

# STUDY INFORMATION AND DATA USE AGREEMENT FOR THE RESEARCH PROJECT: "CROSS CULTURAL COMPARISON OF HUMAN GROWTH TRAJECTORIES"

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## 1. Brief Description

In this project, we will apply the growth model recently developed by Bunce, Fernández, and Revilla-Minaya 2022 (see: <u>Causal models of human growth and their estimation using temporally-sparse data</u>) to longitudinal datasets from populations inhabiting a wide range of ecoregions with the aim of distinguishing and describing general patterns and global variation in human growth trajectories. By using this model, we expect to be able to estimate, and hopefully disentangle, the relative contributions of different factors affecting growth in height and weight around the world, as well as to compare specific components of growth among diverse populations.

## 2. Intended use of the Data

The data provided by the collaborators in this project will be used for the sole purpose of this study on the comparison of growth trajectories. The final product will be one or more scientific articles on this topic. Data will not be used for any commercial purposes. We will not distribute these data to third parties under any circumstance. We will make no attempt to identify or contact research participants, their households or communities. Collaborators will have the option of making the de-identified data they contribute publicly available as an accompaniment to the published article(s) that result from this project (see below).

# 3. Dataset and population (s)

Please provide the following characteristics of the dataset (s), indicating population, location (s), a brief description of data collection methods and/or a reference for them, and indicate whether data was collected after IRB approval or other ethics review (please, indicate review board and protocol number). If other ethical protocols were followed, please specify.

• Population name(s) and brief description (e.g., subsistence practices, refugees from another region, access to Western healthcare systems, common health challenges, etc.):

These data include data from a village in the northern Republic of the Congo where people from two ethnic groups reside. The BaYaka are culturally and economically forest-oriented and procure most of their food from the forest or via trading gathered forest foods, manual labor, or services for horticultural produce or market-goods with the Bandongo. The Bandongo are Bantu-speakers whose subsistence economy largely involves fishing, horticultural production, and fishing. Primary crops include manioc, plantains, and corn, while several minor crops are also grown. These two communities have dense economic and social ties but remain residentially segregated and do not inter-marry. They generally face the same health challenges, including high mortality due to infectious disease. There is no regular stock of Western medicines in the village. A pharmacy and clinic are available by boat around three hours from the village.

• Location (s) (please be as specific as possible, and include GPS coordinates if available):

The village of Djoube is located in the Congo Basin tropical forest and is situated along the banks of the Motaba river in the District of Dongou, Likouala Department, the Republic of the Congo, at the GPS coordinates 2.437314, 17.46918.

• Data collection methods (please include reference if available):

Data collection is described in Boyette et al. (2018, 2020). In brief, Bandongo reported their birth years, which are recorded in village registries. BaYaka do not tend to know the years they are born, so approximate birth year was determined by ranking all individuals in the community relative to each other and then following the estimation procedure described in Diekmann et al. (2017). Height was measured with a Seca stadiometer and weight with a standing scale. Bandongo data was collected in June-July, 2016 and BaYaka data in June-July, 2017.

• IRB/ethical review board approval (yes/no, explain):

Permission to conduct research in the Republic of the Congo was granted by the Centre de Recherche et des Etudes en Sciences Sociales et Humaines (currently the Institute National de Recherche en Sciences Sociales et Humaines), and the study was approved of by the Institutional Review Boards of Duke University (Protocol # 2017-0038) and the University of Notre Dame (# 18-02-4397).

# 4. Full name, affiliation and e-mail of all contributors/collaborators for this dataset (s):

Adam Howell Boyette Senior Researcher Department of Human Behavior, Ecology, and Culture Max Planck Institute for Evolutionary Anthropology adam boyette@eva.mpg.de

Lee T. Gettler
Rev. John A. O'Brien Associate Professor
Department Chairperson
Department of Anthropology
Director, Hormones, Health, and Human Behavior Lab
Faculty affiliate, Eck Institute for Global Health
University of Notre Dame
Lee.T.Gettler.1@nd.edu

Sheina Lew-Levy Assistant Professor Department of Psychology Durham University sheina.lew-levy@durham.ac.uk

#### 5. Data submission

Please send a <u>link to a folder</u> containing a file or several files (in case of more than one dataset) in **CSV format** to <u>catalina\_fernandez@eva.mpg.de</u>.

- Please include the **name of the population in the file name**, and an additional dataset identifier in case of multiple datasets.
- If the dataset also contains single (as opposed to longitudinal) measures for some individuals, please do not remove them from the file. We will also use those observations in the analysis.
- If you have data on adult individuals, whether or not they were also measured as children/adolescents, please include these observations as part of your dataset.

Please, do not include any direct identifying information in the dataset relating to the individuals, their relatives or household members, such as names, addresses, telephone numbers, e-mail addresses or social media identifiers, etc.

For each dataset, we request collaborators on this project to share a <u>deidentified</u> data file (.csv) containing the following information and columns:

**Column 1**: Individual identification code or number that is consistent across longitudinal observations of the same individual.

Column 2: Date of birth in dd/mm/yyyy format.

**Column 3**: Date of data collection event in **dd/mm/yyyy** format.

- \* If date of birth is not available, please provide **age** in days, whenever possible, for each data collection event.
- \* If known age is uncertain, if possible, please provide a **range** of minimum, estimated, and maximum age, based on your best approximations.

Column 4: Sex; coded as m or f.

Column 4: Height in cm Column 5: Weight in kg

\*Note that, for longitudinal measures, you will have multiple rows with the same individual identification code, date of birth, and sex, but with different dates of data collection, heights, and weights.

# 6. Long term archiving and data access

One of the objectives of this project is to make the deidentified datasets and code (s) used in the data cleaning and analyses available in an open-access repository. We encourage all collaborators to commit to share and deposit deidentified data in a public repository created for this project, once they have checked that, by doing so, they are not violating compliance with their ethical review protocols (e.g., IRB) or other agreements with research participants and community members/leaders. Making data open-access is not a requirement to participate in this study, but we believe that by granting public access to the code and data used we will allow other researchers to check our analyses and reproduce our findings.

If you agree that data from your research site can be made publicly available for this purpose, we will create an appropriate repository (e.g., on Github) to curate this dataset and indicate the contact information for the researchers responsible for each field site, in case someone wishes to use the dataset for a purpose other than simply checking the results of our analysis. If for any reason, in the future, you wish to change your decision regarding data access and usage, we can either include or remove the data from this repository.

#### Use of the data and access on a public repository (choose one option):

X Restricted use. I agree that deidentified data can be made publicly available for the spurpose of reproducing this specific analysis. Data cannot be used by the scientific communior the public for any other purpose.	
Unrestricted. I agree that deidentified data can be made publicly available without barriers to access or use.	
I do not consent to making this data available in a public repository. I am sharthis data only for the purpose of this study. I do not agree that this dataset(s) or any portion it/them can be shared publicly under any circumstances.	_

#### 7. Data sharing information to the communities involved

We believe that one of our responsibilities as scientists is to communicate research findings with the communities and individuals who provided their time and biometric information in order to make this study possible. We are aware that most, if not all, collaborators on this project have active field sites and engage regularly with the communities they work with for the purpose of informing and sharing research findings and other initiatives alike. We ask collaborators to this project to share the results of this study with the contributing communities, particularly the parts that concern the specific population that they work with. If a collaborator is no longer in contact with the study population, please let the project leaders know so that together we can potentially brainstorm an alternative solution. Ideally, presenting results to participants will occur prior to publication, so that participating communities have the opportunity to (re)express their permission for us to publish the results. Sharing the results may take the form of a live Power Point presentation, a video, a written information sheet, or any other format that the collaborator believes is most culturally appropriate to present this kind of information to the communities and participants who provided the data. If it is of interest as a model or guide, the project leaders (Catalina Fernández, Caissa Revilla-Minaya and John Bunce) can share with the contributors the materials that they will design for the Matsigenka population that they work with.

## 8. Manuscript authorship

The project leaders (Catalina Fernández, Caissa Revilla-Minaya and John Bunce) will draft the main manuscript and decide the order of authorship and co-authorship. We will ask all collaborators for feedback once analyses are completed and while we draft the discussion. We are currently considering organizing a workshop at MPI in Leipzig for project collaborators, after analyses are completed, but prior to publication, in order to share the results of the analyses, discuss the main findings and potential causal mechanisms contributing to variation in growth in different populations. This can also be an opportunity for planning further analyses and related projects for the future. More information about this potential meeting/workshop will be shared in the next months.

#### 9. Tentative timeline

- 30 June 2023: All datasets have been submitted to the project leaders.
- 30 September 2023: All datasets have been checked for errors and formatted for analysis.
- 30 March 2024: Data analysis is complete; manuscript is drafted by project leaders.
- Mid-May 2024: Potential workshop for contributors/collaborators at MPI Leipzig.
- 30 July 2024: Results have been presented to participating communities.
- 30 August 2024: Main manuscript is submitted for publication.

#### 10. Questions

<sup>\*</sup>Note that this is a very tentative timeline.

For questions regarding data submission, data archiving, and other aspects of this study, or additional information, please contact Catalina Fernández by e-mail at <a href="mailto:catalina\_fernandez@eva.mpg.de">catalina\_fernandez@eva.mpg.de</a>.

Name of the person in charge of the dataset:

Date: