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2	Geometric morphometrics and Finite elements analysis: Assessing the functional	
3	implications of differences in craniofacial form in the hominin fossil record	
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26 Abstract:

The study of morphological variation in the hominin fossil record has been transformed in 27 28 recent years by the advent of high resolution 3D imaging combined with improved geometric morphometric (GM) toolkits. In parallel, increasing numbers of studies have applied finite 29 30 elements analysis (FEA) to the study of skeletal mechanics in fossil and extant hominoid material. While FEA studies of fossils are becoming ever more popular they are constrained 31 by the difficulties of reconstruction and by the uncertainty that inevitably attaches to the 32 estimation of forces and material properties. Adding to these modelling difficulties it is still 33 34 unclear how FEA analyses should best deal with species variation.

Comparative studies of skeletal form and function can be further advanced by applying tools 35 from the GM toolkit to the inputs and outputs of FEA studies. First they facilitate virtual 36 reconstruction of damaged material and can be used to rapidly create 3D models of skeletal 37 structures. Second, GM methods allow variation to be accounted for in FEA by warping 38 models to represent mean and extreme forms of interest. Third, GM methods can be applied 39 to compare FEA outputs - the ways in which skeletal elements deform when loaded. Model 40 comparisons are hampered by differences in material properties, forces and size among 41 models but how deformations from FEA are impacted by these parameters is increasingly 42 well understood, allowing them to be taken into account in comparing FEA outputs. 43

In this paper we review recent advances in the application of GM in relation to FEA studies
of craniofacial form in hominins, providing examples from our recent work and a critical
appraisal of the state of the art.

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48 Keywords: Form-function; Geometric Morphometrics; Finite Element Analysis; Craniofacial
49 form; Functional performance.

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1. Introduction

In this paper we consider how the mechanical performance of crania in biting can be estimated and compared among fossils, paying particular attention to how the methods of geometric morphometrics (GM) can facilitate such analyses in combination with biomechanical modelling using finite elements analysis (FEA).

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As with other skeletal elements, crania fulfil mechanical functions, such as housing and 58 protection of organs and provision of a rigid framework for food acquisition and intra-oral 59 processing by the masticatory apparatus (Lieberman, 2011), which comprises jaws, teeth and 60 soft tissues. Thus, much research has focused on the association between cranial form and 61 masticatory function, with the aims of understanding how crania function and how their 62 functional abilities (performances) differ among related species. These differences have 63 underpinned investigations of how skeletal form, function, ecology and behaviour interrelate 64 (Groning, et al., 2011b, Rayfield, 2005, Rayfield, 2007, Rayfield, et al., 2001, Strait, et al., 65 66 2010, Strait, et al., 2007, Strait, et al., 2009, Wroe, et al., 2010, Wroe, et al., 2007). In turn, knowledge of these interrelationships has been used to infer ecology and behaviour from 67 skeletal remains of extinct taxa (Attard, et al., 2014, Cox, et al., 2015, Degrange, et al., 2010, 68 69 Ledogar, et al., 2016, Oldfield, et al., 2012, Rayfield, 2005, Rayfield, et al., 2001, Smith, et al., 2015b, Strait, et al., 2010, Strait, et al., 2009, Wroe, 2008). 70

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One aspect of performance, bite force, can be directly measured in extant species using force 72 73 transducers. These have been widely used to measure bite forces in living humans (Braun, et 74 al., 1995, Kikuchi, et al., 1997, Paphangkorakit and Osborn, 1997, Sinn, et al., 1996). In 75 extinct material alternative approaches are required to estimate forces from skeletal evidence, using bony proxies to approximate lever arm lengths and maximum muscle forces based on 76 the relationships between muscle area, intrinsic muscle fibre strength and force production 77 (Gans and de Vree, 1987, Josephson, 1975, Weijs, 1980). Although this provides an estimate 78 of the force produced by muscles it ignores pennation and depends on the validity of the 79 estimates of muscle areas from bony proxies. Muscle force is converted into bite force 80 through the masticatory lever arm system. By measuring in and out-levers of the masticatory 81 system and computing their ratios, it is possible to estimate the mechanical efficiencies of 82 each muscle and to estimate maximal bite forces (Antón, 1990, Demes and Creel, 1988, Eng, 83

et al., 2013, O'Connor, et al., 2005). However, this approach has limitations due to 84 differences between the cross sectional areas estimated via bony proxies and actual 85 physiological muscle cross sectional areas (Eng, et al., 2013, Toro-Ibacache, et al., 2015), and 86 because the lever system of the jaws is often simplified to two dimensions to ease 87 calculations. Bite force can also be predicted using FEA (Wroe, et al., 2010) and by 88 multibody dynamic analysis (MDA) (Bates and Falkingham, 2012, Curtis, et al., 2008, Shi, et 89 al., 2012). MDA can also be used to infer muscle activation patterns given a particular load. 90 These approaches take full account of the three dimensional geometry of the masticatory 91 92 lever system but retain dependence on the accuracy of input of variables, such as muscle 93 forces, force vector directions and cranial geometry.

Bite forces are transmitted to items held between the teeth, and the teeth and cranium 94 experience the bite reaction force. Thus, crania have to be adapted to withstand masticatory 95 forces. In order to assess, explain and compare how the cranium resists occlusal forces, 96 researchers have used several approaches. These include the analysis of simplified 97 biomechanical models of craniofacial anatomy considered in terms of vertical and horizontal 98 99 column-like structures that buttress the face and channel bite reaction forces (Görke, 1904; Richter, 1920; Endo, 1965; Endo, 1966) and models that consider crania as a cylinder that is 100 101 twisted during biting (Greaves, 1985; Greaves and Mucci, 1997; Demes, 1987). These models and their underlying assumptions have been tested through the application of strain 102 gauges to directly measure the surface strains experienced during biting (Hylander et al., 103 1991; Hylander et al., 1992; Ross and Hylander, 1996; Ravosa et al., 2000a; Ravosa et al., 104 105 2000b; Ross, 2001; Ross et al., 2011). FEA has also been applied to this task (Ledogar, et al., 2016, Smith, et al., 2015b, Strait, et al., 2010, Strait, et al., 2007, Strait, et al., 2009, Wroe, et 106 al., 2010) but an important issue in such studies is validity of FEA results, do they match 107 reality? 108

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For this reason validation studies have been carried out to assess the accuracy of prediction of 110 111 cranial and mandibular deformations, comparing measured with predicted strains (Bright and Groning, 2011, Groning, et al., 2009, Kupczik, et al., 2007, Ross, 2005, Toro-Ibacache, et al., 112 2016). In so doing, researchers can assess how various input parameters including skeletal 113 geometry, material properties, constraints, applied forces etc. impact FE model results in 114 order to create more realistic models (Ross, 2005, Strait, et al., 2005, Toro-Ibacache, et al., 115 2016). In general validation studies tell us that accurate strain prediction in any one specimen 116 is difficult and requires careful adjustment of model parameters to achieve valid results. 117

The accuracy of predictions achieved by FEA is, however, entirely dependent on these input 119 parameters, so, acknowledging modelling limitations, researchers have sought to understand 120 the impact of variations and simplifications in FE modelling. This has led to sensitivity 121 studies in which the impact on predicted deformations of varying specific parameters is 122 123 assessed. Such parameters include the muscle force magnitudes, directions and activation patterns (Cox, et al., 2011, Fitton, et al., 2012, Groning, et al., 2012, Ross, 2005, Sellers and 124 Crompton, 2004), variations in material properties (Cox, et al., 2011, Groning, et al., 2012, 125 126 Kupczik, et al., 2007, Reed, et al., 2011, Strait, et al., 2005, Toro-Ibacache, et al., 2016), modelling cranial sutures (Kupczik, et al., 2007, Reed, et al., 2011, Wang, et al., 2010), 127 simplifications in model geometry (Fitton, et al., 2015, Toro-Ibacache, et al., 2016), 128 modelling the periodontal ligament (Groning, et al., 2012, Holland, 2013, McCormack, et al., 129 2014, Wood, et al., 2011), impact of variations in modelling of trabecular bone (Parr, et al., 130 2013) and model constraints (Cox, et al., 2011). Validation and sensitivity studies have 131 shown that modelling variations that affect model stiffness (e.g. bone thicknesses, how 132 133 cancellous bone is represented, material properties, etc.) or total applied force tend to lead to differences in magnitude rather than mode of deformation, while variations in relative muscle 134 135 activations, muscle vectors and constraints tend to impact mode of deformation, how the cranium deforms (Fitton, et al., 2015, Godinho, et al., 2017, Parr, et al., 2012, Toro-Ibacache, 136 et al., 2016). 137

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139 Researchers have tried to predict how fossil hominin crania resist biting. Such analyses have until recently relied on geometrical simplifications of crania (Demes, 1987, Rak, 1983, Rak, 140 1986, Trinkaus, 1987). More recently, FEA has been used to more fully model fossil hominin 141 masticatory biomechanics with the aim of improving prediction of the stresses and strains 142 experienced by fossil crania as they deform during biting (Ledogar, et al., 2016, Smith, et al., 143 2015b, Strait, et al., 2010, Strait, et al., 2009, Wroe, et al., 2010). Sensitivity studies are of 144 particular relevance here as validation is not possible for fossils. FEA applied to fossils has 145 become popular (Cox, et al., 2011, Cox, et al., 2012, Rayfield, 2007, Strait, et al., 2013, 146 Strait, et al., 2010, Strait, et al., 2007, Strait, et al., 2009, Wroe, et al., 2010, Wroe, et al., 147 2007) and models are frequently based on medical CT scans. However, another challenge of 148 fossils arises because they are often fragmented and invaded by sedimentary matrix that, due 149 to mineralization processes, is undistinguishable, or at least very difficult to distinguish, from 150 bone in scans. This often precludes, for example, segmentation of sedimentary matrix from 151

bone and does not allow fossils to be modelled reliably in terms of their full anatomical complexity. Indeed, given the multiscale organisation of bone, teeth and soft tissues, it is not within the reach of present technology to produce an accurately realistic model. Moreover, increasing model complexity demands higher computational power for solution (Groning, et al., 2012).

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Model simplification in geometry is therefore useful and necessary to overcome these 158 limitations (Fitton, et al., 2015). Assessment of the impact of simplifications typically relies 159 160 on comparison of variables of interest in subsequent FEA. Thus, researchers commonly focus on stress/strain magnitudes and directions and compare how different modelling decisions 161 impact on those variables (Groning, et al., 2012, Reed, et al., 2011, Strait, et al., 2005, 162 Szwedowski, et al., 2011, Wood, et al., 2011), although bite force has also been used to 163 assess model sensitivity (Fitton, et al., 2012, Sellers and Crompton, 2004). Beyond this, the 164 165 methods of geometric morphometrics have recently been applied to this task; to compare the deformations of variant models and estimate the impact of such simplifications on results 166 167 (Fitton, et al., 2015, Fitton, et al., 2012, Godinho, et al., 2017, Toro-Ibacache, et al., 2016).

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169 The application of GM methods to FEA output is discussed below in more detail, as it is

used for the reconstruction of fossils for FEA and in the creation of models of interesting realand hypothetical forms.

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2. How GM can help in reconstruction

Once the segmentation process is finished, reconstruction of missing anatomical regions 174 begins. This usually combines imaging software 175 process (e.g. Avizo/Amira/Mimics/Geomagic) and GM to approximately restore the original geometry of 176 an incomplete or distorted specimen (Weber, 2015, Weber and Bookstein, 2011). In 177 specimens that preserve one side intact, the most straightforward approach is to use bilateral 178 symmetry (Gunz, et al., 2009). In such cases it is possible to reflect the preserved regions 179 180 onto the incomplete side and use them to replace the missing areas (Gunz, et al., 2009). However, no skeletal structures are completely symmetric and they present different 181 magnitudes of asymmetry (Quinto-Sánchez, et al., 2015). Thus, reflected regions will not 182 perfectly fit the remaining preserved anatomy. To overcome this mismatch, and account for 183 asymmetry, it is possible to use the thin plate spline (TPS) function to warp the reflected 184

structure onto the remaining preserved anatomy (Gunz, et al., 2009). Even though this is a 185 desirable approach, fossils often lack preserved structures on both sides or along the midline, 186 thus precluding reflection. In these cases reference based reconstruction (Gunz, et al., 2004, 187 Gunz, et al., 2009) should be used. The choice of reference specimen should be considered 188 carefully so as to not bias the reconstruction and it has been suggested that references should 189 190 be species specific (Gunz, et al., 2009, Senck, et al., 2015, Zollikofer and Ponce de León, 2005). Such reconstructions may be statistical or geometric (Gunz, et al., 2004, Gunz, et al., 191 2009, Neeser, et al., 2009). Statistical reconstruction uses covariances among landmarks in a 192 193 given sample to predict the location of missing landmarks via multivariate regression (Gunz, et al., 2009, Neeser, et al., 2009). Geometric reconstruction uses the TPS function to estimate 194 the position of the missing landmarks based on known ones (Gunz, et al., 2004, Gunz, et al., 195 2009). The latter has the advantage of requiring one single specimen, which may be a 196 particular individual or a mean specimen calculated from a given sample using GM (Gunz, et 197 al., 2009) but omits information on intra specific covariations. However, Senck and 198 Coquerelle (2015) show that using mean specimens yields good results when reconstructing 199 200 large portions of incomplete specimens. Further where sample sizes are limited to one or a few specimens, as with fossils, TPS based warping can be applied, whereas statistical 201 202 approaches cannot.

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3. How GM can generate interesting hypothetical forms

Transforming an existing model into a target specimen is of significant value in allowing us 205 to visualise the results of GM analyses. To that end an original specimen may be landmarked 206 densely and then warped into a target that was landmarked similarly (O'Higgins, et al., 2011, 207 Stayton, 2009). Models that represent extremes of morphological variation within a taxon 208 may be created applying a similar approach. Such models can readily be used to simulate 209 mechanical loading and examine the impact of intra-specific morphological variance on 210 mechanical function (Smith, et al., 2015a). One major obstacle to using such an approach is 211 that accurate warping of one specimen into another requires many landmarks and 212 semilandmarks and even then, internal structures such as tooth roots, sinuses and cancellous 213 bone are unlikely to be warped to the form they would have in the target specimen. This is 214 because such internal architecture is very finely detailed and sinuses and cancellous bone 215 architecture are, to great extent, the result of adaptation in the specific individual to habitual 216 loading, this is not accounted for by warping alone. Any errors in warping will therefore 217

likely impact the resulting deformations of the FE model. An alternative is to warp 'solid' models, ones in which all that is represented is the geometry of the cranial surfaces with the spaces in between infilled with homogenous material. Cancellous bone and sinuses are filled and teeth are not represented as distinct structures, but merely as material with the properties of bone, and with roots merged with the surrounding bone. This is a drastic manoeuvre and a gross simplification. As such, the question arises as to what solid simplified models can tell us?

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226 Parr et al. (2012) examined the impact of infilling mandibular cavities on the deformations (bending displacements and strain magnitudes frequency) experienced by the mandible of a 227 varanoid lizard during simulated loading. They show that models with infilled cavities 228 deform less than, but generally similarly to, models with preserved cavities. Likewise, Fitton 229 et al. (2015) investigated the effects of simplifying details of internal anatomy 230 231 (presence/absence of the maxillary sinus) and material properties of teeth in a Macaca fascicularis cranium, concluding that it does not impact significantly on large scale 232 deformations but it does have localized effects in strain distributions. Toro-Ibacache (2016) 233 addressed the impact of segmentation protocols and of simplifying material properties of a 234 235 cadaveric human cranium. They concluded that segmentation protocols can have a significant impact on large scale deformations but that simplifying material properties (differentiating 236 trabecular bone from cortical bone vs not differentiating between the two) had little impact on 237 mode of deformation. Thus, if constraints and loads are held constant, solid models behave 238 239 similarly to much more detailed ones. The key difference emerging from these studies is that the solid models deform less, and so absolute magnitudes of deformation (measured as strains 240 or in terms of changes in size and shape; see below) are not accurately predicted while the 241 mode of deformation (how it deforms) is more consistent. This leads to the realisation that 242 solid models are useful in studies where absolute magnitudes of strains are of no interest but 243 rather, the focus is on mode, how a cranium deforms. 244

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These findings open up the possibility of carrying out many interesting 'virtual experiments' by warping or modifying skeletal anatomy to predict the functional role of particular features (O'Higgins, et al., 2011). Strait et al. (2007) applied this virtual experiment approach to infer the relevance of thick palates in Austalopiths by experimentally thickening the hard palate of a *Macaca fascicularis* and measure resulting strains. Fitton et al. (2009) reconstructed a specimen of *Austrolopithecus africanus* (STS 5) and warped the zygomatic region to that of a *Paranthropus boisei* (OH 5) while maintaining the remaining anatomy constant. In both
 cases, the impact of such modifications was assessed in terms of their impact on stresses and
 strains in the face.

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4. How GM can be used to compare FEA results

While stresses and strains from FEA are informative with regard to how skeletal structures bear loads and where they are likely to fail at a localized level (elements or nodes of elements) they do not allow ready assessment of how the model deforms as a whole, for instance, how it bends, twists and undergoes other changes in size and shape. Rather, such modes have to be inferred from strain contour maps based on expertise and knowledge.

GM, on the other hand, uses configurations of landmark coordinates and multivariate 262 statistics to assess how specimens differ in form, thereby quantifying morphological 263 differences in size and shape. Thus, it has been proposed that GM can be used to measure and 264 265 describe global deformations (defined here as changes in size and shape) of models under loading (O'Higgins, et al., 2011, O'Higgins, et al., 2012). This approach differs from GM 266 267 shape analyses in that size is also simultaneously considered, because loadings change the shape and the size of objects. The basis and application of this approach is described more 268 269 fully in section 5.3.

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5. Example Studies

To illustrate how the approaches described above are applied in practice, example studies arepresented and reviewed, below.

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5.1 Reconstruction of crania

We illustrate the application of GM to reconstruction using a CT scan of the cranium of Kabwe 1, which is remarkably well preserved but still presents missing and damaged anatomy due to taphonomic and pathological processes (Schwartz and Tattersall, 2003). Missing anatomical regions include a large portion of the right side of the cranial vault and base (affecting parts of the right temporal, parietal, zygomatic and occipital bone), a small region of the alveolus of the maxilla, teeth and small portions of the orbital cavities (Figure 1A). The reconstruction was based on a CT scan (courtesy of Robert Kruszynski, Natural History Museum, London) performed using a Siemens Somatom Plus 4 CT scanner, with voxel size of 0.47 x 0.47 x 0.50 mm and 140 kVp. Reconstruction started with the segmentation of the existing anatomy from the volume. Reconstruction of the left side of the vault followed, and this was later used to restore the large region missing from the right side of the cranium. Lastly, all remaining missing anatomical regions were reconstructed.

Segmentation was performed in Avizo 7.0 (Visualization Sciences Group Inc.) and used a 288 variety of approaches. The initial segmentation applied a half maximum height value 289 (HMHV; Spoor et al., 1993) to the whole volume to threshold segment it. Regional 290 291 thresholds were subsequently calculated and applied to specific anatomical regions as a second step because the HMHV did not segment thin bones. Manual segmentation was also 292 applied for fine details of thin bones that were not picked up by the two previous approaches. 293 Because teeth present clearly different grey values specific thresholds were calculated and 294 applied so as to not overestimate their dimensions. Last, existing sedimentary matrix was 295 296 removed manually.

298 Once the segmentation of existing structures was complete, the large missing region of the right half of the cranium was restored by reflecting the existing contralateral half and fitting 299 300 (warping) it to the existing structures. This last step used the TPS function and is necessary because crania are not absolutely symmetric. This warping is achieved by placing matching 301 302 landmarks on the damaged region and reflected fragment and then deforming (warping) the fragment to the cranium using the 'Bookstein' warping function. This resulted in an almost 303 304 perfect fit between the restored and preserved anatomy, requiring only minimal manual editing. Restoration of the damaged alveolar process of the right hemi-maxilla was also 305 achieved by reflecting the preserved left region. Existing gaps (such as the one present in the 306 orbital surfaces of the maxilla and ethmoid, internal nasal walls, maxilla, occipital bone, left 307 temporal bone, ethmoid bone and vomer) were restored using a combination of manual 308 editing and the software Geomagic Studio 2011 (courtesy of DR W. Sellers, University of 309 Manchester) to interpolate between existing bone edges. The missing posterior region of the 310 occipital bone was reconstructed using the occipital of a modern human cranium, which was 311 manually edited using Geomagic to adjust its morphology. Teeth were preferentially restored 312 by reflecting existing antimeres. When this was not possible portions of teeth from a modern 313 human were used to reconstruct incomplete teeth (final result of reconstruction in Figure 1B). 314

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317 Figure 1

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5.2 Hypothetical forms

321 FEA may be applied to any model, whether it represents a real specimen or not. For instance a prior GM analysis may have established the mean form and limits of variation of a 322 landmark configuration taken on a sample of crania. Rather than be interested in how any 323 particular specimens perform, we may be interested in the range of performances represented 324 by the sample. Earlier we noted that solid models, in which internal detail is grossly 325 simplified and filled, provide a reasonable basis for experimental manipulation of FE models 326 to assess specific questions such as the effects of varying palatal thickness or maxillary 327 morphology. This same principle can be extended to whole landmark configurations such as 328 those representing the limits of variation of a sample. By using triplets of thin plate splines to 329 330 warp whole crania between the mean and these limits of variation hypothetical crania can be created. They do not represent real crania but rather a statistical result from prior 331 morphometric analyses, in this case limits of variation but also, feasibly, through regression 332 333 or partial least squares (PLS), they could represent forms at the limits of cranial covariation with some interesting ecological or functional variables (e.g. climatic or dietary data, 334 335 measured bite forces etc.). FEA is then carried out on these hypothetical forms to see how the modes of variation of cranial form identified in the analysis impact performance when the 336 cranium is loaded. 337

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339 This warping approach is illustrated here using a simple example; the Kabwe 1 cranium warped into a mean Neanderthal (model 2) using thin plate splines based on classical and 340 sliding semi-landmarks. Classical landmarks (Figure 2, red spheres) of the mean Neanderthal 341 were calculated from 4 Neanderthal crania (Gibraltar 1, Guattari, La Chapelle-aux-Saints, La 342 Ferrassie). The sliding semi-landmarks on the maxilla and brow-ridge (yellow spheres) were 343 calculated from the 4 specimens and the sliding semi-landmarks of the vault and zygoma 344 (light blue spheres) were calculated from the 2 crania in which these structures are almost 345 completely preserved (Guattari and La Ferrassie). In Figure 2, the original model of Kabwe is 346 shown on the left (Figure 2A) and the warped 'mean Neanderthal' on the right (Figure 2B). It 347 is clear that a visually satisfactory result is obtained but of course internal architecture (tooth 348

roots, cortical thicknesses, cavities, sinuses and cancellous bone) will also be warped, not necessarily in such a way that they reasonably represent the average form in Neanderthals. However, by using 'solid' models as described above such errors are avoided. The FEA in such a circumstance does not aim to predict and compare actual deformations but rather it provides an answer to a different type of question: how do the differences in external form between these models impact mode and magnitude of deformation?

This approach is more limited than we may wish but it is useful in many contexts, for instance in considering how facial retraction vs projection, or brachycephaly vs dolichocephaly, or the mode of form variation predicted by e.g. climate or diet etc. impact on model performance. These are more general questions whose answer does not rest on study of single specimens, but rather on consideration of general modes of variation and their general effects.

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363 Figure 2

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5.3 Application of GM methods to the comparison of FEA results

367 As noted earlier a third way in which GM methods complement FEA is through comparison of deformations that occur due to loading (O'Higgins, et al., 2011, O'Higgins, et al., 2012). 368 This approach has been applied in several studies (Cox, et al., 2011, Fitton, et al., 2015, 369 Groning, et al., 2012, Groning, et al., 2011a, Holland, 2013, Prôa, 2013, Toro-Ibacache, et al., 370 2016). Such analyses of deformation rely on assessment of changes in model size and shape, 371 rather than of shape alone as is common in GM studies of organismal variation. This is 372 because as a model is loaded it changes in both size and shape, and it makes no sense to focus 373 on one aspect alone (shape or size). In consequence size and shape are analysed jointly, using 374 rescaled shape coordinates from GPA. The resulting size and shape distances between 375 unloaded and loaded forms describe the magnitudes of deformation and the direction of the 376 vector connecting unloaded and loaded forms in the size and shape space describes the mode 377 of deformation. These vectors can be compared among different load cases applied to the 378 same model or among different models by ignoring the differences in size and shape among 379 unloaded forms and focussing in the vectors connecting unloaded and loaded forms. 380

We illustrate the application of this approach by summarizing a study (Godinho, et al., 2017, 381 Toro-Ibacache, et al., 2016) that examines the impact of simplifications of a cadaveric Homo 382 sapiens cranium on the resulting modes of deformation predicted by FEA. Specifically, it 383 assesses the impact of simplifications among a three materials model (cortical bone, 384 cancellous bone and teeth; model 3), a two materials (cortical bone and teeth; model 2) and a 385 386 one material model (everything with material properties of cortical bone; model 1). Thus model 1 is a simple 'solid' model (see above) and model 3 is a much more anatomically 387 accurate model. The models are loaded to simulate a bite on the first molar, although the 388 389 applied forces are not physiological, rather they replicate the loading of an accompanying validation study to facilitate comparison with that in ongoing work. 390

The results in terms of strain contour plots (not shown), suggest that variations among models 391 generally impact on magnitudes of strains but not so much on the distribution of regions of 392 high and low strain throughout the model. The GM analysis of deformations complements 393 394 these findings. Size and shape distances are calculated by multiplying the shape coordinates (from GPA) of each specimen by that specimen's original centroid size. This results in the 395 396 specimens being represented by points in a (size and shape) space that can be thought of as the space of GPA aligned coordinates (Slice 2001), an approximation of Kendall's shape 397 398 space, with size as an additional dimension. The vector of centroid size (the additional dimension) at any point on the manifold can be visualised as passing radially from the 399 400 centroid of the manifold of this space (zero size), through the manifold (centroid size = 1) and beyond to infinity (infinite size). When centroid size is 1, the objects lie on the manifold of 401 the space of GPA aligned coordinates (Figure 3). The resulting space differs from the classic 402 size and shape space (Dryden and Mardia, 1998) that results from translating and rotating, but 403 not scaling landmark configurations. In particular, the rotations of configurations with respect 404 to each other differ because size influences rotation. In consequence the estimates of mean 405 size and shape (the size and shape variables; translated and rotated coordinates) obtained by 406 these two approaches, the resulting covariance matrix, and so PCA, also differ. However in 407 408 the application to FEA, where deformations are extremely small, the resulting size and shape differences are negligible. Either space could be used, with almost no difference in results, 409 410 but the approach we adopt is useful in understanding how shape analysis and size and shape analysis are related (Figure 3). Thus, simply making the model stiffer or less stiff (material 411 properties) or applying the same force vectors but varying their magnitudes results in greater 412 or less deformation; the vectors connecting unloaded and loaded models simply scale directly 413 with force or inversely with Young's modulus (a measure of stiffness). Deformations (size 414

and shape distances) also scale inversely with model centroid size if loads, geometry and 415 material properties are held constant. In contrast Procrustes distances scale inversely with the 416 square of centroid size. Figure 3 illustrates these scaling relationships with centroid size. 417 Thus if we take the shape of the black point (a; on the GPA hemisphere, centroid size =1; 418 Fig. 3) to be the unloaded form and the grey point (b) as the shape of the loaded form, the 419 distance between them represents the deformation in shape and approximates Procrustes 420 distance when variations are small. If size also differs due to loading, then the loaded form 421 does not lie on the hemisphere but is above (grey point, c) or below it depending on if it 422 423 increased or decreased in centroid size. The distance (a-c) between loaded and unloaded forms is the size and shape distance and is a measure of deformation (change in size and 424 shape with loading). 425

426

If the same forces and same material properties apply but the unloaded form is larger (black 427 428 point, d, on the outer semicircle representing the GPA hemisphere with centroid size >1 in Fig 3) the resulting deformation in size and shape is less (distance d-f; which in this diagram 429 430 is shown larger than in reality to facilitate visualisation). The physics dictate that the size and shape distances (deformations) between unloaded and loaded objects scale inversely with 431 432 centroid sizes of the unloaded objects; bigger forms deform less under the same load. However, shape change due to loading (Procrustes distance as opposed to size and shape 433 distance) scales inversely with the square of the centroid size of the unloaded object. Thus, 434 scaling the unloaded large object, d, to centroid size 1, results in it overlying point a, the 435 unloaded object with centroid size 1 (these are identical in shape but differ only in size). 436 Scaling the loaded large object, f, to centroid size 1 projects it along a radius (dashed line in 437 Fig. 3) through point e to an intersection, g, with the arc of the GPA hemisphere. As a result 438 of this scaling, the ratio of Procrustes distances (a-b)/(a-g) is the inverse of the ratio of the 439 squares of centroid sizes of the unloaded forms, a and d. 440

441

These scaling ratios are important because they allow us to account (at least approximately) for differences in size when comparing deformations predicted by FEA among similar objects using geometric morphometric methods. Such scaling is inevitably an approximation unless the objects whose deformations are being compared are the same shape, differing only in size. As shapes become more different, it makes less sense to compare deformations and the degree of approximation in scaling increases.

449 Figure 3

450 451

452 Principal components analysis using the covariance matrix among size and shape variables can be used to visualise and compare deformations. Figure 4 presents the first two principal 453 components from the sensitivity study we conducted on a Homo sapiens of model 454 simplification with regard to segmentation and allocation of material properties. These 455 456 account for some 99% of the total variance and so fairly represent the results. Included in the analysis are the unloaded model and the three variants of model segmentation (model 1, 457 458 whole model as cortical bone = 17 GPa; model 2, bone = 17 GPa and teeth = 50 GPa; model 3, cortical bone = 17GPa, cancellous bone = 56 MPa, teeth = 50 GPa; all materials allocated a 459 Poisson's ratio of 0.3) after loading in a simulated first molar bite. Model 3 is also loaded in a 460 simulated incisor bite. This allows the effects of simplification to be compared against the 461 effect of varying bite point. The molar bites cluster away from the incisor bite, indicating 462 they are more similar in mode of deformation. The modes of variation are represented by the 463 vectors connecting unloaded and loaded models. They are visualised by the inset warped 464 surface models and transformation grids, computed using thin plate splines and magnified 465 500 times to facilitate interpretation. Models 2 and 1 overlie each other and so are represented 466 467 by a single point. This implies that representing dental roots as cortical bone has little effect. Modes of deformation differ greatly between incisor and molar bites and consist mainly of 468 upwards deflection of the anterior maxilla in the former and of the lateral maxilla in the latter. 469 With regard to the molar bites, simplification has its greatest impact when cancellous bone is 470 allocated the material properties of cortical, effectively making a 'solid' model. The effect is 471 to reduce the degree of deformation as is reflected in the shorter vector connecting models 1 472 and 2 with the unloaded than that connecting model 3. Similarly the degree of deformation 473 evident in the inset warpings is reduced. There is a difference in mode of deformation as 474 evidenced by the angle between these vectors but the difference in mode is less obvious in 475 comparing the inset warping for models 2 and 3 with that for model 1. 476

The impact of simplification on mode of deformation is small compared to the largedifference between molar and incisor bites.

479

481 Figure 4

482 483

This simple analysis can be extended to more complex and interesting questions concerning 484 485 multiple variants of models and to the comparison of deformations among models (O'Higgins and Milne, 2013) by focusing on differences among vectors of deformation rather than 486 differences among unloaded forms. This application of GM is particular useful in relation to 487 FEA sensitivity studies providing an easily visible and quantifiable approach to the 488 assessment of model "error" and sensitivity. Comparisons can be made within and between 489 models to assess whether differences in performance due to modelling assumptions are 490 491 drowning out any meaningful biological signals.

492

493 **6.** Discussion

The last three decades have seen an explosion of advances in techniques pertinent to the 494 495 study of skeletal change through time. Morphometrics underwent a revolution (Adams, et al., 2004, Rohlf and Marcus, 1993) beginning in the late 1970's (Bookstein, 1978) and gathering 496 497 pace through the next three decades. This, in common with the tools for high resolution 498 imaging, visualisation and manipulation of images, took great advantage of the advances in computing that occurred over the same period. These same advances in computational power 499 led to the development of increasingly sophisticated software tools for FEA, to simulate and 500 501 predict the effects of loadings on structures.

502

All of these tools are in common use today in the field of Archaeology, in particular they 503 have been driven by work on fossil material, but increasingly they are applied to more recent 504 skeletal finds, archaeobotany (García-Granero, et al., 2016, Ros, et al., 2014), zooarchaeology 505 (Cucchi, et al., 2011, Evin, et al., 2013, Owen, et al., 2014) and to material culture such as 506 ancient architecture (Levy and Dawson, 2009), stone tools (Buchanan and Collard, 2010, 507 Buchanan, et al., 2011, Okumura and Araujo, 2014), and pottery (Hein, et al., 2008, 508 Kilikoglou and Vekinis, 2002, Wilczek, et al., 2014). As these tools have become more 509 commonly applied, useful ways of combining them have come to the fore. Thus, GM 510 methods combined with tools for imaging and image manipulation can play an important role 511 in the reconstruction of skeletal material, as illustrated by the first example we present in this 512 513 paper.

Reconstruction provides data for morphometric analyses and GM has proven powerful in this 515 domain. Beyond the fact that these methods provide approaches that are statistically robust 516 and well understood, GM's great advantage for many workers lies in the ability to visualise 517 the results of statistical analyses as warpings of the mean form. These visualisations close the 518 loop between measurement, statistics and interpretation of results in terms of changes in size 519 and shape. They also can be used to produce 3D models of the results of statistical analyses 520 such as mean forms or forms representing extremes of variation or extremes of interesting 521 522 modes of variation, such as the limits of regressions of form on ecological, behavioural and functional data. In turn, such forms are potentially interesting targets for FEA. For instance, 523 the results of a study of how cranial form covaries with the toughness of diet might be 524 actualised as a pair of 3D surfaces representing crania at the limits of the regression; those 525 suited to tough and those suited to less tough diets. These can be submitted to FEA to explore 526 how each responds to loading and, in this way, link modes of morphological variation to load 527 resistance. This kind of analysis provides a very direct way of exploring how form and 528 function interact. 529

530

These kinds of studies depend critically on the validity of FEA modelling and on how sensitive such modelling is to errors in model building and loading. Validity is assessed by comparing predicted strains with directly measured, real strains. Sensitivity, on the other hand is assessed by varying model parameters to replicate likely errors and comparing results among (often many) variant models as in the example we provide earlier. In this endeavour, GM has been usefully combined with FEA.

537

538 Size and shape analysis allows ready understanding of the effects of different model building 539 decisions in terms of how the models deform and how they differ in deformation. It leads to 540 statements about changes in form such as how a skull twists or bends under loading. If, 541 instead of landmarks the coordinates of all nodes of the finite element mesh are submitted to 542 analysis, strains can be computed from the coordinates of the unloaded and loaded meshes.

543 On the other hand the concentrations of stresses or strains in localised regions are useful 544 predictors of failure and while the overall distribution of strains can be used to infer global 545 modes of deformation this requires simultaneous interpretation of one contour map for each 546 principal strain (2 in 2D analyses and 3 in 3D). Thus the approaches can be considered 547 complementary. GM analyses inform with regard to how loading deforms an object, in terms

of changes in size and shape. On the other hand, stresses and strains inform with regard to the 548 likelihood of failure in particular anatomical regions and how distributions of high and low 549 strains may eventually relate to (re)modelling fields. Each can be used to infer the other, thus 550 GM analyses that use the coordinates of the nodes of an FE mesh result in visualisations of 551 deformations of that mesh and strains can be computed and displayed in these meshes, while 552 strains can be used to infer global degrees and modes of deformation. It should be noted that 553 the metric of the GM analysis, the Procrustes size and shape distance relates to changes in 554 size and shape, not the risk of failure. It differs from metrics derived from strains. These 555 556 describe different aspects of deformation that are complementary, but not coincident, in 557 interpreting FEA results.

558

Beyond sensitivity analyses, GM methods have been extended to the task of comparing 559 deformations among different specimens modelled and loaded in equivalent ways (Milne and 560 O'Higgins, 2012, O'Higgins and Milne, 2013). In this case the analysis focuses on differences 561 in vectors of deformation rather than the differences between unloaded forms. This is 562 563 achieved by computing deformations as differences between registered loaded and unloaded forms. They can be visualised by adding these vectors to a convenient unloaded form such as 564 565 the mean of all unloaded specimens. Beyond comparisons of deformation, it is also possible to use GM approaches to assess the association between deformation under loading and other 566 interesting factors such as skeletal form or ecological variables through regression of PLS 567 analyses. This has not yet been much exploited in the literature but is likely to increasingly be 568 taken up as a useful approach to the interpretation of the biological significance of differences 569 in modes and magnitudes of variation. Differences among models in size, applied forces and 570 material properties can be taken into account by 'correcting' the magnitudes of deformations 571 according to the known scaling relationships described above. 572

573

The application of GM methods in conjunction with FEA is as yet in its infancy. There are 574 clear roles for GM methods in reconstruction, the production of models with modified 575 geometry to explore how form and function interact and in comparing the results of FEAs 576 among models and load cases. Each of these approaches has potential benefits and pitfalls 577 and, in time, with increasing numbers of studies applying these methods we will better 578 understand where they are applicable and where not. At present these combined GM/FEA 579 approaches are still fairly novel and some years of methodological development can be 580 581 anticipated.

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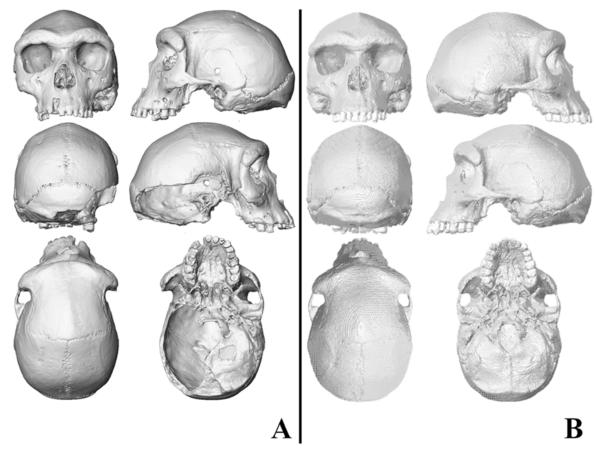
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863 Figure 1: Cranium of Kabwe 1 (A) before and (B) after reconstruction.

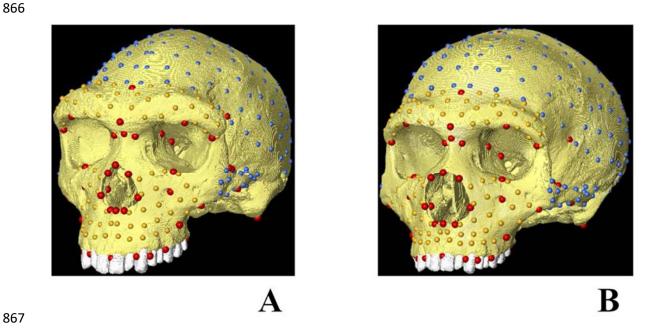
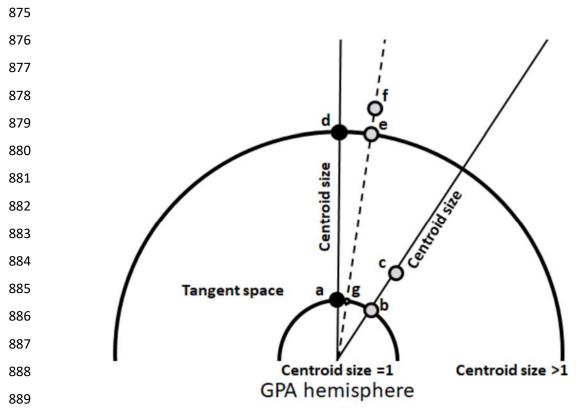


Figure 2: A reconstructed Kabwe 1, *Homo heidelbergensis* cranium model (A) and a hypothetical *Homo neanderthalensis* (B). The hypothetical Neanderthal was created via surface warping model A into a mean Neanderthal landmark data set (B) using thin plate splines based on classical and sliding semi-landmarks. Conventional landmarks (red spheres); sliding semi-landmarks on the maxilla and brow-ridge (yellow spheres); sliding semilandmarks of the vault and zygoma (light blue spheres).



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Figure 3: A schematic illustrating the hemisphere of GPA aligned coordinates (Slice 2001) for triangles, showing the tangent space, the vectors of centroid size, and the size and shape space resulting from rescaling of GPA registered coordinates to centroid size >1. The black points, a and b, represent the same unloaded forms with different centroid sizes. The grey points c and f represent the loaded forms and the grey points b and e represent their projections onto the GPA hemispheres with centroid sizes =1 and >1. G is the projection of f and e onto the GPA hemisphere with centroid size =1. See text for explanation.

Figure 4: PCA of large scale deformations of a sensitivity study assessing the impact of
simplifications of material properties in a modern human cranium. Model 1 is not visible
because it is in the same location in the plot as model 2. Deformations are magnified by a
factor of 500 to facilitate visualisation.