AN APPLICATION OF THE RELATIVE WARPS ANALYSIS TO PROBLEMS IN HUMAN PALEONTOLOGY - WITH NOTES ON RAW DATA QUALITY

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ABSTRACT

This study investigates the use of geometric morphometrics as well as methodological aspects specifically related to its application in paleoanthropology. Based on lateral photographs taken from a fossil sample of 58 specimens, a relative warps analysis was computed in order to assess the variation of cranial shape among various hominin groups. The fossil sample represents Middle and Late Pleistocene populations commonly assigned to H. erectus, H. neanderthalensis, archaic H. sapiens as well as anatomically modern H. sapiens. The preliminary results indicate that several fossil skulls considered as belonging to archaic H. sapiens have a distinct shape compared to modern humans. The results suggest that these hominins are not as closely related to modern humans as previously thought. Instead, their morphometric affinities suggest that they are as distinct from modern humans as are the Neandertals. Methodological aspects, such as raw data accuracy and the use of type 2 and 3 landmarks that are directly related to this type of quantitative analysis and that potentially affect their results, are discussed.

Keywords: cranial shape, deficient landmarks, geometric morphometrics, paleoanthropology, photographic distortion.

INTRODUCTION

Questions related to modern human origins have largely dominated the paleoanthropological output of the last decade (Nitecki and Nitecki, 1994; Wood, 1994). Despite ongoing discussions, the idea of an African origin of modern humans appears to emerge as a general consensus, although certain details about this process remain open for debate. For instance, several authors claim a single African origin of modern humans in the sense of a speciation event (Cann et al., 1987; Stringer, 1994; Chaline, 1998), while Bräuer (1984; Bräuer et al., 1997) suggested a model of an African origin with more gradual elements. According to this model, modern humans evolved around 200,000 to 100,000 years bp (before present) in South and East Africa from Homo erectus-like, archaic populations that are represented by specimens like Nduhu, Kabwe, Omo 2, Eley Springs, Singa and others and referred to ‘archaic Homo sapiens’. Around 100,000 years bp, specimens like Omo 1 (Ethiopia) as well as Skhul and Qafzeh (Israel) are thought to represent the earliest anatomically modern human forms. This series of fossils is referred to as the African transition (from archaic to modern humans).

In opposition to the multiregional model (Wolpoff et al., 1984), which will not be addressed here, both African scenarios consider that Neandertal populations from Europe and the Near East were marginal to or even excluded from the origin and dispersal of modern humans.

The purpose of this paper is to investigate the variation of cranial size and shape in Middle and Late Pleistocene fossil hominins from Europe, Asia and Africa using geometric morphometrics. Although uncertainties often remain regarding the absolute age of fossil remains, the sample of this study covers a time range between ca. 1 million years and 10,000 years bp. Within the background of the modern human origins debate, the initial idea of this study is to test the hypothesis of a gradual evolution from early archaic Homo sapiens to anatomically modern humans in Africa, the model proposed by Bräuer (1984; Bräuer et al., 1997). Here, in addition to reporting results and discussing their phylogenetic implications, we will emphasize a few methodological
issues that in our view are often associated with shape analysis of fossil hominins.

The morphometric analysis of biological shapes is a common approach to assessing affinities of organisms in general. According to Rohlf and Marcus (1993), geometric morphometric techniques are particularly powerful in that they combine the usual multivariate statistics with the geometry of the object and therefore graphically illustrate shape differences. Its application in paleoanthropology should therefore contribute to the quantification of evolutionary trends among fossil populations.

**MATERIAL AND METHODS**

In order to study the variation of cranial shape during the end of the Pleistocene, raw data were collected from 58 fossils, dating from the Middle Pleistocene at or slightly below 1 million years (*Homo erectus*) to the Late Pleistocene, about 10,000 bp (European and Near Eastern Upper and Epipaleolithic crania, Table 1). During this relatively long time range, non-modern hominin forms existed throughout the Old World, as is witnessed by Neandertals in Europe, and *Homo heidelbergensis* or archaic *Homo sapiens* in Africa and Europe. It is also the time period, during which the earliest modern Humans are believed to have appeared, somewhere by the end of the Middle Pleistocene, beginning of the Late Pleistocene (dated to 125,000 bp).

**SHAPE ANALYSIS USING GEOMETRIC MORPHOMETRICS**

Bookstein (1991) introduced a series of techniques for the statistical analysis of shape variation, now commonly referred to as geometric morphometrics. Within this framework, a biological shape is assessed through homologous landmarks that summarize the geometry or form of the organism under study. The landmark configuration of each specimen is then normalized for translation, rotation and scaled to a common size, so that the remaining differences between individual or mean shapes are differences in shape only. This is achieved by Procrustes superimposition, a least squares type of registration method that minimizes the distance between pairs of homologous landmarks. Once landmark configurations are Procrustes aligned, shape variables can be extracted either by a GPA (Generalized Procrustes Analysis, Rohlf, 2000) or, as in this paper, through a relative warps analysis (Bookstein, 1991). These shape

<table>
<thead>
<tr>
<th>Group</th>
<th>Relative age</th>
<th>General morphology</th>
<th>N</th>
<th>Original specimens and casts (*)</th>
</tr>
</thead>
<tbody>
<tr>
<td>Asian <em>H. erectus</em> (&lt;1 MY)</td>
<td>Middle morphology</td>
<td>1</td>
<td>Sangiran 2</td>
<td></td>
</tr>
<tr>
<td>Archaic <em>Homo sapiens</em> / <em>H. heidelbergensis</em> (90,000 – 400,000 BP)</td>
<td>Non-modern groups</td>
<td>6</td>
<td>Eliye Springs*, Kabwe, LH 18*, Ndutu*, Omo 2*, Singa</td>
<td></td>
</tr>
<tr>
<td><em>H. neanderthalensis</em> (34,000 – 170,000 BP)</td>
<td>Middle – Late Pleistocene</td>
<td>12</td>
<td>Amud 1, La Chapelle-aux-Saints, La Ferrassie 1 Forbes’ Quarry, Guattari 1, Neandertal, La Quina H5, Saccopastore 1+2, Saint-Césaire, Shanidar 1*, Tabun C</td>
<td></td>
</tr>
<tr>
<td>Earliest anatomically modern <em>Homo sapiens</em> (90,000 – 130,000 BP)</td>
<td>Middle – Late Pleistocene</td>
<td>6</td>
<td>Omo 1, Qafzeh 6+9, Skhul 4+5+9</td>
<td></td>
</tr>
<tr>
<td>Early Upper Paleolithic (35,000 – 20,000 BP)</td>
<td>Modern groups</td>
<td>8</td>
<td>Abri Pataud; Brno 2; Cro-Magnon 1-3; Mladec 1-3</td>
<td></td>
</tr>
<tr>
<td>Late Upper Paleolithic/Epipaleolithic (≤20,000 BP)</td>
<td>Late Pleistocene</td>
<td>25</td>
<td>Arene Candide 1, 4, 5, &quot;Prince&quot;, n. num; Avelines Hole; Barma Grande 2, 3; Cheddar; Nahal Ein Gev 1; Eraq el-Almar H2; Fallah 16, 23; Hayonim 4; Langwith Cave; Le Bichon; Mallaha 37; Oberkassel 1, 2; Ohalo II; Rond du Barry; San Teodoro 3; Uzzo IVa, b; Veyrier 1</td>
<td></td>
</tr>
<tr>
<td>Total</td>
<td></td>
<td>58</td>
<td></td>
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</tr>
</tbody>
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Table 1. Fossil specimens, and their time range, used in this study.
variables can further be used as input to standard uni-
and multivariate analyses and the observed variation
and mean differences can be visually represented in
the picture plane in which the raw data (i.e.,
landmarks) were originally recorded. Besides the
improved statistical parameterization of size and
shape, the approach also strongly enhances the
biological interpretation of statistical observations.

The relative warps analysis (Bookstein, 1991) was
performed using the software tpsRelw (Rohlf, 1997).
Relative warps analysis corresponds to a Principal
Components analysis of the covariance matrix of the
partial warp scores, which are different scales of a
thin-plate spline transformation of landmarks. The thin-
plate spline is a smooth interpolation function that
computes and visualizes transformations of Cartesian
Coordinates in a way similar to D’Arcy Thompson’s
transformation grids (Thompson, 1917). A rectangular
grid is projected over Procrustes aligned landmark
configurations and the bending of the grid visually
depicts the difference in landmark locations between
2 configurations. The Thin-plate spline uses the
analogy of an infinitely thin metal sheet that is bent
so that points on it move from their original position
to a defined target position. The closer two positions
are, the more bending energy is required, whereas
large scale transformations require less energy. Shape
differences between any two landmark configurations
are visualized as landmark displacement, while the
changes in the space between landmarks are
interpolated. Every specimen of a sample is compared
to the mean (or consensus) configuration of the
sample, and shape changes along any axis of interest
(here, the relative warps, see Figs. 4-7) are visualized
as deviations from the mean configuration. Given its
similarity with a Principal Components Analysis,
relative warps are a tool for exploring within-sample
variation serving to reduce the total variation to a
smaller number of independent dimensions. Typically,
the first few components or relative warps summarize
most of the variation of a sample.

Subsequently, size allometry (i.e. shape change
as a function of size) was estimated by calculating
 correlations between relative warp scores and
centroid size. Centroid size, the square root of the
sum of the squared distances between all homologous
landmarks and the center of gravity of the landmarks,
is commonly used as general size measure in
geometric morphometrics and was computed with
GRF-ND (Slice, 1994). For all statistics, except the
relative warps, a standard package of statistical
software was used (Statistica version 5.1, StatSoft
France, 1998). The landmarks were all collected in
the lateral view and correspond to six different data
sets (Frieß, 1999). This procedure takes into account
specific variation of fossil humans, and is the best
way to cope with their generally poor state of
preservation, since estimating missing values remains
critical. Furthermore, some of the landmarks included
are bilateral points, and by default they were taken on
one side only. For modern humans, this assumes
symmetry which is known to hardly apply to any
living organism having a symmetry axis. However,
given that in fossils both sides are virtually never
preserved, it was impossible to address the question
of asymmetry in this study.

Fig. 1. Schematic illustration of the landmarks used for this study. These landmarks correspond to standard
cranioemetric points except nos. 3, 5 and 7 in the vault configuration, which are so-called deficient landmarks.
Most of the raw data were recorded from original specimens, but six original specimens were not available and had to be replaced with casts. Based on lateral outlines of these crania, up to 19 landmarks were defined and digitized using a PC and the tpsdig software (Rohlf, 1996) (type 1 to 3, see Fig. 1). A type 1 landmark is a point that is unambiguously defined (typically by the juxtaposition of tissues) and therefore can be claimed to be homologous. A type 2 landmark commonly is a curvature maximum, its homology is therefore based on the geometry rather than the anatomy. Finally, a type 3 landmark is an extremal landmark, e.g. one end of a diameter etc. It is considered to be a deficient landmark, because its coordinates depend on some kind of orientation or construction. And therefore considered to be of limited use in geometric morphometrics (Bookstein, 1991).

**RAW DATA ACCURACY**

As was emphasized in the introduction, an additional goal of this paper is to raise a few methodological questions with regard to the use of 2D representations of 3D objects (i.e., pictures of skulls) as well as previous applications of geometric morphometrics in paleoanthropology (Yaroch, 1996).

As far as the use of 2D geometric morphometrics is concerned, one possible way of obtaining raw data for such analyses is by using drawings or photographs commonly used to document the fossil record. Photographs, however, may not always be suitable for metric comparisons because of the well known principle of optical distortion (Fig. 2, cf. Jacobshagen et al., 1988). Any optical lens leads to distortion of the absolute and relative dimensions of a spherical object like a skull. The outer edge of the sphere will be invisible in a central projection, affecting absolute dimensions. Secondly, objects that are closer to the focal point will appear relatively bigger than they are compared to portions that are further away, affecting thereby the proportions of the object. In practical terms this means that taking a picture of a human cranium with a distance of 1 meter, not uncommon for published high-quality illustrations, results in a distortion of about 7.5%, which is hardly acceptable. These distortions can be mostly avoided by stereophotographic techniques. Alternatively, if no 3D devices are available, increasing the distance between the camera and the object can significantly reduce the photographic error. In order to obtain accurate cranial outlines in 2D, a simple standardized photogrammetric protocol with a maximal distance of 6.5 meters to the specimen was used. The specimen was positioned exactly in the optical axis of the camera, and its symmetry axis was aligned perpendicular to the optical axis. Camera and center of volume of the specimen were at level along 2 axes, the third being defined by the distance between them. This distance varied between 5 m and 6.5 m and was merely dependent on local settings, knowing that the optical distortion at 5 m and more is below 2%. The mean deviation between standard measurements taken on the crania and the same set of measurements taken on the photographs was found to be less than a millimeter (Frieß, 1997), which is considered as an acceptable error in anthropometry. These considerations allow us to hypothesize that raw data that are derived from pictures and taken without the necessary respect of basic photogrammetry are less reliable. The following example (Fig. 3) which refers to a study based on published photographs of fossil hominins (Yaroch, 1996) may serve to illustrate this issue. This figure shows the thin-plate spline from a standardized picture (Frieß, 1999) of a Neandertal cranium (Amud 1) to a published picture of the same specimen (Suzuki and Takai, 1970). In this specific case the optical distortion tends to reduce certain traits that are usually common in Neandertals, such as (Fig. 3):

- the alveolar prognathism, i.e., projecting upper jaw,
- the strong supraorbital torus (brow ridges),
- the relatively flat vault, especially in the parietal region and
- the occipital bun in the back of the skull.

Fig. 2. (after Jacobshagen et al. 1988): Optical distortion due to a central projection affects measurement accuracy at two levels: The maximum diameter of a given object (dimension G1) is only partly visible to the camera on the right, and therefore erroneously measured smaller than in reality (dimension G2). The ratio between any dimension situated between the maximum diameter and the camera (G3/G1), is incorrectly reproduced on
the film (B3/B2). The values for the expected error are based on the dimensions of a human skull.

These distortions could then lead to the, in our view erroneous, impression that Neandertals are actually ‘less Neandertal-like’ than generally thought, simply because their cranial shape was insufficiently reproduced in these pictures (see c.f., Yaroch, 1996 who noted an absence of a clearly distinct Neandertal cranial shape compared to modern humans).

CRANIOFACIAL SHAPE VARIATION DURING THE MIDDLE AND LATE PLEISTOCENE

The first analysis is based on 10 landmarks of the vault. The first and second axis of the relative warps analysis account for 60% of the variation, with the first component explaining 49.1% (Fig. 4a). This first axis allows for a very clear distinction between two poles: The modern cranial shape, represented by Upper Paleolithic and Epipaleolithic fossils (recovered from Europe and the Near East and dated between 35,000 and 10,000 years) as well as the Middle Paleolithic Skhul/Qafzeh group (Israel, ca. 100,000 years bp); and a non-modern group including Neandertals (170,000 – 35,000) and archaic Homo sapiens (400,000 – 90,000). The only H. erectus specimen (Sangiran 2) is clearly part of this group. Note that some of the specimens grouped here as archaic Homo sapiens have also been classified as Homo heidelbergensis, and are often seen as ancestors of Neandertals and/or Homo sapiens (Tattersall, 1995; Rightmire, 1996). The second axis separates the Neandertal lineage from the rest of the archaic specimens, although some overlap exists along this second axis.

The main shape difference in the vault that is associated with the relative warps is visualized as landmark displacement using the thin-plate spline function (Fig. 5a). This reveals for the whole non-modern group along the first axis a flat vault, which is horizontally elongated and exhibits a strong supraorbital torus and a pronounced occipital convexity. The second axis (Fig. 5b) seems to differentiate among the more archaic (non-modern) crania those who show an occipital bun, often considered a Neandertal character, from those who do not exhibit such a bun.

Judging from the correlations between these warps and centroid size (Fig. 4b), the first axis appears to depend moderately on size ($r = 0.35$, $p = 0.009$). Given that significant size differences are only observed between 2 major clusters, modern and non-modern fossils, the observed allometry therefore mainly concerns differences between Homo erectus, Neandertals and archaic Homo sapiens on the one hand and anatomically modern humans on the other. A
strong correlation ($r = 0.6$, $p = 0.002$) among the non-modern groups was also observed and may reflect the size differences between early and classic Neandertals. No significant correlations ($r = -0.34$, $p > 0.05$) were observed among the fossil crania with modern shape (dating from the Upper Paleolithic and Epipaleolithic).

As far as facial variation is concerned, the differences between modern humans and more archaic populations are much weaker, as is shown in Fig. 6a. In fact, although archaic faces all cluster together due to negative scores on both first and second relative warps (33 and 17%), they overlap with clearly modern faces, such as the Natoufian Mallaha 37 or the Skhul/Qafzeh group, which is considered to represent some of the earliest anatomically modern humans. Here, they are grouped closely together with Neandertals, a non-modern form that occupied Europe between 170,000 and 35,000 years ago and that is believed to have disappeared without descendents.

The shape change associated with these axes (Figs. 5c and 5d) confirms for the group of non-modern faces the presence of a general prognathism (a projecting upper jaw), a strong supraorbital torus (brow ridges) as well as a zygomatic retreat (cheek). Such a facial morphology has been described as typical for Neandertals (Rak, 1986; Trinkaus, 1987). Our data show that fossils of clearly modern vault shape tend to exhibit this supposedly typical Neandertal facial morphology, which leads us to reconsider its status as ‘Neandertal-like’. Furthermore, this combination of facial characters is also present in Kabwe, considered by some authors as archaic $H. sapiens$, while some others associate it with $H. heidelbergensis$, a possible ancestor of Neandertals. Our findings support the latter model. The overlap of modern and non-modern facial morphologies can be explained by allometry, as is illustrated in the biplot of relative warp 1 against centroid size. This relation is seen in Fig. 6b, and one can observe that the correlation is very strong ($r = -0.78$) and highly significant ($p < 0.001$). From this, it can be proposed that some aspects of Neandertal facial morphology are merely a result of allometry (shape change due to size increase), which at the same time could explain the presence of this plesiomorphic (ancestral) trait in clearly modern fossils like the Skhul/Qafzeh group, as these specimens have higher scores for centroid size.

Certain phylogenetic considerations of this study can be emphasized. Comparing the evolution of size and shape among later Pleistocene hominins by performing a relative warps analysis and subsequent allometry study indicates that most of the so-called African transitional specimens (except for Omo 1 and Singa, see below) are very distinct from the modern human cranial shape (represented by Skhul/Qafzeh and the Upper to Epipaleolithic samples) and do not show any trend towards the anatomically modern populations, except for size (Frieß, 1999). In fact, when plotted against generally accepted absolute dates, archaic $Homo sapiens$ was found to exhibit an increase in overall size. Size increase, however, is of less phylogenetic relevance because it is also observed in the Neandertal lineage. Therefore these results do not support a model of gradual evolution of $Homo sapiens$ in Africa. Most of the so-called transitional African hominins appear to retain a generally plesiomorphic craniofacial shape and are as distinct from modern humans as are Neandertals. Conversely, the data are consistent with either a more rapid speciation event during the African later Pleistocene (Stringer, 1994), or with a Near Eastern origin of modern humans (Vandermeersch, 1981; Mann, 1995), as exemplified by the Skhul/Qafzeh hominins. However, support for the former model from this study can only be seen in a fairly limited number of fossils (Omo 1, Singa) and therefore requires further investigation using the approach presented here. Moreover, as far as facial morphology is concerned, essential aspects of Neandertal facial morphology can be considered as widely
plesiomorphic and dependent on size, rather than a derived character.

Fig. 6. Relative warps analysis of the face of Middle and Late Pleistocene hominins, M37 = Mallaha 37, Sk4 = Skhul 4, Sk5 = Skhul 5, Q6 = Qafzeh 6, Q9 = Qafzeh 9. a) individual scores on relative warps 1 and 2, b) biplot of relative warp 1 against centroid size (highly significant overall allometry, $r = -0.78$).

METHODOLOGICAL ASPECTS OF LANDMARK-BASED METHODS APPLIED TO FOSSIL CRANIA

These preliminary results need to be checked by extending the sample as well as the analysis itself to 3-dimensional comparisons. A further problem, specific to the study of the human cranium, lies in the use of landmarks for the description of shape. Bookstein's recommendation to use type 1 landmarks whenever possible (or type 4 landmarks, see Bookstein, 1997), i.e., points that can be defined unambiguously and without any geometric or orientational reference, is difficult to respect while studying human crania. In fact, one needs more than points of suture intersections (type 1 landmarks), in particular because certain characters, for instance the occipital bun or the parietal flattening, are not only highly relevant from the phylogenetic point of view, but also impossible to describe with type 1 landmarks. This consideration leads to the question whether the use of type 2 and 3 landmarks alters the shape change described in this study.

In order to evaluate potential effects of deficient landmarks, different configurations of a reduced set of fossil specimens ($n = 15$) were explored using the relative warps. In total, 4 different sets with a variable proportion of the three landmark types were analyzed, adding more and more deficient landmarks to the successive configurations (for details and landmark definitions, see Frieß, 1999). The first configuration, using type 1 and 2 landmarks, covers points across the complete skull. However, the overall coverage of vault shape is rather incomplete, because the vault itself provides very few type 1 landmarks. Judging from the plot of the first two axes and the associated shape change (Fig. 7a), it is confirmed that Neandertals have a very distinct basic cranial geometry, mainly due to a vertical extension of the anterior portion relative to the posterior parts. There is also a clear flattening of the cranium in the bregma region and at lambda. In the second configuration (analysis 2), more landmarks were added, such as maximum curvature points, referred to as type 2 as well as deficient or type 3 landmarks (landmarks 5, 6 and 8). As one can notice in Fig. 7b, the variation among this restricted fossil sample remains the same, with Neandertals showing very different scores in the first plane from modern humans. The thin-plate spline confirms what is found in the first configuration, i.e. a vertical extension of the face and frontal, with a clear flattening around bregma. It is evident that the newly added points supply additional information on other portions of the cranium, but the basic shape change as well as population variation remains unchanged. The third configuration is identical to the one used for the main study in this paper. It describes the cranial vault with all three landmark types, including deficient landmarks on the frontal (landmarks 2 and 3), the parietal (landmark 5) and the occipital (landmark 7). There is a persistent lambdoid flattening in Neandertals (landmark 6), as well as the general flattening of the vault, while a vertical extension is this time restricted to the supraorbital region (Fig. 7c). Finally, the full cranial shape variation including the lower jaw (landmarks 16 to 19), with about half type 1 and half deficient landmarks (Fig. 7d), provides a very similar result in terms of group differences and shape change compared to analyses 1 to 3. All analyses indicate that our subsample contains two
major poles of cranial morphology, Neandertals on the one hand and modern humans on the other, which is only the consequence of our non random selection for this subsample. What is noteworthy here is that all four analyses reveal the same major within-sample variations, despite the fact that they are based on different landmark types and configurations. Therefore, these comparative analyses indicate that the use of deficient landmarks in these particular configurations does not alter the results, as far as observable shape change and variations among populations are concerned. Whether the same can be said for the statistical power of these analyses remains to be tested in a differently designed study.

Fig. 7. Relative warps analyses with a variable number of type 1, 2 and 3 landmarks, using a restricted sample of Neandertals and Upper/Epipaleolithic modern Humans a) cranium b) portions of the vault and the face using type 1 to 3 landmarks c) cranial vault d) cranium including the lower jaw, using type 1 to 3 landmarks.
CONCLUSION

The purpose of this paper was twofold: First, to provide new insight into the variation of craniofacial variation among later Pleistocene hominins. Second, to discuss methodological issues inherent to the use of photography on the one hand and the use of deficient landmarks on the other, both being common in paleoanthropology. As far as the phylogenetic aspects of this study are concerned, two points can be emphasized: Although the term "archaic Homo sapiens" implies a relatively close relationship to modern Homo sapiens, various fossil remains assigned to this taxon do not show any particular affinity to later anatomically modern humans. Instead, their cranial and facial shape appears closer to Neandertals, which is consistent with assigning some of them to the species Homo heidelbergensis (Rightmire, 1996; Tattersall, 1995). Very few African fossils (Omo 1, Singa) as well as the specimens from Skhul and Qafzeh in Israel possess a clearly modern cranial shape. Given their clear morphometric distance to other "archaic Homo sapiens", the concept of gradual evolution to modern humans in Africa is not supported.

The methodological aspects of this study allow us to suggest that the relative warps analysis appears to be rather robust even when geometrically dependent landmarks are used. However, this conclusion is only preliminary, since it is restricted to a very specific sample of human crania. Other biological objects may give very different results, but they may not share the crucial lack of clearly defined landmarks. Furthermore, with the recently introduced concept of sliding landmarks (Bookstein, 1997) the problem of deficient landmarks may become obsolete, once the methods are implemented in the available software. Insufficient raw data accuracy, on the other hand, appears to be a more important, yet apparently insufficiently recognized source of biased results.

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Softwares

