## nature portfolio

Corresponding author(s):	Rebecca Rimbach
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## **Reporting Summary**

Nature Portfolio wishes to improve the reproducibility of the work that we publish. This form provides structure for consistency and transparency in reporting. For further information on Nature Portfolio policies, see our Editorial Policies and the Editorial Policy Checklist.

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FOL	an statistical analyses, confirm that the following items are present in the figure legend, table legend, main text, or Methods section.
n/a	Confirmed
	The exact sample size (n) for each experimental group/condition, given as a discrete number and unit of measurement
	🗶 A statement on whether measurements were taken from distinct samples or whether the same sample was measured repeatedly
	The statistical test(s) used AND whether they are one- or two-sided  Only common tests should be described solely by name; describe more complex techniques in the Methods section.
	🗶 A description of all covariates tested
	🗶 A description of any assumptions or corrections, such as tests of normality and adjustment for multiple comparisons
	A full description of the statistical parameters including central tendency (e.g. means) or other basic estimates (e.g. regression coefficient) AND variation (e.g. standard deviation) or associated estimates of uncertainty (e.g. confidence intervals)
	For null hypothesis testing, the test statistic (e.g. <i>F</i> , <i>t</i> , <i>r</i> ) with confidence intervals, effect sizes, degrees of freedom and <i>P</i> value noted <i>Give P values as exact values whenever suitable.</i>
	For Bayesian analysis, information on the choice of priors and Markov chain Monte Carlo settings
X	For hierarchical and complex designs, identification of the appropriate level for tests and full reporting of outcomes
×	Estimates of effect sizes (e.g. Cohen's <i>d</i> , Pearson's <i>r</i> ), indicating how they were calculated
	Our web collection on <u>statistics for biologists</u> contains articles on many of the points above.

## Software and code

Policy information about availability of computer code

Data collection

No software was used to collect data.

Data analysis

 $R \ (version 3.6.2), R \ package \ rptR \ (version 0.9.22), R \ package \ MCMCglmm \ (version 2.29), R \ package \ coda \ (version 0.19-3). We provide the source code used to perform analysis and output files through the OSF repository (https://osf.io/6q2kz/).$ 

For manuscripts utilizing custom algorithms or software that are central to the research but not yet described in published literature, software must be made available to editors and reviewers. We strongly encourage code deposition in a community repository (e.g. GitHub). See the Nature Portfolio guidelines for submitting code & software for further information.

## Data

Policy information about availability of data

All manuscripts must include a <u>data availability statement</u>. This statement should provide the following information, where applicable:

- Accession codes, unique identifiers, or web links for publicly available datasets
- A description of any restrictions on data availability
- For clinical datasets or third party data, please ensure that the statement adheres to our policy

All data supporting the analyses and results in this paper are available from the Doubly Labeled Water Database (https://doubly-labelled-water-database.iaea.org/home, https://www.dlwdatabase.org/) upon reasonable request. Because of subject confidentiality the database is not open access. However, access to components of the data is freely available to perform novel and approved analyses. Details of the application process are available at https://www.dlwdatabase.org. We provide a minimum dataset through the OSF repository to demonstrate the code (https://osf.io/6q2kz/).

Field-spe	ecific re	porting				
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Life scier	nces stu	dy design				
All studies must dis	close on these p	points even when the disclosure is negative.				
Sample size		Il individuals in the IAEA DLW database (https://doubly-labelled-water-database.iaea.org/ home) which had their TEE measured and that were at least 1 year old. This resulted in 696 TEE measurements of 348 adults and 114 TEE measurements of 47				
Data exclusions		TEE measurements from children younger than 1 year because the relationship between TEE and FFM in this age group appears in that for older children.				
Replication		nain purposes of this study is to assess repeatability of TEE measurements. TEE of adults was measured twice. For 27 children, two ements were available and for 20 children, three repeated measurements were available.				
Randomization	Not applicable, t	ble, there are not experimental groups.				
Blinding	Not applicable, there are not experimental groups.					
We require information	on from authors a	ecific materials, systems and methods bout some types of materials, experimental systems and methods used in many studies. Here, indicate whether each material, our study. If you are not sure if a list item applies to your research, read the appropriate section before selecting a response.				
•						
Materials & exp		n/a Involved in the study				
n/a   Involved in the study		ChiP-seq				
Eukaryotic	cell lines	Flow cytometry				
Palaeontology and archaeology  MRI-based neuroimaging						
Animals and other organisms						
	Clinical data   Dual use research of concern					
Dual use le	esearch of concern					
Human rese	arch partio	ipants				
Policy information a	about <u>studies in</u>	volving human research participants				
Population characte	eristics	This sutdy is a secondary analysis of a sub-population of DLW data compiled into the IAEA DLW database. The data pertaining				

database management group and the prior approval posted on the database website (www.dlwdatabase.org).

This sutdy is a secondary analysis of a sub-population of DLW data compiled into the IAEA DLW database and as such we did not recruit participants for this study.

All of the studies that provided data into the IAEA DLW database were locally ethically reviewed and approved. The present paper is based on a secondary analysis of these compiled data and such analyses do not require ethical permission.

Note that full information on the approval of the study protocol must also be provided in the manuscript.

Recruitment

Ethics oversight