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Modularity of the wrist in extant hominines

Ana Bucchi^{1,2,*}, Thomas A. Püschel^{3,4}, Carlos Lorenzo^{5,2}

- ¹ Centro de Investigación en Odontología Legal y Forense (CIO), Facultad de Odontología, Universidad de La Frontera, Temuco, Chile
- ² Institut Català de Paleoecologia Humana i Evolució Social (IPHES), Zona Educacional 4, Campus Sescelades URV (Edifici W3), 43007 Tarragona, Spain
- ³ Ecology and Evolutionary Biology Division, School of Biological Sciences, Philip Lyle Building, University of Reading, RG6 6BX, Reading, United Kingdom
- ⁴ Primate Models for Behavioural Evolution Lab, Institute of Cognitive and Evolutionary Anthropology, School of Anthropology, University of Oxford, 64 Banbury Road, OX2 6PN, Oxford, United Kingdom
- ⁵ Universitat Rovira i Virgili, Departament d'Història i Història de l'Art, Avinguda de Catalunya 35, 43002 Tarragona, Spain

* Corresponding author: ana.bucchi@ufrontera.cl

With 4 figures and 4 tables

Abstract: Wrist shape varies greatly across primates and previous studies indicate that the numerous morphological differences among them are related to a complex mixture of phylogeny and function. However, little is known about whether the variation in these various anatomical differences is linked and to what extent the wrist bones vary independently. Here, we used 3D geometric morphometrics on a sample of extant hominines (Homo sapiens, Pan troglodytes, Gorilla gorilla, and Gorilla beringei), to find the model that best describes the covariation patterns among four of the eight carpals (i.e., capitate, lunate, scaphoid, and trapezium). For this purpose, 15 modular hypotheses were tested using the Covariance Ratio. Results indicate that there is a covariation structure common to all hominines, which corresponds to stronger covariation within each carpal as compared to the covariation between carpals. However, the results also indicate that that there is a degree of codependence in the variation of some carpals, which is unique in humans, chimpanzees, and gorillas, respectively. In humans there is evidence of associated shape changes between the lunate and capitate, and between the scaphoid and trapezium. This covariation between lunate and capitate is also apparent in gorillas, while chimpanzees display the greatest disassociation among carpals, showing low covariation values in all pairwise comparisons. Our analyses indicate that carpals have an important level of variational independence which might suggest a high degree of independent evolvability in the wrists of hominines, and that although weak, the structure of associated changes of these four carpals varies across genera. To our knowledge this is the first report on the patterns of modularity between these four wrist bones in the Homininae and future studies might attempt to investigate whether the anatomical shape associations among carpals are functionally related to locomotion and manipulation.

Keywords: modularity; trait covariation; wrist; hominines

Introduction

The wrist in hominines is composed of eight bones with complex shapes and numerous joint surfaces, which allow the hand to move along multiple axes (Kivell et al. 2016). Genetically, a common Hox gene expression regulates the development of the hand in anthropoids (Reno et al. 2008), yet carpals also have a degree of functional and evolutionary independence (Tocheri et al. 2003; Kivell et al. 2013). This functional and evolutionary independence may explain why carpal morphology varies so greatly across taxa (Tocheri et al. 2005; Marzke et al. 2010; Orr 2017).

Among primates, humans exhibit a derived carpal morphology (Kivell et al. 2016), which previous studies suggest evolved as a consequence of relaxed locomotor pressures with the advent of bipedalism and as an adaptation to tool making and use (Hamrick et al. 1998; Williams et al. 2010; Key & Dunmore 2015; Skinner et al. 2015; Kivell et al. 2016). Wrist morphology in humans contributes significantly to stone tool-making performance (Tocheri et al. 2003; Marzke et al. 2010; Williams et al.; Williams et al. 2010; Williams et a

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of joint surfaces at the trapezium, capitate, and radiocarpal joints (Marzke 1983; Marzke 1997; Niewoehner et al. 1997; Richmond & Strait 2000; Tocheri et al. 2003; Tocheri et al. 2005; Marzke et al. 2010; Williams et al. 2010; Williams et al. 2014; Orr 2017). The characteristic joint surfaces in the human wrist allow for increased accuracy (Williams et al. 2014) and mechanical work at the joint during stone tool production (Williams et al. 2010). They also allow toolmakers to effectively resist and transmit both axial and oblique joint reaction forces generated by power and precision grips as compared to the rest of the extant apes (Marzke 1983; Niewoehner et al. 1997). Conversely, the wrist in chimpanzees and gorillas seems better adapted to locomotor demands, by contributing to better stabilization at the joint (Tuttle 1967; Richmond & Strait 2000) and by allowing the joint to better withstand the stresses imposed by knuckle walking (Püschel et al. 2020).

Several previous studies have analyzed single bones and specific joint surfaces with the aim of inferring the functional capabilities that set apart hominins from non-human primates (e.g., Tocheri et al. 2003; Tocheri et al. 2005; Marzke et al. 2010; Kivell 2011). However, with some exceptions (Williams 2010; Peña 2018; Bardo et al. 2020), there are almost no studies analyzing whether the numerous shape variations in wrist bones are associated or independent with respect to each other. Peña (2018) proposes that the level of integration of the wrist is higher in some primate genera (i.e., Pongo) than others, suggesting that specific covariation patterns may be shaping the evolution of this structure in primates. For humans, previous studies indicate that the morphological integration of autopods is lower than in quadrupeds, making the human hand more evolvable (Rolian 2009; Rolian et al. 2010; Young et al. 2010). In particular, the covariance structure observed between the shape and orientation in the facets present on the trapezium and the first metacarpal suggest they are biomechanically adventageous for the habitual use of forceful precision grips (Bardo et al. 2020). However, Williams (2010) indicates that the patterns of integration of the capitate and third metacarpal are more similar between humans and gorillas than between gorillas and chimpanzees, and that knuckle-walkers are not characterized by highly integrated morphologies.

The mutual relationships between bony elements of a single structure are best studied within the framework of modularity as they allow us to know how flexible the evolution of this anatomical region is under differing functional demands. If all carpals behave as a single entity that is tightly integrated by strong interactions, they should comprise a module (Klingenberg 2008; Esteve-Altava 2017), thus causing wrist bones to covary strongly. Conversely, if more than one module is present in the wrist, this should cause carpals in different modules to vary independently. It is currently unknown how many modules there are in the primate wrist, and how strong the modular signal is.

Our analysis intends to address the question of how independent the variation within the wrist is by analyzing the modularity pattern of four carpals in extant hominines (i.e., the capitate, trapezium, lunate, and scaphoid). As far as we know, this is the first time that the covariation structure for these bones has been reported for modern humans (Homo sapiens), chimpanzees (Pan troglodytes), and gorillas (Gorilla gorilla and Gorilla beringei). 3D models and geometric morphometrics were used for this purpose, and modularity was investigated through the testing procedure proposed by Adams and Collyer (2019), known as the covariance ratio effect sizes (Z_{CR} and \hat{Z}_{12}). We tested 15 different modular hypotheses combining all possible partitions of the wrist bones and selected the one that best describes the covariation structure in hominines as a whole, and in humans, chimpanzees, and gorillas in particular. In doing so, we try to answer two main questions: a) what is the modularity pattern of these four bones in living hominines? and b) is the observed covariation pattern shared across the analyzed taxa? We hypothesize that humans exhibit a pattern of covariation that distinguishes them from African apes, based on previous studies suggesting that manipulation has driven the evolution of the wrist in humans (e.g., Williams et al. 2010; Key & Dunmore 2015; Skinner et al. 2015), while in apes its better adapted for locomotion (e.g., Richmond & Strait 2000; Püschel et al. 2020).

Materials and methods

Primate sample

The sample comprises 478 bones from three primate genera: 50 modern humans (*Homo sapiens*), 41 chimpanzees (*Pan troglodytes*), and 41 gorillas (19 *Gorilla gorilla* and 22 *Gorilla beringei*) (Table 1). 3D models came from different sources. All human surface models were obtained using a Breuckmann SmartSCAN structured light scanner (Breuckmann Inc.). Most non-human primate surface models were generated via photogrammetry (further details can be found in Bucchi et al. 2020), while CT scans of 23 ape hands were accessed from two different digital repositories: Morphosource (www.morphosource.org) and the Museum of Primatology (https://carta.anthropogeny.org/).

The resolutions of micro-CT, surface scanner, and photogrammetric models have been previously tested and found to be comparable (Giacomini et al. 2019) thus allowing us to combine these data types in our analyses. The human hands belonged to a medieval cemetery (Burgos, Spain) (Casillas García & Adán Álvarez 2005) and the non-human sample were of different origins (wild shot, in captivity, and of unknown provenance). Right hands were preferred. Most of the wrists included the four carpals under analysis, and when there were some missing bones, their antimeres, when present, were reflected using the 'Flip and/or Swap

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Table 1. The study sample. * TM = trapezium, SC = scaphoid, CA = capitate, LU = lunate. ** UBU: Universidad de Burgos, AM: AfricaMuseum, IPHES: Catalan Institute of Human Palaeoecology and Social Evolution, MZB: Natural Sciences Museum in Barcelona, and ZSM: Zoological State Collection in Munich.

Species	Specimens	Carpal bones*			Sex			Collection**		
		ТМ	SC	СА	LU	Male	Female	Unknown		
Homo sapiens	50	40	39	41	42	25	25	0	UBU	
Pan troglodytes	41	38	41	40	38	16	14	11	AM, ZMB, ZCM,	
Gorilla beringei	22	22	22	22	21	10	9	3	Morphosource, Museum of	
Gorilla gorilla	19	17	18	19	18	5	5	9	Primatology	
Total	110	117	120	122	119	56	53	24		

axis' and 'Invert faces orientation' tools in Meshlab software (v. 2020.02) (Cignoni et al. 2008).

These carpals (i.e., the capitate, trapezium, lunate, and scaphoid) were chosen because they were best represented in the collections, although not all individuals had all of these bones (some elements were missing in some cases; further details can be found in Table 1 and in Suppl.-Table S1). Although studying all 8 carpals would give a broader pattern of modularity, it would decrease the probability of finding individuals with all carpals present in already limited primate collections, and it also would affect the statistical power of the analyses (see below).

Landmark configuration

We acquired five fixed landmarks per bone (Fig. 1, Table 2). We decided to collect a limited number of landmarks based on the known problems that occur in geometric morphometrics when the ratio of variables to cases is extremely high (e.g., when using excessive semi-landmarks and a limited sample size) (e.g. Cardini 2019). To assess whether sufficient number of landmarks have been sampled to characterize shape variation, we used the *lasec()* function of the LaMBDA R package (Watanabe 2018). We randomly sampled 10 capitates and collected additional 30 landmarks (i.e., 35 landmarks in total per specimen). Our results show that an adequate shape characterization is already achieved with 5 landmarks (see Suppl. Fig. 1). This means that our landmarking protocol is sufficient to characterize carpal shape.

As some of the selected carpals articulate (capitate, lunate and scaphoid and scaphoid and trapezium; Fig. 1), careful was taken in no to duplicate information when designing the landmark configuration, so landmarks representing shared facets were placed only in one of the two bones. In order to not distort the covariation strength among carpals that articulate, the same numbers of landmarks (two) were place in each of the facets of the articulating bones. On the other hand, trapezium does not articulate with capitate and lunate and this raises the question of whether the shared facets among the other carpals drive the covariation patterns observed in our study. This possibility was taken into consideration when analyzing the results. Landmark coordinates were imported into R using the *Arothron* package version 1.1.1 (Profico et al. 2018) in R 1.2.5019 (R Core Team 2019). A generalized Procrustes analysis (GPA) was then performed separately for each bone in order to normalize for location, rotation, and scale. Corrected coordinates were then compiled into a new dataframe, and hypotheses of modularity were tested (see below).

Allometry

Taxonomic differences in size can affect the pattern and magnitude of modularity (Klingenberg & Marugán-Lobón 2013). Therefore, we tested for allometric signals in the data by using a regression of Procrustes shape variables on centroid size. This test was performed with the *procD.lm()* function of the *geomorph* package, version 3.2.1 (Adams et al. 2019). A MANOVA test revealed that there are no significant differences in the shape of none of the carpals studied here (p > 0.05) between *G. gorilla* and *G. beringei*, thus in order to have balanced sample sizes, this and all further statistical analyses were carried out by pooling both gorilla species together.

Modular hypotheses

We tested 15 different hypotheses of modularity corresponding to all possible partitions of the sample (Table 3). We defined one four-module model (H1), seven two-module models (H2-8), six three-module models (H9-H14), and one single-module model (H15). The optimal modular hypothesis for the wrist was assessed by measuring the strength of covariation for each modular hypothesis with the covariance ratio (CR) (Adams 2016) and then statistically comparing alternative modular hypotheses with the covariance ratio effect sizes (Z_{CR} and \hat{Z}_{12}) (Adams & Collyer 2019).

Covariance ratio (CR)

The covariance ratio (CR) (Adams 2016) was computed to measure the degree of modular signal in two or more *a priori* modules of Procrustes shape variables. The CR coefficient calculates the ratio of the overall covariation between modules relative to the overall covariation within modules. The



Fig. 1. The landmark configuration shown on specimen AM 998 (*Gorilla beringei*) for the capitate, trapezium, lunate, and scaphoid bones. Landmark definitions are provided in Table 2.

CR coefficient ranges from 0 to positive values. CR values lower than 1 indicates low covariation between modules, and strong covariation otherwise. The significance of the CR coefficient is assessed via permutations. At each repetition, landmarks are randomly assigned to different modules and the CR coefficient is calculated. Finally, the original CR value is then compared to the CR distribution (Adams 2016).

Comparing the strengths of the modular signals $(Z_{CR} \text{ and } \hat{Z}_{12})$

The covariance ratios effect size (Z_{CR}) is derived from the CR and is a standardized test statistic which ensures statis-

tical compatibility with the CR (Adams & Collyer 2019) (Table 1). When the observed CR is larger than expected under the null hypothesis of no modularity, the Z_{CR} exhibits greater negative values which indicates a stronger modular signal. Here, whether Z_{CR} values are statistically different from each other was evaluated using a two sample Z-score for comparing modular signals (\hat{Z}_{12}). Both metrics are needed to compare alternative modular hypotheses. Z_{CR} was calculated for all modular hypotheses and the model presenting the strongest modular signal (i.e., the lowest Z_{CR}) was selected as the optimal modular hypothesis for all samples, and for each genus separately. Once the best hypothesis was

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Table 2. Definitions of landmarks digitalized in this study.

Bone	Landmark	Position					
Capitate	1	Most anterior point of the union between the facets for the second and third metacarpals.					
	2	Most distal and posterior point of the union of the facets for the hamate and the lunate.					
	3	Most distal point of the facet for the hamate.					
	4	Most inferior and anterior point of the union of facets for the hamate and the lunate.					
	5	Point of maximum curvature of the lunate-scaphoid facet.					
Trapezium	1	Point of maximum curvature of the ridge of the trapezium.					
	2	Most anterior point of the facet for the second metacarpal.					
	3	Most lateral and proximal point of the facet for the first metacarpal.					
	4	Most anterior point of the union between the facets for the trapezoid and the scaphoid.					
	5	Most posterior point of the union between the facets for the trapezoid and the scaphoid.					
Lunate	1	Most posterior and distal point of the facet for the scaphoid.					
	2	Most anterior and distal point of the facet for the scaphoid.					
	3	Most anterior point of the intersection between the facets for the triquetral and the hamate.					
	4	Most posterior point of the intersection between the facets for the hamate and capitate.					
	5	Point of maximum curvature of the facet for the radius.					
Scaphoid	1	Most posterior point of the facet for the radius.					
	2	Most anterior point for the facet for the radius.					
	3	Point of maximum curvature of the tubercle of the scaphoid.					
	4	Most medial point of the facet for the capitate.					
	5	Most lateral point of the facet for the capitate.					

Table 3.	The 15 modular hypotheses te	ted in this study. CA = capitate, L	U = lunate, SC = scaphoid, and	d TZM = trapezium.
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Model hypotheses	Modules	Description
H1	CA-LU-SC-TZM	All carpals belong to different modules.
H2	CALU-SCTZM	The capitate and lunate belong to one module and the scaphoid and trapezium to another.
H3	CASC-LUTZM	The capitate and scaphoid belong to one module and the lunate and trapezium to another.
H4	CATZM-LUSC	The capitate and trapezium belong to one module and the lunate and scaphoid to another.
H5	CA-LUSCTZM	The capitate belongs to one module and the lunate, scaphoid, and trapezium to another.
Н6	LU-CASCTZM	The lunate belongs to one module and the capitate, scaphoid, and trapezium to another.
H7	SC-CALUTZM	The scaphoid belongs to one module and the capitate, lunate, and trapezium to another.
H8	TZM-CALUSC	The trapezium belongs to one module and the capitate, lunate, and scaphoid to another.
Н9	CALU-SC-TZM	There are three modules: one includes the capitate and lunate, the second includes the scaphoid, and the third includes the trapezium.
H10	CASC-LU-TZM	There are three modules: one includes the capitate and scaphoid, the second includes the lunate, and the third includes the trapezium.
H11	CATZM-LU-SC	There are three modules: one includes the capitate and trapezium, the second includes the lunate, and the third includes the scaphoid.
H12	LUSC-CA-TZM	There are three modules: one includes the lunate and scaphoid, the second includes the capitate, and the third includes the trapezium.
H13	LUTZM-CA-SC	There are three modules: one includes the lunate and trapezium, the second includes the capitate, and the third includes the scaphoid.
H14	SCTZM-CA-LU	There are three modules: one includes scaphoid and trapezium, the second includes the capitate, and the third includes the lunate.
H15	CALUSCTZM	All carpals belong to one module

Table 4. Covariance ratio (CR) and effect sizes (Z_{CR}) for the modularity hypotheses in the hominid wrist. All CR are statistically significant at p < 0.01. The Z_{CR} values are depicted in Fig. 2 and the pairwise differences in ZCR (\hat{Z}_{12}) are in Tables S3–S6. Hypotheses are described in Table 3.

Hypotheses	All		Human		Chimp	anzees	Gorillas	
	CR	Z _{CR}	CR	Z _{CR}	CR	Z _{CR}	CR	Z _{CR}
H1	0.64	-8.9	0.55	-8.6	0.57	-8.5	0.53	-8.3
H2	0.75	-7.4	0.56	-9.1	0.7	-6.9	0.63	-7.3
H3	0.81	-5.8	0.76	-5.4	0.76	-5.6	0.63	-7.2
H4	0.80	-5.9	0.74	-5.9	0.66	-7.4	0.64	-7.2
H5	0.75	-5.2	0.61	-6.3	0.67	-5.8	0.66	-5.3
H6	0.60	-8.1	0.62	-6.2	0.71	-5.2	0.76	-3.9
H7	0.77	-4.9	0.7	-5.1	0.71	-5.2	0.48	-7.7
H8	0.81	-4.5	0.66	-5.7	0.65	-6.1	0.56	-6.7
H9	0.81	-6.6	0.57	-7.7	0.6	-7.6	0.48	-8.6
H10	0.73	-7.9	0.59	-7.6	0.63	-7.3	0.61	-6.7
H11	0.65	-8.4	0.6	-7.1	0.61	-7.3	0.57	-7.3
H12	0.63	-6.2	0.59	-7.6	0.6	-7.4	0.56	-7.4
H13	0.74	-6.5	0.6	-7.1	0.61	-7.5	0.51	-8.1
H14	0.63	-8.4	0.58	-7.8	0.63	-7.2	0.66	-6
H15	0	0	0	0	0	0	0	0

identified, we also tested whether some genera displayed a greater degree of modularity than others. The CR, Z_{CR} , and \hat{Z}_{12} were also calculated using the *modularity.test()* and *compare.CR()* functions of the *geomorph* R package (Adams et al. 2019).

Nor the RV coefficient (Klingenberg 2009) nor the Partial-Least squared, commonly used for testing modular hypotheses (e.g., Mitteroecker & Bookstein 2008; Klingenberg 2009; Klingenberg & Marugán-Lobón 2013) were used here because they are adversely affected the sample size and the number of variables. Instead, the CR and \hat{Z} 12 coefficients were used as they are not affected by these attributes of the data (Adams 2016).

All the data used in this study are available in Supplementary Material 1 (Table S1). These data comprise the landmark coordinates after Procrustes superimposition.

Results

Allometry

Regression analyses of Procrustes coordinates on centroid size produced non-significant results in all cases (p > 0.05). Therefore, we excluded size as a factor contributing to variation in shape among the taxa studied here, and the following analyses were carried out using Procrustes coordinates and not 'size-corrected' variables (i.e., the residuals from the regressions of shape on centroid size).

Optimal modular hypotheses for hominines.

The CRs of all hypotheses were significantly less than 1 (Table 4), indicating that regardless of how the bones are combined to create the alternative modular hypotheses there is a strong modular signal in the sample. When comparing all hypotheses, H1 for the whole sample exhibited the largest negative Z_{CR} (Fig. 2, Table 4) which was significantly different (p < 0.05) from all the remaining hypotheses (Fig. 2, Table S6). H1 was thus selected as the best modularity model for hominines, which implies that each carpal represented is its own modular unit. However, except in chimpanzees (Figs 2 and 3, Table 4), H1 was not the best modular model for each genus individually. In humans, H2 showed a larger negative Z_{CR} than H1 (Fig. 2, Table 4), although this difference was not significant ($\hat{Z}_{12} = 0.63$, p = 0.53) (Table S3). Model H2 implies that the capitate and lunate form a different module than that of the scaphoid and trapezium. In gorillas, H9 yielded a larger negative Z_{CR} than H1 (Fig. 2, Table 4), yet this difference was not statistically significant either ($\hat{Z}_{12} = 0.43$, p = 0.67) (Table S5). H9 groups the capitate and lunate in the same module, while the scaphoid and trapezium each belong to their own modules. Fig. 3 depicts the optimal modular hypothesis for each genus.

To further explore the previous finding indicating possible variation in the modularity structure across taxa (Fig. 2), a pairwise modularity score (\hat{Z}_{12}) was calculated for every pair of carpals within each genus (Fig. 4). In humans, the



Fig. 2. Effect sizes (Z_{CR}) for the covariance ratio (CR) for the 15 modular hypotheses for all samples, and for each genus separately. Hypotheses are described in Table 3. The exact Z_{CR} values are in Table 4 and the pairwise differences in ZCR (\hat{Z}_{12}) are in Tables S3–S6.



Fig. 3. Illustration of the dorsal view of a left wrist showing the optimal modular hypothesis for humans (H2), chimpanzees (H1) and gorillas (H9). They were selected as they have the largest negative Z_{CR} values (Table 4). Hypotheses are described in Table 3.

modular signals between capitate and lunate, and between trapezium and scaphoid, was significantly lower (p < 0.05) than those of the remaining pairs of carpals (capitate and trapezium, and lunate and trapezium). This might suggest that the lunate and capitate have a degree of morphological integration, as do the trapezium and scaphoid. Additionally, the modular signals between capitate and lunate in one module, and trapezium and scaphoid in another, were statistically similar ($\hat{Z}_{12} = 0.26$, p = 0.28) (Fig. 4). These findings are consistent with H2 being the model with the best fit for humans (Figs 2 and 3). In chimpanzees, no pair of carpals exhibits a greater Z_{CR} than any other, which is also expected given that H1 is the optimal modular hypothesis for this genus. As for gorillas, the capitate and trapezium show a significantly higher modular signal than the lunate and scaphoid $(\hat{Z}_{12} = 2.14, p = 0.03)$, which is consistent with the capitate belonging to a different module than the trapezium, as indicated by the hypothesis with the most negative Z_{CR} value (H9). Similarly, the only other significantly different modular signal in gorillas was between the capitate and trapezium, which is higher than that found for the capitate and lunate $(\hat{Z}_{12} = 1.90, p = 0.05)$. Both results for gorillas are consistent with H9 being the best model for this genus. However, these results for gorillas do not exclude other hypotheses from being the best modular hypothesis (H1, H8, H10, H12, and H13, Table S5).



Fig. 4. Effect sizes (Z_{CR}) for the optimal modular hypothesis for the wrist in hominines (H1), and for each genus separately.

Discussion

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In this study we aimed to describe the modular pattern in the wrist of hominines and determine whether the pattern and strength of covariation across carpals is shared in humans, chimpanzees, and gorillas. To do this, we used the covariance ratio (CR) (Adams 2016; Adams & Collyer 2019) to test the degree to which changes in the capitate, lunate, scaphoid, and trapezium are associated with changes in each of the other bones. Our results indicate that the best fit for the covariation patterns in the wrist of hominines is the hypothesis that indicates that each carpal is its own modular unit (H1), as the level of covariation between carpals was always smaller than the covariation within carpals (CR in Table 4). This results supports previous evidence demonstrating great variability in the shape of carpals across primates (Lewis 1972; Corruccini 1978; Kivell et al. 2013). It also indicates that although the hands of humans have become less integrated with the feet in comparison to species with functionally similar use of both structures (Rolian 2009), it may not mean that the strength of reciprocal relationships across carpals is lower than in apes (H1 in Fig. 2). On the other hand, this result is intriguing given midcarpal joints are tightly supported by and array of ligaments (e.g. scaphotrapeziotrapezoid, scaphocapitate, dorsal intercarpal ligament that units the lunate scaphoid and trapezium), the flexor retinaculum and the limited range of movements of plane joint such as most of the intercarpacal joints are, and thus a coordinate covariation pattern would be intuitively expected in all hominines. Orr (2017) has also found that mobility in extension in chimpanzees scaphocapitate joint and in a lesser extent lunatecapitate joint, are relatively restricted when compared with other primates, yet this species showed the higher level of autonomy among the taxa studied here. Our results thus do not seem to indicate that a

low range of motion of carpal joints would necessarily result in a lower degree of integration.

The high level of autonomy of these four carpals indicated by our results requires some caution. First, the generalized Procrustes superimposition procedure, in which each bone was subject to a separate GPA, reduces the possible inflation of the covariance pattern between bones, as compared to the approach that uses one common superimposition and then splits the dataset to assess modularity hypotheses (Cardini 2019). However, the applied approach (i.e., separate superimpositions) may overestimate modularity, as it discards information related to the relative size and position of the modules (Cardini 2019). Second, it is also probable that the different covariation structure in the wrist found in some of our analyses for humans, chimpanzees, and gorillas (Figs 2, 3, Table 4), favors the simplest of all available hypotheses (H1), particularly when the entire sample is pooled (in terms that suggest no covariation between any of the carpals). In relation to the latter, although H1 was selected as the best model explaining the covariation structure of hominines, the different behavior of the genera when analyzed separately (Fig. 2) and the Z_{CR} comparison between carpal pairs indicate otherwise: that the level of association between some of them vary across taxa. This is true for the levels of covariation between the capitate and lunate, and the trapezium and scaphoid, which are higher for humans when compared to other pairs of carpals (Fig. 4), while for chimpanzees carpal pairs do not present different strengths of covariation. This makes H1 the optimal modular hypothesis for chimpanzees (in which each carpal corresponds to its own modular unit), while in the case of humans H2 is a better fit (i.e., the capitate and lunate belong to the same modular unit, and the trapezium to another) (Fig. 2 and 3). Gorillas share with humans that the capitate and lunate exhibit a degree of covariation

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and that the capitate and trapezium belong to different modules (as indicated by H9). However, results were less conclusive for this genus than for the others, as H9 presented the lowest Z_{CR} yet these results could not be confirmed when a pairwise modularity score (\hat{Z}_{12}) was calculated for every pair of carpals (Table S5).

As expected, bones that articulate present higher degrees of covariation (Fig. 1). However, our results also indicate that shape covariation is not just driven by the mere connection of carpals, as for instance trapezium and scaphoid, and capitate and lunate articulate in all hominines, but they do not covary with the same strength (Fig. 1). This raises the possibility that the covariance structure we observe here might be influenced by other factors, such as functional demands (see e.g., Püschel et al. 2020). According to our analysis, what separates humans from African apes is a stronger degree of covariation between the trapezium and the scaphoid. It is interesting that the radial side of the wrist separates these two groups, as a large proportion of studies dealing with manual differences between apes and humans have focused on the thumb, including the trapeziometacarpal joint, and point to enhanced manipulative capabilities in the former (Hamrick et al. 1998; Marzke et al. 1999; Marzke et al. 2010; Tocheri et al. 2008; Feix et al. 2015; Key & Dunmore 2015; Bardo et al. 2020; Püschel et al. 2022). Also, the radio-carpal joint (which involves the scaphoid) has been related to mechanical advantages in accuracy and force generation for the use of tools in humans (Williams et al. 2010; Williams et al. 2014). Even though functional interpretations are beyond the scope of our aims, the higher covariance we found between the scaphoid and the trapezium is consistent with Bardo's et al. (2020) findings regarding the covariance in all facets of the trapezium and the first metacarpal. Most of the facets of these bones are obliquely oriented relative to the transverse plane, which would be advantageous for the habitual use of forceful precision grips. Further analyses should estimate whether other associated changes of the carpals can be informative with respect to functional requirements in humans as compared to the African apes (Tocheri et al. 2005; Tocheri et al. 2008; Marzke et al. 2010; Feix et al. 2015). This would require a more detailed landmark configuration and a different statistical approach than the one presented here, as the CR cannot be used to describe specific shape changes, as principal component analysis and/or partial least squares analysis would do (although see Cardini 2019).

The presence of different modular strengths in the wrist bones of gorillas and chimpanzees (higher modular strength in the latter) is also noteworthy, as the presence of a knucklewalking complex, common to chimpanzees and gorillas, has long been discussed (Corruccini 1978; Begun 1992; Richmond & Strait 2000; Kivell & Schmitt 2009; Williams 2010; Püschel et al. 2020). For instance, Richmond & Strait (2000) proposed that African apes have a unique suite of skeletal traits involving the radiocarpal joint, which is adapted to stabilize the wrist during knuckle-walking, yet others argue that this type of locomotion is not the same biomechanical phenomenon in chimpanzees and gorillas (Inouye 1994; Kivell & Schmitt 2009). Our analysis does not indicate that there is a common covariation pattern for chimpanzees and gorillas, different from that of humans, that could allow us to define a potential knuckle-walking complex. This is consistent with Williams' (2010) conclusion that there is not a unique pattern of integration between the capitate and third metacarpal that distinguishes knuckle-walkers from nonknuckle-walking taxa.

9

Conclusions

Hominines have in common that each carpal covaries mainly with itself (scaphoid, lunate, trapezium and capitate) and with other carpals to a lesser extent. However, there are differences in the covariation strength that they exhibit with other wrist bones. In humans, the trapezium and scaphoid present a significantly lower modular signal with one another than with the remaining bones, and this also occurs with the capitate and lunate. This suggests that there may be associated shape changes between the scaphoid and trapezium, and between the capitate and lunate in humans. In gorillas there are also significant differences in the covariation structure across carpals, which indicates that the capitate and trapezium vary more independently than other pairs of carpals, and that the capitate and lunate covary as they do in humans. Of the three genera, chimpanzees presented the lowest interaction among carpals.

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Fig. S1. Sampling curve from performing LaSEC on the capitate dataset. Each grey line indicates fit values from one iteration of sub-sampling. The thick red line corresponds to the median fit value at each number of landmarks. The dashed line and white dot correspond to the median fit value obtained using 5 landmarks. The presence of a plateau indicates convergence, and hence oversampling if more landmarks are included.

Table S2. Effect sizes for the covariance ratio (Z_{CR}) for the modular hypotheses in the wrist of humans, chimpanzees and gorillas. All CR values were significant at p < 0.01.

Table S3. Matrix of pairwise differences in effect sizes (Z_{12}) (lower left triangle) between the modular hypotheses in humans and their associated *p*-values (upper right triangle). Significant values are in bold.

Table S4. Matrix of pairwise differences in effect sizes (Z_{12}) (lower left triangle) between the modular hypotheses in chimpanzees and their associated *p*-values (upper right triangle). Significant values are in bold.

Table S5. Matrix of pairwise differences in effect sizes (Z_{12}) (lower left triangle) between the modular hypotheses in gorillas and their associated *p*-values (upper right triangle). Significant values are in bold.

Table S6. Matrix of pairwise differences in effect sizes (Z_{12}) (lower left triangle) between the modular hypotheses in hominids (genera pooled) and their associated *p*-values (upper right triangle). Significant values are in bold.