DGHT Key Population Surveillance Seminar Series

Novel approaches to population size estimation synthesis using Bayesian statistics









The Anchored Multiplier calculator

A Bayesian tool to reconcile discrepant population size estimates

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8/17/2023



Population Size Estimation (PSE) is fundamental to public health surveillance:

- Apportionment/Equitable distribution of services and resources
- Determine denominators for calculating measures of disease
- Establish sampling frame for studies
- Quantify a potential public health issue
- Evaluate the reach and coverage of outreach programs and social services
- Set concrete targets for program services



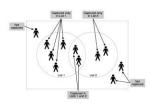
Methods for Population Size Estimation











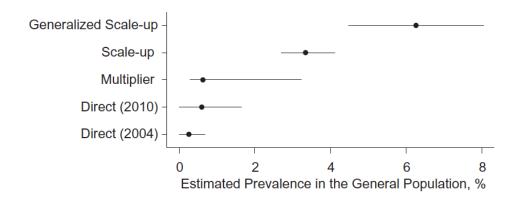


Opinion	General Population	Venue	Literature	Multiple- Sample	Network
Programme Estimates	Population survey	Mapping and Enumeration	Literature review	Capture- Recapture	Network Scale- up
Wisdom of the Crowd		Reverse Tracking Method		Multiplier Method	Sequential Sampling (SS- PSE)
Modified Delphi					





How reliable (consistent) are population size estimates?



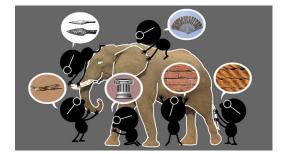
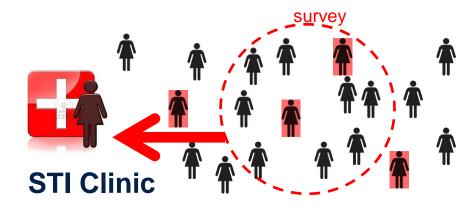


Figure 2. Five estimates of the prevalence of heavy drug use in Curitiba, Brazil, 2004 and 2009–2010. Scale-up and generalized scale-up estimates were substantially higher than those obtained from standard methods (direct estimation and the multiplier method). Estimates of the number of heavy drug users in Curitiba ranged from 4,700 to 114,000. Bars, estimated 95% confidence interval.



Multiplier Method

- Data
 - 2 data sources
- Estimation
 - N = M/P
- Assumptions
 - Two data sources must be independent (count and survey)
 - Two data sources must be specific to the population (correct person, place, time)
 - At least one source should be representative
 - Limited in- and out-migration
 - No duplicate count (within data source)



$$N = \frac{1}{P} \times M = \frac{M}{P}$$



Member of Hidden P.



Member of Hidden P. who received STI service (in last year) from the clinic

M: total recruited people in your hidden population survey

P: proportion of hidden population who received STI services from the clinic

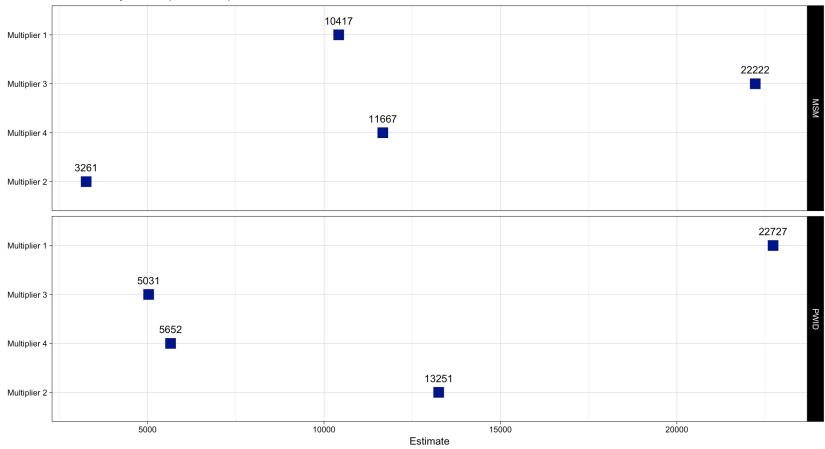


Multiplier Method – Potential Sources of Bias

- Sample from target population may not be representative
- Service providers may report number of visits (not number of unique people)
- Service providers may report total number of people seen (including those who are not members of the key population)
- Study sample participants may misremember receiving specific service
- Dependency (positive or negative) between the benchmark and the multiplier









How do you arrive at a single "best" estimate?

Take the median

- Pro: Simple. Presumed to "balance" biases that drive estimates in different directions
- Con: Assumes individual estimates are equally biased (or valid)

Delphi approach

- **Pro**: Incorporate local knowledge. Final estimate is more "acceptable" to stakeholders
- Con: Vulnerable to subjective biases to increase or decrease PSE



The Anchored Multiplier

- Bayesian framework to synthesize multiple data points (namely, estimates from multiple multipliers) and stakeholder knowledge (or estimates from published literature) as a priori belief.
- Estimates from multiplier methods (likelihood) and prior belief (prior) converted to probability distributions and combined according to Bayes Theorem.
- Posterior probability distribution reflects updated knowledge of the population size, driven by the strength (i.e. precision) of the individual inputs.

$$p(\theta|x) \propto p(x|\theta)p(\theta)$$



Anchored Multiplier – Variance Adjusted

- Incorporating more data results in increasingly strong priors and extremely narrow 95% credible intervals
- Borrow methodology from meta-analysis literature for calculating random effects variance to estimate additional variance:

$$\tau^2 = \frac{Q - df}{C}$$

• Where...

$$Q = \sum_{i=1}^{k} W_i Y_i^2 - \frac{\left(\sum_{i=1}^{k} W_i Y_i\right)^2}{\sum_{i=1}^{k} W_i}; \qquad C = \sum_{i=1}^{k} W_i - \frac{\sum_{i=1}^{k} W_i^2}{\sum_{i=1}^{k} W_i}; \qquad df = k-1$$



Case study: PWID in San Francisco

- Data Source
 - National HIV Behavioral Surveillance (NHBS) data for PWID in San Francisco
 - 3 rounds of data collection 2005, 2009, 2012
 - Published estimates from Multiplier Methods (Chen et al. 2016)
- Priors
 - Lifetime experience = 2.6% (1.8% 3.3%) ... Lansky et al. 2014
 - Recent drug use = 0.3% (0.19% 0.41%) ... Lansky et al. 2014
 - Vaguely informative = Uniform (0, 12.88%)
- Model

 $y \sim dbinom(theta, n)$

Prior

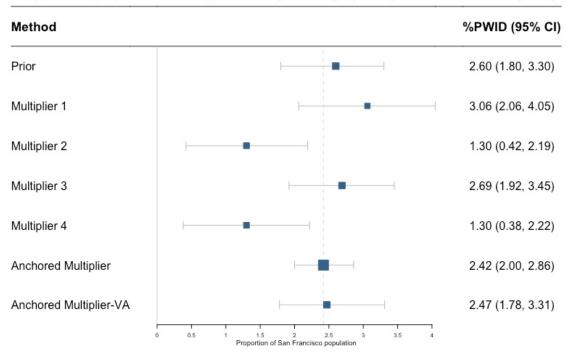
theta \sim dbeta(a, b)

y = population proportion of PWID



Results: Comparison of estimates (2005)

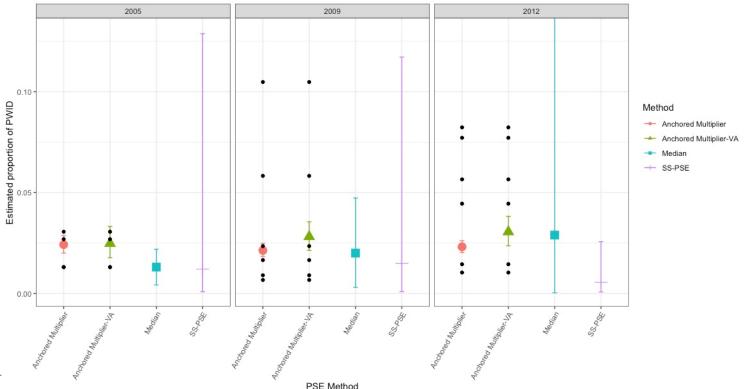
Population proportion of People Who Inject Drugs (PWID) in San Francisco (2005)





Results: Comparison of estimates over time

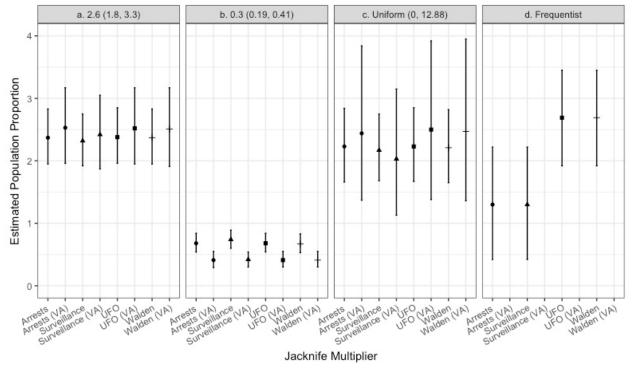
Comparison of the proportion of People Who Inject Drugs (PWID) in San Francisco at three cross-sections using different PSE methods





Results: Sensitivity Analysis

Anchored Multiplier Sensitivity Analysis



Jackknife_Multiplier

- Arrests
- Surveillance
- UFO
- + Walden



Strengths

- Accounts for "strength" of individual estimates, does not treat all estimates as equally precise/valid
- Systematically synthesizes data in way that is transparent and replicable
- Posterior "anchored" by the prior, down-weighted (imprecise) estimates do not pull final estimate towards zero
- Does not require advanced quantitative background

Limitations

 Not a guaranteed fix to biases from the multiplier method (or other PSE methods)



Appropriate Usage

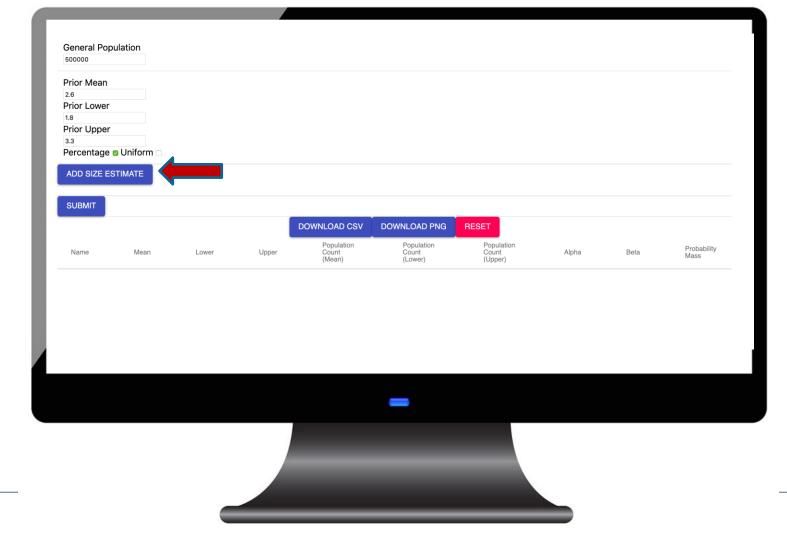
- Population size estimates
 - Using ANY PSE method (not just the multiplier method)
 - Population percentage or absolute counts as input
 - At least two population size estimates required (including the prior)
- Prevalence estimates
- The online calculator is freely available at

https://globalhealthsciences.ucsf.edu/resources/tools

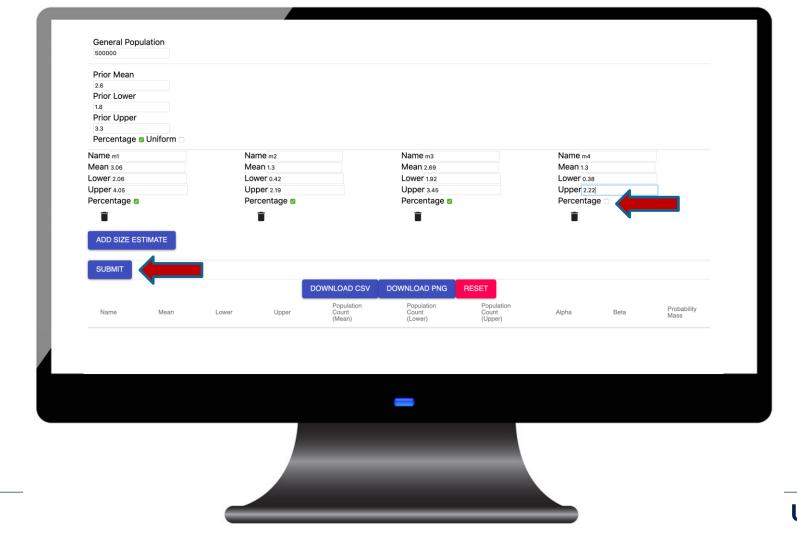






















Questions?

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Acknowledgments

- This work was supported by: NIH/NIMH T32 MH19105
- Ali Mirzazadeh, PhD, MD, MPH
- Willi McFarland, MD, PhD, MPH, TM

Publications:

Wesson PD, Mirzazadeh A, McFarland W. A Bayesian approach to synthesize estimates of the size of hidden populations: the Anchored Multiplier. Int J Epidemiol. 2018 Oct 1; 47(5):1636-1644.

Wesson PD, McFarland W, Qin CC, Mirzazadeh A. Software Application Profile: The Anchored Multiplier calculator-a Bayesian tool to synthesize population size estimates. Int J Epidemiol. 2019 Dec 1; 48(6): 1744-1749.

Rasheed A, Sharifi H, Wesson P, Pashtoon SJ, Tavakoli F, Ghalekhani N, Haghdoost AA, Atarud A, Banehsi MR, Hamdard N, Sadaat SI, McFarland W, Mirzazadeh A. Mapping and population size estimates of people who inject drugs in Afghanistan in 2019: Synthesis of multiple methods. PLoS One. 2022 Jan 28; 17(1):e0262405.



Triangulating Truth

A Flexible Statistical Tool for Reaching Consensus on Population Size, Prevalence and More Ian Fellows, Carl Corcoran and Anne McIntyre

Key Population Surveillance Team Webinar Series August 15, 2023







Motivation

- Population size, prevalence, incidence, etc... are critical metrics
- Experts are faced with multiple estimates of varying quality, and estimates are often incongruent with one-another
- Combining estimates is often done in informal, non-transparent ways
 - 'Pick one' from multiple estimates at a point in time from one city
 - Group of experts often pick one that makes 'most sense'
- Synthesizing expert knowledge with empirical estimates can be tricky



Goal

Create a tool for stakeholders to use that:

- 1. Synthesizes multiple possibly contradictory estimates of a single population quantity.
- 2. Allows the incorporation of stakeholder knowledge about the population and estimates.
- Is fully transparent with triangulations that are auditable by any third party.
- 4. Can be used to triangulate different types of population quantities (e.g. means, proportions, population sizes)
- 5. Is easy to use and intuitive.



Background

Inspiration

Inspired by meta-analysis models

	Fixed Effects	Random Effects	Quality Effects
Weights	$w_j = \frac{v_j^{-1}}{\sum_{j=1}^k v_j^{-1}}$	$w'_{j} = \frac{(v_{j} + \tau^{2})^{-1}}{\sum_{j=1}^{k} (v_{j} + \tau^{2})^{-1}}$	$w''_{j} = \frac{\left(\frac{Q_{j}}{v_{j}} + \hat{\tau}_{j}\right)}{\sum_{j=1}^{k} \left(\frac{Q_{j}}{v_{j}} + \hat{\tau}_{j}\right)}$
Estimator	$\hat{\theta}_{FE} = \sum_{j=1}^{k} w_j \ \hat{\delta}_j$	$\hat{\theta}_{RE} = \sum_{j=1}^{k} w'_{j} \hat{\delta}_{j}$	$\hat{\theta}_{QE} = \sum_{j=1}^{k} w^{\prime\prime}{}_{j} \ \hat{\delta}_{j}$

Notation: $\hat{\delta}_j$ is the estimated effect size of *j*th study, v_j is the sampling error variance of *j*th study, τ^2 is the DerSimonian and Laird estimator of the between-study variance, Q_j is the *j*th study rank (scaled between 0 to 1), and $\hat{\tau}_j = f(Q_j, v_j, N)$ is a bias correction (where *N* is the number of studies).

Background

Broad Strokes

- Create a Bayesian model similar to the "quality effects" model
 - Scale the study error by a "confidence" score, similar to "quality" score
 - Incorporate between-study variance as a source of variation



Background

Broad Strokes

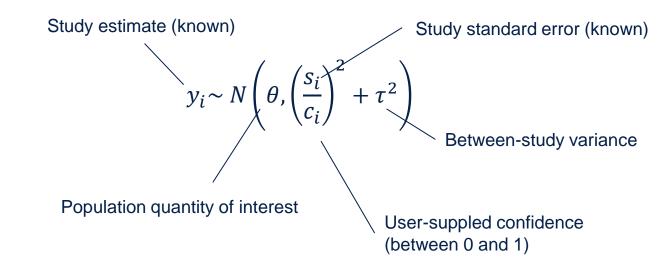
- Bayesian hierarchical model: statistically rigorous framework that allows for input of user knowledge of population
 - **Empirical inputs**: *all* estimates of population quantity (and their standard errors)
 - **Expert inputs**: prior knowledge of the population quantity, confidence in estimate quality



Begin Math

Model Structure

Data Model



Priors

$$\theta \sim N(\mu_0, \sigma_0^2)$$

the population quantity Prior for the unaccounted for between study variation. The prior is set to be

Stakeholder knowledge about

 $\tau \sim HC(0.01 \cdot Var(y_i))$ very close to zero most of the time.



Detail: Theta Transformations

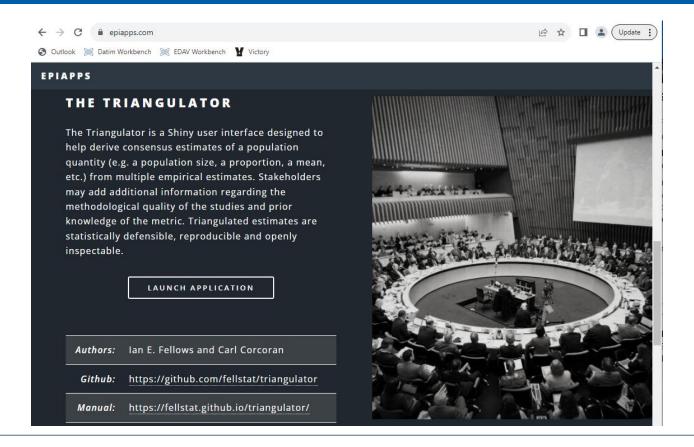
- It is often better to triangulate a transformed population quantity (e.g. triangulate log(Population Size) rather than Population Size directly) because
 - Invalid ranges are prevented (e.g. a proportion of -.05).
 - The normal approximation of the estimates may be more valid.
 - The functional form of the prior may be better.
- This is all handled behind the scenes and is transparent to the user.

Quantity	Transformati on	Formula	Untransforme d Range	Motivating Model
Mean or Other	None	y = x	$[-\infty,\infty]$	Normal
Proportion	Logit	$y = \log(\frac{x}{1-x})$	[0,1]	Logistic
Population Size	Log	$y = \log(x)$	(0,∞]	Log-linear

UCSF

End Math

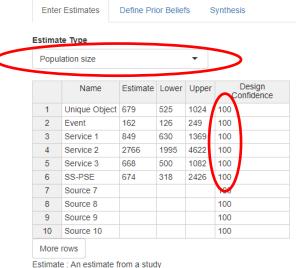
Web Application



Example 1 - PSE

The App

The Triangulator - A Consensus Estimate Calculator



Lower: 95% confindence interval lower bound

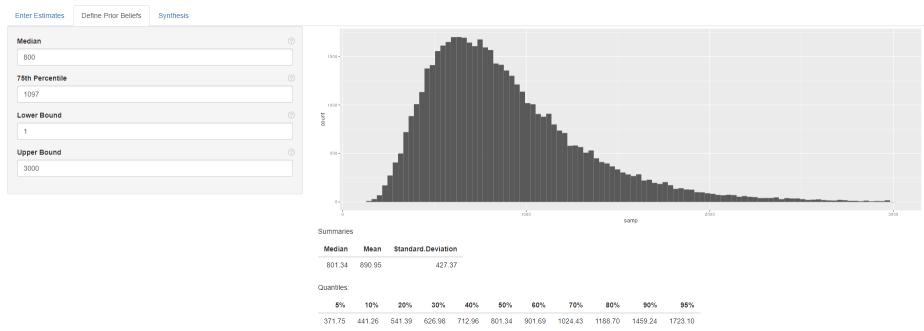
Upper: 95% confindence interval upper bound

Design Confidence: Expert confidence in the design / implementation of the study. This scales the standard error such that a value of 50 will double the standard error.

Example 1 - PSE

The App

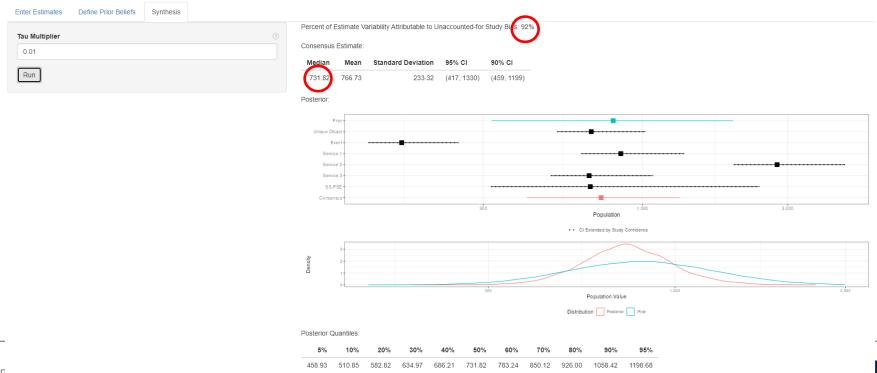
The Triangulator - A Consensus Estimate Calculator



Example 1 - PSE

The App

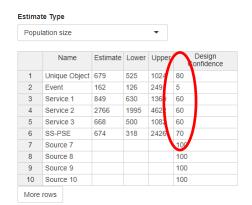
The Triangulator - A Consensus Estimate Calculator

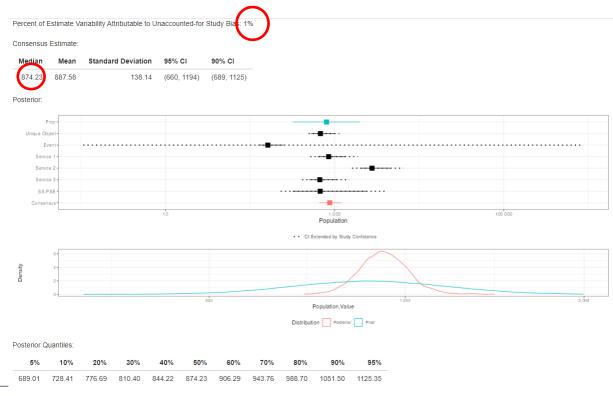


Division TR

Example 1 - PSE

The App

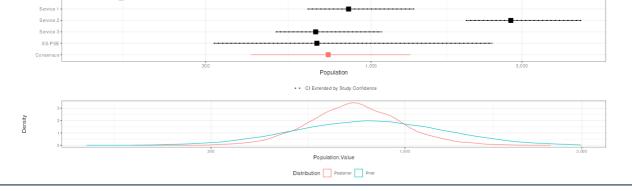




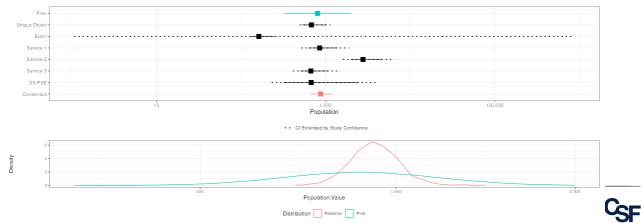
Example 1 - PSE

Unique Object

Unscaled: 732 (417, 1330)



Confidence Scaling: 847 (660, 1194)



Division of Global HIV &

J.

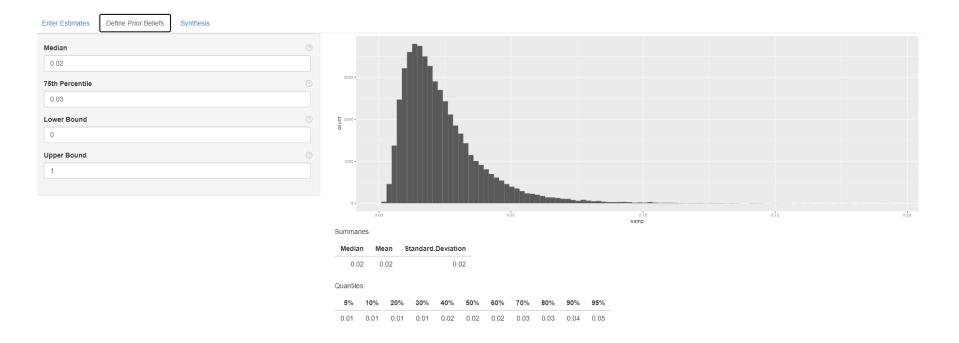
FSW in Mombasa, Kenya

The Triangulator - A Consensus Estimate Calculator

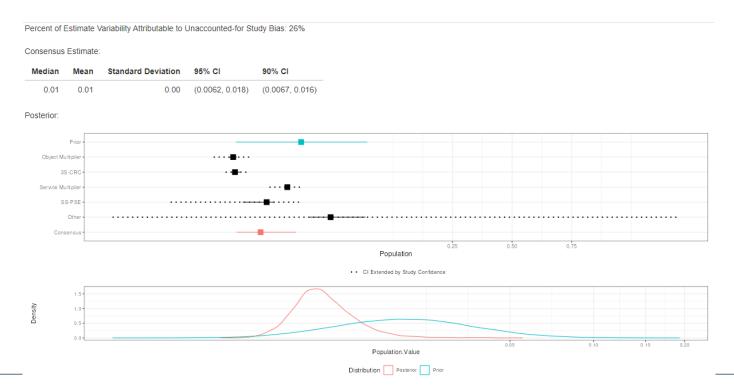


Design Confidence: Expert confidence in the design / implementation of the study. This scales the standard error such that a value of 50 will double the standard error.

Prior 1: Informative

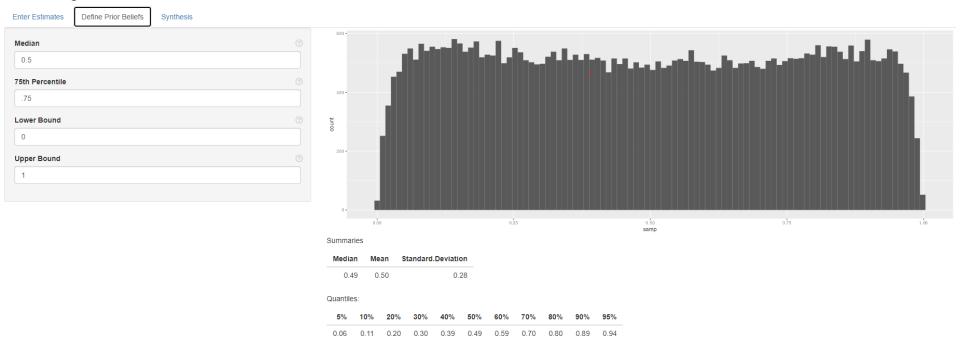


Prior 1: Informative



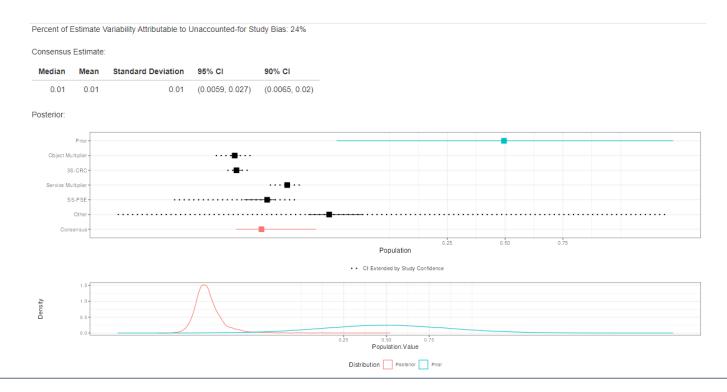
Prior 2: Less Informative

The Triangulator - A Consensus Estimate Calculator

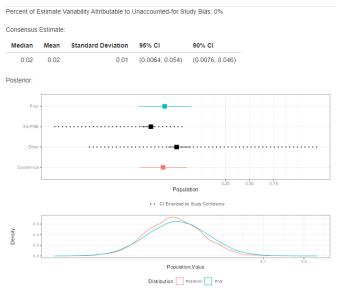


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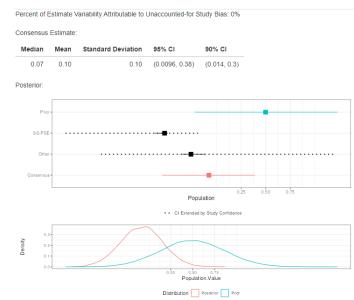
Prior 2: Less Informative



When does the prior have a big impact?



Informative Prior



Less Informative Prior

When uncertainty is high (and/or confidence is low)



Benefits and Limitations

- Statistically rigorous framework
- Combine empirical evidence and expert knowledge
- Ease of use
 - https://epiapps.com/shiny/app_direct/shinyproxy_combine_estimates/
 - https://github.com/fellstat/triangulator + manual

- Requires confidence bounds on estimates
 - Triangulator manual on Github has techniques to reconstruct intervals in some cases



Acknowledgements

- Original Paper
- Triangulating Truth: A Flexible Statistical Tool for Reaching
- Consensus on Population Size, Prevalence and More
- Ian E Fellows 1,2*, PhD; Carl Corcoran 15*, PhD; Anne F McIntyre 1, MPH, PhD
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- Abstract

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- Background: Population size, prevalence, and incidence are essential metrics that influence public health
- policy. However, stakeholders are frequently tasked with designing policy based on multiple (often
- incongruous) estimates of these variables, and often do so in the absence of a formal, transparent
- framework for reaching a consensus estimate.
- Objective: This study aims to describe a model to synthesize multiple study estimates while
- 23 incorporating stakeholder knowledge, introduce an R Shiny application to implement the model, and
- demonstrate the model and application using real data.
- Methods: This study develops a Bayesian hierarchical model to synthesize multiple study estimates that
- allows the user to incorporate the quality of each estimate as a 'confidence' score. The model is
- 27 implemented as a user-friendly R Shiny app, aimed at practitioners of population size estimation. The
- underlying Bayesian model is programmed in STAN for efficient sampling and computation.
- Results: The application is demonstrated using population size estimates (and accompanying confidence
- scores) of female sex workers and men who have sex with men in a country in sub-Saharan Africa. The
- consensus results incorporating confidence scores are compared to the case where they are absent, and the
- results with confidence scores are shown to perform better according to an application-supplied metric for
- Conclusions: The utility of the model, including the incorporation of confidence scores, is demonstrated
- by a use case example.

Special thanks to Anne McIntyre, Ray Shiraishi and Oli Stevens



The Triangulator

Explained Variance

- Bayesian Measures of Explained Variance and Pooling in Multilevel (Hierarchical) Models (Gelman and Pardoe, 2006)
- Bayesian hierarchical model equivalent of R²

$$R^2 = 1 - rac{E\Bigl(igvee_{j=1}^N y_j -
u_j\Bigr)}{E\Bigl(igvee_{j=1}^N y_j\Bigr)}$$

Fraction of estimate variability due to unaccounted-for study bias



The Triangulator

Explained Variance

$$R^2 = 1 - rac{Eig(igee_{j=1}^N y_j -
u_jig)}{Eig(igee_{j=1}^N y_jig)}$$

$$R^{2} = \frac{E\left(\bigvee_{j=1}^{N} \hat{v}_{j} + \frac{1}{N} \sum_{j=1}^{N} V_{j}\right)}{E\left(\bigvee_{j=1}^{N} y_{j}\right)},$$

$$\hat{v}_j = \frac{\tau^2}{\sigma_j^2 + \tau^2} y_j + \frac{\sigma_j^2}{\sigma_j^2 + \tau^2} \theta \text{ and } V_j = \frac{\sigma_j^2 \tau^2}{\sigma_j^2 + \tau^2}.$$

