

Conditional vs. marginal estimators of within-pair regression effects in individually-matched case-control studies and twin cohorts

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Acknowledgements

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Acknowledgements No. 1

Jack and Jill...

John Carlin

Jonathan Sterne

John Hopper

and

Gillian Dite

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Collaborators on the more recent work on binary data...

Martin Hazelton

Fizz Williamson

Sabria Khan

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Outline

- ▶ Background and motivation
- ▶ Regression models for continuously-valued paired exposure and outcome data
- ▶ History
- ▶ Conditional estimators
- ▶ Binary data (both exposure and outcome)
- ▶ Estimators of within-pair effect
- ▶ Simulation results
- ▶ Extensions

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Paired data

- ▶ Twins provide naturally matched pairs for studies of human health, although paired data goes beyond just twins.
- ▶ We can exploit within-pair comparisons of data to avoid confounding associations between outcomes and exposures by shared factors.
- ▶ Specific assumptions about shared factors allow the determination of genetic and environmental contributions to disease risk.

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Twin Studies - 1

- ▶ Tradition of focussing on genetic hypotheses
- ▶ Decompose variation in a quantitative trait
- ▶ Compare within-pair correlation of DZ with MZ ($\frac{1}{2}$ under the additive genetic model)
- ▶ *Classical Twin Model* assumes that variation attributable common or shared environment is the same for DZ and MZ twins
- ▶ Lower DZ than MZ within-pair correlation provides evidence that a trait is determined by genetic factors

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Twin Studies - 2

- ▶ Can the twin context provide greater insight on associations?
 - ▶ Cardiovascular risk (blood pressure) with birthweight
 - ▶ Cancer risk (breast density) with physical measures (height, weight, BMI)
- ▶ Ideally like to separate the effect of shared and individual factors (eg maternal versus placental)
- ▶ When can a regression relationship be said to have a genetic basis?

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Individual twin regression

Exposure variable x_{ij} and binary outcome y_{ij} for $i = 1, \dots, n$ and $j = 1, 2$. A cross-sectional or individual-level regression model might propose that

$$E(y_{i1}) = \alpha + \beta x_{i1}$$

$$E(y_{i2}) = \alpha + \beta x_{i2}$$

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Individual twin regression

$$E(y_{i1}) = \alpha + \beta x_{i1}$$

$$E(y_{i2}) = \alpha + \beta x_{i2}$$

If we take the difference between the two equations we get

$$E(y_{i1} - y_{i2}) = \beta(x_{i1} - x_{i2})$$

If we take the average between the two equations we get

$$E(\bar{y}_i) = \alpha + \beta \bar{x}_i$$

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Between- and within-pair regression

These are special cases of a model general model

$$E(y_{i1}) = \beta_0 + \beta_w(x_{i1} - \bar{x}_i) + \beta_b\bar{x}_i$$

$$E(y_{i2}) = \beta_0 + \beta_w(x_{i2} - \bar{x}_i) + \beta_b\bar{x}_i$$

where $\bar{x}_i = (x_{i1} + x_{i2})/2$. Since

$$x_{i1} - \bar{x}_i = (x_{i1} - x_{i2})/2$$

$$\begin{aligned}x_{i2} - \bar{x}_i &= (x_{i2} - x_{i1})/2 \\ &= -(x_{i1} - x_{i2})/2\end{aligned}$$

we can re-write the multivariable between- and within-pair model as...

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Between- and within-pair regression

These are special cases of a model general model

$$E(y_{i1}) = \beta_0 + \beta_w(x_{i1} - x_{i2}) + \beta_b \bar{x}_i$$

$$E(y_{i2}) = \beta_0 + \beta_w(x_{i2} - x_{i1}) + \beta_b \bar{x}_i$$

Univariate regressions of the within-pair differences and within-pair means yield estimates of β_w and β_b respectively.

Simultaneous estimation of β_w and β_b from the multivariable model generates the same estimates for OLS and GLS (but standard errors will differ).

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Interpretation - 1

- ▶ β_w is the expected change in the the outcome y for a unit change in the *deviation* of the exposure x from the pair mean, holding this pair mean constant.
- ▶ β_b is the expected change in the outcome y for a unit change in the *pair mean* \bar{x} , holding the within-pair deviation (difference) constant.

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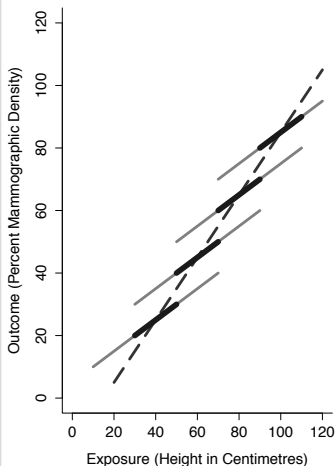
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Illustration of between- and within-pair effects

Between- and within-pair regression effects



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Interpretation - 2

What we are postulating is

- ▶ A model for the expected value of the outcome y can be improved by using data from pairs.
- ▶ A good way to do this is to relate the expected value of y_{i1} not just to x_{i1} (the twin's own exposure value) but also to their co-twins exposure value x_{i2} .
- ▶ The expected difference in outcome y comparing between two x values may depend on whether we are comparing (i) co-twins with each other within-pair; or (ii) unrelated twins between pairs.

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Twin – Co-Twin regression

The multivariable model re-expressed

$$E(y_{i1}) = \beta_0 + \beta_t x_{i1} + \beta_c x_{i2}$$

$$E(y_{i2}) = \beta_0 + \beta_t x_{i2} + \beta_c x_{i1}$$

where

$$\beta_t = (\beta_w + \beta_b)/2$$

$$\beta_c = (\beta_w - \beta_b)/2$$

from which we can see that $\beta_c = 0$ ($E(y_{i1})$ does not depend on x_{i2} and *vice versa*) is equivalent to $\beta_w = \beta_b$ (between- & within-pair reg. effects are the same).

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Individual twin regression

Recall the individual-level regression model

$$E(y_{i1}) = \alpha + \beta x_{i1}$$

$$E(y_{i2}) = \alpha + \beta x_{i2}$$

If the multivariable between- and within-pair regression model is correct, then fitting the individual-level regression (again, by either OLS or GLS) produces an estimate of β that is a weighted average of the corresponding estimates of β_w and β_b with weights that depend on ρ_x and ρ_y , the observed within-pair correlation of x and y respectively.

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Weighted average estimates of β

This results first came to my attention through **biostatistics** via the seminal paper by Neuhaus & Kalbfleisch (1998) in *Biometrics*.

Neuhaus & Kalbfleisch (1998), however, quote Scott & Holt (1982) in *J. Amer. Stat. Assoc.*, a paper on two-stage **sample surveys**.

Scott & Holt (1982) in turn trace the result back to Maddala (1971) in *Econometrica*, so we're now in **economics** where the interest at the time was “pooling cross section and time series data”.

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Weighted average estimates of β

There's more: Maddala (1971) references this

Wallace TD & Hussain A (1969). The use of error components models in combining cross section with time series data. *Econometrica*, **37**, 55–72,

which in turn refers to this

Hildreth C (1950). Combining Cross-Section Data and Time Series. Cowles Commission Discussion Paper: Statistics No. 347, May 15, 1950.

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Weighted average estimates of β

This led me to propose **Gurrin's Law**: One can always find a reference to the between- and within-cluster “beta is weighted average” result published before one was born *regardless* of how old one is!

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Papers by John Neuhaus et al.

- ▶ **Neuhaus JM & Jewell N** (1990). The effect of retrospective sampling on binary regression models for clustered data. *Biometrics*, **46**, 977 – 990.
- ▶ **Neuhaus JM & Kalbfleisch JD** (1998). Between- and within-cluster covariate effect in the analysis of clustered data. *Biometrics*, **54**, 638 – 645.
- ▶ **Neuhaus JM & McCulloch CE** (2006). Separating between- and within-cluster covariate effects by using conditional and partitioning methods. *J. R. Statist. Soc. B.*, **68**, 859 – 872.

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Papers on interpretation

- ▶ **Begg MD & Parides MK (2003)**. Separation of individual-level and cluster-level covariate effects in regression analysis of correlated data. *Statistics in Medicine*, **22**, 2591 – 2602.
- ▶ **Carlin JB, Gurrin LC, Sterne JAC, Morley R & Dwyer T (2005)**. Regression models for twin studies: a critical review. *Int. J. Epidemiol.*, **34**, 1089 – 1099.
- ▶ **Dwyer T, Blizzard CL (2005)**. A discussion of some statistical methods for separating within-pair associations among all twins in research on fetal origins of disease. *Paediatric and Perinatal Epidemiology*, **19**(1), 48 – 53.
- ▶ **Gurrin LC, Carlin JB, Sterne JAC, Dite GS & Hopper JL (2006)**. Using bivariate models to understand between- and within-cluster regression coefficients with application to twin data. *Biometrics*, **62**, 745 – 751.

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The Sjolander Show

- ▶ **Sjolander A, Lichtenstein P, Larsson H & Pawitan Y.** (2012). Between-within models for survival analysis. *Statistics in Medicine*.
- ▶ **Sjolander A, Frisell T & Oberg S.** (2012). Causal interpretation of between-within models for twin research. *Epidemiologic Methods*, **1**(1), No. 10.
- ▶ **Sjolander A, Johansson ALV, Lundholm C, Altman D, Almqvist C & Pawitan Y.** (2012). Analysis of 1:1 matched cohort studies and twin studies, with binary exposures and binary outcomes. *Statistical Science*, **27**(3), 395-411.

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Binary x and y

- ▶ We now consider the scenario where both x_{ij} and y_{ij} are binary 0/1 variables.
- ▶ $x_{i1} - \bar{x}_i = (x_{i1} - x_{i2})/2 = -(x_{i2} - x_{i1})/2$ can take only three values: $-\frac{1}{2}$, 0 or $\frac{1}{2}$.
- ▶ \bar{x}_i can take only three values: 0, $\frac{1}{2}$ or 1.
- ▶ A pair is *exposure-concordant* if $x_{i1} = x_{i2}$, so $\bar{x}_i = 0$ or $\bar{x}_i = 1$, otherwise it is *exposure-discordant* where $\bar{x}_i = \frac{1}{2}$.
- ▶ A pair is *outcome-concordant* if $y_{i1} = y_{i2}$, otherwise it is *outcome-discordant*.

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Individual twin regression

Exposure variable x_{ij} and binary outcome y_{ij} with expectation p_{ij} for pair i and individual j .
An ordinary logistic regression model implies:

$$\log \left(\frac{p_{ij}}{1 - p_{ij}} \right) = \beta_0 + \beta x_{ij}$$

So for each pair we have:

$$\log \left(\frac{p_{i1}}{1 - p_{i1}} \right) = \beta_0 + \beta x_{i1}$$

$$\log \left(\frac{p_{i2}}{1 - p_{i2}} \right) = \beta_0 + \beta x_{i2}$$

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Individual twin regression

Within-pair difference in the two regression equations:

$$\log\left(\frac{p_{i1}}{1-p_{i1}}\right) - \log\left(\frac{p_{i2}}{1-p_{i2}}\right) = \beta(x_{i1} - x_{i2})$$

Average the two regression equations:

$$\frac{1}{2} \left[\log\left(\frac{p_{i1}}{1-p_{i1}}\right) + \log\left(\frac{p_{i2}}{1-p_{i2}}\right) \right] = \beta_0 + \beta \bar{x}_i$$

- ▶ Both equations depend on the exposure x through the regression coefficient β .
- ▶ We can, however, generalise to allow the exposure effect on the within-pair log odds ratio and the between-pair average log odds to be distinct.

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Between- and within-pair regression

- ▶ The model proposed by Neuhaus & Kalbfleisch (1998) where $p_{ij} = E(y_{ij})$ is

$$\log\left(\frac{p_{ij}}{1 - p_{ij}}\right) = \beta_0 + \beta_w(x_{ij} - \bar{x}_i) + \beta_b\bar{x}_i$$

for $j = 1, 2$ and $\bar{x}_i = (x_{i1} + x_{i2})/2$.

- ▶ We can also write

$$\log\left(\frac{p_{ij}}{1 - p_{ij}}\right) = \beta_0 + \beta_w\frac{1}{2}(x_{ij} - x_{ik}) + \beta_b\bar{x}_i$$

where $k = (3 - j)$, indicating that this model has terms for both between- and within-pair regression effects.

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Paired data in matched studies

- ▶ Binary outcome data from individual matched pairs is typically analysed for associations with exposures using conditional logistic regression (CLR).
- ▶ CLR uses the likelihood *conditional* on the sum of the pair's outcome (0, $\frac{1}{2}$ or 1) to estimate the regression parameter β .

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Paired data in matched studies

- ▶ Pairs that are either outcome-concordant or exposure-concordant do not contribute to the conditional likelihood.
- ▶ So in an individually matched case-control study, where $y_{i1} \neq y_{i2}$ by design, only exposure-discordant pairs contribute to the estimation of β , an inherently within-pair regression parameter.

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Paired data in cohort studies

- ▶ But what about outcome-concordant twin- or sib-pairs appearing in cohort studies?
- ▶ Can their inclusion improve the precision of estimation of regression parameters while still maintaining the benefits of a paired analysis?

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Paired data in cohort studies

- ▶ We know their inclusion does not influence the results of CLR.
- ▶ But what about the multivariable between- and within-pair regression model?
- ▶ What is the relationship between estimates of β from CLR and estimates of β_w from ordinary logistic regression with the multivariable model?

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Estimation of β_b and β_w

- ▶ Use ordinary logistic regression (OLR) on *all* pairs to estimate β_0 , β_w and β_b , and take β_w as our association parameter. The OLR estimate of β_w depends on the assumptions for β_0 & β_b (ie whether we estimate them or set them to zero).
- ▶ Use conditional logistic regression (CLR) on pairs that are both exposure-discordant *and* outcome-discordant (so “doubly-discordant”) pairs to estimate β_w . The CLR model does not have parameters β_0 and β_b .

There is a “close empirical correspondence” (Neuhaus & Kalbfleisch (1998), Ten Have *et al.* (1995), Sjolander *et al.* (2012)) between the OLR and CLR estimates...
...but they are **not** formally identical.

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Table : Summary of paired binary data for one pair member exposed and one pair member unexposed

	Unexposed member ($x_2 = 0$)		
	Event ($y_2 = 1$)	No event ($y_2 = 0$)	Totals
Exposed member ($x_1 = 1$)			
Event ($y_1 = 1$)	n_{11}	n_{10}	$n_{11} + n_{10}$
No event ($y_1 = 0$)	n_{01}	n_{00}	$n_{01} + n_{00}$
Totals	$n_{11} + n_{01}$	$n_{10} + n_{00}$	$\sum \sum n_{ij}$

CLR estimator of β_w

For a single binary exposure the CLR estimate of β_w is the log ratio of number of exposure-outcome concordant pairs (n_{10}) to exposure-outcome discordant pairs (n_{01}) among **outcome-discordant** pairs:

$$\hat{\beta}_w(\text{CLR}) = \log(n_{10}/n_{01})$$

with standard error

$$\text{s.e.}(\hat{\beta}_w(\text{CLR})) = \sqrt{n_{10}^{-1} + n_{01}^{-1}}.$$

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OLR estimators of β_w : $\beta_b = \beta_0 = 0$

For OLR there is an additional contribution from the $n_{11} + n_{00}$ exposure-discordant pairs that are **outcome-concordant** pairs. For $\beta_b = \beta_0 = 0$ we have

$$\hat{\beta}_w(\text{OLR}, \beta_b = \beta_0 = 0) = 2 \log \left(\frac{n_{10} + \frac{1}{2}(n_{11} + n_{00})}{n_{01} + \frac{1}{2}(n_{11} + n_{00})} \right)$$

with s.e. $(\hat{\beta}_w(\text{OLR}), \beta_b = \beta_0 = 0) =$

$$\sqrt{\left(n_{10} + \frac{1}{2}(n_{11} + n_{00}) \right)^{-1} + \left(n_{01} + \frac{1}{2}(n_{11} + n_{00}) \right)^{-1}}.$$

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OLR estimators of β_w : $\beta_b \neq 0, \beta_0 \neq 0$

For \bar{x}_i categorical, β_b and β_0 replaced with their MLE's we have

$$\hat{\beta}_w(\text{OLR}) = \log \left(\frac{n_{10} + n_{11}}{n_{01} + n_{11}} \times \frac{n_{10} + n_{00}}{n_{01} + n_{00}} \right)$$

with $\text{s.e.}(\hat{\beta}_w(\text{OLR}))^2 =$

$$(n_{10} + n_{11})^{-1} + (n_{01} + n_{11})^{-1} + (n_{10} + n_{00})^{-1} + (n_{01} + n_{00})^{-1}$$

where

$$n = n_{00} + n_{01} + n_{10} + n_{11}.$$

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OLR estimators of β_w : General Form

More generally we have $\hat{\beta}_w =$

$$\log \left[\frac{n_{10}^{1-\alpha_1/2} + \alpha_2(\alpha_3 n_{11} + (1-\alpha_3)n_{00})}{n_{01}^{1-\alpha_1/2} + \alpha_2(\alpha_3 n_{11} + (1-\alpha_3)n_{00})} \right]$$
$$\times \left[\frac{n_{10}^{1-\alpha_1/2} + \alpha_2((1-\alpha_3)n_{11} + \alpha_3 n_{00})}{n_{01}^{1-\alpha_1/2} + \alpha_2((1-\alpha_3)n_{11} + \alpha_3 n_{00})} \right]$$

where $(\alpha_1, \alpha_2, \alpha_3) =$

$(1, 0, 0)$ for CLR;

$(0, \frac{1}{2}, \frac{1}{2})$ for OLR with $\beta_b = \beta_0 = 0$; and

$(0, 1, 0)$ for OLR with $\beta_b = \hat{\beta}_b$ and $\beta_0 = \hat{\beta}_0$.

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Unanswered questions

- ▶ The result when $\beta_b \neq 0$ and $\beta_0 \neq 0$ is from Sjolander *et al.* (*Stat. Sci.*, 2012). They state “...the decomposition into within- and between-effects is a legitimate method for binary exposures, which was questioned by Carlin *et al.* (2005)”.
- ▶ **BUT** this result only applies when we model the mean effect $m(\bar{x}_i)$ as

$$\beta_0 \mathbf{I}(\bar{x}_i = 0) + \beta_{0.5} \mathbf{I}(\bar{x}_i = 0.5) + \beta_1 \mathbf{I}(\bar{x}_i = 1)$$

that is, the mean appears in the regression equation as a categorical variable.

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Unanswered questions

- ▶ In this case only exposure-concordant pairs contribute to estimating β_w , so it's not much of a “decomposition into within- and between-effects”.
- ▶ Explicit results for \bar{x}_i as a continuously valued exposure (albeit with only three possible values) are not available, but one can show the precision of $\hat{\beta}_w$ is (slightly) greater than for the above categorical model.

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Simulation: $\beta_0 = \beta_b = 0, \beta_w = 1$

Table : Summary of simulation results for 100 datasets with 500 twin pairs. True values are $\beta_0 = 0, \beta_b = 0, \beta_w = 1$. $z \sim N(0, \sigma^2)$ with between- and within-pair std dev $\sigma = 2$. The observed binary covariate is $x = I(z > 0)$.

	Mean Estimate	Empirical Std Err	Model Std Err
CLR	0.98	0.25	0.28
OLR ($\beta_b = \beta_0 = 0$)	0.96	0.24	0.26
OLR ($\beta_b = \hat{\beta}_b, \beta_0 = \hat{\beta}_0$)	0.97	0.24	0.26

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Simulation: $\beta_0 = 0, \beta_b = \beta_w = 1$

Table : Summary of simulation results for 100 datasets with 500 twin pairs. True values are $\beta_0 = 0, \beta_b = 1, \beta_w = 1$. $z \sim N(0, \sigma^2)$ with between- and within-pair std dev $\sigma = 2$. The observed binary covariate is $x = I(z > 0)$.

	Mean Estimate	Empirical Std Err	Model Std Err
CLR	1.02	0.28	0.29
OLR	0.92	0.24	0.26
$(\beta_b = \beta_0 = 0)$			
OLR	0.99	0.26	0.27
$(\beta_b = \hat{\beta}_b, \beta_0 = \hat{\beta}_0)$			

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Conclusions

- ▶ CLR and OLR estimators of the within-pair regression effect are specific examples of a general estimator that assigns weights to count data from outcome-concordant exposure-discordant pairs.
- ▶ OLR estimators are potentially more efficient than CLR estimators since they use data from all exposure-concordant pairs, rather than just those that are outcome-discordant, an assertion that is born out by our simulation studies.

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Extension 1: Two time-points

Let y_{ijk} (x_{ijk}) be the outcome (exposure) at time $k = 1, 2$ for the j^{th} twin in the i^{th} twin pair.

Then we can decompose x_{ijk} as using a two-way analysis of variance

$$\begin{aligned}x_{ijk} = & 0.5(x_{ijk} - \bar{x}_{ij.}) + 0.5(x_{ijk} - \bar{x}_{i.k}) \\ & + 0.5(\bar{x}_{ij.} - \bar{x}_{i..}) + 0.5(\bar{x}_{i.k} - \bar{x}_{i..}) \\ & + \bar{x}_{i..}\end{aligned}$$

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Extension 2: Count data

Between- and within-pair effects for **Poisson regression of count outcomes**. The unconditional estimate (OPR?) of β_w is the difference in the log of the mean count between exposed and unexposed.

The conditional estimate (CPR?) of β_w is (approximately) the log of the mean difference in counts d between exposed and unexposed (if $d > 0$). More formally

$$\exp(\hat{\beta}_w) = d/2 + \sqrt{(d/2)^2 + 1}$$

so $\exp(\hat{\beta}_w) > 0$ even if $d < 0$.

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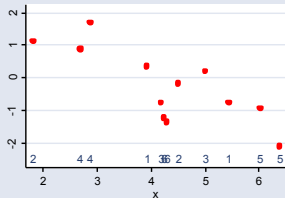
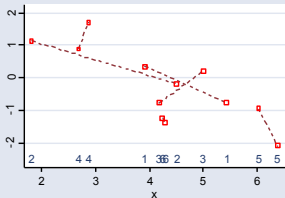
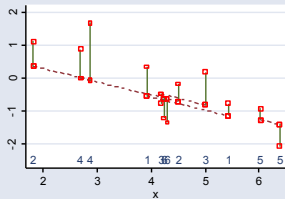
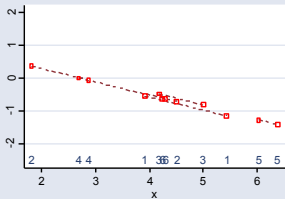
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Graph 1

model (1) $\beta_C = -0.4$



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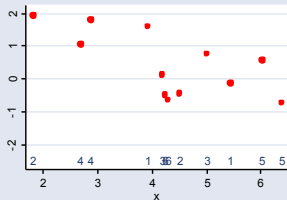
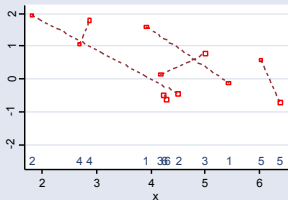
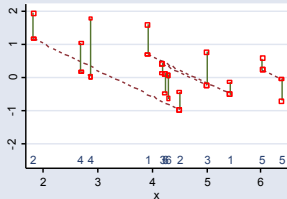
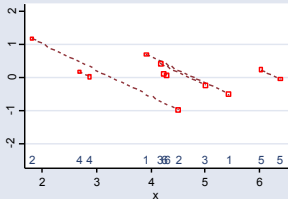
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Graph 2

model (2a) $\beta_W = -0.8, \beta_B = 0$



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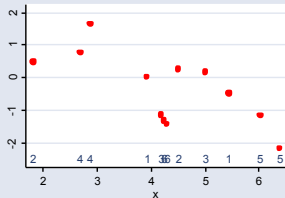
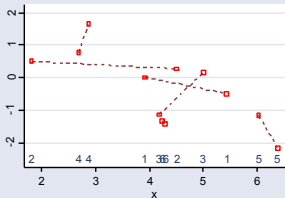
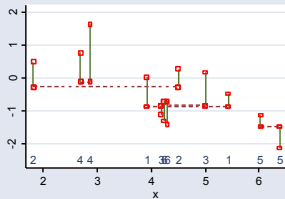
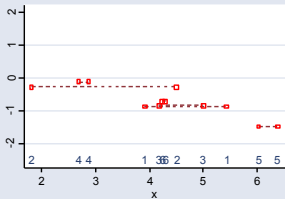
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Graph 3

model (2b) $\beta_W = 0$, $\beta_B = -0.4$



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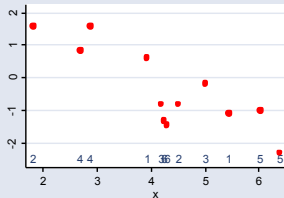
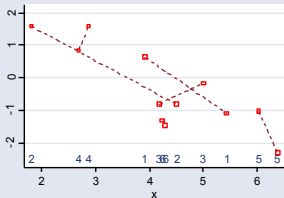
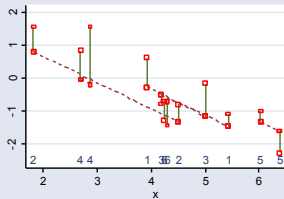
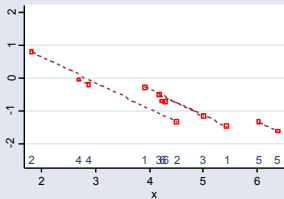
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Graph 4

model (2c) $\beta_W = -0.8, \beta_B = -0.4$



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