

Estimating repeatability and reproducibility with limited replications

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Tuesday 4th August, 2020

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Background

- The justice system relies heavily on forensic science.
- However, innocent people have been held or imprisoned because of incorrect application of forensic science. This is a serious problem.

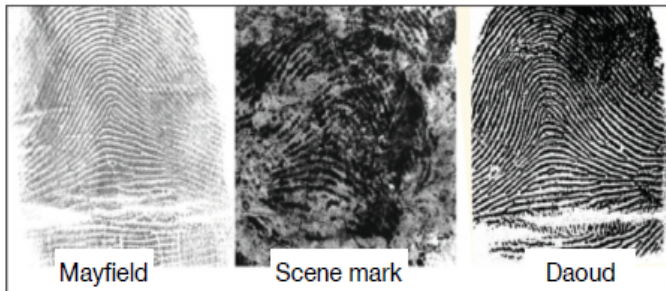


Figure: Brandon Mayfield's fingerprint compared to the mark found at the scene.

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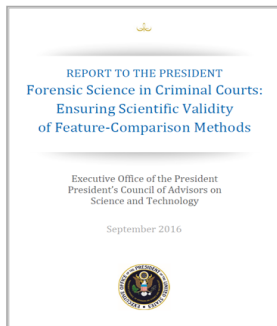
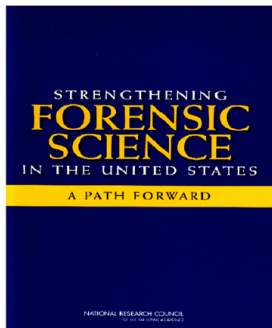
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Background



- National Academy of Sciences (NAS) prepared a report in 2009 emphasizing the need for formal assessment of scientific foundations of forensic science.
- In 2016, President's Council of Advisors on Science and Technology (PCAST) prepared a report assessing progress and highlighting the need to ensure validity and reliability of feature-based comparison methods.

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Important definitions

- Reliability: Related to the consistency of a measure or a procedure. Two types of reliability:
 - Repeatability: Consistency of repeated measurements obtained from the same examiner under the same conditions. Intra-examiner reliability.
 - Reproducibility: Consistency of measurements obtained from different examiners under the same conditions. Inter-examiner reliability.
- Validity: Related to the correctness (accuracy) of the procedure.

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- An important goal is to establish reliability and validity for feature-based comparison methods through empirical studies.
- For objective measurement methods, it is easier to establish reliability and validity.
- For subjective feature-based comparison methods, “black-box” studies.
- The decision process is subjective and treated like a black-box.

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Reliability of fingerprint evidence

- In late 2000's, the FBI conducted a study to establish reliability of latent fingerprint examination decisions (Ulery et al., 2011, 2012).
- 169 examiners, 744 latent prints and corresponding exemplars (mated and non-mated), ≈ 100 pairs per examiner.
- Prints selected to contain features that mimic real cases.
- About 7 months after the first study, a subset of prints were re-examined by a subset of these examiners.
- Studies with large repeatability components are rare because they are expensive and use precious examiner time. Hence, this incomplete design is common in many forensic studies.

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Reliability of signature complexity decisions

- Signature complexity is an important factor for an examiner to differentiate between a real signature and a simulation.
- Data collected by LAPD/ LASD: 123 signatures, 5 forensic document examiners (FDEs). Repetitions on $n = 7$ signatures. Signature complexity assessed on two different scales (a 1-5 point scale and a 1-3 point scale).
- Stern et al. (2018) analyzed reliability for handwritten signature complexity scores based on this study.
- 1-5 point scale data was treated as continuous. Also the data collected in the repeatability study was analyzed separately.

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- Develop methods that allow for assessing reliability that:
 - Combine data from reproducibility studies and (incomplete) repeatability studies.
 - Allow for investigating examiner-sample interactions.

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Continuous data model

- Let Y_{ijk} be continuous measurements (e.g., signature complexity) for sample j by examiner i on the k^{th} trial.
- Assume that α_i is a random effect for examiner ability, γ_j is random effect for sample difficulty, and δ_{ij} is an interaction between sample and examiner.

$$p(Y_{ijk}|\beta_0, \alpha_i, \gamma_j, \delta_{ij}) \sim N(\beta_0 + \alpha_i + \gamma_j + \delta_{ij}, \sigma_\epsilon^2)$$

$$p(\alpha_i|\sigma_\alpha^2) \stackrel{i.i.d.}{\sim} N(0, \sigma_\alpha^2)$$

$$p(\gamma_j|\sigma_\gamma^2) \stackrel{i.i.d.}{\sim} N(0, \sigma_\gamma^2)$$

$$p(\delta_{ij}|\sigma_\delta^2) \stackrel{i.i.d.}{\sim} N(0, \sigma_\delta^2)$$

- A Bayesian approach was used to fit these models.

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- A diffuse prior was used on β_0 and a half-t prior was used on standard deviations $\sigma_\alpha, \sigma_\gamma, \sigma_\delta$.
- One way of assessing reliability in this setting, would be to calculate correlation: $\sigma_\alpha^2, \sigma_\gamma^2, \sigma_\delta^2$.

$$\text{Reproducibility} = \text{corr}(Y_{ijk}, Y_{i'jk'}) = \frac{\sigma_\gamma^2}{\sigma_\alpha^2 + \sigma_\gamma^2 + \sigma_\delta^2 + \sigma_\epsilon^2}$$

$$\text{Repeatability} = \text{corr}(Y_{ijk}, Y_{ijk'}) = \frac{\sigma_\alpha^2 + \sigma_\gamma^2 + \sigma_\delta^2}{\sigma_\alpha^2 + \sigma_\gamma^2 + \sigma_\delta^2 + \sigma_\epsilon^2}$$

- Computation for the reliability components is straightforward through MCMC. Credible intervals are also easily obtained.

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Continuous Data Simulations

- For simulation studies we try to replicate studies with smaller repeatability studies as compared to the reproducibility study.
- We simulate 100 datasets for fixed parameters β_0 , σ_α^2 , σ_γ^2 , σ_δ^2 for total number of examiners $I = 10$ and total number of samples $J = 120$.
- Four scenarios:
 - 100% replicates - ($120 \times 10 \times 2$)
 - 50% replicates - (60 samples re-examined)
 - 25% replicates - (30 samples re-examined)
 - 12.5% replicates - (15 samples re-examined)
- We report average posterior median and average lower 2.5% quantiles and average upper 97.5% quantiles.

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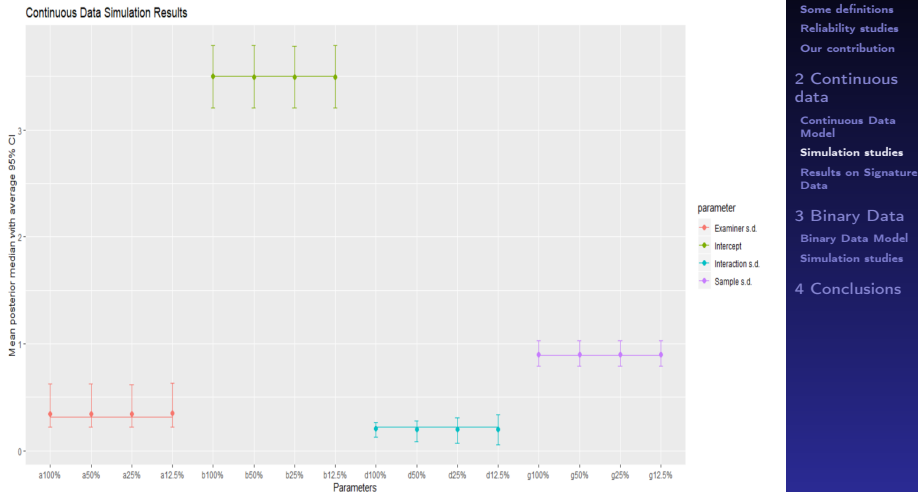


Figure: Simulation results from 100 datasets (continuous data) for each setting of replicates.

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Signature data results

- Analyzed the signature 1-5 point complexity data.

Parameters	σ_{α}^2 Examiner variation	σ_{γ}^2 Signature variation	σ_{δ}^2 Interaction variation
Estimates	0.09 (0.01, 0.40)	0.80 (0.61, 1.05)	0.02 (0.00, 0.09)

Table: Posterior means with 95% credible intervals. A low interaction is a positive for an measurement process.

Model type	Reproducibility $\text{corr}(Y_{ijk}, Y_{i'jk'})$	Repeatability $\text{corr}(Y_{ijk}, Y_{ijk'})$
Separate Analysis	0.65 (0.58, 0.72)	0.67 (0.36, 0.85)
Combining datasets (using posterior mean)	0.63 (0.49, 0.71)	0.71 (0.64, 0.80)

Table: Changes in reliability after combining datasets and accounting for interaction variance. Note that the credible interval is much narrower for repeatability.

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- Assume Y_{ijk} are binary measurements (e.g., match/non-match or value/no-value) on sample j by examiner i on the k^{th} trial.
- Albert and Chib (1993) proposed an MCMC algorithm for Bayesian analysis for binary and polychotomous data based on latent variables Z_{ijk} .

$$Y_{ijk} = \mathbb{1}(Z_{ijk} > 0)$$

$$p(Z_{ijk} | \beta_0, \alpha_i, \gamma_j, \delta_{ij}) \sim N(\beta_0 + \alpha_i + \gamma_j + \delta_{ij}, 1)$$

$$p(\alpha_i | \sigma_\alpha^2) \stackrel{i.i.d.}{\sim} N(0, \sigma_\alpha^2)$$

$$p(\gamma_j | \sigma_\gamma^2) \stackrel{i.i.d.}{\sim} N(0, \sigma_\gamma^2)$$

$$p(\delta_{ij} | \sigma_\delta^2) \stackrel{i.i.d.}{\sim} N(0, \sigma_\delta^2)$$

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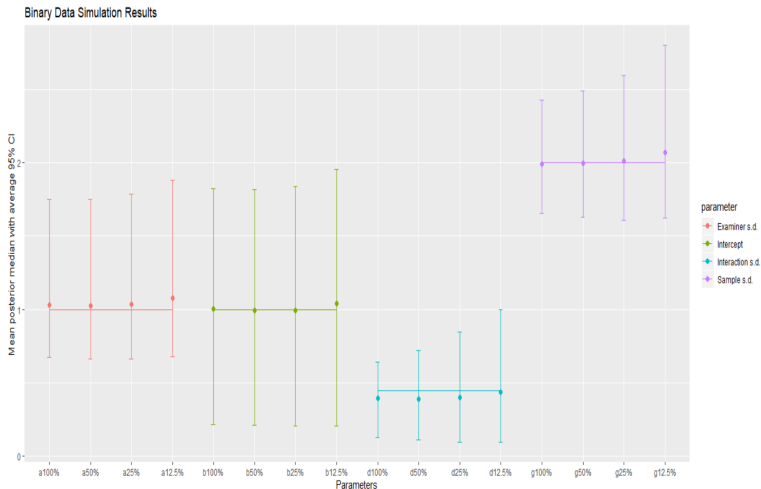


Figure: Simulation results from 100 datasets (binary data) for each setting of replicates.

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- A single model can be used to simultaneously analyze data from reproducibility and repeatability studies.
- Have been able to show through simulation studies that parameters can be recovered even with 25% repeated decisions.
- In forensic signature analysis no evidence for examiner-signature interactions.
- Some convergence issues with small interaction variance and higher computational time for large data sets.

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