

## Bias-corrected Cancer Incidence Rate to account for Differential Privacy Error in the Population Data

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#### Disclaimer:

The opinions expressed in this presentation are the author's own and do not reflect the view of the National Cancer Institute, National Institutes of Health, the Department of Health and Human Services, or the United States government.



## **Outline**

- Background
  - NCI's SEER program and Cancer Rates
  - Differential Private (DP) Census Population Estimates
  - Bias-corrected Rate Estimator
- Study Objective performance of correcting for bias due to DP error in Population data
- Discussion



# **Cancer Surveillance and SEER Areas**

- Surveillance, Epidemiology, and End Results (SEER) program is established and supported by the NCI since 1973.
- SEER collects and publishes cancer incidence and survival data from population-based registries across the nation.
- Information include patient demographics, primary tumor site, tumor morphology and stage at diagnosis, first course of treatment, and follow-up for vital status (survival)



SEER registries cover ~48% of the U.S. population.



#### **Age-Adjusted Rate of Cancer Incidence**





Data Source:

 SEER Incidence Data, November 2022 Submission (1975-2020), SEER 22 registries [https://seer.cancer.gov/registries/terms.html]. Methodology: Rates are per 100,000 and are age-adjusted to the 2000 US Std Population (19 age groups - Census P25-1130).
 The Annual Percent Change (APC) and Average Annual Percent Change (AAPC) estimates were calculated from the underlying

rates using the Joinpoint Trend Analysis Software (https://surveillance.cancer.gov/joinpoint), Version 4.9, March 2021, National Cancer Institute using the default settings

Lancer instance study of the data serings. The APCS/AAPCS direction is "Bsing" (1) when the entire 95% confidence interval (C.1) is above 0, "Falling" (1) when the entire 95% C.1 is lower than 0, otherwise, the trend is considered "Not Significant". The 2020 incidence rate is displayed but not used in the fit of the trend line(s), impact of COVID on SEER Cancer incidence 2020 data

[https://seer.cancer.gov/data/covid-impact.html] Race/Ethnicity Coding:

. For more details on SEER race/ethnicity groupings and changes made to the grouping for this year's data release, please see Race and Hispanic Ethnicity Changes [https://seer.cancer.gov/seerstat/variables/seer/race\_ethnicity/]. • Rates for American Indians/Alaska Natives only include cases that are in a Purchased/Referred Care Delivery Area (PRCDA) Incidence data for Hispanics and Non-Hispanics are based on the NAACCR Hispanic Latino Identification Algorithm (NHA)

Cancer Site Coding: · See SEER\*Explorer Cancer Site Definitions [https://seer.cancer.gov/statistics-netw/ork/explorer/cancer-sites.html] for details about the cancer site coding used for SEER Incidence data.

Created by https://seer.cancer.gov/statistics-network/explorer on Tue Oct 10 2023

Age Adjusted Rate = 
$$\sum_{J} w_j \frac{c_j}{N_j}$$

 $c_i$  = number of tumors in age group j  $w_i$  = age-adjusting weight in age group j  $N_i$  = at risk population in age group j, and it is assumed to be error-free (despite of small enumeration error)

### **Bias-corrected Rate**



Age Adjusted Rate<sub>bc</sub> = 
$$\sum_{J} w_{j} \frac{c_{j}}{\widehat{N}_{j}} (1 - CV^{2})$$

 $\widehat{N}_{j}$  = estimated population in age group j

 $CV_i$  = estimated Coefficient of Variation of population in age group j

> Stat Methods Med Res. 2021 Feb;30(2):535-548. doi: 10.1177/0962280220962516. Epub 2020 Oct 15.

Inference about age-standardized rates with sampling errors in the denominators

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# **Differential Private Population Data**



- A Topdown algorithm to adds noise or variations from the actual count – to the collected data
- Geographic pop control totals were assigned a privacy budget (state totals are exactly matched)
- However, pop totals by demographics, such as age and race, are not controlled for and may be subject to systematic deviations
- Size of deviation tends to be greater for small groups despite of iterations of improvements from Census Bureau on the Topdown algorithms

#### **A Few Key Challenges for Calculating Cancer Rates**

Inaccurate

AGE data

for Age-

Adjusting



Race/Ethnicity data may be compromised

# **Study Objectives and Methods**

- Objectives:
  - 1. Impact of DP on the validity of rates
  - 2. Performance of bias-correction to alleviate the DP impact
- Use California data as a test case
  - Stratification variables: county and 5-year age group
  - Outcome: age-adjusted rates for counties
- Most challenging part is how to simulate DP population estimates
  - Detailed algorithms are confidential and kept within Census Bureau
  - Size of DP errors derived from Census 2010 demonstration dataset
  - A TopDown approach similar in principle to Census's algorithm



# **Population Simulation-A TopDown Appraoch**

- Add normal-distributed noise to pops totals
- With variance of noise proportional to the size of population totals (p=0.2)
- Resulted noise is similar to observed differences between the demonstration and real 2010 census data
- Optimized noise size using a two-step approach



# **Simulate Study**

- Questions to be answered
  - Whether Bias-correction helps to adjust for DP error?
  - What is the population cutoffs for DP errors to be negligible for ageadjusted rates of incidence (AAR)?
- Study Methods:
  - Simulate Poisson incidence county by age and county (since cancers are random events)
  - Calculate AARs using DP-pop and Real-pop
  - Calculate % Relative Bias of AAR from simulation studies



# **Results: % Relative Bias in AAR (county-level)**



DATA

<b>3: Summary of % RB for Different ASR Estimators (TopDown Algo</b>			
Summary	Naive No DP	Naive with DP	Bias-correction with DP
Min.	-4.413	-2.238	-4.338
Q1	-0.110	-0.110	-0.111
Median	0.025	0.05	0.023
Mean	-0.098	-0.042	-0.102
Q3	0.263	0.283	0.282

2.071

Naïve No DP: Population does not have DP errors

1.603

Max.

Naïve with DP: Population has DP errors and AAR is calculated using the standard method (i.e. without bias correction) Bias-correction with DP: Population has DP errors and AAR is bias-corrected

1.310

### **Boxplots of % Relative Bias**





Note: Each dot corresponds to one county



### % Relative Bias by County Population Size (log scale)





# A further simulation of Sampling Error+ DP Error

- The case when DP error is added to survey samples
- E.g., DP modified American Community Survey estimates.
- *ρ* is the measure of sampling fraction and gauges sampling error
- q gauges DP error

Figure 3: Boxplots of %RB (TopDown DP algorithm): q = 0.05; (a)  $\rho = 0.05$ , (b)  $\rho = 0.2$ ; PRB\_Rub: Naive No DP; PRB\_Rhat: Naive with DP; PRB\_Rbc: Bias-correction with DP









- The magnitude of DP bias is not comparable to that of sampling error
- The impact of bias-correct on AARs is small in relative to sampling error
- DP error's impacts become small/negligible only if the population is at least 100K or greater limit the ability for detailed disparity analysis
- Current simulation is limited to Age variable and the next step is to consider race/ethnicity

# **Policy Implications**



- Health burden studies for small subpopulations or at local geographic level are becoming impossible, however policies are mostly 'local' and 'specific'
- Noise metrics released by Census Bureau are not detailed enough to help understand the extend of impact on cancer rates
- User community would benefit a guidance from Census Bureau regarding to how to use the noise metrics, e.g., how to relate noise measure to variance for common statistics



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