U.S. Cancer Statistics Public Use Database Technical Documentation

U.S. Data
November, 2020 Submission
Diagnosis Years 2001–2018
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Researchers can access and analyze high-quality population-based cancer incidence data on the entire United States population.

De-identified cancer incidence data reported to CDC’s National Program of Cancer Registries (NPCR) and the National Cancer Institute’s (NCI’s) Surveillance, Epidemiology, and End Results (SEER) Program are available to researchers for free in public use databases that can be analyzed using software developed by NCI’s SEER Program.

Cancer surveillance data from CDC and NCI are combined to become U.S. Cancer Statistics, the official source for federal cancer data. U.S. Cancer Statistics public use databases include cancer incidence and population data for all 50 states, the District of Columbia, and Puerto Rico, providing information on more than 31 million cancer cases.

U.S. Cancer Statistics Public Use Database


- Includes race and ethnicity variables.
- Does not include Puerto Rico data.
- The population denominators are race-specific, ethnicity-specific, and sex-specific county population estimates from the U.S. Census (July 1, 2010–2019 bridged–race vintage 2019 population estimates), modified by SEER and aggregated to the state and national levels.

Population Coverage by Diagnosis Year

For cases diagnosed from 2003 through 2017, 100% of the population is covered for all 50 states and the District of Columbia. In 2001 and 2002, cases that were diagnosed in Mississippi are not available and in 2018, cases that were diagnosed in Nevada are not available. The U.S. population covered for each of those three years is 99%. The U.S. population coverage for 2001 through 2018 is 98%.

Suggested Citations

Please use these standard citations for tables and figures when presented in presentations or publications.

For population coverage: Data are from population-based registries that participate in CDC’s National Program of Cancer Registries and/or NCI’s Surveillance, Epidemiology, and End Results Program and meet high-quality data criteria. These registries cover approximately [XX]% of the U.S. population.

For age-adjusted rates: Rates are per 100,000 persons and are age-adjusted to the 2000 U.S. standard population (19 age groups – Census P25–1130).


U.S. Cancer Statistics Public Use Database

Before using the U.S. (2001–2018) data, analysts should read and understand the following information. If you have questions, please contact CDC at uscsdata@cdc.gov.

Case Inclusions and Exclusions

Cancer registries that are supported by CDC’s National Program of Cancer Registries (NPCR) or the National Cancer Institute’s (NCI’s) Surveillance, Epidemiology, and End Results (SEER) program report all incident cases coded as in situ (non-malignant), invasive (malignant; primary site only), and non-malignant (including borderline and benign) central nervous system tumors according to the International Classification of Diseases for Oncology, Third Edition (ICD-O-3), with the following exceptions—

- In situ cancers of the cervix are not reported.
- Basal and squamous cell carcinomas of the skin are not reported, except when these occur on the skin of the genital organs.
- In situ cancers of the urinary bladder are re-coded as invasive behavior because the information needed to distinguish between in situ and invasive bladder cancers is not always available or reliable. Stage for these cases remains coded as in situ.  

Additionally, in these public use databases—

- Cases with an unknown age or with sex other than male or female have been excluded from the database. The frequency counts will not change based on whether Known Age or Male or Female Sex is checked on the SEER*Stat Selection tab.
- Malignant Behavior is a default selection for this database, as this restriction is used by CDC’s NPCR and NCI’s SEER Program for generating most official cancer statistics. Malignant behavior is defined by the variable Behavior Code ICD-O-3. This database includes in situ and nonmalignant central nervous system (CNS) cases. These nonmalignant cases can be analyzed by unselecting the Malignant Behavior check box on the SEER*Stat Selection tab.

Changes Made to the Database

In the June 2021 release, the following changes were made to the data—

- To protect confidentiality further while allowing access to all cases, the Type of Reporting Source variable is not included in the database. Excluding this variable allows cases identified through death certificates and autopsy reports to be included in the database. This change means that statistics calculated using the public use database now match statistics in the U.S. Cancer Statistics Data Visualizations and CDC WONDER tools.
- Two revised site recode variables were added—
  o AYA site recode 2020
  o Lymphoma neoplasm recode 2021
- The following variables were added—
  o Grade clinical (available for cases diagnosed in 2018)
  o Grade pathological (available for cases diagnosed in 2018)
o Merged estrogen receptor (available for breast cancer cases diagnosed in 2004 or later). This variable replaces CS Site Specific Factor 1 for breast cancers.

o Merged progesterone receptor (available for breast cancer cases diagnosed in 2004 or later). This variable replaces CS Site Specific Factor 2 for breast cancers.

o Merged HER2 receptor (available for breast cancer cases diagnosed in 2010 or later). This variable replaces CS Site Specific Factor 15 for breast cancers.

Suppression Rules

Suppressing Fewer Than 16 Cases

The suppression rule is fewer than 16 cases for the time period based on rate stability. This suppression rule is enforced automatically in these databases.

When the number of cases used to compute the incidence rates is small, those rates tend to have poor reliability. Therefore, to discourage misinterpretation and misuse of counts, rates, and trends that are unstable because of the small number of cases, these statistics are not shown in tables and figures if the counts are fewer than 16 for the time period. A count of fewer than about 16 in a numerator results in a standard error of the rate that is about 25% or more as large as the rate itself. Equivalently, a count of fewer than about 16 results in the width of the 95% confidence interval around the rate being at least as large as the rate itself. These relationships were derived under the assumption of a Poisson process and with the standard population age distribution close to the observed population age distribution.

Another important reason for employing a cell suppression threshold value is to protect the confidentiality of patients whose data are included in a report by reducing or eliminating the risk of identity disclosure. The cell suppression threshold value of 16 is recommended to protect patient confidentiality given the low level of geographic and clinical detail provided.

Complementary Cell Suppression

Complementary cell suppression is necessary to prevent users from subtracting to find suppressed counts. This practice should be employed when any suppression occurs in the data presentation. In addition, when information from other cells, tables, or figures can be used to determine a suppressed cell, at least one other cell must also be suppressed. When analyzing data at the state or regional levels, counts for national and regional data must be suppressed if a single state in a region is suppressed. Rates, confidence intervals (CIs), and populations can be shown at the national and regional levels. This suppression should occur when a single or multiple years of data are being presented.

Race and Ethnicity Suppression

States have the option to suppress race-specific and Hispanic ethnicity-specific data every submission year. While these states can be included in an aggregated analysis, the affected state’s race and ethnicity information cannot be reported at the state level.

The merged system-supplied variable, state race eth suppress, can be used to restrict your analysis to the states that are eligible to be included in a state-level analysis of race and ethnicity combinations. If conducting a state-level analysis of race or ethnicity only, manually make restrictions in the SEER*Stat Selection tab.

The following states have race or ethnicity data presentation restrictions—

- Data for American Indian and Alaska Native people cannot be displayed for Delaware, Illinois, Kansas, Kentucky, New Jersey, and New York.
- Data for Asian and Pacific Islander people cannot be displayed for Delaware, Kansas, and Kentucky.
- Hispanic ethnicity data cannot be displayed for Delaware, Kentucky, and Massachusetts.
• Race and ethnicity combinations—White Hispanic, White non-Hispanic, Black Hispanic, and Black non-Hispanic—cannot be displayed for Delaware, Kansas, Kentucky, Massachusetts, and Pennsylvania.

For more information, please refer to the Race Recode for USCS, Origin recode NHIA (Hispanic, Non-Hisp) variable description.

**Case-Level Data**

As a further mechanism to protect data confidentiality and due to data sharing agreements with some states, the case listing function in SEER*Stat has been disabled for this database.

**Benign Central Nervous System (CNS) Tumors**

Cancer registries began collecting information on nonmalignant brain and other central nervous system tumors with cases diagnosed in 2004. Collection of these tumors is in accordance with Public Law 107-260, the Benign Brain Tumor Cancer Registries Amendment Act, which mandates that NPCR registries collect data on all brain and other central nervous system tumors with a behavior code of 0 (benign) or 1 (borderline), in addition to *in situ* and malignant tumors. Data for nonmalignant brain and other nervous system tumors were available from all registries contributing to this report.

**Behavior**

The behavior variable in the current database is Behavior Code ICD-O-3. Previous database releases included the variable Behavior Recode for Analysis.

The database’s default is to restrict analyses to malignant cases. This restriction is used by CDC’s NPCR and NCI’s SEER Program for generating most official cancer statistics. To analyze benign, borderline, or in situ cases, uncheck the “Malignant Behavior” box in the SEER*Stat Selection tab.

To create comparable analyses using a database with data from submission years 2018 and earlier—

- Uncheck the “Malignant Behavior” box in the SEER*Stat Selection tab.
- Add the following selection criteria: {Site and Morphology.Behavior recode for analysis} = 'Malignant','Only malignant in ICD-O-3','Only malignant 2010+'.

**Primary Site Variables**

Beginning in diagnosis year 2010, some lymphoma and leukemia ICD-O-3 codes were updated based on changes from the World Health Organization. The appropriate site recode variables to include these updates are Site recode ICD-O-3/WHO 2008 for all ages and International Classification of Childhood Cancer (ICCC) site recode ICD-O-3/WHO 2008 and ICCC site rec extended ICD-O-3/WHO 2008 for the childhood cancer recodes.

Consider reviewing the variable Site recode ICD-O-3/WHO 2008 before using the directly coded primary site. See more information on the SEER primary site recodes.

**Stage**

A merged variable, Merged Summary Stage, is provided to span time periods when three different staging schemes are used. The coding logic for this merged variable is—

- For NPCR-registries—
  - If a case was diagnosed in 2001, 2002, 2003, 2016 or 2017, stage at diagnosis is recorded using the SEER Summary Stage 2000 variable value.
  - If a case was diagnosed in or between 2004 and 2015, stage at diagnosis is recorded using the Derived SEER Summary Stage 2000 variable value. If the Derived SEER Summary Stage 2000 variable is blank or unstaged, and the SEER Summary Stage 2000 variable has a valid value, that value is used to populate the merged variable.
If a case was diagnosed in 2018, stage at diagnosis is recorded using the *Summary Stage 2018* variable value.

- For SEER-only registries (Connecticut, Hawaii, Iowa, and New Mexico)—
  - If a case was diagnosed in 2001, 2002, or 2003, stage at diagnosis is recorded using the *SEER Summary Stage 2000* variable value.
  - If a case was diagnosed in or between 2004 and 2015, stage at diagnosis is recorded using the *Derived SEER Summary Stage 2000* variable value.
  - If a case was diagnosed in 2016 or 2017, the best available data from either *Derived SEER Summary Stage 2000* or *SEER Summary Stage 2000* is used.
  - If a case was diagnosed in 2018, stage at diagnosis is recorded using the *Summary Stage 2018* variable value.

Users of this variable should be aware that—

- Due to changes made in the *Summary Stage 2018 Coding Manual*, for cases diagnosed in 2018,
  - the category *Regional, NOS* (code 5) is no longer used.
  - There is an artificial increase in the category *Regional by Direct Extension Only* (code 2) for brain, CNS Other, and lymphoma cases. This is because *Regional, NOS* for these cases changed from code 5 to code 2.

- *Merged Summary Stage* data are not available for testis, leukemia, or myeloma cases.

### Reporting Delay

NPCR and SEER registries annually submit all eligible years of data to CDC and NCI, respectively. As a result, cases submitted in previous years may be deleted, and new cases diagnosed in previous years may be added. The addition of new cases is called a *reporting delay*. This reporting delay may cause an appearance of decreasing trends. For example, reporting of melanoma cases diagnosed in an outpatient facility may be delayed. As a result, the trend in incident melanoma cases might superficially appear to have dropped in the most recent year.

### References


U.S. Cancer Statistics Public Use Database

Multi-Year Analyses

The database includes variables that can be used to restrict analyses to the states meeting U.S. Cancer Statistics publication criteria during the most commonly analyzed multi-year time periods, specifically—

- All years of data in the database (variable USCS0118 for diagnosis years 2001–2018).
- The most recent 10 years of data (USCS0918 for diagnosis years 2009–2018).
- The most recent 5 years of data (USCS1418 for diagnosis years 2014–2018).

If you are conducting a multi-year analysis and want to restrict it to the states that met publication criteria during each of the years, did you use variable USCS0118, USCS0918, or USCS1418 and also use the Year of Diagnosis variable to restrict to the corresponding year range on the SEER*Stat Selection tab?

- This is important for trend analyses, as the same states need to be included for each year being analyzed for comparisons.

- The Year of Diagnosis variable is used in combination with the predefined USCS variable to exclude the non-relevant years. For example, if USCS1418 is used, then Year of Diagnosis should also be restricted to diagnosis years 2014–2018 in the SEER*Stat Selection tab.

- If you would like to analyze a range of years other than those predefined variables, please contact CDC at uscsdata@cdc.gov and we will create a new variable for you.

Single-Year Analyses

If you are analyzing just one year of data, did you use the variable USCS Standard and restrict the analysis to the specific Year of Diagnosis in the SEER*Stat Selection tab?

Common Selection and Reporting Considerations

- If you are reporting state-level race, ethnicity or race/ethnicity combinations, have you suppressed data from the registries that opted out of reporting these data items? Race and ethnicity combinations can be excluded using the state race eth suppress variable; race-only or ethnicity-only suppressions should be done manually in the SEER*Stat Selection tab.

- If a user-defined primary site variable was created (rather than using the Site recode ICD-O-3/WHO 2008 variable)—
  - Did you exclude leukemias and lymphomas (9590–9992)?
  - Did you consider excluding Kaposi sarcoma (9140) and mesothelioma (9050–9055)?

For more information, see Primary Site Variables.

- If your analysis includes histology, and if appropriate for the cancer site, did you use the Diagnostic Confirmation variable to specify the analysis be limited to microscopically confirmed cases?

- If you are analyzing sex-specific cancers (such as prostate cancer or female breast cancer), did you limit the analysis to the appropriate sex to get the correct population denominator?

- When reporting rates, have you included the label "per 100,000 persons," "per 100,000 women," or "per 100,000 men"?

- Have you included citations for the—
  - Percentage of United States population coverage provided by the database?
U.S. Data Variables

U.S. Cancer Statistics Public Use Database

The following variables are available in the U.S. Cancer Statistics Public Use Database, U.S. data (2001–2018). They are listed by SEER*Stat category.

Age at Diagnosis

- Age recode with <1 year olds

Race, Sex, Year of Diagnosis, Registry, and County

- Sex
- Year of diagnosis
- Addr at DX – state
- USCS standard
- Race recode for USCS
- Program
- Region
- USCS0118
- USCS0918
- USCS1418
- Origin recode NHIA (Hispanic, Non-Hisp)

Site and Morphology

- Primary site – labeled
- Histologic type ICD-O-3 (International Classification of Diseases for Oncology, Third Edition)
- Behavior code ICD-O-3
- Grade
  - Restricted to diagnosis years 2001—2017
- Grade clinical
  - Restricted to diagnosis year 2018
- Grade pathological
  - Restricted to diagnosis year 2018
- Diagnostic confirmation
- ICD-O-3 histology/behavior, labeled
- Site recode ICD-O-3/World Health Organization 2008
- International Classification of Childhood Cancer (ICCC) site recode ICD-O-3/WHO 2008
- ICCC site recode extended ICD-O-3/WHO 2008
- Adolescent and young adult (AYA) site recode 2020
- Lymphoma neoplasm recode 2020

Stage – Local, Regional, Distant (LRD) [Summary and Historic]

- Merged summary stage
Therapy

- Rx summary – surgery primary site
  Restricted to female breast only and diagnosis years ≥2003

Extent of Disease – Collaborative Stage (CS)

- CS site-specific factor 1 (WHO Grade Classification)
  Restricted to brain/central nervous system and diagnosis years 2011—2017
- Laterality
- Merged estrogen receptor
  Restricted to female breast and diagnosis years 2004 or later
- Merged progesterone receptor
  Restricted to female breast and diagnosis years 2004 or later
- Merged HER2 summary
  Restricted to female breast and diagnosis years 2010 or later

Multiple Primary Fields

- Sequence number – central

Dates

- Year of birth
- Month of diagnosis

User-Specified

- Rural-urban Continuum 2013, grouped

Merged System-Supplied

- Alcohol-related cancers
- Human papillomavirus (HPV)-related cancers
- Obesity-related cancers
- Physical inactivity-related cancers
- Tobacco-related cancers
- State race ethnicity suppress