

# A joint mean-correlation multilevel model with grouped random effects: application to analysis of household effects in longitudinal studies

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# Household effects in health and social research

- ▶ Substantial interest in accounting for and measuring dependencies among household members in health behaviours and outcomes, social attitudes etc.
- ▶ Within-household correlation due to:
  - ▶ Shared environment, e.g. economic circumstances
  - ▶ Selection of individuals with similar characteristics into co-residence (homophily)
  - ▶ Reciprocal influences over time
- ▶ In epidemiology, interest in whether area effects can be explained by household effects

# Data sources for estimation of household effects

1. **Household panel surveys** track individuals and their coresidents over time
  - ▶ Use to study impact of individual and household characteristics on variety of outcomes, e.g. social inequalities in health
  - ▶ Allow separation of individual, household and area effects
  - ▶ Increasingly linked to administrative data
2. **Linked population registers**

Both types of data source are **longitudinal**.

## Challenges with longitudinal data

How to handle changes in household membership over time, e.g. after adult child leaves parental home or after partnership breakdown?

- ▶ *“Efforts to define a longitudinal household are bound to be futile”* (Duncan & Hill 1985)
- ▶ *“The UK Household Longitudinal Study is not a longitudinal study of households, since arguably households have no coherent existence over time”* (Buck & McFall 2012)

## Previous attempts to estimate household effects

- ▶ Most have studied household effects at a cross-section (one wave of a panel study)
- ▶ Approaches using longitudinal data:
  - ▶ Restrict to 'intact' households (e.g. Keizer & Schenk 2012)
  - ▶ Multiple-membership multilevel model (Goldstein et al. 2000)
  - ▶ Marginal model (Steele, Clarke & Kuha 2019)

## General panel model

Linear model for outcome at wave  $t$  for individual  $i$ :

$$Y_{ti} = \mathbf{x}'_{ti}\beta + r_{ti}$$

where  $r_{ti}$  is a residual.

Within- and **between-individual** covariances:

$$\text{cov}(Y_{ti}, Y_{t'i'} | \mathbf{x}_{ti}, \mathbf{x}_{t'i'})$$

### Approaches to covariance modelling

- ▶ **Marginal:** direct parameterisation of covariance matrix
- ▶ **Random effects:** decompose  $r_{ti}$

## A marginal mean-correlation model for clustered panel data

Joint GLM for marginal expectation and correlation of  $\mathbf{Y}_k$ ,  
response vector for cluster  $k$ :

$$\text{Expectation: } g_1(\mu_k) = \mathbf{X}_k \boldsymbol{\beta}$$

$$\text{Correlation: } g_2(\rho_k) = \mathbf{Z}_k \boldsymbol{\alpha}$$

$\mathbf{Z}_k$  contains characteristics of response pairs  $(Y_{tik}, Y_{t'i'k})$  in cluster  $k$ , e.g. relationship between  $i$  and  $i'$ .

Estimate using 2nd-order GEE (e.g. Yan & Fine 2004) .

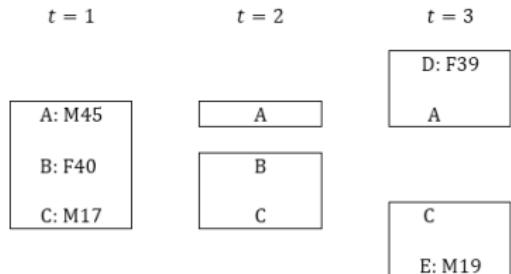
How to define clusters to allow for correlation among individuals connected by coresidence?

## Definition of clusters: “Superhouseholds”

- ▶ View households as evolving social networks
- ▶ A “superhousehold” is a group of individuals linked by pathways of edges in a network graph
- ▶ A superhousehold contains:
  - individuals linked (directly or indirectly) by coresidence
  - observations on same individual over time (autocorrelation)

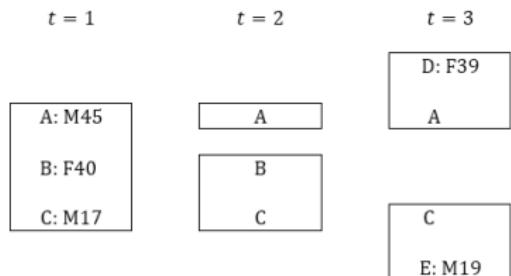
# Evolution of a superhousehold

## Household change over 3 waves

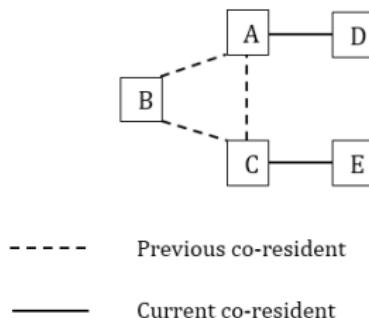


# Evolution of a superhousehold

## Household change over 3 waves



## Superhousehold with coresidence at wave 3



## Common reasons for household change

- ▶ Partnership formation/ dissolution
- ▶ Birth of child
- ▶ Adult child leaves or returns to parental home
- ▶ Parent moves in with adult child
- ▶ Housemate moves in or out (unrelated sharers)

These are not mutually exclusive.

## Marginal model with household/coresidence effects

Joint GLM for marginal expectation and correlation of  $\{Y_{tik}\}$  in superhousehold  $k$  (Steele, Clarke & Kuha 2019):

$$\text{Expectation: } g_1(\mu_{tik}) = \beta' \mathbf{x}_{tik}$$

$$\text{Correlation: } g_2(\text{cor}(Y_{tik}, Y_{t'i'k})) = \alpha' \mathbf{z}_{tik, t'i'k}$$

where  $\mathbf{z}_{tik, t'i'k}$  contains indicators for:

- ▶ Time between waves  $t$  and  $t'$
- ▶ Coresidence status of individuals  $i$  and  $i'$  at  $t$  and  $t'$  (past/present/future)
- ▶ Relationship between  $i$  and  $i'$  (e.g. couple, parent-child)

# Random effects models: motivation

## Marginal model

- ▶ Adjust SEs for clustering in  $\{Y_{tik}\}$  for coresidents
- ▶ Offers insights into nature of between-individual correlations

## BUT

- ▶ Does not provide partitioning of variation
- ▶ Does not extend to additional layers of clustering (e.g. areas)
- ▶ Does not guarantee positive definite estimated within-cluster correlation matrices

## Multiple membership random effects model: idea

Decompose  $r_{ti}$  in general model into 3 terms: individual effects, household effects and residual.

BUT individuals are only nested within households if household membership is fixed over time.

Instead view individuals as members of **multiple** households over time, with appropriate (user-specified) weight attached to each household.<sup>1</sup>

⇒ **Non-hierarchical multilevel model.**

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<sup>1</sup>Goldstein et al. 2000

## Multiple membership random effects model: details

Decompose  $r_{ti}$  into individual and household effects and residual

$$r_{ti} = u_i + v_{ti} + e_{ti}$$

Time-varying household effect:  $v_{ti} = \sum_{h \in \mathcal{H}_i} q_h v_h^*$

$\mathcal{H}_i = (h_{1i}, \dots, h_{Ti})$  is set of households  $i$  belongs to over time,  $v_h^*$  are household effects, and  $q_h$  are (user-specified) weights.

**Problem:** Between-individual covariance structure determined by (arbitrary) choice of weights.

## Proposed alternative: 'grouped' random effects model

Recall linear panel model

$$Y_{ti} = \mathbf{x}'_{ti}\beta + r_{ti}$$

Partition residual  $r_{ti}$  into individual, household and area effects:

$$r_{ti} = \underbrace{u_i}_{\text{ind}} + \underbrace{v_{h(ti)}}_{\text{hh}} + \underbrace{w_{a(ti)}}_{\text{area}} + e_{ti}$$

$h(ti)$  and  $a(ti)$  index household and area of individual  $i$  at wave  $t$ .

Assume all components of  $r_{ti}$  are normally distributed and independent, **except for  $v_{h(ti)}$** .

## Specification of household effects

Household effect for individual  $i$  at wave  $t$  is  $v_{h(ti)}$ .

A new household ID  $h(ti)$  is assigned to  $i$  at  $t$  if **any** change in their coresidents since  $t - 1$ .

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A new household ID  $h(ti)$  is assigned to  $i$  at  $t$  if **any** change in their coresidents since  $t - 1$ .

Households  $h$  and  $h'$  in the same superhousehold are linked through coresidence, so allow for correlation between their random effects:

$$\text{cor}(v_h, v_{h'}) = \gamma' \mathbf{z}_{h,h'} \quad \text{for } h \neq h', \quad s(h) = s(h')$$

$\mathbf{z}_{h,h'}$  are covariates describing link between  $h$  and  $h'$ .

## Random effects model: estimation

Constrained MCMC: block-wise Gibbs/Metropolis-Hastings hybrid.

- ▶ Iterate between sampling random effects and parameters
- ▶ Sample random effects for households in the same superhousehold jointly.

Denote by  $\mathbf{v}_s = (v_1, \dots, v_{m_s})$  the household effects for superhousehold  $s$  with  $m_s$  households.

Assume  $\mathbf{v}_s \sim N(\mathbf{0}, \boldsymbol{\Omega}_{vs})$  where  $\boldsymbol{\Omega}_{vs} = \sigma_v^2 \mathbf{R}_{vs}$ .

- ▶ Sample correlation parameters  $\gamma$  using M-H step to ensure  $\mathbf{R}_{vs}$  is pos. def. for all  $s$  (extension of Zhang, Kuha & Steele 2024).

## Selected simulation results

Impact of misspecification of household effects on random effect variance estimates. Data generated from M3 with 20k superhouseholds. Mean estimates from 200 replications.

	Wave $\sigma_e^2$	Ind $\sigma_u^2$	hh $\sigma_v^2$
True	0.4	0.3	0.3
M1: No hh effects	0.439	0.561	–
M2: Independent hh effects	0.398	0.361	0.209
M3: Correlated hh effects	0.400	0.300	0.300

## Other results

- ▶ Underestimated SEs for household-level covariates
- ▶ Biases for M1 and M2 decrease with intra-household correlation

## UK Household Longitudinal Study (UKHLS)

- ▶ Began 2009–10 with ~ 40k households, annual interviews
- ▶ Two Y at 11 waves: impact of individual's health on everyday **physical** and **mental functioning** (SF-12)
- ▶ Covariates  $x_{ti}$ : age, gender, ethnicity, partnership status, number/age of children, education, employment status, household income, housing tenure
- ▶ Clusters defined as superhouseholds at wave 11<sup>2</sup>
- ▶ Areas are Lower Super Output Areas
- ▶ Analysis sample: 387,238 person-wave observations on 76,053 individuals in 63,400 households; 37,867 superhouseholds

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<sup>2</sup>Superhousehold and hh IDs based on complete enumeration of households.

## Covariates for between-hh random effect correlations

Covariates ( $\mathbf{z}_{h_1 h_2}$ ) describe connection between pair of households  $(h_1, h_2)$  formed at waves  $(t_1, t_2)$ ,  $t_1 \leq t_2$ .<sup>3</sup>

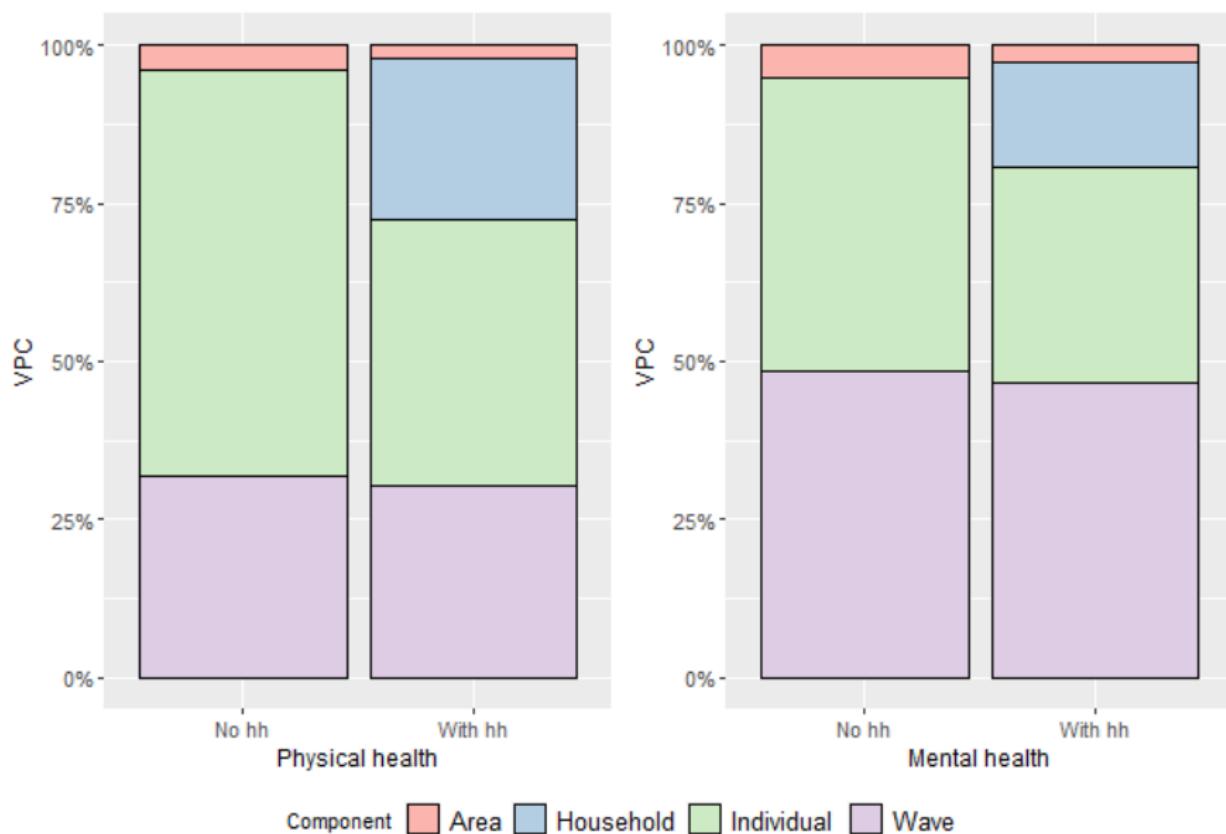
- (i) Indicators of whether  $(h_1, h_2)$  share past/current/future partners
- (ii) Indicators of whether a member of one household is the parent of a member of the other
- (iii) Proportion of total number of individuals across  $h_1$  and  $h_2$  who are in both

Partnership and parent-child links (i) and (ii) account for 91% of all household pairs; remainder mainly from unrelated sharers.

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<sup>3</sup>Using data from complete enumeration of households.

## Unconditional variance decomposition without and with household effects



## Examples of predicted household random effect correlations

Connection between $h_1$ and $h_2$	$\text{cor}(v_{h_1}, v_{h_2})$		Overlap	% pairs
	Phys.	Mental		
Parent and parent-child hhs ( $C \geq 16$ ) i.e. child leaves/enters parent hh	0.737	0.700	0.653	14.2

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Parent and parent-child hhs ( $C \geq 16$ ) i.e. child leaves/enters parent hh	0.737	0.700	0.653	14.2
Separate parent and child hhs ( $C \geq 16$ )	0.439	0.439	0	12.6

## Conditional intra-class correlations, $\text{cor}(r_{ti}, r_{t'i'})$

Type of ICC	Physical	Mental
ICC <sub>ind</sub> : Same ind., diff. waves	0.596	0.504
ICC <sub>hh</sub> : Diff. ind., same hh	0.136	0.177
ICC <sub>shh</sub> : Diff. hhs, same super-hh	0.077 (0.018)*	0.096 (0.024)*
ICC <sub>area</sub> : Diff. super-hhs, same area	0.021	0.022

\*Mean (st. dev.) of ICC<sub>shh</sub>. Variation between hh pairs as between-household correlation depends on covariates.

## Other applications of grouped random effects model

- ▶ Longitudinal data or cross-sectional multivariate data on grouped individuals
- ▶ Groups could be couples, families, schools, workplaces etc
- ▶ Within-group correlations between pairs of individual random effects (or latent traits) depend on covariates
- ▶ Pairwise covariates could be individuals' respective roles (e.g. parent-child, manager-employer), age difference, gender

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