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1 Morphometric human embryonic brain features according to developmental 2 stage 3 ¹Ami Kobayashi, ¹Koichi Ishizu, ^{1,2}Shigehito Yamada, ²Chigako Uwabe, 5 ³Katsumi Kose, ¹Tetsuya Takakuwa $6\overline{6}$ 1) Human Health Science, Graduate School of Medicine, Kyoto University, 8 Kyoto, Japan 2) Congenital Anomaly Research Center, Graduate School of Medicine, Kyoto 10 University, Kyoto, Japan 3) Institute of Applied Physics, University of Tsukuba, Ibaragi, Japan 12 **Running title**: Human embryonic brain measurements **Word count**: 2237 words **Number of Tables**; 4, **Figures**; 5 **Corresponding author**: Dr. Tetsuya Takakuwa, Human Health Science,

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Abstract

42 Objectives: The present study investigated linear, area, and volume measurements of human brain samples according to Carnegie stages (CS) in an attempt to select suitable morphometric features that reflect embryonic 45 development. Methods: Using magnetic resonance (MR) imaging, we measured seven linear segments, three separate areas, and three regional volumes in 101 samples 48 between CS 13 and CS 23. Brain volume was determined via manual segmentation of the MR image, whereby a formula was generated to estimate 50 the volume of each linear measurement. Results: All parameters correlated with crown-rump length. Bitemporal length (BT) and mesencephalic height increased linearly according to the CS, and a high correlation between BT and both whole-brain (r = 0.98) and 54 prosencephalon volume ($r = 0.99$) was found when brain cavity volume was 55 excluded.

Conclusion: Morphometric data related to human embryonic stages are valuable

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*T***2** Introduction

102 few morphometric studies of the human embryonic brain have employed the 103 $CS^{14,15}$.

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Materials and methods

Embryonic specimens

 Approximately 44000 human embryos (comprising the Kyoto 117 Collection) are stored at the Congenital Anomaly Research Center at Kyoto

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Morphometric analysis

133 MR imaging datasets for each embryo were initially obtained as 256 \times

134 256×512 voxel data¹⁹. The midsagittal and transverse planes were used 135 according to previous studies, with minor modifications^{14,15} (Figure 1). The planes for length and area measurements were digitally resectioned using OsiriX™ software (ver. 4.0, Pixmeo SARL, Geneva, Switzerland).

138 The following seven segments were measured, which were consistent 139 with a previous study¹⁴: fronto-occipital diameters (FOa and FOb) and 140 bitemporal diameter (BT) as substitutions for the prosencephalon, 141 mesencephalon length (ML) and mesencephalon height (MH) as substitutions 142 for the mesencephalon, and cerebellar length (CL) and cerebellar height (CH) as 143 substitutions for the rhombencephalon and cerebellum, respectively (Figure 1A). Area measurements were obtained for three regions in the midsagittal section 145 that corresponded to the prosencephalon, mesencephalon, and 146 rhombencephalon¹⁵.

 Brain volume was measured by manual segmentation, as described 148 previously¹⁰. Briefly, brains and ventricles were segmented for 3D reconstruction 149 using the FSL view of the FMRIB Software LibraryTM (ver. 4.1.9, Analysis Group,

150 FMRIB, Oxford, UK). Three-dimensional brain morphology was computationally 151 reconstructed with AmiraTM software (ver. 5.4.0, Visage Imaging, Berlin, 152 Germany).

153 Brain and whole embryo volumes were calculated using OsiriX[™] software (ver. 4.0, Pixmeo SARL, Geneva, Switzerland). Vesicles were divided into three regions according to the following anatomical landmarks: the 156 supramammillary recess and posterior commissure were used to define the 157 prosencephalon and mesencephalon; the isthmic recess and the isthmic groove 158 were used to define the mesencephalon and rhombencephalon; and the C1 159 vertebral level was used to define the separation between the rhombencephalon 160 and spinal cord¹⁰. After dimensional matching, a formula was derived to estimate 161 brain volume from the linear measurements. We then analyzed both total brain 162 volume (with the ventricles) and brain volume without the ventricles, since an increase in ventricular volume – a feature of the embryonic period – could have 164 influenced our measurements^{5,6,10,15}. The Ethics Committee of the Kyoto University Graduate School and Faculty of Medicine (E986) approved this study.

167 **Results**

168 *Linear, area, and volume measurements*

169 **1. Linear measurements**

176 The seven length segments were plotted by CS group (Figure 2). BT and 177 MH exhibited a nearly linear increase, while other segments increased 178 non-monotonically during CS13 and CS23. A plateau phase between CS19 and CS20 was observed in several segments (FOa, FOb, ML, and CL). It should be 180 noted that we were unable to compare our data with that of a previous study 181 because the segment and CS correlations in that study were not precisely

166

 182 analyzed¹⁴.

183

184 **2. Area measurements**

185 Areas in the midsagittal section that corresponded to the whole brain, as 186 well as the prosencephalon, mesencephalon, and rhombencephalon, were 187 measured (Figure 3). Each area increased between CS13 and CS23. 188 Exponential curves demonstrated a good fit to the data $(R^2 = 0.92 - 0.97)$, except 188 189 for the rhombencephalon area (R^2 = 0.85); this finding was consistent with that of 190 a previous study (Table $2)^{15}$.

191

3. Volume measurements

 When brain volume was measured without the ventricles (i.e., brain tissue only), we determined that it increased exponentially until CS23. On the other hand, when ventricular volume was included in the measurement, we 196 determined that brain volume increased and then plateaued between CS19 and CS20 (Table 3).

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Predicting brain volume from linear data and measurements

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228 **Discussion**

 The human embryonic brain develops in a complicated manner over a 230 $\,$ short period¹⁰, with growth speed and increases in measurements varying according to developmental stage. Thus, the determination of developmental 232 stage is necessary for a more precise detection of embryonic abnormalities. 233 Improvements in sonographic resolution may contribute to our ability to observe detailed and precise embryonic morphology. Nevertheless, it remains unclear how measurements, which reflect developmental features, relate to developmental stages. Classic embryology using histological techniques has 237 revealed morphological features according to developmental stage, and this 238 forms the current basis of sonoembryology. However, classic embryology has 239 provided relatively little morphometric data^{14.15}. The present study improved 240 upon previous methods by providing a suitable plane for a morphometric 241 analysis from staged human embryos using MR microscopic data.

242 **In the current study, the increases in brain cavity and whole body** 243 volume (in relation to the CRL) were comparable to those reported by Blaas' 244 sonoembryologic study²⁰ (Figure 4). Regarding the prosencephalon, the volume

261 maximum observed c. CS19−20). Rousian et al.⁸ measured the relationship of numerous parameters with CS using 3D sonography and found that brain cavity 263 volume increased with the quadratic function of the CS. The mean brain cavity 264 volume in our study was comparable to that reported by Rousian et al. from CS13 to 17, but not with the quadratic function they used after CS 18 (Figure 5). Analyzing why such discrepancies arise may provide clues for further understanding stage-specific morphological features during development. 268 Linear measurements, which can also be used to estimate volume, may be preferable to other measurements because they are related to both brain 270 development and growth. Thus, BT is a good marker for estimating human 271 embryonic brain development and growth, as has been mentioned in previous 272 studies^{1,4,7,13, 21}. The present study included embryos at earlier stages (between 273 CS13 and CS15) in which the telencephalon was not prominent; thus, our results demonstrated that linear measurement values, such as BT and volume, 275 are useful between CS13 and CS15.

276 **It is difficult to compare our linear measurements with most previous**

277 sonographic measurements^{3,5,6,7}, as the measurements used in the present 278 study were selected from classical embryonic studies^{14,15}. Thus, with the 279 exception of MH and ML, most of the linear measurement definitions we 280 employed differed from those of recent studies^{5,9}. Using sonography, Tanaka et 280 281 \cdot al.⁹ presented length measurements (including ML and MH) at every day of 282 gestation between days 49 and 69. The length of ML in our study was shorter 283 than that of the Tanaka at all stages between CS 15 and 23. The MH length in 284 our study was also shorter than that of Tanaka et al. between CS 15 and 20, but 285 larger after CS 21 (data not shown). Note that the comparison may not be 286 entirely reliable as we compared the ML and MH data using the predicted CS 287 from the day of gestation¹³. Selection and reevaluation of linear measurements 288 that allow comparisons of data among sonoembryonic studies will be necessary 289 for future studies. Further, the relationship between CS and embryo age is a 290 critical issue to consider. Embryological studies demonstrate that considerable 291 variability is observed in the size and developmental stage among human 292 embryos at a given gestational age²². Thus, the embryonic age that is used in

 clinics cannot be used to determine the developmental stage (CS), as prenatal development may not proceed at the same speed in every embryo. The present study provided a morphometric analysis from staged human embryos using MR microscopic data. Morphometric data, according to human 297 embryonic stages, are valuable for correcting and comparing sonographic data. 298 The present approach may contribute to improvements in prenatal diagnostics 299 by enabling the selection of more suitable measurements during earlier 300 embryonic stages. 301 302 References 303 1 Blaas HG. Detection of structural abnormalities in the first trimester using 304 ultrasound. Best Pract Res Clin Obstet Gynaecol 2014;28:341–53. 305 2 Pooh RK, Shiota K, Kurjak A. Imaging of the human embryo with magnetic 306 resonance imaging microscopy and high-resolution transvaginal 3-dimensional 307 sonography: human embryology in the $21st$ century. Am J Obste Gynecol 308 2011;204:77.e1-16.

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360 Figure legends

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Figure 1. MR images of midsagittal and transverse sections

 drawing a perpendicular line connecting the dorsal and ventral primary fissures,

- 378 Pr: prosencephalon, Me: mesencephalon, Rh: rhombencephalon, Md: sum of Pr,
- 379 Me, and Rh.
- 380 Measurements were obtained from two previous studies (Desmond and
- 381 O'Rahilly, 1981; Levitan and Desmond, 2009).
- 382 **(B)** Human brain at CS16.
- 383 **(C)** Human brain at CS13.
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- 385 Figure 2. Embryonic brain length measurements according to Carnegie stages.

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- 387 Figure 3. Area measurements between CS13 and CS23.
- 388 The rhombencephalon in our data is compared with "whole cerebellum" defined 389 in the previous study¹⁵.

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- Figure 4. Comparison of the present volume measurements with Blaas' study.
- (A) Increase in brain cavity and whole body volume in relation to the CRL.

 (B) Prosencephalon, mesencephalon, and rhombencephalon volumes in relation 394 to the CRL.

- Blue: the present study, red: data from Blaas' study (pp 76 in Ref. 20). The prosencephalon volume was calculated as the sum of the hemispheres and the
- 397 diencephalon.
- 398

Figure 5. Brain cavity volume in relation to the CS.

400 The present volume measurements were compared to those reported by 401 Rousian et al.

402 The mean volume of our data overlapped, from CS13 to 17, with the quadratic

403 function employed by Rousian et al⁸.

404

405 Supplementary Figure 1. Correlation between the seven length segments 406 measured and the CRL. The data imply that brain size measurements are useful 407 indicators of linear embryonic neural development.

CRL; Crown-Rump length, r; correlation coefficient

CS = Carnegie stages; *; whole cerebellum in Levitan & Desmond (2009)

		Whole brain					Prosencephalon					Mesencephalon				Rhombencephalon					
Carnegie		mean volume (mm ³)				mean volume $(mm3)$					mean volume (mm ³)				mean volume (mm ³)						
stage	n n	total	(SD)	tissue	cavity	ratio	total	(SD)	tissue	cavity	ratio	total	(SD)	tissue	cavity	ratio	total	(SD)	tissue	cavity	ratio
13	5	2.1	0.7	1.2	0.9	0.82	0.5	0.2	0.3	0.2	0.77	0.3	0.1	0.2	0.1	0.40	1.3	0.4	0.7	0.7	0.96
14	9	5.7	$\mathsf{.4}$	2.6	3.1	.16	1.2	0.4	0.7	0.6	0.88	0.7	0.1	0.4	0.3	0.60	3.8	0.9	1.6	2.3	1.43
15	9	11.4	.6	5.5	6.0	.10	2.6	0.4	1.5	1.2	0.79	1.3	0.2	0.8	0.5	0.67	7.5	1.2	3.2	4.3	1.34
16	10	16.3	2.6	7.4	8.9	.20	4.1	1.1	2.1	2.0	0.96	2.0	0.4	1.2	0.9	0.74	10.1	1.3	4.2	6.0	1.45
17	10 [°]	29.9	6.2	12.2	17.8	.46	9.0	2.5	4.3	4.7	1.11	3.9	1.0	1.9	2.0	1.04	17.1	3.0	6.0	11.1	1.84
18	9	57.7	7.5	27.6	30.1	.09	19.3	2.7	9.8	9.6	0.98	8.1	0.9	4.1	4.0	0.98	30.3	4.8	13.7	16.6	1.21
19	10	79.0	8.9	43.9	35.1	0.80	29.9	4.6	17.1	12.8	0.75	11.6	1.2	6.6	4.9	0.74	37.5	4.9	20.1	17.4	0.87
20	10	.9 81	19.0	52.8	29.1	0.55	36.8	8.2	23.0	13.8	0.60	11.3	2.6	6.8	4.5	0.66	33.8	9.0	22.9	10.8	0.47
21	10 [°]	145.1	16.6	95.3	49.8	0.52	74.0	14.5	47.5	26.5	0.56	19.0	2.8	12.5	6.5	0.52	52.1	6.2	35.3	16.9	0.48
22	9.	206.4	39.0	135.2	71.1	0.53	115.5	29.0	70.5	45.0	0.64	25.7	4.1	17.9	7.8	0.44	65.2	10.2	46.8	18.3	0.39
23	10	286.3	61.7	189.1	97.2	0.51	184.5	50.5	111.0	73.5	0.66	28.9	4.0	21.9	7.0	0.32	73.1	10.0	56.3	16.7	0.30

Table 3. Brain volume of human embryos measured by manual segmentation

Formula										
V (mm ³) = a $[X][Y][Z]+b$	a	[X][Y][Z]	+b	r						
Volume with ventricles										
Whole brain	0.52	$[BT]^3$	26.2	0.97						
Prosencephalon	0.92	[FOa] ³	-9.13	0.86						
	0.49	$[FOb]$ ³	-8.21	0.93						
	0.34	$[BT]^3$	4.05	0.98						
	0.59	[FOa][FOb][BT]	-7.50	0.97						
Mesencephalon	0.70	$[MH]$ ³	3.27	0.94						
	0.28	$[ML]$ ³	0.90	0.95						
Rhombencephalon	18.1	[CH] ³	19.8	0.80						
	11.2	$[CL]^3$	9.87	0.92						
Volume without ventricles										
Whole brain	0.35	$[BT]^3$	12.7	0.98						
Prosencephalon	0.57	[FOa] ³	-6.39	0.88						
	0.30	$[FOb]$ ³	5.18	0.94						
	0.21	$[BT]^3$	2.11	0.99						
	0.36	[FOa][FOb][BT]	-5.03	0.97						
Mesencephalon	0.53	$[MH]$ ³	1.42	0.96						
	0.21	$[ML]$ ³	-0.21	0.95						
Rhombencephalon	15.1	[CH] ³	10.3	0.85						
	8.99	$[CL]^3$	2.68	0.94						

Table 4. Calculation of brain volume from length measurements

Prosencephalon

