

PROPOSAL

Bayesian Evidence Synthesis

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In recent years, a meta-analytic way of thinking has been advocated in the scientific community (Cumming, 2014). Additionally, the importance of replication has been legitimately supported (e.g., Open Science Collaboration, 2015; Baker, 2016; Brandt et al., 2014). However, most of the attention has been focused on studies that are highly similar, using an identical methodology and research design. These studies, commonly referred to as exact, direct or close replications, are merely concerned with the statistical reliability of the results. Unfortunately, if the results of these studies depend on methodological flaws, inferences from all studies will lead to suboptimal or invalid conclusions (Munafò & Smith, 2018). A safeguard against this deficiency is available in the form of conceptual replications, which primarily assess the validity of a study. That is, conceptual replications are a way of investigating whether the initial conclusions hold under different conditions, using varying measurement instruments or choosing different operationalizations. Voor mijn gevoel kan de relevantie nog verder benadrukt worden door te zeggen dat onderzoekers ook onbewust verschillende (te rechtvaardigen) keuzes maken mbt de analyse (*garden of forking paths*), maar ik vind dat ook weer niet zo goed passen bij de rest van dit stuk.

As a consequence, multiple studies regarding the same hypotheses arise and as per the cumulative nature of science, synthesizing the results is required to build a robust and solid body of evidence. When the studies are highly similar, established methods as (Bayesian) meta-analysis and Bayesian updating can be used to pool the results (Glasauer, 2019; Lipsey & Wilson, 2001; Sutton & Abrams, 2001) Heel vreemd, maar ik kan echt zo goed als niks over Bayesian Sequential Updating vinden. Ondanks dat het in deze bron (Glasauer) wel oke uitgelegd wordt, is het niet echt ons interessegebied geloof ik.

1 Introduction

2 References

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