



# A beginners guide to (growth trajectory analysis using) multilevel models

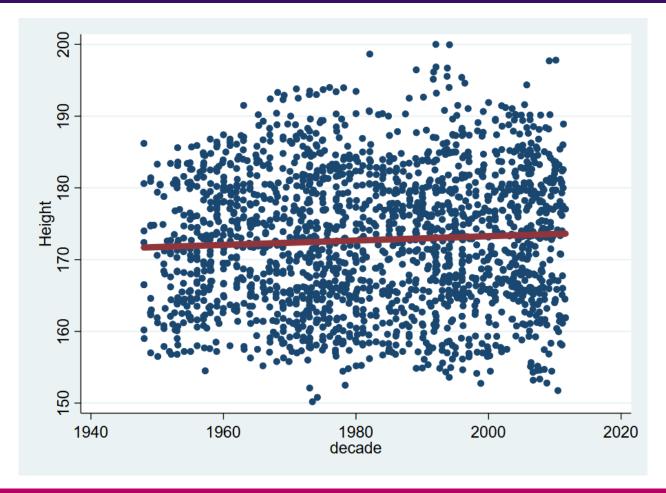
# Will Johnson

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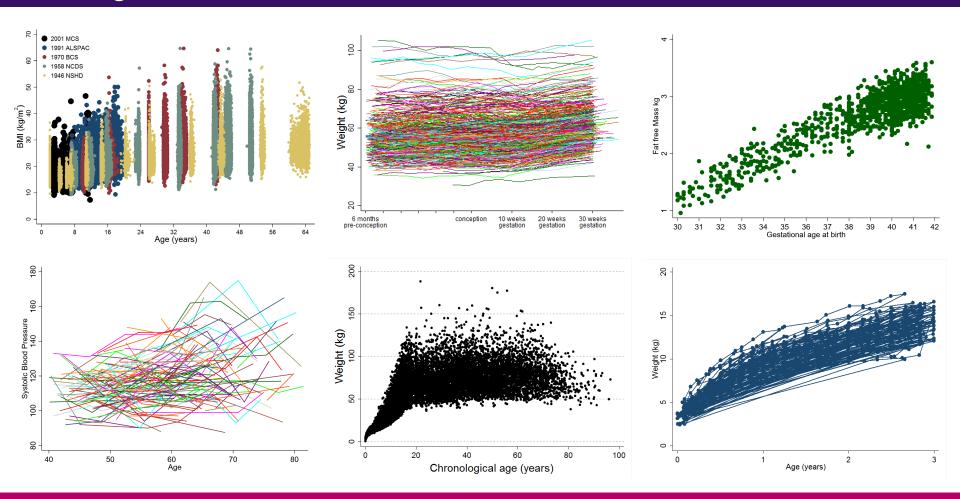


# What is a trajectory?



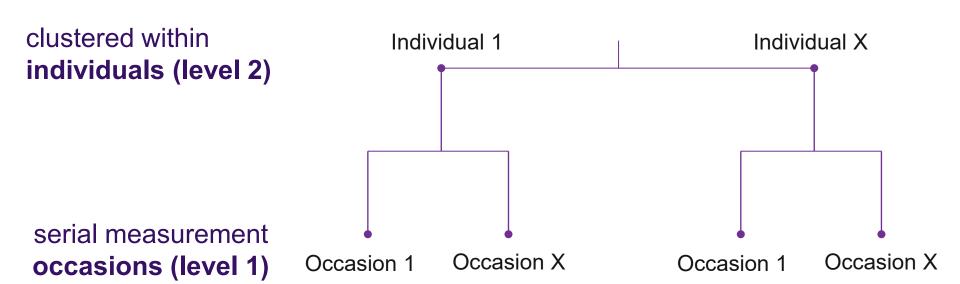


#### Longitudinal data





#### Structure of longitudinal data





#### Structure of longitudinal data

	Occasion			
Individual	1	2	3	4
1	<b>y</b> <sub>11</sub>	<b>y</b> <sub>21</sub>		<b>y</b> <sub>41</sub>
2	<b>y</b> <sub>12</sub>	<b>y</b> <sub>22</sub>	<b>y</b> <sub>32</sub>	<b>y</b> <sub>42</sub>
3	<b>y</b> <sub>13</sub>		<b>y</b> <sub>33</sub>	<b>y</b> <sub>43</sub>

 $y_{ij}$  is the response at occasion i for individual j

Individual	Occasion	
1	1	<b>y</b> <sub>11</sub>
1	2	<b>y</b> <sub>21</sub>
1	3	
1	4	<b>y</b> <sub>41</sub>
2	1	<b>y</b> <sub>12</sub>
2	2	Y <sub>22</sub>
2	2	Y <sub>32</sub>
2	2	Y <sub>42</sub>
3	1	Y <sub>13</sub>
3	2	
3	2	Y <sub>33</sub>
3	2	Y <sub>43</sub>



# Other examples of multilevel data





#### How many upper-level units?



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Elff, Martin; Heisig, Jan Paul; Schaeffer, Merlin; Shikano, Susumu

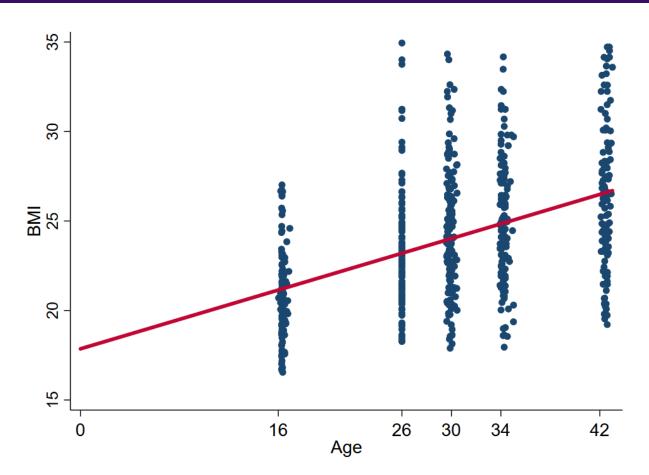
Article — Accepted Manuscript (Postprint)

Multilevel Analysis with Few Clusters: Improving
Likelihood-based Methods to Provide Unbiased
Estimates and Accurate Inference

British Journal of Political Science



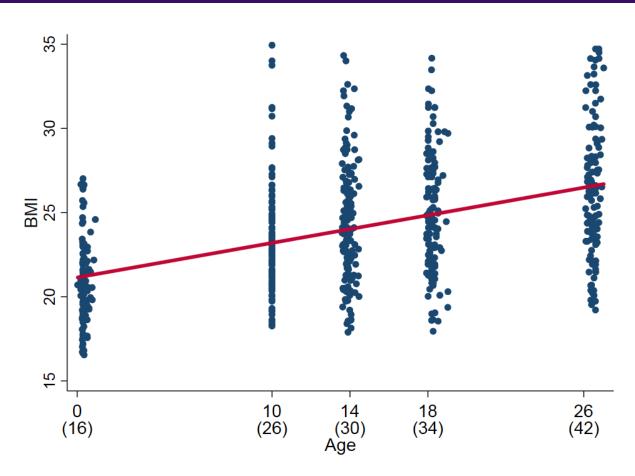
## Regression



bmi<sub>ij</sub> =  $\beta_{0i}$ cons + 0.205(0.016)age<sub>ij</sub>  $\beta_{0i}$  = 17.857(0.511) +  $e_{0ij}$ 



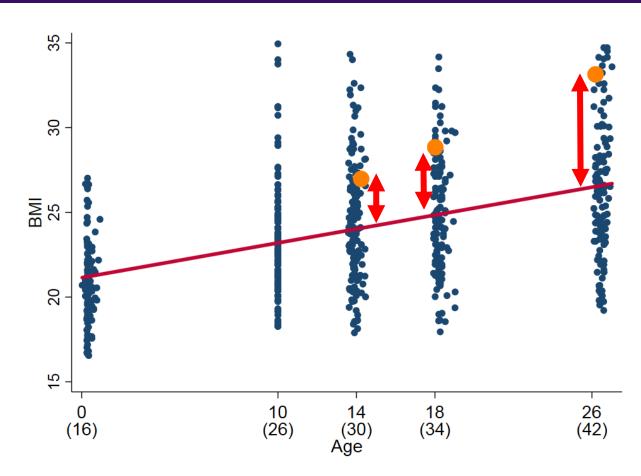
## Regression



bmi<sub>ij</sub> =  $\beta_{0i}$ cons + 0.205(0.016)agec<sub>ij</sub>  $\beta_{0i}$  = 21.144(0.267) +  $e_{0ij}$ 



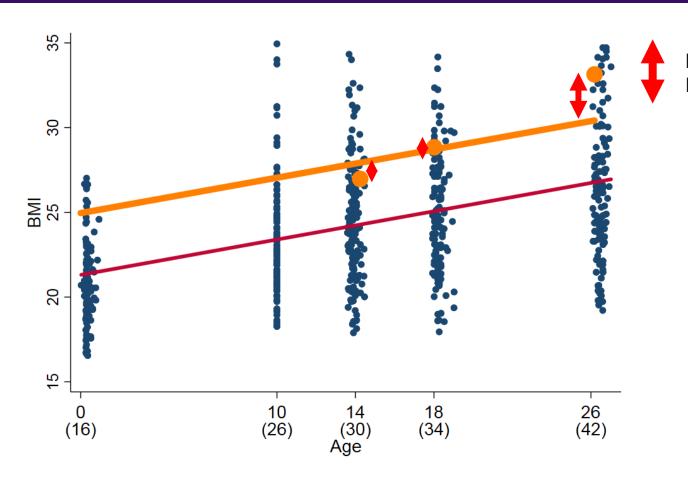
## Regression



bmi<sub>ij</sub> =  $\beta_{0i}$ cons + 0.205(0.016)agec<sub>ij</sub>  $\beta_{0i}$  = 21.144(0.267) +  $e_{0ij}$ 

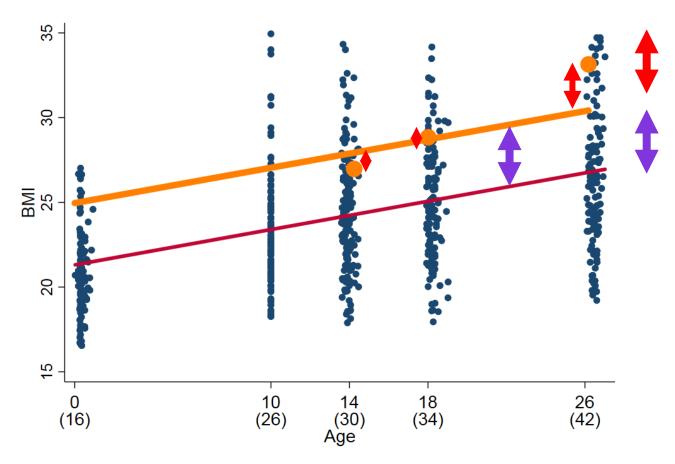
 $\left[ e_{0ij} \right] \sim N(0, \ \Omega_e) \ : \ \Omega_e = \left[ 11.674(0.661) \right]$ 





Level 1 or within-individual Residuals/ variance

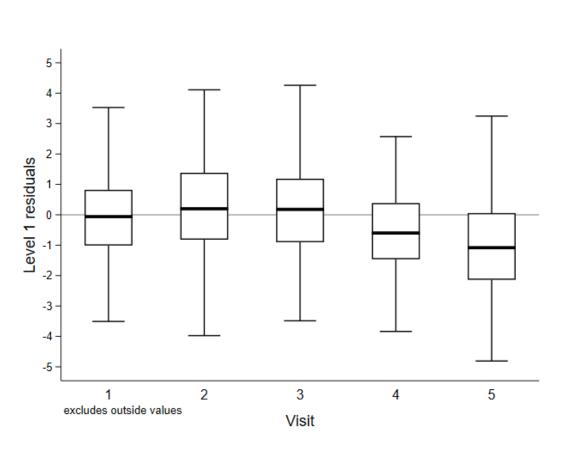




Level 1 or within-individual Residuals/ variance

Level 2 or between-individual Residuals/ variance





$$bmi_{ij} = \beta_{0ij}cons + \beta_1 agec_{ij}$$
$$\beta_{0ij} = \beta_0 + u_{0j} + e_{0ij}$$





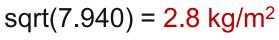
$$\begin{bmatrix} e_{0ij} \end{bmatrix} \sim N(0, \Omega_e) : \Omega_e = \begin{bmatrix} \sigma_{e0}^2 \end{bmatrix}$$



bmi<sub>ij</sub> = 
$$\beta_{0ij}$$
cons + 0.209(0.011)agec<sub>ij</sub>  
 $\beta_{0ij}$  = 21.311(0.263) +  $u_{0j}$  +  $e_{0ij}$ 



$$\left[u_{0j}\right] \sim N(0, \ \Omega_u) : \ \Omega_u = \left[7.940(0.952)\right]$$





$$\left[e_{0ij}\right] \sim N(0, \Omega_e) : \Omega_e = \left[4.104(0.282)\right]$$
 sqrt(

 $sqrt(4.104) = 2.0 kg/m^2$ 



bmi<sub>ij</sub> = 
$$\beta_{0ij}$$
cons + 0.209(0.011)agec<sub>ij</sub>  
 $\beta_{0ij}$  = 21.311(0.263) +  $u_{0j}$  +  $e_{0ij}$ 



$$\left[u_{0j}\right] \sim N(0, \ \Omega_u) : \ \Omega_u = \left[7.940(0.952)\right]$$



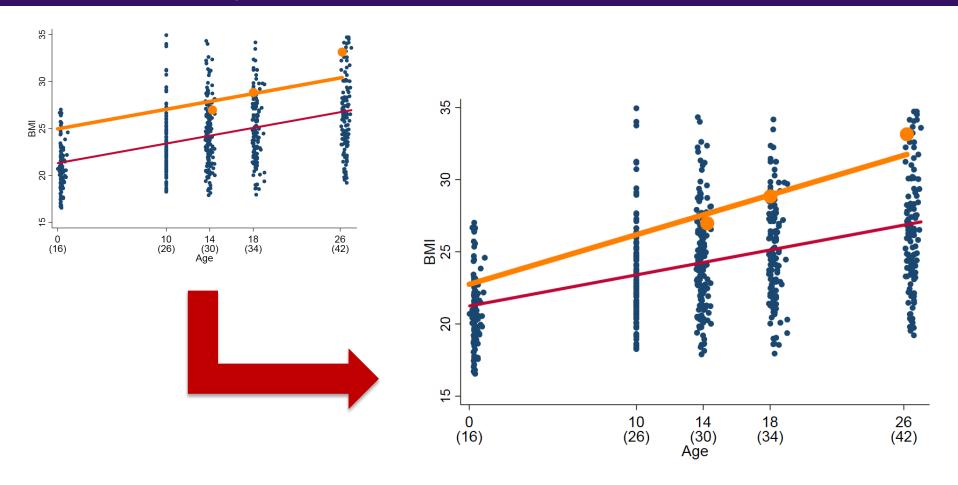
$$\left[e_{0ij}\right] \sim N(0, \ \Omega_e) : \ \Omega_e = \left[4.104(0.282)\right]$$

Variance partitioning coefficient Proportion of total variance that is due to between-individual differences (i.e., level 2 variance)

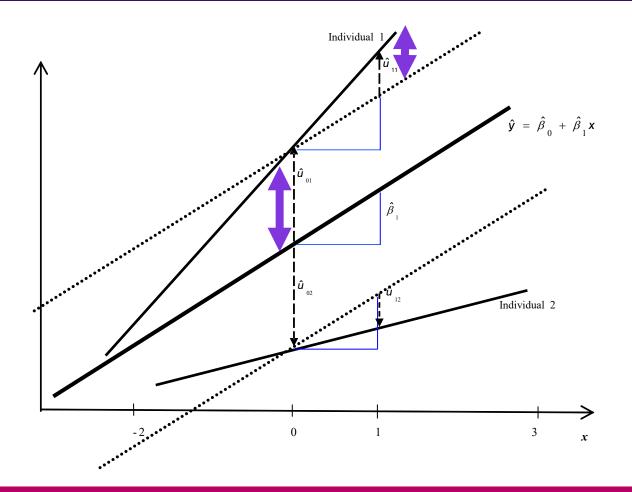
$$sqrt(4.104) = 2.0 kg/m^2$$

 $sqrt(7.940) = 2.8 kg/m^2$ 







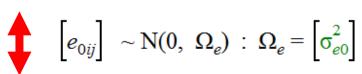




$$bmi_{ij} = \beta_{0ij}cons + \beta_{1j}agec_{ij}$$
$$\beta_{0ij} = \beta_0 + u_{0j} + e_{0ij}$$
$$\beta_{1j} = \beta_1 + u_{1j}$$

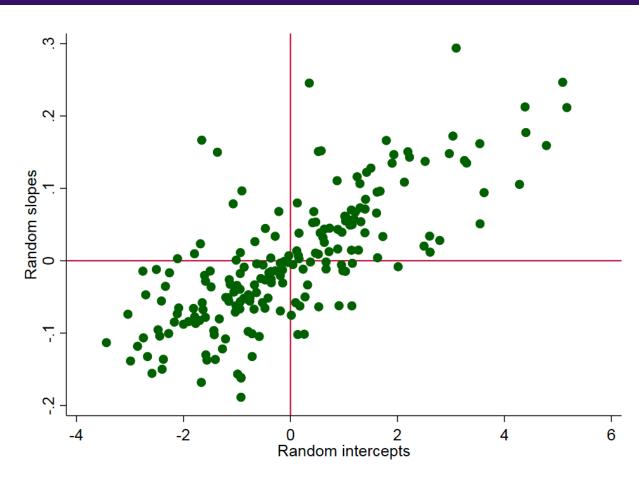


$$\begin{bmatrix} u_{0j} \\ u_{1j} \end{bmatrix} \sim \mathcal{N}(0, \ \Omega_u) : \ \Omega_u = \begin{bmatrix} \sigma_{u0}^2 \\ \sigma_{u01} \ \sigma_{u1}^2 \end{bmatrix}$$



2.967(0.246)

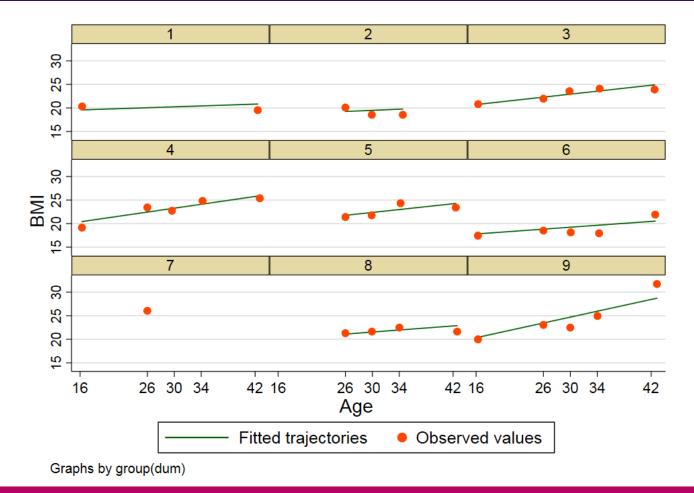




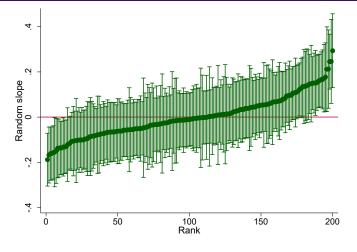
Covariance 0.071

Correlation 0.298

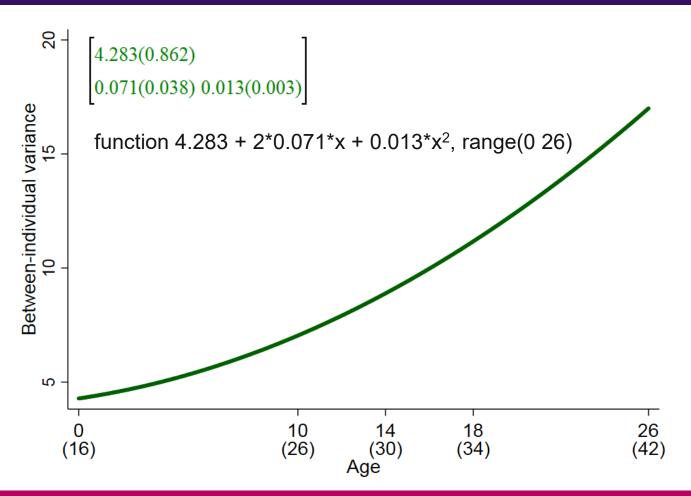








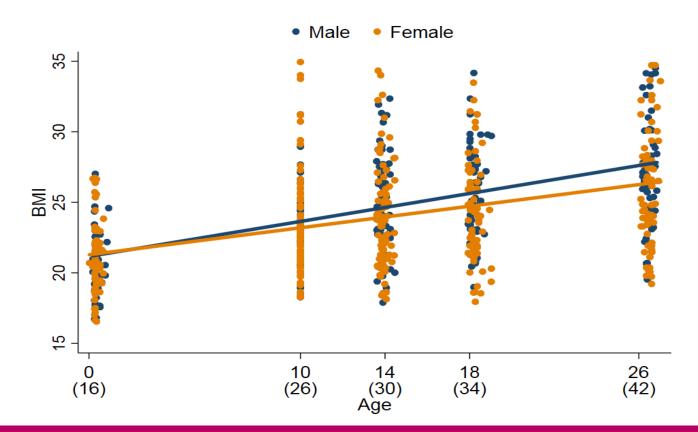






#### Including explanatory variables – level 2

 $bmi_{ij} = \beta_{0ij}cons + \beta_{1j}agec_{ij} + 0.118(0.427)sex_j + -0.057(0.026)agec\_sex_{ij}$ 



# Including explanatory variables – level 1

$$\begin{split} \text{bmi}_{ij} &= \beta_{0j} \text{cons} + \beta_{1j} \text{agec}_{ij} + 0.115(0.423) \text{sex}_j + -0.057(0.025) \text{agec\_sex}_{ij} + e_{4ij} \text{male}_j \\ &+ e_{5ij} \text{female}_j \\ \beta_{0j} &= 21.171(0.319) + u_{0j} \\ \beta_{1j} &= 0.248(0.019) + u_{1j} \end{split}$$

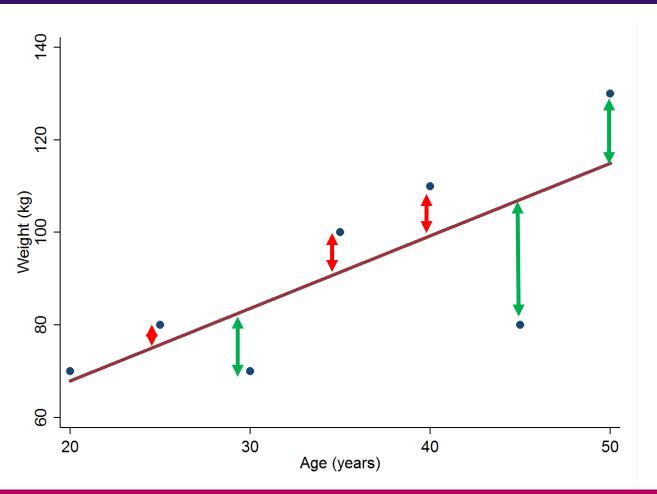
$$\begin{bmatrix} u_{0j} \\ u_{1j} \end{bmatrix} \sim N(0, \ \Omega_u) : \ \Omega_u = \begin{bmatrix} 4.189(0.851) \\ 0.074(0.037) \ 0.012(0.003) \end{bmatrix}$$

$$\begin{bmatrix} e_{4ij} \\ e_{5ij} \end{bmatrix} \sim N(0, \ \Omega_e) : \ \Omega_e = \begin{bmatrix} 2.699(0.343) \\ 0 & 3.202(0.322) \end{bmatrix}$$





#### Including explanatory variables – level 1



#### **Measured**

**Self-reported** 



#### With longitudinal data:

- Multilevel models account for clustering by partitioning variance into within-individual (level 1) and between-individual (level 2)
- A separate trajectory, describing how y changes over age/time, is essentially estimated for each individual
  - Fixed intercept/slope = sample average trajectory
  - Fixed intercept/slope + random intercept/slope = individual trajectories
- Additional explanatory variables at level 1 (self-report) or 2 (sex) can be included to model systematic differences between-individuals
- Additional explanatory variables at level 1 (self-report) or 2 (sex) can be included to model systematic differences within-individuals



### Does accounting for clustering really matter?

Fixed intercept & slope  $\beta_1 = 0.205(0.016)$ 

Vs.

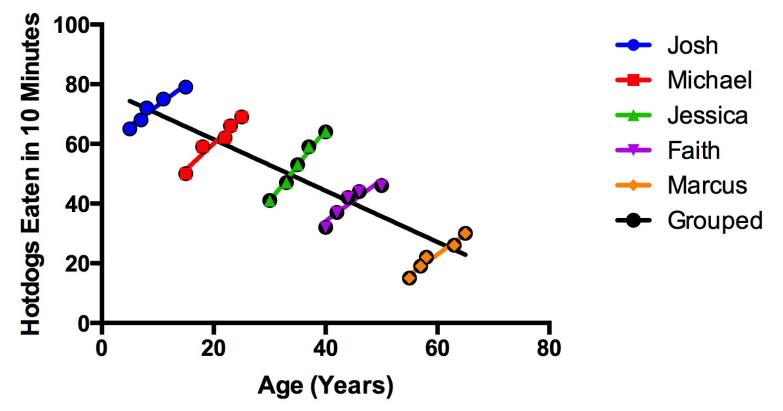
Random intercept & slope  $\beta_1 = 0.216(0.013)$ 





#### Does accounting for clustering really matter?

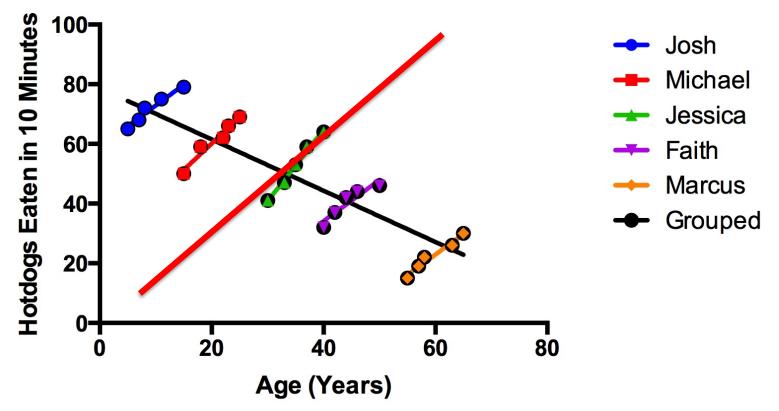
#### **Grouped Hotdogs Eaten in 10 Minutes vs. Age**





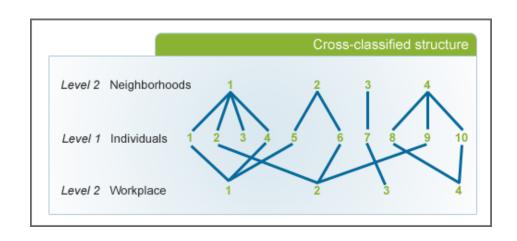
#### Does accounting for clustering really matter?

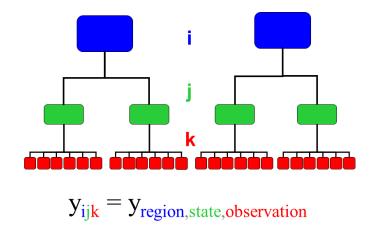
#### Grouped Hotdogs Eaten in 10 Minutes vs. Age





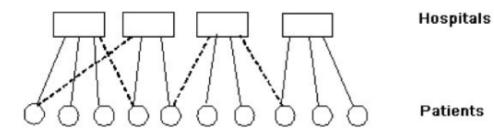
#### Other structures





Multiple membership model: some patients

attend more than one hospital





#### Other structures

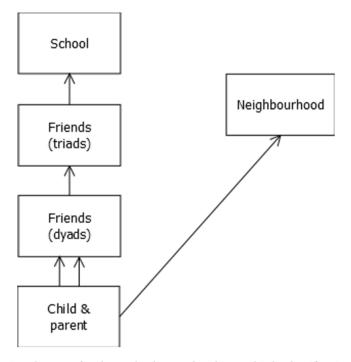




Article

A Multilevel Analysis of Neighbourhood, School, Friend and Individual-Level Variation in Primary School Children's Physical Activity

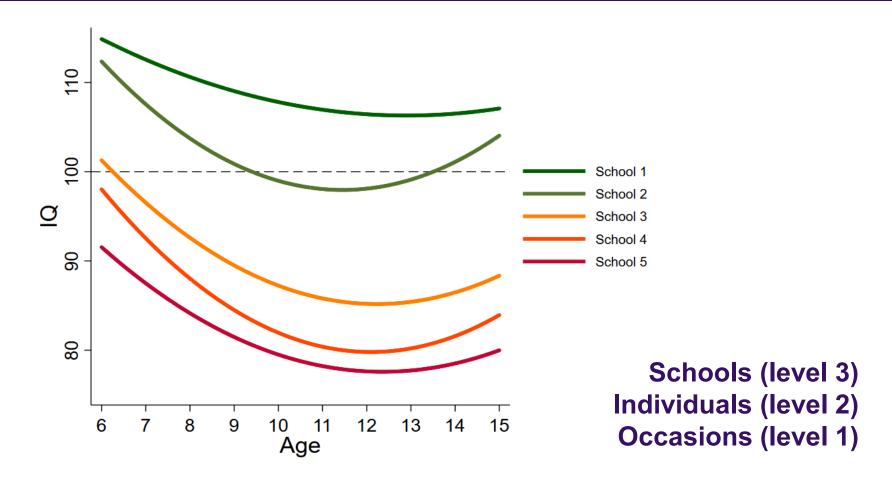
Ruth Salway <sup>1</sup>, Lydia Emm-Collison <sup>1</sup>, Simon J. Sebire <sup>1</sup>, Janice L. Thompson <sup>2</sup>, Deborah A. Lawlor <sup>3,4</sup> and Russell Jago <sup>1</sup>,\*



**Figure 1.** Classification diagram for the multiple-membership multiple-classification model. An arrow indicates a nested relationship; a double arrow indicates multiple membership.

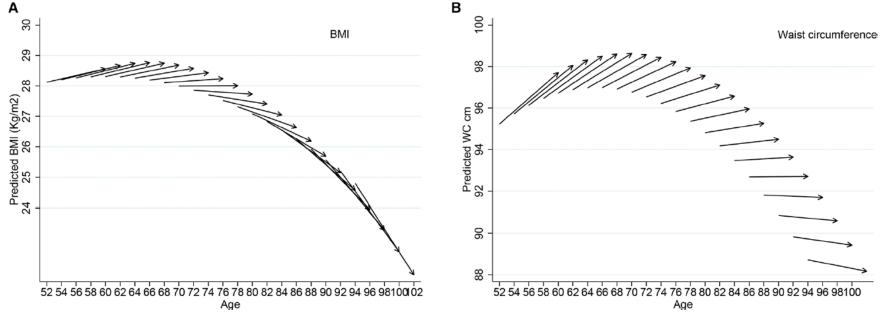


#### Non-linear trajectories





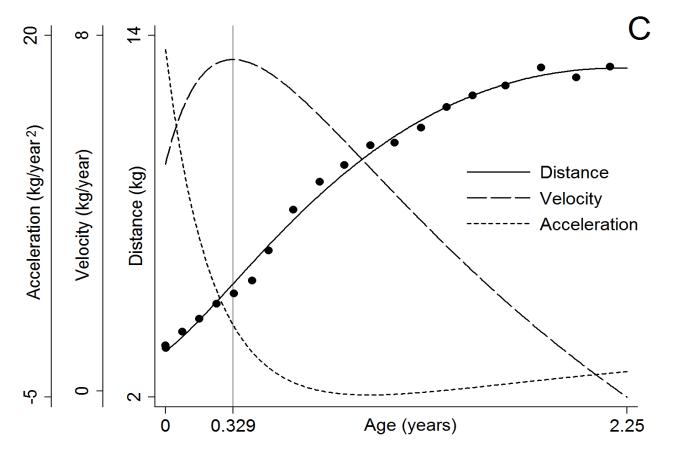
#### Non-linear trajectories



**Figure 1** Vector graph showing 8-year ageing vectors of anthropometric markers, ELSA 2004–2005 to 2012–2013. BMI, body mass index; ELSA, English Longitudinal Study of Ageing; WC, waist circumference.



#### Derivatives and traits





#### Derivatives and traits

Journal of Youth and Adolescence (2019) 48:815–827 https://doi.org/10.1007/s10964-018-0976-5

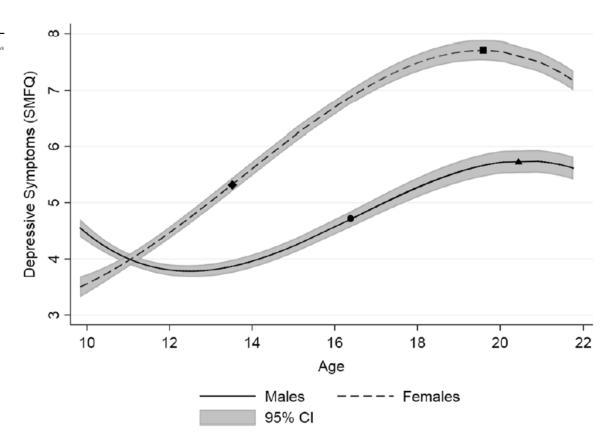
#### EMPIRICAL RESEARCH



#### Identifying Critical Points of Trajectories of Depressive Symptoms from Childhood to Young Adulthood

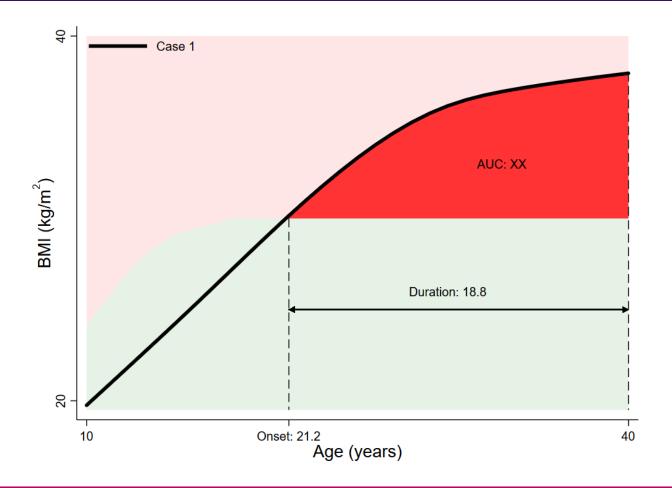
Alex S. F. Kwong 6<sup>1,2,3</sup> · David Manley<sup>1,2</sup> · Nicholas J. Timpson<sup>3,4</sup> · Rebecca M. Pearson<sup>3,4,5</sup> · Jon Heron<sup>3,4,5</sup> · Hannah Sallis<sup>3,4,5,6</sup> · Evie Stergiakouli<sup>3,4,7</sup> · Oliver S. P. Davis<sup>3,4</sup> · George Leckie<sup>2,8</sup>

Received: 7 December 2018 / Accepted: 8 December 2018 / Published online: 22 January 2019 © The Author(s) 2019





#### Derivatives and traits

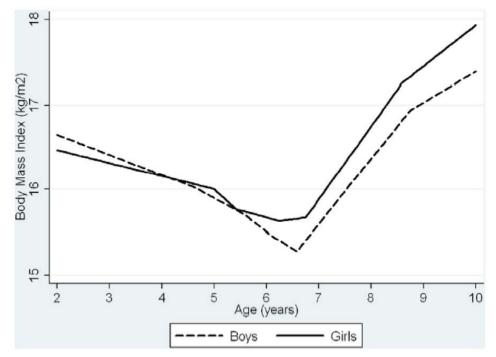




#### Relating level 2 residuals to distal outcomes

Changes in Ponderal Index and Body Mass Index across Childhood and Their Associations with Fat Mass and Cardiovascular Risk Factors at Age 15

Laura D. Howe<sup>1,2</sup>\*, Kate Tilling<sup>2</sup>, Li Benfield<sup>1,2</sup>, Jennifer Logue<sup>3</sup>, Naveed Sattar<sup>3</sup>, Andy R. Ness<sup>4</sup>, George Davey Smith<sup>1,2</sup>, Debbie A. Lawlor<sup>1,2</sup>



Article



Linear spline multilevel models for summarising childhood growth trajectories: A guide to their application using examples from five birth cohorts Statistical Methods in Medical Research 2016, Vol. 25(5) 1854–1874 © The Author(s) 2013 Reprints and permissions: sagepub.co.uk/journalsPermissions.nav DOI: 10.1.177/0962280213503925



Laura D Howe, <sup>1</sup> Kate Tilling, <sup>2</sup> Alicia Matijasevich, <sup>3</sup> Emily S Petherick, <sup>4</sup> Ana Cristina Santos, <sup>5</sup> Lesley Fairley, <sup>4</sup> John Wright, <sup>4</sup> Iná S Santos, <sup>3</sup> Aluísio JD Barros, <sup>3</sup> Richard M Martin, <sup>2,6</sup> Michael S Kramer, <sup>7</sup> Natalia Bogdanovich, <sup>8</sup> Lidia Matush, <sup>8</sup> Henrique Barros, <sup>5</sup> and Debbie A Lawlor, <sup>1</sup>



#### Relating level 2 residuals to distal outcomes

SMMR STATISTICAL METHODS IN MEDICAL RESEARCH

Article

Joint modelling compared with two stage methods for analysing longitudinal data and prospective outcomes: A simulation study of childhood growth and BP Statistical Methods in Medical Research 2017, Vol. 26(1) 437–452 © The Author(s) 2014 Reprints and permissions: sagepub.co.uk/journalsPermissions.nav DOI: 10.1177/0962280214548822 smm.sagepub.com

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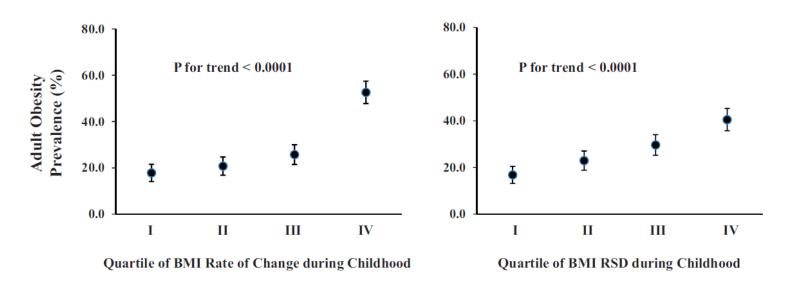
A Sayers, <sup>1</sup> J Heron, <sup>1</sup> ADAC Smith, <sup>1,2</sup> C Macdonald-Wallis, <sup>1,2</sup> MS Gilthorpe, <sup>3</sup> F Steele <sup>4</sup> and K Tilling <sup>1,2</sup>



#### Relating level 1 residuals to distal outcomes

# Variability and rapid increase in body mass index during childhood are associated with adult obesity

Shengxu Li,<sup>1\*</sup> Wei Chen,<sup>1</sup> Dianjianyi Sun,<sup>1,2</sup> Camilo Fernandez,<sup>1</sup> Jian Li,<sup>3</sup> Tanika Kelly,<sup>1</sup> Jiang He,<sup>1</sup> Marie Krousel-Wood<sup>1,4,5</sup> and Paul K Whelton<sup>1</sup>





#### Relating level 1 residuals to distal outcomes

#### RESEARCH ARTICLE



# Estimating the association between blood pressure variability and cardiovascular disease: An application using the ARIC Study

Jessica K. Barrett<sup>1,2</sup> Raphael Huille<sup>2,3</sup> | Richard Parker<sup>4</sup> | Yuichiro Yano<sup>5</sup> | Michael Griswold<sup>6</sup>

<sup>1</sup>MRC Biostatistics Unit, University of Cambridge, Cambridge, UK

<sup>2</sup>Department of Public Health and Primary Care, University of Cambridge, Cambridge, UK

<sup>3</sup>École Nationale de la Statistique et de l'Administration Économique, Malakoff, France

<sup>4</sup>School of Social and Community Medicine, University of Bristol, Bristol, UK

<sup>5</sup>Department of Preventive Medicine, University of Mississippi Medical Center, Jackson, Mississippi

<sup>6</sup>Center of Biostatistics and Bioinformatics, University of Mississippi Medical Center, Jackson, Mississippi

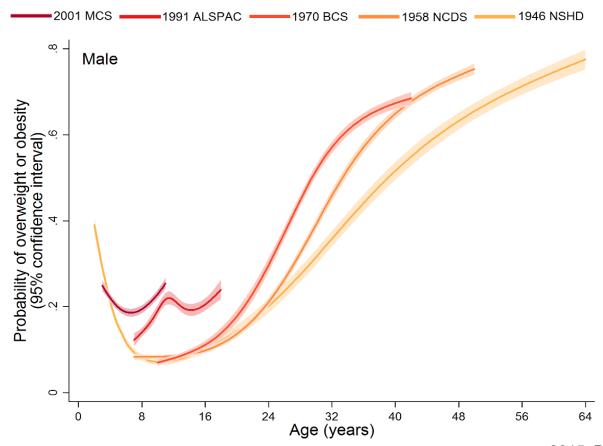
#### Correspondence

Jessica K. Barrett, MRC Biostatistics Unit,

The association between visit-to-visit systolic blood pressure variability and cardiovascular events has recently received a lot of attention in the cardiovascular literature. But, blood pressure variability is usually estimated on a person-by-person basis and is therefore subject to considerable measurement error. We demonstrate that hazard ratios estimated using this approach are subject to bias due to regression dilution, and we propose alternative methods to reduce this bias: a two-stage method and a joint model. For the two-stage method, in stage one, repeated measurements are modelled using a mixed effects model with a random component on the residual standard deviation (SD). The mixed effects model is used to estimate the blood pressure SD for each individual, which, in stage two, is used as a covariate in a time-to-event model. For the joint model, the mixed effects submodel and time-to-event submodel are fitted simultaneously using shared random effects. We illustrate the methods using data from the Atherosclerosis Risk in Communities study.



#### Other outcome distributions



2015. Plos Med 12: e1001828



#### Parallel processes

# Statistics in Medicine

#### **Research Article**

Received 8 March 2011,

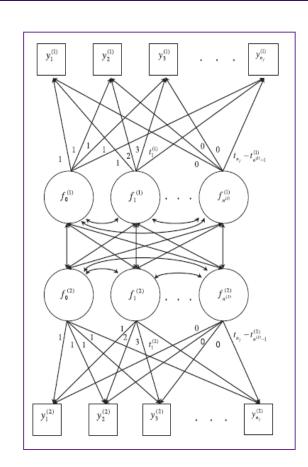
Accepted 6 March 2012

Published online 26 June 2012 in Wiley Online Library

(wileyonlinelibrary.com) DOI: 10.1002/sim.5385

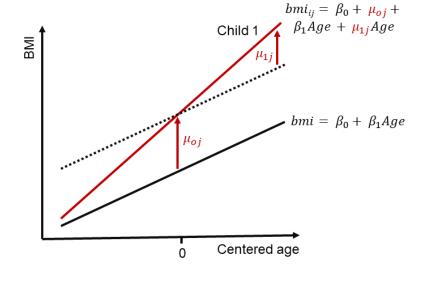
#### Multivariate multilevel spline models for parallel growth processes: application to weight and mean arterial pressure in pregnancy

Corrie Macdonald-Wallis, \*\* Debbie A. Lawlor, \*Tom Palmer\* and Kate Tilling\*





#### Extend to SEM framework



Multilevel (growth curve) model

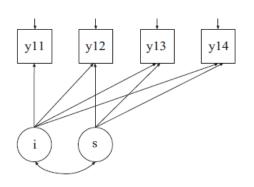
TITLE: this is an example of a linear growth model for a continuous outcome

DATA: FILE IS ex6.1.dat;

VARIABLE: NAMES ARE y11-y14 x1 x2 x31-x34;

USEVARIABLES ARE y11-y14;

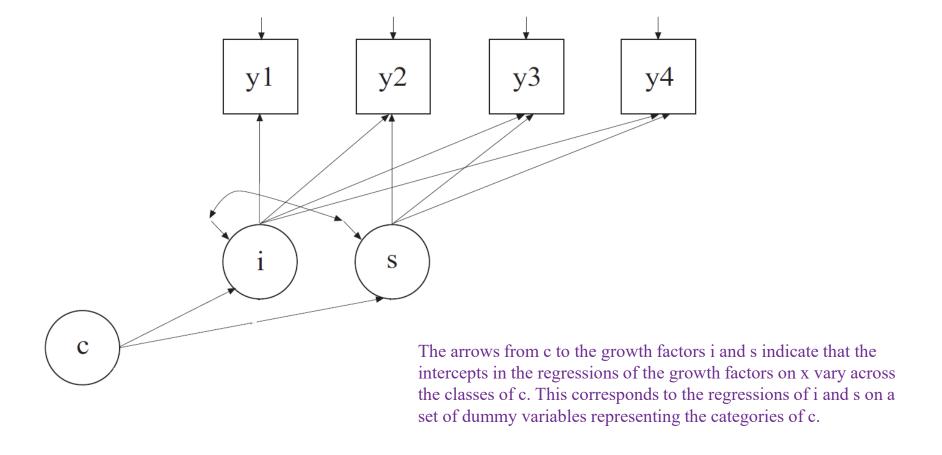
MODEL: i s | y11@0 y12@1 y13@2 y14@3;



Latent (growth curve) model

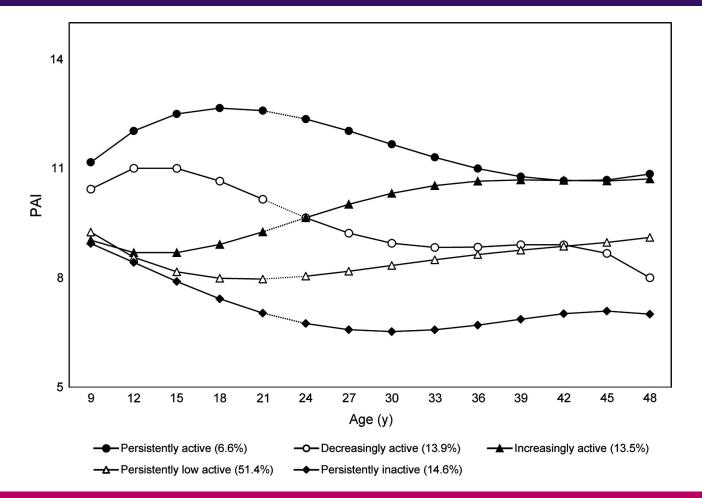


#### Growth mixture model





#### Growth mixture model





#### Resources



#### Centre for Multilevel Modelling



The Centre for Multilevel Modelling (CMM) is a research centre based at the University of Bristol. Our researchers are drawn from the <u>School of Education</u> and <u>School of Geographical Sciences</u>. We collaborate with a range of researchers working with <u>multilevel models</u>.

Johnson W. 2015. Analytical strategies in human growth research. Am J Hum Biol 27(1): 69-83

Tu YK, Tilling K, Sterne JA, Gilthorpe MS. 2013. A critical evaluation of statistical approaches to examining the role of growth trajectories in the developmental origins of health and disease. Int J Epidemiol 42:1327-1339

