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Abstract

Previous research has demonstrated the potential of adversarial collaboration (AC) in empirical studies to reduce ambiguity in findings and facilitate a truthful representation of opposing viewpoints. In this study, we propose an AC framework specifically designed for Monte Carlo simulation studies to enhance methodological rigor and generalizability in this line of research. Our framework involves two parts: conducting independent preliminary studies, followed by a joint study to synthesize and debate the results, utilizing tools to structure, document, and evaluate the AC process. To test this framework, we conducted a focused case study comparing iterative structural after measurement (SAM) estimation and traditional structural equation model (SEM) estimation. After conducting our individual studies, we integrated our results. The findings revealed no substantial disagreements between the studies, and left us unable to conduct a joint study. Still, we found some preliminary evidence that AC can increase transparency, rigor, and epistemic accountability in simulation research. Furthermore, our study provides initial insights into the practical applicability and technical feasibility of AC, supported by tools like Github for collaborative work. Although we were unable to test our AC framework in its entirety, our findings suggest that AC holds significant potential for improving the quality of simulation studies. Future research should further explore AC in simulations to validate and extend these findings.

Introduction

Monte Carlo simulations serve as a widely adopted method for evaluating and comparing statistical methods in quantitative science. They are used to assess the performance of estimation and inference methods by analyzing them with knowledge of simulated population models and values. Although simulations provide insight into underlying data structures, they are not without flaws and are subject to pitfalls similar to those implicated in the replication crisis of empirical research (Koehler et al., 2009; Lohmann et al., 2022). A major challenge in simulation studies is to ensure the generalizability of the results to real-world contexts (Feinberg & Rubright, 2016). It is crucial for these studies to use performance metrics, experimental factors, and inference models that accurately mirror and test practical scenarios. Given the impracticality of simulating every conceivable model, method, and use case, these studies necessarily involve numerous decision-making elements that may introduce bias: The decisions of selecting evaluation criteria for competing methods (for example absolute versus relative bias of performance), selecting which models or methods to compare in the first place, and which simulation specifications to run (for example deciding which sample sizes to simulate), can have substantial impact on the results (Kulinskaya et al., 2020; Pawel et al., 2023). Evidently, many issues of open science in the empirical realm are present in the context of simulation studies as well.

One approach that addresses these issues at the level of value selection suggests to sample all relevant values from existing results published in the empirical literature of interest (Bollmann et al., 2015). An example application could be to decide on a simulated sample size, not just based on a hunch or prior experience, but based on previous empirical studies conducted in the field. Even though this idea is fruitful and improves upon the problem of generalizability, the argument with regards to the arbitrariness of decision-making still remains. For example, deciding on which papers, or even which models within a paper to select (and which not), can still be subject to individual bias or simply chance. Additionally, conducting extensive literature reviews for all parameter values chosen is time-consuming and difficult to apply in practice.

Other promising approaches focus on applying well-known concepts of open science to simulation studies: Multiple frameworks propose using preregistration in simulation studies and call for increased transparency in reporting of key aspects of simulation studies (Boulesteix, 2015; Morris et al., 2019; Siepe et al., 2023). While increasing transparency and accountability in this way is important for reducing research ambiguity, these approaches do not necessarily incline scientists to refrain from choosing values at will, as long as they are preregistered and reported transparently.

Another approach to tackle these challenges that has been proposed in the empirical context

is Adversarial Collaboration (AC). Initially tested by Mellers et al. (2001), adversarial collaboration is gaining recognition within the quantitative empirical research community for its potential to enhance scientific rigor. Touted as “The next science reform” adversarial collaboration involves scholars with opposing viewpoints working together to resolve scientific disputes (Clark & Tetlock, 2023). An ongoing effort by Melloni et al. (2023), for example, conducts adversarial collaboration between proponents of two different theories on the relationship between consciousness and brain activity, hoping to advance research in this field. Adversarial Collaboration involves pinpointing empirical disagreements, designing studies that are mutually agreed upon to test competing hypotheses, and jointly publishing the results regardless of the outcome. The goal is to ensure fair comparison and truthful representation of opposing views, thereby enhancing epistemic accountability and reducing ambiguity in scientific decision-making (Clark et al., 2022a). Moreover, juxtaposing and integrating competing positions in this manner can improve the generalizability of results.

Our objective is to apply the advantages of adversarial collaboration to simulation studies. To achieve this, we design and implement a focused case study, utilizing an exemplary topic in the literature of simulation studies: Comparing the newly proposed iterative structural after measurement (SAM) estimation approach to structural equation model (SEM) estimation with traditional, non-iterative SEM estimation (Dhaene & Rosseel, 2023; Robitzsch, 2022; Rosseel & Loh, 2022). In their respective simulations, the authors’ results differed up to the point of contradiction, providing us with ideal grounds for adversarial collaboration. While Dhaene & Rosseel (2023) concluded that SAM estimation generally outperforms non-iterative SEM in small samples, Robitzsch (2022) did not find the methods to differ. Similarly, whereas the former found SAM to be more robust against model misspecification, the latter argued the opposite to be true. Applying adversarial collaboration, we, the authors of this paper, each represent one of these competing positions, as detailed later.

This allows us to examine the applicability of adversarial collaboration in Monte Carlo simulation studies as well its potential to enhance methodological rigor and generalizability in this domain of research. Thus, this investigation leads us to address the following research question:

Can adversarial collaboration be applied to simulation studies, in terms of practical applicability and technical feasibility?

Methods

This section contains two parts that represent our proposed AC framework. First, we describe the procedure of the AC with its two parts and the tools it uses. Then, we propose a technical setup to facilitate its implementation.

Procedure of the Adversarial Collaboration

To answer our research question, we created and implemented the following procedure for the adversarial collaboration process: The procedure consists of two parts. In the first part, each researcher conducts an independent preliminary study. In the second part, a joint study, including a joint simulation protocol, should be pursued.

Part one: Individual studies

As a first step of the individual studies, the two adversarial sides have to agree on one or more research questions that should be as specific as possible and directly tackle the disagreement. These individual studies should include the generation of individual simulation protocols, as suggested by previous work (Morris et al., 2019; Siepe et al., 2023). To facilitate a structured comparison and integration of studies throughout the collaboration process, we agreed on a structure for conducting our individual simulation studies in advance. This structure (see Table 1) is based on relevant literature in the field of simulation studies (Boomsma, 2013; Morris et al., 2019; Paxton et al., 2001).

This structure allows for the expected divergences in results of the individual simulations, but at the same time provides a basis for systematic comparison and synthesis for the joint simulation.

Part two: Joint study

In the second part, a joint study including a joint simulation protocol should be pursued. This process starts based on the results of Part one. Here, the main part of the adversarial collaboration takes place. It contains multiple parts I will explain in the following sections: Each step of the individual simulation studies in Table 1 (from the substantive research question up until the interpretation of the results) is scrutinized and debated between collaborators, using adversarial collaboration techniques. Decisions are made and documented based on the most convincing argument presented, if possible.

Table 1*Structure of simulation studies*

Step	Phase
1	Defining Aims and Objectives <ul style="list-style-type: none"> - Verbal description of the research question, making it specific. - Examples include examining goodness-of-fit statistics and comparing ML to 2SLS.
2	Specification of Population Model <ul style="list-style-type: none"> - Optional and depending on the field. - Modularities include: <ul style="list-style-type: none"> - Structure: e.g., CFA or SEM - Size: number of latents and indicators - Complexity: cross-loaded indicators, reciprocal paths, exogenous predictors - Selection of target model.
3	Data Generation Mechanism <ul style="list-style-type: none"> - Resampling vs. parametric model draw. - Random number draw for data generation.
4	Experimental Design Simulation Procedures <ul style="list-style-type: none"> - Determine factors to vary, levels, and whether fully, partly factorial, or one at a time. - Examples include: <ul style="list-style-type: none"> - Sample size - Distribution of the observed variables - Extent of misspecification
5	Method Selection <ul style="list-style-type: none"> - Varies depending on research question. - Examples include type and number of estimation methods to be compared.
6	Defining Estimands / Population Level Values <ul style="list-style-type: none"> - Should reflect values commonly encountered in applied research. - Considerations for power issues and bias due to misspecification. - Example: Parameters should be statistically significant, even at the smallest sample size of the simulation.
7	Performance Measures <ul style="list-style-type: none"> - Selection and justification of use of measures such as bias, sensitivity/specificity, predictive accuracy. - Decision on number of simulations for acceptable Monte Carlo SE for these measures.

Step	Phase
8	Software Selection <ul style="list-style-type: none"> - Software to run simulation using specific packages & functions.
9	Analysis and Interpretation Plan <ul style="list-style-type: none"> - Analysis: descriptive vs. inferential. - Interpretation: decision criteria that evaluate performance (e.g., if $1-SS > 90\%$, the method performs well).
10	Coding and Execution <ul style="list-style-type: none"> - Amount and content of scripts. - Include (sanity) checks, setting seeds, troubleshooting & verification.
11	Analyzing Results <ul style="list-style-type: none"> - Descriptive, graphical, inferential, and exploration.
12	Reporting & Presentation <ul style="list-style-type: none"> - Provide rationales for each choice made in previous steps. - Publishing code and simulated data.

If an interest in the evaluation of the collaboration exists, diary entries can be written after each step. In the end, each collaborator reports the results in their own paper. If desired, a joint paper can be published as well.

Adversarial collaboration techniques

We are aware that adversarial collaboration has its limitations when it comes to decision making in joint studies, and has lead to unresolvable disagreements in previous studies (Cowan et al., 2020; Mellers et al., 2001). In order to mitigate this risk, next to each collaborator publishing their own paper, we propose to conduct the joint study in a structured and formalized manner. To this end, we identified several collaboration techniques presented in Table 2. These were also meant to facilitate the evaluation of the adversarial collaboration with regards to our main research question.

Documentation - Decision log

As the number of decisions made and their documentation is large, only the most important results should be presented in the respective papers. In addition, the appendix should contain a separate decision log with a detailed and complete documentation of all decisions made.

Table 2

List of adversarial collaboration techniques.

Core Disagreements: Arrive at clearly defined core disagreements that might be the origin for conflicts. (Clark et al., 2022b)

Assumption check: List, question, and categorize assumptions (Kardos & Dexter, 2017)

Red-Teaming: Generating what if scenarios, to identify limitations of the adversaries' approach (Kardos & Dexter, 2017)

Quality of information (Kardos & Dexter, 2017): checking the adversaries' quality of evidence based on literature.

Third neutral arbiter: In case of fundamentally unresolvable disagreements, a third neutral arbiter (Aaron Peikert) will be consulted to try to resolve them.

Here, the summarized results of the AC-techniques used for each step of the simulation study, as well as their consequence for the decision-making process should be detailed.

Another aspect of structuring the collaboration within the decision log, lies in the way decision making is implemented in the joint study of our framework. In our mind, decision making can be based on four distinct grounds: Evidence-based, pragmatic reasons, arbitrary reasons or other reasons (e.g. personal values, political issues). Firstly, we deem a decision evidence based, if one can find a clear answer for a disagreement, based on empirical evidence or in the literature more general. Secondly, to be able to keep the scope of this project, decisions can be made for pragmatic reasons, as for example in the presence of time constraints. Thirdly, arbitrary reasons could be any agreement where the first two grounds are not present, but still a decision has to be made. While this reason does not help in deciding for an option directly, it helps in understanding how often there is no substantive or pragmatic reason for decisions in simulation studies. If this is the case, one might resort to deciding at random. Lastly, there might be other reasons that lead to a decision, that can not be anticipated but should still be captured.

These four reasons ground decision-making and aim to ease the collaborative process by giving a structure. In some cases, more than one of these reasons can be present. In such cases, their relevance to a decision should be ranked, if possible. An example is the following mock log entry:

Decision element: factor of sample sizes

Result: c(50, 100, 200)

Grounds: Primary - Evidence-based, as per Peikert et al. (2020);

Secondary pragmatic reason, average of individual suggestions.

Diary

The adversarial collaboration in Part 2 (Joint study) can be documented and evaluated from each collaborators subjective perspective using semi-structured diary entries after each step. We provided an exemplary diary template that includes questions based on Shah & Leeder (2016) in our Github repository under this link: <https://github.com/lkosanke/AdversarialSimulation/blob/main/diary-template.md>. The diary has the purpose of accumulating data for the evaluation of the AC and the collaboration procedure we propose. In our case, collecting this qualitative data was supposed to help us answer our research question. The evaluation of the diary entry can be conducted in a semi-structured and comparative manner, resorting for example to analysis of quantity of words to identify common themes of the collaboration.

Technical setup

As simulation studies are computationally intensive and require a structured approach to save, share and review code, we propose using Github to ease collaboration (Peikert et al., 2021). Github allows to set up and synchronize multiple repositories to allow both individual and collaborative work. After setting up a first repository at one adversaries user account, the other party can fork that repository to have a copy at their own account. These forks can be synchronized at any time, but also left separate. This functionality aligns well with the different parts of our procedure: For the individual studies, each adversary can work in their own repository. For the joint study, both can work in one repository, and synchronize the other one at will. Additionally, the Github releases feature allows to align milestones such as the publishing of the simulation protocols or the finished result analysis of all studies. These can even be assigned a DOI via Zenodo (European Organization For Nuclear Research & OpenAIRE, 2013).

Results

The structure of this section mirrors our AC procedure. I will start presenting the preparations, methods and results of the individual studies and their substantive topics. I will then integrate the results of these studies to form a substantive conclusion. Lastly, I will analyze the results of the collaboration and thus answer our main research question.

Preparation of the individual studies

For our substantive topic of interest, Leonard Kosanke replicated relevant parts of the study by Robitzsch (2022), and Valentin Kriegmair replicated parts of the studies by Rosseel & Loh (2022) and Dhaene & Rosseel (2023).

As a first step, after reading the relevant literature in the field and informing us thoroughly on our side of the argument, we agreed on two substantive research questions to depict the disagreement of the results as precisely as possible:

1. How do SAM and traditional SEM methods (including maximum-likelihood (ML) and unweighted-least-squares) compare in terms of bias, Root Mean Squared Error (RMSE), and convergence rates in small to moderate samples?
2. What is the impact of model misspecifications, such as residual correlations and cross-loadings, on the performance of SAM compared to traditional SEM methods?

Next, we started the AC by conducting our individual studies. We replicated and in some parts extended on the results of Robitzsch (2022) and Dhaene & Rosseel (2023). The next sections presents the methods and results of these studies, as well as their integration.

Individual study by Kriegmair

Methods

The following two sections contain the methods and results for the studies implemented by Valentin Kriegmair, including the tables and figures. I added them italicized, as they are direct quotes.

The following description of the simulation studies is based on the established structure for simulation studies to co-align with the other conducted studies to facilitate a potential collaboration.

Aims, objectives and research questions

Both studies aimed to evaluate the performance of traditional maximum-likelihood SEM (SEM), compared to global SAM (GSAM), local SAM with maximum likelihood (LSAM-ML), and local SAM with unweighted least squares (LSAM-ULS) estimation under various conditions. The two research questions we established prior to conducting the studies served as basis for both studies.

Population Models and Data Generation Mechanisms

Study 1:

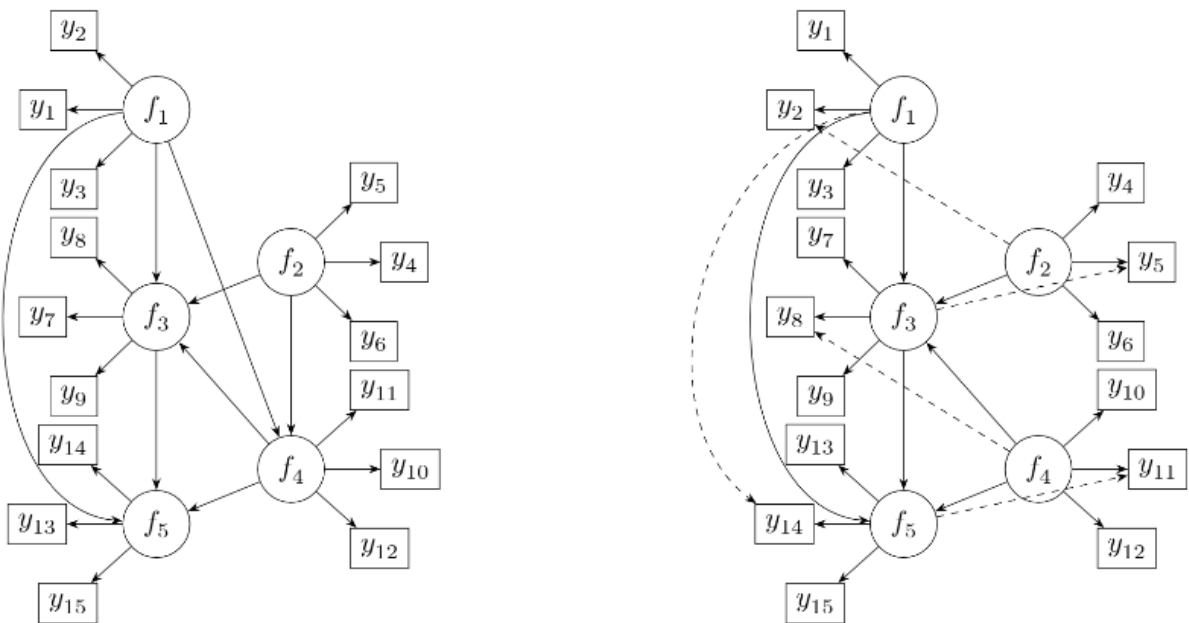
Data was generated based on a 5-factor population structural model with 3 indicators for each factor. Four different models were simulated (see Figures 1-2).

- Model 1.1: Correctly specified model.
- Model 1.2: Misspecified with cross-loadings ignored in the estimation model.
- Model 1.3: Misspecified with wrong direction of path and omitted correlated residuals.
- Model 1.4: Misspecified with multiple omitted structural paths.

For all models, the population-level values of the structural regression parameters were set to 0.1. Indicator reliability levels were manipulated with factor loadings set at low ($\lambda = 0.3$), moderate ($\lambda = 0.5$), or high ($\lambda = 0.7$).

Figure 1

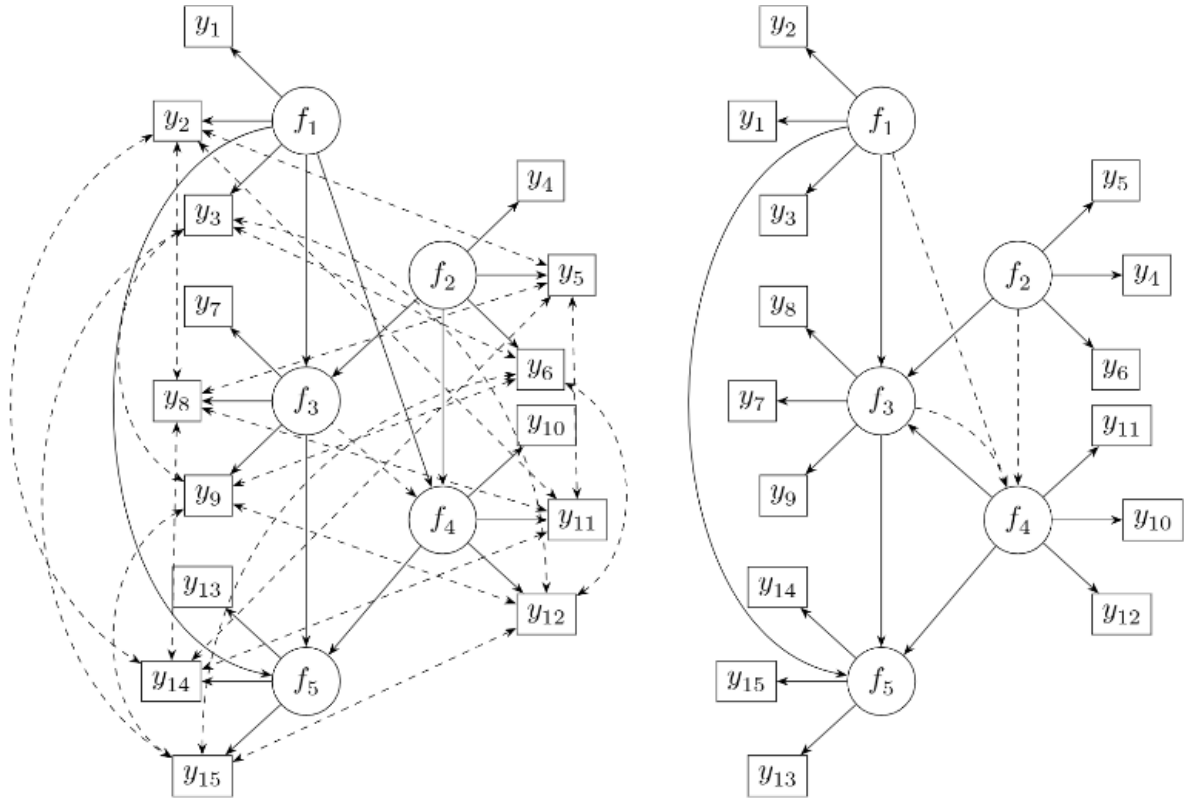
Study 1: Models 1.1 (left) and 1.2 (right).



Note. Error terms are not shown in the figure.

Figure 2

Study 1: Models 1.3 (left) and 1.4 (right).



Note. Error terms are not shown in the figure.

Study 2:

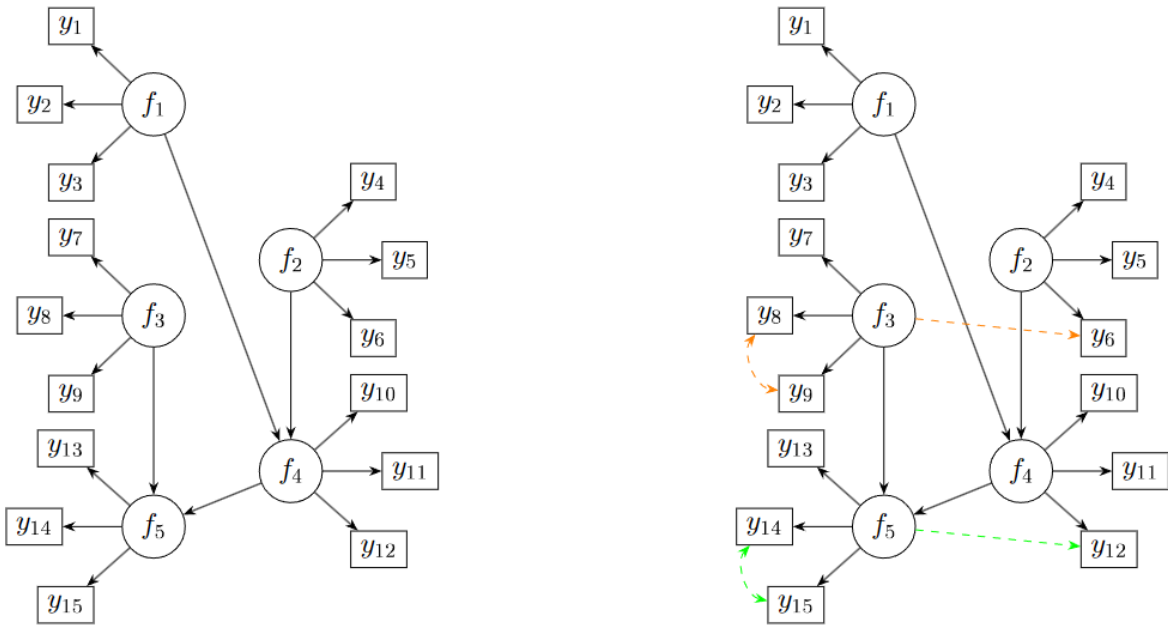
Data were generated based on a 5-factor population structural model with 3 indicators for each factor, similar to Study 1, but with the additional condition of varying variance explained (R^2) by the endogenous factors was set at low ($R^2 = 0.1$) or medium ($R^2 = 0.4$), so that the regression weights were either between 0.183 and 0.224 (low) or between 0.365 and 0.447 (medium). Note however that the computation of this was a simplification and did not accurately result in said R^2 values. The aim here was only generally to modulate between lower and higher regression weights. In study 2, two different models were simulated (see Figure 3).

- Model 2.1: Correctly specified model.
- Model 2.2: Misspecified in either exogenous or endogenous parts of the model

The Misspecifications in Model 2.2 included omitting a residual covariance and a factor cross-loading in either the exogenous or endogenous part of the model, or both. These types of misspecifications resulted in different realizations of model 2.2 when they were modulated as factors as described below.

Figure 3

Study 1: Models 2.1 (left) and 2.2 (right).



Note. Error terms are not shown in the figure. Orange lines represent misspecifications in the exogenous part of the model. Green lines represent misspecifications in the endogenous part of the model.

Experimental Design of simulation procedures

Study 1

The study varied three main conditions:

- *Sample sizes: small ($N = 100$), moderate ($N = 400$), and large ($N = 6400$).*
- *Indicator reliability: low ($\lambda = 0.3$), moderate ($\lambda = 0.5$), high ($\lambda = 0.7$).*
- *Model specifications: correctly specified model and misspecified models (1.2, 1.3, and 1.4).*

Study 2

The study varied five main conditions:

- *Sample sizes: small ($N = 100$), medium ($N = 400$), and large ($N = 6400$).*
- *Variance explained by endogenous factors: low ($R^2 = 0.1$) and medium ($R^2 = 0.4$).*
- *Indicator reliability: low ($\lambda = 0.3$), moderate ($\lambda = 0.5$), and high ($\lambda = 0.7$).*
- *Model misspecifications: varying the population model by omitting a residual covariance and a factor cross-loading in different parts of the model.*

- *Number of measurement blocks: separate measurement model per latent variable ($b = 5$) and joint measurement model for all exogenous variables ($b = 3$).*

Both studies were conducted with 10000 replications for each condition.

Method Selection

In both studies, the performance of four estimation methods was compared: Traditional SEM, GSAM, LSAM-ML and LSAM-ULS estimation.

Performance Measures

The following performance measures were captured: Convergence rates, empirical relative biases, empirical coverage levels of 95% confidence intervals (CIs) and Root Mean Squared Errors (RMSE).

Software

All analyses were conducted in R (R Core Team, 2023). See <https://github.com/valentinkm/AdversarialSimulation> for more details.

Analysis and Interpretation plan

The results were interpreted by descriptively comparing the performance measures (bias, MSE, convergence rates) of the different estimation methods under varying sample sizes, indicator reliability levels, and model misspecifications.

Results

Below are the results (including the tables and figures) by Valentin Kriegmair as a direct quotation or copy. This is why they are italicized.

In the following the main patterns and trends of the results are reported and only exemplary. The full results can be found in the Github repository.

Study 1

Convergence Rate

For moderate and large sample sizes ($N = 400, 6400$), all methods achieved 100% convergence. At a small sample size ($N = 100$), SEM showed lower convergence rates, especially at lower reliability (0.3), while GSAM, LSAM-ML, and LSAM-ULS maintained high convergence rates. SEM faced significant convergence drops at the lowest reliability (0.3), particularly with misspecifications present.

Average Relative Biases

Here this study diverged from the original of Rosseel & Loh (2022) by focusing exclusively on the average relative bias under a single type of correctly specified model (model 1), which did not include any cross loadings or correlated residuals. This simplification allowed to concentrate on the core advantage of SAM over traditional SEM: its more robust estimation of structural parameters, especially when measurement models are misspecified. For the correct model (1.1), SEM showed higher biases at smaller sample sizes and lower reliability, while GSAM, LSAM-ML, and LSAM-ULS exhibited lower biases.

Table 3 shows average relative bias percentages for the different estimation methods under varying model specifications and sample sizes in presence of omitted correlated residuals (model 1.2) and Table 4 under correlated residuals and structural misspecification. Under omitted cross loadings (model 1.2) and structural misspecification (model 1.4), all methods demonstrated substantial biases, particularly SEM at smaller sample sizes and lower reliability. With omitted correlated residuals and structural misspecification (1.3), GSAM, LSAM-ML, and LSAM-ULS showed more negative biases compared to SEM. Overall, GSAM, LSAM-ML, and LSAM-ULS were more robust with lower biases compared to SEM, especially under challenging conditions with smaller sample sizes and lower reliability.

Coverage

For the correct model (1.1), SEM exhibited undercoverage at smaller sample sizes and lower reliability, indicating narrow CI's and less reliable estimates. GSAM, LSAM-ML, and LSAM-ULS showed slight overcoverage, reflecting more conservative estimates. Under the cross loadings model (1.2), all methods had substantial undercoverage, especially SEM. The correlated residuals model (1.3) consistently revealed undercoverage across all methods, with SEM showing the highest biases. For the structural model (1.4), SEM suffered from undercoverage at lower sample sizes and reliability, while GSAM, LSAM-ML, and LSAM-ULS performed better with relatively consistent biases.

Table 3

Study 1: Average relative bias (in percentage) under omitted cross-loadings (Model 1.2)

N	Reliability	SEM	GSAM	LSAM-ML	LSAM-ULS
100	0.3	141.771	43.908	43.075	45.883
100	0.5	83.748	44.476	44.696	45.435
100	0.7	48.378	37.107	37.257	37.057
400	0.3	92.112	52.356	52.824	52.500
400	0.5	77.962	48.421	48.775	48.136
400	0.7	48.172	38.923	39.097	38.427
6400	0.3	81.625	54.110	54.676	53.948
6400	0.5	66.948	48.572	48.948	48.164
6400	0.7	46.667	39.142	39.316	38.585

Note. N = Sample size, SEM = Maximum-likelihood estimation, GSAM = Global-SAM-maximum-likelihood-estimation, LSAM-ML = Local-SAM-maximum-likelihood-estimation, LSAM-ULS = Local-SAM-unweighted-least-squares estimation.

Table 4

Study 1: Average relative bias (in percentage) under omitted residual correlations (Model 1.3)

N	Reliability	SEM	GSAM	LSAM-ML	LSAM-ULS
100	0.3	-39.158	-56.090	-56.109	-55.530
100	0.5	-27.914	-32.240	-32.251	-30.610
100	0.7	-14.898	-16.446	-16.453	-15.897
400	0.3	-51.468	-53.692	-53.703	-53.543
400	0.5	-30.509	-31.317	-31.329	-31.132
400	0.7	-15.524	-15.859	-15.864	-15.711
6400	0.3	-52.334	-52.484	-52.500	-52.396
6400	0.5	-31.140	-31.225	-31.238	-31.117
6400	0.7	-15.893	-15.933	-15.938	-15.853

Note. N = Sample size, SEM = Maximum-likelihood estimation, GSAM = Global-SAM-maximum-likelihood-estimation, LSAM-ML = Local-SAM-maximum-likelihood-estimation, LSAM-ULS = Local-SAM-unweighted-least-squares estimation.

Overall, GSAM, LSAM-ML, and LSAM-ULS demonstrated better empirical coverage than SEM but the overcoverage indicates too wide confidence intervals. Table 5 show the average difference between empirical coverage levels of the 95% confidence intervals (CIs) and their nominal level (95%) using each method for different reliability values and sample sizes under omitted cross loadings (model 1.2).

RMSE

For the correct model (1.1), SEM had higher RMSE values at smaller sample sizes and lower reliability, while GSAM, LSAM-ML, and LSAM-ULS demonstrated lower RMSE values. Under omitted cross loadings (1.2) and correlated residuals (1.3), SEM showed significantly higher RMSE values, although all methods exhibited increased RMSE under these conditions. The structural model (1.4) revealed that SEM had higher RMSE values at lower sample sizes and reliability, whereas GSAM, LSAM-ML, and LSAM-ULS showed better performance with lower RMSE. Overall, GSAM, LSAM-ML, and LSAM-ULS demonstrated more robustness with lower RMSE values compared to SEM, particularly in challenging conditions with smaller sample sizes and lower reliability.

Table 5

Study 1: Average coverage difference (in percentage) under omitted cross-loadings (Model 1.2)

N	Reliability	SEM	GSAM	LSAM-ML	LSAM-ULS
100	0.3	-1.008	3.671	3.650	3.451
100	0.5	-12.684	-2.277	-2.301	-3.112
100	0.7	-8.435	-4.447	-4.411	-5.164
400	0.3	-21.617	-6.320	-6.097	-6.914
400	0.5	-29.866	-16.801	-16.647	-17.457
400	0.7	-22.129	-15.936	-15.939	-16.359
6400	0.3	-65.231	-52.064	-51.036	-53.229
6400	0.5	-65.266	-54.583	-53.841	-55.944
6400	0.7	-55.541	-50.556	-50.237	-52.093

Note. N = Sample size, SEM = Maximum-likelihood estimation, GSAM = Global-SAM-maximum-likelihood-estimation, LSAM-ML = Local-SAM-maximum-likelihood-estimation, LSAM-ULS = Local-SAM-unweighted-least-squares estimation.

Improper Solutions

Improper solutions occur when parameter estimates fall outside the boundary of the parameter space, such as when variances are estimated to be negative or correlations exceed an absolute value of one. These issues are more likely to arise in smaller sample sizes and can indicate problems with model estimation.

SEM exhibited a significantly higher count of improper solutions compared to GSAM, LSAM-ML, and LSAM-ULS, particularly in small sample sizes and lower reliability conditions. GSAM, LSAM-ML, and LSAM-ULS consistently showed minimal improper solutions across all scenarios, highlighting their robustness and reliability in parameter estimation.

Study 2

Convergence

In Study 2, all methods achieved a 100% convergence rate for moderate and large sample sizes ($N = 400, 6400$). For small sample sizes ($N = 100$), SEM showed lower convergence rates, especially at lower reliability levels (0.3), while GSAM, LSAM-ML, and LSAM-ULS maintained high convergence rates.

R-squared values and measurement block size had minimal impact on convergence rates, indicating robustness of GSAM, LSAM-ML, and LSAM-ULS across various conditions. In contrast, SEM estimation struggled particularly in the conditions with lower reliability and smaller sample sizes.

The Influence of Number of Measurement Blocks on Bias and RMSE

For low reliability ($\lambda = 0.3$) and small sample size ($N = 100$), using 3 measurement blocks generally resulted in less negative bias compared to 5 blocks, although 5 blocks had lower RMSE. As reliability and sample sizes increased, the differences in bias and RMSE between 3 and 5 blocks diminished. Thus, for low reliability and small sample sizes, 3 measurement blocks were preferable for reducing bias, while 5 blocks performed better for RMSE.

For subsequent comparisons, we considered LSAM with 3 measurement blocks as this difference was more pronounced especially in low sample size and reliability conditions, which we were most interested in.

Average Relative Bias

For the correctly specified model with low reliability ($\lambda = 0.3$) and small sample size ($N = 100$), traditional SEM showed a larger average relative bias compared to SAM methods. GSAM and LSAM-ML had small negative biases, while LSAM-ULS had a positive bias in challenging conditions. Higher R^2 had little impact on SEM's bias as absolute values, but SAM methods showed increased bias. For higher reliability and sample sizes, biases decreased similarly for all estimation methods.

Under misspecification (correlated residuals and crossloadings in the exogenous model), SEM had an even greater positive bias in challenging conditions, while SAM methods showed consistent negative biases, with LSAM-ULS performing best. In the endogenous misspecification model, SEM exhibited relatively small biases compared to SAM methods, which showed substantial negative biases. SEM's bias got considerably worse with larger sample sizes and lower reliability, while SAM methods' biases remained similarly large and negative.

For misspecifications in both endogenous and exogenous parts, SAM methods had consistent negative biases (see Table 6). Here, SEM showed a positive bias with low reliability and sample size, turning negative with increasing sample size and higher R^2 values.

RMSE

For the correctly specified model with low reliability ($\lambda = 0.3$) and small sample size ($N = 100$), traditional SEM showed higher RMSE compared to SAM methods. GSAM and LSAM-ML had lower RMSEs, while LSAM-ULS had a slightly higher RMSE. Increasing regression weights (R^2) improved RMSE for both SEM and SAM methods.

Under exogenous misspecifications, SEM had higher RMSE compared to SAM methods at low reliability and small sample size. This pattern persisted for endogenous misspecifications and misspecifications in both exogenous and endogenous factors, as GSAM and LSAM-ML consistently showed lower RMSEs and LSAM-ULS performed slightly worse. RMSE improved for all methods with higher R^2 .

Improper Solutions

In conditions with small sample sizes ($N = 100$) and lower reliability ($\lambda = 0.3$), traditional SEM exhibited a high percentage of improper solutions. GSAM, LSAM-ML, and LSAM-ULS, on the other hand, consistently showed very low or zero improper solutions across all conditions investigated.

Table 6

Study 2: Mean relative bias in percentage under cross-loadings and residual correlations in Exo- and Endogenous factors of the model (Model 2.2-both).

N	Reliability	R2	SEM	GSAM	LSAM-ML	LSAM-ULS
100	0.3	0.1	9.39	-24.32	-16.81	-15.15
100	0.3	0.4	6.44	-25.21	-22.61	-21.45
100	0.5	0.1	5.56	-9.54	2.57	3.01
100	0.5	0.4	-4.46	-12.24	-10.20	-10.51
100	0.7	0.1	3.73	-1.85	6.55	6.26
100	0.7	0.4	-2.76	-5.81	-4.33	-5.09
400	0.3	0.1	-12.30	-22.13	-16.74	-17.62
400	0.3	0.4	-17.65	-22.40	-22.73	-23.34
400	0.5	0.1	1.59	-8.36	1.20	-0.23
400	0.5	0.4	-8.48	-11.26	-10.92	-12.13
400	0.7	0.1	2.93	-0.95	3.85	2.93
400	0.7	0.4	-3.40	-5.10	-4.56	-5.83
6400	0.3	0.1	-16.73	-21.19	-17.64	-18.85
6400	0.3	0.4	-20.17	-21.73	-23.11	-24.04
6400	0.5	0.1	-1.78	-8.08	-1.13	-2.76
6400	0.5	0.4	-9.66	-11.15	-11.28	-12.74
6400	0.7	0.1	2.17	-0.79	2.13	1.27
6400	0.7	0.4	-3.49	-5.00	-5.21	-6.97

Note. N = Sample size, R^2 = R^2 , SEM = Maximum-likelihood estimation, GSAM = Global-SAM-estimation, LSAM-ML = Local-SAM-maximum-likelihood-estimation, LSAM-ULS = Local-SAM-unweighted-least-squares estimation.

Individual study by Kosanke

Methods

The structure of this section closely aligns to our agreed upon structure of simulation studies in Table 1.

In a first step, I published a simulation protocol containing all the planned analysis to be replicated from the original paper by Robitzsch (2022). This protocol can be accessed here: https://github.com/lkosanke/AdversarialSimulation/blob/main/LK/simulation_protocol.pdf.

Aims, objectives and research questions

For my individual study, I replicated parts of Robitzsch (2022) that were relevant to our two substantive research questions. Overall, I conducted 6 simulation studies.

Population Models and Data Generation Mechanisms

The most important details with regards to the population models and data-generating mechanisms are visible in Table 7. With regards to the population models, all factors in all studies loaded onto 3 indicators each. I chose the population values to align to the original paper by Robitzsch (2022). The multivariate normally distributed data was generated parametrically, based on a specified population model. All simulations were conducted using seeds to allow for the reproducibility of results.

For more details on the exact values of each study, see the simulation scripts in the Github repository.

Table 7

Overview of the individual simulation studies by Leonard Kosanke.

Study	Model	Correct model included?	Unmodelled RC	Unmodelled CL	N Sizes	φ / β	λ
Study 1	2-factor-CFA	Yes	1 and 2, both pos. and neg.	x	7	$\varphi = 0.6$	Fixed
Study 1b	2-factor-CFA	Yes	x	x	2	$\varphi = 0.2 - 0.8$	Varied
Study 2	2-factor-CFA	x	x	1 and 2, both pos. and neg.	7	$\varphi = 0.6$	Fixed
Study 3	2-factor-CFA	x	1, pos.	1, pos.	7	$\varphi = 0.6$	Fixed
Study 4	5-factors	Yes	20, all pos.	5, all pos.	7	$\beta = 0.1$	Fixed
Study 4a	5-factors	x	20, all pos.	5, all pos.	7	$\beta = 0.1 - 0.4$	Fixed

Experimental Design of simulation procedures

Overall, 3 different types of factors were varied that can be deduced from Table 7 and are detailed again in the simulation scripts provided.

Firstly, I varied the sample size in all studies, ranging from $N = 50$ to 100.000. I included a smaller sample size $N=50$ for all studies, to be able to answer our substantive research questions in more detail. Study 1b explicitly investigated the small sample bias of LSAM estimation in low sample sizes. Thus, only $N=50$ and $N=100$ were present in this study.

Additionally, I varied the amount of misspecification in all studies, either via different numbers of unmodelled residual correlations, cross-loadings, or both.

Thirdly, in Studies 1b and 4a, I varied the population values for three model parameters (ϕ , β and/ or λ).

Besides studies 1 and 2, I implemented full factorial designs. In Studies 1 and 2 I omitted conditions where both one positive and one negative value would be present. I hypothesize that this was done in Robitzsch (2022) to avoid cancellation of biases, but the authors did not give reasoning for this decision themselves.

In Studies 4 and 4a I looked at the differential performance of the estimators in a model that included a non-saturated structural model (i.e. regressions between some of the factors). These studies were replications not only of the paper by Robitzsch (2022), but of the first paper on the SAM approach by Rosseel & Loh (2022). In contrast to the other studies, studies 4 and 4a differed in the way the misspecification variation was labelled in Robitzsch (2022). Instead of varying a factor misspecification as in the previous study, they varied 3 different data-generating mechanisms (DGM's) as a whole. Thus the conditions are labelled differently: DGM 1 contained no misspecification. DGM 2 contained 5 cross-loadings in the data-generating model, that were not modelled in the estimated models. DGM 3 contained 20 residual correlations that were not modelled in the models. I extended them to investigate the interaction of β and N for the 5-factor regression model, as this again was of interest for our substantial research questions. Additionally, I omitted the inclusion of DGM 1 in Study 4a, as it neither contained misspecification (which is central to our research question), nor did it lead to interesting results in the original study.

Method Selection

In terms of estimation methods, I used constrained SEM maximum-likelihood (SEM-ML) and unweighted-least-squares estimation (SEM-ULS), so that loadings and variance parameters were given the constraints that they had to be positive and larger than 0.01. Additionally, I implemented

local-SAM (LSAM) and global-SAM (GSAM) estimation, in both maximum-likelihood (LSAM-ML/ GSAM-ML), and unweighted-least-squares estimation (GSAM-ML/ GSAM-ULS) contexts. Exceptions were studies 1b, 4 and 4a, where only LSAM was investigated, as results did not really differ between the two different SAM-methods (Robitzsch, 2022).

Performance Measures

I calculated the bias and RMSE of the estimated factor correlations in all studies, as well as the standard deviation of the one factor correlation present in Studies 1,2 and 3. For the type of bias calculated, I oriented on Robitzsch (2022), besides in Study 1b. Thus, I calculated average relative bias in Studies 1, 2 and 3, and average absolute bias in Studies 1b, 4 and 4a. In Study 1b, I took the absolute value to see if negative and positive biases canceled each other out in the original study for conditions with lower phi values. In addition to what was done in Robitzsch (2022), I calculated confidence intervals for the bias estimates, but omitted them in the results tables for presentation purposes. The exact computation of the performance measures is detailed in the simulation scripts and results.pdf file in my sub-folder of the Github repository.

I did not include a detailed mechanism to capture model convergence as detailed in the first substantive research question. As Robitzsch (2022) argued in their paper, and was shown already in other simulations, using constrained maximum likelihood estimation should resolve convergence issues of classical maximum likelihood estimation in smaller samples (Lüdtke et al., 2021; Ulitzsch et al., 2023). I did include, however, a mechanism to track the total number of warnings for each estimation and compare it to the total number of estimations as a sanity check.

Software

All analyses were conducted in R (R Core Team, 2023). I used the packages lavaan, purrr, tidyverse, furrr to conduct the simulations, as well as knitr and kableExtra for presenting the results (Rosseel, 2012; Vaughan & Dancho, 2022; Wickham et al., 2019; Wickham & Henry, 2023; Xie, 2024; Zhu et al., 2024) .

Analysis and Interpretation plan

For the interpretation of results, I oriented on cut-offs that were used in the original paper by Robitzsch (2022). For bias, I interpreted differences of 0.05 or higher as substantial. For SD, I explicitly mentioned percentage reductions of more or equal to 5%. For RMSE, the same interpretation was used for differences of 0.03 or higher. The simulation was repeated 1500 times for each Study.

Results

The full result analysis for my individual study is available here: <https://github.com/lkosanke/AdversarialSimulation/blob/main/LK/results.pdf>. The repository readme.md contains a detailed explanation of how the analyses were implemented and how they can be reproduced. In this section, I will focus on the most important results only. For the most part, results from Robitzsch (2022) have been successfully replicated: I did not observe substantial convergence issues in any study. Across studies, as in the original paper, SAM did not generally outperform SEM in small to moderate samples. SAM exhibited a negative small sample bias that made SAM appear superior in conditions with unmodelled positive cross-loadings and residual correlations. This bias was especially strong for lower lambda and higher phi or beta values. Going ahead of what was investigated in Robitzsch (2022), I found that this bias is also present in models with lower phi or beta values. Thus, it cannot be concluded that SAM is more robust in models with non-saturated structural parameters. If there was no misspecification or unmodelled negative cross-loadings and residual correlations, SAM tended to perform worse than traditional SEM, as far as can be concluded from my results.

Convergence

As Robitzsch (2022) argued in their paper, I did not expect convergence issues due to constrained ML estimation that only allows for positive variances and loadings. Nevertheless, I captured all messages, warnings and errors that occurred during the simulations. No messages and errors were present in any of the studies. Multiple warnings were observed in the first 4 simulations, some of them referring to potential problems with convergence. Overall, the number of these warnings was very small compared to the total number of estimations performed. They amounted to between 0.5-1.8%. In studies 4 and 4a, an even smaller number of warnings was present, amounting to problems in 0.02% of estimations in study 4 and 0.1% in study 4a. These warnings referred to potential problems with positive definite matrices and model identification. In total, these numbers are negligible in size and align with the report of Robitzsch (2022), that convergence issues were not substantial for my estimations. Additionally, a larger number of warnings was present with regards to the computation of fit indices in these final two studies. As we were not interested in fit indices in our research question, they were not relevant for our purposes. A detailed analysis of all the warnings was conducted in the *results.pdf* document in my sub-folder of the Github repository.

Conditions without misspecification

Tables 8 and 9 show the most relevant results of Studies 1 and 4 were I investigated the comparative performance of SAM vs. traditional SEM estimation under correctly specified models. Here it became apparent, that in absence of misspecification, none of the two estimation methods clearly outperformed the other. In Study 4a, only slight differences could be observed in terms of bias and RMSE between LSAM- and classical ML-estimation. In Study 1, both SEM outperformed all SAM estimators in samples of N=50-500. This was true for both relative bias and RMSE, and visible for the former in Table 9. Here, SAM's negative small sample bias is already visible as well.

Table 8

Study 4 (Kosanke): RMSE for DGM 1 (without misspecification).

Method/Metric	Sample Size						
	50	100	250	500	1000	2500	100000
SEM_ML	0.188	0.123	0.075	0.051	0.037	0.023	0.004
SEM_ULS	1.062	0.128	0.077	0.053	0.037	0.023	0.004
LSAM_ML	0.165	0.115	0.072	0.050	0.036	0.023	0.004

Note. SEM_ML = Maximum-likelihood estimation, SEM_ULS = Unweighted-least-squares estimation, LSAM_ML= Local-SAM-maximum-likelihood estimation.

Table 9

Study 1 (Kosanke): Relative bias in conditions without misspecification.

Method/Metric	Sample Size						
	50	100	250	500	1000	2500	100000
SEM_ML	-0.045	-0.011	0.001	-0.003	0.003	-0.002	-0.000
SEM_ULS	0.024	0.022	0.012	0.002	0.006	-0.001	-0.000
LSAM_ML	-0.394	-0.270	-0.111	-0.056	-0.022	-0.011	-0.000
LSAM_ULS	-0.393	-0.270	-0.111	-0.056	-0.022	-0.011	-0.000
GSAM_ML	-0.394	-0.270	-0.111	-0.056	-0.022	-0.011	-0.000
GSAM_ULS	-0.393	-0.270	-0.111	-0.056	-0.022	-0.011	-0.000

Note. SEM_ML = Maximum-likelihood estimation, SEM_ULS = Unweighted-least-squares estimation, LSAM_ML = Local-SAM-maximum-likelihood estimation, LSAM_ULS = Local-SAM-unweighted-least-squares estimation, GSAM_ML = Global-SAM-maximum-likelihood estimation, GSAM_ULS = Global-SAM-unweighted-least-squares estimation.

Conditions with negatively valenced unmodelled parameters

Studies 1 and 2 explicitly investigated negatively valenced unmodelled parameters in the generating model. In these studies, it became apparent that traditional SEM outperformed SAM estimation.

As can be seen in Table 10, both SEM estimators outperformed all four SAM estimators in terms of relative bias with two negative residual correlations present. The same was true in Study 2, in the presence of two negative cross-loadings. In both these cases, bias values overall remained high but substantially less so in the traditional SEM methods. when comparing them in small to moderate sample sizes. Importantly, no differences between the two approaches arose in these two examples in terms of RMSE, as can be seen in Table 11 for the negative cross-loadings in study 2.

Conditions with positively valenced unmodelled parameters

In terms of performance for positively valenced cross-loadings and residual correlations, SAM appeared to slightly outperform traditional SEM estimation, but not in all scenarios. Table 12 shows this finding in Study 3, in conditions with both one unmodelled residual correlation and one cross-loading. Only from N=100-1000 did SAM outperform SEM.

Table 10

Study 1 (Kosanke): Relative bias in conditions with two negative unmodelled residual correlations.

Method/Metric	Sample Size						
	50	100	250	500	1000	2500	100000
SEM_ML	-0.205	-0.175	-0.166	-0.168	-0.161	-0.166	-0.164
SEM_ULS	-0.139	-0.145	-0.159	-0.167	-0.163	-0.170	-0.169
LSAM_ML	-0.498	-0.385	-0.272	-0.225	-0.196	-0.189	-0.180
LSAM_ULS	-0.497	-0.385	-0.272	-0.225	-0.196	-0.189	-0.180
GSAM_ML	-0.497	-0.385	-0.272	-0.225	-0.196	-0.189	-0.180
GSAM_ULS	-0.496	-0.385	-0.272	-0.225	-0.196	-0.189	-0.180

Note. SEM_ML = Maximum-likelihood estimation, SEM_ULS = Unweighted-least-squares estimation, LSAM_ML = Local-SAM-maximum-likelihood estimation, LSAM_ULS = Local-SAM-unweighted-least-squares estimation, GSAM_ML = Global-SAM-maximum-likelihood estimation, GSAM-ULS = Global-SAM-unweighted-least-squares estimation.

Table 11

Study 2 (Kosanke): RMSE in conditions with two negative unmodelled cross-loadings.

Method/Metric	Sample Size						
	50	100	250	500	1000	2500	100000
SEM_ML	0.480	0.382	0.257	0.211	0.182	0.172	0.161
SEM_ULS	0.477	0.367	0.241	0.205	0.182	0.175	0.166
LSAM_ML	0.486	0.421	0.323	0.269	0.232	0.216	0.201
LSAM_ULS	0.487	0.421	0.323	0.269	0.232	0.216	0.201
GSAM_ML	0.490	0.422	0.323	0.269	0.232	0.216	0.201
GSAM_ULS	0.490	0.421	0.323	0.269	0.232	0.216	0.201

Note. SEM_ML = Maximum-likelihood estimation, SEM_ULS = Unweighted-least-squares estimation, LSAM_ML = Local-SAM-maximum-likelihood estimation, LSAM_ULS = Local-SAM-unweighted-least-squares estimation, GSAM_ML = Global-SAM-maximum-likelihood estimation, GSAM-ULS = Global-SAM-unweighted-least-squares estimation.

Table 12

Study 3 (Kosanke): Relative bias in conditions with each one positive unmodelled cross-loading and residual correlation.

Method/Metric	Sample Size						
	50	100	250	500	1000	2500	100000
SEM_ML	0.209	0.270	0.283	0.289	0.289	0.282	0.284
SEM_ULS	0.250	0.284	0.277	0.280	0.279	0.271	0.272
LSAM_ML	-0.232	-0.061	0.127	0.211	0.246	0.261	0.276
LSAM_ULS	-0.229	-0.060	0.127	0.211	0.246	0.261	0.276
GSAM_ML	-0.230	-0.060	0.127	0.211	0.246	0.261	0.276
GSAM_ULS	-0.228	-0.060	0.127	0.211	0.246	0.261	0.276

Note. SEM_ML = Maximum-likelihood estimation, SEM_ULS = Unweighted-least-squares estimation, LSAM_ML = Local-SAM-maximum-likelihood estimation, LSAM_ULS = Local-SAM-unweighted-least-squares estimation, GSAM_ML = Global-SAM-maximum-likelihood estimation, GSAM-ULS = Global-SAM-unweighted-least-squares estimation.

In Study 4, a comparative advantage of LSAM compared to SEM-ML was present, but only for smaller samples.

With regards to RMSE, results were mixed as well. LSAM appeared to outperform in Table 13 for DGM 2 of Study 4. In other conditions, however, no substantial differences arose in terms of RMSE.

Small sample bias in LSAM estimation

The small sample bias of LSAM estimation in Study 1b revealed that in smaller samples ranging from N=50 to N=100, both LSAM-ML and LSAM-ULS estimation were biased. Table 14 shows this was especially apparent in a sample size of 50.

Table 13

Study 4 (Kosanke): RMSE in DGM 2 (conditions with five positive unmodelled cross-loadings).

Method/Metric	Sample Size						
	50	100	250	500	1000	2500	100000
SEM_ML	0.373	0.257	0.166	0.124	0.107	0.100	0.095
SEM_ULS	2.070	0.373	0.320	0.306	0.300	0.298	0.296
LSAM_ML	0.188	0.141	0.103	0.089	0.080	0.075	0.071

Note. SEM_ML = Maximum-likelihood estimation, SEM_ULS = Unweighted-least-squares estimation, LSAM_ML = Local-SAM-maximum-likelihood estimation.

Table 14

Study 1b (Kosanke): Absolute Bias of LSAM-ML for N=50.

Lambda	Phi values				
	phi=0	phi=0.2	phi=0.4	phi=0.6	phi=0.8
0.4	0.202	0.202	0.258	0.346	0.444
0.5	0.187	0.179	0.203	0.245	0.305
0.6	0.176	0.165	0.166	0.170	0.190
0.7	0.164	0.150	0.139	0.122	0.116
0.8	0.150	0.135	0.119	0.098	0.074

Note that the absolute values of bias were calculated in this study. Consequently, the values of the bias should be interpreted as negative, as follows from the results of the original paper (Robitzsch, 2022). The bias persisted, but to a lesser degree in samples of 100. Thus, as expected, a clear effect of sample size was present. Overall, comparing LSAM-ML and -ULS estimation, results were very similar.

Importantly, differential effects due to lambda and phi were present in Study 1b. The small sample bias was especially strong for lower lambda and higher phi values, thus in contexts of low reliability and high factor correlations. Also, a new insight is that the bias remained relevant for low values of phi, unlike in the original paper by Robitzsch (2022). In consequence, there seemed to be no conditions where SAM's small sample bias was negligible.

Another new insight lied in the presence of what could be called a reversal effect: For higher values of lambda, the bias did not increase for higher values of phi. On the contrary, absolute bias values decreased for higher phi values, when looking at the conditions with lambda = 0.7-0.8.

As an additional investigation of the small sample bias, I included Study 4a to see its effect come to play in a 5-factor-model with regressions. Table 15 shows the performance of SEM-ML, whereas Table 16 shows the performance of LSAM-ML in DGM 2 (in presence of unmodelled cross-loadings).

Aligning with the findings of Study 1b, the results suggested an even better relative performance of LSAM- over traditional SEM-ML estimation for smaller N and higher beta. Thus, the negative small sample bias came into play in this study as well. Results looked very similar with regards to RMSE. Note that this trend was less strong, but still present in the conditions of DGM 3 when looking at residual correlations.

Table 15

Study 4a (Kosanke): Absolute bias of SEM-ML for DGM 2 (conditions with five positive unmodelled cross-loadings).

beta	Sample size						
	50	100	250	500	1000	2500	100.000
0.1	0.253	0.172	0.115	0.095	0.085	0.080	0.075
0.2	0.327	0.220	0.162	0.139	0.122	0.114	0.109
0.3	0.545	0.270	0.218	0.206	0.195	0.190	0.180
0.4	0.981	0.336	0.270	0.257	0.252	0.251	0.254

Table 16

Study 4a (Kosanke): Absolute bias of LSAM-ML for DGM 2 (conditions with five positive unmodelled cross-loadings).

beta	Sample size						
	50	100	250	500	1000	2500	100.000
0.1	0.150	0.112	0.083	0.072	0.065	0.061	0.057
0.2	0.145	0.108	0.077	0.066	0.059	0.056	0.053
0.3	0.141	0.104	0.073	0.060	0.052	0.047	0.043
0.4	0.151	0.115	0.088	0.076	0.070	0.067	0.065

One aspect to mention is that lambda values were quite high in study 4a ($\lambda=0.7$). Matching the results from Study 1b, SAM's bias did not increase for higher values of beta, unlike SEM's. This effect could hint at a stronger robustness of SAM in contexts of higher correlations with misspecifications present. But, as the effect could not be observed in other conditions (e.g. in DGM 3 with residual correlations), I did not deem it substantial.

Summary of results

For the most part, I successfully replicated the results from Robitzsch (2022): I did not observe substantial convergence issues in any study. Across studies, as in the original paper, SAM did not generally outperform SEM in small to moderate samples. SAM exhibited a negative small sample bias that made SAM appear superior in conditions with unmodelled positive cross-loadings and residual correlations. This bias was especially strong for lower lambda and higher phi or beta values. Going ahead of what was investigated in Robitzsch (2022), I found that this bias is also present in models with lower phi or beta values. Thus, it cannot be concluded that SAM is more robust in models with non-saturated structural parameters. If there was no misspecification or unmodelled negative cross-loadings and residual correlations, SAM tended to perform worse than traditional SEM, as far as can be concluded from my results.

Conclusions from both studies

Comparing the result from both individual studies, it became evident that they are at no point contradictory or in disagreement. This is due to some central distinctions between the studies. Firstly, same as in Robitzsch (2022), the use of constrained ML estimation solves both the convergence

issues in small samples, as well as the positive small sample bias of the unconstrained version of the estimator. Additionally, the observed performance differences disappeared, when accounting for SAM's negative small sample bias.

Secondly, the results with regards to negatively valenced unmodelled residual correlations and cross-loading paint a more complete picture of the comparative performance of the two estimation methods. Unlike Kosanke, Kriegmair did not include these conditions. The apparent performance advantages for traditional SEM in these conditions balanced the advantages of SAM in positive conditions. In the context of empirical studies, the valence of potential residual correlations and cross-loadings is not known a priori. If that would be the case, one could simply turn to an estimator that is biased in the appropriate direction, to compensate this misspecification. Thus, achieving unbiased estimates is desirable in application contexts.

The evident performance differences observed between traditional SEM and SAM estimation in the study by Kriegmair are thus to be interpreted differently for all effects, but align with the results of Kosanke: This is true for all performance measures, even for coverage, which was not investigated in the study by Kosanke. Coverage is a measure that is partly determined by bias. Results in the studies by Kriegmair showed, that SAM had overcoverage due to its wider confidence intervals. Consequently, SAM was less biased but also less accurate than traditional SEM. Hence, SAM only partly outperformed traditional SEM, even in the unconstrained version.

Kriegmair's results also hinted at the presence of SAM's negative small sample bias, as the largest performance differences were observed in conditions of low reliability, i.e. λ , where the bias was especially pronounced (see Table 3). Moreover, in some conditions of Kriegmair's studies, SAM performed even worse than unconstrained SEM for conditions of low reliability and sample size (see Tables 4 and 6).

One set of findings, that hinted towards the possibility of SAM outperforming SEM was the slightly stronger robustness of SAM over constrained SEM for misspecifications along higher values of ϕ in Kosanke's Study 4a (see Tables 15-16). Only traditional SEM seemed to be negatively affected for higher beta values, compared to lower beta values in conditions with unmodelled cross-loadings (DGM 2). This showed slightly, but not substantially in conditions with residual correlations as well (DGM 3), in the same study by Kosanke. Kriegmair's study 2 showed this finding to not be consistent in the end, as in his conditions for model 2.2 in the endogenous conditions, for low sample size and reliability, the opposite trend was visible. Here, SEM performed better, even in the unconstrained version in the presence of both cross-loadings and residual correlations (see Table 6). All together, having in mind the differential effects for conditions of negative unmodelled parameters as well, we do not assume SAM to be consistently preferable in

this specific context.

Importantly, besides the use of coverage as a performance measure, other differences in the design and implementations were present when comparing our respective studies. These have to be addressed:

Unlike Kosanke, Kriegmair investigated additional effects with regards to endo- and exogeneity of predictors. Results showed, that the positive Bias of unconstrained SEM is stronger, if misspecification is present in exogenous predictors, if compared to it being present in endogenous and both endo- and exogenous predictors. In these latter conditions, SAM performed even worse. Interestingly, this finding suggests that the negative impacts of using unconstrained SEM-ML are limited to an even more specific set of model misspecifications, than initially suggested (Dhaene & Rosseel, 2023). Thus, unconstrained SEM can be viewed in an even better light relatively, for the endogenous as well as both endo- and exogenous predictor misspecifications. This does not warrant further investigation for our research question.

Furthermore, only Kriegmair investigated the differential effects of using a different number of measurement blocks in LSAM estimation. Results were mixed, as for Bias, using less measurement blocks appeared to lead to better performance, while the opposite was true in terms of RMSE. In consequence, no clear effect, that would suggest a differential performance of LSAM was present. We concluded that these findings have no relevance for our investigations on the relative comparison of SAM vs. SEM.

The difference of observing effects of regression weights directly in the study by Kosanke, and total variance explained by Kriegmair, are also minor, as both are aimed at modifying the regression weights and lead to comparable parameter values. For Kosanke's studies 4 and 4a, they were fixed between 0.1-0.4, while they varied between 0.183 and 0.447 in Kriegmair's study 2.

Another difference is the inclusion of conditions that examine misspecifications of the structural regression coefficient, both in terms of directionality (i.e. switching predictor and outcome variable incorrectly in the estimation model) and by omitting them in Kriegmair's studies. We want to highlight that we did not observe any clear differences depending on the type of misspecification looking across all studies, and Kriegmair's studies in specific. As the same was true even in the original papers for our estimators of interest, we saw no reason to suspect that results should be different if looking at constrained ML and negative misspecified parameter values (Dhaene & Rosseel, 2023; Robitzsch, 2022; Rosseel & Loh, 2022). Here, the same argument as before can be made: Only unconstrained ML-estimation and positive values of misspecified parameters were investigated. The observed effects should disappear when accounting for this as well as the negative small sample bias of SAM.

The results of the individual studies leave us with no imminent call for additional investigation in the joint study. What we and others viewed as a disagreement up to the point of contradiction, did not turn out to be one, upon closer examination. In consequence, we did not conduct a joint study and stopped the collaboration at this point.

Results of the collaboration

Going back to the outline of our collaboration framework, we successfully agreed on two research questions, conducted individual studies and integrated their results. Consequently, we did not find remaining disagreement or additional scenarios of interest that warranted the implementation of the second part of our AC procedure, the joint study. This means that we were unable to completely test our framework. Still, we have gained some insights with regards to our research question of whether adversarial collaboration can be applied to simulation studies, in terms of practical applicability and technical feasibility.

Practical applicability

The first step of agreeing on a joint research questions is already a collaborative effort that requires to change perspective, and to understand more deeply what the adversaries did in their previous studies. We see its main success in the focus on small to moderate sample sizes and on a limited number of estimators. This allowed us to focus on specific parts of our replications and led me, for example, to include an additional sample size in all my studies, as well as omit studies 5 and 6 that were conducted on the population level in the paper by Robitzsch (2022). For the same reasons, Kriegmair focussed on less correctly specified conditions than the original paper in his study 1 (Rosseel & Loh, 2022). Because of this our general focus became much narrow, as we did not include other discussions in the original papers, as for example on whether two-step-estimation is generally desirable as it separates the definition from the measurement of latent variables (Robitzsch, 2022).

When conducting the individual studies, we knew that we would have to be completely transparent with our results and had to give explanations for our decisions, as they would be scrutinized by the adversary later on. This increased quality of reporting, rigor and transparency in our individual studies. After the individual studies, we integrated our results and looked for disagreement and points for further investigation. We achieved a fair and truthful comparison of views and could agree that they were not opposing each other. We also believe that we increased generalizability simply by integrating the findings of two studies into one conclusion for more data generating scenarios and conditions than one individual study conducted.

With regards to the practical applicability of the collaboration of the second part, we want to emphasize that the reason for not conducting the second part was not that it was not practically applicable, but that there was no substantial reason to do it. Applying our framework, we observed the possibility to converge on research questions and found our individual simulations to be generally very compatible. This can be seen as preliminary evidence, that the second part of the collaboration should be practically applicable as well. Nevertheless, we were unable to conduct the second part of our collaboration framework.

Technical feasibility

Technical feasibility in the collaboration was achieved for the parts we could implement in our case study. Utilizing tools like Github with its functionalities to review code and jointly work on the same project allowed us to conduct simulations collaboratively in terms of comparing results, giving feedback and having access to the adversaries individual studies within the same Github repository.

Next to these aspects, which we integrated in our AC framework, two other aspect of technical feasibility were present for us while conducting our individual simulation studies. Firstly, we had access to a computing cluster. This allowed a higher limit on computational complexity, and thus more freedom to include additional conditions to test, without fearing too much runtime or having to limit the number of repetitions. Additionally, feasibility was facilitated by the use of the `furrr` package in R, which allowed for parallelisation of the code and significantly reduced the runtime of our simulations (Vaughan & Dancho, 2022). Again, as we could not conduct the second part of our collaboration framework, the technical feasibility of the collaborative study can not be judged conclusively.

Discussion

In this focused case study, we aimed to conduct adversarial collaboration to investigate the question if adversarial collaboration can be applied to simulation studies, in terms of practical applicability and technical feasibility. To do so, we created and tested a collaboration framework that starts with each adversary conducting an individual study, before collaborating in a joint study. For this, we agreed on two substantial research questions to judge the comparative performance of traditional SEM vs. SAM estimation in small to moderate samples and under misspecifications.

After finishing our individual studies, the supposed disagreement turned out not to be one, as results aligned well. Thus, we did not conduct a joint study and were only partly able to answer our research question.

For the first part of our framework, we found some evidence for practical applicability and technical feasibility: We were able to agree on two focused research question, that nudged our individual studies to focus on points of potential disagreement. Additionally, we were both able to successfully plan and conduct our individual simulations in a way that allowed integration of our results and to converge on a shared understanding of their consequences. Hence, we achieved a fair and truthful representation of both sides views. Technical feasibility was present with the use of Github for sharing and accessing our own and the adversaries results. Additionally, usage of a computing cluster and parallelisation allowed us to freely include conditions where we deemed it necessary, in our individual studies.

For the second part of our framework, it remains unclear whether collaboration would be applicable and technically feasible. Another attempt should be pursued to be able to answer our research question in the second part of our AC framework. Facilitating advancement in discussion between adversaries with the structured use of the tools we presented, we are confident that the necessary convergence for a joint study is achievable. This is underlined by the fact that there are some key differences between empirical and simulation studies: Firstly, opposing opinions in simulation studies are less incommensurable in terms of testing. In empirical studies, opposing theories might have completely different paradigms, measures and even construct definitions to answer the same research question (Rakow, 2022). In simulation studies, these differences are smaller. There is more consensus on what data generating scenarios of interest could be, how performance is measured, and what the opposing methods are. Another difference is the luxury of simulation studies to have much fewer resource restrictions. The main resources are time and computation power, which usually are not too limited. This allows adversaries to not have to find compromise in all cases, and, if in doubt, just include all conditions that might be relevant. In empirical studies, there are much more limited resources such as number of participants, money, labs and much more. In consequence, aspects of power have to always be considered and this leads to a need of more compromise in AC.

In terms of technical feasibility, we believe that using tools like Github with its functionalities to review code and jointly work on the same project, should allow to conduct simulations collaboratively. We see a lot of potential for these tools to be used in the second part of our collaboration framework, as they allow to split work within one study, give feedback on each others implementations via pull-requests, and much more. Ideally, additional resources like a computing cluster should be present to guarantee a higher limit on computational load, and thus more freedom to

include additional conditions to test. This, however, should not be a problem for research conducted in universities or institutes that usually have access to computing clusters. Still, besides these assumptions and learnings from our first case study, we were unable to test the main part of collaboration and can not conclude that the practical applicability and technical feasibility is completely given.

With regards to limitations there are several other aspects to consider. We are aware that we tested a very superficial “simulation” of adversariality. Neither side had long scientific experience in our substantial field of interest, and thus the stakes were much lower than as if the original authors would have collaborated. In that case, collaboration most likely would have been more difficult. Nevertheless, including the individual studies as a first step we think we succeeded priming ourselves to the views of our respective sides of the argument.

Another important aspect to discuss is whether we should have been able to see the lack of disagreement earlier, even before conducting our individual studies. As most of our findings were replications of the original papers, we might have been able to see their alignment if our preparation of the substantive topic would have been more in depth. To avoid this, collaborators should invest enough time and attention to detail into the evaluation of all the papers of interest before deciding on adversarial collaboration, to make sure that there is actually conflicting evidence present.

Another question is in how far the advantages we observed with regards to increased rigor, transparency and open access were incremental benefits of adversarial collaboration, over and above the usage of simulation protocols and our open access approach to science. We believe this not to be problematic, as AC and other aspects of open science can complement each very well and reinforce positive effects. Thus, researchers who are not generally interested in open science, but in collaboration, would still be nudged to work more openly, which we view as a positive outcome.

We believe that it would be worthwhile for future research to conduct another case study like ours. Science is, in the end, a collaborative effort. Exploring the limits of collaboration is thus valuable to learn more about opportunities to advance scientific progress.

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