Let's talk about Thurstone & Co.: An information-theoretical model

for comparative judgments, and its statistical translation

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Abstract

This study revisits Thurstone's law of comparative judgments (CJ) by addressing two key limitations

in traditional approaches. Firstly, it addresses the overreliance on the assumptions of Thurstone's

Case V in the statistical analysis of CJ data. Secondly, it addresses the apparent disconnect

between CJ's approach to trait measurement and hypothesis testing. We put forward a systematic

approach based on causal analysis and Bayesian statistical methods, which results in a model that

facilitates a more comprehensive understanding of the factors influencing CJ experiments while

offering a robust statistical translation. The new model accommodates unequal dispersions and

correlations between stimuli, enhancing the reliability and validity of CJ's trait estimation, thereby

ensuring the accurate measurement and interpretation of comparative data. The paper highlights

the relevance of this updated framework for modern empirical research, particularly in education

and social sciences. This contribution advances current research methodologies, providing a robust

foundation for future applications in diverse fields.

Keywords: causal inference, directed acyclic graphs, structural causal models, bayesian statistical

methods, thurstonian model, comparative judgement, probability, statistical modeling

1. Introduction

In comparative judgment (CJ) studies, judges assess a specific trait or attribute across different

stimuli by performing pairwise comparisons (Thurstone, 1927b,a). Each comparison produces a

dichotomous outcome, indicating which stimulus is perceived to have a higher trait level. For

example, when assessing writing quality, judges compare pairs of written texts (the stimuli) to

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Email addresses: JoseManuel.RiveraEspejo@uantwerpen.be (Jose Manuel Rivera Espejo), tine.vandaal@uantwerpen.be (Tine van Daal), sven.demaeyer@uantwerpen.be (Sven De Maeyer), determine the relative writing quality each text exhibit (the trait) (Laming, 2004; Pollitt, 2012b; Whitehouse, 2012; van Daal et al., 2016; Lesterhuis, 2018a; Coertjens et al., 2017; Goossens and De Maeyer, 2018; Bouwer et al., 2023).

Numerous studies have documented the effectiveness of CJ in assessing traits and competencies over the past decade. These studies have highlighted three aspects of the method's effectiveness: its reliability, validity, and practical applicability. Research on reliability suggests that CJ requires a relatively modest number of pairwise comparisons (Verhavert et al., 2019; Crompvoets et al., 2022) to generate trait scores that are as precise and consistent as those generated by other assessment methods (Coertjens et al., 2017; Goossens and De Maeyer, 2018; Bouwer et al., 2023). In addition, the evidence suggests that the reliability and time efficiency of CJ are comparable, if not superior, to those of other assessment methods when employing adaptive comparison algorithms (Pollitt, 2012b; Verhavert et al., 2022; Mikhailiuk et al., 2021). Meanwhile, research on the validity of CJ scores indicates their capacity to represent the traits under measurement accurately (Whitehouse, 2012; van Daal et al., 2016; Lesterhuis, 2018a; Bartholomew et al., 2018; Bouwer et al., 2023). Moreover, research on CJ's practical applicability highlights its versatility across both educational and noneducational contexts (Kimbell, 2012; Jones and Inglis, 2015; Bartholomew et al., 2018; Jones et al., 2019; Marshall et al., 2020; Bartholomew and Williams, 2020; Boonen et al., 2020).

Nevertheless, despite the increasing number of CJ studies, the prevalence of unsystematic and fragmented research approaches has left several critical issues unaddressed. The present study primarily focuses on two issues: the overreliance on Thurstone's Case V assumptions in the statistical analysis of CJ data and the apparent disconnect between CJ's approach to trait measurement and hypothesis testing. The following sections begin with a brief overview of Thurstone's theory followed by a detailed examination of these issues. Subsequently, the study introduces a theoretical model for CJ that builds upon Thurstone's theory, alongside its statistical translation, designed to address the two concerns simultaneously.

2. Thurstone's theory

In its most general form, Thurstone's theory addresses pairwise comparisons wherein a single judge evaluates multiple stimuli (Thurstone, 1927a, pp. 267). The theory posits that two key factors determine the dichotomous outcome of these comparisons: the discriminal process of each stimulus and their discriminal difference. The discriminal process captures the psychological impact each stimulus exerts on the judge or, more simply, his perception of the stimulus trait. The theory

assumes that the discriminal process for any given stimulus forms a Normal distribution along the trait continuum (Thurstone, 1927a, pp. 266). The mode (mean) of this distribution, known as the modal discriminal process, indicates the stimulus position on this continuum, while its dispersion, referred to as the discriminal dispersion, reflects variability in the perceived trait of the stimulus.

Figure 1a illustrates hypothetical discriminal processes along a quality trait continuum for two written texts. The figure indicates that the modal discriminal process for Text B is positioned further along the continuum than that of Text A $(T_B > T_A)$, suggesting that Text B exhibits higher quality. Additionally, the figure highlights that Text B has a broader distribution compared to Text A, which arises from its larger discriminal dispersion $(\sigma_B > \sigma_A)$.

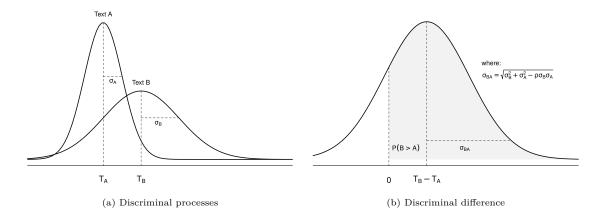


Figure 1: Hypothetical discriminal processes and discriminant difference along a quality trait continuum for two written texts.

However, given that the individual discriminal processes of the stimuli are not directly observable, the theory introduces the *law of comparative judgment*. This law posits that in pairwise comparisons, a judge perceives the stimulus with a discriminal process positioned further along the trait continuum as possessing more of the trait (Bramley, 2008, pp. 251). This suggests that the relative distance between stimuli, rather than their absolute positions on the continuum, likely defines the outcome of pairwise comparisons. Indeed, the theory assumes that the difference between the underlying discriminal processes of the stimuli, referred to as the *discriminal difference*, determines the observed dichotomous outcome. Furthermore, the theory assumes that because the individual discriminal processes form a Normal distribution on the continuum, the discriminal difference will also conform to a Normal distribution (Andrich, 1978). In this distribution, the mode (mean) represents the relative separation between the stimuli, and its dispersion indicates the variability of that separation.

Figure 1b illustrates the distribution of the discriminal difference for the hypothetical texts depicted in Figure 1a. The figure indicates that the judge perceives Text B as having significantly higher quality than Text A. This conclusion is supported by two key observations: the positive difference between their modal discriminal processes $(T_B - T_A > 0)$ and the probability area where the discriminal difference distinctly favors Text B over Text A, represented by the shaded gray area denoted as P(B > A). As a result, the dichotomous outcome of this comparison is more likely to favor Text B over Text A.

3. The two critical issues in CJ literature

This section examines the two critical issues in the CJ literature that serve as the primary focus of the present study. The first is related to the overreliance on Thurstone's Case V assumptions in the statistical analysis of CJ data. The second concern with the apparent disconnect between CJ's approach to trait measurement and hypothesis testing.

3.1. The Case V and the statistical analysis of CJ data

Thurstone noted from the outset that the general form of the theory, as outlined in Section 2, gave rise to a problem of trait scaling. The model required estimating more "unknown" parameters than the available pairwise comparisons (Thurstone, 1927a, pp. 267). To address this issue and facilitate the practical implementation of the theory, he developed five cases derived from this general form, each case progressively incorporated additional simplifying assumptions into the model.

In Case I, Thurstone postulated that pairs of stimuli would maintain a constant correlation across all comparisons. In Case II, he allowed multiple judges to undertake comparisons instead of confining evaluations to a single judge. In Case III, he posited that there was no correlation between stimuli. In Case IV, he assumed that the stimuli exhibited similar dispersions. Finally, in Case V, he replaced this assumption with the condition that stimuli had equal discriminal dispersions. Table 1 summarizes the assumptions of the general form and the five cases. For a detailed discussion of these cases and their progression, refer to Thurstone (1927a) and Bramley (2008, pp. 248–253).

Notably, despite relying on the most extensive set of simplifying assumptions (Bramley, 2008, pp. 253; Kelly et al., 2022, pp. 677), Case V remains the most widely used case in the CJ literature. This popularity stems mainly from its simplified statistical representation in the Bradley-Terry-Luce (BTL) model (Bradley and Terry, 1952; Luce, 1959). The BTL model mirrors the assumptions of Case V, with one notable distinction: whereas Case V assumes a Normal distribution for the

Table 1: Thurstones cases and their asumptions

	$\mathbf{General}$	ral Thurstone's					BTL
${f Assumption}$	\mathbf{form}	Case I	Case II	Case III	Case IV	Case V	\mathbf{model}
Discriminal process (distribution)	Normal	Normal	Normal	Normal	Normal	Normal	Logistic
Discriminal dispersion (between stimuli)	Different	Different	Different	Different	Similar	Equal	Equal
Correlation (between stimuli)	One per pair	Constant	Constant	Zero	Zero	Zero	Zero
How many judges compare?	Single	Single	Multiple	Multiple	Multiple	Multiple	Multiple

stimuli's discriminal processes, the BTL model uses the more mathematically tractable Logistic distribution (Andrich, 1978; Bramley, 2008, pp. 254) (see Table 1). This substitution has little impact on the model's estimation or interpretation, as the Normal and Logistic distributions exhibit analogous statistical properties, differing only by a scaling factor of approximately 1.7 (van der Linden, 2017a, pp. 16).

However, Thurstone originally developed Case V to provide a "rather coarse scaling" of traits (Thurstone, 1927a, pp. 269), prioritizing statistical simplicity over precision in trait measurement (Kelly et al., 2022, pp. 677). He explicitly warned against its untested application, stating that its use "should not be made without (an) experimental test" (Thurstone, 1927a, pp. 270). Furthermore, he acknowledged that some assumptions could prove problematic when researchers assess complex traits or heterogeneous stimuli (Thurstone, 1927b, pp. 376). Consequently, given that modern CJ applications frequently involve such traits and stimuli, two main assumptions of Case V and, by extension, of the BTL model may not consistently hold in theory or practice, namely the assumption of equal dispersion and zero correlation between stimuli.

3.1.1. The assumption of equal dispersions between stimuli

According to the theory, discrepancies in the discriminal dispersions of stimuli shape the distribution of the discriminal difference, exerting a direct influence on the outcome of pairwise comparisons. Figure 2a presents a thought experiment to illustrate this idea. In this experiment, a researcher can observe the discriminal processes for the texts depicted in Figure 1a. Furthermore, the figure assumes that the discriminal dispersion for Text A remains constant and that the texts are uncorrelated ($\rho = 0$). The figure reveals that an increase in the uncertainty associated with the perception of Text B in comparison to Text A, ($\sigma_B - \sigma_A$), broadens the distribution of their discriminal difference. This broadening affects the probability area where the discriminal difference distinctly favors Text B over Text A, expressed as P(B > A), ultimately influencing the compari-

son outcome. Additionally, the figure reveals that when the discriminal dispersions of the texts are equal $(\sigma_B - \sigma_A = 0)$, the discriminal difference is more likely to favor Text B over Text A (shaded gray area), compared to situations where their dispersions differ.

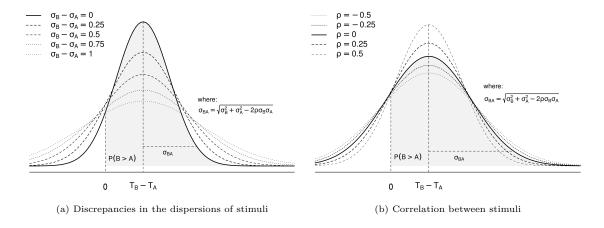


Figure 2: The effect of dispersion discrepancies and stimulus correlation on the distribution of the discriminal difference.

In experimental practice, however, this process occurs in reverse. Researchers first observe the comparison outcome and then use the BTL model to infer the discriminal difference between the stimuli and their respective discriminal processes (Thurstone, 1927b, pp. 373). Therefore, the outcome's ability to reflect the "true" differences between stimuli largely depends on the validity of the model's assumptions (Kohler et al., 2019, pp. 150), particularly the assumption of equal dispersions. For instance, when researchers observe a sample of outcomes favoring Text B over Text A and correctly assume equal dispersions between the texts, the BTL model estimates a discriminal difference distribution that accurately represents the "true" discriminal difference of the texts. This scenario is illustrated in Figure 2a, where the model's discriminal difference distribution aligns with the "true" distribution, represented by the thick continuous line corresponding to $\sigma_B - \sigma_A = 0$. The accuracy of these discriminal difference ensures reliable estimates for the texts' discriminal processes (citation needed?).

However, Thurstone argued that the assumption of equal dispersions may not be applicable when researchers assess complex traits or heterogeneous stimuli (Thurstone, 1927b, pp. 376), as these traits and stimuli can introduce judgment discrepancies due to their unique features (van Daal et al., 2016; Lesterhuis, 2018b; Chambers and Cunningham, 2022). Indeed, evidence of this violation may already be present in the CJ literature in the form of misfit statistics, which measure judgment discrepancies associated with specific stimuli (Pollitt, 2004, pp. 12; Goossens and De Maeyer, 2018,

pp. 20). For example, labeling texts as "misfits" indicates that comparisons involving these texts result in more judgment discrepancies than those involving other texts (Pollitt, 2012a,b; van Daal et al., 2016; Goossens and De Maeyer, 2018). These discrepancies, in turn, suggest that the discriminal differences for "misfit" texts have broader distributions, indicating that their discriminal processes may also exhibit more variation than that of other texts. A similar line of reasoning applies to the concept of "misfit" judges, whose evaluations deviate substantially from the shared consensus due to the unique characteristics of the stimuli or the judges themselves. These "misfit" judges and their associated deviations can give rise to additional statistical and measurement issues, which we discuss in more detail in Section 3.1.2.

Thus, model misspecification, in the form of an erroneous assumption of equal dispersions between stimuli, can give rise to significant statistical and measurement issues. For instance, the model may overestimate the degree to which the outcome accurately reflects the "true" discriminal differences between stimuli. This overestimation can result in researchers drawing spurious conclusions about these differences (McElreath, 2020, pp. 370) and, by extension, about the underlying discriminal processes of stimuli. Figure 2a also illustrates this issue when the model's discriminal difference distribution aligns with the thick continuous line for $\sigma_B - \sigma_A = 0$, while the "true" discriminal difference follows any discontinuous line where $\sigma_B - \sigma_A \neq 0$. Additionally, if researchers recognize that misfit statistics highlight these critical differences in dispersions, the conventional CJ practice of excluding stimuli based on these statistics (Pollitt, 2012a,b; van Daal et al., 2016; Goossens and De Maeyer, 2018) can unintentionally discard valuable information. Such exclusions can introduce bias into trait estimates (Zimmerman, 1994; McElreath, 2020, chap. 12). The direction and magnitude of these biases are often unpredictable, as they depend on which stimuli are excluded from the analysis.

3.1.2. The assumption of zero correlation between stimuli

The correlation, represented by the symbol ρ , measures how much a judge's perception of a specific trait in one stimulus depends on their perception of the same trait in another. As with the discriminal dispersions, this correlation shapes the distribution of the discriminal difference, directly impacting the outcomes of pairwise comparisons. Figure 2b presents a similar thought experiment as in Section 3.1.1 to illustrate this idea. The illustration now assumes that the discriminal dispersions for both texts remain constant. The figure reveals that as the correlation between the texts increases, the distribution of their discriminal difference becomes narrower. This narrowing affects the area under the curve where the discriminal difference distinctly favors Text B over Text A, denoted as

P(B > A), thus influencing the comparison outcome. Furthermore, the figure shows that when two texts are independent or uncorrelated ($\rho = 0$), their discriminal difference is less likely to favor Text B over Text A (shaded gray area) compared to scenarios where the texts are highly correlated.

Off course, in experimental practice, researchers approach this process in reverse. They begin by observing the sample of outcomes favoring Text B over Text A and then use the BTL model to estimate the discriminal difference and the discriminal processes of the stimuli. Given that the BTL model assumes independent discriminal processes across comparisons, if this assumption holds, then the model estimates a discriminal difference distribution that accurately reflects the "true" discriminal difference of the texts. This scenario is also illustrated in Figure 2b when the discriminal difference distribution of the model aligns with the "true" distribution, represented by the thick continuous line corresponding to $\rho = 0$. Once more, the estimation accuracy of the discriminal difference ensures reliable estimates for the discriminal processes of the texts (citation needed?).

Notably, Thurstone attributed the independence of stimuli to the cancellation of potential judges' biases. He argued that this cancellation resulted from two opposing and equally weighted effects occurring during pairwise comparisons (Thurstone, 1927a, pp. 268). Andrich (1978) provided a mathematical demonstration of this cancellation using the BTL model under the assumption of discriminal processes with additive biases. However, it is easy to imagine at least two scenarios in which the zero correlation assumption is almost certainly invalid: when the pairwise comparison involves multidimensional, complex traits with heterogeneous stimuli and when an additional hierarchical structure is relevant to the stimuli.

In the first scenario, the intricate aspects of multidimensional, complex traits may introduce dependencies between the stimuli due to certain judges' biases that resist cancellation. Research on text quality suggests that when judges evaluate these traits, they often rely on various intricate characteristics of the stimuli to form their judgments (van Daal et al., 2016; Lesterhuis, 2018b; Chambers and Cunningham, 2022). These additional relevant characteristics, which are unlikely to be equally weighted or opposing, can exert an uneven influence on judges' perceptions, creating biases in their judgments and, ultimately, introducing dependencies between stimuli (van der Linden, 2017b, pp. 346). For example, this could occur when a judge assessing the argumentative quality of a text places more weight on its grammatical accuracy than other judges, thereby favoring texts with fewer errors but weaker arguments. While direct evidence for this particular scenario is lacking,

studies such as Pollitt and Elliott (2003) demonstrate the presence of such biases, supporting the notion that the factors influencing pairwise comparisons may not always cancel out.

In the second scenario, the shared context or inherent connections created by additional hierarchical structures may further introduce dependencies between stimuli, a statistical phenomenon commonly known as clustering (Everitt and Skrondal, 2010). Despite the CJ literature acknowledging the existence of such hierarchical structures, the statistical handling of this additional source of dependence between stimuli has been inadequate. For instance, when CJ data incorporates multiple samples of stimuli from the same individuals, researchers frequently rely on (average) estimated BTL scores to conduct subsequent analyses and tests at the individual hierarchical level (Bramley and Vitello, 2019; Boonen et al., 2020; Bouwer et al., 2023; van Daal et al., 2017; Jones et al., 2019; Gijsen et al., 2021). However, this approach can introduce additional statistical and measurement issues, which we discuss in greater detail in Section 3.2.

In any case, similar to Section 3.1.1, model misspecification due to an erroneous assumption of zero correlation between stimuli can lead to significant statistical and measurement issues. For instance, the model may over- or underestimate how accurately the outcome reflects the "true" discriminal differences between stimuli. Such inaccuracies can result in spurious inferences about these differences and, by extension, about the stimuli's discriminal processes. This scenario is also illustrated by Figure 2b, when the model's discriminal difference distribution aligns with the thick continuous line for $\rho = 0$, while the "true" discriminal difference follows any discontinuous line where $\rho \neq 0$.

The misspecification may arise from neglecting additional relevant traits, excluding judges based on misfit statistics, or ignoring hierarchical (grouping) structures. Neglecting relevant traits, such as judges' biases, can cause dimensional mismatches in the BTL model, artificially inflating the trait's reliability (Hoyle, 2023, pp. 341) or, worse, introducing bias into the trait's estimates (Ackerman, 1989). Excluding judges based on misfit statistics risks discarding valuable information, which may further bias the trait's estimates (Zimmerman, 1994; McElreath, 2020, chap. 12). Finally, ignoring hierarchical structures may reduce the precision of model parameter estimates, potentially amplifying the overestimation of the trait's reliability (Hoyle, 2023, pp. 482).

3.2. The disconnect between trait measurement and hypothesis testing

Building on the previous section, it is clear that, despite its limitations, the BTL model is commonly used as a measurement model in CJ assessments. A measurement model specifies how

manifest variables contribute to the estimation of latent variables (Everitt and Skrondal, 2010). For example, when evaluating writing quality, researchers use the BTL model to process the dichotomous outcomes resulting from the pairwise comparisons (the manifest variables) to estimate scores that reflect the underlying level of writing quality (the latent variable) (Laming, 2004; Pollitt, 2012b; Whitehouse, 2012; van Daal et al., 2016; Lesterhuis, 2018a; Coertjens et al., 2017; Goossens and De Maeyer, 2018; Bouwer et al., 2023).

Researchers then typically use these estimated BTL scores, or their transformations, to conduct additional analyses or hypothesis tests. For example, these scores have been used to identify 'misfit' judges and stimuli (Pollitt, 2012b; van Daal et al., 2016; Goossens and De Maeyer, 2018), detect biases in judges' ratings (Pollitt and Elliott, 2003; Pollitt, 2012b), calculate correlations with other assessment methods (Goossens and De Maeyer, 2018; Bouwer et al., 2023), or test hypotheses related to the underlying trait of interest (Bramley and Vitello, 2019; Boonen et al., 2020; Bouwer et al., 2023; van Daal et al., 2017; Jones et al., 2019; Gijsen et al., 2021).

However, the statistical literature advises caution when using estimated scores for additional analyses and tests. A key consideration is that BTL scores are parameter estimates that inherently carry uncertainty. Ignoring this uncertainty can bias the analysis and reduce the precision of hypothesis tests. Notably, the direction and magnitude of such biases are often unpredictable. Results may be attenuated, exaggerated, or remain unaffected depending on the degree of uncertainty in the scores and the actual effects being tested (Kline, 2023, pp. 25; Hoyle, 2023, pp. 137). Finally, the reduced precision in hypothesis tests diminishes their statistical power, increasing the likelihood of committing type-I or type-II errors (McElreath, 2020).

In aggregate, researchers' inadequate handling of violations to the assumptions of equal dispersion and zero correlation between stimuli, coupled with the apparent disconnect between CJ's approach to trait measurement and hypothesis testing, can potentially compromise the reliability of the trait estimates and, by extension, their validity (Perron and Gillespie, 2015, pp. 2). Consequently, adopting a more systematic and integrated approach to handling these assumptions and examining the factors influencing CJ experiments could offer several statistical and measurement benefits, including the ability to address these issues.

4. Updating CJ's theoretical and statistical model

This section presents a theoretical model for CJ that extends Thurstone's theory. It uses causal inference and, in particular, the structural approach where structural causal models (SCMs) and directed acyclic graphs (DAGs) articulate the core theoretical principles of CJ theory. The model also incorporates several assessment design features that influence CJ experiments, such as the selection of judges, stimuli, and comparisons. In addition, the study uses Bayesian statistical methods to translate these theoretical and practical elements into a statistical model that facilitates the analysis of pairwise comparison data.

4.1. The theoretical model

4.1.1. The population model

structural causal models (SCMs) and directed acyclic graphs (DAGs) (Pearl, 2009; Pearl et al., 2016; Gross et al., 2018; Neal, 2020)

causal analysis (Pearl, 2009; Pearl et al., 2016; Morgan and Winship, 2014)

Assuming population data or more commonly known as census data, we ...

The (latent) discriminal difference of the stimuli directly determines the (manifest) outcome of the pairwise comparisons

The (latent) "perceived" discriminal processes for the stimuli directly determines their discriminal difference

The (latent) "true" discriminal processes for the stimuli and the judges' biases directly determines their (latent) "perceived" discriminal processes

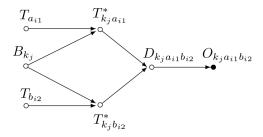


Figure 3

without loosing generality, the (latent) "perceived" and "true" discriminal processes for the stimuli can be depicted in a vector for each judge, as in

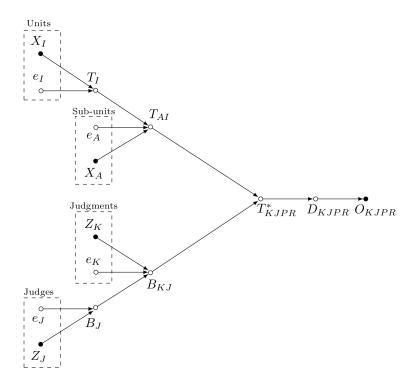


Figure 4

 ${\it 4.1.2. \ The \ assessment \ design \ features}$

Considering the sampling mechanism $\,$

 ${\bf Considering\ comparison\ mechanisms}$

- 4.2. From theory to statistics
- 5. Discussion
- 5.1. Findings
- 5.2. Limitations and further research
- 6. Conclusion

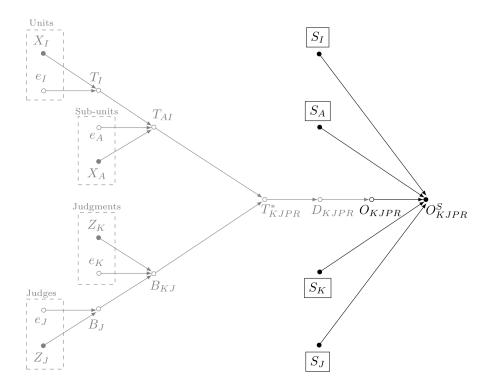


Figure 5

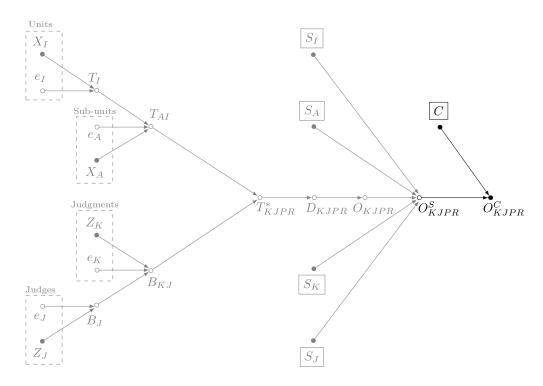


Figure 6

Declarations

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Code availability: All the code utilized in this research is available in the digital document located at: https://jriveraespejo.github.io/paper2_manuscript/.

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7. Appendix

This section introduces fundamental statistical and causal inference concepts necessary for understanding the core theoretical principles described in Section 4. It does not, however, offer a comprehensive overview of statistical and causal inference methods. Readers seeking more in-depth understanding may wish to explore introductory papers such as Pearl (2010), Rohrer (2018), Pearl (2019), and Cinelli et al. (2020). They may also find it helpful to consult introductory books like Pearl and Mackenzie (2018), Neal (2020), and McElreath (2020). For more advanced study, readers may refer to seminal intermediate papers such as Neyman (1923), Rubin (1974), Spirtes et al. (1991), and Sekhon (2009), as well as books such as Pearl (2009), Morgan and Winship (2014), and Hernán and Robins (2020).

7.1. Empirical research and randomized experiments

Empirical research uses evidence from observation and experimentation to address real-world challenges. In this context, researchers typically formulate their research questions as estimands or targets of inference, i.e., the specific quantities they seek to determine (Everitt and Skrondal, 2010). For instance, researchers might be interested in answering the following question: "To what extent do different teaching methods (T) influence students' ability to produce high-quality written texts (Y)?" To investigate this, researchers could randomly assign students to two groups, each exposed to a different teaching method $(T_i = \{1,2\})$. Then, they would perform pairwise comparisons, generating a dichotomous outcome $(Y_i = \{0,1\})$ showing which student exhibits more of the ability. In this scenario, the research question can be rephrased as the estimand, "On average, is there a difference in the ability to produce high-quality written texts between the two groups of students?" and this estimand can be mathematically represented by the random quantity $E[Y_i|T_i=1]-E[Y_i|T_i=2]$, where $E[\cdot]$ denotes the expected value.

Researchers would then proceed to identify the estimands. *Identification* refers to the process of accurately computing an estimand using an estimator. An *estimator* is a method or function that transforms data into an estimate (Neal, 2020). *Estimates* are numerical values that approximate the estimand and are derived through *estimation*, which refers to the process of integrating data with an estimator (Everitt and Skrondal, 2010). The Identification-Estimation flowchart (McElreath, 2020; Neal, 2020) in Figure 7 provides a visual representation of the process of transitioning from estimands to estimates.

While numerous methods can approximate an estimand, researchers prioritize estimators with

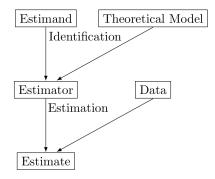


Figure 7: Identification-Estimation flowchart. Extracted and slightly modified from Neal (2020, pp. 32)

desirable properties that ensure the accuracy of estimates. For instance, the Z-test is an estimator known for its effectiveness in comparing groups' proportions, yielding accurate estimates when the underlying assumptions of the statistic are met (Kanji, 2006). If this is the case, the Z-test is expressed as a signal-to-noise statistic $Z=(\hat{p}_1-\hat{p}_2)/\hat{s}_p$. The signal is defined as the difference between the groups' sample proportions, $\hat{p}_1=\sum_{i=1}^{n_1}Y_i/n_1$ and $\hat{p}_2=\sum_{i=1}^{n_2}Y_i/n_2$, analogous to $E[Y_i|T_i=1]$ and $E[Y_i|T_i=2]$, respectively. The noise, represented by \hat{s}_p , is defined as the unpooled sample variability observed between the two groups.

However, researchers often seek to uncover the mechanisms underlying specific data and establish causal relationships rather than simply identify associations. In the example, researchers can interpret the associational estimate represented by the Z-statistic as causal. This interpretation relies on the data meeting the assumptions of the Z-test and the data being collected through a randomized experiment.

Randomized experiments are widely recognized as the gold standard in evidence-based science (Hariton and Locascio, 2018; Hansson, 2014). This recognition stems from their ability to enable researchers to interpret associational estimates as causal. They achieve this by ensuring data, and by extension an estimator, satisfies several key properties, such as common support, no interference, and consistency (Morgan and Winship, 2014; Neal, 2020). The most critical property, however, is the elimination of confounding. Confounding occurs when an external variable X simultaneously influences the outcome Y and the variable of interest T, resulting in spurious associations (Everitt and Skrondal, 2010). Randomization addresses this issue by decoupling the association between the intervention allocation T from other variables X and the outcome Y (Morgan and Winship, 2014; Neal, 2020).

Nevertheless, researchers often face constraints that limit their ability to conduct randomized

experiments. These constraints include ethical concerns, such as the assignment of individuals to potentially harmful interventions, and practical limitations, such as the infeasibility of, for example, assigning individuals to genetic modifications or physical impairments (Neal, 2020). In these cases, causal inference offers a valuable alternative for generating causal estimates and understanding the mechanisms underlying specific data. In addition, the framework can provide significant theoretical insights that can enhance the design of experimental and observational studies (McElreath, 2020).

7.2. Empirical research under causal inference

Unlike classical statistical modeling, which focuses primarily on summarizing data and inferring associations, the *causal inference* framework is designed to identify causes and estimate their effects using data (Shaughnessy et al., 2010; Neal, 2020). The framework uses rigorous mathematical techniques to address the *fundamental problem of causality* (Pearl, 2009; Pearl et al., 2016; Morgan and Winship, 2014). This problem revolves around the question, "What would have happened 'in the world' under different circumstances?" This question introduces the concept of counterfactuals, which are instrumental in understanding and defining causal effects.

Counterfactuals represent hypothetical scenarios that are contrary to fact, where alternative outcomes resulting from a given cause are neither observed nor observable (Neal, 2020; Counterfactual, 2024). Although a comprehensive discussion of causes and counterfactuals is beyond the scope of this document, it is possible to provide a brief overview of how the framework addresses the fundamental problem of causality. Using the example outlined in Section 7.2, the framework begins by defining the individual causal effect (ICE) as the difference between the students' potential outcomes: $\tau_i = Y_i^1 - Y_i^2$. Here, Y_i^1 represents the potential outcome observed under $T_i = 1$, while Y_i^2 represents the potential outcome observed under $T_i = 2$. Note that when a student is assigned to $T_i = 1$, the potential outcome under $T_i = 2$ is no longer observed nor observable, making it a counterfactual. To address the challenge posed by these counterfactuals, the framework extends the ICE to the average causal effect (ACE). Researchers define the ACE as $\tau = E[\tau_i] = E[Y_i^1] - E[Y_i^2]$, which represents the difference between the average of observed potential outcomes and counterfactuals.

Even when data originates from an observational study, researchers can identify the ACE from associational estimates through causal inference. They achieve this by performing statistical conditioning on a *sufficient adjustment set* of variables X (Pearl, 2009; Pearl et al., 2016; Morgan and Winship, 2014). This *sufficient* set (potentially an empty set) should block all (and not open any) non-causal paths between T to Y, thereby ensuring that the estimator for the ACE satisfies several

key properties, including eliminating confounding. If such a set exists, researchers can express the ACE from associational random quantities as $\tau = E_X \left[E[Y_i | T_i = 1, X] - E[Y_i | T_i = 2, X] \right]$, where $E_X[\cdot]$ denotes the marginal expected value over X (Morgan and Winship, 2014). Of course, the validity of the claims about the effects of T on Y now hinges on the assumption that X serves as a sufficient adjustment set. However, as Kohler et al. (2019, pp. 150) noted, "statements about the real world from observed (not observational) data require assumptions (and) this is true no matter what kind of data we have". For example, if researchers cannot conduct the randomized experiments described in Section 7.1 and must rely on observational data, they can still estimate τ if a variable X, such as the socio-economic status of the school, blocks all (and not open any) non-causal paths from the teaching method T to the outcome Y.

Several approaches to causal inference and counterfactuals exist, but two are particularly prominent: the potential outcomes approach, also known as the Neyman-Rubin causal model (Neyman, 1923; Rubin, 1974), and the structural approach (Pearl, 2009; Pearl et al., 2016). Both approaches use rigorous mathematical notation to characterize the ACE, not only for dichotomous but also for continuous variables T, though they do so in different ways (Neal, 2020). The potential outcomes approach relies on counterfactual notation, as seen in the above examples, whereas the structural approach uses do-calculus (Pearl, 2009). Despite these differences, both notations can be expressed in terms of the other, and both approaches provide methods for using experimental and observational data to identify and estimate causal effects (Pearl, 2010).

7.3. SCMs and DAGs

The structural approach to causal inference, however, offers a notable advantage over the potential outcomes approach. It enables researchers to formally and graphically represent a system using SCMs and DAGs (Pearl, 2009; Pearl et al., 2016; Gross et al., 2018; Neal, 2020). SCMs and DAGs function as conceptual models describing the presumed causal structure underlying the ACE. In essence, these models act as theoretical models that guide researchers in determining which statistical models can yield valid causal inferences, assuming the depicted causal structure of the models is correct (McElreath, 2020). Notably, every SCM has a associated DAG (Cinelli et al., 2020). Figure 7 provides a visual representation of the role of theoretical models in the inference process.

SCMs and DAGs offer three key advantages for modeling causal structures. First, they enable researchers to represent causal relations in a non-parametric and fully interactive manner. This fea-

ture allows for feasible ACE identification strategies without specifying the data type or the nature of the functional dependence among variables (Morgan and Winship, 2014). Second, regardless of complexity, they can represent a wide range of causal structures using only five fundamental building blocks (Neal, 2020; McElreath, 2020). This feature allows researchers to decompose complex structures, facilitating their analysis by focusing on the causal assumptions associated with each building block (McElreath, 2020). Finally, they help to formalize (causal) theoretical models into (probabilistic) statistical models for identification purposes (McElreath, 2020) (McElreath, 2020).

Figures 8, 9, 10, 11, and 12 display the five fundamental building blocks of SCMs and DAGs. The left panels of the figures show the formal mathematical models, represented by the SCMs, defined in terms of a set of endogenous variables $V = \{X, Z, Y\}$, a set of exogenous variables $E = \{e_X, e_Z, e_Y\}$, and a set of functions $F = \{f_X, f_Z, f_Y\}$ (Pearl, 2009; Cinelli et al., 2020; Neal, 2020). Endogenous variables are those whose causal mechanisms a researchers chooses to model (Neal, 2020). In contrast, exogenous variables represent errors or disturbances arising from omitted factors that the investigator chooses not to model explicitly (Pearl, 2009, pp. 27,68). Lastly, the functions, referred to as structural equations, express the endogenous variables as non-parametric functions of other variables. These functions use the symbol ':=' to denote the asymmetrical causal dependence of the variables and the symbol ' \perp ' to represent d-separation, a concept akin to (conditional) independence.

The right panels of the figures display the complementary DAGs. A DAG consists of nodes connected by edges, where nodes represent random variables. The term directed means the edges extend from one node to another, with arrows indicating the direction of causal influence. The term acyclic signifies that the causal influences do not form loops, ensuring the influences do not cycle back on themselves (McElreath, 2020). DAGs represent observed variables as solid black circles, while they use open circles for unobserved (latent) variables (Morgan and Winship, 2014). Finally, the arrows in the graphs show the expected direction of causal influences among these variables. Often, DAGs omit the exogenous variables for simplicity (the standard representation). However, including exogenous variables in a graph (the magnified representation) can be beneficial, as their presence can reveal potential issues related to conditioning and confounding (Cinelli et al., 2020).

$$X:=f_X(e_X)$$

$$Y:=f_Y(e_Y)$$

$$\stackrel{e_X}{\circ} X \qquad Y \qquad \stackrel{e}{\circ}$$

$$e_X \perp\!\!\!\perp e_Y \qquad \qquad \text{(b) DAG}$$

Figure 8: Two unconnected nodes

$$X:=f_X(e_X)$$

$$Y:=f_Y(X,e_Y)$$

$$e_X \perp \!\!\! \perp e_Y$$
 (b) DAG
$$(a) \text{ SCM}$$

Figure 9: Two connected nodes or descendant

$$X := f_X(e_X)$$

$$Z := f_Z(X, e_Z)$$

$$Y := f_Y(Z, e_Y)$$

$$e_X \perp \!\!\!\perp e_Y$$

$$e_X \perp \!\!\!\perp e_Z$$
 (b) DAG
$$e_Z \perp \!\!\!\perp e_Y$$
 (a) SCM

Figure 10: Chain or mediator

$$X := f_X(Z, e_X)$$

$$Z := f_Z(e_Z)$$

$$Y := f_Y(Z, e_Y)$$

$$e_X \perp \!\!\!\perp e_Y$$

$$e_X \perp \!\!\!\perp e_Z$$

$$e_Z \perp \!\!\!\perp e_Y$$
(b) DAG

Figure 11: Fork or confounder

$$X := f_X(e_X)$$

$$Z := f_Z(X, Y, e_Z)$$

$$Y := f_Y(e_Y)$$

$$e_X \perp \!\!\!\perp e_Y$$

$$e_X \perp \!\!\!\perp e_Z$$
 (b) DAG
$$e_Z \perp \!\!\!\perp e_Y$$
 (a) SCM

Figure 12: Collider or inmorality

A careful examination of the figures highlights the assumptions underlying these building blocks. Figures 8a and 8b depict two unconnected nodes, representing a scenario where variables X and Y are not causally related. Figures 9a and 9b illustrate two connected nodes, representing a scenario where a parent node X exerts a causal influence on a child node Y. Consequently, Y is considered a descendant of X. Figures 10a and 10b depict a chain, where X influences X, and X influences X. In this configuration, X is a parent node of X, which is a parent node of X. This structure creates a directed path between X and Y. Consequently, X is an ancestor of Y, and X fully mediates the relationship between the two. Figures 11a and 11b illustrate a fork, where variables X and Y are both influenced by X. Here, X is a parent node that confounds the relationship between X and Y. Finally, Figures 12a and 12b show a collider, where variables X and Y are concurrent causes

of Z. In this configuration, X and Y are not causally related to each other but both influence Z (an *inmorality*). Notably, in all SCMs, the errors are assumed to be independent of each other and from all other variables in the graph, as evidenced by the pairwise relations $e_X \perp\!\!\!\perp e_Y$, $e_X \perp\!\!\!\perp e_Z$, and $e_Z \perp\!\!\!\perp e_Y$.

Researchers can use these building blocks to depict the scenario described in Section 7.2. In this scenario, they rely on observational data because it is not feasible to run a randomized experiment. Furthermore, the scenario indicates that researchers have measured a variable X (the socio-economic status of the school) that is assumed to block all non-causal paths from the teaching method T to the outcome Y. Figures 13a and 13b illustrate the plausible causal structure for this example. The figures highlight the presence of at least four of the five fundamental building blocks. The figures display multiple descendants, as shown by pairwise relationships such as $X \to T$, $X \to Y$, and $T \to Y$. They also feature multiple pairs of unconnected nodes, evident from the relationships $e_T \perp \!\!\!\perp e_X$, $e_T \perp \!\!\!\perp e_Y$, and $e_X \perp \!\!\!\perp e_Y$. Finally, the figures illustrate one fork, $X \to \{T,Y\}$, and two colliders: $\{X,e_T\} \to T$ and $\{X,T,e_Y\} \to Y$.

$$X := f_X(e_X)$$

$$T := f_Z(X, e_T)$$

$$Y := f_Y(T, X, e_Y)$$

$$e_T \perp \!\!\!\perp e_X$$

$$e_T \perp \!\!\!\perp e_Y$$
 (b) DAG
$$e_X \perp \!\!\!\perp e_Y$$

Figure 13: Plausible causal structure of the example.

7.4. The flow of association and causation in DAGs



Figure 14: The flow of association and causation in graphs. Extracted and slightly modified from Neal (2020, pp. 31)

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