Statistical Privacy in the 21st Century: Census and Consensus

An Overview of the 2020 Census Disclosure Avoidance System and Noisy Measurement Files

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Disclosure Avoidance for the 2020 Census

The 2020 Census improves on the noise injection methods of the 1990-2010 Censuses by employing a risk assessment framework based on Differential Privacy (DP) to assess and quantify disclosure risk and confidentiality protection.

Every individual that is reflected in a particular statistic contributes towards that statistic's value.

Every statistic that you publish "leaks" a small amount of private information.

DP as a framework allows you to assess each individual's contribution to the statistic, and to measure (and thus, limit) how much information about them will leak.





The 2020 Census Disclosure Avoidance System (DAS)



TopDown Algorithm (TDA)

Produces privacy-protected microdata (Microdata Detail File) that is ingested by Decennial tabulation system

- Redistricting Data (P.L. 94-171)
 Summary File
- Demographic Profile
- Demographic and Housing Characteristics File (DHC)
- Congressional District Summary Files



SafeTab PHSafe

Produce privacy-protected tabulations

- Detailed DHC-A
- Detailed DHC-B
- Supplemental DHC



The TopDown Algorithm

Input Microdata (CEF) & Tabulation Geographic Reference File (Tab GRF-C)

Conversion to Histogram*

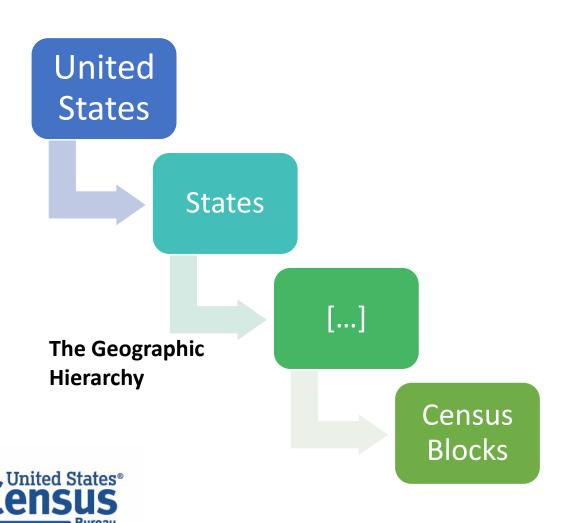
Noisy Measurements Postprocessing Conversion to Microdata (MDF)

*A histogram, in this context, is a tabular representation of the microdata with counts of records for each possible combination of values for each attribute in the microdata.

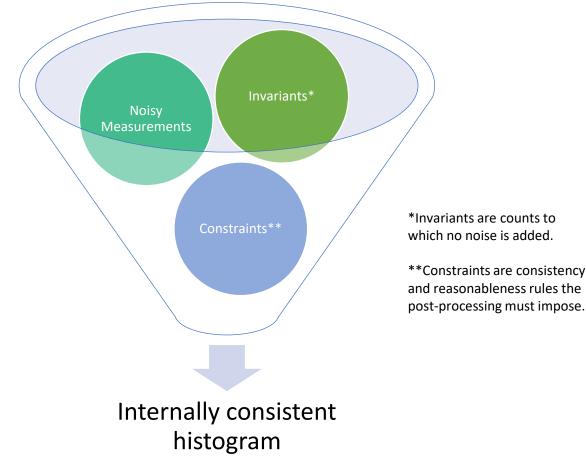
For complete details see: Abowd, J., Ashmead, R., Cumings-Menon, R., Garfinkel, S., Heineck, M., Heiss, C., Johns, R., Kifer, D., Leclerc, P., Machanavajjhala, A., Moran, B., Sexton, W., Spence, M., & Zhuravlev, P. (2022). The 2020 Census Disclosure Avoidance System TopDown Algorithm. *Harvard Data Science Review*. (June) https://doi.org/10.1162/99608f92.529e3cb9



The TopDown Algorithm

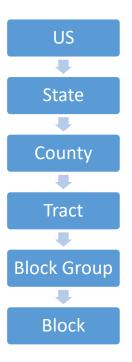


At each geographic level:

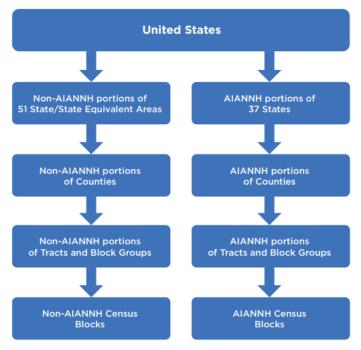


The Geographic Hierarchy ("Spine")

Standard Spine Tabulation Hierarchy

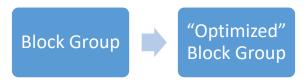


TDA's American Indian/Alaska Native/Native Hawaiian (AIANNH) Spine for Redistricting



Geographic Optimization

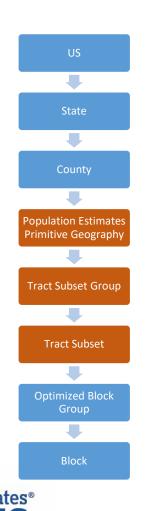
(for Redistricting Data)



Reconfigured TDA's definition of block groups to optimize accuracy in statistics for certain types of geographies, including Minor Civil Divisions, Places, and individual AIANNH areas. Optimized block groups are used inside TDA. Tabulation block groups from the standard hierarchy are used for all published data tables.



Population Estimates Primitive Geographies



Population Estimates Primitive Geographies are the most granular geographic areas that are required in order to derive tables for every geography for which official Population Estimates are produced.

The Population Estimates Primitive Geographies form a complete, mutually exclusive partition of the U.S.

Tract Subsets are defined as the intersection of Population Estimates Primitive Geographies with census tabulation tracts.

Tract Subset Groups are defined as the union of multiple tract subsets that are all within the same Population Estimates primitive geography.

For the DHC, the TDA Geographic Hierarchy was further modified to include "Population Estimates Primitive Geographies" on the spine.

Queries and Privacy-loss Budget Allocation

Global rho	2.56
Global epsilon	17.90
delta	10 ⁻¹⁰

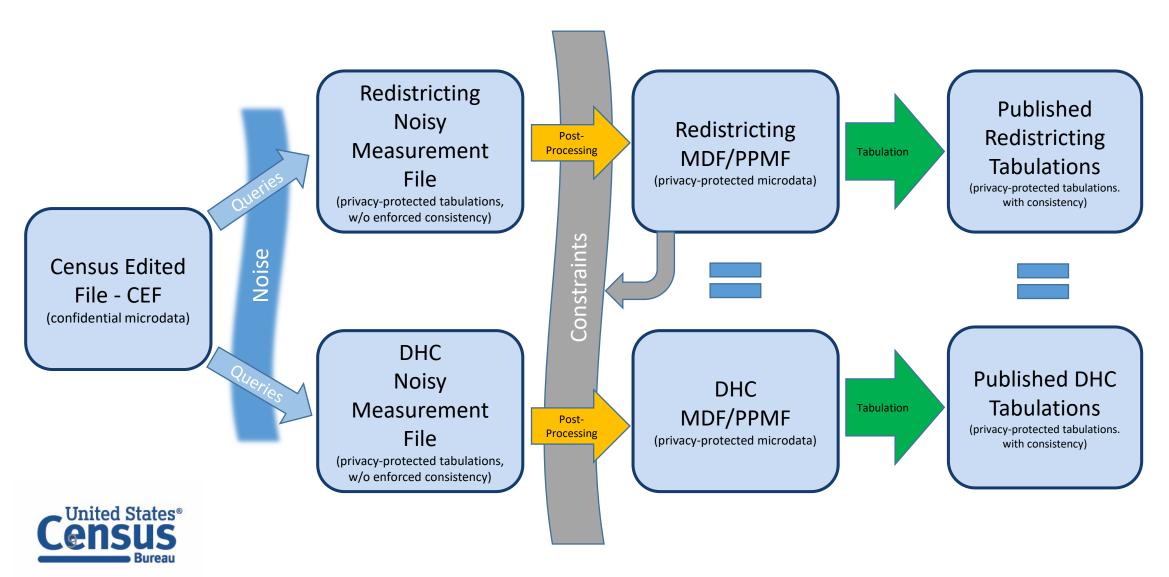
	rho Allocation by Geographic Level
US	2.54%
State	35.13%
County	10.91%
Tract	16.76%
Optimized Block Group*	30.64%
Block	4.03%

Production settings for the 2020 Census Redistricting Data (P.L. 94-171)
Summary File (Persons tables P1-P5)

	Per Query rho Allocation by Geographic Level					
Query	US	State	County	Tract	Optimized Block Group*	Block
TOTAL (1 cell)		32.35%	8.32%	6.40%	12.75%	0.00%
CENRACE (63 cells)	0.03%	0.05%	0.03%	0.03%	0.02%	0.01%
HISPANIC (2 cells)	0.02%	0.05%	0.03%	0.02%	0.02%	0.00%
VOTINGAGE (2 cells)	0.02%	0.05%	0.03%	0.02%	0.02%	0.00%
HHINSTLEVELS (3 cells)	0.02%	0.05%	0.03%	0.02%	0.02%	0.00%
HHGQ (8 cells)	0.02%	0.05%	0.03%	0.02%	0.02%	0.00%
HISPANIC*CENRACE (126 cells)	0.08%	0.10%	0.07%	7.90%	7.89%	0.02%
VOTINGAGE*CENRACE (126 cells)	0.08%	0.10%	0.07%	0.08%	0.07%	0.02%
VOTINGAGE*HISPANIC (4 cells)	0.02%	0.05%	0.03%	0.02%	0.02%	0.00%
VOTINGAGE*HISPANIC*CENRACE (252						
cells)	0.27%	0.29%	0.27%	0.27%	0.18%	0.07%
HHGQ*VOTINGAGE*						
HISPANIC*CENRACE (2,016 cells)	1.99%	1.97%	2.01%	1.97%	9.63%	3.88%



Noisy Measurement Files (NMFs), Privacy-Protected Microdata Files (PPMFs), Published Tabulations

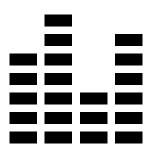


Should I Use the NMF, the PPMF, or the Tabulations?

• There are two sources of error in the published statistics (PPMF and Tabulations):

Differentially private noise

- Unbiased
- Known distribution
- Reflected in the noisy measurements



Post-processing

- Data dependent
 - While the nonnegativity requirement decreases error in the detailed cell counts, it also introduces a <u>positive bias</u> in small counts and an offsetting <u>negative bias</u> in large counts.
 - TDA also reduces the amount of error for many statistics relative to their corresponding noisy measurements.
- Block-level statistics will often have a <u>lower expected variation</u> than you would expect based solely on the amount of PLB assigned to that query at the block level.



Should I Use the NMF, the PPMF, or the Tabulations?







2020 Census Redistricting and DHC Tabulations

- Official 2020 Census Statistics
- Higher Accuracy (feature of TDA)
- Does include bias due to post-processing

2020 Census PPMF

- 100% microdata file
- Consistent with published tabulations
- Useful for special tabulations and microdata analysis

2020 Census NMF

- Can be used to produce unbiased estimates and confidence intervals
- Can be used to evaluate alternate post-processing mechanisms
- Research product

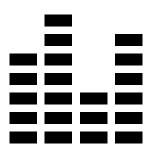


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How to Access the NMFs

There are approximately 25 trillion numbers associated with the combined redistricting data and DHC Noisy Measurement Files, consuming 33 terabytes of space in a compressed format. Given their large size, the files are housed and accessible via off-site locations.

Use the links below to access the files:

2010 Demonstration Data	Release Date	Download L	ocations
Redistricting Data (P.L. 94-171)	April 3, 2023	Inter-university Consortium for Political and Social Research (ICPSR)*	Harvard Dataverse
Demographic and Housing Characteristics File (DHC)	June 30, 2023	<u>ICPSR*</u>	N/A
2020 Research Data	Release Date	Download Lo	ocations
Redistricting Data (P.L. 94-171)	June 15, 2023	<u>ICPSR*</u>	<u>Harvard Dataverse</u>
Demographic and Housing Characteristics File (DHC)	Fall 2023		



^{*} Requires registering for a free Globus account to access and download.

Content of the NMF

	geocode	query_name	hhgq	votingage	hispanic	cenrace	query_shape	value	variance	plb
0	1100890045	total_dpq	*	*	*	*	[1 1 1 1]	[2980]	6.110105	[52707577526 322048826043]
1	1100890045	cenrace_dpq	*	*	*	cenrace	[1 1 1 63]	[1975 837 22 6 3 51 9 -9 26 7 -20 -40 72 47 -55 -9 44 -52 -44 44 55 -49 11 -39 -41 -1 -70 -20 12 20 22 -17 51 45 34 76 23 20 0 -27 5 48 6 22 -66 -24 -41 20 -6 -4 -15 -12 -7 -72 -42 1 -26 -14 16 -30 -14 26 -48]	1196.8168	[269087824 322048826043]
2	1100890045	hispanic_dpq	*	*	hispanic	*	[1 1 2 1]	[2890 123]	1914.9069	[168179890 322048826043]
3	1100890045	votingage_dpq	*	votingage	*	*	[1 2 1 1]	[379 2645]	1914.9069	[168179890 322048826043]
4	1100890045	hhinstlevels_dpq	hhinstlevels	*	*	*	[3 1 1 1]	[1781 1127 29]	1914.9069	[168179890 322048826043]
5	1100890045	hhgq_dpq	hhgq	*	*	*	[8 1 1 1]	[1754 1017 59 120 6 -68 -84 -10]	1914.9069	[168179890 322048826043]
6	1100890045	hispanic * cenrace_dpq	*	*	hispanic	cenrace	[1 1 2 63]	[1933 856 24 18 2 2 11 15 3 1 1 7 1 -2 -4 -1 2 -1 5 3 2 -1 -3 -3 1 -2 2 0 2 0 -1 0 -1 0 1 2 -6 -3 -1 3 1 0 -1 -1 1 -2 0 0 -3 -5 1 2 1 1 -2 -1 0 7 -3 -4 0 1 2 67 8 10 -2 -1 13 0 4 0 0 3 -1 0 -2 3 1 1 7 6 2 0 2 0 4 -2 -1 -1 3 0 2 -2 3 3 -2 4 3 4 -1 1 -3 -5 2 -2 -2 1 2 0 -1 4 0 1 -1 -1 4 3 2 -2 4 2 0 0 3 -1]	4.9531994	[65018345474 322048826043]



Using the NMFs to Generate Unbiased Confidence Intervals

Recorded webinar on how to use the NMFs available here: <u>Noisy</u>
 <u>Measurement Files for the Redistricting and DHC Data Products</u>
 <u>(census.gov)</u>

Sample code, examples, and useful tips:
 <u>Computing Confidence Intervals Using the 2010 Census Redistricting</u>
 (P.L. 94-171) Demonstration Data Noisy Measurement File (2023-04-03) by Ryan Cumings-Menon, Michael Hawes, and Matthew Spence (April 2023)



Generating Confidence Intervals

Generating a more precise county-level estimate and CI by leveraging multiple noisy measurements

Bullock County, AL (00110011):

total_dpq: 10911 variance: 4.70162740

Tract 1 (001100110001):

total_dpq: 1433

variance: 6.11010487

Tract 2 (001100110002):

total_dpq: 7105

variance: 6.11010487

Tract 3 (001100110003):

total_dpq: 2376

variance: 6.11010487

```
In [6]: # Constants in this example:
    coef_mat = np.vstack((np.ones((1, 3)), np.eye(3)))
    var_county = 4.70162740
    var_tracts = 6.11010487
    sigma_sqrds = np.array([var_county] + [var_tracts] * 3)
    dp_answers = np.array([10911, 1433, 7105, 2376])

    weights = np.diag(1 / sigma_sqrds)
    point_estimate, ci_half_width = simulation_based_ci(dp_answers, coef_mat, weights, sigma_sqrds, prng, num_sims, alpha)
    print(f"The confidence interval is: {point_estimate} +/- {ci_half_width}")

The confidence interval is: 10911.612405249798 +/- 3.193797375100286
```



Tip: Even when the statistic of interest is measured directly in the NMFs, it can be advantageous to incorporate additional information (e.g., noisy measurements for the child geounits) in order to obtain more precise estimates and smaller CIs.

How do these estimates compare?

2010 Total Population	SF1	2010 PPMF	2010 NMF
Bullock County, AL	10,914	10,912	10,911.61 +/- 3.19
Menominee, WI	4,232	4,230	4,232 +/- 5
Redfield, IA	835	835	835 +/- 4

			Error: middle 90 percent (counts	
Counties by size	Number of	absolute error (counts of	of peop	le)
_	counties	people)	Minus	Plus
All counties.	3,143	117.27	-248	+230
Counties with housing unit population between	37	10.03	-10	+2
1,000-9,999.	691	28.23	-38	+7
Counties with housing unit population between	1,849	74.45	-131	+17
100,000-999,999 Counties with housing unit population at or	527	292.21	-784	+54
above 1,000,000	39	1.463.12	-3,659	+1.35

Counties by size		Mean absolute error	Error: middle 90 percent (counts of people)	
	Number of counties	(counts of people)	Minus	Plus
All counties. Counties with housing unit population between	3,143	964.00	-1,841	+2,048
O-999 Counties with housing unit population between	37	23.00	-22	+54
1,000-9,999. Counties with housing unit population between	691	121.00	-146	+284
10,000-99,999	1,849	446.00	-832	+1,053
100,000-999,999	527	2,930.00	-7,222	+6,278
1,000,000.	39	14,848.00	-44,833	+20,007



research and 2010 Census housing unit population

Source: Understanding Disclosure Avoidance-Related Variability 2020 Census



What additional resources would you like to see?