces at the 50% level. Each parameter displayed capability in distinguishing between experimental groups previously in the study, and each examined a unique property of the bone. Because data collected from the SSMD group appeared to be more

similar to the C group throughout this study, and because the nature of logistic regression models only allows for a dichotomous response, only the C group and the combined LB / MC3 group were included in this model to achieve maximum potential predictive power.

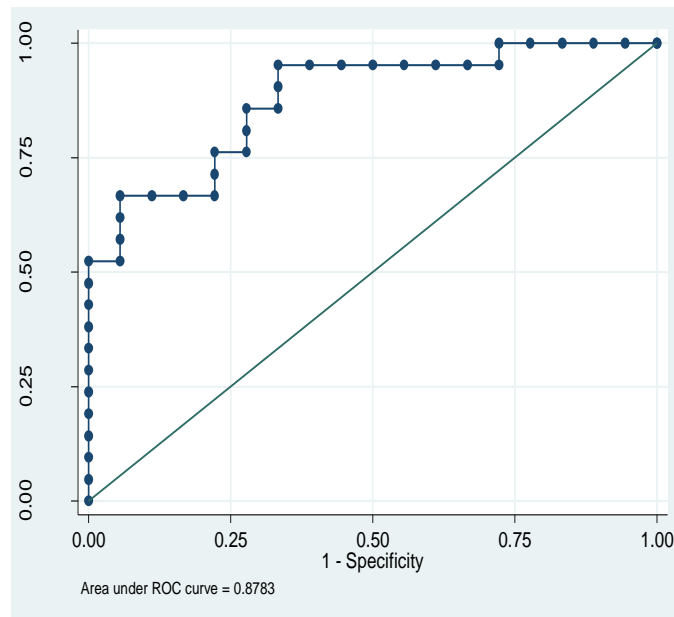


Figure 3.27. When the Control group and LB / MC3 group are included in the regression model with cortical thickness at the 50% dorsal surface, BMD at the 90% plane, and difference in BMSi between the dorsolateral and medial surfaces at 50% length, an area under the ROC curve of 0.88 is achieved.

This model returned an area under the ROC curve of 0.8783, with optimal sensitivity is 95.24% and specificity at 66.67%. Though the SSMD group is excluded from the model, it shows considerable promise in distinguishing between third metacarpal bones from horses that have experienced a long bone fracture and those that have not.

4. DISCUSSION

4.1 X-ray

Though prevalent differences in both geometry and fracture occurrence between right and left long bones in horses have been recorded [35][63][64], few discrepancies emerged between right and left cortical thicknesses at a given site within each group. As mentioned previously, the likelihood of type I errors increases when a high volume of hypothesis tests are performed concurrently; therefore, with the knowledge that the data's variance is large and that common controlling procedures would eliminate the significance of each observed difference, the mean of the data from each analogous location on right and left limbs was used in further analyses.

Cortical thickness was found to be significantly different between fracture groups along the dorsal surface of the bone, a clinically relevant site relating to the MC3's ability to adapt to exposure of compressive forces and strain [30]. Though the difference is indeed significant, it remains unclear how practical these findings will be in a clinical setting as the difference in means was on the order of 1 – 2 mm. Due to the potential for operator variation in the imaging and analysis of x-rays and the inherent high variability in MC3 geometry among racehorses [41], it is likely that the observed differences will have only marginal predictive applications when utilized in a univariate model. When considered in conjunction with data from other tests, however, it very well may bolster the predictive capability of the model.

4.2 OsteoProbe

Control bones did not distinguish themselves from any fracture group (or combination thereof) in pure BMSi measures through the skin or directly on the surface of the bone. However, the study provided insight to different test methods and potential differences in equine and human models.

The average standard deviation of measurements when taken percutaneously more than doubled that when taken on the same samples after having removed the skin and periosteum. While this doesn't disqualify the OsteoProbe as a potentially useful tool to clinically collect information on the material properties of bone, it does suggest that its capacity to distinguish between healthy and at-risk bones may be inversely related to its degree of noninvasiveness. Alternatively, it may

be that factors other than the bone surface itself (such as the periosteum, which has been shown to exhibit different properties in patients with and without skeletal pathologies [65]), help differentiate between healthy and pathologic bones. This may be supported by the differences seen between the medial-minus-dorsolateral differences between the Control group, LB group, and MC3 group. Although the standard deviations are strikingly high, the results are consistent with what is expected in the Control group: a large discrepancy between the dorsal surface, which presumably may have a layer of more easily-indented primary bone, and the medial and lateral surfaces, which are not thought to be undergoing modeling. While the fact that the opposite trend was seen in the MC3 group cannot be ignored, the immense standard deviation values compared to the sample size ($n = 5$) does render the data unreliable.

It may also be that the OsteoProbe is a more useful tool in certain situations than others. While our data exhibits BMSi trending lower in the control group than the fracture group, human studies have concluded the opposite in studies examining both pathology and response to loading [55,56,66]. We hypothesize that a lower BMSi on the dorsal surface—an area associated with exercise-induced bone modeling—may be due to the indentation probe encountering woven bone. Woven bone is characterized by disorderly structure and is mechanically weaker than mature, lamellar bone. While its presence on the surface of bone may reduce its resistance to indentation, it is also likely an indicator of healthy response to loading. A surface more resistant to indentation may reveal a lack of immature bone and therefore a potential dysregulation in response to exercise, which may help explain the occurrence of fracture experienced in these horses.

OsteoProbe, despite not displaying significant differences in comparing means between control and fracture groups, did greatly enhance the apparent predictive capability of a logistic regression model including x-ray data. While BMSi and cortical thickness returned ROC area under the curve (AUC) values of 0.637 and 0.663, respectively, individually, the two parameters taken together with mass created a model with an AUC of 0.76. Though this AUC would not generally be regarded by clinicians as robust enough to use as a reliable diagnostic method, it lends credence to the possibility of creating a predictive model from a combination of subtle changes that might otherwise go unnoticed.

4.3 BioDent

While no significant differences emerged when comparing control bones to the three fracture groups, some trends did appear. Total indentation distance, for example, was 28% greater in the control group than the MC3 group at the midshaft dorsal site, though statistical significance was likely barred by the small sample size of the MC3 group.

When the three fracture groups were combined into one and compared against the control group, differences along the dorsal surface persisted. At the 50% dorsal site, total indentation distance was 11% greater in the control group than the fracture group and indentation distance increase was 6% greater. At the 75% dorsal site, average energy dissipated (avgED) was significantly greater in the control group. avgED is calculated by measuring the area under the force-displacement curve generated during testing. Consistent with the aforementioned BioDent trends and the BMSi trends seen in OsteoProbe measurements, a greater displacement per unit force may suggest a layer on the surface of bone that is less resistant to indentation, as may be expected from woven bone deposited on the periosteal surface.

Though the BioDent elucidated a significant difference between the control and fracture groups where the OsteoProbe did not, the OsteoProbe is still considered to be the more clinically-relevant microindenter between the two. The correlations between BMSi and most BioDent parameters are modest, though the trends among experimental groups are similar. Replacing 2 BioDent parameters in the multivariable logistic regression model with mass—a much more accessible metric—resulted in only a minor decrease in diagnostic ability of the model, according to the area under their respective ROC curves.

4.4 pQCT

Among the host of parameters that pQCT measures, many had significant site * group interactions when the four experimental groups were compared against one another. Among these, certain parameters can intuitively be expected to contribute more to the multivariable model than others. For example, geometric measurements may be somewhat redundant in the model if cortical thickness derived from x-rays is included. Alternatively, however, were pQCT to provide more reliable insight to fracture susceptibility than x-ray, a metric such as cortical area may be used in place of x-ray data. Bone mineral density (BMD) is looked upon as a valuable contribution from

pQCT because of how distinct in nature it is from indentation and cortical thickness. Within the realm of BMD, cortical & subcortical measures at distal levels are deemed relevant due to the mechanical importance and common fracture incidence of these areas of bone.

It remains unclear whether pQCT should be classified as a clinically relevant tool. While standing pQCT has been successfully performed in standing horses before [67], it is not currently common practice and would likely be difficult to attain. Though it is not as practical as x-ray or OsteoProbe, it will not be disqualified from the potentially clinically applicable methodologies for the time being.

4.5 μ CT

μ CT testing was performed on a smaller and more selective sample size than the other test methods, but results were consistent with the running theory of bone modeling being evident on the dorsal surface of bones from horses without fracture diagnoses. Low BMD, in some cases, can be an indicator of pathology such as osteoporosis. While this may seem counterintuitive, as it was found that bones from the Control group had a lower surface BMD than bones from the fracture-afflicted group, the location of testing may play a significant role in interpreting the results. Just as the indentation testing and x-rays have suggested that modeling may be occurring on the dorsal surface of Control bones, the nature of μ CT testing seems to also detect this immature bone. Woven bone has reduced mineral content compared to mature lamellar bone, which may explain why the BMD at the dorsal periosteal surface in Control bones is lower than that of LB bones despite LB bones having significantly lower cortical BMD per the pQCT results in this study (Table A.24).

The low-density layers found adjacent to the surface on the affected bones remain to be identified with certainty. Once again, it may be the case that disorganized, comparatively low-mineral primary bone that forms in response to exercise is lining the dorsal surface of the bone, creating an area of low attenuation that appears as a black strip in the radiographic image. The porous appearance of the bone pictured in Figure 3.23, however, likely exhibited another metabolic mechanism. It is possible that the increased porosity may be due to the resorption phase of the remodeling process, in which case the afflicted horse may have been undergoing healthy adaptation in the bone tissue but was simply left vulnerable to fracture due to decreased bone density and volume. Alternatively, the apparent porosity may have been pathological, making fracture essentially imminent. While the possibility of the apparent spots being caused by signal

noise is not an impossibility, it is not being significantly considered given the surrounding portions of the image in conjunction with the other well-processed images in the set.

4.6 Multivariable Regression Model

The logistic regression analysis performed in this study was meant only to explore the field of predictive multivariable models, not to act as one. Predictive models of the sort for which we are aiming require statistical training sets much larger than our current sample size—depending on how many predictor variables and factors the model is to account for, hundreds of samples may be necessary. It is also worth noting that ROC curves and their corresponding AUC values are typically regarded as better predictors of diagnostic ability than comparison-of-mean tests such as t-tests or ANOVA. Because a large enough sample size could grant statistical significance to a difference on the order of a few osteons in a measure like cortical thickness, it is important not to draw any conclusions that are inappropriate for the nature of our testing.

At the same time, the results obtained from these preliminary models should not be discounted. Which predictive variables and potential covariates to use are yet to be optimized, and the working sample size is still relatively small. An AUC of 1.0 will never be achieved, but if the current model can be adjusted to a point where it can reliably predict the state of a bone, or the skeletal system as a whole, a vast majority of the time, it may find some important applications in equine or human fracture risk prediction.

When optimizing the model, sensitivity and specificity should also be considered. Horse owners, for example, may prefer a high sensitivity in the model, as allowing a period of rest may amount to a more economic decision than risking catastrophic breakdown and euthanasia.

4.7 Raman Spectroscopy and MRI

In a previous study that included a number of bones among this sample, Raman spectroscopy was used to determine the concentration of inorganic components in bone including phosphate, carbonate, and amide groups.

Raman spectroscopy was not utilized in this analysis due to the finding that bones exhibited different spectra than they did when measured in a previous study. Recent spectra suggest that bones have significantly higher carbonate than phosphate concentrations, when the opposite is

known to be true and has been displayed in previous analyses of the same bones. While undergoing freeze-thaw cycles may have an effect on the organic composition of bone, it is not expected to cause any changes in the representation of inorganic components.

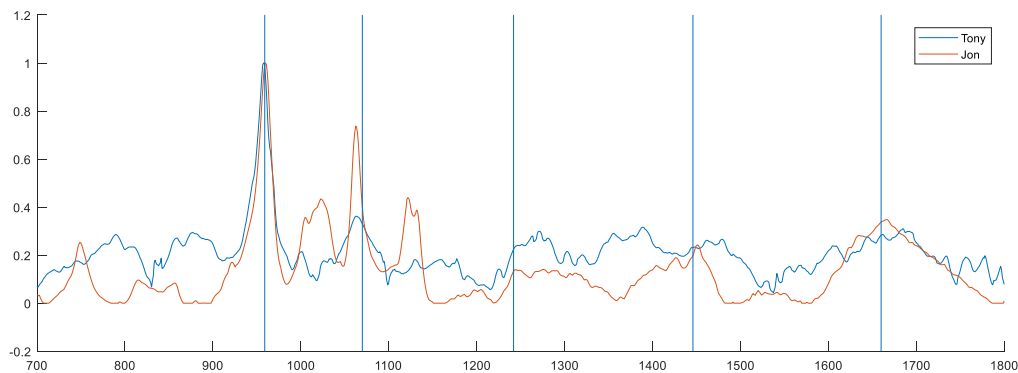


Figure 4.1. When previously obtained spectra (“Tony”) were compared with recent samples (“Jon”) and normalized by the height of the peak at $\sim 970\text{ cm}^{-1}$, other expected peaks (noted by vertical blue lines) were notably different between samples

A myriad of methods to remedy the discrepancies between previous and recent data were attempted. These revisions yielded spectra with distinct peaks at expected locations, though the magnitude of the peaks does not appear to be consistent with the known composition of bone.

A select number of bones in this study also underwent MRI testing. While cost and accessibility of MRI testing may exclude it from being considered as a clinically-relevant parameter in an equine model, it may be implemented in future iterations when human subjects are being considered.

4.8 Limitations

One conspicuous limitation of this study was the sample size of the MC3 experimental group. While grouping it together with the LB group didn’t appear to have any adverse effects on the results, this experiment could not properly compare this subset of fractures to the other experimental groups. The overall sample size was also a limitation to the logistic regression model. In order to move forward with the model, a substantially greater sample size in each of the relevant fracture groups must be collected.

The relatively high standard deviations among the x-ray data may have stemmed from multiple factors. Bone geometries varied substantially: the length of the bones being analyzed

ranged over 40 mm and standard deviations at certain sites within groups amounted to over 30% of the measurements themselves. Bone orientation during imaging may have also played a role, as small and potentially undetectable rotations of the bone would result in the projection of slightly different planes in the x-ray image. Additionally, human factors should be considered, as the calibration process allowed room for variability and the measurement process inherently relied on some judgement in distinguishing the interface between cortical and trabecular bone at the endosteal surface of the cortex.

While no significant results were discovered while using the OsteoProbe on a bare bone surface compared to when it was used percutaneously, the data collected percutaneously is less consistent. This tradeoff with clinical relevancy may be a drawback for the argument of the use of OsteoProbe in clinic.

5. CONCLUSIONS

Much was discovered in this study regarding which clinical and preclinical methods may be best suited for a predictive model and how to best focus future efforts. A lack of compelling evidence was found suggesting that any parameters significantly differ between left and right limbs, potentially eliminating the need for a dual-limb paradigm for each horse.

Cortical thickness, as determined by x-ray imaging, was found to be significantly different between the Control group and Fracture group at the mid- and distal-dorsal surfaces of third metacarpal bones. Bone material strength index (BMSi), while not significantly different between fracture groups, was found to potentially be able to detect primary bone deposition on the dorsal surface of healthy bones. BioDent measurements were able to detect that the average energy dissipated during cyclic indentation is higher in the Control group than in Fracture group on the distal dorsal surface, suggesting a less elastic surface more prone to permanent deformation. pQCT data suggested that overall cortical bone mineral density (BMD) may be lower in certain fracture groups than in the control group, and μ CT found a significantly lower BMD on the dorsal surface—but not lateral or medial surfaces—of bones in the Control group compared to bones in the Long Bone group. Some μ CT images show what appears to be immature primary bone on the surface of Control bones, which is consistent with other methodologies and suggests a healthy ability of the bone to adapt to training, which is consistent with the fracture history (or lack thereof) in this group.

While individual parameters exhibit poor ability to distinguish between control or fracture bones in a regression model, utilizing multiple parameters can establish a model with intriguing predictive potential. With a large sample size and an optimized set of independent variables, the prospect of a regression model with a clinically relevant ability to distinguish between healthy and at-risk bones appears promising.

6. FUTURE DIRECTIONS

The primary objective for the future of this project is to continue to increase the sample size. Once an adequate amount of data has been collected, a reliable predictive model can be created. In order to obtain the necessary sample size within practical bounds of time and cost, however, data will have to be collected more discriminately. A simple way to eliminate data collection resources by half is to only analyze one limb per horse. Given that the existing data does not exhibit any considerable differences between right and left limbs, a single-limb protocol can be implemented. Similarly, a majority of differences between the Control group and fracture groups are manifesting themselves at a select number of sites on the bone; if only these sites are tested, the process again becomes more efficient. Lastly, it may be prudent to only consider methodologies that are clinically relevant in either horses or humans. While data from sources such as Raman spectroscopy provide valuable insight to the composition of bones, the testing protocols are not at all conducive to or logistically feasible for live patients.

Considering the results from this study, it may be beneficial to move forward with the model without including the SSMD fracture group. Sesamoid bones are pointedly different from third metacarpals and other long bones, and sesamoid fractures may involve a distinct pathogenesis from their long bone counterparts. At many parameters and locations, bones from the SSMD group resemble the control bones more than the other two fracture groups; therefore, excluding them from the study may reveal more differences between bones in the Control group and those in the LB and MC3 groups. The inclusion of only the C, LB, and MC3 groups would also leave the regression model to distinguish only between bones from control horses and those from horses with long-bone fractures, which may lead to higher predictive values. To optimize which variables are included in the model, sensitivity analyses should be performed on each parameter collected thus far to establish which may have the best predictive capability when included in the model.

Additionally, a goal much further in the future is to translate a successful equine goal into one for humans. It will likely be most intuitive to transition the model from equine athletes to human athletes or soldiers, but being able to detect at-risk bones from pathologies such as osteoporosis are not out of the realm of this project in the long-term.

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APPENDIX

Horse Information List

Table A.1. Sex, age, mass, fracture group, and cause of death listed for each horse. (Sex: M = male, F = female, G = gelding, C = colt).

| Horse ID | Sex | Age (yrs) | Mass (kg) | Fx Group | Cause of Death |
|-----------|-----|-----------|-----------|----------|---|
| A13-13503 | M | 3 | 514 | C | Exercise-induced pulmonary embolism |
| A13-14144 | F | 4 | 410 | C | |
| A14-0419 | F | 3 | 398 | C | Diarrhea, Lethargy |
| A14-14702 | F | 5 | 488 | C | Possible ruptured aorta |
| A14-15808 | F | 4 | 425 | C | Ruptured right-front superior digital flexor tendon |
| A15-1229 | F | 3 | 485 | C | Right-hind fetlock laceration / soft tissue injury |
| A15-1432 | G | 5 | 518 | C | Severe osteoarthritis, right-front radiocarpal joint injury |
| A15-2920 | F | 3 | | C | Pneumonia, severe colitis |
| A15-4789 | F | 2 | 433 | C | Colic |
| A16-1177 | G | 4 | 458 | C | Exercise-induced pulmonary hemorrhage |
| A16-2118 | G | 4 | 432 | C | Enterocolitis |
| A16-2293 | M | 2 | 494 | C | Pastern joint luxation |
| A16-3336 | G | 3 | 532 | C | Sudden death at end of race |
| A17-3709 | G | 4 | 527 | C | Suspected colic |
| A18-15218 | F | 3 | 491 | C | Possible colic |
| A18-17727 | F | 3 | 527 | C | Sudden death at end of race |
| A18-584 | G | 3 | 450 | C | Right-hind hoof avulsion |
| A19-4449 | C | 4 | 503 | C | Laminitis |
| A19-5963 | G | 5 | 491 | C | Open dislocation of left carpus |
| A19-7032 | G | 3 | 548 | C | Suspected cardiovascular event |
| A14-1356 | F | 2 | 459 | LB | Left radius fracture |
| A14-15954 | G | 5 | 514 | LB | Left ulna olecranon tubercle fracture |
| A14-1818 | F | 3 | 525 | LB | Comminuted right carpus fracture |
| A15-13734 | F | 3 | 480 | LB | Comminuted right scapula fracture |
| A15-14441 | G | 3 | 515 | LB | Comminuted left tibia fracture |
| A15-4293 | F | 2 | 459 | LB | Right front long pastern bone fracture |
| A15-4869 | M | 4 | 493 | LB | Left radius fracture |
| A15-5258 | M | 3 | 517 | LB | 3rd and 4th carpal fractures |
| A16-16656 | F | 6 | 552 | LB | Scapula fracture |
| A16-2964 | G | 3 | 541 | LB | Scapula fracture |
| A16-647 | F | 3 | 503 | LB | Radial carpal bone fracture |

Table A.1, continued

| | | | | | |
|-----------|---|---|-----|------|---|
| A18-1274 | G | 3 | 500 | LB | Left scapula fracture |
| A18-15243 | G | 4 | 505 | LB | Left 3rd carpal bone fracture |
| A18-15426 | F | 3 | 495 | LB | Comminuted left humerus fracture |
| A18-3846 | G | 3 | 442 | LB | Left-front scapula fracture |
| A18-6100 | F | 4 | 450 | LB | Left humerus fracture |
| A19-16901 | F | 4 | 465 | LB | Right-hind third metatarsal & long pastern bone fractures |
| A19-4350 | G | 4 | 622 | LB | Comminuted left-front third carpal fracture |
| A19-998 | G | 3 | 476 | LB | Comminuted right scapula fracture |
| A13-13148 | F | 3 | 433 | SSMD | Both left-front sesamoid fracture |
| A14-0498 | F | 5 | 490 | SSMD | Both left-front sesamoid fracture |
| A14-14416 | M | 4 | 526 | SSMD | Both left-front sesamoid fracture |
| A14-15391 | F | 3 | 425 | SSMD | Both left-front sesamoid fracture |
| A14-15834 | G | 4 | 503 | SSMD | Both left-front sesamoid fracture |
| A14-1972 | G | 4 | 570 | SSMD | Both left-front sesamoid fracture |
| A14-3323 | M | 5 | 575 | SSMD | Both left-front sesamoid fracture |
| A14-4991 | F | 4 | 485 | SSMD | Both left-front sesamoid fracture |
| A14-4992 | G | 5 | 441 | SSMD | Both left-front sesamoid fracture |
| A16-1925 | F | 5 | 514 | SSMD | Both right-front sesamoid fracture |
| A16-2635 | G | 4 | 541 | SSMD | Both right-front sesamoid fracture |
| A16-9 | F | 6 | 463 | SSMD | Both left-front sesamoid fracture |
| A17-13458 | F | 4 | 468 | SSMD | Left-front sesamoid fracture, luxated fetlock |
| A17-18109 | G | 3 | 527 | SSMD | Both left-front sesamoid fracture |
| A17-5101 | G | 3 | 440 | SSMD | Left-front medial sesamoid fracture, flexor tendon rupture, DLS rupture |
| A19-1548 | G | 3 | 548 | SSMD | Both right-front sesamoid fracture |
| A19-1718 | G | 3 | 491 | SSMD | Both right-front sesamoid fracture |
| A19-2036 | F | 4 | 527 | SSMD | Both right-front sesamoid fracture |
| A19-4679 | F | 6 | 489 | SSMD | Left-front medial sesamoid fracture |
| A19-488 | G | 6 | 584 | SSMD | Both left-front sesamoid fracture |
| 391-187 | G | 5 | 480 | MC3 | Right condylar MC3 fracture |
| A14-14505 | G | 5 | 498 | MC3 | Left comminuted midshaft MC3 fracture |
| A15-4375 | F | 2 | 439 | MC3 | Left comminuted MC3 fracture |
| A17-14797 | G | 2 | 498 | MC3 | Left transverse MC3 fracture |
| A18-6468 | F | 3 | 481 | MC3 | Left comminuted midshaft MC3 fracture |

Mixed Linear Model Result Tables

X-ray

Table A.2. Cortical thickness was not found to be significant ($p < 0.05$) between groups or in site * group interactions.

Type III Tests of Fixed Effects^a

| Source | Numerator df | Denominator df | F | Sig. |
|--------------|--------------|----------------|----------|------|
| Intercept | 1 | 61.082 | 6359.164 | .000 |
| site | 11 | 665.502 | 148.665 | .000 |
| group | 3 | 61.175 | 1.226 | .308 |
| site * group | 33 | 665.587 | 1.282 | .136 |

a. Dependent Variable: thickness.

Table A.3. Cortical thickness (normalized by mass of horse) was not found to be significant ($p < 0.05$) between groups or in site * group interactions.

Type III Tests of Fixed Effects^a

| Source | Numerator df | Denominator df | F | Sig. |
|--------------|--------------|----------------|----------|------|
| Intercept | 1 | 60.062 | 3293.937 | .000 |
| site | 11 | 654.304 | 146.181 | .000 |
| group | 3 | 60.120 | 1.112 | .351 |
| site * group | 33 | 654.357 | 1.186 | .221 |

a. Dependent Variable: thickness_norm_mass.

Table A.4. Cortical thickness (normalized by length of MC3 bone) was not found to be significant ($p < 0.05$) between groups or in site * group interactions.

Type III Tests of Fixed Effects^a

| Source | Numerator df | Denominator df | F | Sig. |
|--------------|--------------|----------------|----------|------|
| Intercept | 1 | 61.058 | 6716.450 | .000 |
| site | 11 | 665.495 | 149.268 | .000 |
| group | 3 | 61.154 | 1.628 | .192 |
| site * group | 33 | 665.582 | 1.274 | .142 |

a. Dependent Variable: thickness_norm_length.

Table A.5. Cortical thickness was found to be significant ($p < 0.05$) in site * group interactions when the LB and MC3 experimental groups were combined.

Type III Tests of Fixed Effects^a

| Source | Numerator df | Denominator df | F | Sig. |
|--------------|--------------|----------------|----------|------|
| Intercept | 1 | 62.319 | 9199.826 | .000 |
| site | 11 | 676.729 | 213.207 | .000 |
| group | 2 | 62.304 | 1.708 | .190 |
| site * group | 22 | 676.708 | 1.560 | .049 |

a. Dependent Variable: thickness.

Table A.6. Cortical thickness was found to be significant ($p < 0.05$) in site * group interactions when the LB, SSMD, and MC3 experimental groups were combined and compared against the C group in the mixed linear model. Post-hoc analysis revealed significant differences at the 50% and 75% dorsal sites.

Type III Tests of Fixed Effects^a

| Source | Numerator df | Denominator df | F | Sig. |
|--------------|--------------|----------------|----------|------|
| Intercept | 1 | 62.976 | 7667.165 | .000 |
| site | 11 | 687.389 | 186.144 | .000 |
| group | 1 | 62.976 | .161 | .690 |
| site * group | 11 | 687.389 | 1.889 | .038 |

a. Dependent Variable: thickness.

OsteoProbe

Skin-on

Table A.7. BMSi was found to be significantly ($p < 0.05$) in site * group interactions.

Type III Tests of Fixed Effects^a

| Source | Numerator df | Denominator df | F | Sig. |
|--------------|--------------|----------------|----------|------|
| Intercept | 1 | 58.944 | 1579.594 | .000 |
| site | 11 | 625.195 | 6.167 | .000 |
| group | 3 | 59.143 | .573 | .635 |
| site * group | 33 | 625.456 | 1.957 | .001 |

a. Dependent Variable: bmsi.

Table A.8. BMSi was found to be significantly ($p < 0.05$) in site * group interactions when the LB and MC3 experimental groups were combined.

Type III Tests of Fixed Effects^a

| Source | Numerator df | Denominator df | F | Sig. |
|--------------|--------------|----------------|----------|------|
| Intercept | 1 | 60.475 | 2253.061 | .000 |
| site | 11 | 636.824 | 11.272 | .000 |
| group | 2 | 60.433 | .745 | .479 |
| site * group | 22 | 636.746 | 1.978 | .005 |

a. Dependent Variable: bmsi.

No Skin

Table A.9. BMSi was not found to be significant ($p < 0.05$) between groups or in site * group interactions.

Type III Tests of Fixed Effects^a

| Source | Numerator df | Denominator df | F | Sig. |
|--------------|--------------|----------------|-----------|------|
| Intercept | 1 | 60.000 | 19440.801 | .000 |
| site | 11 | 660.000 | 50.837 | .000 |
| group | 3 | 60 | 1.489 | .227 |
| site * group | 33 | 660.000 | 1.312 | .116 |

a. Dependent Variable: bmsi.

Table A.10. BMSi was not found to be significant ($p < 0.05$) between groups or in site * group interactions when the LB and MC3 experimental groups were combined.

Type III Tests of Fixed Effects^a

| Source | Numerator df | Denominator df | F | Sig. |
|--------------|--------------|----------------|-----------|------|
| Intercept | 1 | 61 | 27482.411 | .000 |
| site | 11 | 671.000 | 75.605 | .000 |
| group | 2 | 61 | 1.624 | .205 |
| site * group | 22 | 671.000 | 1.518 | .061 |

a. Dependent Variable: bmsi.

BioDent

Table A.11. Total Indentation Distance was not found to be significant ($p < 0.05$) between groups or in site * group interactions.

Type III Tests of Fixed Effects^a

| Source | Numerator df | Denominator df | F | Sig. |
|--------------|--------------|----------------|----------|------|
| Intercept | 1 | 63.209 | 2666.509 | .000 |
| site | 8 | 455.805 | 24.535 | .000 |
| group | 3 | 60.669 | .497 | .686 |
| site * group | 24 | 454.052 | .859 | .660 |

a. Dependent Variable: TID.

Table A.12. Indentation Distance Increase was not found to be significant ($p < 0.05$) between groups or in site * group interactions.

Type III Tests of Fixed Effects^a

| Source | Numerator df | Denominator df | F | Sig. |
|--------------|--------------|----------------|----------|------|
| Intercept | 1 | 63.927 | 1545.501 | .000 |
| site | 8 | 456.493 | 23.262 | .000 |
| group | 3 | 61.350 | 1.823 | .152 |
| site * group | 24 | 454.752 | .773 | .772 |

a. Dependent Variable: IDI.

Table A.13. Average Energy Dissipated was not found to be significant ($p < 0.05$) between groups or in site * group interactions.

Type III Tests of Fixed Effects^a

| Source | Numerator df | Denominator df | F | Sig. |
|--------------|--------------|----------------|----------|------|
| Intercept | 1 | 64.354 | 4163.483 | .000 |
| site | 8 | 456.941 | 37.704 | .000 |
| group | 3 | 61.752 | .746 | .529 |
| site * group | 24 | 455.206 | 1.103 | .336 |

a. Dependent Variable: avgED.

Table A.14. Initial Indentation Distance was not found to be significant ($p < 0.05$) between groups or in site * group interactions.

Type III Tests of Fixed Effects^a

| Source | Numerator df | Denominator df | F | Sig. |
|--------------|--------------|----------------|----------|------|
| Intercept | 1 | 62.973 | 2683.871 | .000 |
| site | 8 | 455.614 | 23.430 | .000 |
| group | 3 | 60.438 | .456 | .714 |
| site * group | 24 | 453.853 | .858 | .661 |

a. Dependent Variable: ID1.

Table A.15. Average Unloading Slope was not found to be significant ($p < 0.05$) between groups or in site * group interactions.

Type III Tests of Fixed Effects^a

| Source | Numerator df | Denominator df | F | Sig. |
|--------------|--------------|----------------|-----------|------|
| Intercept | 1 | 61.258 | 18926.768 | .000 |
| site | 8 | 453.257 | 14.346 | .000 |
| group | 3 | 59.564 | 1.026 | .388 |
| site * group | 24 | 451.972 | 1.189 | .246 |

a. Dependent Variable: avgUS.

Table A.16. 1st-Cycle Unloading Slope was not found to be significant ($p < 0.05$) between groups or in site * group interactions.

Type III Tests of Fixed Effects^a

| Source | Numerator df | Denominator df | F | Sig. |
|--------------|--------------|----------------|-----------|------|
| Intercept | 1 | 62.440 | 19112.689 | .000 |
| site | 8 | 454.145 | 16.735 | .000 |
| group | 3 | 60.626 | 1.549 | .211 |
| site * group | 24 | 452.816 | 1.377 | .112 |

a. Dependent Variable: US1.

Table A.17. 1st-Cycle Creep Indentation Distance was not found to be significant ($p < 0.05$) between groups or in site * group interactions.

Type III Tests of Fixed Effects^a

| Source | Numerator df | Denominator df | F | Sig. |
|--------------|--------------|----------------|----------|------|
| Intercept | 1 | 63.526 | 1986.372 | .000 |
| site | 8 | 456.114 | 30.539 | .000 |
| group | 3 | 60.968 | .934 | .430 |
| site * group | 24 | 454.366 | .892 | .614 |

a. Dependent Variable: CID1.

Table A.18. Average Creep Indentation Distance was not found to be significant ($p < 0.05$) between groups or in site * group interactions.

Type III Tests of Fixed Effects^a

| Source | Numerator df | Denominator df | F | Sig. |
|--------------|--------------|----------------|----------|------|
| Intercept | 1 | 63.598 | 3146.191 | .000 |
| site | 8 | 456.246 | 40.305 | .000 |
| group | 3 | 61.028 | .354 | .786 |
| site * group | 24 | 454.495 | .946 | .538 |

a. Dependent Variable: avgCID.

Table A.19. Average Loading Slope was not found to be significant ($p < 0.05$) between groups or in site * group interactions.

Type III Tests of Fixed Effects^a

| Source | Numerator df | Denominator df | F | Sig. |
|--------------|--------------|----------------|-----------|------|
| Intercept | 1 | 62.392 | 18424.770 | .000 |
| site | 8 | 454.674 | 44.277 | .000 |
| group | 3 | 60.295 | 1.915 | .137 |
| site * group | 24 | 453.160 | 1.140 | .295 |

a. Dependent Variable: avgLS.

Table A.20. BioDent summary table including all p-values for site, group, and site * group interactions when all experimental groups are included in the linear mixed model.

| Metric | Site | Group | Site * Group |
|----------------|-------------|--------------|---------------------|
| TID | 0.000 | 0.686 | 0.660 |
| IDI | 0.000 | 0.152 | 0.772 |
| avg ED | 0.000 | 0.529 | 0.336 |
| ID1 | 0.000 | 0.714 | 0.661 |
| avg US | 0.000 | 0.388 | 0.246 |
| US1 | 0.000 | 0.211 | 0.112 |
| CID1 | 0.000 | 0.430 | 0.614 |
| avg CID | 0.000 | 0.786 | 0.538 |
| avg LS | 0.000 | 0.137 | 0.295 |

Table A.21. BioDent summary table including all p-values for site, group, and site * group interactions when the LB and MC3 groups are combined and compared against the C and SSMD groups in the linear mixed model

| Metric | Site | Group | Site * Group |
|----------------|-------------|--------------|---------------------|
| TID | 0.000 | 0.666 | 0.622 |
| IDI | 0.000 | 0.073 | 0.655 |
| avg ED | 0.000 | 0.321 | 0.093 |
| ID1 | 0.000 | 0.721 | 0.593 |
| avg US | 0.000 | 0.348 | 0.212 |
| US1 | 0.000 | 0.153 | 0.212 |
| CID1 | 0.000 | 0.266 | 0.317 |
| avg CID | 0.000 | 0.586 | 0.231 |
| avg LS | 0.000 | 0.064 | 0.217 |

Table A.22. BioDent summary table including all p-values for site, group, and site * group interactions when all the LB, MC3, and SSMD groups are combined and compared against the C group in the linear mixed model

| Metric | Site | Group | Site * Group |
|---------------|-------------|--------------|---------------------|
| TID | 0.000 | 0.547 | 0.126 |
| IDI | 0.000 | 0.939 | 0.395 |
| avg ED | 0.000 | 0.433 | 0.046 |
| ID1 | 0.000 | 0.494 | 0.107 |

| | | | |
|----------------|-------|-------|-------|
| avg US | 0.000 | 0.368 | 0.359 |
| US1 | 0.000 | 0.500 | 0.330 |
| CID1 | 0.000 | 0.538 | 0.065 |
| avg CID | 0.000 | 0.884 | 0.095 |
| avg LS | 0.000 | 0.760 | 0.252 |

BioDent / OsteoProbe Correlations

Table A.23. R² values of no-skin BMSi compared with each BioDent parameter at each site. The 50% dorsal site, achieved by averaging together dorsolateral and dorsomedial BMSi measurements, consistently displays relatively high correlation values.

| | 25 Med | 50 Med | 75 Med | 25 Dors | 50 Dors | 75 Dors | 25 Lat | 50 Lat | 75 Lat |
|-------------------------------|---------------|---------------|---------------|----------------|-----------------|----------------|---------------|---------------|---------------|
| Initial Indentation Distance | 0.305862 | 0.075273 | 0.157515 | 0.283154 | 0.561892 | 0.250936 | 0.042826 | 0.002449 | 0.032099 |
| Total Indentation Distance | 0.320588 | 0.078343 | 0.145904 | 0.29009 | 0.582224 | 0.271427 | 0.062476 | 0.004231 | 0.031863 |
| Indentation Distance Increase | 0.246358 | 0.059795 | 0.080039 | 0.317028 | 0.537587 | 0.334848 | 0.131853 | 0.062217 | 0.034911 |
| Avg. CID | 0.349406 | 0.235461 | 0.145012 | 0.369369 | 0.575954 | 0.382922 | 0.233675 | 0.132688 | 0.075463 |
| Avg. ED | 0.227203 | 0.188592 | 0.02954 | 0.200816 | 0.446797 | 0.097634 | 0.050333 | 0.003218 | 0.016083 |
| Avg. US | 0.222036 | 0.225556 | 0.004757 | 0.018106 | 0.032446 | 0.003313 | 0.039977 | 0.05237 | 0.065617 |
| Avg. LS | 0.304272 | 0.291006 | 0.062505 | 0.086505 | 0.253977 | 0.09293 | 0.048363 | 0.027182 | 0.098403 |

Detailed pQCT Results

Table A.24. Group differences in cortical BMD. 1 = Control, 2 = LB, 3 = SSMD, 4 = MC3.

| Estimates ^a | | | | | |
|-----------------------------|---------|------------|--------|-------------------------|-------------|
| 1:CTRL, 2:LB, 3:SSMD, 4:MC3 | Mean | Std. Error | df | 95% Confidence Interval | |
| | | | | Lower Bound | Upper Bound |
| 1 | 969.171 | 5.297 | 61 | 958.579 | 979.763 |
| 2 | 947.783 | 5.297 | 61 | 937.191 | 958.376 |
| 3 | 971.834 | 5.297 | 61 | 961.242 | 982.426 |
| 4 | 922.329 | 10.594 | 61.000 | 901.145 | 943.514 |

a. Dependent Variable: CRT_DEN.

| Pairwise Comparisons ^a | | | | | | | |
|-----------------------------------|---------------------------------|-----------------------|------------|--------|-------------------|---|-------------|
| (I) 1:CTRL, 2:LB, 3:SSMD, 4:MC3 | (J) 1:CTRL, 2:LB, 3:SSMD, 4:MC3 | Mean Difference (I-J) | Std. Error | df | Sig. ^c | 95% Confidence Interval for Difference ^c | |
| | | | | | | Lower Bound | Upper Bound |
| 1 | 2 | 21.388* | 7.491 | 61 | .035 | .959 | 41.816 |
| | 3 | -2.663 | 7.491 | 61 | 1.000 | -23.092 | 17.766 |
| | 4 | 46.842* | 11.845 | 61.000 | .001 | 14.541 | 79.142 |
| 2 | 1 | -21.388* | 7.491 | 61 | .035 | -41.816 | -.959 |
| | 3 | -24.051* | 7.491 | 61 | .013 | -44.479 | -3.622 |
| | 4 | 25.454 | 11.845 | 61.000 | .214 | -6.846 | 57.755 |
| 3 | 1 | 2.663 | 7.491 | 61 | 1.000 | -17.766 | 23.092 |
| | 2 | 24.051* | 7.491 | 61 | .013 | 3.622 | 44.479 |
| | 4 | 49.505* | 11.845 | 61.000 | .001 | 17.204 | 81.805 |
| 4 | 1 | -46.842* | 11.845 | 61.000 | .001 | -79.142 | -14.541 |
| | 2 | -25.454 | 11.845 | 61.000 | .214 | -57.755 | 6.846 |
| | 3 | -49.505* | 11.845 | 61.000 | .001 | -81.805 | -17.204 |

Based on estimated marginal means

*. The mean difference is significant at the .05 level.

a. Dependent Variable: CRT_DEN.

c. Adjustment for multiple comparisons: Bonferroni.

Table A.25. Group differences in cortical and subcortical mineral BMC. 1 = Control, 2 = LB, 3 = SSMD, 4 = MC3.

| Estimates ^a | | | | | |
|-----------------------------|---------|------------|----|-------------------------|-------------|
| 1:CTRL, 2:LB, 3:SSMD, 4:MC3 | Mean | Std. Error | df | 95% Confidence Interval | |
| | | | | Lower Bound | Upper Bound |
| 1 | 794.596 | 15.278 | 61 | 764.047 | 825.146 |
| 2 | 760.486 | 15.278 | 61 | 729.937 | 791.036 |
| 3 | 806.557 | 15.278 | 61 | 776.008 | 837.107 |
| 4 | 712.293 | 30.555 | 61 | 651.194 | 773.392 |

a. Dependent Variable: CRTSUB_CNT.

| Pairwise Comparisons ^a | | | | | | | |
|-----------------------------------|---------------------------------|-----------------------|------------|----|-------------------|---|-------------|
| (I) 1:CTRL, 2:LB, 3:SSMD, 4:MC3 | (J) 1:CTRL, 2:LB, 3:SSMD, 4:MC3 | Mean Difference (I-J) | Std. Error | df | Sig. ^c | 95% Confidence Interval for Difference ^c | |
| | | | | | | Lower Bound | Upper Bound |
| 1 | 2 | 34.110 | 21.606 | 61 | .717 | -24.809 | 93.029 |
| | 3 | -11.961 | 21.606 | 61 | 1.000 | -70.880 | 46.958 |
| | 4 | 82.303 | 34.162 | 61 | .114 | -10.856 | 175.463 |
| 2 | 1 | -34.110 | 21.606 | 61 | .717 | -93.029 | 24.809 |
| | 3 | -46.071 | 21.606 | 61 | .222 | -104.990 | 12.848 |
| | 4 | 48.193 | 34.162 | 61 | .980 | -44.966 | 141.353 |
| 3 | 1 | 11.961 | 21.606 | 61 | 1.000 | -46.958 | 70.880 |
| | 2 | 46.071 | 21.606 | 61 | .222 | -12.848 | 104.990 |
| | 4 | 94.264* | 34.162 | 61 | .046 | 1.105 | 187.424 |
| 4 | 1 | -82.303 | 34.162 | 61 | .114 | -175.463 | 10.856 |
| | 2 | -48.193 | 34.162 | 61 | .980 | -141.353 | 44.966 |
| | 3 | -94.264* | 34.162 | 61 | .046 | -187.424 | -1.105 |

Based on estimated marginal means

*. The mean difference is significant at the .05 level.

a. Dependent Variable: CRTSUB_CNT.

c. Adjustment for multiple comparisons: Bonferroni.

Table A.26. Group differences in cortical and subcortical BMD. 1 = Control, 2 = LB, 3 = SSMD, 4 = MC3.

| Estimates^a | | | | | |
|------------------------------|----------|------------|--------|-------------------------|-------------|
| 1:CTRL, 2:LB, 3:SSMD, 4:MC3 | Mean | Std. Error | df | 95% Confidence Interval | |
| | | | | Lower Bound | Upper Bound |
| 1 | 1114.151 | 5.439 | 61.000 | 1103.275 | 1125.028 |
| 2 | 1105.256 | 5.439 | 61.000 | 1094.379 | 1116.133 |
| 3 | 1121.785 | 5.439 | 61.000 | 1110.908 | 1132.661 |
| 4 | 1080.226 | 10.879 | 61 | 1058.472 | 1101.979 |

a. Dependent Variable: CRTSUB_DEN.

| Pairwise Comparisons^a | | | | | | | |
|---|---------------------------------|-----------------------|------------|--------|-------------------|---|-------------|
| (I) 1:CTRL, 2:LB, 3:SSMD, 4:MC3 | (J) 1:CTRL, 2:LB, 3:SSMD, 4:MC3 | Mean Difference (I-J) | Std. Error | df | Sig. ^c | 95% Confidence Interval for Difference ^c | |
| | | | | | | Lower Bound | Upper Bound |
| 1 | 2 | 8.895 | 7.692 | 61.000 | 1.000 | -12.082 | 29.873 |
| | 3 | -7.633 | 7.692 | 61.000 | 1.000 | -28.611 | 13.344 |
| | 4 | 33.926* | 12.163 | 61 | .042 | .757 | 67.094 |
| 2 | 1 | -8.895 | 7.692 | 61.000 | 1.000 | -29.873 | 12.082 |
| | 3 | -16.528 | 7.692 | 61.000 | .214 | -37.506 | 4.449 |
| | 4 | 25.030 | 12.163 | 61 | .263 | -8.138 | 58.199 |
| 3 | 1 | 7.633 | 7.692 | 61.000 | 1.000 | -13.344 | 28.611 |
| | 2 | 16.528 | 7.692 | 61.000 | .214 | -4.449 | 37.506 |
| | 4 | 41.559* | 12.163 | 61 | .007 | 8.390 | 74.727 |
| 4 | 1 | -33.926* | 12.163 | 61 | .042 | -67.094 | -7.757 |
| | 2 | -25.030 | 12.163 | 61 | .263 | -58.199 | 8.138 |
| | 3 | -41.559* | 12.163 | 61 | .007 | -74.727 | -8.390 |

Based on estimated marginal means

*. The mean difference is significant at the .05 level.

a. Dependent Variable: CRTSUB_DEN.

c. Adjustment for multiple comparisons: Bonferroni.

Table A.27. Group differences in total BMD. 1 = Control, 2 = LB, 3 = SSMD, 4 = MC3.

| Estimates ^a | | | | | |
|-----------------------------|---------|------------|--------|-------------------------|-------------|
| 1:CTRL, 2:LB, 3:SSMD, 4:MC3 | Mean | Std. Error | df | 95% Confidence Interval | |
| | | | | Lower Bound | Upper Bound |
| 1 | 885.170 | 7.490 | 61 | 870.191 | 900.148 |
| 2 | 862.322 | 7.490 | 61.000 | 847.344 | 877.301 |
| 3 | 884.962 | 7.490 | 61.000 | 869.984 | 899.940 |
| 4 | 832.537 | 14.981 | 61 | 802.580 | 862.493 |

a. Dependent Variable: TOT_DEN.

| Pairwise Comparisons ^a | | | | | | | |
|-----------------------------------|---------------------------------|-----------------------|------------|--------|-------------------|---|-------------|
| (I) 1:CTRL, 2:LB, 3:SSMD, 4:MC3 | (J) 1:CTRL, 2:LB, 3:SSMD, 4:MC3 | Mean Difference (I-J) | Std. Error | df | Sig. ^c | 95% Confidence Interval for Difference ^c | |
| | | | | | | Lower Bound | Upper Bound |
| 1 | 2 | 22.847 | 10.593 | 61.000 | .210 | -6.041 | 51.735 |
| | 3 | .207 | 10.593 | 61.000 | 1.000 | -28.680 | 29.095 |
| | 4 | 52.633 [*] | 16.749 | 61 | .016 | 6.958 | 98.308 |
| 2 | 1 | -22.847 | 10.593 | 61.000 | .210 | -51.735 | 6.041 |
| | 3 | -22.640 | 10.593 | 61.000 | .220 | -51.527 | 6.248 |
| | 4 | 29.786 | 16.749 | 61 | .482 | -15.889 | 75.461 |
| 3 | 1 | -.207 | 10.593 | 61.000 | 1.000 | -29.095 | 28.680 |
| | 2 | 22.640 | 10.593 | 61.000 | .220 | -6.248 | 51.527 |
| | 4 | 52.426 [*] | 16.749 | 61 | .016 | 6.750 | 98.101 |
| 4 | 1 | -52.633 [*] | 16.749 | 61 | .016 | -98.308 | -6.958 |
| | 2 | -29.786 | 16.749 | 61 | .482 | -75.461 | 15.889 |
| | 3 | -52.426 [*] | 16.749 | 61 | .016 | -98.101 | -6.750 |

Based on estimated marginal means

*. The mean difference is significant at the .05 level.

a. Dependent Variable: TOT_DEN.

c. Adjustment for multiple comparisons: Bonferroni.

Table A.28. Site * group interactions in cortical BMD. 1 = Control, 2 = LB, 3 = SSMD, 4 = MC3.

| | | | Mean Difference (I-J) | | | | 95% Confidence Interval for Difference ^a | |
|------|---------------------------------|---------------------------------|-----------------------|------------|---------|-------------------|---|-------------|
| site | (I) 1:CTRL, 2:LB, 3:SSMD, 4:MC3 | (J) 1:CTRL, 2:LB, 3:SSMD, 4:MC3 | | Std. Error | df | Sig. ^c | Lower Bound | Upper Bound |
| 10 | 1 | 2 | 46.096 [*] | 13.506 | 284.412 | .004 | 10.212 | 81.981 |
| | | 3 | 15.195 | 13.506 | 284.412 | 1.000 | -20.690 | 51.079 |
| | | 4 | 48.435 | 21.356 | 284.412 | .144 | -8.304 | 105.174 |
| | 2 | 1 | -46.096 [*] | 13.506 | 284.412 | .004 | -81.981 | -10.212 |
| | | 3 | -30.902 | 13.506 | 284.412 | .137 | -66.786 | 4.983 |
| | | 4 | 2.339 | 21.356 | 284.412 | 1.000 | -54.400 | 59.077 |
| | 3 | 1 | -15.195 | 13.506 | 284.412 | 1.000 | -51.079 | 20.690 |
| | | 2 | 30.902 | 13.506 | 284.412 | .137 | -4.983 | 66.786 |
| | | 4 | 33.240 | 21.356 | 284.412 | .724 | -23.498 | 89.979 |
| | 4 | 1 | -48.435 | 21.356 | 284.412 | .144 | -105.174 | 8.304 |
| | | 2 | -2.339 | 21.356 | 284.412 | 1.000 | -59.077 | 54.400 |
| | | 3 | -33.240 | 21.356 | 284.412 | .724 | -89.979 | 23.498 |
| 25 | 1 | 2 | 3.510 | 13.506 | 284.412 | 1.000 | -32.375 | 39.395 |
| | | 3 | -1.260 | 13.506 | 284.412 | 1.000 | -37.145 | 34.625 |
| | | 4 | 13.650 | 21.356 | 284.412 | 1.000 | -43.089 | 70.389 |
| | 2 | 1 | -3.510 | 13.506 | 284.412 | 1.000 | -39.395 | 32.375 |
| | | 3 | -4.770 | 13.506 | 284.412 | 1.000 | -40.655 | 31.115 |
| | | 4 | 10.140 | 21.356 | 284.412 | 1.000 | -46.599 | 66.879 |
| | 3 | 1 | 1.260 | 13.506 | 284.412 | 1.000 | -34.625 | 37.145 |
| | | 2 | 4.770 | 13.506 | 284.412 | 1.000 | -31.115 | 40.655 |
| | | 4 | 14.910 | 21.356 | 284.412 | 1.000 | -41.829 | 71.649 |
| | 4 | 1 | -13.650 | 21.356 | 284.412 | 1.000 | -70.389 | 43.089 |
| | | 2 | -10.140 | 21.356 | 284.412 | 1.000 | -66.879 | 46.599 |
| | | 3 | -14.910 | 21.356 | 284.412 | 1.000 | -71.649 | 41.829 |
| 50 | 1 | 2 | -1.225 | 13.506 | 284.412 | 1.000 | -37.110 | 34.660 |
| | | 3 | -5.080 | 13.506 | 284.412 | 1.000 | -40.965 | 30.805 |
| | | 4 | 8.950 | 21.356 | 284.412 | 1.000 | -47.789 | 65.689 |
| | 2 | 1 | 1.225 | 13.506 | 284.412 | 1.000 | -34.660 | 37.110 |
| | | 3 | -3.855 | 13.506 | 284.412 | 1.000 | -39.740 | 32.030 |
| | | 4 | 10.175 | 21.356 | 284.412 | 1.000 | -46.564 | 66.914 |
| | 3 | 1 | 5.080 | 13.506 | 284.412 | 1.000 | -30.805 | 40.965 |
| | | 2 | 3.855 | 13.506 | 284.412 | 1.000 | -32.030 | 39.740 |
| | | 4 | 14.030 | 21.356 | 284.412 | 1.000 | -42.709 | 70.769 |
| | 4 | 1 | -8.950 | 21.356 | 284.412 | 1.000 | -65.689 | 47.789 |
| | | 2 | -10.175 | 21.356 | 284.412 | 1.000 | -66.914 | 46.564 |
| | | 3 | -14.030 | 21.356 | 284.412 | 1.000 | -70.769 | 42.709 |
| 75 | 1 | 2 | 25.755 | 13.506 | 284.412 | .345 | -10.130 | 61.640 |
| | | 3 | -22.470 | 13.506 | 284.412 | .584 | -58.355 | 13.415 |
| | | 4 | 82.255 [*] | 21.356 | 284.412 | .001 | 25.516 | 138.994 |
| | 2 | 1 | -25.755 | 13.506 | 284.412 | .345 | -61.640 | 10.130 |
| | | 3 | -48.225 [*] | 13.506 | 284.412 | .003 | -84.110 | -12.340 |
| | | 4 | 56.500 | 21.356 | 284.412 | .052 | -.239 | 113.239 |
| | 3 | 1 | 22.470 | 13.506 | 284.412 | .584 | -13.415 | 58.355 |
| | | 2 | 48.225 [*] | 13.506 | 284.412 | .003 | 12.340 | 84.110 |
| | | 4 | 104.725 [*] | 21.356 | 284.412 | .000 | 47.986 | 161.464 |
| | 4 | 1 | -82.255 [*] | 21.356 | 284.412 | .001 | -138.994 | -25.516 |
| | | 2 | -56.500 | 21.356 | 284.412 | .052 | -113.239 | -.239 |
| | | 3 | -104.725 [*] | 21.356 | 284.412 | .000 | -161.464 | -47.986 |
| 90 | 1 | 2 | 32.802 | 13.506 | 284.412 | .095 | -3.083 | 68.686 |
| | | 3 | .300 | 13.506 | 284.412 | 1.000 | -35.585 | 36.185 |
| | | 4 | 80.919 [*] | 21.356 | 284.412 | .001 | 24.180 | 137.657 |
| | 2 | 1 | -32.802 | 13.506 | 284.412 | .095 | -68.686 | 3.083 |
| | | 3 | -32.502 | 13.506 | 284.412 | .101 | -68.386 | 3.383 |
| | | 4 | 48.117 | 21.356 | 284.412 | .150 | -8.621 | 104.856 |
| | 3 | 1 | -.300 | 13.506 | 284.412 | 1.000 | -36.185 | 35.585 |
| | | 2 | 32.502 | 13.506 | 284.412 | .101 | -3.383 | 68.386 |
| | | 4 | 80.619 [*] | 21.356 | 284.412 | .001 | 23.880 | 137.357 |
| | 4 | 1 | -80.919 [*] | 21.356 | 284.412 | .001 | -137.657 | -24.180 |
| | | 2 | -48.117 | 21.356 | 284.412 | .150 | -104.856 | 8.621 |
| | | 3 | -80.619 [*] | 21.356 | 284.412 | .001 | -137.357 | -23.880 |

Based on estimated marginal means

*. The mean difference is significant at the .05 level.

a. Dependent Variable: CRT_DEN.

c. Adjustment for multiple comparisons: Bonferroni.

Table A.29. Site * group interactions in cortical and subcortical BMC. 1 = Control, 2 = LB, 3 = SSMD, 4 = MC3.

| | | | | Mean Difference (I-J) | | | | 95% Confidence Interval for Difference ^a | |
|------|---------------------------------|---------------------------------|--|-----------------------|------------|---------|-------------------|---|-------------|
| site | (I) 1:CTRL, 2:LB, 3:SSMD, 4:MC3 | (J) 1:CTRL, 2:LB, 3:SSMD, 4:MC3 | | | Std. Error | df | Sig. ^c | Lower Bound | Upper Bound |
| 10 | 1 | 2 | | 64.597 | 26.030 | 122.277 | .087 | -5.211 | 134.405 |
| | | 3 | | 10.780 | 26.030 | 122.277 | 1.000 | -59.028 | 80.588 |
| | | 4 | | 100.398 | 41.156 | 122.277 | .097 | -9.977 | 210.774 |
| | 2 | 1 | | -64.597 | 26.030 | 122.277 | .087 | -134.405 | 5.211 |
| | | 3 | | -53.817 | 26.030 | 122.277 | .245 | -123.625 | 15.991 |
| | | 4 | | 35.801 | 41.156 | 122.277 | 1.000 | -74.574 | 146.177 |
| | 3 | 1 | | -10.780 | 26.030 | 122.277 | 1.000 | -80.588 | 59.028 |
| | | 2 | | 53.817 | 26.030 | 122.277 | .245 | -15.991 | 123.625 |
| | | 4 | | 89.618 | 41.156 | 122.277 | .188 | -20.757 | 199.994 |
| | 4 | 1 | | -100.398 | 41.156 | 122.277 | .097 | -210.774 | 9.977 |
| | | 2 | | -35.801 | 41.156 | 122.277 | 1.000 | -146.177 | 74.574 |
| | | 3 | | -89.618 | 41.156 | 122.277 | .188 | -199.994 | 20.757 |
| 25 | 1 | 2 | | 11.555 | 26.030 | 122.277 | 1.000 | -58.253 | 81.363 |
| | | 3 | | -7.736 | 26.030 | 122.277 | 1.000 | -77.543 | 62.072 |
| | | 4 | | 54.216 | 41.156 | 122.277 | 1.000 | -56.160 | 164.591 |
| | 2 | 1 | | -11.555 | 26.030 | 122.277 | 1.000 | -81.363 | 58.253 |
| | | 3 | | -19.291 | 26.030 | 122.277 | 1.000 | -89.098 | 50.517 |
| | | 4 | | 42.661 | 41.156 | 122.277 | 1.000 | -67.715 | 153.036 |
| | 3 | 1 | | 7.736 | 26.030 | 122.277 | 1.000 | -62.072 | 77.543 |
| | | 2 | | 19.291 | 26.030 | 122.277 | 1.000 | -50.517 | 89.098 |
| | | 4 | | 61.951 | 41.156 | 122.277 | .809 | -48.425 | 172.327 |
| | 4 | 1 | | -54.216 | 41.156 | 122.277 | 1.000 | -164.591 | 56.160 |
| | | 2 | | -42.661 | 41.156 | 122.277 | 1.000 | -153.036 | 67.715 |
| | | 3 | | -61.951 | 41.156 | 122.277 | .809 | -172.327 | 48.425 |
| 50 | 1 | 2 | | 35.466 | 26.030 | 122.277 | 1.000 | -34.342 | 105.274 |
| | | 3 | | -1.069 | 26.030 | 122.277 | 1.000 | -70.877 | 68.739 |
| | | 4 | | 83.946 | 41.156 | 122.277 | .261 | -26.430 | 194.322 |
| | 2 | 1 | | -35.466 | 26.030 | 122.277 | 1.000 | -105.274 | 34.342 |
| | | 3 | | -36.535 | 26.030 | 122.277 | .978 | -106.343 | 33.273 |
| | | 4 | | 48.480 | 41.156 | 122.277 | 1.000 | -61.896 | 158.856 |
| | 3 | 1 | | 1.069 | 26.030 | 122.277 | 1.000 | -68.739 | 70.877 |
| | | 2 | | 36.535 | 26.030 | 122.277 | .978 | -33.273 | 106.343 |
| | | 4 | | 85.015 | 41.156 | 122.277 | .246 | -25.361 | 195.391 |
| | 4 | 1 | | -83.946 | 41.156 | 122.277 | .261 | -194.322 | 26.430 |
| | | 2 | | -48.480 | 41.156 | 122.277 | 1.000 | -158.856 | 61.896 |
| | | 3 | | -85.015 | 41.156 | 122.277 | .246 | -195.391 | 25.361 |
| 75 | 1 | 2 | | 54.863 | 26.030 | 122.277 | .223 | -14.945 | 124.671 |
| | | 3 | | -37.756 | 26.030 | 122.277 | .897 | -107.564 | 32.052 |
| | | 4 | | 133.295 [*] | 41.156 | 122.277 | .009 | 22.919 | 243.670 |
| | 2 | 1 | | -54.863 | 26.030 | 122.277 | .223 | -124.671 | 14.945 |
| | | 3 | | -92.619 [*] | 26.030 | 122.277 | .003 | -162.427 | -22.811 |
| | | 4 | | 78.432 | 41.156 | 122.277 | .354 | -31.944 | 188.807 |
| | 3 | 1 | | 37.756 | 26.030 | 122.277 | .897 | -32.052 | 107.564 |
| | | 2 | | 92.619 [*] | 26.030 | 122.277 | .003 | 22.811 | 162.427 |
| | | 4 | | 171.051 [*] | 41.156 | 122.277 | .000 | 60.675 | 281.426 |
| | 4 | 1 | | -133.295 [*] | 41.156 | 122.277 | .009 | -243.670 | -22.919 |
| | | 2 | | -78.432 | 41.156 | 122.277 | .354 | -188.807 | 31.944 |
| | | 3 | | -171.051 [*] | 41.156 | 122.277 | .000 | -281.426 | -60.675 |
| 90 | 1 | 2 | | 4.069 | 26.030 | 122.277 | 1.000 | -65.739 | 73.877 |
| | | 3 | | -24.024 | 26.030 | 122.277 | 1.000 | -93.832 | 45.783 |
| | | 4 | | 39.663 | 41.156 | 122.277 | 1.000 | -70.713 | 150.038 |
| | 2 | 1 | | -4.069 | 26.030 | 122.277 | 1.000 | -73.877 | 65.739 |
| | | 3 | | -28.093 | 26.030 | 122.277 | 1.000 | -97.901 | 41.714 |
| | | 4 | | 35.594 | 41.156 | 122.277 | 1.000 | -74.782 | 145.969 |
| | 3 | 1 | | 24.024 | 26.030 | 122.277 | 1.000 | -45.783 | 93.832 |
| | | 2 | | 28.093 | 26.030 | 122.277 | 1.000 | -41.714 | 97.901 |
| | | 4 | | 63.687 | 41.156 | 122.277 | .746 | -46.689 | 174.063 |
| | 4 | 1 | | -39.663 | 41.156 | 122.277 | 1.000 | -150.038 | 70.713 |
| | | 2 | | -35.594 | 41.156 | 122.277 | 1.000 | -145.969 | 74.782 |
| | | 3 | | -63.687 | 41.156 | 122.277 | .746 | -174.063 | 46.689 |

Based on estimated marginal means

*. The mean difference is significant at the .05 level.

a. Dependent Variable: CRTSUB_CNT.

c. Adjustment for multiple comparisons: Bonferroni.

Table A.30. Site * group interactions in cortical and subcortical BMD. 1 = Control, 2 = LB, 3 = SSMD, 4 = MC3.

| | | | Mean Difference (I-J) | | | | 95% Confidence Interval for Difference ^a | |
|------|---------------------------------|---------------------------------|-----------------------|------------|---------|-------------------|---|-------------|
| site | (I) 1:CTRL, 2:LB, 3:SSMD, 4:MC3 | (J) 1:CTRL, 2:LB, 3:SSMD, 4:MC3 | | Std. Error | df | Sig. ^c | Lower Bound | Upper Bound |
| 10 | 1 | 2 | 4.814 | 13.951 | 285.668 | 1.000 | -32.249 | 41.877 |
| | | 3 | -.995 | 13.951 | 285.668 | 1.000 | -38.058 | 36.068 |
| | | 4 | 6.969 | 22.058 | 285.668 | 1.000 | -51.633 | 65.571 |
| | 2 | 1 | -4.814 | 13.951 | 285.668 | 1.000 | -41.877 | 32.249 |
| | | 3 | -5.808 | 13.951 | 285.668 | 1.000 | -42.872 | 31.255 |
| | | 4 | 2.155 | 22.058 | 285.668 | 1.000 | -56.447 | 60.757 |
| | 3 | 1 | .995 | 13.951 | 285.668 | 1.000 | -36.068 | 38.058 |
| | | 2 | 5.808 | 13.951 | 285.668 | 1.000 | -31.255 | 42.872 |
| | | 4 | 7.963 | 22.058 | 285.668 | 1.000 | -50.639 | 66.565 |
| | 4 | 1 | -6.969 | 22.058 | 285.668 | 1.000 | -65.571 | 51.633 |
| | | 2 | -2.155 | 22.058 | 285.668 | 1.000 | -60.757 | 56.447 |
| | | 3 | -7.963 | 22.058 | 285.668 | 1.000 | -66.565 | 50.639 |
| 25 | 1 | 2 | 3.510 | 13.951 | 285.668 | 1.000 | -33.553 | 40.573 |
| | | 3 | -1.260 | 13.951 | 285.668 | 1.000 | -38.323 | 35.803 |
| | | 4 | 13.650 | 22.058 | 285.668 | 1.000 | -44.952 | 72.252 |
| | 2 | 1 | -3.510 | 13.951 | 285.668 | 1.000 | -40.573 | 33.553 |
| | | 3 | -4.770 | 13.951 | 285.668 | 1.000 | -41.833 | 32.293 |
| | | 4 | 10.140 | 22.058 | 285.668 | 1.000 | -48.462 | 68.742 |
| | 3 | 1 | 1.260 | 13.951 | 285.668 | 1.000 | -35.803 | 38.323 |
| | | 2 | 4.770 | 13.951 | 285.668 | 1.000 | -32.293 | 41.833 |
| | | 4 | 14.910 | 22.058 | 285.668 | 1.000 | -43.692 | 73.512 |
| | 4 | 1 | -13.650 | 22.058 | 285.668 | 1.000 | -72.252 | 44.952 |
| | | 2 | -10.140 | 22.058 | 285.668 | 1.000 | -68.742 | 48.462 |
| | | 3 | -14.910 | 22.058 | 285.668 | 1.000 | -73.512 | 43.692 |
| 50 | 1 | 2 | -1.225 | 13.951 | 285.668 | 1.000 | -38.288 | 35.838 |
| | | 3 | -5.080 | 13.951 | 285.668 | 1.000 | -42.143 | 31.983 |
| | | 4 | 8.950 | 22.058 | 285.668 | 1.000 | -49.652 | 67.552 |
| | 2 | 1 | 1.225 | 13.951 | 285.668 | 1.000 | -35.838 | 38.288 |
| | | 3 | -3.855 | 13.951 | 285.668 | 1.000 | -40.918 | 33.208 |
| | | 4 | 10.175 | 22.058 | 285.668 | 1.000 | -48.427 | 68.777 |
| | 3 | 1 | 5.080 | 13.951 | 285.668 | 1.000 | -31.983 | 42.143 |
| | | 2 | 3.855 | 13.951 | 285.668 | 1.000 | -33.208 | 40.918 |
| | | 4 | 14.030 | 22.058 | 285.668 | 1.000 | -44.572 | 72.632 |
| | 4 | 1 | -8.950 | 22.058 | 285.668 | 1.000 | -67.552 | 49.652 |
| | | 2 | -10.175 | 22.058 | 285.668 | 1.000 | -68.777 | 48.427 |
| | | 3 | -14.030 | 22.058 | 285.668 | 1.000 | -72.632 | 44.572 |
| 75 | 1 | 2 | 2.720 | 13.951 | 285.668 | 1.000 | -34.343 | 39.783 |
| | | 3 | -5.240 | 13.951 | 285.668 | 1.000 | -42.303 | 31.823 |
| | | 4 | 26.600 | 22.058 | 285.668 | 1.000 | -32.002 | 85.202 |
| | 2 | 1 | -2.720 | 13.951 | 285.668 | 1.000 | -39.783 | 34.343 |
| | | 3 | -7.960 | 13.951 | 285.668 | 1.000 | -45.023 | 29.103 |
| | | 4 | 23.880 | 22.058 | 285.668 | 1.000 | -34.722 | 82.482 |
| | 3 | 1 | 5.240 | 13.951 | 285.668 | 1.000 | -31.823 | 42.303 |
| | | 2 | 7.960 | 13.951 | 285.668 | 1.000 | -29.103 | 45.023 |
| | | 4 | 31.840 | 22.058 | 285.668 | .900 | -26.762 | 90.442 |
| | 4 | 1 | -26.600 | 22.058 | 285.668 | 1.000 | -85.202 | 32.002 |
| | | 2 | -23.880 | 22.058 | 285.668 | 1.000 | -82.482 | 34.722 |
| | | 3 | -31.840 | 22.058 | 285.668 | .900 | -90.442 | 26.762 |
| 90 | 1 | 2 | 34.657 | 13.951 | 285.668 | .081 | -2.406 | 71.720 |
| | | 3 | -25.591 | 13.951 | 285.668 | .406 | -62.655 | 11.472 |
| | | 4 | 113.459 ^a | 22.058 | 285.668 | .000 | 54.857 | 172.061 |
| | 2 | 1 | -34.657 | 13.951 | 285.668 | .081 | -71.720 | 2.406 |
| | | 3 | -60.248 ^a | 13.951 | 285.668 | .000 | -97.311 | -23.185 |
| | | 4 | 78.802 ^a | 22.058 | 285.668 | .002 | 20.200 | 137.404 |
| | 3 | 1 | 25.591 | 13.951 | 285.668 | .406 | -11.472 | 62.655 |
| | | 2 | 60.248 ^a | 13.951 | 285.668 | .000 | 23.185 | 97.311 |
| | | 4 | 139.050 ^a | 22.058 | 285.668 | .000 | 80.448 | 197.652 |
| | 4 | 1 | -113.459 ^a | 22.058 | 285.668 | .000 | -172.061 | -54.857 |
| | | 2 | -78.802 ^a | 22.058 | 285.668 | .002 | -137.404 | -20.200 |
| | | 3 | -139.050 ^a | 22.058 | 285.668 | .000 | -197.652 | -80.448 |

Based on estimated marginal means

*. The mean difference is significant at the .05 level.

a. Dependent Variable: CRTSUB_DEN.

c. Adjustment for multiple comparisons: Bonferroni.

Table A.31. Site * group interactions in total BMD. 1 = Control, 2 = LB, 3 = SSMD, 4 = MC3.

| site | (I) 1:CTRL, 2:LB, 3:SSMD, 4:MC3 | (J) 1:CTRL, 2:LB, 3:SSMD, 4:MC3 | Mean Difference (I-J) | Std. Error | df | Sig. ^a | 95% Confidence Interval for Difference ^b | |
|------|---------------------------------|---------------------------------|-----------------------|------------|---------|-------------------|---|-------------|
| | | | | | | | Lower Bound | Upper Bound |
| 10 | 1 | 2 | 47.819 ^c | 14.918 | 193.214 | .009 | 8.054 | 87.585 |
| | | 3 | 16.203 | 14.918 | 193.214 | 1.000 | -23.562 | 55.969 |
| | | 4 | 50.476 | 23.587 | 193.214 | .202 | -12.399 | 113.352 |
| | 2 | 1 | -47.819 ^c | 14.918 | 193.214 | .009 | -87.585 | -8.054 |
| | | 3 | -31.616 | 14.918 | 193.214 | .212 | -71.382 | 8.150 |
| | | 4 | 2.657 | 23.587 | 193.214 | 1.000 | -60.218 | 65.532 |
| | 3 | 1 | -16.203 | 14.918 | 193.214 | 1.000 | -55.969 | 23.562 |
| | | 2 | 31.616 | 14.918 | 193.214 | .212 | -8.150 | 71.382 |
| | | 4 | 34.273 | 23.587 | 193.214 | .887 | -28.602 | 97.148 |
| | 4 | 1 | -50.476 | 23.587 | 193.214 | .202 | -113.352 | 12.399 |
| | | 2 | -2.657 | 23.587 | 193.214 | 1.000 | -65.532 | 60.218 |
| | | 3 | -34.273 | 23.587 | 193.214 | .887 | -97.148 | 28.602 |
| 25 | 1 | 2 | 6.240 | 14.918 | 193.214 | 1.000 | -33.526 | 46.006 |
| | | 3 | 3.970 | 14.918 | 193.214 | 1.000 | -35.796 | 43.736 |
| | | 4 | 17.935 | 23.587 | 193.214 | 1.000 | -44.940 | 80.810 |
| | 2 | 1 | -6.240 | 14.918 | 193.214 | 1.000 | -46.006 | 33.526 |
| | | 3 | -2.270 | 14.918 | 193.214 | 1.000 | -42.036 | 37.496 |
| | | 4 | 11.695 | 23.587 | 193.214 | 1.000 | -51.180 | 74.570 |
| | 3 | 1 | -3.970 | 14.918 | 193.214 | 1.000 | -43.736 | 35.796 |
| | | 2 | 2.270 | 14.918 | 193.214 | 1.000 | -37.496 | 42.036 |
| | | 4 | 13.965 | 23.587 | 193.214 | 1.000 | -48.910 | 76.840 |
| | 4 | 1 | -17.935 | 23.587 | 193.214 | 1.000 | -80.810 | 44.940 |
| | | 2 | -11.695 | 23.587 | 193.214 | 1.000 | -74.570 | 51.180 |
| | | 3 | -13.965 | 23.587 | 193.214 | 1.000 | -76.840 | 48.910 |
| 50 | 1 | 2 | 1.910 | 14.918 | 193.214 | 1.000 | -37.856 | 41.676 |
| | | 3 | .960 | 14.918 | 193.214 | 1.000 | -38.806 | 40.726 |
| | | 4 | 37.065 | 23.587 | 193.214 | .706 | -25.810 | 99.940 |
| | 2 | 1 | -1.910 | 14.918 | 193.214 | 1.000 | -41.676 | 37.856 |
| | | 3 | -.950 | 14.918 | 193.214 | 1.000 | -40.716 | 38.816 |
| | | 4 | 35.155 | 23.587 | 193.214 | .826 | -27.720 | 98.030 |
| | 3 | 1 | -.960 | 14.918 | 193.214 | 1.000 | -40.726 | 38.806 |
| | | 2 | .950 | 14.918 | 193.214 | 1.000 | -38.816 | 40.716 |
| | | 4 | 36.105 | 23.587 | 193.214 | .765 | -26.770 | 98.980 |
| | 4 | 1 | -37.065 | 23.587 | 193.214 | .706 | -99.940 | 25.810 |
| | | 2 | -35.155 | 23.587 | 193.214 | .826 | -98.030 | 27.720 |
| | | 3 | -36.105 | 23.587 | 193.214 | .765 | -98.980 | 26.770 |
| 75 | 1 | 2 | 27.205 | 14.918 | 193.214 | .418 | -12.561 | 66.971 |
| | | 3 | -22.125 | 14.918 | 193.214 | .838 | -61.891 | 17.641 |
| | | 4 | 85.695 ^c | 23.587 | 193.214 | .002 | 22.820 | 148.570 |
| | 2 | 1 | -27.205 | 14.918 | 193.214 | .418 | -66.971 | 12.561 |
| | | 3 | -49.330 ^c | 14.918 | 193.214 | .007 | -89.096 | -9.564 |
| | | 4 | 58.490 | 23.587 | 193.214 | .084 | -4.385 | 121.365 |
| | 3 | 1 | 22.125 | 14.918 | 193.214 | .838 | -17.641 | 61.891 |
| | | 2 | 49.330 ^c | 14.918 | 193.214 | .007 | 9.564 | 89.096 |
| | | 4 | 107.820 ^c | 23.587 | 193.214 | .000 | 44.945 | 170.695 |
| | 4 | 1 | -85.695 ^c | 23.587 | 193.214 | .002 | -148.570 | -22.820 |
| | | 2 | -58.490 | 23.587 | 193.214 | .084 | -121.365 | 4.385 |
| | | 3 | -107.820 ^c | 23.587 | 193.214 | .000 | -170.695 | -44.945 |
| 90 | 1 | 2 | 31.061 | 14.918 | 193.214 | .232 | -8.705 | 70.827 |
| | | 3 | 2.028 | 14.918 | 193.214 | 1.000 | -37.737 | 41.794 |
| | | 4 | 71.993 ^c | 23.587 | 193.214 | .016 | 9.118 | 134.869 |
| | 2 | 1 | -31.061 | 14.918 | 193.214 | .232 | -70.827 | 8.705 |
| | | 3 | -29.032 | 14.918 | 193.214 | .319 | -68.798 | 10.734 |
| | | 4 | 40.933 | 23.587 | 193.214 | .506 | -21.943 | 103.808 |
| | 3 | 1 | -2.028 | 14.918 | 193.214 | 1.000 | -41.794 | 37.737 |
| | | 2 | 29.032 | 14.918 | 193.214 | .319 | -10.734 | 68.798 |
| | | 4 | 69.965 ^c | 23.587 | 193.214 | .020 | 7.090 | 132.840 |
| | 4 | 1 | -71.993 ^c | 23.587 | 193.214 | .016 | -134.869 | -9.118 |
| | | 2 | -40.933 | 23.587 | 193.214 | .506 | -103.808 | 21.943 |
| | | 3 | -69.965 ^c | 23.587 | 193.214 | .020 | -132.840 | -7.090 |

Based on estimated marginal means

*. The mean difference is significant at the .05 level.

a. Dependent Variable: TOT_DEN.

c. Adjustment for multiple comparisons: Bonferroni.

Table A.32. Site * group interactions in axial area moment of inertia. 1 = Control, 2 = LB, 3 = SSMD, 4 = MC3.

| | | | Mean Difference (I-J) | | | | 95% Confidence Interval for Difference ^c | |
|------|---------------------------------|---------------------------------|-------------------------|------------|---------|-------------------|---|-------------|
| site | (I) 1:CTRL, 2:LB, 3:SSMD, 4:MC3 | (J) 1:CTRL, 2:LB, 3:SSMD, 4:MC3 | | Std. Error | df | Sig. ^c | Lower Bound | Upper Bound |
| 10 | 1 | 2 | -3771.700 | 10279.633 | 257.400 | 1.000 | -31103.226 | 23559.826 |
| | | 3 | -3484.599 | 10279.633 | 257.400 | 1.000 | -30816.126 | 23846.927 |
| | | 4 | 12217.317 | 16253.526 | 257.400 | 1.000 | -30997.621 | 55432.255 |
| | 2 | 1 | 3771.700 | 10279.633 | 257.400 | 1.000 | -23559.826 | 31103.226 |
| | | 3 | 287.101 | 10279.633 | 257.400 | 1.000 | -27044.426 | 27618.627 |
| | | 4 | 15989.017 | 16253.526 | 257.400 | 1.000 | -27225.921 | 59203.955 |
| | 3 | 1 | 3484.599 | 10279.633 | 257.400 | 1.000 | -23846.927 | 30816.126 |
| | | 2 | -287.101 | 10279.633 | 257.400 | 1.000 | -27618.627 | 27044.426 |
| | | 4 | 15701.916 | 16253.526 | 257.400 | 1.000 | -27513.021 | 58916.854 |
| | 4 | 1 | -12217.317 | 16253.526 | 257.400 | 1.000 | -55432.255 | 30997.621 |
| | | 2 | -15989.017 | 16253.526 | 257.400 | 1.000 | -59203.955 | 27225.921 |
| | | 3 | -15701.916 | 16253.526 | 257.400 | 1.000 | -58916.854 | 27513.021 |
| 25 | 1 | 2 | 474.700 | 10279.633 | 257.400 | 1.000 | -26856.826 | 27806.226 |
| | | 3 | -1999.850 | 10279.633 | 257.400 | 1.000 | -29331.376 | 25331.676 |
| | | 4 | 5726.450 | 16253.526 | 257.400 | 1.000 | -37488.488 | 48941.388 |
| | 2 | 1 | -474.700 | 10279.633 | 257.400 | 1.000 | -27806.226 | 26856.826 |
| | | 3 | -2474.550 | 10279.633 | 257.400 | 1.000 | -29806.076 | 24856.976 |
| | | 4 | 5251.750 | 16253.526 | 257.400 | 1.000 | -37963.188 | 48466.688 |
| | 3 | 1 | 1999.850 | 10279.633 | 257.400 | 1.000 | -25331.676 | 29331.376 |
| | | 2 | 2474.550 | 10279.633 | 257.400 | 1.000 | -24856.976 | 29806.076 |
| | | 4 | 7726.300 | 16253.526 | 257.400 | 1.000 | -35488.638 | 50941.238 |
| | 4 | 1 | -5726.450 | 16253.526 | 257.400 | 1.000 | -48941.388 | 37488.488 |
| | | 2 | -5251.750 | 16253.526 | 257.400 | 1.000 | -48466.688 | 37963.188 |
| | | 3 | -7726.300 | 16253.526 | 257.400 | 1.000 | -50941.238 | 35488.638 |
| 50 | 1 | 2 | 4770.800 | 10279.633 | 257.400 | 1.000 | -22560.726 | 32102.326 |
| | | 3 | 80.450 | 10279.633 | 257.400 | 1.000 | -27251.076 | 27411.976 |
| | | 4 | 7905.300 | 16253.526 | 257.400 | 1.000 | -35309.638 | 51120.238 |
| | 2 | 1 | -4770.800 | 10279.633 | 257.400 | 1.000 | -32102.326 | 22560.726 |
| | | 3 | -4690.350 | 10279.633 | 257.400 | 1.000 | -32021.876 | 22641.176 |
| | | 4 | 3134.500 | 16253.526 | 257.400 | 1.000 | -40080.438 | 46349.438 |
| | 3 | 1 | -80.450 | 10279.633 | 257.400 | 1.000 | -27411.976 | 27251.076 |
| | | 2 | 4690.350 | 10279.633 | 257.400 | 1.000 | -22641.176 | 32021.876 |
| | | 4 | 7824.850 | 16253.526 | 257.400 | 1.000 | -35390.088 | 51039.788 |
| | 4 | 1 | -7905.300 | 16253.526 | 257.400 | 1.000 | -51120.238 | 35309.638 |
| | | 2 | -3134.500 | 16253.526 | 257.400 | 1.000 | -46349.438 | 40080.438 |
| | | 3 | -7824.850 | 16253.526 | 257.400 | 1.000 | -51039.788 | 35390.088 |
| 75 | 1 | 2 | 2261.100 | 10279.633 | 257.400 | 1.000 | -25070.426 | 29592.626 |
| | | 3 | 1904.600 | 10279.633 | 257.400 | 1.000 | -25426.926 | 29236.126 |
| | | 4 | 3347.800 | 16253.526 | 257.400 | 1.000 | -39867.138 | 46562.738 |
| | 2 | 1 | -2261.100 | 10279.633 | 257.400 | 1.000 | -29592.626 | 25070.426 |
| | | 3 | -356.500 | 10279.633 | 257.400 | 1.000 | -27688.026 | 26975.026 |
| | | 4 | 1086.700 | 16253.526 | 257.400 | 1.000 | -42128.238 | 44301.638 |
| | 3 | 1 | -1904.600 | 10279.633 | 257.400 | 1.000 | -29236.126 | 25426.926 |
| | | 2 | 356.500 | 10279.633 | 257.400 | 1.000 | -26975.026 | 27688.026 |
| | | 4 | 1443.200 | 16253.526 | 257.400 | 1.000 | -41771.738 | 44658.138 |
| | 4 | 1 | -3347.800 | 16253.526 | 257.400 | 1.000 | -46562.738 | 39867.138 |
| | | 2 | -1086.700 | 16253.526 | 257.400 | 1.000 | -44301.638 | 42128.238 |
| | | 3 | -1443.200 | 16253.526 | 257.400 | 1.000 | -44658.138 | 41771.738 |
| 90 | 1 | 2 | -25804.108 | 10279.633 | 257.400 | .076 | -53135.634 | 1527.419 |
| | | 3 | 2366.934 | 10279.633 | 257.400 | 1.000 | -24964.593 | 29698.460 |
| | | 4 | -58758.550 [*] | 16253.526 | 257.400 | .002 | -101973.488 | -15543.613 |
| | 2 | 1 | 25804.108 | 10279.633 | 257.400 | .076 | -1527.419 | 53135.634 |
| | | 3 | 28171.041 [*] | 10279.633 | 257.400 | .039 | 839.515 | 55502.568 |
| | | 4 | -32954.443 | 16253.526 | 257.400 | .262 | -76169.380 | 10260.495 |
| | 3 | 1 | -2366.934 | 10279.633 | 257.400 | 1.000 | -29698.460 | 24964.593 |
| | | 2 | -28171.041 [*] | 10279.633 | 257.400 | .039 | -55502.568 | -839.515 |
| | | 4 | -61125.484 [*] | 16253.526 | 257.400 | .001 | -104340.422 | -17910.546 |
| | 4 | 1 | 58758.550 [*] | 16253.526 | 257.400 | .002 | 15543.613 | 101973.488 |
| | | 2 | 32954.443 | 16253.526 | 257.400 | .262 | -10260.495 | 76169.380 |
| | | 3 | 61125.484 [*] | 16253.526 | 257.400 | .001 | 17910.546 | 104340.422 |

Based on estimated marginal means

*. The mean difference is significant at the .05 level.

a. Dependent Variable: I_CIRC.

c. Adjustment for multiple comparisons: Bonferroni.

Table A.33. Site * group interactions in total bone area. 1 = Control, 2 = LB, 3 = SSMD, 4 = MC3.

| site | (I) 1:CTRL, 2:LB, 3:SSMD, 4:MC3 | (J) 1:CTRL, 2:LB, 3:SSMD, 4:MC3 | Mean Difference (I-J) | Std. Error | df | Sig. ^a | 95% Confidence Interval for Difference ^b | |
|------|---------------------------------|---------------------------------|-----------------------|------------|---------|-------------------|---|---------|
| 10 | 1 | 2 | -17.180 | 44.280 | 210.473 | 1.000 | -135.117 | 100.757 |
| | | 3 | -16.070 | 44.280 | 210.473 | 1.000 | -134.007 | 101.867 |
| | | 4 | 64.890 | 70.013 | 210.473 | 1.000 | -121.585 | 251.364 |
| | 2 | 1 | 17.180 | 44.280 | 210.473 | 1.000 | -100.757 | 135.117 |
| | | 3 | 1.110 | 44.280 | 210.473 | 1.000 | -116.827 | 119.047 |
| | | 4 | 82.070 | 70.013 | 210.473 | 1.000 | -104.405 | 268.545 |
| | 3 | 1 | 16.070 | 44.280 | 210.473 | 1.000 | -101.867 | 134.007 |
| | | 2 | -1.110 | 44.280 | 210.473 | 1.000 | -119.047 | 116.827 |
| | | 4 | 80.960 | 70.013 | 210.473 | 1.000 | -105.515 | 267.434 |
| | 4 | 1 | -64.890 | 70.013 | 210.473 | 1.000 | -251.364 | 121.585 |
| | | 2 | -82.070 | 70.013 | 210.473 | 1.000 | -268.545 | 104.405 |
| | | 3 | -80.960 | 70.013 | 210.473 | 1.000 | -267.434 | 105.515 |
| 25 | 1 | 2 | 5.280 | 44.280 | 210.473 | 1.000 | -112.657 | 123.217 |
| | | 3 | -12.280 | 44.280 | 210.473 | 1.000 | -130.217 | 105.657 |
| | | 4 | 36.794 | 70.013 | 210.473 | 1.000 | -149.681 | 223.269 |
| | 2 | 1 | -5.280 | 44.280 | 210.473 | 1.000 | -123.217 | 112.657 |
| | | 3 | -17.560 | 44.280 | 210.473 | 1.000 | -135.497 | 100.377 |
| | | 4 | 31.514 | 70.013 | 210.473 | 1.000 | -154.961 | 217.989 |
| | 3 | 1 | 12.280 | 44.280 | 210.473 | 1.000 | -105.657 | 130.217 |
| | | 2 | 17.560 | 44.280 | 210.473 | 1.000 | -100.377 | 135.497 |
| | | 4 | 49.074 | 70.013 | 210.473 | 1.000 | -137.401 | 235.549 |
| | 4 | 1 | -36.794 | 70.013 | 210.473 | 1.000 | -223.269 | 149.681 |
| | | 2 | -31.514 | 70.013 | 210.473 | 1.000 | -217.989 | 154.961 |
| | | 3 | -49.074 | 70.013 | 210.473 | 1.000 | -235.549 | 137.401 |
| 50 | 1 | 2 | 31.634 | 44.280 | 210.473 | 1.000 | -86.303 | 149.571 |
| | | 3 | .934 | 44.280 | 210.473 | 1.000 | -117.003 | 118.871 |
| | | 4 | 44.812 | 70.013 | 210.473 | 1.000 | -141.663 | 231.287 |
| | 2 | 1 | -31.634 | 44.280 | 210.473 | 1.000 | -149.571 | 86.303 |
| | | 3 | -30.700 | 44.280 | 210.473 | 1.000 | -148.637 | 87.237 |
| | | 4 | 13.178 | 70.013 | 210.473 | 1.000 | -173.297 | 199.653 |
| | 3 | 1 | -.934 | 44.280 | 210.473 | 1.000 | -118.871 | 117.003 |
| | | 2 | 30.700 | 44.280 | 210.473 | 1.000 | -87.237 | 148.637 |
| | | 4 | 43.878 | 70.013 | 210.473 | 1.000 | -142.597 | 230.353 |
| | 4 | 1 | -44.812 | 70.013 | 210.473 | 1.000 | -231.287 | 141.663 |
| | | 2 | -13.178 | 70.013 | 210.473 | 1.000 | -199.653 | 173.297 |
| | | 3 | -43.878 | 70.013 | 210.473 | 1.000 | -230.353 | 142.597 |
| 75 | 1 | 2 | 15.530 | 44.280 | 210.473 | 1.000 | -102.407 | 133.467 |
| | | 3 | 11.506 | 44.280 | 210.473 | 1.000 | -106.431 | 129.443 |
| | | 4 | 17.308 | 70.013 | 210.473 | 1.000 | -169.167 | 203.783 |
| | 2 | 1 | -15.530 | 44.280 | 210.473 | 1.000 | -133.467 | 102.407 |
| | | 3 | -4.024 | 44.280 | 210.473 | 1.000 | -121.961 | 113.913 |
| | | 4 | 1.778 | 70.013 | 210.473 | 1.000 | -184.697 | 188.253 |
| | 3 | 1 | -11.506 | 44.280 | 210.473 | 1.000 | -129.443 | 106.431 |
| | | 2 | 4.024 | 44.280 | 210.473 | 1.000 | -113.913 | 121.961 |
| | | 4 | 5.802 | 70.013 | 210.473 | 1.000 | -180.673 | 192.277 |
| | 4 | 1 | -17.308 | 70.013 | 210.473 | 1.000 | -203.783 | 169.167 |
| | | 2 | -1.778 | 70.013 | 210.473 | 1.000 | -188.253 | 184.697 |
| | | 3 | -5.802 | 70.013 | 210.473 | 1.000 | -192.277 | 180.673 |
| 90 | 1 | 2 | -98.644 | 44.280 | 210.473 | .162 | -216.581 | 19.292 |
| | | 3 | 6.746 | 44.280 | 210.473 | 1.000 | -111.191 | 124.683 |
| | | 4 | -220.116 ^c | 70.013 | 210.473 | .011 | -406.590 | -33.641 |
| | 2 | 1 | 98.644 | 44.280 | 210.473 | .162 | -19.292 | 216.581 |
| | | 3 | 105.391 | 44.280 | 210.473 | .109 | -12.546 | 223.328 |
| | | 4 | -121.471 | 70.013 | 210.473 | .505 | -307.946 | 65.003 |
| | 3 | 1 | -6.746 | 44.280 | 210.473 | 1.000 | -124.683 | 111.191 |
| | | 2 | -105.391 | 44.280 | 210.473 | .109 | -223.328 | 12.546 |
| | | 4 | -226.862 ^c | 70.013 | 210.473 | .008 | -413.337 | -40.388 |
| | 4 | 1 | 220.116 ^c | 70.013 | 210.473 | .011 | 33.641 | 406.590 |
| | | 2 | 121.471 | 70.013 | 210.473 | .505 | -65.003 | 307.946 |
| | | 3 | 226.862 ^c | 70.013 | 210.473 | .008 | 40.388 | 413.337 |

Based on estimated marginal means

^a. The mean difference is significant at the .05 level.

^a. Dependent Variable: TOT_A

^c. Adjustment for multiple comparisons: Bonferroni.

Table A.34. Site * group interactions in cortical thickness (circular ring model). 1 = Control, 2 = LB, 3 = SSMD, 4 = MC3.

| site | (I) 1:CTRL, 2:LB, 3:SSMD, 4:MC3 | (J) 1:CTRL, 2:LB, 3:SSMD, 4:MC3 | Mean Difference (I-J) | Std. Error | df | Sig. ^a | 95% Confidence Interval for Difference ^b | |
|------|---------------------------------|---------------------------------|-----------------------|------------|---------|-------------------|---|-------------|
| | | | | | | | Lower Bound | Upper Bound |
| 10 | 1 | 2 | -.128 | .341 | 234.231 | 1.000 | -1.035 | .778 |
| | | 3 | -.122 | .341 | 234.231 | 1.000 | -1.029 | .785 |
| | | 4 | .535 | .539 | 234.231 | 1.000 | -.898 | 1.969 |
| | 2 | 1 | .128 | .341 | 234.231 | 1.000 | -.778 | 1.035 |
| | | 3 | .006 | .341 | 234.231 | 1.000 | -.900 | .913 |
| | | 4 | .664 | .539 | 234.231 | 1.000 | -.770 | 2.097 |
| | 3 | 1 | .122 | .341 | 234.231 | 1.000 | -.785 | 1.029 |
| | | 2 | -.006 | .341 | 234.231 | 1.000 | -.913 | .900 |
| | | 4 | .657 | .539 | 234.231 | 1.000 | -.776 | 2.091 |
| | 4 | 1 | -.535 | .539 | 234.231 | 1.000 | -1.969 | .898 |
| | | 2 | -.664 | .539 | 234.231 | 1.000 | -2.097 | .770 |
| | | 3 | -.657 | .539 | 234.231 | 1.000 | -2.091 | .776 |
| 25 | 1 | 2 | .123 | .341 | 234.231 | 1.000 | -.784 | 1.029 |
| | | 3 | .046 | .341 | 234.231 | 1.000 | -.861 | .952 |
| | | 4 | .349 | .539 | 234.231 | 1.000 | -1.084 | 1.783 |
| | 2 | 1 | -.123 | .341 | 234.231 | 1.000 | -1.029 | .784 |
| | | 3 | -.077 | .341 | 234.231 | 1.000 | -.984 | .830 |
| | | 4 | .227 | .539 | 234.231 | 1.000 | -1.207 | 1.660 |
| | 3 | 1 | -.046 | .341 | 234.231 | 1.000 | -.952 | .861 |
| | | 2 | .077 | .341 | 234.231 | 1.000 | -.830 | .984 |
| | | 4 | .304 | .539 | 234.231 | 1.000 | -1.130 | 1.737 |
| | 4 | 1 | -.349 | .539 | 234.231 | 1.000 | -1.783 | 1.084 |
| | | 2 | -.227 | .539 | 234.231 | 1.000 | -1.660 | 1.207 |
| | | 3 | -.304 | .539 | 234.231 | 1.000 | -1.737 | 1.130 |
| 50 | 1 | 2 | .278 | .341 | 234.231 | 1.000 | -.629 | 1.185 |
| | | 3 | .065 | .341 | 234.231 | 1.000 | -.841 | .972 |
| | | 4 | .825 | .539 | 234.231 | .761 | -.608 | 2.259 |
| | 2 | 1 | -.278 | .341 | 234.231 | 1.000 | -1.185 | .629 |
| | | 3 | -.213 | .341 | 234.231 | 1.000 | -1.119 | .694 |
| | | 4 | .547 | .539 | 234.231 | 1.000 | -.886 | 1.981 |
| | 3 | 1 | -.065 | .341 | 234.231 | 1.000 | -.972 | .841 |
| | | 2 | .213 | .341 | 234.231 | 1.000 | -.694 | 1.119 |
| | | 4 | .760 | .539 | 234.231 | .958 | -.674 | 2.194 |
| | 4 | 1 | -.825 | .539 | 234.231 | .761 | -2.259 | .608 |
| | | 2 | -.547 | .539 | 234.231 | 1.000 | -1.981 | .886 |
| | | 3 | -.760 | .539 | 234.231 | .958 | -2.194 | .674 |
| 75 | 1 | 2 | .144 | .341 | 234.231 | 1.000 | -.763 | 1.051 |
| | | 3 | .101 | .341 | 234.231 | 1.000 | -.806 | 1.008 |
| | | 4 | .140 | .539 | 234.231 | 1.000 | -1.294 | 1.574 |
| | 2 | 1 | -.144 | .341 | 234.231 | 1.000 | -1.051 | .763 |
| | | 3 | -.043 | .341 | 234.231 | 1.000 | -.950 | .864 |
| | | 4 | -.004 | .539 | 234.231 | 1.000 | -1.438 | 1.430 |
| | 3 | 1 | -.101 | .341 | 234.231 | 1.000 | -1.008 | .806 |
| | | 2 | .043 | .341 | 234.231 | 1.000 | -.864 | .950 |
| | | 4 | .039 | .539 | 234.231 | 1.000 | -1.395 | 1.473 |
| | 4 | 1 | -.140 | .539 | 234.231 | 1.000 | -1.574 | 1.294 |
| | | 2 | .004 | .539 | 234.231 | 1.000 | -1.430 | 1.438 |
| | | 3 | -.039 | .539 | 234.231 | 1.000 | -1.473 | 1.395 |
| 90 | 1 | 2 | -.655 | .341 | 234.231 | .335 | -1.562 | .252 |
| | | 3 | .036 | .341 | 234.231 | 1.000 | -.871 | .942 |
| | | 4 | -1.416 | .539 | 234.231 | .055 | -2.850 | .017 |
| | 2 | 1 | .655 | .341 | 234.231 | .335 | -.252 | 1.562 |
| | | 3 | .691 | .341 | 234.231 | .263 | -.216 | 1.597 |
| | | 4 | -.761 | .539 | 234.231 | .954 | -2.195 | .672 |
| | 3 | 1 | -.036 | .341 | 234.231 | 1.000 | -.942 | .871 |
| | | 2 | -.691 | .341 | 234.231 | .263 | -1.597 | .216 |
| | | 4 | -1.452 [*] | .539 | 234.231 | .045 | -2.886 | -.018 |
| | 4 | 1 | 1.416 | .539 | 234.231 | .055 | -.017 | 2.850 |
| | | 2 | .761 | .539 | 234.231 | .954 | -.672 | 2.195 |
| | | 3 | 1.452 [*] | .539 | 234.231 | .045 | .018 | 2.886 |

Based on estimated marginal means

*. The mean difference is significant at the .05 level.

a. Dependent Variable: CRT_THK_C.

c. Adjustment for multiple comparisons: Bonferroni.

Table A.35. Site * group interactions in periosteal circumference (circular ring model). 1 = Control, 2 = LB, 3 = SSMD, 4 = MC3.

| site | (I) 1:CTRL, 2:LB, 3:SSMD, 4:MC3 | (J) 1:CTRL, 2:LB, 3:SSMD, 4:MC3 | Mean Difference (I-J) | Std. Error | df | Sig. ^c | 95% Confidence Interval for Difference ^a | |
|------|---------------------------------|---------------------------------|-----------------------|------------|---------|-------------------|---|-------------|
| | | | | | | | Lower Bound | Upper Bound |
| 10 | 1 | 2 | -.816 | 2.128 | 187.068 | 1.000 | -6.491 | 4.859 |
| | | 3 | -.773 | 2.128 | 187.068 | 1.000 | -6.448 | 4.903 |
| | | 4 | 3.352 | 3.365 | 187.068 | 1.000 | -5.621 | 12.325 |
| | 2 | 1 | .816 | 2.128 | 187.068 | 1.000 | -4.859 | 6.491 |
| | | 3 | .043 | 2.128 | 187.068 | 1.000 | -5.632 | 5.719 |
| | | 4 | 4.168 | 3.365 | 187.068 | 1.000 | -4.805 | 13.141 |
| | 3 | 1 | .773 | 2.128 | 187.068 | 1.000 | -4.903 | 6.448 |
| | | 2 | -.043 | 2.128 | 187.068 | 1.000 | -5.719 | 5.632 |
| | | 4 | 4.125 | 3.365 | 187.068 | 1.000 | -4.849 | 13.098 |
| | 4 | 1 | -3.352 | 3.365 | 187.068 | 1.000 | -12.325 | 5.621 |
| | | 2 | -4.168 | 3.365 | 187.068 | 1.000 | -13.141 | 4.805 |
| | | 3 | -4.125 | 3.365 | 187.068 | 1.000 | -13.098 | 4.849 |
| 25 | 1 | 2 | .370 | 2.128 | 187.068 | 1.000 | -5.305 | 6.045 |
| | | 3 | -.660 | 2.128 | 187.068 | 1.000 | -6.335 | 5.015 |
| | | 4 | 2.066 | 3.365 | 187.068 | 1.000 | -6.907 | 11.039 |
| | 2 | 1 | -.370 | 2.128 | 187.068 | 1.000 | -6.045 | 5.305 |
| | | 3 | -1.030 | 2.128 | 187.068 | 1.000 | -6.705 | 4.645 |
| | | 4 | 1.696 | 3.365 | 187.068 | 1.000 | -7.277 | 10.669 |
| | 3 | 1 | .660 | 2.128 | 187.068 | 1.000 | -5.015 | 6.335 |
| | | 2 | 1.030 | 2.128 | 187.068 | 1.000 | -4.645 | 6.705 |
| | | 4 | 2.726 | 3.365 | 187.068 | 1.000 | -6.247 | 11.699 |
| | 4 | 1 | -2.066 | 3.365 | 187.068 | 1.000 | -11.039 | 6.907 |
| | | 2 | -1.696 | 3.365 | 187.068 | 1.000 | -10.669 | 7.277 |
| | | 3 | -2.726 | 3.365 | 187.068 | 1.000 | -11.699 | 6.247 |
| 50 | 1 | 2 | 1.825 | 2.128 | 187.068 | 1.000 | -3.851 | 7.500 |
| | | 3 | .065 | 2.128 | 187.068 | 1.000 | -5.611 | 5.740 |
| | | 4 | 2.470 | 3.365 | 187.068 | 1.000 | -6.503 | 11.443 |
| | 2 | 1 | -1.825 | 2.128 | 187.068 | 1.000 | -7.500 | 3.851 |
| | | 3 | -1.760 | 2.128 | 187.068 | 1.000 | -7.435 | 3.915 |
| | | 4 | .646 | 3.365 | 187.068 | 1.000 | -8.327 | 9.619 |
| | 3 | 1 | -.065 | 2.128 | 187.068 | 1.000 | -5.740 | 5.611 |
| | | 2 | 1.760 | 2.128 | 187.068 | 1.000 | -3.915 | 7.435 |
| | | 4 | 2.406 | 3.365 | 187.068 | 1.000 | -6.567 | 11.379 |
| | 4 | 1 | -2.470 | 3.365 | 187.068 | 1.000 | -11.443 | 6.503 |
| | | 2 | -.646 | 3.365 | 187.068 | 1.000 | -9.619 | 8.327 |
| | | 3 | -2.406 | 3.365 | 187.068 | 1.000 | -11.379 | 6.567 |
| 75 | 1 | 2 | .906 | 2.128 | 187.068 | 1.000 | -4.769 | 6.581 |
| | | 3 | .634 | 2.128 | 187.068 | 1.000 | -5.041 | 6.309 |
| | | 4 | .878 | 3.365 | 187.068 | 1.000 | -8.095 | 9.852 |
| | 2 | 1 | -.906 | 2.128 | 187.068 | 1.000 | -6.581 | 4.769 |
| | | 3 | -.272 | 2.128 | 187.068 | 1.000 | -5.947 | 5.403 |
| | | 4 | -.028 | 3.365 | 187.068 | 1.000 | -9.001 | 8.945 |
| | 3 | 1 | -.634 | 2.128 | 187.068 | 1.000 | -6.309 | 5.041 |
| | | 2 | .272 | 2.128 | 187.068 | 1.000 | -5.403 | 5.947 |
| | | 4 | .244 | 3.365 | 187.068 | 1.000 | -8.729 | 9.218 |
| | 4 | 1 | -.878 | 3.365 | 187.068 | 1.000 | -9.852 | 8.095 |
| | | 2 | .028 | 3.365 | 187.068 | 1.000 | -8.945 | 9.001 |
| | | 3 | -.244 | 3.365 | 187.068 | 1.000 | -9.218 | 8.729 |
| 90 | 1 | 2 | -4.336 | 2.128 | 187.068 | .258 | -10.011 | 1.339 |
| | | 3 | .226 | 2.128 | 187.068 | 1.000 | -5.449 | 5.901 |
| | | 4 | -9.590 ^a | 3.365 | 187.068 | .029 | -18.564 | -.617 |
| | 2 | 1 | 4.336 | 2.128 | 187.068 | .258 | -1.339 | 10.011 |
| | | 3 | 4.562 | 2.128 | 187.068 | .200 | -1.113 | 10.237 |
| | | 4 | -5.254 | 3.365 | 187.068 | .721 | -14.228 | 3.719 |
| | 3 | 1 | -.226 | 2.128 | 187.068 | 1.000 | -5.901 | 5.449 |
| | | 2 | -4.562 | 2.128 | 187.068 | .200 | -10.237 | 1.113 |
| | | 4 | -9.816 ^a | 3.365 | 187.068 | .024 | -18.790 | -.843 |
| | 4 | 1 | 9.590 ^a | 3.365 | 187.068 | .029 | .617 | 18.564 |
| | | 2 | 5.254 | 3.365 | 187.068 | .721 | -3.719 | 14.228 |
| | | 3 | 9.816 ^a | 3.365 | 187.068 | .024 | .843 | 18.790 |

Based on estimated marginal means

*. The mean difference is significant at the .05 level.

a. Dependent Variable: PERI_C.

c. Adjustment for multiple comparisons: Bonferroni.

Table A.36. Site * group interactions in cortical and subcortical area. 1 = Control, 2 = LB, 3 = SSMD, 4 = MC3.

| | | | Mean | | | | 95% Confidence Interval for | |
|------|---------------------------------|---------------------------------|-----------------------|------------|---------|-------------------|-----------------------------|-------------|
| | | | Difference (I- | | | | Difference ^a | |
| site | (I) 1:CTRL, 2:LB, 3:SSMD, 4:MC3 | (J) 1:CTRL, 2:LB, 3:SSMD, 4:MC3 | J) | Std. Error | df | Sig. ^b | Lower Bound | Upper Bound |
| 10 | 1 | 2 | 55.004 | 22.106 | 122.743 | .085 | -4.276 | 114.285 |
| | | 3 | 10.191 | 22.106 | 122.743 | 1.000 | -49.090 | 69.472 |
| | | 4 | 85.232 | 34.952 | 122.743 | .097 | -8.499 | 178.963 |
| | 2 | 1 | -55.004 | 22.106 | 122.743 | .085 | -114.285 | 4.276 |
| | | 3 | -44.814 | 22.106 | 122.743 | .269 | -104.095 | 14.467 |
| | | 4 | 30.227 | 34.952 | 122.743 | 1.000 | -63.504 | 123.959 |
| | 3 | 1 | -10.191 | 22.106 | 122.743 | 1.000 | -69.472 | 49.090 |
| | | 2 | 44.814 | 22.106 | 122.743 | .269 | -14.467 | 104.095 |
| | | 4 | 75.041 | 34.952 | 122.743 | .203 | -18.690 | 168.772 |
| | 4 | 1 | -85.232 | 34.952 | 122.743 | .097 | -178.963 | 8.499 |
| | | 2 | -30.227 | 34.952 | 122.743 | 1.000 | -123.959 | 63.504 |
| | | 3 | -75.041 | 34.952 | 122.743 | .203 | -168.772 | 18.690 |
| 25 | 1 | 2 | 7.840 | 22.106 | 122.743 | 1.000 | -51.441 | 67.121 |
| | | 3 | -5.800 | 22.106 | 122.743 | 1.000 | -65.081 | 53.481 |
| | | 4 | 36.224 | 34.952 | 122.743 | 1.000 | -57.507 | 129.955 |
| | 2 | 1 | -7.840 | 22.106 | 122.743 | 1.000 | -67.121 | 51.441 |
| | | 3 | -13.640 | 22.106 | 122.743 | 1.000 | -72.921 | 45.641 |
| | | 4 | 28.384 | 34.952 | 122.743 | 1.000 | -65.347 | 122.115 |
| | 3 | 1 | 5.800 | 22.106 | 122.743 | 1.000 | -53.481 | 65.081 |
| | | 2 | 13.640 | 22.106 | 122.743 | 1.000 | -45.641 | 72.921 |
| | | 4 | 42.024 | 34.952 | 122.743 | 1.000 | -51.707 | 135.755 |
| | 4 | 1 | -36.224 | 34.952 | 122.743 | 1.000 | -129.955 | 57.507 |
| | | 2 | -28.384 | 34.952 | 122.743 | 1.000 | -122.115 | 65.347 |
| | | 3 | -42.024 | 34.952 | 122.743 | 1.000 | -135.755 | 51.707 |
| 50 | 1 | 2 | 31.158 | 22.106 | 122.743 | .967 | -28.123 | 90.439 |
| | | 3 | 2.776 | 22.106 | 122.743 | 1.000 | -56.505 | 62.057 |
| | | 4 | 64.428 | 34.952 | 122.743 | .406 | -29.303 | 158.159 |
| | 2 | 1 | -31.158 | 22.106 | 122.743 | .967 | -90.439 | 28.123 |
| | | 3 | -28.382 | 22.106 | 122.743 | 1.000 | -87.663 | 30.899 |
| | | 4 | 33.270 | 34.952 | 122.743 | 1.000 | -60.461 | 127.001 |
| | 3 | 1 | -2.776 | 22.106 | 122.743 | 1.000 | -62.057 | 56.505 |
| | | 2 | 28.382 | 22.106 | 122.743 | 1.000 | -30.899 | 87.663 |
| | | 4 | 61.652 | 34.952 | 122.743 | .481 | -32.079 | 155.383 |
| | 4 | 1 | -64.428 | 34.952 | 122.743 | .406 | -158.159 | 29.303 |
| | | 2 | -33.270 | 34.952 | 122.743 | 1.000 | -127.001 | 60.461 |
| | | 3 | -61.652 | 34.952 | 122.743 | .481 | -155.383 | 32.079 |
| 75 | 1 | 2 | 46.460 | 22.106 | 122.743 | .226 | -12.821 | 105.741 |
| | | 3 | -30.580 | 22.106 | 122.743 | 1.000 | -89.861 | 28.701 |
| | | 4 | 103.870 ^c | 34.952 | 122.743 | .021 | 10.139 | 197.601 |
| | 2 | 1 | -46.460 | 22.106 | 122.743 | .226 | -105.741 | 12.821 |
| | | 3 | -77.040 ^c | 22.106 | 122.743 | .004 | -136.321 | -17.759 |
| | | 4 | 57.410 | 34.952 | 122.743 | .618 | -36.321 | 151.141 |
| | 3 | 1 | 30.580 | 22.106 | 122.743 | 1.000 | -28.701 | 89.861 |
| | | 2 | 77.040 ^c | 22.106 | 122.743 | .004 | 17.759 | 136.321 |
| | | 4 | 134.450 ^c | 34.952 | 122.743 | .001 | 40.719 | 228.181 |
| | 4 | 1 | -103.870 ^c | 34.952 | 122.743 | .021 | -197.601 | -10.139 |
| | | 2 | -57.410 | 34.952 | 122.743 | .618 | -151.141 | 36.321 |
| | | 3 | -134.450 ^c | 34.952 | 122.743 | .001 | -228.181 | -40.719 |
| 90 | 1 | 2 | -15.955 | 22.106 | 122.743 | 1.000 | -75.236 | 43.326 |
| | | 3 | -12.528 | 22.106 | 122.743 | 1.000 | -71.809 | 46.753 |
| | | 4 | -21.244 | 34.952 | 122.743 | 1.000 | -114.976 | 72.487 |
| | 2 | 1 | 15.955 | 22.106 | 122.743 | 1.000 | -43.326 | 75.236 |
| | | 3 | 3.427 | 22.106 | 122.743 | 1.000 | -55.854 | 62.708 |
| | | 4 | -5.289 | 34.952 | 122.743 | 1.000 | -99.021 | 88.442 |
| | 3 | 1 | 12.528 | 22.106 | 122.743 | 1.000 | -46.753 | 71.809 |
| | | 2 | -3.427 | 22.106 | 122.743 | 1.000 | -62.708 | 55.854 |
| | | 4 | -8.716 | 34.952 | 122.743 | 1.000 | -102.447 | 85.015 |
| | 4 | 1 | 21.244 | 34.952 | 122.743 | 1.000 | -72.487 | 114.976 |
| | | 2 | 5.289 | 34.952 | 122.743 | 1.000 | -88.442 | 99.021 |
| | | 3 | 8.716 | 34.952 | 122.743 | 1.000 | -85.015 | 102.447 |

Based on estimated marginal means

*. The mean difference is significant at the .05 level.

a. Dependent Variable: CRTSUB_A.

c. Adjustment for multiple comparisons: Bonferroni.

Table A.37. Site * group interactions in axial moment of inertia of cortical area (x-axis). 1 = Control, 2 = LB, 3 = SSMD, 4 = MC3.

| site | (I) 1:CTRL, 2:LB, 3:SSMD, 4:MC3 | (J) 1:CTRL, 2:LB, 3:SSMD, 4:MC3 | Mean Difference (I-J) | Std. Error | df | Sig. ^a | 95% Confidence Interval for Difference ^b | |
|------|---------------------------------|---------------------------------|-----------------------|------------|---------|-------------------|---|-------------|
| | | | | | | | Lower Bound | Upper Bound |
| 10 | 1 | 2 | -3922.168 | 8690.880 | 269.274 | 1.000 | -27021.585 | 19177.249 |
| | | 3 | -4423.209 | 8690.880 | 269.274 | 1.000 | -27522.625 | 18676.208 |
| | | 4 | 7199.543 | 13741.488 | 269.274 | 1.000 | -29323.841 | 43722.928 |
| | 2 | 1 | 3922.168 | 8690.880 | 269.274 | 1.000 | -19177.249 | 27021.585 |
| | | 3 | -501.041 | 8690.880 | 269.274 | 1.000 | -23600.457 | 22598.376 |
| | | 4 | 11121.711 | 13741.488 | 269.274 | 1.000 | -25401.673 | 47645.096 |
| | 3 | 1 | 4423.209 | 8690.880 | 269.274 | 1.000 | -18676.208 | 27522.625 |
| | | 2 | 501.041 | 8690.880 | 269.274 | 1.000 | -22598.376 | 23600.457 |
| | | 4 | 11622.752 | 13741.488 | 269.274 | 1.000 | -24900.633 | 48146.137 |
| | 4 | 1 | -7199.543 | 13741.488 | 269.274 | 1.000 | -43722.928 | 29323.841 |
| | | 2 | -11121.711 | 13741.488 | 269.274 | 1.000 | -47645.096 | 25401.673 |
| | | 3 | -11622.752 | 13741.488 | 269.274 | 1.000 | -48146.137 | 24900.633 |
| 25 | 1 | 2 | 628.475 | 8690.880 | 269.274 | 1.000 | -22470.942 | 23727.891 |
| | | 3 | -1290.308 | 8690.880 | 269.274 | 1.000 | -24389.724 | 21809.109 |
| | | 4 | 1553.512 | 13741.488 | 269.274 | 1.000 | -34969.872 | 38076.897 |
| | 2 | 1 | -628.475 | 8690.880 | 269.274 | 1.000 | -23727.891 | 22470.942 |
| | | 3 | -1918.782 | 8690.880 | 269.274 | 1.000 | -25018.199 | 21180.634 |
| | | 4 | 925.038 | 13741.488 | 269.274 | 1.000 | -35598.347 | 37448.422 |
| | 3 | 1 | 1290.308 | 8690.880 | 269.274 | 1.000 | -21809.109 | 24389.724 |
| | | 2 | 1918.782 | 8690.880 | 269.274 | 1.000 | -21180.634 | 25018.199 |
| | | 4 | 2843.820 | 13741.488 | 269.274 | 1.000 | -33679.565 | 39367.205 |
| | 4 | 1 | -1553.512 | 13741.488 | 269.274 | 1.000 | -38076.897 | 34969.872 |
| | | 2 | -925.038 | 13741.488 | 269.274 | 1.000 | -37448.422 | 35598.347 |
| | | 3 | -2843.820 | 13741.488 | 269.274 | 1.000 | -39367.205 | 33679.565 |
| 50 | 1 | 2 | 6280.249 | 8690.880 | 269.274 | 1.000 | -16819.168 | 29379.665 |
| | | 3 | 2334.076 | 8690.880 | 269.274 | 1.000 | -20765.341 | 25433.493 |
| | | 4 | 6490.273 | 13741.488 | 269.274 | 1.000 | -30033.112 | 43013.657 |
| | 2 | 1 | -6280.249 | 8690.880 | 269.274 | 1.000 | -29379.665 | 16819.168 |
| | | 3 | -3946.173 | 8690.880 | 269.274 | 1.000 | -27045.590 | 19153.244 |
| | | 4 | 210.024 | 13741.488 | 269.274 | 1.000 | -36313.361 | 36733.409 |
| | 3 | 1 | -2334.076 | 8690.880 | 269.274 | 1.000 | -25433.493 | 20765.341 |
| | | 2 | 3946.173 | 8690.880 | 269.274 | 1.000 | -19153.244 | 27045.590 |
| | | 4 | 4156.197 | 13741.488 | 269.274 | 1.000 | -32367.188 | 40679.581 |
| | 4 | 1 | -6490.273 | 13741.488 | 269.274 | 1.000 | -43013.657 | 30033.112 |
| | | 2 | -210.024 | 13741.488 | 269.274 | 1.000 | -36733.409 | 36313.361 |
| | | 3 | -4156.197 | 13741.488 | 269.274 | 1.000 | -40679.581 | 32367.188 |
| 75 | 1 | 2 | 2526.918 | 8690.880 | 269.274 | 1.000 | -20572.499 | 25626.335 |
| | | 3 | 1141.243 | 8690.880 | 269.274 | 1.000 | -21958.173 | 24240.660 |
| | | 4 | -466.178 | 13741.488 | 269.274 | 1.000 | -36989.562 | 36057.207 |
| | 2 | 1 | -2526.918 | 8690.880 | 269.274 | 1.000 | -25626.335 | 20572.499 |
| | | 3 | -1385.675 | 8690.880 | 269.274 | 1.000 | -24485.091 | 21713.742 |
| | | 4 | -2993.096 | 13741.488 | 269.274 | 1.000 | -39516.481 | 33530.289 |
| | 3 | 1 | -1141.243 | 8690.880 | 269.274 | 1.000 | -24240.660 | 21958.173 |
| | | 2 | 1385.675 | 8690.880 | 269.274 | 1.000 | -21713.742 | 24485.091 |
| | | 4 | -1607.421 | 13741.488 | 269.274 | 1.000 | -38130.806 | 34915.964 |
| | 4 | 1 | 466.178 | 13741.488 | 269.274 | 1.000 | -36057.207 | 36989.562 |
| | | 2 | 2993.096 | 13741.488 | 269.274 | 1.000 | -33530.289 | 39516.481 |
| | | 3 | 1607.421 | 13741.488 | 269.274 | 1.000 | -34915.964 | 38130.806 |
| 90 | 1 | 2 | -24451.318* | 8690.880 | 269.274 | .032 | -47550.734 | -1351.901 |
| | | 3 | 1033.370 | 8690.880 | 269.274 | 1.000 | -22066.047 | 24132.787 |
| | | 4 | -75508.534* | 13741.488 | 269.274 | .000 | -112031.919 | -38985.150 |
| | 2 | 1 | 24451.318* | 8690.880 | 269.274 | .032 | 1351.901 | 47550.734 |
| | | 3 | 25484.688* | 8690.880 | 269.274 | .022 | 2385.271 | 48584.105 |
| | | 4 | -51057.217* | 13741.488 | 269.274 | .001 | -87580.601 | -14533.832 |
| | 3 | 1 | -1033.370 | 8690.880 | 269.274 | 1.000 | -24132.787 | 22066.047 |
| | | 2 | -25484.688* | 8690.880 | 269.274 | .022 | -48584.105 | -2385.271 |
| | | 4 | -76541.904* | 13741.488 | 269.274 | .000 | -113065.289 | -40018.520 |
| | 4 | 1 | 75508.534* | 13741.488 | 269.274 | .000 | 38985.150 | 112031.919 |
| | | 2 | 51057.217* | 13741.488 | 269.274 | .001 | 14533.832 | 87580.601 |
| | | 3 | 76541.904* | 13741.488 | 269.274 | .000 | 40018.520 | 113065.289 |

Based on estimated marginal means

*. The mean difference is significant at the .05 level.

a. Dependent Variable: IX_CRT_A

c. Adjustment for multiple comparisons: Bonferroni.

Table A.38. Site * group interactions in axial moment of inertia of cortical area (y-axis). 1 = Control, 2 = LB, 3 = SSMD, 4 = MC3.

| site | (I) 1:CTRL, 2:LB, 3:SSMD, 4:MC3 | (J) 1:CTRL, 2:LB, 3:SSMD, 4:MC3 | Mean Difference (I-J) | Std. Error | df | Sig. ^c | 95% Confidence Interval for Difference ^a | |
|------|---------------------------------|---------------------------------|--------------------------|------------|---------|-------------------|---|-------------|
| | | | | | | | Lower Bound | Upper Bound |
| 10 | 1 | 2 | -6696.827 | 15143.183 | 241.924 | 1.000 | -46979.638 | 33585.984 |
| | | 3 | -4217.472 | 15143.183 | 241.924 | 1.000 | -44500.283 | 36065.339 |
| | | 4 | 19324.853 | 23943.475 | 241.924 | 1.000 | -44367.863 | 83017.570 |
| | 2 | 1 | 6696.827 | 15143.183 | 241.924 | 1.000 | -33585.984 | 46979.638 |
| | | 3 | 2479.355 | 15143.183 | 241.924 | 1.000 | -37803.456 | 42762.166 |
| | | 4 | 26021.681 | 23943.475 | 241.924 | 1.000 | -37671.036 | 89714.397 |
| | 3 | 1 | 4217.472 | 15143.183 | 241.924 | 1.000 | -36065.339 | 44500.283 |
| | | 2 | -2479.355 | 15143.183 | 241.924 | 1.000 | -42762.166 | 37803.456 |
| | | 4 | 23542.325 | 23943.475 | 241.924 | 1.000 | -40150.391 | 87235.042 |
| | 4 | 1 | -19324.853 | 23943.475 | 241.924 | 1.000 | -83017.570 | 44367.863 |
| | | 2 | -26021.681 | 23943.475 | 241.924 | 1.000 | -89714.397 | 37671.036 |
| | | 3 | -23542.325 | 23943.475 | 241.924 | 1.000 | -87235.042 | 40150.391 |
| 25 | 1 | 2 | -1158.954 | 15143.183 | 241.924 | 1.000 | -41441.765 | 39123.857 |
| | | 3 | -3013.939 | 15143.183 | 241.924 | 1.000 | -43296.750 | 37268.872 |
| | | 4 | 9723.042 | 23943.475 | 241.924 | 1.000 | -53969.675 | 73415.758 |
| | 2 | 1 | 1158.954 | 15143.183 | 241.924 | 1.000 | -39123.857 | 41441.765 |
| | | 3 | -1854.985 | 15143.183 | 241.924 | 1.000 | -42137.796 | 38427.826 |
| | | 4 | 10881.996 | 23943.475 | 241.924 | 1.000 | -52810.721 | 74574.712 |
| | 3 | 1 | 3013.939 | 15143.183 | 241.924 | 1.000 | -37268.872 | 43296.750 |
| | | 2 | 1854.985 | 15143.183 | 241.924 | 1.000 | -38427.826 | 42137.796 |
| | | 4 | 12736.981 | 23943.475 | 241.924 | 1.000 | -50955.736 | 76429.698 |
| | 4 | 1 | -9723.042 | 23943.475 | 241.924 | 1.000 | -73415.758 | 53969.675 |
| | | 2 | -10881.996 | 23943.475 | 241.924 | 1.000 | -74574.712 | 52810.721 |
| | | 3 | -12736.981 | 23943.475 | 241.924 | 1.000 | -76429.698 | 50955.736 |
| 50 | 1 | 2 | 1523.005 | 15143.183 | 241.924 | 1.000 | -38759.806 | 41805.816 |
| | | 3 | -3191.130 | 15143.183 | 241.924 | 1.000 | -43473.941 | 37091.681 |
| | | 4 | 8409.491 | 23943.475 | 241.924 | 1.000 | -55283.226 | 72102.207 |
| | 2 | 1 | -1523.005 | 15143.183 | 241.924 | 1.000 | -41805.816 | 38759.806 |
| | | 3 | -4714.135 | 15143.183 | 241.924 | 1.000 | -44996.946 | 35568.676 |
| | | 4 | 6886.486 | 23943.475 | 241.924 | 1.000 | -56806.231 | 70579.202 |
| | 3 | 1 | 3191.130 | 15143.183 | 241.924 | 1.000 | -37091.681 | 43473.941 |
| | | 2 | 4714.135 | 15143.183 | 241.924 | 1.000 | -35568.676 | 44996.946 |
| | | 4 | 11600.620 | 23943.475 | 241.924 | 1.000 | -52092.096 | 75293.337 |
| | 4 | 1 | -8409.491 | 23943.475 | 241.924 | 1.000 | -72102.207 | 55283.226 |
| | | 2 | -6886.486 | 23943.475 | 241.924 | 1.000 | -70579.202 | 56806.231 |
| | | 3 | -11600.620 | 23943.475 | 241.924 | 1.000 | -75293.337 | 52092.096 |
| 75 | 1 | 2 | 3302.833 | 15143.183 | 241.924 | 1.000 | -36979.978 | 43585.644 |
| | | 3 | 3942.759 | 15143.183 | 241.924 | 1.000 | -36340.053 | 44225.570 |
| | | 4 | -1650.578 | 23943.475 | 241.924 | 1.000 | -65343.295 | 62042.139 |
| | 2 | 1 | -3302.833 | 15143.183 | 241.924 | 1.000 | -43585.644 | 36979.978 |
| | | 3 | 639.925 | 15143.183 | 241.924 | 1.000 | -39642.886 | 40922.736 |
| | | 4 | -4953.411 | 23943.475 | 241.924 | 1.000 | -68646.128 | 58739.306 |
| | 3 | 1 | -3942.759 | 15143.183 | 241.924 | 1.000 | -44225.570 | 36340.053 |
| | | 2 | -639.925 | 15143.183 | 241.924 | 1.000 | -40922.736 | 39642.886 |
| | | 4 | -5593.336 | 23943.475 | 241.924 | 1.000 | -69286.053 | 58099.380 |
| | 4 | 1 | 1650.578 | 23943.475 | 241.924 | 1.000 | -62042.139 | 65343.295 |
| | | 2 | 4953.411 | 23943.475 | 241.924 | 1.000 | -58739.306 | 68646.128 |
| | | 3 | 5593.336 | 23943.475 | 241.924 | 1.000 | -58099.380 | 69286.053 |
| 90 | 1 | 2 | -35967.680 | 15143.183 | 241.924 | .110 | -76250.491 | 4315.131 |
| | | 3 | 8675.536 | 15143.183 | 241.924 | 1.000 | -31607.275 | 48958.347 |
| | | 4 | -109524.166 ^a | 23943.475 | 241.924 | .000 | -173216.882 | -45831.449 |
| | 2 | 1 | 35967.680 | 15143.183 | 241.924 | .110 | -4315.131 | 76250.491 |
| | | 3 | 44643.216 ^a | 15143.183 | 241.924 | .021 | 4360.405 | 84926.027 |
| | | 4 | -73556.486 ^a | 23943.475 | 241.924 | .014 | -137249.202 | -9863.769 |
| | 3 | 1 | -8675.536 | 15143.183 | 241.924 | 1.000 | -48958.347 | 31607.275 |
| | | 2 | -44643.216 ^a | 15143.183 | 241.924 | .021 | -84926.027 | -4360.405 |
| | | 4 | -118199.702 ^a | 23943.475 | 241.924 | .000 | -181892.418 | -54506.985 |
| | 4 | 1 | 109524.166 ^a | 23943.475 | 241.924 | .000 | 45831.449 | 173216.882 |
| | | 2 | 73556.486 ^a | 23943.475 | 241.924 | .014 | 9863.769 | 137249.202 |
| | | 3 | 118199.702 ^a | 23943.475 | 241.924 | .000 | 54506.985 | 181892.418 |

Based on estimated marginal means

*. The mean difference is significant at the .05 level.

a. Dependent Variable: IV_CRT_A.

c. Adjustment for multiple comparisons: Bonferroni.

Table A.39. Site * group interactions in polar moment of inertia of cortical area. 1 = Control, 2 = LB, 3 = SSMD, 4 = MC3.

| site | (I) 1:CTRL, 2:LB, 3:SSMD, 4:MC3 | (J) 1:CTRL, 2:LB, 3:SSMD, 4:MC3 | Mean Difference (I-J) | Std. Error | df | Sig. ^a | 95% Confidence Interval for Difference ^b | |
|------|---------------------------------|---------------------------------|--------------------------|------------|---------|-------------------|---|-------------|
| | | | | | | | Lower Bound | Upper Bound |
| 10 | 1 | 2 | -10618.995 | 22985.249 | 253.415 | 1.000 | -71739.744 | 50501.755 |
| | | 3 | -8640.680 | 22985.249 | 253.415 | 1.000 | -69761.430 | 52480.069 |
| | | 4 | 26524.395 | 36342.870 | 253.415 | 1.000 | -70115.995 | 123164.785 |
| | 2 | 1 | 10618.995 | 22985.249 | 253.415 | 1.000 | -50501.755 | 71739.744 |
| | | 3 | 1978.314 | 22985.249 | 253.415 | 1.000 | -59142.435 | 63099.064 |
| | | 4 | 37143.390 | 36342.870 | 253.415 | 1.000 | -59497.000 | 133783.780 |
| | 3 | 1 | 8640.680 | 22985.249 | 253.415 | 1.000 | -52480.069 | 69761.430 |
| | | 2 | -1978.314 | 22985.249 | 253.415 | 1.000 | -63099.064 | 59142.435 |
| | | 4 | 35165.075 | 36342.870 | 253.415 | 1.000 | -61475.315 | 131805.465 |
| | 4 | 1 | -26524.395 | 36342.870 | 253.415 | 1.000 | -123164.785 | 70115.995 |
| | | 2 | -37143.390 | 36342.870 | 253.415 | 1.000 | -133783.780 | 59497.000 |
| | | 3 | -35165.075 | 36342.870 | 253.415 | 1.000 | -131805.465 | 61475.315 |
| 25 | 1 | 2 | -530.479 | 22985.249 | 253.415 | 1.000 | -61651.228 | 60590.270 |
| | | 3 | -4304.247 | 22985.249 | 253.415 | 1.000 | -65424.996 | 56816.502 |
| | | 4 | 11276.554 | 36342.870 | 253.415 | 1.000 | -85363.836 | 107916.944 |
| | 2 | 1 | 530.479 | 22985.249 | 253.415 | 1.000 | -60590.270 | 61651.228 |
| | | 3 | -3773.768 | 22985.249 | 253.415 | 1.000 | -64894.517 | 57346.982 |
| | | 4 | 11807.033 | 36342.870 | 253.415 | 1.000 | -84833.357 | 108447.423 |
| | 3 | 1 | 4304.247 | 22985.249 | 253.415 | 1.000 | -56816.502 | 65424.996 |
| | | 2 | 3773.768 | 22985.249 | 253.415 | 1.000 | -57346.982 | 64894.517 |
| | | 4 | 15580.801 | 36342.870 | 253.415 | 1.000 | -81059.589 | 112221.191 |
| | 4 | 1 | -11276.554 | 36342.870 | 253.415 | 1.000 | -107916.944 | 85363.836 |
| | | 2 | -11807.033 | 36342.870 | 253.415 | 1.000 | -108447.423 | 84833.357 |
| | | 3 | -15580.801 | 36342.870 | 253.415 | 1.000 | -112221.191 | 81059.589 |
| 50 | 1 | 2 | 7803.254 | 22985.249 | 253.415 | 1.000 | -53317.495 | 68924.003 |
| | | 3 | -857.053 | 22985.249 | 253.415 | 1.000 | -61977.803 | 60263.696 |
| | | 4 | 14899.764 | 36342.870 | 253.415 | 1.000 | -81740.626 | 111540.154 |
| | 2 | 1 | -7803.254 | 22985.249 | 253.415 | 1.000 | -68924.003 | 53317.495 |
| | | 3 | -8660.307 | 22985.249 | 253.415 | 1.000 | -69781.057 | 52460.442 |
| | | 4 | 7096.509 | 36342.870 | 253.415 | 1.000 | -89543.881 | 103736.899 |
| | 3 | 1 | 857.053 | 22985.249 | 253.415 | 1.000 | -60263.696 | 61977.803 |
| | | 2 | 8660.307 | 22985.249 | 253.415 | 1.000 | -52460.442 | 69781.057 |
| | | 4 | 15756.817 | 36342.870 | 253.415 | 1.000 | -80883.573 | 112397.207 |
| | 4 | 1 | -14899.764 | 36342.870 | 253.415 | 1.000 | -111540.154 | 81740.626 |
| | | 2 | -7096.509 | 36342.870 | 253.415 | 1.000 | -103736.899 | 89543.881 |
| | | 3 | -15756.817 | 36342.870 | 253.415 | 1.000 | -112397.207 | 80883.573 |
| 75 | 1 | 2 | 5829.751 | 22985.249 | 253.415 | 1.000 | -55290.998 | 66950.500 |
| | | 3 | 5084.002 | 22985.249 | 253.415 | 1.000 | -56036.748 | 66204.751 |
| | | 4 | -2116.755 | 36342.870 | 253.415 | 1.000 | -98757.145 | 94523.635 |
| | 2 | 1 | -5829.751 | 22985.249 | 253.415 | 1.000 | -66950.500 | 55290.998 |
| | | 3 | -745.749 | 22985.249 | 253.415 | 1.000 | -61866.499 | 60375.000 |
| | | 4 | -7946.506 | 36342.870 | 253.415 | 1.000 | -104586.896 | 88693.884 |
| | 3 | 1 | -5084.002 | 22985.249 | 253.415 | 1.000 | -66204.751 | 56036.748 |
| | | 2 | 745.749 | 22985.249 | 253.415 | 1.000 | -60375.000 | 61866.499 |
| | | 4 | -7200.757 | 36342.870 | 253.415 | 1.000 | -103841.147 | 89439.633 |
| | 4 | 1 | 2116.755 | 36342.870 | 253.415 | 1.000 | -94523.635 | 98757.145 |
| | | 2 | 7946.506 | 36342.870 | 253.415 | 1.000 | -88693.884 | 104586.896 |
| | | 3 | 7200.757 | 36342.870 | 253.415 | 1.000 | -89439.633 | 103841.147 |
| 90 | 1 | 2 | -60418.997 | 22985.249 | 253.415 | .055 | -121539.747 | 701.752 |
| | | 3 | 9708.908 | 22985.249 | 253.415 | 1.000 | -51411.841 | 70829.657 |
| | | 4 | -185032.699 [*] | 36342.870 | 253.415 | .000 | -281673.089 | -88392.309 |
| | 2 | 1 | 60418.997 | 22985.249 | 253.415 | .055 | -701.752 | 121539.747 |
| | | 3 | 70127.905 [*] | 22985.249 | 253.415 | .015 | 9007.156 | 131248.655 |
| | | 4 | -124613.702 [*] | 36342.870 | 253.415 | .004 | -221254.092 | -27973.312 |
| | 3 | 1 | -9708.908 | 22985.249 | 253.415 | 1.000 | -70829.657 | 51411.841 |
| | | 2 | -70127.905 [*] | 22985.249 | 253.415 | .015 | -131248.655 | -9007.156 |
| | | 4 | -194741.607 [*] | 36342.870 | 253.415 | .000 | -291381.997 | -98101.217 |
| | 4 | 1 | 185032.699 [*] | 36342.870 | 253.415 | .000 | 88392.309 | 281673.089 |
| | | 2 | 124613.702 [*] | 36342.870 | 253.415 | .004 | 27973.312 | 221254.092 |
| | | 3 | 194741.607 [*] | 36342.870 | 253.415 | .000 | 98101.217 | 291381.997 |

Based on estimated marginal means

*. The mean difference is significant at the .05 level.

a. Dependent Variable: IP_CRT_A.

c. Adjustment for multiple comparisons: Bonferroni.

Table A.40. Site * group interactions in cortical moment of resistance (x-axis). 1 = Control, 2 = LB, 3 = SSMD, 4 = MC3.

| site | (I) 1:CTRL, 2:LB, 3:SSMD, 4:MC3 | (J) 1:CTRL, 2:LB, 3:SSMD, 4:MC3 | Mean Difference (I-J) | Std. Error | df | Sig. ^c | 95% Confidence Interval for Difference ^c | |
|------|---------------------------------|---------------------------------|------------------------|------------|---------|-------------------|---|-------------|
| | | | | | | | Lower Bound | Upper Bound |
| 10 | 1 | 2 | -148.681 | 317.560 | 264.808 | 1.000 | -992.828 | 695.465 |
| | | 3 | -174.594 | 317.560 | 264.808 | 1.000 | -1018.740 | 669.552 |
| | | 4 | 421.302 | 502.107 | 264.808 | 1.000 | -913.411 | 1756.014 |
| | 2 | 1 | 148.681 | 317.560 | 264.808 | 1.000 | -695.465 | 992.828 |
| | | 3 | -25.913 | 317.560 | 264.808 | 1.000 | -870.059 | 818.234 |
| | | 4 | 569.983 | 502.107 | 264.808 | 1.000 | -764.729 | 1904.695 |
| | 3 | 1 | 174.594 | 317.560 | 264.808 | 1.000 | -669.552 | 1018.740 |
| | | 2 | 25.913 | 317.560 | 264.808 | 1.000 | -818.234 | 870.059 |
| | | 4 | 595.896 | 502.107 | 264.808 | 1.000 | -738.817 | 1930.608 |
| | 4 | 1 | -421.302 | 502.107 | 264.808 | 1.000 | -1756.014 | 913.411 |
| | | 2 | -569.983 | 502.107 | 264.808 | 1.000 | -1904.695 | 764.729 |
| | | 3 | -595.896 | 502.107 | 264.808 | 1.000 | -1930.608 | 738.817 |
| 25 | 1 | 2 | 43.698 | 317.560 | 264.808 | 1.000 | -800.449 | 887.844 |
| | | 3 | -35.820 | 317.560 | 264.808 | 1.000 | -879.966 | 808.327 |
| | | 4 | 81.091 | 502.107 | 264.808 | 1.000 | -1253.622 | 1415.803 |
| | 2 | 1 | -43.698 | 317.560 | 264.808 | 1.000 | -887.844 | 800.449 |
| | | 3 | -79.517 | 317.560 | 264.808 | 1.000 | -923.664 | 764.629 |
| | | 4 | 37.393 | 502.107 | 264.808 | 1.000 | -1297.319 | 1372.106 |
| | 3 | 1 | 35.820 | 317.560 | 264.808 | 1.000 | -808.327 | 879.966 |
| | | 2 | 79.517 | 317.560 | 264.808 | 1.000 | -764.629 | 923.664 |
| | | 4 | 116.910 | 502.107 | 264.808 | 1.000 | -1217.802 | 1451.623 |
| | 4 | 1 | -81.091 | 502.107 | 264.808 | 1.000 | -1415.803 | 1253.622 |
| | | 2 | -37.393 | 502.107 | 264.808 | 1.000 | -1372.106 | 1297.319 |
| | | 3 | -116.910 | 502.107 | 264.808 | 1.000 | -1451.623 | 1217.802 |
| 50 | 1 | 2 | 233.181 | 317.560 | 264.808 | 1.000 | -610.965 | 1077.327 |
| | | 3 | 70.812 | 317.560 | 264.808 | 1.000 | -773.334 | 914.959 |
| | | 4 | 218.238 | 502.107 | 264.808 | 1.000 | -1116.474 | 1552.951 |
| | 2 | 1 | -233.181 | 317.560 | 264.808 | 1.000 | -1077.327 | 610.965 |
| | | 3 | -162.368 | 317.560 | 264.808 | 1.000 | -1006.515 | 681.778 |
| | | 4 | -14.943 | 502.107 | 264.808 | 1.000 | -1349.655 | 1319.770 |
| | 3 | 1 | -70.812 | 317.560 | 264.808 | 1.000 | -914.959 | 773.334 |
| | | 2 | 162.368 | 317.560 | 264.808 | 1.000 | -681.778 | 1006.515 |
| | | 4 | 147.426 | 502.107 | 264.808 | 1.000 | -1187.287 | 1482.138 |
| | 4 | 1 | -218.238 | 502.107 | 264.808 | 1.000 | -1552.951 | 1116.474 |
| | | 2 | 14.943 | 502.107 | 264.808 | 1.000 | -1319.770 | 1349.655 |
| | | 3 | -147.426 | 502.107 | 264.808 | 1.000 | -1482.138 | 1187.287 |
| 75 | 1 | 2 | 139.938 | 317.560 | 264.808 | 1.000 | -704.209 | 984.084 |
| | | 3 | 88.413 | 317.560 | 264.808 | 1.000 | -755.733 | 932.559 |
| | | 4 | 87.786 | 502.107 | 264.808 | 1.000 | -1246.926 | 1422.499 |
| | 2 | 1 | -139.938 | 317.560 | 264.808 | 1.000 | -984.084 | 704.209 |
| | | 3 | -51.525 | 317.560 | 264.808 | 1.000 | -895.671 | 792.621 |
| | | 4 | -52.151 | 502.107 | 264.808 | 1.000 | -1386.864 | 1282.561 |
| | 3 | 1 | -88.413 | 317.560 | 264.808 | 1.000 | -932.559 | 755.733 |
| | | 2 | 51.525 | 317.560 | 264.808 | 1.000 | -792.621 | 895.671 |
| | | 4 | -626 | 502.107 | 264.808 | 1.000 | -1335.339 | 1334.086 |
| | 4 | 1 | -87.786 | 502.107 | 264.808 | 1.000 | -1422.499 | 1246.926 |
| | | 2 | 52.151 | 502.107 | 264.808 | 1.000 | -1282.561 | 1386.864 |
| | | 3 | 626 | 502.107 | 264.808 | 1.000 | -1334.086 | 1335.339 |
| 90 | 1 | 2 | -899.889 ^a | 317.560 | 264.808 | .030 | -1744.035 | -55.742 |
| | | 3 | -79.859 | 317.560 | 264.808 | 1.000 | -924.005 | 764.287 |
| | | 4 | -2153.299 ^a | 502.107 | 264.808 | .000 | -3488.012 | -818.587 |
| | 2 | 1 | 899.889 ^a | 317.560 | 264.808 | .030 | 55.742 | 1744.035 |
| | | 3 | 820.030 | 317.560 | 264.808 | .062 | -24.117 | 1664.176 |
| | | 4 | -1253.411 | 502.107 | 264.808 | .079 | -2588.123 | 81.302 |
| | 3 | 1 | 79.859 | 317.560 | 264.808 | 1.000 | -764.287 | 924.005 |
| | | 2 | -820.030 | 317.560 | 264.808 | .062 | -1664.176 | 24.117 |
| | | 4 | -2073.440 ^a | 502.107 | 264.808 | .000 | -3408.153 | -738.728 |
| | 4 | 1 | 2153.299 ^a | 502.107 | 264.808 | .000 | 818.587 | 3488.012 |
| | | 2 | 1253.411 | 502.107 | 264.808 | .079 | -81.302 | 2588.123 |
| | | 3 | 2073.440 ^a | 502.107 | 264.808 | .000 | 738.728 | 3408.153 |

Based on estimated marginal means

^a. The mean difference is significant at the .05 level.

a. Dependent Variable: RX_CRT_A.

c. Adjustment for multiple comparisons: Bonferroni.

Table A.41. Site * group interactions in cortical moment of resistance (y-axis). 1 = Control, 2 = LB, 3 = SSMD, 4 = MC3.

| site | (I) 1:CTRL, 2:LB, 3:SSMD, 4:MC3 | (J) 1:CTRL, 2:LB, 3:SSMD, 4:MC3 | Mean Difference (I-J) | Std. Error | df | Sig. ^a | 95% Confidence Interval for Difference ^b | |
|------|---------------------------------|---------------------------------|------------------------|------------|---------|-------------------|---|-------------|
| | | | | | | | Lower Bound | Upper Bound |
| 10 | 1 | 2 | -238.960 | 420.329 | 235.009 | 1.000 | -1357.364 | 879.443 |
| | | 3 | -102.133 | 420.329 | 235.009 | 1.000 | -1220.536 | 1016.270 |
| | | 4 | 532.460 | 664.599 | 235.009 | 1.000 | -1235.891 | 2300.810 |
| | 2 | 1 | 238.960 | 420.329 | 235.009 | 1.000 | -879.443 | 1357.364 |
| | | 3 | 136.827 | 420.329 | 235.009 | 1.000 | -981.576 | 1255.230 |
| | | 4 | 771.420 | 664.599 | 235.009 | 1.000 | -996.930 | 2539.771 |
| | 3 | 1 | 102.133 | 420.329 | 235.009 | 1.000 | -1016.270 | 1220.536 |
| | | 2 | -136.827 | 420.329 | 235.009 | 1.000 | -1255.230 | 981.576 |
| | | 4 | 634.593 | 664.599 | 235.009 | 1.000 | -1133.757 | 2402.944 |
| | 4 | 1 | -532.460 | 664.599 | 235.009 | 1.000 | -2300.810 | 1235.891 |
| | | 2 | -771.420 | 664.599 | 235.009 | 1.000 | -2539.771 | 996.930 |
| | | 3 | -634.593 | 664.599 | 235.009 | 1.000 | -2402.944 | 1133.757 |
| 25 | 1 | 2 | -83.810 | 420.329 | 235.009 | 1.000 | -1202.213 | 1034.593 |
| | | 3 | -70.038 | 420.329 | 235.009 | 1.000 | -1188.441 | 1048.365 |
| | | 4 | 269.921 | 664.599 | 235.009 | 1.000 | -1498.429 | 2038.272 |
| | 2 | 1 | 83.810 | 420.329 | 235.009 | 1.000 | -1034.593 | 1202.213 |
| | | 3 | 13.772 | 420.329 | 235.009 | 1.000 | -1104.632 | 1132.175 |
| | | 4 | 353.731 | 664.599 | 235.009 | 1.000 | -1414.620 | 2122.081 |
| | 3 | 1 | 70.038 | 420.329 | 235.009 | 1.000 | -1048.365 | 1188.441 |
| | | 2 | -13.772 | 420.329 | 235.009 | 1.000 | -1132.175 | 1104.632 |
| | | 4 | 339.959 | 664.599 | 235.009 | 1.000 | -1428.391 | 2108.310 |
| | 4 | 1 | -269.921 | 664.599 | 235.009 | 1.000 | -2038.272 | 1498.429 |
| | | 2 | -353.731 | 664.599 | 235.009 | 1.000 | -2122.081 | 1414.620 |
| | | 3 | -339.959 | 664.599 | 235.009 | 1.000 | -2108.310 | 1428.391 |
| 50 | 1 | 2 | 57.478 | 420.329 | 235.009 | 1.000 | -1060.925 | 1175.881 |
| | | 3 | -67.884 | 420.329 | 235.009 | 1.000 | -1186.287 | 1050.519 |
| | | 4 | 289.133 | 664.599 | 235.009 | 1.000 | -1479.217 | 2057.484 |
| | 2 | 1 | -57.478 | 420.329 | 235.009 | 1.000 | -1175.881 | 1060.925 |
| | | 3 | -125.362 | 420.329 | 235.009 | 1.000 | -1243.765 | 993.041 |
| | | 4 | 231.656 | 664.599 | 235.009 | 1.000 | -1536.695 | 2000.006 |
| | 3 | 1 | 67.884 | 420.329 | 235.009 | 1.000 | -1050.519 | 1186.287 |
| | | 2 | 125.362 | 420.329 | 235.009 | 1.000 | -993.041 | 1243.765 |
| | | 4 | 357.017 | 664.599 | 235.009 | 1.000 | -1411.333 | 2125.368 |
| | 4 | 1 | -289.133 | 664.599 | 235.009 | 1.000 | -2057.484 | 1479.217 |
| | | 2 | -231.656 | 664.599 | 235.009 | 1.000 | -2000.006 | 1536.695 |
| | | 3 | -357.017 | 664.599 | 235.009 | 1.000 | -2125.368 | 1411.333 |
| 75 | 1 | 2 | 212.521 | 420.329 | 235.009 | 1.000 | -905.882 | 1330.924 |
| | | 3 | 216.508 | 420.329 | 235.009 | 1.000 | -901.895 | 1334.911 |
| | | 4 | 260.749 | 664.599 | 235.009 | 1.000 | -1507.602 | 2029.099 |
| | 2 | 1 | -212.521 | 420.329 | 235.009 | 1.000 | -1330.924 | 905.882 |
| | | 3 | 3.987 | 420.329 | 235.009 | 1.000 | -1114.416 | 1122.390 |
| | | 4 | 48.228 | 664.599 | 235.009 | 1.000 | -1720.123 | 1816.578 |
| | 3 | 1 | -216.508 | 420.329 | 235.009 | 1.000 | -1334.911 | 901.895 |
| | | 2 | -3.987 | 420.329 | 235.009 | 1.000 | -1122.390 | 1114.416 |
| | | 4 | 44.241 | 664.599 | 235.009 | 1.000 | -1724.110 | 1812.591 |
| | 4 | 1 | -260.749 | 664.599 | 235.009 | 1.000 | -2029.099 | 1507.602 |
| | | 2 | -48.228 | 664.599 | 235.009 | 1.000 | -1816.578 | 1720.123 |
| | | 3 | -44.241 | 664.599 | 235.009 | 1.000 | -1812.591 | 1724.110 |
| 90 | 1 | 2 | -977.021 | 420.329 | 235.009 | .126 | -2095.424 | 141.382 |
| | | 3 | -29.280 | 420.329 | 235.009 | 1.000 | -1147.683 | 1089.123 |
| | | 4 | -2105.764 ^a | 664.599 | 235.009 | .010 | -3874.114 | -337.413 |
| | 2 | 1 | 977.021 | 420.329 | 235.009 | .126 | -141.382 | 2095.424 |
| | | 3 | 947.742 | 420.329 | 235.009 | .150 | -170.662 | 2066.145 |
| | | 4 | -1128.742 | 664.599 | 235.009 | .545 | -2897.093 | 639.608 |
| | 3 | 1 | 29.280 | 420.329 | 235.009 | 1.000 | -1089.123 | 1147.683 |
| | | 2 | -947.742 | 420.329 | 235.009 | .150 | -2066.145 | 170.662 |
| | | 4 | -2076.484 ^a | 664.599 | 235.009 | .012 | -3844.834 | -308.133 |
| | 4 | 1 | 2105.764 ^a | 664.599 | 235.009 | .010 | 337.413 | 3874.114 |
| | | 2 | 1128.742 | 664.599 | 235.009 | .545 | -639.608 | 2897.093 |
| | | 3 | 2076.484 ^a | 664.599 | 235.009 | .012 | 308.133 | 3844.834 |

Based on estimated marginal means

^a. The mean difference is significant at the .05 level.

a. Dependent Variable: RY_CRT_A

c. Adjustment for multiple comparisons: Bonferroni.

Table A.42. Site * group interactions in polar moment of resistance. 1 = Control, 2 = LB, 3 = SSMD, 4 = MC3.

| site | (I) 1:CTRL, 2:LB, 3:SSMD, 4:MC3 | (J) 1:CTRL, 2:LB, 3:SSMD, 4:MC3 | Mean Difference (I-J) | Std. Error | df | Sig. ^e | 95% Confidence Interval for Difference ^e | |
|------|---------------------------------|---------------------------------|------------------------|------------|---------|-------------------|---|-----------|
| 10 | 1 | 2 | -223.949 | 605.161 | 236.675 | 1.000 | -1834.051 | 1386.152 |
| | | 3 | -185.107 | 605.161 | 236.675 | 1.000 | -1795.209 | 1424.994 |
| | | 4 | 863.631 | 956.843 | 236.675 | 1.000 | -1682.163 | 3409.425 |
| | 2 | 1 | 223.949 | 605.161 | 236.675 | 1.000 | -1386.152 | 1834.051 |
| | | 3 | 38.842 | 605.161 | 236.675 | 1.000 | -1571.259 | 1648.943 |
| | | 4 | 1087.580 | 956.843 | 236.675 | 1.000 | -1458.214 | 3633.374 |
| | 3 | 1 | 185.107 | 605.161 | 236.675 | 1.000 | -1424.994 | 1795.209 |
| | | 2 | -38.842 | 605.161 | 236.675 | 1.000 | -1648.943 | 1571.259 |
| | | 4 | 1048.738 | 956.843 | 236.675 | 1.000 | -1497.056 | 3594.532 |
| | 4 | 1 | -863.631 | 956.843 | 236.675 | 1.000 | -3409.425 | 1682.163 |
| | | 2 | -1087.580 | 956.843 | 236.675 | 1.000 | -3633.374 | 1458.214 |
| | | 3 | -1048.738 | 956.843 | 236.675 | 1.000 | -3594.532 | 1497.056 |
| 25 | 1 | 2 | -18.967 | 605.161 | 236.675 | 1.000 | -1629.068 | 1591.134 |
| | | 3 | -38.033 | 605.161 | 236.675 | 1.000 | -1648.134 | 1572.068 |
| | | 4 | 357.260 | 956.843 | 236.675 | 1.000 | -2188.534 | 2903.054 |
| | 2 | 1 | 18.967 | 605.161 | 236.675 | 1.000 | -1591.134 | 1629.068 |
| | | 3 | -19.066 | 605.161 | 236.675 | 1.000 | -1629.167 | 1591.035 |
| | | 4 | 376.227 | 956.843 | 236.675 | 1.000 | -2169.567 | 2922.021 |
| | 3 | 1 | 38.033 | 605.161 | 236.675 | 1.000 | -1572.068 | 1648.134 |
| | | 2 | 19.066 | 605.161 | 236.675 | 1.000 | -1591.035 | 1629.167 |
| | | 4 | 395.293 | 956.843 | 236.675 | 1.000 | -2150.501 | 2941.087 |
| | 4 | 1 | -357.260 | 956.843 | 236.675 | 1.000 | -2903.054 | 2188.534 |
| | | 2 | -376.227 | 956.843 | 236.675 | 1.000 | -2922.021 | 2169.567 |
| | | 3 | -395.293 | 956.843 | 236.675 | 1.000 | -2941.087 | 2150.501 |
| 50 | 1 | 2 | 313.898 | 605.161 | 236.675 | 1.000 | -1296.203 | 1923.999 |
| | | 3 | 127.034 | 605.161 | 236.675 | 1.000 | -1483.068 | 1737.135 |
| | | 4 | 616.480 | 956.843 | 236.675 | 1.000 | -1929.314 | 3162.273 |
| | 2 | 1 | -313.898 | 605.161 | 236.675 | 1.000 | -1923.999 | 1296.203 |
| | | 3 | -186.864 | 605.161 | 236.675 | 1.000 | -1796.966 | 1423.237 |
| | | 4 | 302.582 | 956.843 | 236.675 | 1.000 | -2243.212 | 2848.375 |
| | 3 | 1 | -127.034 | 605.161 | 236.675 | 1.000 | -1737.135 | 1483.068 |
| | | 2 | 186.864 | 605.161 | 236.675 | 1.000 | -1423.237 | 1796.966 |
| | | 4 | 489.446 | 956.843 | 236.675 | 1.000 | -2056.348 | 3035.240 |
| | 4 | 1 | -616.480 | 956.843 | 236.675 | 1.000 | -3162.273 | 1929.314 |
| | | 2 | -302.582 | 956.843 | 236.675 | 1.000 | -2848.375 | 2243.212 |
| | | 3 | -489.446 | 956.843 | 236.675 | 1.000 | -3035.240 | 2056.348 |
| 75 | 1 | 2 | 342.107 | 605.161 | 236.675 | 1.000 | -1267.994 | 1952.209 |
| | | 3 | 301.281 | 605.161 | 236.675 | 1.000 | -1308.821 | 1911.382 |
| | | 4 | 509.914 | 956.843 | 236.675 | 1.000 | -2035.880 | 3055.707 |
| | 2 | 1 | -342.107 | 605.161 | 236.675 | 1.000 | -1952.209 | 1267.994 |
| | | 3 | -40.827 | 605.161 | 236.675 | 1.000 | -1650.928 | 1569.275 |
| | | 4 | 167.806 | 956.843 | 236.675 | 1.000 | -2377.988 | 2713.600 |
| | 3 | 1 | -301.281 | 605.161 | 236.675 | 1.000 | -1911.382 | 1308.821 |
| | | 2 | 40.827 | 605.161 | 236.675 | 1.000 | -1569.275 | 1650.928 |
| | | 4 | 208.633 | 956.843 | 236.675 | 1.000 | -2337.161 | 2754.427 |
| | 4 | 1 | -509.914 | 956.843 | 236.675 | 1.000 | -3055.707 | 2035.880 |
| | | 2 | -167.806 | 956.843 | 236.675 | 1.000 | -2713.600 | 2377.988 |
| | | 3 | -208.633 | 956.843 | 236.675 | 1.000 | -2754.427 | 2337.161 |
| 90 | 1 | 2 | -1425.749 | 605.161 | 236.675 | .116 | -3035.851 | 184.352 |
| | | 3 | -11.525 | 605.161 | 236.675 | 1.000 | -1621.627 | 1598.576 |
| | | 4 | -3724.253 [†] | 956.843 | 236.675 | .001 | -6270.046 | -1178.459 |
| | 2 | 1 | 1425.749 | 605.161 | 236.675 | .116 | -184.352 | 3035.851 |
| | | 3 | 1414.224 | 605.161 | 236.675 | .122 | -195.878 | 3024.325 |
| | | 4 | -2298.503 | 956.843 | 236.675 | .102 | -4844.297 | 247.290 |
| | 3 | 1 | 11.525 | 605.161 | 236.675 | 1.000 | -1598.576 | 1621.627 |
| | | 2 | -1414.224 | 605.161 | 236.675 | .122 | -3024.325 | 195.878 |
| | | 4 | -3712.727 [†] | 956.843 | 236.675 | .001 | -6258.521 | -1166.933 |
| | 4 | 1 | 3724.253 [†] | 956.843 | 236.675 | .001 | 1178.459 | 6270.046 |
| | | 2 | 2298.503 | 956.843 | 236.675 | .102 | -247.290 | 4844.297 |
| | | 3 | 3712.727 [†] | 956.843 | 236.675 | .001 | 1166.933 | 6258.521 |

Based on estimated marginal means

*. The mean difference is significant at the .05 level.

a. Dependent Variable: RP_CRT_A.

c. Adjustment for multiple comparisons: Bonferroni.

Table A.43. Site * group interactions in cortical area. 1 = Control, 2 = LB, 3 = SSMD, 4 = MC3.

| site | (I) 1:CTRL, 2:LB, 3:SSMD, 4:MC3 | (J) 1:CTRL, 2:LB, 3:SSMD, 4:MC3 | Mean Difference (I-J) | Std. Error | df | Sig. ^a | 95% Confidence Interval for Difference ^b | |
|------|---------------------------------|---------------------------------|-----------------------|------------|---------|-------------------|---|-------------|
| | | | | | | | Lower Bound | Upper Bound |
| 10 | 1 | 2 | -20.176 | 44.285 | 221.349 | 1.000 | -138.070 | 97.718 |
| | | 3 | -18.253 | 44.285 | 221.349 | 1.000 | -136.147 | 99.641 |
| | | 4 | 68.561 | 70.021 | 221.349 | 1.000 | -117.846 | 254.968 |
| | 2 | 1 | 20.176 | 44.285 | 221.349 | 1.000 | -97.718 | 138.070 |
| | | 3 | 1.923 | 44.285 | 221.349 | 1.000 | -115.971 | 119.817 |
| | | 4 | 88.737 | 70.021 | 221.349 | 1.000 | -97.671 | 275.144 |
| | 3 | 1 | 18.253 | 44.285 | 221.349 | 1.000 | -99.641 | 136.147 |
| | | 2 | -1.923 | 44.285 | 221.349 | 1.000 | -119.817 | 115.971 |
| | | 4 | 86.814 | 70.021 | 221.349 | 1.000 | -99.593 | 273.221 |
| | 4 | 1 | -68.561 | 70.021 | 221.349 | 1.000 | -254.968 | 117.846 |
| | | 2 | -88.737 | 70.021 | 221.349 | 1.000 | -275.144 | 97.671 |
| | | 3 | -86.814 | 70.021 | 221.349 | 1.000 | -273.221 | 99.593 |
| 25 | 1 | 2 | 7.840 | 44.285 | 221.349 | 1.000 | -110.054 | 125.734 |
| | | 3 | -5.800 | 44.285 | 221.349 | 1.000 | -123.694 | 112.094 |
| | | 4 | 36.224 | 70.021 | 221.349 | 1.000 | -150.183 | 222.631 |
| | 2 | 1 | -7.840 | 44.285 | 221.349 | 1.000 | -125.734 | 110.054 |
| | | 3 | -13.640 | 44.285 | 221.349 | 1.000 | -131.534 | 104.254 |
| | | 4 | 28.384 | 70.021 | 221.349 | 1.000 | -158.023 | 214.791 |
| | 3 | 1 | 5.800 | 44.285 | 221.349 | 1.000 | -112.094 | 123.694 |
| | | 2 | 13.640 | 44.285 | 221.349 | 1.000 | -104.254 | 131.534 |
| | | 4 | 42.024 | 70.021 | 221.349 | 1.000 | -144.383 | 228.431 |
| | 4 | 1 | -36.224 | 70.021 | 221.349 | 1.000 | -222.631 | 150.183 |
| | | 2 | -28.384 | 70.021 | 221.349 | 1.000 | -214.791 | 158.023 |
| | | 3 | -42.024 | 70.021 | 221.349 | 1.000 | -228.431 | 144.383 |
| 50 | 1 | 2 | 31.158 | 44.285 | 221.349 | 1.000 | -86.736 | 149.052 |
| | | 3 | 2.776 | 44.285 | 221.349 | 1.000 | -115.118 | 120.670 |
| | | 4 | 64.428 | 70.021 | 221.349 | 1.000 | -121.979 | 250.835 |
| | 2 | 1 | -31.158 | 44.285 | 221.349 | 1.000 | -149.052 | 86.736 |
| | | 3 | -28.382 | 44.285 | 221.349 | 1.000 | -146.276 | 89.512 |
| | | 4 | 33.270 | 70.021 | 221.349 | 1.000 | -153.137 | 219.677 |
| | 3 | 1 | -2.776 | 44.285 | 221.349 | 1.000 | -120.670 | 115.118 |
| | | 2 | 28.382 | 44.285 | 221.349 | 1.000 | -89.512 | 146.276 |
| | | 4 | 61.652 | 70.021 | 221.349 | 1.000 | -124.755 | 248.059 |
| | 4 | 1 | -64.428 | 70.021 | 221.349 | 1.000 | -250.835 | 121.979 |
| | | 2 | -33.270 | 70.021 | 221.349 | 1.000 | -219.677 | 153.137 |
| | | 3 | -61.652 | 70.021 | 221.349 | 1.000 | -248.059 | 124.755 |
| 75 | 1 | 2 | 14.942 | 44.285 | 221.349 | 1.000 | -102.952 | 132.836 |
| | | 3 | 10.616 | 44.285 | 221.349 | 1.000 | -107.278 | 128.510 |
| | | 4 | 4.188 | 70.021 | 221.349 | 1.000 | -182.219 | 190.595 |
| | 2 | 1 | -14.942 | 44.285 | 221.349 | 1.000 | -132.836 | 102.952 |
| | | 3 | -4.326 | 44.285 | 221.349 | 1.000 | -122.220 | 113.568 |
| | | 4 | -10.754 | 70.021 | 221.349 | 1.000 | -197.161 | 175.653 |
| | 3 | 1 | -10.616 | 44.285 | 221.349 | 1.000 | -128.510 | 107.278 |
| | | 2 | 4.326 | 44.285 | 221.349 | 1.000 | -113.568 | 122.220 |
| | | 4 | -6.428 | 70.021 | 221.349 | 1.000 | -192.835 | 179.979 |
| | 4 | 1 | -4.188 | 70.021 | 221.349 | 1.000 | -190.595 | 182.219 |
| | | 2 | 10.754 | 70.021 | 221.349 | 1.000 | -175.653 | 197.161 |
| | | 3 | 6.428 | 70.021 | 221.349 | 1.000 | -179.979 | 192.835 |
| 90 | 1 | 2 | -114.671 | 44.285 | 221.349 | .062 | -232.565 | 3.224 |
| | | 3 | 8.473 | 44.285 | 221.349 | 1.000 | -109.422 | 126.367 |
| | | 4 | -304.282 ^c | 70.021 | 221.349 | .000 | -490.690 | -117.875 |
| | 2 | 1 | 114.671 | 44.285 | 221.349 | .062 | -3.224 | 232.565 |
| | | 3 | 123.143 ^c | 44.285 | 221.349 | .035 | 5.249 | 241.038 |
| | | 4 | -189.612 ^c | 70.021 | 221.349 | .044 | -376.019 | -3.204 |
| | 3 | 1 | -8.473 | 44.285 | 221.349 | 1.000 | -126.367 | 109.422 |
| | | 2 | -123.143 ^c | 44.285 | 221.349 | .035 | -241.038 | -5.249 |
| | | 4 | -312.755 ^c | 70.021 | 221.349 | .000 | -499.162 | -126.348 |
| | 4 | 1 | 304.282 ^c | 70.021 | 221.349 | .000 | 117.875 | 490.690 |
| | | 2 | 189.612 ^c | 70.021 | 221.349 | .044 | 3.204 | 376.019 |
| | | 3 | 312.755 ^c | 70.021 | 221.349 | .000 | 126.348 | 499.162 |

Based on estimated marginal means

*. The mean difference is significant at the .05 level.

a. Dependent Variable: CRT_A

c. Adjustment for multiple comparisons: Bonferroni.