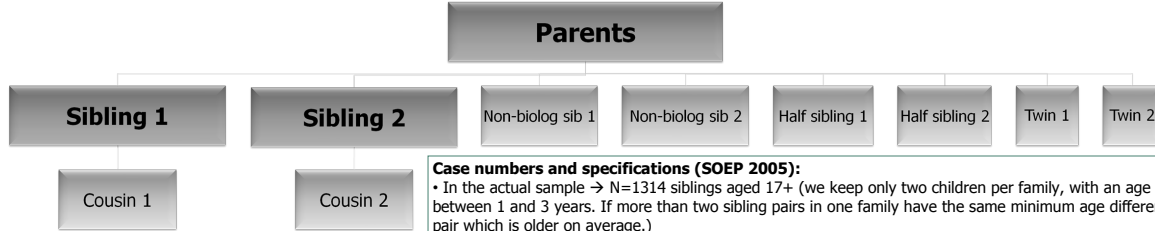


Feasibility Study for an Extended Twin-Family Design Related Study based on SOEP

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Family Data in the SOEP



Case numbers and specifications (SOEP 2005):

- In the actual sample → N=1314 siblings aged 17+ (we keep only two children per family, with an age difference between 1 and 3 years. If more than two sibling pairs in one family have the same minimum age difference, we keep the pair which is older on average.)
- Non-biological children → N=53 (15 mothers have one biological and one non-biological child and 19 mothers have 2 non-biological children)
- Half siblings, foster children and cousins yet to be identified. Potential spouses of both generations are possible to identify.

Extended Twin-Family-Design by an extra-sampling of genetically informative groups Test with a related sample is underway

Extended Twin-Family Design

Combination of offspring and parent data to separate different genetic and environmental factors

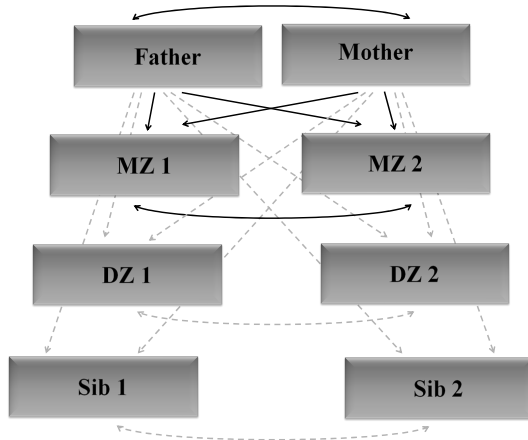
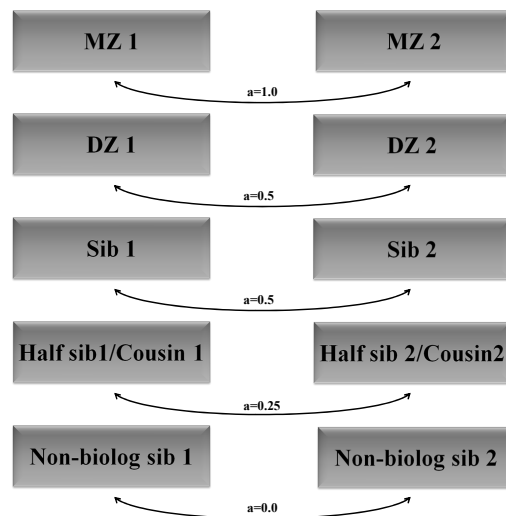


Figure caption:

MZ= monozygotic twins; DZ= dizygotic twins; Sib= full siblings; Half sib= half siblings; Cousin= children of siblings; Non-biolog sib= adoptive children, step children or foster children; 1 and 2= first and second member of a pair; a= additive genetic relatedness

Multi-Group Design

Simultaneous comparison of covariances within different groups varying in genetic and environmental similarity



Actual sample size:

- MZ → N=358 individuals, 179 pairs (Jena, TwinPaw, Infratest)
- DZ → N=268 individuals, 134 pairs (Jena, TwinPaw, Infratest)
- Sib → N=1314 individuals, 657 pairs (SOEP)

Methodological possibilities

Extended Twin-Family Design

- Differentiation of common environmental and non-additive genetic effects
- Increased power to discriminate between additive and non-additive genetic effects
- Increased power to detect the effect of assortative mating
- Reliable estimation of cultural transmission

SOEP and Registries

- Measurement in the home (incl. possibilities for tests)
- Greater flexibility regarding psychometric and diagnostic requirements
- Representative longitudinal design (incl. all members of a family)
- Tailoring of assessments along the line of content requirements
- Interdisciplinary data sets (psychological, social and economic research topics)

Multi-Group Design

- Consideration of gene-environment (G-E) correlation
- Investigation of gene-environment (G×E) interaction
- More robust estimates of genetic and environmental components than classical twin design
- Greater generalizability to population
- Possibility of examining the sensitivity of results to variations in modeling assumptions
- Possibility of testing the equal environment assumption and specific twin environments

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