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|  | **The Clinical and Community Data Initiative** |
| **Sponsor:** Centers for Disease Control and Prevention  **Dept. No.:** P351  **Contract No.:** 75FCMC18D0047  **Project No.:** 37208164 | **Clinical and Community Data Initiative Prevalence Queries  Implementation Guide  for Youth and Teen Data**  **Version 1.1** |
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Record of Changes

Implementation Guide Changes

|  |  |  |  |
| --- | --- | --- | --- |
| Version | Date | Author / Owner | Description of Change |
| DRAFT | July 9, 2020 | Erin Tanenbaum / Health FFRDC | Initial Version |
| DRAFT | July 23, 2020 | Erin Tanenbaum / Health FFRDC | Edited |
| DRAFT | August 11, 2021 | Erin Tanenbaum/Health FFRDC | Added Social Determinants of Health, Glossary, and additional context |
| DRAFT | October 25, 2021 | Erin Tanenbaum and Melissa Garcia/Health FFRDC | Edited |
| DRAFT | December 23, 2021 | Erin Tanenbaum and the Health FFRDC | Edited draft |

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# Introduction

As part of the Centers for Disease Control and Prevention’s (CDC) efforts to promote health, prevent disease, injury, and disability, and prepare for emerging health threats, the Division of Nutrition, Physical Activity, and Obesity, and the Center for Surveillance, Epidemiology, and Laboratory Services partnered with the Centers for Medicare & Medicaid Services Alliance to Modernize Healthcare federally funded research and development center (Health FFRDC) on the [Clinical and Community Data Initiative (CODI)](https://www.cdc.gov/obesity/initiatives/codi/community-and-clinical-data-initiative.html). CODI will expand the ability to capture, standardize, integrate, and query existing patient-level electronic health record (EHR) and non-medical community data. To accomplish this, open-access programs were created to support research and program evaluation on prevalence of youth and teen weight-status categories per BMI percentiles, using a distributed health data network (DHDN) infrastructure. CODI will expand the traditional DHDN infrastructure[[1]](#footnote-2) to include data from both clinical data partners and non-clinical, community-based data partners.

The Health FFRDC developed open-access programs, referenced here as the CODI prevalence queries (CODI-PQ) to generate weight-status-category prevalence based on body mass index (BMI) percentiles[[2]](#footnote-3) in youth and teens, aged 2 through 19 stratified by age, sex, and geography. CODI-PQ were designed to use data from the CODI DHDN and other non-probability samples derived from EHRs.

## Background

Public health surveillance of youth and teen obesity often relies on self-report surveys by child or parent proxy, which are subject to bias. These surveys can be expensive to administer, limited in geographic specificity, and may struggle with response rates and timeliness. Data from EHRs have the potential to play a significant role in obesity population health surveillance, programs, interventions, and evaluations. While EHR data are non-probability samples of health outcomes among the care-seeking population, the data – measurements, diagnoses, observations, prescriptions, and procedures – offer an opportunity to provide decision makers with detailed, timely, and accurate information at the local, state, and regional levels for surveillance, research, and evaluation.

Population estimates for surveillance can be obtained by applying statistical weights, imputation, and suppression criteria to EHR data. Applying statistical weights can reduce potential biases introduced by the EHR sampling. CODI-PQs provide a tool for calculating prevalence estimates by weighting user provided EHR or DHDN data for a target population. The CODI-PQ algorithms can generate stable prevalence estimates at state, county, and zip code geographies from EHR data, depending on the data provided by the use, with the aim to improve access to timely data on local disease burden to inform prevention and other public health activities.

## Purpose

The purpose of the CODI-PQ Implementation Guide for Youth and Teen Data is to provide a guide for CODI data partners[[3]](#footnote-4) or data users to run the CODI-PQ. The Implementation Guide covers the following:

* CODI-PQ data inputs and link population data (pre-processing)
* Generating results in CODI-PQ
* Understanding the CODI-PQ results
* Methodological details

## Scope

CODI-PQ were created and tested with IQVIA's Ambulatory Electronic Medical Record (AEMR-US)[[4]](#footnote-5) data and synthetic data generated for CODI using Synthea.™ [[5]](#footnote-6) CODI-PQ require patient level records for children ages 2 through 19. Each record must include year of medical encounter, demographic information, and BMI percentile category. Patient-level records must include residential address information at the level of state, county, and zip code, or at the level of state and the ZIP Code Tabulation Area’s first three digits (ZCTA-3). CODI-PQ leverages population counts from the American Community Survey. CODI-PQ assumes end users include all available EHR data for a geography and/or subpopulation.

All statistical programs described in this document were created and tested using SAS 9.4 software (SAS Institute, Inc., Cary, North Carolina). The guidance provided in this document is implemented through open-access programs.

## Audience

The audience for this IG is CODI data partners and data users. The user should have a working knowledge of SAS language and macros. Those interested in statistical analysis details used in CODI-PQ can refer to Appendix A for more information. Technical staff preparing datasets for CODI-PQ can refer to Appendices C and D for detailed descriptions of the format required for input data. Explanation of CODI-PQ results can be found in Appendix E.

## Document Organization

This document is organized as follows:

|  |  |  |
| --- | --- | --- |
| **Section** |  | **Purpose** |
| **Section 1** | Introduction | Provides a background for CODI-PQ |
| **Section 2** | User’s Guide | Provides a general guide for users |
| **Appendix A** | Analysis Details | Provides detailed description of analysis |
| **Appendix B** | Social Determinants of Health | Provides methodology for prioritization of social determinants of health for statical weighting |
| **Appendix C** | ACS File Layouts | Table outlining the required ACS input file layouts |
| **Appendix D** | EHR File Layouts | Table outlining the required EHR input file layouts |
| **Appendix E** | CODI-PQ-GEO3 Example SAS Programs | Provides example SAS program for generating results by county or ZCTA-3 |
| **Appendix F** | CODI-PQ Results Example | Provides CODI-PQ results data dictionary and example results |
| **Appendix G** | State FIPS codes | Provides list of state abbreviations |
| **Appendix H** | Glossary | Defines terms used in this document |
| **Appendix I** | Abbreviations and Acronyms | Defines acronyms used in this document |
| **Appendix J** | Bibliography | Lists sources used in preparing this document |

# User’s Guide

The User’s Guide section describes:

Organizing the CODI-APQ folders and properly preparing the data based on specifications are key to a successful implementation.

## CODI Concept

Figure 1 shows how CODI users (e.g., researchers, community-based program evaluators) interact with the data coordinating center, which distributes their research queries to data partners. The data coordinating center assembles the results into longitudinal records, which are sent to the CODI end-users. CODI end-users use the patient-level longitudinal records to create prevalence estimates with CODI-PQ. Additional CODI details can be found in the documentation available through GitHub at https://github.com/mitre/codi.

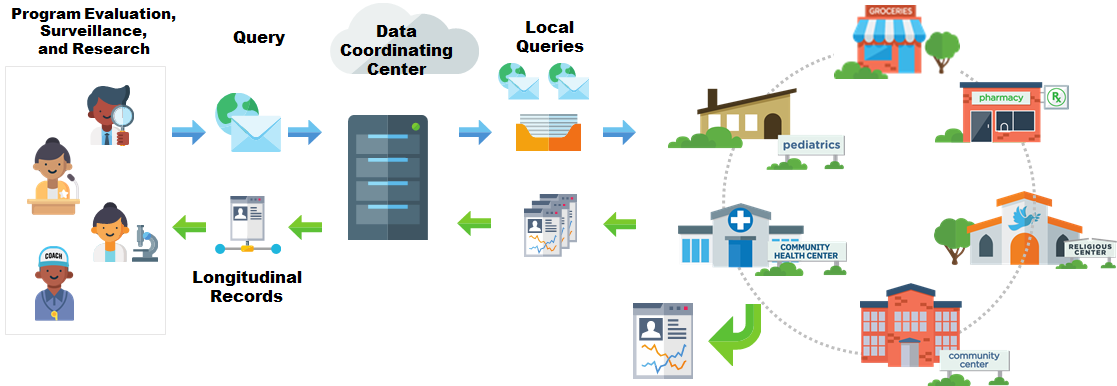


Figure 1. Data Partners with a Common Data Coordinating Center

## About CODI-PQ

CODI-PQ is a set of programs that calculate youth and teen BMI percentile prevalence estimates from a non-probability sample[[6]](#footnote-7) of EHRs. CODI-PQ is divided into two parts: pre-processing, and the prevalence query. First, patient data is imported into SAS, linked to the American Community Survey (ACS), and race imputation is conducted in the pre-processing steps (PRE\_PROCESSING\_GEO3). Next, within the prevalence query steps, patients are selected based on user specifications, statistically weighted, variance estimates are calculated, results are suppressed (if needed), and prevalence results are output (CODI\_PQ\_GEO3).

CODI-PQ programs require careful review of the methodological details (described in appendices), EHRs supplied by the user, and ACS data from 2019 supplied by the Health FFRDC[[7]](#footnote-8). Results can be calculated for a specific geography (e.g., state, state and county, state, and ZCTA-3), subpopulation (e.g., age group, sex, race), or geography and subpopulation.

Results are suppressed[[8]](#footnote-9) if the user selects a geography or subpopulation with an insufficient number of patients for statistical weighting or if results do not meet NCHS suppression criteria. The CODI-PQ programs user should have a working knowledge of SAS language and macros to select the population of interest, execute CODI-PQ, and review the SAS log.

The programs described in the User’s Guide are designed to:

* Impute race for youth and teens missing race information (optional)
* Calculate statistical weights with an EHR non-probability sample
* Calculate age-adjusted prevalence results (optional)
* Calculate youth and teen BMI percentile prevalence by BMI percentile, including:
* **Underweight**: BMI less than the 5th percentile
* **Healthy Weight**: BMI between the 5th and 85th percentile
* **Overweight**: BMI between the 85th and 95th percentile
* **Obesity**: 95th percentile to less than 120 percent of the body mass index (BMI) value for the 95th percentile
  + **Severe Obesity**: 120 percent or greater of the BMI value for the 95th percentile
* Suppress prevalence estimates based on the National Center for Health Statistics (NCHS) Data Presentation Standards for Proportions

## SAS Setup

All statistical programs described in the User’s Guide were created and tested using SAS 9.4 software (SAS Institute, Inc., Cary, North Carolina) in a Windows environment. CODI-PQ requires the following SAS features:

* BASE SAS
* SAS STAT
* The ability to import a file from csv into SAS
* The ability to export a file from SAS into csv

## Step-By-Step Process to Run CODI-PQ

The four-step process to run the CODI-PQ is outlined below:

Figure 2. CODI-PQ Process

### STEP 1: Download and Unzip CODI-PQ-master.zip File

Access CODI-PQ programs on GitHub: <https://github.com/NORC-UChicago/CODI-PQ>.

Select and download “CODI-PQ-GEO3-master.zip.” Note that “GEO3” refers to the program’s options to estimate prevalence at the county or ZCTA3 level.

Use Winzip or other available software to unzip the files. Be sure the option is selected to unzip both files and folders and **preserve the folder names (Full Path Information set to on)**.

Unzip with Folders
After downloading CODI-PQ, unzip the SAS programs, and preserve the folder names by setting Full Path Information on. CODI-PQ includes “Quickstart” programs that automatically execute additional programs based on the folder structure in the zipped file and folders.


CODI-PQ-GEO3’s folder structure is shown in the figure below. Note that folders and subfolders have been created and structured in a way to make it easier for the user to organize the input and results files.

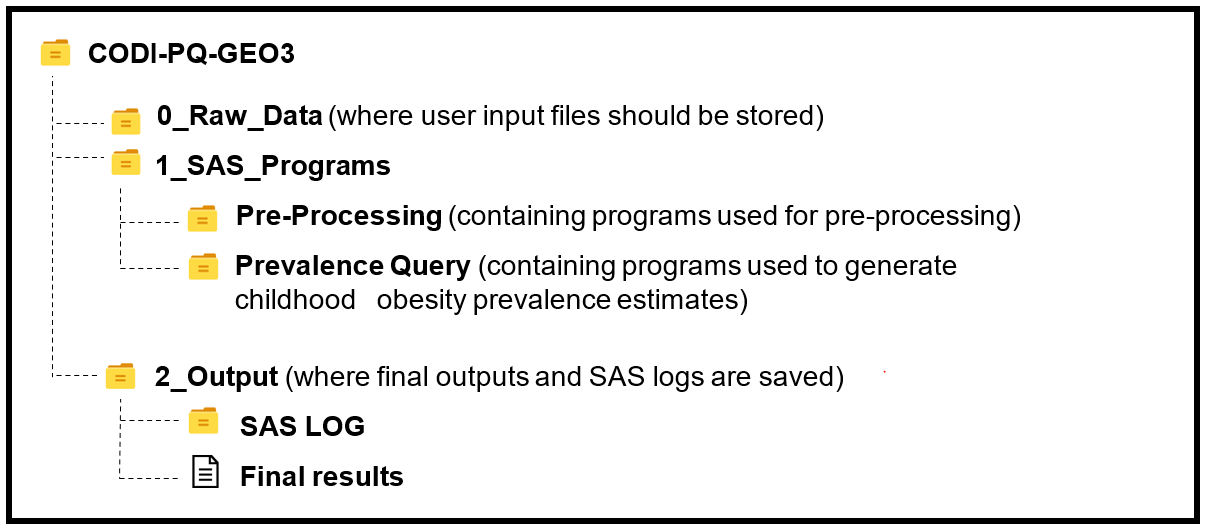


Figure 3. CODI-PQ-GEO3 Folder Structure[[9]](#footnote-10)

### STEP 2: Obtain Input Files and Store Them in the ‘0\_Raw\_Data’ Folder

Required input files include:

1. **ACS data file** (downloaded from <https://sft.mitre.org/#/folder/6281923>) The 2019 ACS data can be downloaded from the Health FFRDC Secured File Transfer Protocol (SFTP) site to ensure consistency with the models embedded into the SAS programs. For variable names, variable order, and a description of the file, see Appendix C.
2. **EHR file** supplied by the end user in comma separated values (.csv). The EHR file must:

* Contain all variables in the order (sequence) expected. Variable names and order can be found in Appendix D.
* Contain valid variable values as anticipated. Variable values can be found in Appendix D.
* Have a unique identifier for all patients and the identifier is consistent between years.
* Include a **maximum** of one record per patient per year. The user can choose the record kept so it aligns with analysis goals. For testing purposes, the event date closest to July 1 of each year was kept prior to executing pre-processing.
* Include a valid height and weight value obtained on the same day which was used to calculate the BMI percentile for all patients (underweight, healthy weight, etc.)
* Have a geographic location of the patient’s residency either as the state and ZCTA-3 or the state, county, and ZIP code.
* Users may also wish to reconcile demographic characteristics across years for each patient, including:
  + Sex
  + Race

Additional variables on the file must be included even though their inclusion in the prevalence query is optional. If the file does not include these variables, researcher can add two columns to the end of their file with blank values. These variables include:

* A sickle cell disease[[10]](#footnote-11) indicator
* A pregnancy indicator

A full description of the EHRs file format is available in [Appendix](#EHRfilelayouts) D.

### STEP 3: Link Population (Pre-Processing)

Open the “Quickstart-Pre\_Processing\_CODI\_PQ\_GEO3” SAS program stored in “\1\_SAS\_Programs” and change the selection per the steps outlined in the tables below.

Note that the pre-prevalence program should be submitted once and only once per file. As such, include the start and end years for the full file. The programs also impute the race of those with unknown race and each time the program is submitted, new imputed race values are created and stored for each patient. For consistency, we encourage submitting the pre-processing programs only once for each EHR file. If additional data is later processed for the same patient, we encourage 1) replacing the race of all patients who were imputed before but their race is now known, 2) keeping the imputed race value consistent for patients who were imputed before and their race value is still unknown.

A new folder (“\2\_Output\Pre-Processed\_…”) will be created upon completion of the programs. In this folder, two SAS7bdat files (user input ACS file and pre-processed CODI file) will be generated. Once pre-processing is complete, the user can submit an unlimited number of prevalence queries using the same pre-processed file each time.

Table 1. Change Specifications, Pre-Processing Steps

| **Order** | **Description** | **Details** |
| --- | --- | --- |
| **1** | Open the Pre-processing Quickstart program. | The Quickstart program is stored in the folder:  "..\1\_SAS\_Programs” |
| **2** | Edit the SAS program within “SECTION 1: Input Folder and file names.” | Follow the SAS programs and update the macro variable specifications, in particular (see Table 2) |

Table 2. Change SAS Specifications, Section 1

|  |  |  |
| --- | --- | --- |
| **SAS Macro Variable** | **Details** | **Example** |
| ROOT\_PRE | The core folder name where CODI-PQ-GEO3 are saved (see part 2.1.1). The SAS Programs folder and all other folders and files are stored in this directory. | %let ROOT\_PRE = P:\CODI-PQ-master; |
| PRE\_DEST | Following the example, the results from pre-processing will be generated and stored in folder P:\CODI-PQ-master \2\_Output\Pre\_Processed\_***CODI\_PQ\_GEO3*** | %let PRE\_DEST = CODI\_PQ\_GEO3; |
| ACS\_FILENAME | The American Community Survey file name from part 2.4.2. The file is in csv format. Do not include the extension in the file name. Important: the csv file must have all variables in the order specified, with the correct variable name, and with expected values. See C.1. | %let ACS\_FILENAME = ACS\_State\_COUNTY; |
| EHR\_FileNAME | The youth and teen level EHR file name (file described in 2.4.2). Do not include the extension. Important: the csv file must have all variables in the order specified, with the correct variable name, and with expected values. See D.1.  For the example, the user will have stored their raw data in:  P:\CODI-PQ-master\0\_Raw\_Data\**EHR\_csv\_filename**.csv | %let EHR\_FileNAME = EHR\_csv\_filename; |
| EHR\_PRE\_OUT | Optional, user can name the pre-processing output file (ACCEPTABLE VALUES: file name (no punctuations)).  The example would be stored in: P:\CODI-PQ-master \2\_Output\Pre\_Processed\_*CODI\_PQ\_GEO3*\**SAVES\_EHR\_HERE** | %let EHR\_PRE\_OUT = SAVES\_EHR\_HERE; |
| LOG\_NAME\_PRE | The name of the SAS log file. Quickstart\_Pre\_Processing\_CODI\_PQ. Users have the option to rename the log file name before it is created.  The example SAS log would be stored in: P:\CODI-PQ-master \2\_Output\SAS LOG\***LogName***<Date and Time>.log. Note, the program automatically includes the date and time in all log file names\*/ | %let LOG\_NAME\_PRE = LogName; |

Table 3. Change Specifications, Pre-Processing Steps, Continued

| **Order** | **Description** | **Example or Detail** |
| --- | --- | --- |
| **3** | Section 2, edit the SAS program within “SECTION 2: Beginning and End Year of longitudinal EHR data.”  **In pre-processing, the start and end year includes all years within your patient level file (e.g., 2014 – 2019)**. In contrast, when generating prevalence estimate results, only include the specific year(s) requested for analysis (Part 4, e.g., 2019 if prevalence estimates with 2019 records are requested). | .  /\*\*\*/ %LET BEGIN\_YEAR = **2014**;  /\*\*\*/ %LET END\_YEAR = **2019**; |
| **4** | Edit the SAS program within “Section 4: County or ZCTA3 data (REQUIRED)”  Edit with a Y or N. For example, Y for County level data, **N for ZCTA3 level data**.  The ACS data file and EHR data file must have a variable GEO3 which includes three digits for either ZCTA3 or County codes. Both files must have equivalent geographic types. Thus, the files can have either have State+County or State+ZCTA3, but not both. | %LET COUNTY=**N**; |
| **5** | Save the Quickstart program. | SAS encourages saving all files before submitting the program. |

Table 4. Pre-Processing CODI-PQ Program Execution Steps

| Order | Description | Details |
| --- | --- | --- |
| 1 | Submit the Quickstart program. | Submit the Quickstart program. The program completes all linkage and pre-processing tasks by looping through all needed SAS programs automatically. |
| 2 | Review the log. | Review the log for possible errors including words such as error, and uninitialized. Assuming no errors, continue to Part 4. In the event of errors, reassess the location of the files and the file formats. |

### STEP 4: Generate Prevalence Estimate Results

Open the “Quickstart-CODI\_PQ\_GEO3” SAS program stored in “\1\_SAS\_Programs” and change the selections outlined in the tables below.

The final results (CODI-PQ results) will be generated in .csv or Excel format and saved in “\2\_Output.” Appendix F provides an example of the results. Note that results are for the group of patients selected by the user. To calculate results for multiple geographic or demographic characteristics (e.g., first for youths ages 2 to 4 and then for teens aged 18 and 19), the user will need to update and execute the programs multiple times.

Note: the age ranges, sex, and races selected must match the data on the EHRs. For example, if all age ranges are selected by the user and the file has youths ages 2 to 12 but does not have teens aged 13 to 19, then the program will fail with an error message caused by insufficient sample size for patients aged 15 to 19.

Table 5. Change Specifications, Processing Steps

|  |  |  |
| --- | --- | --- |
| **Order** | **Description** | **Details** |
| **1** | Open the Quickstart program. | The Quickstart program is stored in the folder:  "..\1\_SAS\_Programs” |
| **2** | Edit the SAS program within  “SECTION 1: Folder and file names”  “SECTION 2: Subset data based on specifications INCLUDING YEAR, GEOGRAPHY, STATE, STATE/ZCTA3, or STATE/COUNTY” | Follow the SAS programs and update the macro variable specifications. |

Table 6. Change Specifications, Processing Steps

| **SAS Macro Variable** | **Details** | **Example** |
| --- | --- | --- |
| **SECTION 1: Folder and file names** |  |  |
| ROOT\_PQ | The core folder name, same as in pre-processing. | %let ROOT\_PRE = P:\CODI-PQ-master; |
| PRE\_DEST | The value from pre-processing quickstart variable pre\_dest. See 2.4.3, Table 2.  Following the example, the results from pre-processing were previously generated and stored in P:\CODI-PQ-master\2\_Output\ Pre\_Processed\_**CODI\_PQ\_GEO3** | %let PRE\_DEST = CODI\_PQ\_GEO3; |
| EHR\_PRE\_OUT | The youth and teen level EHR file from pre-processing. The example was stored in: P:\CODI-PQ-master \2\_Output\Pre\_Processed\_*CODI\_PQ\_GEO3*\**SAVES\_EHR\_HERE** | %let EHR\_PRE\_OUT = SAVES\_EHR\_HERE; |
| LOG\_NAME | The name of the resulting SAS log. Users have the option to rename the log file name before it is created.  Following the example syntax, the SAS log will be stored in: P:\CODI-PQ-master\2\_Output \SAS LOG\**LogName**<Date and Time>.log. Note, the program automatically includes the date and time in all log file names. | %let LOG\_NAME = LogName; |
| FileOUT\_Name | The prefix for the resulting .csv or Excel file.  Following the example syntax, the .csv or Excel file will be stored in: P:\CODI-PQ-master\ \2\_Output\**File\_name**<Date and Time>.csv. Note, the program automatically includes the date and time in all results file names. | %LET FileOUT\_Name = File\_name; |
| **SECTION 2: Subset data based on specifications INCLUDING YEAR, GEOGRAPHY, STATE, STATE/ZCTA3, or STATE/COUNTY** |  |  |
| BEG\_YEAR | Subsets the prevalence to medical encounters in this year for BMI prevalence. The prevalence will include adult EHRs from this year and after.  Acceptable values must be a 4 digit year and the year must be present on the user’s EHR file. | /\*\*\*/ %LET BEG\_YEAR = 2016; |
| END\_YEAR | Subsets the prevalence to medical encounters through this year for BMI prevalence. The prevalence will include adult EHRs from this year and before.  Acceptable values must be a 4 digit year and the year must be present on the user’s EHR file.  If the end year is not equal to the beginning year, then each patient’s most recent record will be kept. Patients are not included multiple times within analytic results. | /\*\*\*/ %LET END\_YEAR = 2018; |
| ALL\_STATES | Includes all states and the District of Columbia (national estimate) in the prevalence based on the geographic location of the youth or teen. If ALL\_STATES = N; then by default the program will subset the prevalence based on the individual state or state+GEO3 values specified (in future step). Note that the EHR file must have sufficient sample size in all states if set to yes (Y). | /\*@Note: Include all geographical locations in file? (ACCEPTED VALUES: Y/N) \*\*\*/ |
| ALL\_AGES | Subsets the prevalence based on the age of the youth or teen. The user may either select to include all youth and teens aged 2 to 19 (Y) or alternatively may select age groups (N). Note: if ALL\_AGES = Y; then by default the program will include all youth and teens aged 2 to 19. If ALL\_AGES = N; then by default the program will subset the prevalence based on the individual age ranges selected (in a future step). | /\*\*\*/ %LET ALL\_AGES = Y; |
| ALL\_SEXES | Subsets the prevalence based on the sex of the youth or teen. The user may either select to include all male and female youth and teens (Y) or alternatively may select either males or females (N). Note: if ALL\_SEXES = Y; then by default the program will include both males and females. If ALL\_SEXES = N; then by default the program will subset the prevalence based on the individual sex(es) selected (in a future step). | /\*\*\*/ %LET ALL\_SEXES = Y; |
| ALL\_RACES | Subsets the prevalence based on the race of the youth or teen. The user may either select to include all races (Y) or alternatively may select race(s) (N). Inclusion or exclusion of imputed race is not impacted by the choice made in this step. Note: if ALL\_RACES = Y; then by default the program will include all races (White, Black, Asian, Other). If ALL\_RACES = N; then by default the program will subset the prevalence based on the individual races selected (in a future step). | /\*\*\*/ %LET ALL\_RACES = Y; |
| **SECTION 3: Additional Flags** |  |  |
| ACSCOUNTY | Specifies the geographic level of the ACS data. | /\*\*\*/ %LET ACSCOUNTY = N; /\*@Note: Is the ACS data at the county or ZCTA3 level? (ACCEPTABLE VALUES: Y for County level data, N for ZCTA3 level data) \*\*\*/ |
| INCLUDE\_PREGNANCY | Subsets EHR records based on pregnancy flag.  **If set to yes (Y), then patients will be included regardless of pregnancy status.** If set to no (N,) then only patients with a pregnancy status set to no (0) will be included. | /\*\*\*/ %LET INCLUDE\_PREGNANCY = Y; |
| SAMPLE\_CHECK | Executes an optional review of the EHR counts by age, race, and sex based on user defined criteria. All demographic categories selected by the user (e.g., sex male, etc.) will be displayed in the SAS output or results window. Each factor will include the factor, value (e.g., sex, male) and either “Sample Size Is Insufficient” if n <20 or “Sample Size Is Sufficient. | /\*\*\*/ %LET SAMPLE\_CHECK = Y; |

Table 7. Change Specifications, Processing Steps, Continued

|  |  |  |
| --- | --- | --- |
| **Order** | **Description** | **Details** |
| **3** | Edit the SAS program within  “SECTION 4: Only complete section 4 for any "N" values listed in section 2“  “SECTION 5: Methodological option selections” | **Review specifications below.** |

Table 8. Change Specifications, Processing Steps, Continued

| **SAS Macro Variable Category** | **Details** | **Example** |
| --- | --- | --- |
| **SECTION 4: Only complete section 4 for any "N" values listed in section 2** |  |  |
| If ALL\_STATES = N | GEO\_GROUP informs the program the level of geography in the EHR and ACS data as well as in the GEO\_LIST macro variable. The level of geography must match in all three locations.  GEO\_LIST subsets the EHR used in prevalence results based on the location of the patients.  GEO\_GROUP can take the value of a) STATE, b) ZCTA3, or c) COUNTY. Syntax for all three scenarios is described and shown in examples. Of note, values should be surrounded by single quotes and comma delimited if more than one geography is to be included in the results.  If ALL\_STATES is set to yes (Y), then the SAS program does not review the user’s responses to the GEO\_LIST or GEO\_GROUP.  Example one selects patients in Colorado and Delaware.  Example two selects patients living in Virginia, within the ZIP-3 of 221 as well as patients living in Michigan with the ZIP-3 of 486.  Example three selects patients living in Virginia, within Fauquier County and patients living in Virginia within Fairfax County. | /\*Example 1:\*/  /\*\*\*/ %LET GEO\_GROUP = STATE; /\*\*\*/ %LET GEO\_LIST = %STR('08', ‘10’);  /\*Example 2:\*/  /\*\*\*/ %LET GEO\_GROUP = ZCTA3; /\*\*\*/ %LET GEO\_LIST = %STR('51221', ‘26486’);  /\*Example 3:\*/  /\*\*\*/ %LET GEO\_GROUP = COUNTY; /\*\*\*/ %LET GEO\_LIST = %STR('51061', ‘51059’); |
| If ALL\_AGES = N and CO\_OCCURRING = N; | If ALL\_AGES is set to no and CO\_OCCURRING is set to no, the age macros (2-4, 5-9, 10-14, 15-17, 18-19) subset the prevalence based on the age of the youth or teen and the responses to each individual age macro. Note that if ALL\_AGES is set to yes, then the SAS program does not review the age-specific macros. | %LET WGT\_AGE\_2\_4 = N;  %LET WGT\_AGE\_5\_9 = N;  %LET WGT\_AGE\_10\_14 = Y;  %LET WGT\_AGE\_15\_17 = Y;  %LET WGT\_AGE\_18\_19 = Y; |
| If ALL\_RACES = N; | If ALL\_RACES is set to no, the race macros (White, Black, Asian, Other) subset the prevalence based on the race or imputed race of the youth or teen and the responses to each individual age macro. Note that if ALL\_RACES is set to yes, then the SAS program does not review the race-specific macros. | %LET RACE\_WHITE = N;  %LET RACE\_BLACK = Y;  %LET RACE\_ASIAN = Y;  %LET RACE\_OTHER = Y; |
| If ALL\_SEXES = N; | If ALL\_SEXES is set to no, the sex macros (male, female) subset the prevalence based on the sex of the youth or teen and the responses to each individual sex macro. Note that if ALL\_SEXES is set to yes, then the SAS program does not review the sex-specific macros | %LET SEX\_MALE = N;  %LET SEX\_FEMALE = Y; |
| **SECTION 5: Methodological option selections** |  |  |
| IMP\_RACES | If IMP\_RACES is set to yes (Y), then the program includes youth and teens with imputed race values. Otherwise, if IMP\_RACES is set to no (N), then the patients with imputed races are excluded. | %LET IMP\_RACES = Y; |
| AGE\_ADJ | If AGE\_ADJ is set to yes (Y), then the program generates age adjusted prevalence and standard errors. Otherwise, if AGE\_ADJ is set to no (N), age adjusted prevalence is not generated. Crude and weighted estimates are generated by default. | %LET AGE\_ADJ = Y; |

Table 9. Change Specifications, Processing Steps, Continued

|  |  |  |
| --- | --- | --- |
| Order | Description | Details |
| 4 | Save the Quickstart program. | It is encouraged to save the Quickstart **program before submitting in SAS.** |

Table 10. CODI-PQ Execution Processing Steps

| **Order** | **Description** | **Details** |
| --- | --- | --- |
| **1** | Submit CODI-PQ Quickstart program. | Submit the Quickstart program. The program completes all tasks within the data sets and proc statements in the Quickstart program and moves to the next SAS program automatically through an include statement. |
| **2** | Review the log. | Review the log for possible errors including words such as error, and uninitialized. Assuming no errors, continue to the next step. In the event of errors, reassess the location of the files and the file formats. |
| **3** | Review the results. | Review the results for possible data suppression or errors. Consider a statistical review based on the NCHS data presentation standards. In the event of errors reassess the choices described above and re-submit. In the event of data suppression, consider expanding your selection criteria and re-submit. For example, if prevalence results cannot be created with a single year, consider using two or three years of data[[11]](#footnote-12). |

### Review BMI Percentile Prevalence Results

CODI-PQ generates prevalence outputs as a csv file. Table 11 provides an overview of the variables included for BMI percentile prevalence. Note, descriptive information about CODI-PQ user inputs, error codes, sources of technical documentation, caveats, and a possible citation begins with the rows labeled Order 3 and beyond. The exact notes displayed vary.

Table 11. CODI-PQ BMI Percentile Prevalence Results Data Dictionary

| Column | Description |
| --- | --- |
| Order | Row order |
| Weight Category | The weight category based on BMI percentile. |
| Sample | The observed (or unadjusted, or crude) count of youth and teens in the study population. |
| Population | The weighted (or adjusted) count of the study population. |
| Crude Prevalence | The observed (or unadjusted, or crude) prevalence in the study population. |
| Crude Prevalence Standard Error | The observed (or unadjusted, or crude) standard error in the study population. |
| Weighted Prevalence | Prevalence based on weighted counts. A sample weight is assigned to each sampled patient. It is a measure of the number of youth and teens in the population represented by that sample patient. See implementation guide, Appendix A. Sample Weights for more information. |
| Weighted Prevalence Standard Error | Standard error based on weighted counts. See implementation guide, Appendix A. Variance for more information. |
| Age-Adjusted Prevalence | Prevalence based on weighted, age-adjusted counts (optional). See implementation guide, Appendix A. Age Adjustment for more information. |
| Age-Adjusted Prevalence Standard Error | Standard error based on weighted, age-adjusted counts. See implementation guide, See implementation guide, Appendix A. Age Adjustment for more information. |

## Additional Details for Users

Further detail on file layouts for input and results is provided in the following appendices:

* Appendix C – ACS File Layouts
* Appendix D – EHRs File Layouts
* Appendix E – CODI-PQ-GEO3 Example SAS Programs
* Appendix F – CODI-PQ Results
* Appendix G – State FIPS Codes

###### Analysis Details

Age Adjustment

Data are age-adjusted to eliminate differences in observed results that result from differences in the age distribution of the population among geographies. The projected 2000 U.S. population was used as the standard population.[[12]](#footnote-13) The specific age groups used for age adjustment are 2 to 4 years, 5 to 14 years, and 15 to 19 years. Age-adjusted values may differ from weighted values even though age is used within the weighting program since the age distribution within a geography (GEO3) may differ from the nation.

Age adjustment, using the direct method, is the application of age-specific results in a population of interest to a standardized age distribution to eliminate differences in observed results that result from age differences in population composition. This adjustment is usually done when comparing two or more populations at one point in time or one population at two or more points in time.

Age-adjusted proportions are calculated by the direct method as follows:

where mi = measure of the proportion in age group *i* in the population of interest, pi = standard population in age group *i*, and n = total number of age groups over the age range of the age-adjusted prevalence.

Age adjustment by the direct method requires use of a standard age distribution. The standard for age adjusting proportions for data occurring after year 2000 is the year 2000 projected U.S. resident population.

Table 12. Projected Year 2000 U.S. Population Proportion Distribution by Age for Age Adjusting

|  |  |
| --- | --- |
| Age | Proportion Distribution (weights) |
| All ages\* | 1 |
| 2 to 4 | 0.1605 |
| 5 to 9 | 0.2796 |
| 10 to 14 | 0.2815 |
| 15 to 17 | 0.1659 |
| 18 and 19 | 0.1123 |

\*Figure is rounded up instead of down to force total to 1.0.

Age-adjusted prevalence results and standard errors will typically be similar or identical to the weighted prevalence and standard errors. Age-adjusted results may differ from weighted results if one or more age group weighting cell was aggregated.

Body Mass Index

Body mass index (BMI) is a patient’s weight in kilograms divided by the square of height in meters. A high BMI can be an indicator of high body fatness. BMI can be used to screen for weight categories that may lead to health problems, but it is not diagnosis of a patient’s body fatness or health.

For youth and teens, BMI is age- and sex-specific and is often referred to as BMI-for-age. In youth and teens, a high amount of body fat can lead to weight-related diseases and other health issues. Being underweight can also put patients at risk for health issues.

Youth and teen BMI Percentile categories are described in section A.11.

For more information, see: <https://www.cdc.gov/healthyweight/assessing/bmi/childrens_bmi/about_childrens_bmi.html>.

Data Sources (Inputs)

This document provides an implementation guide for CODI-PQs on youth and teen data. Required input files are the following:

* EHRs (data in csv format, provided by user) provided by the user
* American Community Survey (ACS) data file (provided by the Health FFRDC[[13]](#footnote-14))

CODI-PQ are intended for use with all available EHRs for a geography or subpopulation. The programs were created and tested with the Ambulatory Electronic Medical Record (AEMR)[[14]](#footnote-15) data and synthetic data generated for CODI using Synthea.[[15]](#footnote-16) The guide provided in this document is implemented through open-access programs.

The programs were tested using EHRs and AEMRs. Both provide a non-probability sample of longitudinally linked patients’ medical records from within the U.S. CODI-PQ subsets the file to youth and teens aged 2 to 19 years of age for BMI percentile prevalence. The programs assume a maximum of one record per year per patient. Data should include patient identifiers that link medical encounters to demographic and geographic characteristics including year of birth, race, ethnicity (when race is not available), sex, state, and either county or the first three digits of the ZIP Code Tabulation Area (ZCTA-3)[[16]](#footnote-17) associated with the patient’s address. Patients are excluded from the analysis if their state and county or ZCTA-3 does not exist or if the ACS estimated population count within their county or ZCTA-3 equals 0.

Testing of CODI-PQ included EHRs pre-processed using ‘growthcleanr.’ The ‘growthcleanr’ package is a publicly available program for identifying biological implausible height and weight measurements in longitudinal files at <https://github.com/mitre/growthcleanr-web>. The program evaluates data against published growth trajectory charts for youth, teens and adults and flags measurements for plausibility ([Daymont et al., 2017](#_ENREF_9)).

To statistically weight EHRs to the general population, the 2015-2019 American Community Survey (ACS) 5-year, population estimates by age, race, sex, and community educational attainment are used. Population counts are available by state and county or state and ZCTA-3.

Testing Data

A large sample of EHRs was used to build CODI-PQ. The sample was compared to Census estimates[[17]](#footnote-18) to determine possible biases associated with the sample. Similar biases maybe in other EHRs, although users are encouraged to perform a similar analysis.

To obtain reliable BMI percentile results from EHRs, the data should be free of potential bias. BMI percentile prevalence results using incomplete EHRs could be biased if there are systematic differences in characteristics associated with BMI percentile between measured and non-measured patients. It is necessary to identify the factors leading to missingness (lack of weight and height measurements) and apply strategies, such as inverse-probability weighting (IPW) or imputation to obtain corrected results.

When using IPW, EHRs are weighted by the inverse of their probability of being sampled. The validity of IPW relies on including variables associated with missingness. In the same context, imputation generates unbiased results for the variables of interest through a program that replaces each missing value with a plausible value, thus creating complete data sets.

The purpose of reviewing the sample distribution to the Census population distribution was to review possible selection bias associated with missing weight and height data.

Table 13. American Community Survey Census Sample by Year for 2015-2019

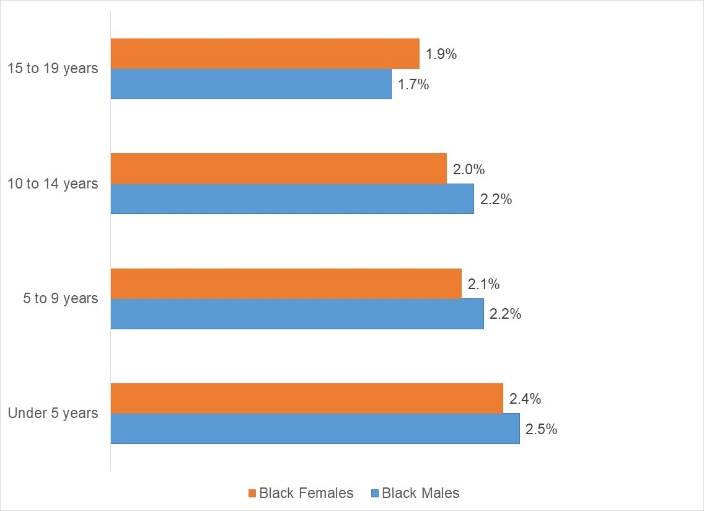
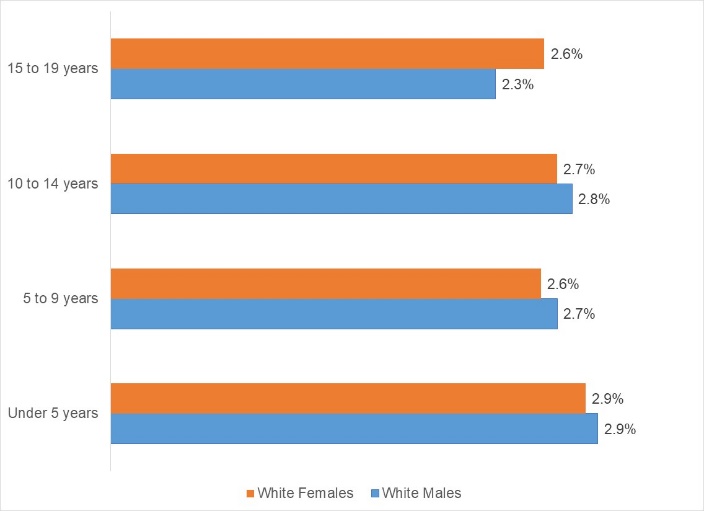
for Youth and Teens Aged 2 to 19

|  |  |  |  |  |
| --- | --- | --- | --- | --- |
| 2015 | 2016 | 2017 | 2018 | 2019 |
| 2,433,287 | 2,395,712 | 2,259,231 | 1,986,787 | 1,703,432 |

In 2015, there were 2,433,287 patients ages 2 to 19 in our sample EHRs file with plausible height and weight values from a total 74,123,130 patients living in the U.S. of that age, representing 3.3% of the population. This count of patients decreased with time to 1,703,432 patients in 2019 out of 74,012,765 patients living the U.S., representing 2.3% of the population. Weight and height measurements data stored in EHRs and pre-processed by growthcleanr were used to calculate the percentages by age with at least one measurement within a one-year period. The greatest percentages of measurements were seen for 2-year-olds (3.87%) and the lowest percentages of measurements were seen for 19-year-olds (2.61%). Counts were also compared by ACS age groups including aged 2 to 4 years, 5 to 9 years, 10 to 14 years, 15 to 17 years, and 18 to 19 years old. The lowest percentages of measures were seen for teens 18 to 19 years old (2.80%) and youth 5 to 9 (3.25%), and the greatest percentages of measures were seen for 2- to 4- year-olds (3.57%) and 10- to 14- year-olds (3.34%). This pattern remained consistent through 2019.

Table 14. Comparison of Patients with Plausible Weight and Height Data in the Electronic Health Records and the U.S. Population by Age, 2015

|  |  |  |  |
| --- | --- | --- | --- |
| **Age** | **Census 2015** | **EHRs 2015** |  |
| 2 years | 3,966,321 | 153,465 | 3.87% |
| 3 years | 3,974,351 | 135,299 | 3.40% |
| 4 years | 4,020,292 | 138,829 | 3.45% |
| 5 years | 4,017,589 | 143,688 | 3.58% |
| 6 years | 4,017,388 | 134,812 | 3.36% |
| 7 years | 4,145,872 | 129,926 | 3.13% |
| 8 years | 4,165,033 | 130,063 | 3.12% |
| 9 years | 4,130,887 | 127,399 | 3.08% |
| 10 years | 4,118,721 | 125,318 | 3.04% |
| 11 years | 4,126,769 | 136,027 | 3.30% |
| 12 years | 4,096,602 | 145,489 | 3.55% |
| 13 years | 4,080,092 | 139,636 | 3.42% |
| 14 years | 4,182,193 | 141,656 | 3.39% |
| 15 years | 4,247,757 | 143,304 | 3.37% |
| 16 years | 4,181,741 | 138,049 | 3.30% |
| 17 years | 4,189,329 | 133,084 | 3.18% |
| 18 years | 4,210,553 | 126,365 | 3.00% |
| 19 years | 4,251,640 | 110,878 | 2.61% |
| Total | 74,123,130 | 2,433,287 | 3.28% |
| 2 to 4 years | 11,960,964 | 427,593 | 3.57% |
| 5 to 9 years | 20,476,769 | 665,888 | 3.25% |
| 10 to 14 years | 20,604,377 | 688,126 | 3.34% |
| 15 to 17 years | 12,618,827 | 414,437 | 3.28% |
| 18 to 19 years | 8,462,193 | 237,243 | 2.80% |



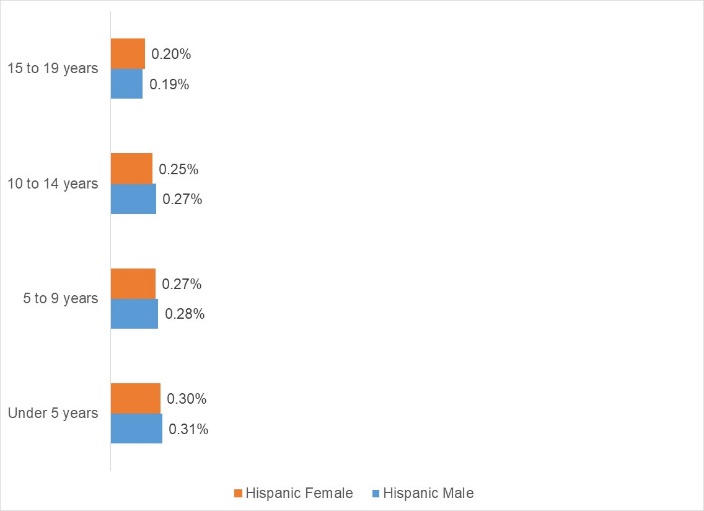
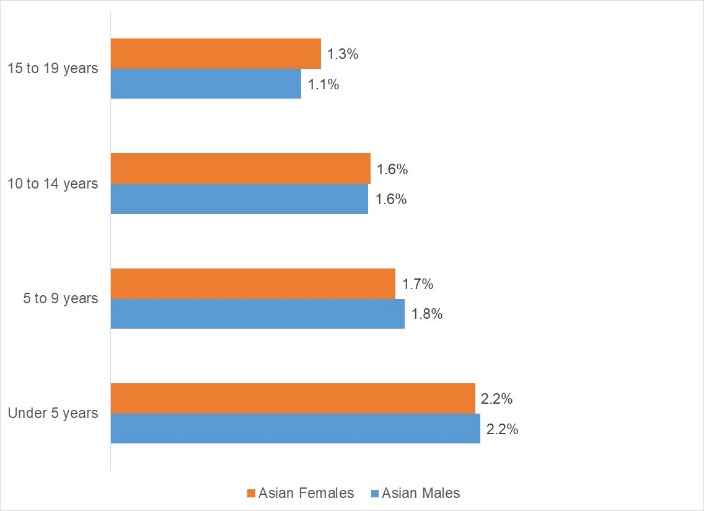


Figure 4. Comparison of Patients with Plausible Weight and Height Data in the Electronic Health Records and the U.S. Population by Race, Sex, and Age Group, 2015

For valid health prevalence results in youth and teens, we reviewed coverage of relevant characteristics related to BMI percentile, specifically sex and race/ethnicity.

Weight and height measurements data extracted from EHR systems and pre-processed by growthcleanr were used to calculate the number of patients with plausible values per 1,000 persons in the population by age and sex (Figure 5. Comparison of Patients with Plausible Weight and Height Data in the Electronic Health Records and the US Population by Age and Sex, 2015). Both males and females were included in the EHR, most often at the age of 2, with more males per 1,000 in the population being included than females. Males continued to outpace females for plausible values until puberty, when the rate of males with plausible height and weight data decreased with age, whereas females with plausible data remained relatively stable. By the age of 19, males were in the data at a rate of 21.2 per 1,000 persons, whereas females were in the data at a rate of 31.2 per 1,000 persons.

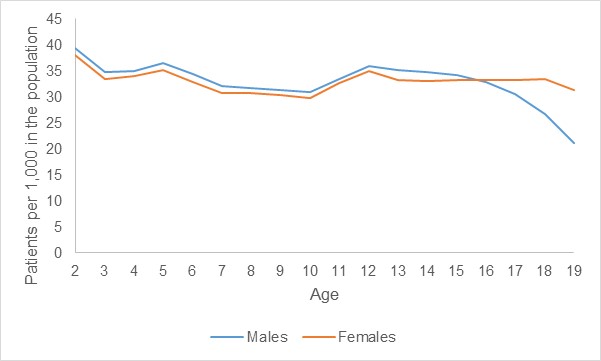


Figure 5. Comparison of Patients with Plausible Weight and Height Data in the Electronic Health Records and the U.S. Population by Age and Sex, 2015

Prevalence

A prevalence is either:

* **Crude**: the proportion of the sample that has a health condition (BMI percentile) at a point in time.
* **Weighted**: the proportion of the population that has a health condition at a point in time. See the Appendix A section “Statistical Weights” for more information.
* **Age-Adjusted**: the proportion of the population (adjusted by national distribution of age) that has a health condition at a point in time. See Appendix A section “Age Adjustment” for more information.

Race

Race is defined by one of the following categories: White, Black, Asian (including Native Hawaiian and other Pacific Islanders), and Other (including American Indian and Alaskan Native, some other race, two or more races).

These racial categories conform to previous work using a sample EHRs file. These categories are used because they are the race breakdowns available when CODI-PQ was created, though we recognize that these categories may not accurately reflect the way that patients would self-identify and may conceal important differences within groups.

Race Exclusion

Youth and teens from one or more races may be excluded from prevalence results. Statistical weighting programs require a large sample size (20 or more) in each stratum. If one or more racial groups has an insufficient sample size, the patients in the racial group impacted will automatically be excluded by the program.

Sickle Cell Disease

The race imputation uses presence of sickle cell disease (optional) to aid in imputing a patient’s race due to its strong correlation with race. The sickle cell disease phenotype used in the development phase includes the following:

**ICD-10 Codes**: D57, D57.0, D57.00, D57.01, D57.02, D57.1, D57.2, D57.20, D57.21, D57.211, D57.212, D57.219, D57.3 (sickle cell trait), D57.4\* (thalassemias), D57.40, D57.41, D57.411, D57.412, D57.419, D57.8, D57.80, D57.81, D57.811, D57.812, D57.819

**ICD-9 Codes**: 282.6, 282.60, 282.61, 282.62, 282.63, 282.64, 282.68, 282.69, 282.4\* (thalassemias), 282.40, 282.41, 282.42, 282.43, 282.44, 282.45, 282.46, 282.47, 282.49, 282.5 (sickle cell trait)

**SNOMED**: Concept ID 22281 (CC 127040003), Concept ID 26942 (CC 417425009), Concept ID 40485018 (CC 444108000), Concept ID 4213628 (CC 417357006), Concept ID 4216915 (CC 417279003), Concept ID 30683 (CC 416180004), Concept ID 315523 (CC 36472007), Concept ID 443738 (CC 416826005), Concept ID 321263 (CC 417048006), Concept ID 25518 (CC- 16402000 sickle cell trait), Concept ID 24006 (CC 35434009), Concept ID 443721 (CC 417517009), Concept ID 443726 (CC 417683006)

The probability of each race, given presence of sickle cell disease was calculated from a combination of published incidence rates as well as verified with AEMR where race and sickle cell disease were available.

Table 15. Proportions of Sickle Cell Disease Used to Impute Race

|  |  |
| --- | --- |
| **Race** | **Sickle Cell Proportion** |
| African American | 94.49% |
| White | 3.94% |
| Other | 1.14% |
| Asian | 0.42% |

Race Imputation

Race is a required input for CODI-PQ. The data inputs and link population data (pre-processing) program inputs race for each youth or teen missing race information. The program operates sequentially in three phases, imputing race for youth and teens in one of the following three phases, those who:

1. Have sickle cell disease
2. Are identified as Hispanic and do not have sickle cell disease
3. Neither have sickle cell disease nor are identified as Hispanic

The race imputation relies on a combination of medical and ACS data.

Once complete, the results from each phase are aggregated with each youth or teen with an EHR-provided race, an imputed race, or categorized as “unknown.”

A patient’s race may be missing after race imputation for one of four reasons:

1. The patient’s geography is either invalid or did not have a population count in the 2019 ACS.
2. The patient’s age is outside of the scope of the program or is unknown. Only persons ages 2 to 19 are in scope.
3. The sex of the patient is unknown.

CODI-PQ assigns a value for race if a patient does not have a known racial value through statistical imputation. In testing, approximately 27% of the records were missing race (values of “unknown”), yet biases by race were found when compared to the national distribution. Specifically, from a national file, white was overrepresented, and all non-white races were underrepresented. In addition, some electronic records do not store both race and ethnicity separately, thus CODI-PQ reassigns all records that are assigned a “race” of Hispanic (note: Hispanic is an ethnicity, not a race).

As of 2019, racial and ethnic disparities exist in youth and teen BMI percentile prevalence in the U.S. To reduce these disparities, high-quality data on race are needed. However, these data are often missing in some portion of EHRs. CODI-PQ imputes race for those with unknown race using programs based on race and ethnicity of surrounding the community, ethnicity of the patient (where available if race is unavailable), sickle cell disease, age, and height. Statistical weights are calculated (based on each patient’s age, sex, race, geography, and community characteristics) and used to adjust the EHRs non-probability sample to the population of interest. Weights are derived from individual-level demographic and social determinant of health (SDOH) data available in the EHR, as well as population-level SDOH proxies derived from the ACS data. Calculated prevalence is included as crude, weighted, and age-adjusted weighted results.

For records lacking race information, automated race imputation is employed in CODI-PQ data inputs and linked population data (pre-processing). Within the final program to calculate prevalence, the user specifies whether patients with imputed race should be included in the results. Records with a race value are included in the prevalence independent of whether imputed race is assigned as “yes” or “no.”

Race imputation occurs for each patient with an unknown race in three phases:

1. Patients with sickle cell disease
2. Patients identified as Hispanic but not identified with sickle cell disease
3. All other patients (neither have sickle cell disease nor are identified as Hispanic)

Table 16. Percentage of Patients Imputed for Each Phase in the Race Imputation Using AEMR Data

| **Phase** | **Percent of those Imputed** |
| --- | --- |
| Phase 1: Imputed based on known chronic condition | 6.8% |
| Phase 2: Imputed based on ethnicity | 7.0% |
| Phase 3: All other patients with unknown race | 86.3% |

Statistical Weights

CODI and National AEMR data are derived from EHRs. As described in Appendix A, applying statistical weights is often used to reduce potential biases introduced by the EHRs sampling methodology. Ratio adjustments are applied to all sampled youth and teens. Ratio adjustment is a statistical weighting technique aimed to improve the accuracy of survey results by both reducing bias and increasing precision.[[18]](#footnote-19) One way to accomplish this goal is known as iterative proportional fitting or raking. Raking adjusts the data so that groups that are underrepresented in the sample can be accurately represented in the final data set. Raking accurately matches sample distributions to known demographic characteristics of populations. The use of raking reduces nonresponse bias and has been shown to reduce error within sample results.

Implementing raking programs require the specification of appropriate weighting classes or cells. Data used to form classes for adjustments must be available for both sample and the population. CODI-PQ raking includes social determinant of health categories – age, sex, race, and education categories in the surrounding area (based on percentage of adults in the community with a bachelor’s degree or higher). Once formed, the weighting classes are assessed, and cells with small sample counts are aggregated with their nearest neighbor to reduce prevalence variability. The collapsing follows these guide points:

**Age** = age category less than or greater than current

**Sex** = do not aggregate

**Race** = do not aggregate, instead exclude small cell categories from prevalence results

**Education** = community with a similar education category

Raking is completed by adjusting for one demographic variable (or dimension) at a time. For example, when weighting by age and sex, weights would first be adjusted for age groups, then those results would be adjusted by sex groups. The calculations continue in an iterative process until all group proportions in the sample approach those of the population, or after a set number of iterations. Once raked, weight trimming is used to reduce errors in the outcome caused by unusually high or low weights in some categories.

The fundamental objective of CODI-PQ is to generate statistics that reduce bias and are sufficiently precise to satisfy the goals of the expected analyses of the data. In general, the goal is to keep the mean squared error (MSE) of the primary statistics of interest as low as possible. The MSE of a survey result is:

MSE = Variance + (Bias)2

The purpose of weighting adjustments is to reduce bias. Thus, the application of weighting adjustments usually results in lower bias in the associated survey statistics, but at the same time adjustments may result in some increases in variances of the survey results when compared with crude variances.

The increases in variance result from the added variability in the sampling weights due to the adjustments. Thus, the user who uses the weights should review the variability in the sampling weights caused by these adjustments. A trade-off is made between variance and bias to keep the MSE as low as possible. There is no exact rule for this trade-off because the amount of bias is unknown.

The five-year estimates of ACS do not include an age group of 2 to 4 years. Thus, CODI-PQ calculates the population of this age group by multiplying the count of persons aged 5 and under within each geography by the percentage of the national population that is aged 2 to 4, given that they are under the age of 5, based on Annual Estimates of the Resident Population by Single Year of Age and Sex for the U.S., July 1, 2018, U.S. Census Bureau. Population counts by race for those age less than 5 are also adjusted from Annual Estimates of the Resident Population using the same adjustment.

ACS race is categorized to match the EHRs file and grouped as White, African American, Asian (including Native Hawaiian and other Pacific Islanders), and other (including American Indian and Alaskan Native, some other race, two or more races).

ACS educational attainment (bachelor’s degree or more) is linked by geography (state and GEO3) based on the patient’s residential address. Once linked, education is calculated as the percent of the population aged 25 to 64 who have earned a bachelor’s degree or more within the youth or teen’s geography. Educational attainment is then dichotomized based on the value: 20% of the population with a bachelor’s degree or more. Approximately 52% of counties in the U.S. fall above 20%, and 48% fall below.

Prevalence Calculations

Crude prevalence is calculated as the count of the sample within each BMI percentile category.

To calculate the weighted prevalence of the population the sum of statistical weights within each BMI percentile is divided by the sum of statistical weights within the EHR. To control extreme weights which may increase the variance, extreme weights are trimmed. To calculate the variance of BMI percentile, a Taylor-series approximation is used.[[19]](#footnote-20)

Users are provided crude (unweighted) population, prevalence, and standard error, weighted population, prevalence, and standard error, and an optional age-adjusted prevalence and standard error. Age-adjusting aims to eliminate differences in results that result from differences in the age distribution of the population among geographies. The projected 2000 U.S. population was used as the standard population per current guide.

Sample Check

If SAMPLE\_CHECK = Y, then the CODI-PQ executes an optional review of the sample size by age, race, and sex based on user defined criteria. All demographic categories selected by the user (e.g., sex male, etc.) will be displayed in the SAS output or results window. Each will include the factor, value (e.g., sex, male) and either “Sample Size Is Insufficient” if n <20 or “Sample Size Is Sufficient.Executes. This optional check is included to pinpoint potential sample size issues. For example, if the sample size is insufficient for males, the user may choose to execute CODI-PQ again after excluding males.

Standard Error

The precision of a sample can be measured using a variety of calculations, including the standard error, confidence interval, and the margin of error. The standard error is the most commonly used measure of the precision of a value and provides a gauge of how close a value is likely to be to the true population value in the absence of any bias. See Appendix A.11 Variance for more information.

Suppression Criteria

Prevalence may be suppressed. CODI-PQ data suppression is adapted from the NCHS data presentation standards for reporting proportions in NCHS reports and data products,[[20]](#footnote-21) developed by the Data Suppression Workgroup at NCHS.

The multistep NCHS Data Presentation Standards for Proportions are based on a minimum denominator sample size and on the absolute and relative widths of a confidence interval calculated using the Clopper-Pearson method. The National Center for Health Statistics (NCHS) Data Presentation Standards for Proportions are applied to all CODI-PQ results. The Presentation Standards also provide guidance for identifying results for statistical review, CODI-PQ does not identify records for statistical review and leave this step for the user. The data presentation standards are described in Table 17 and Figure 6.

If one or more rows are suppressed, the user may select to increase their research criteria by including additional years of data, increasing the geography, or including more age, race, or sex categories. The suppression thresholds may also be altered by the user in the Quickstart program.

Table 17. NCHS Data Presentation Standards for Proportions

| **Statistic** | **Standard** |
| --- | --- |
| **Sample size** | Proportions should be based on a minimum denominator sample size and effective denominator sample size (when applicable) of 30. Results with either a denominator sample size or an effective denominator sample size (when applicable) less than 30 should be suppressed. If the number of encounters is 0 (or its complement[[21]](#footnote-22)), then the denominator sample size should be used to obtain confidence intervals. If all other criteria are met for presentation, a result based on 0 encounters (or its complement) should be flagged for statistical review by the clearance official. The review could result in either the presentation or the suppression of the proportion. |
| **Confidence interval** | If the sample size criterion is met, calculate a 95% two-sided confidence interval using the Clopper-Pearson method, or the Korn-Graubard method for complex surveys, and obtain its width. |
| **Small absolute confidence interval width** | If the absolute confidence interval width is greater than 0.00 and less than or equal to 0.05, then the proportion can be presented if the number of encounters is greater than 0 and the degrees of freedom criterion (below) is met. If the number of encounters is 0 (or its complement) or the degrees of freedom criterion is not met, then the result should be flagged for statistical review by the clearance official. The review could result in either the presentation or the suppression of the proportion. |
| **Large absolute confidence interval width** | If the absolute confidence interval width is greater than or equal to 0.30, then the proportion should be suppressed. |
| **Relative confidence interval width** | If the absolute confidence interval width is between 0.05 and 0.30 and the relative confidence interval width is more than 130%, then the proportion should be suppressed. |
| **Relative confidence interval width** | If the absolute confidence interval width is between 0.05 and 0.30 and the relative confidence interval width is less than or equal to 130%, then the proportion can be presented if the degrees of freedom criterion below is met. If the degrees of freedom criterion is not met, then the result should be flagged for statistical review by the clearance official. The review could result in either the presentation or the suppression of the proportion. |
| **Degrees of freedom** | When applicable for complex surveys, if the sample size and confidence interval criteria are met for presentation and the degrees of freedom are fewer than 8, then the proportion should be flagged for statistical review. This review may result in either the presentation or the suppression of the proportion. |
| **Complementary proportions** | If all criteria are met for presenting the proportion but not for its complement, then the proportion should be shown. A footnote indicating that the complement of the proportion may be unreliable should be provided. |

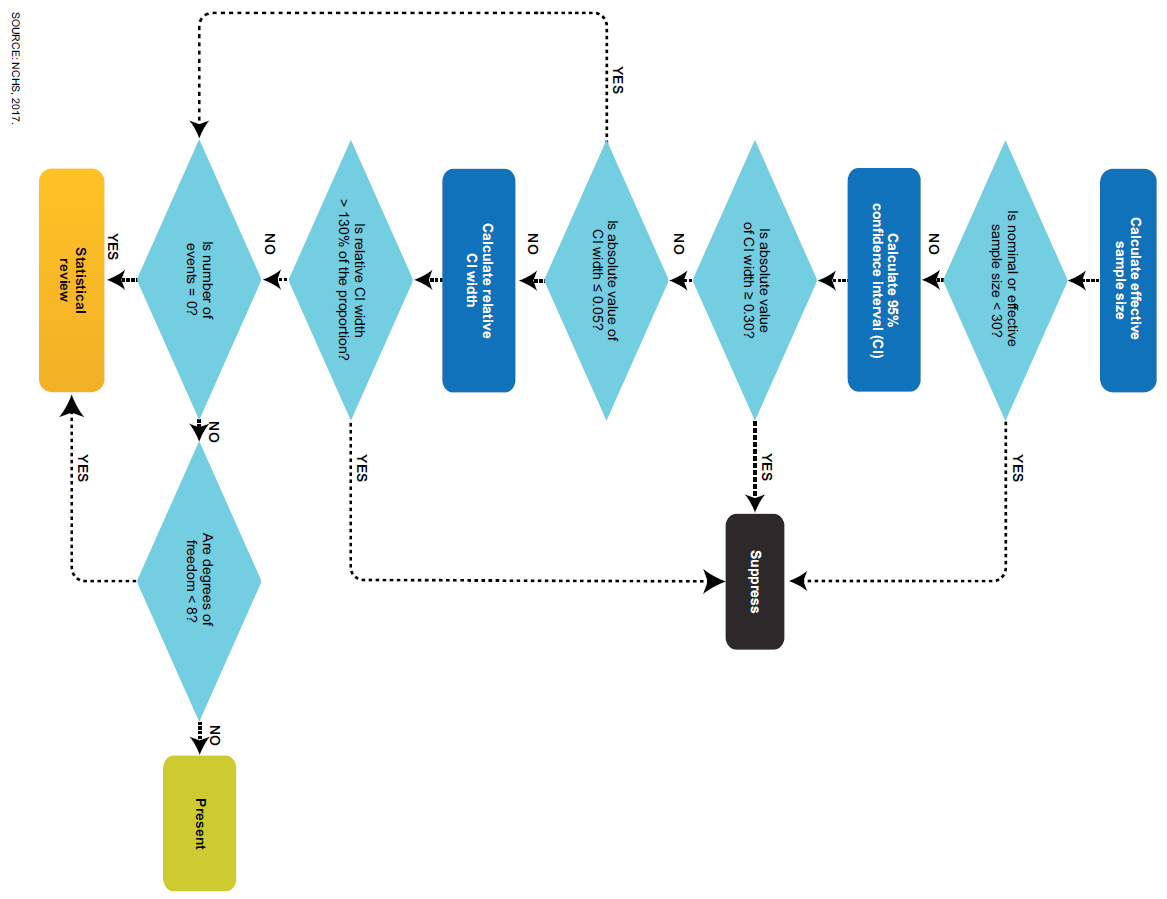


Figure 6. NCHS Suppression Standards

Variance

BMI percentile prevalence is derived using the sample weights and data on BMI percentile. BMI percentile ratios, and the ratio estimator, , corresponds to a population parameter, , such as the true but unknown BMI percentile prevalence. To define the population parameter, let:

= the number of youth and teens in stratum (), where stratum refers to state-GEO3

= the value of for youth or teen of stratum (often the possible values of are 0 and 1, as when indicates whether a youth or teen is or in a specified BMI percentile)

= 0 or 1, indicating whether youth or teen of stratum belongs to a particular domain (such as a specified race)

Then, adding the subscript to indicate the role of the domain, the ratio is the parameter of interest.

In the sample, let:

= the number of sample youth and teens in stratum

= the sampling weight for youth and teens in stratum

= the value of for youth and teen in stratum

= the value of the domain indicator for youth and teen in stratum

The distinction between and and between and is merely that for and the subscript refers to sampled youth and teens within stratum *,* whereas for and they refer to youth and teens in the population in stratum . Then, the ratio estimator for is:   
To calculate the variance of, a Taylor-series approximation is used.[[22]](#footnote-23) Within stratum, linearization yields the new variable.

Then, letting:

the Taylor-series approximation to the variance of is:

BMI Percentile

BMI percentile prevalence is calculated from a patient’s BMI. EHRs included for analysis should have at most one BMI percentile assigned to each patient within a calendar year. Percentiles are based on a patient’s age-sex BMI percentile value. Based on the 2000 CDC Growth Chart, the percentile for prevalence are as follows:

1. **Underweight**: BMI less than 5th percentile
2. **Healthy Weight**: BMI 5th percentile to less than the 85th percentile
3. **Overweight**: BMI 85th to less than the 95th percentile
4. **Obesity**[[23]](#footnote-24): BMI 95th percentile to less than 120 percent of the BMI value for the 95th percentile
   1. **Severe Obesity**: 120 percent or greater of the BMI value for the 95th percentile

For more information, visit: <https://www.cdc.gov/healthyweight/assessing/bmi/childrens_bmi/about_childrens_bmi.html>.

ZCTA-3

A ZCTA is a statistical geographic entity that approximates the delivery area for a U.S. Postal Service five-digit (ZCTA) ZIP code. ZCTAs are aggregations of census blocks that have the same predominant ZIP code associated with the residential mailing addresses in the U.S. Census Bureau’s Master Address File. ZCTAs do not precisely depict ZIP code delivery areas, and do not include all ZIP codes used for mail delivery. The U.S. Census Bureau has established ZCTAs as a new geographic entity similar to, but replacing, data tabulations for ZIP codes undertaken in conjunction with the 1990 and earlier censuses. For more information, refer to census.gov.[[24]](#footnote-25)

A ZCTA-3 includes the first three digits of a five-digit ZCTA. Three-digit ZCTAs (ZCTA-3), representing the first three digits of a ZIP code, were generated from the AEMR and ACS data. Once ZCTA’s are aggregated as ZCTA-3’s, then the first three digits of a residential ZIP code is equivalent to a ZCTA-3 in over 99% of the population.

Limitations

CODI-PQ users should consider the following limitations related to the program development, the data inputs required, and the results:

* Representativeness of CODI-PQ results – CODI-PQ results may differ from those based on a probability-based survey that could be more representative of the general population.
* Inclusion in EHRs – EHRs represent the care-seeking population for all medical providers included within a sample.
* Random missingness of plausible height or weight – CODI-PQ patient inclusion requires a plausible height and weight value. It is assumed that if patients are missing height and weight from EHRs, it is missing at random.
* Random missingness of demographic and geographic characteristics – CODI-PQ patient inclusion requires a valid and known age, sex, and geographic location to be reported. The race of each patient is also needed, although the program imputes race for patients missing race. It is assumed that if patients are missing age, sex, and/or geographic location from EHRs, it is missing at random.
* Race imputation – Race imputation assigns one value of race per patient. Multiple-imputation of race is not employed in CODI-PQ to allow for a) analysis of large EHR files without the need for increasing the length of the original file and b) ease in counting number of respondents in the crude results. Variance for those with imputed race is likely smaller than those with known race. Also, race imputation does not analyze a patient’s first and last name. Other EHR race imputation methodologies have utilized the patient’s first and last name with positive results.
* Probabilistic record linkage strategies include false links and missed matches.
  + It is recommended that the user become familiar with any record linkage strategy and its limitations.
  + If the linkage errors are not properly taken into account, biased estimates and mis-relationships between variables recorded in different sources (i.e., household linkage, person 1 in source A and person 2 in source B) may result (Di Consiglio and Tuoto, 2018).
  + If the user has information about how linkage error affects the distribution of household obesity, consider using techniques for quantitative bias analysis, to adjust for these errors. (Lash, 2011, Schneeweiss, 2006).
* Measurement error – Height and weight measurement protocols may differ between medical providers, even with clear protocols aimed to increase consistency between medical professionals,[[25]](#footnote-26) leading to potential measurement error. Additionally, height and weight values in EHRs are subject to data entry errors or software glitches. All CODI-PQ EHR data were cleaned using growthcleanr. Growthcleanr scans all available height and weight values and flags values that are implausible; however, users must decide to exclude the implausible values, recognizing that biologically acceptable values may still have errors. See Methods for more information about growthcleanr.
* Small sample sizes – A small number of patient-level records (encounters) could result in unstable results and reflect poor EHR coverage, a small underlying population, and/or a rare encounter. CODI-PQ suppresses results based on published small sample guidelines using the National Center for Health Statistics Data Presentation Standards for Proportions[[26]](#footnote-27).

###### Social Determinants of Health

Data from CODI and AEMR are not a random sample, making adjusting the non-probability sample on observed differences between the sample and the target population beneficial. Therefore, SDOH will be used to statistically weight those non-probability samples. To identify the best SDOH for weighting samples of U.S. youth and teens aged 2-19 years, the Health FFRDC identified and prioritized SDOH associated with youth and teen BMI percentile in the U.S. and by subpopulations within the U.S., including age, race, sex, and sub-geography.

Prioritizing Social Determinants of Health

Thirty-six SDOH concepts were considered for analysis. The list of concepts was a result of subject matter expert brainstorm sessions in 2019 between Health FFRDC and CDC staff. All concepts readily available at the geography of interest were included in the analysis. Table 18 provides a list of concepts included in the SDOH analysis. Table 19 outlines additional concepts considered for inclusion that did not meet our final criteria for inclusion.

Table 18. Included Concepts: Social Determinants of Health

|  |  |
| --- | --- |
| Approved Concepts | Source(s) with Concept Available |
| Age | CODI, AEMR, ACS |
| Region | Census, U.S. Department of Health and Human Services |
| Education | ACS |
| Employment/Unemployment | ACS |
| Ethnicity | AEMR, ACS |
| Head-of-household | ACS |
| Housing | ACS |
| Housing Stability | ACS |
| Income | ACS |
| Income Inequality | ACS |
| Poverty | ACS |
| Primary Language | ACS |
| Medical Insurance | CODI, ACS |
| Race | CODI, AEMR, ACS |
| Sex | CODI, AEMR, ACS |
| Transportation | ACS |
| Adult BMI Percentile (underweight, healthy weight, overweight, obesity, severely obesity) | AEMR |

Table 19. Additional Concepts Considered: Social Determinants of Health

|  |
| --- |
| Additional Concepts Considered |
| Pediatric Medical Complexity Algorithm |
| Primary language |
| Urbanicity |
| Walkability |
| Attention Deficit Hyperactivity Disorder (ADHD) |
| Air quality |
| Asthma |
| Deprivation index |
| Domestic violence |
| Food desert/swamp |
| Home value |
| Incarceration history |
| Material security |
| Migrant/seasonal work |
| Park access |
| Refugee status |
| Safety |
| Social integration and support |
| Stress/social vulnerability index |
| Veteran status |

From Concepts to Measures

After SDOH concepts were identified, each data source was reviewed for measures relating to the concept. A comprehensive list of measures was thus identified for analysis. Table 19 provides a full list of measures included in the analysis. In this section we share the methodology and results for both the bivariate and multivariable analyses performed for the SDOH prioritization. Within each methodology section, we share the models, data source used for the analysis, universe (population included in sample), covariate(s) and dependent variable(s) included in the analysis, goodness of fit requirements, and how the results are reported.

Bivariate Analysis

There are multiple ways to characterize some SDOH concepts, leading to multiple candidate measures[[27]](#footnote-28) of a SDOH concept. Up to six possible measures were compared to one another within each SDOH concept. The best measure within each concept was determined using a bivariate analysis. The measure within each concept that best explains the outcome (youth and teen BMI percentile) was considered to outrank other measures. Once the best measure was determined, the concepts were also ranked against one another. These results generate a ranking of measures and concepts without taking other factors into consideration.

SDOH Prioritization Bivariate Methodology

We performed logistic regressions for each approved concept with multiple plausible specifications (measures). The specifications included:

**Models:** Cumulative Logit and Multinomial Logit

When there is a natural ordering to the response categories, a cumulative logit model is appropriate. This logit models the cumulative probability that a response (Y) can be classified at or below a given category. In its simplest form, it assumes that the same proportionality constant applies for each parameter to each cumulative level.

The cumulative logit model has the following form:

Although cumulative logit models rank the dependent variable, a multinomial logit model was also fit. Multinomial logit models are an extension of logistic regression models in which the response can be classified into three or more unordered categories. In these analyses, the response variable is classified into five categories. The multinomial model was fit in addition to or in place of the cumulative logit model for three reasons:

* One or more assumptions were breached thus making a multinomial model the next most appropriate model form.
* The multinomial model took considerably more computing time.
* The rank order of the models was identical in all instances where both models were fit.

The multinomial logit model has the following form. For K classes multinomial problem where labels ranged from [0, K-1], we can generalize it via:

**Data Source:** 1.8 million records from the 2016 AEMR linked to the 2018 ACS by ZIP Code Tabulation Area (3 digit) as defined by the Census Bureau (i.e., ZCTA-3), plus confirmation file from the 2018 AEMR linked to the 2018 ACS by ZCTA-3.

**Universe:** youth and teens, aged 2 to 19.

**Covariate:** one approved measure within a concept.

**Dependent variables:** weight classification (underweight, healthy weight, overweight, obesity, severe obesity).

**Goodness of fit:** McFadden pseudo R-squared. We used pseudo R-squared measures for evaluating “goodness of fit” in regression models with categorical dependent variables. Log-likelihood-based pseudo R-squared represent the improvement in model likelihood over a null model.

**Reporting:** Within each concept, we ranked measures from those with the highest pseudo R-squared to those with the lowest pseudo R-squared. Across concepts, we selected the measure with the highest pseudo R-squared.

Results

Table 20 provides a ranked list of concepts. Age of the patient from AEMR ranked the most predictive followed by adult BMI percentile, educational attainment of population aged 45 to 64 years of age, income of a single parent, race, and medical insurance.

The statistical weights cell formulation is based on the Childhood Obesity Data Initiative – Social Determinants of Health (SDOH) Prioritization report[[28]](#footnote-29) as well as other artifacts. The SDOH report was developed in partnership between the Health FFRDC and CDC, with feedback from leading health researchers. Table 20 includes the finalized list of BMI percentile related social determinants of health used in CODI-PQ.

Table 20. Final List of Prioritized Social Determinants of Health

|  |  |  |
| --- | --- | --- |
| Concept | Measure | Source |
| Age | Age of patient, continuous | EHR |
| Race | Patient race, categorical | EHR |
| Sex | Patient sex, categorical | EHR |
| Education | Educational attainment of population aged 25 to 64 years of age | ACS |

Multivariate Analysis

Seventeen concepts were calculated and ranked for inclusion in the CODI weighting program through bivariate analyses. The top twelve concepts from those bivariate analyses were further calculated in combination using a multivariable method to determine a possible mix of concepts to include in the weighting programs.

SDOH Prioritization Multivariate Methodology

The bivariate analyses ranked concepts against one another but not in combination. A multivariable analysis was needed to determine the best set of SDOH as they relate to youth and teen BMI percentile. Regressions were performed as follows:

**Model**: Stepwise Cumulative Logit and Multicategory Logit. With stepwise regression, effects are entered into and removed from the model in such a way that each forward selection step can be followed by one or more backward elimination steps. The stepwise selection process terminates if no further effect can be added to the model or if the current model is identical to a previously calculated model.

**Data Source:** 1.8 million records from the 2016 AEMR linked to the 2018 ACS by ZCTA-3, plus confirmation file from the 2018 AEMR linked to the 2018 ACS by ZCTA-3.

**Universe:** youth or teen, aged 2 to 19.

**Covariates:** the top 12 ranked concepts from the bivariate analysis.

**Dependent variables:** weight classification (underweight, healthy weight, overweight, obesity, severely obesity).

**Reporting:** Across models, we selected the concepts based on the order in which they are included in the final model. We removed concepts or updated the measure within each concept as appropriate and reported on the updated order.

Results

Three multivariable models were run. In model 1, all concepts were ranked. Results indicate age remained most predictive, followed by race, adult BMI percentile, sex, medical insurance, head-of-household, income, and primary language. Adult BMI percentile ranked third amongst all concepts, but the adult’s data source is not nationally representative because it is also calculated from the non-probability sample of AEMR. As such, inclusion of those data is not possible without applying appropriate statistical weights.

Model 2 excluded the adult BMI percentile. Results indicate age remained most predictive, followed by race, education, sex, primary language, transportation, and income.

Model 3 assumed age, race, and sex were required for subpopulation results and excluded the adult BMI percentile. Results indicate education was next most predictive followed by primary language, transportation, head-of-household, and income.

Summary

Model 3 in Table 23 represents the final ranking of concepts. Within model 3, the educational attainment of the population aged 45 to 64 was found to be most predictive of youth and teen BMI percentile. Although this is the most predictive measure within the concept, this education measure may have limited impact in areas with a small number of persons in this age range. For this reason, a final model was calculated with educational attainment of the population aged 25 to 64, without the inclusion of the adult BMI percentile.

Once the new education measure was found to be equally relevant, a final regression was performed using 2018 AEMR data. The order in which measures were added to the model remained consistent, confirming our findings. The final prioritization is in Table 21.

Table 21. Social Determinants of Health: Summary of Findings

| Rank | Concept | Measure | Source |
| --- | --- | --- | --- |
| 1 | Age | Age of patient, continuous | AEMR |
| 1 | Race | Patient race, categorical | AEMR |
| 1 | Sex | Patient sex, categorical | AEMR |
| 2 | Education | Educational attainment of population aged 25 to 64 years of age | ACS |
| 3 | Primary Language | Language spoken at home, population aged 5 plus | ACS |
| 4 | Transportation | Means of transportation to work | ACS |
| 5 | Head-of-household | Percent of households in geography by head of household (female with children, male with children, married with children, grandparent with children) | ACS |
| 6 | Income | Family income, with kids less than 18 years of age (categorical) | ACS |
| 7 | Poverty | Percent of population who receive supplemental security income (SSI), cash public assistance income, or food stamps/snap in the past 12 months | ACS |
| 8 | Employment/Unemployment | Percent of population that is unemployed, Categorical | ACS |
| 9 | Medical Insurance | Health insurance coverage status of population | ACS |

Conclusion

Bivariate and multivariable analyses provided a systematic method to assess SDOH against youth and teen BMI percentile in the U.S. AEMR’s nationwide data files generate a large number of real-world values of BMI percentile after adjusting for observed differences in demographics relative to the American Community Survey by state and ZCTA-3. The systematic analysis was limited in its knowledge of the data sources available for weighting. Should additional nationally representative files become available, the results could change with the inclusion of different concepts, measures, or sources. For example, the AEMR adult BMI percentile data is a non-probability-based sample and thus requires statistical weights be applied before its use.

Limitations, SDOH Prioritization

The limitations of this analysis include the following identified items as well as others. First, our comprehensive list of SDOH was compiled from a team of subject matter experts in 2019. Future work may consider comparing other concepts to further strengthen the findings or may consider innovative concepts (e.g., use of a phenotype or household BMI percentile).

Second, inclusion of possible SDOH was limited based on readily available data sources, concepts, and measures within those sources. If other SDOH had been available or included in the analyses, the results may have differed. Future work may consider comparing other data sources, concepts, and measures.

Third, our analysis relied on logistic regression, a statistical model commonly used to rank measures against a dependent variable. Other analytic tools are available including use of data science tools such as a random forest model. The results may be consistent with the multinomial logit analysis or may show additional nuances not possible to see with the current analysis. Random Forests are another way to extract information from a set of data. The appeals of this type of model are:

* It does not assume that the model has a linear relationship like regression models do.
* It utilizes ensemble learning. If we were to use just 1 decision tree, we would not be using ensemble learning. A random forest takes random samples, forms many decision trees, and then averages out the leaf nodes to get a clearer model.
* When fine-tuned through careful statistical analysis, it is highly likely to perform better than logistic regression.

As such, random forest models are more time consuming. Logistic regressions tend to be as or more effective than random forests if time to tune is not available so long as the number of covariates is small in comparison to the number of records.

Fourth, analysis of adult data has revealed a strong correlation between the number of medical encounters and the percentage of the population with BMI percentile. A similar trend can also be seen in youth and teen data, although additional research is needed.

Future Research Opportunities

Additional research is warranted, with the following suggestions:

* Include a nationally representative adult BMI percentile file
* Explore how SDOH change as the patient ages
* Explore SDOH of the adult population and the household
* Explore additional data sources when available
* Explore a random forest model or other functional form

Supplemental Tables

Table 22. Additional Concepts and Measures Considered: Social Determinants of Health

|  |  |  |
| --- | --- | --- |
| Concept | Measure | Source (if available) |
| Environmental | Air quality |  |
| Food desert/swamp | Food access research atlas | [https://www.ers.usda.gov/data-products/food-access-research-atlas/  https://www.ers.usda.gov/data-products/food-access-research-atlas/download-the-data/](https://www.ers.usda.gov/data-products/food-access-research-atlas/) |
| Food desert/swamp | Food security | [https://www.ers.usda.gov/webdocs/DataFiles/50764/techdoc2018.pdf?v=7286.8  https://thedataweb.rm.census.gov/ftp/cps\_ftp.html#cpssupps](https://www.ers.usda.gov/webdocs/DataFiles/50764/techdoc2018.pdf?v=7286.8) |
| Food desert/swamp | Food environment atlas | <https://www.ers.usda.gov/data-products/food-environment-atlas/data-access-and-documentation-downloads/> |
| Health | ADHD |  |
| Health | Asthma |  |
| Income inequality | U.S. Census Bureau | <https://www.census.gov/topics/income-poverty/income-inequality/about/metrics.html> |
| Income inequality | Deprivation index | <https://www.neighborhoodatlas.medicine.wisc.edu/> |
| Income inequality | Theil index |  |
| Income inequality | Atkinson index |  |
| Income inequality | Mean Log Deviation (MLD) |  |
| Safety | Uniform Crime Reporting Program | <https://www.fbi.gov/services/cjis/ucr/> |
| Safety | Motor vehicle crashes | <https://one.nhtsa.gov/Data/Fatality-Analysis-Reporting-System-(FARS)> |
| Safety | Domestic violence |  |
| Walkability | Walk Score | [https://www.walkscore.com/cities-and-neighborhoods/  https://www.walkscore.com/methodology.shtml](https://www.walkscore.com/cities-and-neighborhoods/) |
| Walkability | National walkability index | [https://catalog.data.gov/dataset/walkability-index  https://www.epa.gov/sites/production/files/2014-03/documents/sld\_userguide.pdf  https://catalog.data.gov/harvest/object/4f569c8f-4f9a-4d00-bf2b-e242296be0ce/html](https://catalog.data.gov/dataset/walkability-index) |
| Walkability | Park access | National Environmental Public Health Tracking Network (NEPHTN) ephtracking.cdc.gov |
| Wealth | Home value |  |

Table 23. Social Determinants of Health Measures

| Concept | Measure | Source |
| --- | --- | --- |
| Age | Age of patient, categorical | AEMR |
| Age | Age of population in geography, categorical | ACS |
| Age | Age of patient, continuous | AEMR |
| Region | Census division (9) | Census |
| Region | Census region (4) | Census |
| Region | Health and Human Services region (10) | U.S. Department of Health and Human Services |
| Education | Educational attainment of population aged 18 to 24 years of age | ACS |
| Education | Educational attainment of population aged 25 to 34 years of age | ACS |
| Education | Educational attainment of population aged 35 to 44 years of age | ACS |
| Education | Educational attainment of population aged 45 to 64 years of age | ACS |
| Education | Educational attainment of population aged 18 years of age or older | ACS |
| Education | Educational attainment of adult females | ACS |
| Education | Educational attainment of adult males | ACS |
| Employment/Unemployment | Percent of population that is unemployed, categorical | ACS |
| Ethnicity | Percentage of youth or teen in geography who are of Hispanic Origin | ACS |
| Head-of-household | Percent of households in geography by head of household (female with children, male with children, married with children, grandparent with children) | ACS |
| Housing | Tenure of occupied housing (rent, own) | ACS |
| Housing Stability | Percent of households not living in the same house a year ago | ACS |
| Income | Family income, with kids less than 18 years of age (categorical) | ACS |
| Income | Median family income | ACS |
| Income | Median household income | ACS |
| Income | Married family income, with kids less than 18 years of age (categorical) | ACS |
| Income | Income of a single parent (categorical) | ACS |
| Income Inequality | Gini Index | ACS |
| Poverty | Percent of population who receive supplemental security income (SSI), cash public assistance income, or food stamps/snap in the past 12 months | ACS |
| Poverty | Percent of family households living in poverty | ACS |
| Poverty | Percent of population living in poverty | ACS |
| Poverty | Percent of children living in poverty, categorized by age of children | ACS |
| Poverty | Percent of children living in poverty | ACS |
| Poverty | Percent of population living in poverty, by sex | ACS |
| Primary Language | Language spoken at home, population aged 5+ | ACS |
| Medical Insurance | Health Insurance coverage status of population | ACS |
| Race | ACS race distribution | ACS |
| Race | Race, geographic distribution | AEMR |
| Race | Patient race, categorical | AEMR |
| Sex | Patient sex, categorical | AEMR |
| Sex | Sex, geographic distribution | AEMR |
| Sex | Sex of youth or teens, geographic distribution | ACS |
| Transportation | Means of transportation to work | ACS |
| Adult BMI percentile | Adult BMI percentile, categorical | AEMR |

Table 24. Social Determinants of Health: Bivariate Ranked Concepts and Measures

| Rank | Concept | Measure | Source | Measure Rank |
| --- | --- | --- | --- | --- |
| 1 | Age | Age of patient, categorical | AEMR | 1 |
| 1 | Age | Age of patient, continuous | AEMR | 2 |
| 1 | Age | Age of population in geography, categorical | ACS | 3 |
| 2 | Adult BMI percentile | Adult BMI percentile, categorical | AEMR | 1 |
| 3 | Education | Educational attainment of population aged 45 to 64 years of age | ACS | 1 |
| 3 | Education | Educational attainment of adult females | ACS | 2 |
| 3 | Education | Educational attainment of adult males | ACS | 3 |
| 3 | Education | Educational attainment of population aged 35 to 44 years of age | ACS | 4 |
| 3 | Education | Educational attainment of population aged 18 years of age or older | ACS | 5 |
| 3 | Education | Educational attainment of population aged 25 to 34 years of age | ACS | 6 |
| 3 | Education | Educational attainment of population aged 18 to 24 years of age | ACS | 7 |
| 4 | Income | Income of a single parent (categorical) | ACS | 1 |
| 4 | Income | Family income, with kids less than 18 years of age (categorical) | ACS | 2 |
| 4 | Income | Married family income, with kids less than 18 years of age (categorical) | ACS | 3 |
| 4 | Income | Median family income | ACS | 4 |
| 4 | Income | Median household income | ACS | 5 |
| 5 | Race | Patient race, categorical | AEMR | 1 |
| 5 | Race | ACS race distribution | ACS | 2 |
| 5 | Race | Race, geographic distribution | AEMR | 3 |
| 6 | Medical Insurance | Health insurance coverage status of population | ACS | 1 |
| 7 | Poverty | Percent of children living in poverty, categorized by age of children | ACS | 1 |
| 7 | Poverty | Percent of population living in poverty, by sex | ACS | 2 |
| 7 | Poverty | Percent of children living in poverty | ACS | 3 |
| 7 | Poverty | Percent of population living in poverty | ACS | 4 |
| 7 | Poverty | Percent of family households living in poverty | ACS | 5 |
| 7 | Poverty | Percent of population who receive supplemental security income (SSI), cash public assistance income, or food stamps/snap in the past 12 months | ACS | 6 |
| 8 | Transportation | Means of transportation to work | ACS | 1 |
| 9 | Head-of-household | Percent of households in geography by head of household (female with children, male with children, married with children, grandparent with children) | ACS | 1 |
| 10 | Primary Language | Language spoken at home, population aged 5+ | ACS | 1 |
| 11 | Employment/Unemployment | Percent of population that is unemployed, categorical | ACS | 1 |
| 12 | Sex | Patient sex, categorical | AEMR | 1 |
| 12 | Sex | Sex, geographic distribution | AEMR | 2 |
| 12 | Sex | Sex of youth or teen, geographic distribution | ACS | 3 |
| 13 | Ethnicity | Percentage of youth or teen in geography who are of Hispanic Origin | ACS | 1 |
| 14 | Region | Health and Human Services region (10) | U.S. Department of Health and Human Services | 1 |
| 14 | Region | Census region (4) | Census | 2 |
| 14 | Region | Census division (9) | Census | 3 |
| 15 | Housing Stability | Percent of households not living in the same house a year ago | ACS | 1 |
| 16 | Housing | Tenure of occupied housing (rent, own) | ACS | 1 |
| 17 | Income Inequality | Gini Index | ACS | 1 |

Table 25. Social Determinants of Health: Multivariable Ranked Measures

| Model | Rank | Concept | Comments |
| --- | --- | --- | --- |
| 1 | 1 | Age |  |
| 1 | 2 | Race |  |
| 1 | 3 | Adult BMI percentile |  |
| 1 | 4 | Sex |  |
| 1 | 5 | Medical Insurance |  |
| 1 | 6 | Head-of-household |  |
| 1 | 7 | Income |  |
| 1 | 8 | Primary Language |  |
| 1 | 9 | Transportation |  |
| 1 | 10 | Poverty |  |
| 1 | 11 | Employment/Unemployment |  |
| 1 | 12 | Education |  |
| 2 | 1 | Age |  |
| 2 | 2 | Race |  |
| 2 | 3 | Education |  |
| 2 | 4 | Sex |  |
| 2 | 5 | Primary Language |  |
| 2 | 6 | Transportation |  |
| 2 | 7 | Income |  |
| 2 | 8 | Head-of-household |  |
| 2 | 9 | Medical Insurance |  |
| 2 | 10 | Poverty |  |
| 2 | NA | Adult BMI percentile | Excluded |
| 2 | NA | Employment/Unemployment | Excluded |
| 3 | 1 | Age |  |
| 3 | 1 | Race |  |
| 3 | 1 | Sex |  |
| 3 | 2 | Education |  |
| 3 | 3 | Primary Language |  |
| 3 | 4 | Transportation |  |
| 3 | 5 | Head-of-household |  |
| 3 | 6 | Income |  |
| 3 | 7 | Poverty |  |
| 3 | 8 | Employment/Unemployment |  |
| 3 | 9 | Medical Insurance |  |
| 3 | NA | Adult BMI percentile | Excluded |

###### ACS File Layouts

ACS Input File Layout

The following variables are included in both the ZCTA-3 file as well as the County file. ACS data is imported in the CODI-PQ and require a csv file with the following variable names, possible variable values, and in the order listed below. Variable geoid (option 1) is for ZCTA-3 files only and geoid (option 2) is for the County file only.

Table 26. ACS Input File Layout, CSV File

| Variable Name | Label | Description | Format | Example |
| --- | --- | --- | --- | --- |
| Geoid | Geoid (option 1) | 3 digits ZIP Code Tabulation Areas (ZCTAs) followed by two-letter State Abbreviations | Character | 221 |
| Geoid | Geoid (option 2) | 3-digit County FIPS Code | Character | 059 |
| State\_code | State FIPS code | 2-digit State Abbreviation | Character | VA |
| b01001f\_001 | SEX BY AGE (SOME OTHER RACE ALONE); Universe: People who are Some Other Race alone; Total: Total: | Population count | Number | 8199 |
| b01001f\_003 | SEX BY AGE (SOME OTHER RACE ALONE); Universe: People who are Some Other Race alone; Male: Under 5 years | Population count | Number | 256 |
| b01001f\_004 | SEX BY AGE (SOME OTHER RACE ALONE); Universe: People who are Some Other Race alone; Male: 5 to 9 years | Population count | Number | 246 |
| b01001f\_005 | SEX BY AGE (SOME OTHER RACE ALONE); Universe: People who are Some Other Race alone; Male: 10 to 14 years | Population count | Number | 495 |
| b01001f\_006 | SEX BY AGE (SOME OTHER RACE ALONE); Universe: People who are Some Other Race alone; Male: 15 to 17 years | Population count | Number | 297 |
| b01001f\_007 | SEX BY AGE (SOME OTHER RACE ALONE); Universe: People who are Some Other Race alone; Male: 18 and 19 years | Population count | Number | 145 |
| b01001f\_018 | SEX BY AGE (SOME OTHER RACE ALONE); Universe: People who are Some Other Race alone; Female: Under 5 years | Population count | Number | 271 |
| b01001f\_019 | SEX BY AGE (SOME OTHER RACE ALONE); Universe: People who are Some Other Race alone; Female: 5 to 9 years | Population count | Number | 188 |
| b01001f\_020 | SEX BY AGE (SOME OTHER RACE ALONE); Universe: People who are Some Other Race alone; Female: 10 to 14 years | Population count | Number | 267 |
| b01001f\_021 | SEX BY AGE (SOME OTHER RACE ALONE); Universe: People who are Some Other Race alone; Female: 15 to 17 years | Population count | Number | 139 |
| b01001f\_022 | SEX BY AGE (SOME OTHER RACE ALONE); Universe: People who are Some Other Race alone; Female: 18 and 19 years | Population count | Number | 134 |
| b01001e\_001 | SEX BY AGE (NATIVE HAWAIIAN AND OTHER PACIFIC ISLANDER ALONE); Universe: People who are Native Hawaiian and Other Pacific Islander alone; Total: Total: | Population count | Number | 278 |
| b01001e\_003 | SEX BY AGE (NATIVE HAWAIIAN AND OTHER PACIFIC ISLANDER ALONE); Universe: People who are Native Hawaiian and Other Pacific Islander alone; Male: Under 5 years | Population count | Number | 0 |
| b01001e\_004 | SEX BY AGE (NATIVE HAWAIIAN AND OTHER PACIFIC ISLANDER ALONE); Universe: People who are Native Hawaiian and Other Pacific Islander alone; Male: 5 to 9 years | Population count | Number | 28 |
| b01001e\_005 | SEX BY AGE (NATIVE HAWAIIAN AND OTHER PACIFIC ISLANDER ALONE); Universe: People who are Native Hawaiian and Other Pacific Islander alone; Male: 10 to 14 years | Population count | Number | 28 |
| b01001e\_006 | SEX BY AGE (NATIVE HAWAIIAN AND OTHER PACIFIC ISLANDER ALONE); Universe: People who are Native Hawaiian and Other Pacific Islander alone; Male: 15 to 17 years | Population count | Number | 0 |
| b01001e\_007 | SEX BY AGE (NATIVE HAWAIIAN AND OTHER PACIFIC ISLANDER ALONE); Universe: People who are Native Hawaiian and Other Pacific Islander alone; Male: 18 and 19 years | Population count | Number | 0 |
| b01001e\_018 | SEX BY AGE (NATIVE HAWAIIAN AND OTHER PACIFIC ISLANDER ALONE); Universe: People who are Native Hawaiian and Other Pacific Islander alone; Female: Under 5 years | Population count | Number | 0 |
| b01001e\_019 | SEX BY AGE (NATIVE HAWAIIAN AND OTHER PACIFIC ISLANDER ALONE); Universe: People who are Native Hawaiian and Other Pacific Islander alone; Female: 5 to 9 years | Population count | Number | 0 |
| b01001e\_020 | SEX BY AGE (NATIVE HAWAIIAN AND OTHER PACIFIC ISLANDER ALONE); Universe: People who are Native Hawaiian and Other Pacific Islander alone; Female: 10 to 14 years | Population count | Number | 0 |
| b01001e\_021 | SEX BY AGE (NATIVE HAWAIIAN AND OTHER PACIFIC ISLANDER ALONE); Universe: People who are Native Hawaiian and Other Pacific Islander alone; Female: 15 to 17 years | Population count | Number | 0 |
| b01001e\_022 | SEX BY AGE (NATIVE HAWAIIAN AND OTHER PACIFIC ISLANDER ALONE); Universe: People who are Native Hawaiian and Other Pacific Islander alone; Female: 18 and 19 years | Population count | Number | 10 |
| b01001d\_001 | SEX BY AGE (ASIAN ALONE); Universe: People who are Asian alone; Total: Total: | Population count | Number | 14824 |
| b01001d\_003 | SEX BY AGE (ASIAN ALONE); Universe: People who are Asian alone; Male: Under 5 years | Population count | Number | 277 |
| b01001d\_004 | SEX BY AGE (ASIAN ALONE); Universe: People who are Asian alone; Male: 5 to 9 years | Population count | Number | 222 |
| b01001d\_005 | SEX BY AGE (ASIAN ALONE); Universe: People who are Asian alone; Male: 10 to 14 years | Population count | Number | 276 |
| b01001d\_006 | SEX BY AGE (ASIAN ALONE); Universe: People who are Asian alone; Male: 15 to 17 years | Population count | Number | 263 |
| b01001d\_007 | SEX BY AGE (ASIAN ALONE); Universe: People who are Asian alone; Male: 18 and 19 years | Population count | Number | 774 |
| b01001d\_018 | SEX BY AGE (ASIAN ALONE); Universe: People who are Asian alone; Female: Under 5 years | Population count | Number | 101 |
| b01001d\_019 | SEX BY AGE (ASIAN ALONE); Universe: People who are Asian alone; Female: 5 to 9 years | Population count | Number | 237 |
| b01001d\_020 | SEX BY AGE (ASIAN ALONE); Universe: People who are Asian alone; Female: 10 to 14 years | Population count | Number | 355 |
| b01001d\_021 | SEX BY AGE (ASIAN ALONE); Universe: People who are Asian alone; Female: 15 to 17 years | Population count | Number | 242 |
| b01001d\_022 | SEX BY AGE (ASIAN ALONE); Universe: People who are Asian alone; Female: 18 and 19 years | Population count | Number | 1404 |
| b01001c\_001 | SEX BY AGE (AMERICAN INDIAN AND ALASKA NATIVE ALONE); Universe: People who are American Indian and Alaska Native alone; Total: Total: | Population count | Number | 755 |
| b01001c\_003 | SEX BY AGE (AMERICAN INDIAN AND ALASKA NATIVE ALONE); Universe: People who are American Indian and Alaska Native alone; Male: Under 5 years | Population count | Number | 36 |
| b01001c\_004 | SEX BY AGE (AMERICAN INDIAN AND ALASKA NATIVE ALONE); Universe: People who are American Indian and Alaska Native alone; Male: 5 to 9 years | Population count | Number | 44 |
| b01001c\_005 | SEX BY AGE (AMERICAN INDIAN AND ALASKA NATIVE ALONE); Universe: People who are American Indian and Alaska Native alone; Male: 10 to 14 years | Population count | Number | 3 |
| b01001c\_006 | SEX BY AGE (AMERICAN INDIAN AND ALASKA NATIVE ALONE); Universe: People who are American Indian and Alaska Native alone; Male: 15 to 17 years | Population count | Number | 22 |
| b01001c\_007 | SEX BY AGE (AMERICAN INDIAN AND ALASKA NATIVE ALONE); Universe: People who are American Indian and Alaska Native alone; Male: 18 and 19 years | Population count | Number | 37 |
| b01001c\_018 | SEX BY AGE (AMERICAN INDIAN AND ALASKA NATIVE ALONE); Universe: People who are American Indian and Alaska Native alone; Female: Under 5 years | Population count | Number | 0 |
| b01001c\_019 | SEX BY AGE (AMERICAN INDIAN AND ALASKA NATIVE ALONE); Universe: People who are American Indian and Alaska Native alone; Female: 5 to 9 years | Population count | Number | 14 |
| b01001c\_020 | SEX BY AGE (AMERICAN INDIAN AND ALASKA NATIVE ALONE); Universe: People who are American Indian and Alaska Native alone; Female: 10 to 14 years | Population count | Number | 11 |
| b01001c\_021 | SEX BY AGE (AMERICAN INDIAN AND ALASKA NATIVE ALONE); Universe: People who are American Indian and Alaska Native alone; Female: 15 to 17 years | Population count | Number | 9 |
| b01001c\_022 | SEX BY AGE (AMERICAN INDIAN AND ALASKA NATIVE ALONE); Universe: People who are American Indian and Alaska Native alone; Female: 18 and 19 years | Population count | Number | 26 |
| b01001b\_001 | SEX BY AGE (BLACK OR AFRICAN AMERICAN ALONE); Universe: Black or African American alone; Total: Total: | Population count | Number | 13407 |
| b01001b\_003 | SEX BY AGE (BLACK OR AFRICAN AMERICAN ALONE); Universe: Black or African American alone; Male: Under 5 years | Population count | Number | 283 |
| b01001b\_004 | SEX BY AGE (BLACK OR AFRICAN AMERICAN ALONE); Universe: Black or African American alone; Male: 5 to 9 years | Population count | Number | 222 |
| b01001b\_005 | SEX BY AGE (BLACK OR AFRICAN AMERICAN ALONE); Universe: Black or African American alone; Male: 10 to 14 years | Population count | Number | 425 |
| b01001b\_006 | SEX BY AGE (BLACK OR AFRICAN AMERICAN ALONE); Universe: Black or African American alone; Male: 15 to 17 years | Population count | Number | 439 |
| b01001b\_007 | SEX BY AGE (BLACK OR AFRICAN AMERICAN ALONE); Universe: Black or African American alone; Male: 18 and 19 years | Population count | Number | 429 |
| b01001b\_018 | SEX BY AGE (BLACK OR AFRICAN AMERICAN ALONE); Universe: Black or African American alone; Female: Under 5 years | Population count | Number | 485 |
| b01001b\_019 | SEX BY AGE (BLACK OR AFRICAN AMERICAN ALONE); Universe: Black or African American alone; Female: 5 to 9 years | Population count | Number | 254 |
| b01001b\_020 | SEX BY AGE (BLACK OR AFRICAN AMERICAN ALONE); Universe: Black or African American alone; Female: 10 to 14 years | Population count | Number | 359 |
| b01001b\_021 | SEX BY AGE (BLACK OR AFRICAN AMERICAN ALONE); Universe: Black or African American alone; Female: 15 to 17 years | Population count | Number | 189 |
| b01001b\_022 | SEX BY AGE (BLACK OR AFRICAN AMERICAN ALONE); Universe: Black or African American alone; Female: 18 and 19 years | Population count | Number | 561 |
| b01001a\_001 | SEX BY AGE (WHITE ALONE); Universe: People who are White alone; Total: Total: | Population count | Number | 426175 |
| b01001a\_003 | SEX BY AGE (WHITE ALONE); Universe: People who are White alone; Male: Under 5 years | Population count | Number | 9461 |
| b01001a\_004 | SEX BY AGE (WHITE ALONE); Universe: People who are White alone; Male: 5 to 9 years | Population count | Number | 9446 |
| b01001a\_005 | SEX BY AGE (WHITE ALONE); Universe: People who are White alone; Male: 10 to 14 years | Population count | Number | 11409 |
| b01001a\_006 | SEX BY AGE (WHITE ALONE); Universe: People who are White alone; Male: 15 to 17 years | Population count | Number | 7231 |
| b01001a\_007 | SEX BY AGE (WHITE ALONE); Universe: People who are White alone; Male: 18 and 19 years | Population count | Number | 8352 |
| b01001a\_018 | SEX BY AGE (WHITE ALONE); Universe: People who are White alone; Female: Under 5 years | Population count | Number | 8754 |
| b01001a\_019 | SEX BY AGE (WHITE ALONE); Universe: People who are White alone; Female: 5 to 9 years | Population count | Number | 10226 |
| b01001a\_020 | SEX BY AGE (WHITE ALONE); Universe: People who are White alone; Female: 10 to 14 years | Population count | Number | 10640 |
| b01001a\_021 | SEX BY AGE (WHITE ALONE); Universe: People who are White alone; Female: 15 to 17 years | Population count | Number | 7731 |
| b01001a\_022 | SEX BY AGE (WHITE ALONE); Universe: People who are White alone; Female: 18 and 19 years | Population count | Number | 9368 |
| b01001g\_001 | SEX BY AGE (TWO OR MORE RACES); Universe: People who are Two or More Races; Total: Total: | Population count | Number | 12014 |
| b01001g\_003 | SEX BY AGE (TWO OR MORE RACES); Universe: People who are Two or More Races; Male: Under 5 years | Population count | Number | 732 |
| b01001g\_004 | SEX BY AGE (TWO OR MORE RACES); Universe: People who are Two or More Races; Male: 5 to 9 years | Population count | Number | 659 |
| b01001g\_005 | SEX BY AGE (TWO OR MORE RACES); Universe: People who are Two or More Races; Male: 10 to 14 years | Population count | Number | 823 |
| b01001g\_006 | SEX BY AGE (TWO OR MORE RACES); Universe: People who are Two or More Races; Male: 15 to 17 years | Population count | Number | 491 |
| b01001g\_007 | SEX BY AGE (TWO OR MORE RACES); Universe: People who are Two or More Races; Male: 18 and 19 years | Population count | Number | 501 |
| b01001g\_018 | SEX BY AGE (TWO OR MORE RACES); Universe: People who are Two or More Races; Female: Under 5 years | Population count | Number | 683 |
| b01001g\_019 | SEX BY AGE (TWO OR MORE RACES); Universe: People who are Two or More Races; Female: 5 to 9 years | Population count | Number | 650 |
| b01001g\_020 | SEX BY AGE (TWO OR MORE RACES); Universe: People who are Two or More Races; Female: 10 to 14 years | Population count | Number | 652 |
| b01001g\_021 | SEX BY AGE (TWO OR MORE RACES); Universe: People who are Two or More Races; Female: 15 to 17 years | Population count | Number | 410 |
| b01001g\_022 | SEX BY AGE (TWO OR MORE RACES); Universe: People who are Two or More Races; Female: 18 and 19 years | Population count | Number | 651 |
| b03002\_012 | HISPANIC OR LATINO ORIGIN BY RACE; Universe: Total population; Hispanic or Latino: Hispanic or Latino: | Population count | Number | 56886 |
| b03002\_013 | HISPANIC OR LATINO ORIGIN BY RACE; Universe: Total population; Hispanic or Latino: White alone | Population count | Number | 43689 |
| b03002\_014 | HISPANIC OR LATINO ORIGIN BY RACE; Universe: Total population; Hispanic or Latino: Black or African American alone | Population count | Number | 1753 |
| b03002\_015 | HISPANIC OR LATINO ORIGIN BY RACE; Universe: Total population; Hispanic or Latino: American Indian and Alaska Native alone | Population count | Number | 196 |
| b03002\_016 | HISPANIC OR LATINO ORIGIN BY RACE; Universe: Total population; Hispanic or Latino: Asian alone | Population count | Number | 253 |
| b03002\_017 | HISPANIC OR LATINO ORIGIN BY RACE; Universe: Total population; Hispanic or Latino: Native Hawaiian and Other Pacific Islander alone | Population count | Number | 150 |
| b03002\_018 | HISPANIC OR LATINO ORIGIN BY RACE; Universe: Total population; Hispanic or Latino: Some other race alone | Population count | Number | 7913 |
| b03002\_019 | HISPANIC OR LATINO ORIGIN BY RACE; Universe: Total population; Two or more races: Two or more races: | Population count | Number | 2932 |
| b15001\_011 | SEX BY AGE BY EDUCATIONAL ATTAINMENT FOR THE POPULATION 18 YEARS AND OVER; Universe: Population 18 years and over; 25 to 34 years: 25 to 34 years: | Population count | Number | 28989 |
| b15001\_017 | SEX BY AGE BY EDUCATIONAL ATTAINMENT FOR THE POPULATION 18 YEARS AND OVER; Universe: Population 18 years and over; 25 to 34 years: Bachelor's degree | Population count | Number | 7476 |
| b15001\_018 | SEX BY AGE BY EDUCATIONAL ATTAINMENT FOR THE POPULATION 18 YEARS AND OVER; Universe: Population 18 years and over; 25 to 34 years: Graduate or professional degree | Population count | Number | 2604 |
| b15001\_019 | SEX BY AGE BY EDUCATIONAL ATTAINMENT FOR THE POPULATION 18 YEARS AND OVER; Universe: Population 18 years and over; 35 to 44 years: 35 to 44 years: | Population count | Number | 24797 |
| b15001\_025 | SEX BY AGE BY EDUCATIONAL ATTAINMENT FOR THE POPULATION 18 YEARS AND OVER; Universe: Population 18 years and over; 35 to 44 years: Bachelor's degree | Population count | Number | 4891 |
| b15001\_026 | SEX BY AGE BY EDUCATIONAL ATTAINMENT FOR THE POPULATION 18 YEARS AND OVER; Universe: Population 18 years and over; 35 to 44 years: Graduate or professional degree | Population count | Number | 3258 |
| b15001\_027 | SEX BY AGE BY EDUCATIONAL ATTAINMENT FOR THE POPULATION 18 YEARS AND OVER; Universe: Population 18 years and over; 45 to 64 years: 45 to 64 years: | Population count | Number | 62253 |
| b15001\_033 | SEX BY AGE BY EDUCATIONAL ATTAINMENT FOR THE POPULATION 18 YEARS AND OVER; Universe: Population 18 years and over; 45 to 64 years: Bachelor's degree | Population count | Number | 11482 |
| b15001\_034 | SEX BY AGE BY EDUCATIONAL ATTAINMENT FOR THE POPULATION 18 YEARS AND OVER; Universe: Population 18 years and over; 45 to 64 years: Graduate or professional degree | Population count | Number | 8394 |
| b15001\_058 | SEX BY AGE BY EDUCATIONAL ATTAINMENT FOR THE POPULATION 18 YEARS AND OVER; Universe: Population 18 years and over; 25 to 34 years: Bachelor's degree | Population count | Number | 8124 |
| b15001\_059 | SEX BY AGE BY EDUCATIONAL ATTAINMENT FOR THE POPULATION 18 YEARS AND OVER; Universe: Population 18 years and over; 25 to 34 years: Graduate or professional degree | Population count | Number | 5237 |
| b15001\_060 | SEX BY AGE BY EDUCATIONAL ATTAINMENT FOR THE POPULATION 18 YEARS AND OVER; Universe: Population 18 years and over; 35 to 44 years: 35 to 44 years: | Population count | Number | 26186 |
| b15001\_066 | SEX BY AGE BY EDUCATIONAL ATTAINMENT FOR THE POPULATION 18 YEARS AND OVER; Universe: Population 18 years and over; 35 to 44 years: Bachelor's degree | Population count | Number | 6646 |
| b15001\_067 | SEX BY AGE BY EDUCATIONAL ATTAINMENT FOR THE POPULATION 18 YEARS AND OVER; Universe: Population 18 years and over; 35 to 44 years: Graduate or professional degree | Population count | Number | 5657 |
| b15001\_068 | SEX BY AGE BY EDUCATIONAL ATTAINMENT FOR THE POPULATION 18 YEARS AND OVER; Universe: Population 18 years and over; 45 to 64 years: 45 to 64 years: | Population count | Number | 67764 |
| b15001\_074 | SEX BY AGE BY EDUCATIONAL ATTAINMENT FOR THE POPULATION 18 YEARS AND OVER; Universe: Population 18 years and over; 45 to 64 years: Bachelor's degree | Population count | Number | 14062 |
| b15001\_075 | SEX BY AGE BY EDUCATIONAL ATTAINMENT FOR THE POPULATION 18 YEARS AND OVER; Universe: Population 18 years and over; 45 to 64 years: Graduate or professional degree | Population count | Number | 11986 |

ACS for Use with GEO3 Data

Table 27. ACS Pre-Processing Results File Layout – GEO3[[29]](#footnote-30)

| Variable Name | Format | Length |
| --- | --- | --- |
| Geography | Character | 5 |
| GEO3 | Character | 3 |
| State\_FIPS | Character | 2 |
| TOTAL\_ACS\_POPULATION | Number | 8 |
| AGE\_L5\_MALE\_WHITE | Number | 8 |
| AGE\_5\_9\_MALE\_WHITE | Number | 8 |
| AGE\_10\_14\_MALE\_WHITE | Number | 8 |
| AGE\_15\_17\_MALE\_WHITE | Number | 8 |
| AGE\_18\_19\_MALE\_WHITE | Number | 8 |
| AGE\_L5\_FEMALE\_WHITE | Number | 8 |
| AGE\_5\_9\_FEMALE\_WHITE | Number | 8 |
| AGE\_10\_14\_FEMALE\_WHITE | Number | 8 |
| AGE\_15\_17\_FEMALE\_WHITE | Number | 8 |
| AGE\_18\_19\_FEMALE\_WHITE | Number | 8 |
| AGE\_L5\_MALE\_BLACK | Number | 8 |
| AGE\_5\_9\_MALE\_BLACK | Number | 8 |
| AGE\_10\_14\_MALE\_BLACK | Number | 8 |
| AGE\_15\_17\_MALE\_BLACK | Number | 8 |
| AGE\_18\_19\_MALE\_BLACK | Number | 8 |
| AGE\_L5\_FEMALE\_BLACK | Number | 8 |
| AGE\_5\_9\_FEMALE\_BLACK | Number | 8 |
| AGE\_10\_14\_FEMALE\_BLACK | Number | 8 |
| AGE\_15\_17\_FEMALE\_BLACK | Number | 8 |
| AGE\_18\_19\_FEMALE\_BLACK | Number | 8 |
| AGE\_L5\_MALE\_ASIAN | Number | 8 |
| AGE\_5\_9\_MALE\_ASIAN | Number | 8 |
| AGE\_10\_14\_MALE\_ASIAN | Number | 8 |
| AGE\_15\_17\_MALE\_ASIAN | Number | 8 |
| AGE\_18\_19\_MALE\_ASIAN | Number | 8 |
| AGE\_L5\_FEMALE\_ASIAN | Number | 8 |
| AGE\_5\_9\_FEMALE\_ASIAN | Number | 8 |
| AGE\_10\_14\_FEMALE\_ASIAN | Number | 8 |
| AGE\_15\_17\_FEMALE\_ASIAN | Number | 8 |
| AGE\_18\_19\_FEMALE\_ASIAN | Number | 8 |
| AGE\_L5\_MALE\_OTHER | Number | 8 |
| AGE\_5\_9\_MALE\_OTHER | Number | 8 |
| AGE\_10\_14\_MALE\_OTHER | Number | 8 |
| AGE\_15\_17\_MALE\_OTHER | Number | 8 |
| AGE\_18\_19\_MALE\_OTHER | Number | 8 |
| AGE\_L5\_FEMALE\_OTHER | Number | 8 |
| AGE\_5\_9\_FEMALE\_OTHER | Number | 8 |
| AGE\_10\_14\_FEMALE\_OTHER | Number | 8 |
| AGE\_15\_17\_FEMALE\_OTHER | Number | 8 |
| AGE\_18\_19\_FEMALE\_OTHER | Number | 8 |
| AGE\_25\_64\_BACH\_GRAD | Number | 8 |
| AGE\_25\_64\_BACH\_GRAD\_GTR20PERC | Number | 8 |
| TOTAL\_LATIN | Number | 8 |
| LATIN\_WHITE | Number | 8 |
| LATIN\_BLACK | Number | 8 |
| LATIN\_ASIAN | Number | 8 |
| LATIN\_OTHER | Number | 8 |

###### EHR File Layouts

EHR Input File Layout

EHR GEO3 Data

User provided EHRs are imported in CODI-PQ and require a csv file with the following variable names, variable value options (case sensitive), and in the order listed below.

Table 28. EHR Input File Layout for GEO3-Level Programs, CSV File[[30]](#footnote-31)

| Variable Name | Description | Format | Valid values | Example |
| --- | --- | --- | --- | --- |
| SUBJID | Patient Identifier | Character | Character value of maximum length 25 | 123456789 |
| SEX\_NUM | Sex of patient where 0 is male, 1 is female | Number | 0  1 | 0 |
| AGEYEARS | Age of patient in years at the time of the medical encounter | Number | Count of years | 11 |
| RACE\_ETH | Patient’s race if known or ethnicity when race is not known | Character | “AFRICAN AMERICAN”  “Black”  “ASIAN”  “Asian”  “CAUCASIAN”  “White”  “HISPANIC”  “Hispanic”  “OTHER”  “Other”  “UNKNOWN”  “Unknown” | CAUCASIAN |
| STATE\_ABR | Patient’s residential state, two-letter state abbreviations | Character | any postal abbreviation of state (see appendix G for possible values) | VA |
| GEO3 | Either: Patient’s residential a) county code or b) ZIP-3 | Number | Any numeric value | 221 |
| WEIGHT\_CATEGORY | Patient’s BMI percentile. See section A.2 and A.11 for more information. | Character | Normal or Healthy Weight  Obese  Overweight  Severe Obesity  Underweight | Overweight |
| YEAR | Year of the medical encounter | Number | Yyyy | 2018 |
| SCDCNT | Sickle-Cell indicator | Number | Program treats patients with a count of 1 or higher as having sickle cell disease. If sickle cell disease information is not available, set this value to blank or zero. | 2 |
| PREGNANCY\_FLAG | Pregnancy flag | Number | 0/1  If pregnancy information is not available, set this value to zero or blank. | 1 |
| ZIP | 5-digit ZIP code | Character | 5-digit ZIP code. Required for county level records. Optional for ZCTA-3 level records. | 20814 |

EHR Results File Layout for GEO3

Table 29. EHR Pre-Processing Results File Layout – GEO3[[31]](#footnote-32)

|  |  |  |
| --- | --- | --- |
| Variable Name | Format | Example |
| SUBJID | Character | 12345626 |
| Ageyr | Number | 16 |
| AGE\_CATEGORIES | Character | 15 – 17 |
| WTCAT | Character | (2) Healthy Weight (5th to <85th percentile) |
| STATE\_ALPHA | Character | MI |
| STATE\_FIPS | Character | 48 |
| ZIP | Character | 20184 |
| GEO3 | Character | 100 |
| Geography | Character | 48100 |
| SCDCNT | Number | 0 |
| PREGNANCY\_FLAG | Number | 1 |
| Race | Character | Black |
| Sex | Character | Female |
| Year | Number | 2018 |
| SCD | Number | 0 |
| Imputed\_Race | Character | Black |
| Race\_Imputed | Number | 0 |

###### CODI-PQ-GEO3 Example SAS Programs

Data Inputs and Link Population Data (Pre-processing) Quickstart with GEO3 Data

Appendix E.2 includes a program to generate results. This example is for the pre-processing data inputs and link population data step.

Text highlighted in yellow has been reviewed and approved or reviewed and edited from its original values. The program uses the data inputs: ACS\_State\_ZCTA3 and EHR\_filename located in the P:\Example\0\_Raw\_Data folder. The file processes EHRs between 2016 and 2019 and creates a SAS file named CODI\_EHR\_READY stored in the folder P:\Example\2\_Output\Pre\_Processed\_CODI\_PQ. Both the raw data for ACS and EHR file include ZCTA3, not county codes (COUNTY = N). The SAS log is stored in P:\Example\2\_Output\SAS LOG\This\_is\_ the\_Log <plus date and time information>.log. Text between /\* and \*/ are comments in SAS. Comments may vary slightly in CODI-PQ from the example below.

/\*Note: subsection of the full program. Be sure to only edit this section but submit the full program. \*/

/\*\*\*\*\*\*\*\*\*\*\*\*\*\*\*\*\*\*\*\*\*\*\*\*\*\*\*\*\*\*\*\*\*\*\*\*\*\*\*\*\*\*\*\*\*\*\*\*\*\*\*\*\*\*\*\*\*\*\*\*\*\*\*\*\*\*\*\*\*\*\*\*\*\*\*\*\*\*\*\*\*\*\*\*\*\*\*\*\*\*\*\*\*\*\*\*\*\*\*\*\*\*\*\*\*\*\*\*\*\*\*\*\*\*\*\*\*\*\*\*\*\*\*\*\*\*\*\*\*\*\*\*/

/\*\*\*\*\*\*\*\*\*\*\*\*\*\*\*\*\*\*\*\*\*\*\*\*\*\*\*\*\*\*\*\*\*\*\*\*\*\*\*\*\* -- USER SELECTION CRITERA SECTIONS 1 through 4 -- \*\*\*\*\*\*\*\*\*\*\*\*\*\*\*\*\*\*\*\*\*\*\*\*\*\*\*\*\*\*\*\*\*\*\*\*\*\*\*\*/

/\*\*\*\*\*\*\*\*\*\*\*\*\*\*\*\*\*\*\* -- PLEASE UPDATE THE BLACK TEXT AFTER THE EQUAL SIGN (ACCEPTED VALUES LISTED IN SAS NOTE) -- \*\*\*\*\*\*\*\*\*\*\*\*\*\*\*\*\*\*\*/

/\*SECTION 1: Input Folder and file names\*\*\*/

/\*\*\*/ %LET ROOT\_PRE = P:\Example; /\*@Note: base directory (ACCEPTABLE VALUES: computer directory name) \*\*\*/

\*\*\*/

/\*\*\*/ %LET PRE\_DEST = CODI\_PQ; /\*@Note: Suffix name for EHR Output folder (ACCEPTABLE VALUES: folder name (no punctuation) \*\*\*/

/\*\*\*/ %LET ACS\_FILENAME = ACS\_State\_ZCTA3; /\*@Note: ACS file name (ACCEPTABLE VALUES: file name, do not include ".csv") \*\*\*/

/\*\*\*/ %LET EHR\_FILENAME = EHR\_filename; /\*@Note: EHR file name (ACCEPTABLE VALUES: file name, do not include ".csv") \*\*\*/

/\*\*\*/ %LET LOG\_NAME\_PRE = This\_is\_the\_Log; /\*@Note: SAS log file name prefix ACCEPTABLE VALUES: SAS file name (no punctuation) \*\*\*/

/\*SECTION 2: Beginning and End Year of longitudinal EHR data \*\*\*/

/\*\*\*/ %LET BEGIN\_YEAR = 2016; /\*@Note: LONGITUDINAL Start year (ACCEPTABLE VALUES: 4-digit numeric year) \*\*\*/

/\*\*\*/ %LET END\_YEAR = 2019; /\*@Note: LONGITUDINAL End year (ACCEPTABLE VALUES: 4-digit numeric year) \*\*\*/

/\*SECTION 3: OPTIONAL Output File Name Suffix \*\*\*/

/\*\*\*/ %LET EHR\_PRE\_Out = CODI\_EHR\_READY; /\*@Note: EHR output file name (ACCEPTABLE VALUES: SAS file name (no punctuation) \*\*\*/

/\*SECTION 4: County or ZCTA3 data (REQUIRED) \*\*\*/

/\*\*\*/ %LET COUNTY=N; /\*@Note: County/ZCTA3 indicator (ACCEPTABLE VALUES: Y for County level data, N for ZCTA3 level data \*\*\*/ /\*\*\*\*\*\*\*\*\*\*\*\*\*\*\*\*\*\*\*\*\*\*\*\*\*\*\*\*\*\*\*\*\*\*\*\*\*\*\*\*\*\*\*\*\*\*\*\*\*\*\*\*\*\*\*\*\*\*\*\*\*\*\*\*\*\*\*\*\*\*\*\*\*\*\*\*\*\*\*\*\*\*\*\*\*\*\*\*\*\*\*\*\*\*\*\*\*\*\*\*\*\*\*\*\*\*\*\*\*\*\*\*\*\*\*\*\*\*\*\*\*\*\*\*\*\*\*\*\*\*\*\*/

/\*\*\*Note: Root directory includes subfolders: "..\0\_Raw\_Data"

"..\1\_SAS\_Programs"

"..\1\_SAS\_Programs\Pre\_Processing\_GEO3"

"..\1\_SAS\_Programs\CODI\_PQ\_GEO3"

"..\2\_Output" and

"..\2\_Output\SAS LOGS" \*\*\*/

/\*\*\*NOTE: SAS programs must be stored in the PROGS directory including: Macro1-CODI\_PQ.sas,

Macro2-CODI\_PQ.sas,

Macro3-CODI\_PQ.sas,

Macro4-CODI\_PQ.sas,

Module1-CODI\_PQ.sas,

Module2-CODI\_PQ.sas,

Macro1-CODI\_PQ-Co\_occurring.sas,

Macro2-CODI\_PQ-Co\_occurring.sas,

Macro3-CODI\_PQ-Co\_occurring.sas,

Macro4-CODI\_PQ-Co\_occurring.sas,

Module1-CODI\_PQ-Co\_occurring.sas,

Module2-CODI\_PQ-Co\_occurring.sas, \*\*\*/

/\*NOTE: query output is stored as a csv file in "..\2\_Output" named after a time/date stamp and CODI\_Prevalence\_Query\_Report \*\*\*/

/\*\*\*\*\*\*\*\*\*\*\*\*\*\*\*\*\*\*\*\*\*\*\*\*\*\*\*\*\*\*\*\*\*\*\*\*\*\*\*\*\*\*\*\*\*\*\*\*\*\*\*\*\*\*\*\*\*\*\*\*\*\*\*\*\*\*\*\*\*\*\*\*\*\*\*\*\*\*\*\*\*\*\*\*\*\*\*\*\*\*\*\*\*\*\*\*\*\*\*\*\*\*\*\*\*\*\*\*\*\*\*\*\*\*\*\*\*\*\*\*\*\*\*\*\*\*\*\*\*\*\*\*/

/\*\*\*STOP STOP STOP STOP STOP STOP STOP STOP STOP STOP STOP STOP STOP STOP STOP STOP STOP STOP STOP STOP STOP STOP STOP STOP STOP\*/

/\*\*\* DO NOT EDIT BEYOND THIS POINT DO NOT EDIT BEYOND THIS POINT DO NOT EDIT BEYOND THIS POINT \*/

/\*\*\*\*\*\*\*\*\*\*\*\*\*\*\*\*\*\*\*\*\*\*\*\*\*\*\*\*\*\*\*\*\*\*\*\*\*\*\*\*\*\*\*\*\*\*\*\*\*\*\*\*\*\*\*\*\*\*\*\*\*\*\*\*\*\*\*\*\*\*\*\*\*\*\*\*\*\*\*\*\*\*\*\*\*\*\*\*\*\*\*\*\*\*\*\*\*\*\*\*\*\*\*\*\*\*\*\*\*\*\*\*\*\*\*\*\*\*\*\*\*\*\*\*\*\*\*\*\*\*\*\*/

/\*\*\*\*\*\*\*\*\*\*\*\*\*\*\*\*\*\*\*\*\*\*\*\*\*\*\*\*\*\*\*\*\*\*\*\*\*\*\*\*\*\*\*\*\*\*\*\*\*\*\*\*\*\*\*\*\*\*\*\*\*\*\*\*\*\*\*\*\*\*\*\*\*\*\*\*\*\*\*\*\*\*\*\*\*\*\*\*\*\*\*\*\*\*\*\*\*\*\*\*\*\*\*\*\*\*\*\*\*\*\*\*\*\*\*\*\*\*\*\*\*\*\*\*\*\*\*\*\*\*\*\*/

/\*\*\*\*\*\*\*\*\*\*\*\*\*\*\*\*\*\*\*\*\*\*\*\*\*\*\*\*\*\*\*\*\*\*\*\*\*\*\*\*\*\*\*\*\*\*\*\*\*\*\*\*\*\*\*\*\*\*\*\*\*\*\*\*\*\*\*\*\*\*\*\*\*\*\*\*\*\*\*\*\*\*\*\*\*\*\*\*\*\*\*\*\*\*\*\*\*\*\*\*\*\*\*\*\*\*\*\*\*\*\*\*\*\*\*\*\*\*\*\*\*\*\*\*\*\*\*\*\*\*\*\*/

/\*Note: subsection of the full program. Be sure to only edit this section but submit the full program. \*/

Generate Results Example with GEO3 Data

Appendix E.1 includes a program to generate pre-processed results that are then used in this program.

Text highlighted in yellow has been reviewed and approved or reviewed and edited from its original values. The program uses the data inputs: ACS\_State\_ZCTA3 and CODI\_EHR\_READY located in the P:\Example\2\_Output\Pre\_Processed\_CODI\_PQ folder.

Subpopulation selected by the user includes: EHR records from 2017 including patients 2 to 14 years of age who are either white or Asian, living in Jefferson County (059) Colorado (FIPS code = 08) or Yuma County (125) Colorado (FIPS code = 08) see: <https://www.nrcs.usda.gov/wps/portal/nrcs/detail/national/home/?cid=nrcs143_013697> to determine the correct value (08059 is combined state and County). Note that Sex\_Male and Sex\_Female have not been edited since the ALL\_SEXES is turned on to yes.

The file processes a SAS file named CODI\_EHR\_READY stored in the folder P:\Example\2\_Output\Pre\_Processed\_CODI\_PQ. The SAS log is stored in P:\Example\2\_Output\SAS LOG\CODI\_PQ\_ZCTA3\_PEDS<plus date and time information>.log. Text between /\* and \*/ are comments in SAS. Comments may vary slightly in CODI-PQ from the example below.

Methods: Include imputed race information and calculate the age-adjusted prevalence.

/\*Note: subsection of the full program. Be sure to only edit this section, but submit the full program. \*/

/\*\*\*\*\*\*\*\*\*\*\*\*\*\*\*\*\*\*\*\*\*\*\*\*\*\*\*\*\*\*\*\*\*\*\*\*\*\*\*\*\* -- USER SELECTION CRITERA SECTIONS 1 through 5 -- \*\*\*\*\*\*\*\*\*\*\*\*\*\*\*\*\*\*\*\*\*\*\*\*\*\*\*\*\*\*\*\*\*\*\*\*\*\*\*\*\*/

/\*\*\*\*\*\*\*\*\*\*\*\*\*\*\*\*\*\*\* -- PLEASE UPDATE THE BLACK TEXT AFTER THE EQUAL SIGN (ACCEPTED VALUES LISTED IN SAS NOTE) -- \*\*\*\*\*\*\*\*\*\*\*\*\*\*\*\*\*\*\*\*/

/\*SECTION 1: Folder and file names \*\*\*/

/\*\*\*/ %LET ROOT\_PQ = P:\Example; /\*@Note: same as in pre-processing quickstart program, base directory (ACCEPTABLE VALUES: computer directory name) \*\*\*/

/\*\*\*/ %LET PRE\_DEST = CODI\_PQ; /\*@Note: same as in pre-processing quickstart program, Suffix name for EHR Output folder (ACCEPTABLE VALUES: folder name (no puctuation) \*\*\*/

/\*\*\*/ %LET EHR\_PRE\_OUT = CODI\_EHR\_Ready; /\*@Note: same as in pre-processing quickstart program, name the pre-processing output file (ACCEPTABLE VALUES: file name (no punctuations)) \*\*\*/

/\*\*\*/ %LET LOG\_NAME = CODI\_PQ\_ZCTA3\_PEDS; /\*@Note: Name for SAS log storage location \*\*\*/

/\*\*\*/ %LET FileOUT\_Name = Results\_Here; /\*@Note: Output file name; MUST BE FEWER THAN 28 CHARACTERS \*\*\*/

/\*SECTION 2: Subset data based on specifications INCLUDING YEAR, GEOGRAPHY, STATE, STATE/ZCTA3, or STATE/COUNTY \*\*\*/

/\*\*\*/ %LET BEG\_YEAR = 2017; /\*@Note: Beginning year of analysis (ACCEPTED VALUES: 4-Digit numeric, 2015-2019) \*\*\*/

/\*\*\*/ %LET END\_YEAR = 2017; /\*@Note: End year of analysis (ACCEPTED VALUES: 4-Digit numeric, 2015-2019) \*\*\*/

/\*\*\*/ %LET ALL\_STATES = N; /\*@Note: Include all geographical locations in file?(ACCEPTED VALUES: Y/N) \*\*\*/

/\*\*\*/ %LET ALL\_AGES = N; /\*@Note: Include all age ranges? (ACCEPTED VALUES: Y/N) \*\*\*/

/\*\*\*/ %LET ALL\_SEXES = Y; /\*@Note: Include all sex values? (ACCEPTED VALUES: Y/N) \*\*\*/

/\*\*\*/ %LET ALL\_RACES = N; /\*@Note: Include all race categories? (ACCEPTED VALUES: Y/N) \*\*\*/

/\*SECTION 3: Additional flags \*\*\*/

/\*\*\*/ %LET ACSCOUNTY = N; /\*@Note: Is the ACS data at the county or ZCTA3 level? (ACCEPTABLE VALUES: Y for County level data, N for ZCTA3 level data) \*\*\*/

/\*\*\*/ %LET INCLUDE\_PREGNANCY = Y; /\*@Note: Include pregnant individuals and non-pregnant individuals (Y) or non-pregnant individuals only (N) (ACCEPTED VALUES: Y/N) \*\*\*/

/\*\*\*/ %LET SAMPLE\_CHECK = Y; /\*@Note: Run check for sufficient sample size of query (ACCEPTED VALUES: Y/N) \*\*\*/

/\*SECTION 4: Only complete section 4 for any "N" values listed in section 2 \*\*\*/

/\*IF ALLGEOGRAPHIES= N THEN SELECT STATE CODES OR STATE AND COUNTY CODES BELOW: \*\*\*/

/\*\*\* (ACCEPTED VALUES: SINGLE QUOTES SURROUNDING 2 OR 5-Digit CODES w/ "," BETWEEN MULTIPLE SELECTIONS, ) \*\*\*/

/\*IF ALLSTATES = N THEN SELECT ONE OR MORE AGE CATEGORIES BELOW: \*\*\*/

/\*\*\*/ %LET GEO\_GROUP = ZCTA3; /\*@Note: Level of geography (ACCEPTED VALUES: STATE, ZCTA3, COUNTY) \*\*\*/

/\*\*\*/ %LET GEO\_LIST = %STR('08059', '08125'); /\*@Note: IF GEO\_GROUP="STATE" then populate with State FIPS code(s) for example, 36 is New York, If GEO\_GROUP="ZCTA3" then populate with FIPS State+ZCTA3 code(s), \*\*\*/

/\*If Geo\_Group="COUNTY" then populate with FIPS State+FIPS County code(s) (ACCEPTED VALUES: 2-digit state FIPS or 5-digit state FIPS+ZCTA3 \*\*\*/

/\*or 5-digit state FIPS+county FIPS (Must be surrounded by single quotation and comma delimited))\*\*\*/

/\*IF ALL\_AGES = N THEN SELECT ONE OR MORE AGE CATEGORIES BELOW: \*\*\*/

/\*\*\*/ %LET WGT\_AGE\_2\_4 = Y; /\*@Note: Age Range: 2 to 4 (ACCEPTED VALUES: Y/N) \*\*\*/

/\*\*\*/ %LET WGT\_AGE\_5\_9 = Y; /\*@Note: Age Range: 5 to 9 (ACCEPTED VALUES: Y/N) \*\*\*/

/\*\*\*/ %LET WGT\_AGE\_10\_14 = Y; /\*@Note: Age Range: 10 to 14 (ACCEPTED VALUES: Y/N) \*\*\*/

/\*\*\*/ %LET WGT\_AGE\_15\_17 = N; /\*@Note: Age Range: 15 to 17 (ACCEPTED VALUES: Y/N) \*\*\*/

/\*\*\*/ %LET WGT\_AGE\_18\_19 = N; /\*@Note: Age Range: 18 to 19 (ACCEPTED VALUES: Y/N) \*\*\*/

/\*IF ALL\_RACES = N THEN SELECT ONE OR MORE RACE BELOW: \*\*\*/

/\*\*\*/ %LET RACE\_WHITE = Y; /\*@Note: White (ACCEPTED VALUES: Y/N) \*\*\*/

/\*\*\*/ %LET RACE\_BLACK = N; /\*@Note: Black/African American (ACCEPTED VALUES: Y/N) \*\*\*/

/\*\*\*/ %LET RACE\_ASIAN = Y; /\*@Note: Asian (ACCEPTED VALUES: Y/N) \*\*\*/

/\*\*\*/ %LET RACE\_OTHER = N; /\*@Note: Other (ACCEPTED VALUES: Y/N) \*\*\*/

/\*IF ALL\_SEXES = N THEN SELECT MALE OR FEMALE BELOW: \*\*\*/

/\*\*\*/ %LET SEX\_MALE = Y; /\*@Note: Sex: Male (ACCEPTED VALUES: Y/N) \*\*\*/

/\*\*\*/ %LET SEX\_FEMALE = Y; /\*@Note: Sex: Female (ACCEPTED VALUES: Y/N) \*\*\*/

/\*SECTION 5: Methodological option selections \*\*\*/

/\*\*\*/ %LET IMP\_RACES = Y; /\*@Note: Include imputed race values? (ACCEPTED VALUES: Y/N) \*\*\*/

/\*\*\*/ %LET AGE\_ADJ = Y; /\*@Note: Produce age-adjusted estimates? (ACCEPTED VALUES: Y/N) \*\*\*/

/\*\*\*\*\*\*\*\*\*\*\*\*\*\*\*\*\*\*\*\*\*\*\*\*\*\*\*\*\*\*\*\*\*\*\*\*\*\*\*\*\*\*\*\*\*\*\*\*\*\*\*\*\*\*\*\*\*\*\*\*\*\*\*\*\*\*\*\*\*\*\*\*\*\*\*\*\*\*\*\*\*\*\*\*\*\*\*\*\*\*\*\*\*\*\*\*\*\*\*\*\*\*\*\*\*\*\*\*\*\*\*\*\*\*\*\*\*\*\*\*\*\*\*\*\*\*\*\*\*\*\*\*\*/

/\*\*\*Note: Root directory includes subfolders:

"..\0\_Raw\_Data"

"..\1\_SAS\_Programs"

"..\2\_Output" and

"..\2\_Output\SAS LOGS" \*\*\*/

/\*\*\*NOTE: SAS programs must be stored in the PROGS directory including:

Macro1-CODI\_PQ.sas,

Macro2-CODI\_PQ.sas,

Macro3-CODI\_PQ.sas,

Macro4-CODI\_PQ.sas,

Module1-CODI\_PQ.sas,

Module2-CODI\_PQ.sas\*\*\*/

/\*\*\*NOTE: query output is stored as a csv file in "..\2\_Output" named after a time/date stamp and the designated FileOUT\_Name \*\*\*/

/\*\*\*NOTE: to save a SAS file of the weighted observations remove comment from lines 273-276 \*\*\*/

/\*\*\*\*\*\*\*\*\*\*\*\*\*\*\*\*\*\*\*\*\*\*\*\*\*\*\*\*\*\*\*\*\*\*\*\*\*\*\*\*\*\*\*\*\*\*\*\*\*\*\*\*\*\*\*\*\*\*\*\*\*\*\*\*\*\*\*\*\*\*\*\*\*\*\*\*\*\*\*\*\*\*\*\*\*\*\*\*\*\*\*\*\*\*\*\*\*\*\*\*\*\*\*\*\*\*\*\*\*\*\*\*\*\*\*\*\*\*\*\*\*\*\*\*\*\*\*\*\*\*\*\*\*/

/\*\*\*STOP STOP STOP STOP STOP STOP STOP STOP STOP STOP STOP STOP STOP STOP STOP STOP STOP STOP STOP STOP STOP STOP STOP STOP STOP STOP\*/

/\*\*\* DO NOT EDIT BEYOND THIS POINT DO NOT EDIT BEYOND THIS POINT DO NOT EDIT BEYOND THIS POINT DO NOT EDIT BEYOND THIS POINT \*/

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###### CODI-PQ Results

Example BMI Percentile Prevalence

Once complete, CODI-PQ generates prevalence results as a csv file. Table 30 provides an overview of the variables included for BMI percentile prevalence, and Tables 31 and 34 provide example results. User inputs, error codes, sources of technical documentation, caveats, and a possible citation begins with the row labeled order 3 and continues thereafter. The number of rows and exact wording of text will vary based on the criteria selected.

Table 30. CODI-PQ Results Data Dictionary

|  |  |
| --- | --- |
| Column | Description |
| Order | Row order |
| Weight Category | A categorical value based on BMI percentile. |
| Sample | The observed (or unadjusted, or crude) count of youth and teens in the study population. |
| Population | The weighted (or adjusted) count of the study population. |
| Crude Prevalence | The observed (or unadjusted, or crude) prevalence in the study population. |
| Crude Prevalence Standard Error | The observed (or unadjusted, or crude) standard error in the study population. |
| Weighted Prevalence | Prevalence based on weighted counts. A sample weight is assigned to each sample patient. It is a measure of the number of youth and teens in the population represented by that sample patient. See implementation guide, Appendix A. Sample Weights for more information. |
| Weighted Prevalence Standard Error | Standard error based on weighted counts. See implementation guide, Appendix A. Variance for more information. |
| Age-Adjusted Prevalence | Prevalence based on weighted, age-adjusted counts. See implementation guide, Appendix A. Age Adjustment for more information. |

Table 31. Results Example from Synthetic Data[[32]](#footnote-33)

| Order | Weight Category | Sample | Population | Crude Prevalence | Crude Prevalence Standard Error | Weighted Prevalence | Weighted Prevalence Standard Error | Age-Adjusted Prevalence | Age-Adjusted Prevalence Standard Error |
| --- | --- | --- | --- | --- | --- | --- | --- | --- | --- |
| 1 | (1) Underweight (<5th percentile) | 61 | 5,801 | 4.16 | 0.25 | 4.80 | 0.82 | 4.80 | 0.82 |
| 1 | (2) Healthy Weight (5th to <85th percentile) | 824 | 82,020 | 56.17 | 1.3 | 67.88 | 3.58 | 67.88 | 3.58 |
| 1 | (3) Overweight (85th to <95th percentile) | 251 | 14,752 | 17.11 | 0.98 | 12.21 | 2.42 | 12.21 | 2.42 |
| 1 | (4) Obesity (>95th percentile) | 331 | 18,255 | 22.56 | 1.09 | 15.11 | 2.89 | 15.11 | 2.89 |
| 1 | (4b) Severe Obesity (>120% of the 95th percentile) | 100 | 6,085 | 7.52 | 1.01 | 5.25 | 2.22 | 5.25 | 2.22 |
| 2 | Totals: | 1,467 | 120,828 |  |  |  |  |  |  |
| 3 | Dataset: AEMR, 2015-2019 |  |  |  |  |  |  |  |  |
| 4 | Query Parameters: AGE RACE SEX GEOGRAPHY YEAR |  |  |  |  |  |  |  |  |
| 5 | AGE: (10 - 14, 15 - 17, 18 - 19) |  |  |  |  |  |  |  |  |
| 6 | SEX: (Female) |  |  |  |  |  |  |  |  |
| 7 | RACE: (White, Black, Asian, Other) |  |  |  |  |  |  |  |  |
| 8 | RACE Suppressed: No. |  |  |  |  |  |  |  |  |
| 9 | RACE Imputed: People with unknown race were excluded. |  |  |  |  |  |  |  |  |
| 10 | Geography: (08810) |  |  |  |  |  |  |  |  |
| 11 | Year: 2019 |  |  |  |  |  |  |  |  |
| 12 | Weighting cells were collapsed for: (Geography) |  |  |  |  |  |  |  |  |
| 13 | AGE adjusted?: (Yes) |  |  |  |  |  |  |  |  |
| 13 | Error Codes: (None) |  |  |  |  |  |  |  |  |
| 14 | Technical Documentation: See https://github.com/NORC-UChicago/CODI-PQ for more information and full details on data sources and methodologies. |  |  |  |  |  |  |  |  |
| 15 | Query Date: Friday, 1 July 2021 4:09:46 PM |  |  |  |  |  |  |  |  |
| 15 | Suggested Citation: Tanenbaum, E., Campbell, S., Chelluri, D., Zalsha, S., Boim, J., Paddock, S., Copeland, K. (2021). Clinical and Community Data Initiative Prevalence Query (CODI-PQ) SAS programs (version 2015-2019). |  |  |  |  |  |  |  |  |
| 16 | Caveats |  |  |  |  |  |  |  |  |
| 17 | Patients with either missing or invalid age, sex, height, weight, or geography are not included in counts and prevalence results. |  |  |  |  |  |  |  |  |
| 18 | The method used to calculate the standard errors are documented in the technical documentation. |  |  |  |  |  |  |  |  |
| 19 | The population are based on age-race-sex-location specific counts from the 2014-2018 American Community Survey Five-year Estimates released by the Census Bureau on December 19, 2019. |  |  |  |  |  |  |  |  |

Table 32. Example Results with Errors (Insufficient Sample Size), Error Messages Are Shown in Order Row 14[[33]](#footnote-34)

| Order | Weight Category | Sample | Population | Crude Prevalence | Crude Prevalence Standard Error | Weighted Prevalence | Weighted Prevalence Standard Error | Age-adjusted Prevalence | Age-adjusted Prevalence Standard Error |
| --- | --- | --- | --- | --- | --- | --- | --- | --- | --- |
| 1 | (1) Underweight (<5th percentile) |  |  |  |  |  |  |  |  |
| 1 | (2) Healthy Weight (5th to <85th percentile) |  |  |  |  |  |  |  |  |
| 1 | (3) Overweight (85th to <95th percentile) |  |  |  |  |  |  |  |  |
| 1 | (4) Obesity (>95th percentile) |  |  |  |  |  |  |  |  |
| 1 | (4b) Severe Obesity (>120% of the 95th percentile) |  |  |  |  |  |  |  |  |
| 2 | Totals: |  |  |  |  |  |  |  |  |
| 3 | Dataset: IQVIA, 2015-2019 |  |  |  |  |  |  |  |  |
| 4 | Query Parameters: AGE RACE SEX GEOGRAPHY YEAR |  |  |  |  |  |  |  |  |
| 5 | AGE: (02 - 04) |  |  |  |  |  |  |  |  |
| 6 | SEX: (Female) |  |  |  |  |  |  |  |  |
| 7 | RACE: (Other) |  |  |  |  |  |  |  |  |
| 8 | RACE Suppressed: (Error) |  |  |  |  |  |  |  |  |
| 9 | RACE Imputed: (Error) of race values were imputed. Please be advised, prevalence may incur additional bias with imputed race values. Extreme caution is encouraged when the proportion of imputed race values exceeds 40%. |  |  |  |  |  |  |  |  |
| 10 | Geography: (11) District of Columbia |  |  |  |  |  |  |  |  |
| 11 | Years: 2017 - 2018 |  |  |  |  |  |  |  |  |
| 12 | Weighting cells were collapsed for: (Error) |  |  |  |  |  |  |  |  |
| 13 | AGE adjusted: (Yes) |  |  |  |  |  |  |  |  |
| 14 | Error Codes: (Current selection criteria return an insufficient number of youth and teens and do not meet minimum threshold to create sample weights. Ensure that selections are correct (e.g., correct list of state codes or ZCTA-3values) or include additional geographic or demographic categories (e.g., add additional communities or include additional or all races, age groups, sex, etc.).) |  |  |  |  |  |  |  |  |
| 15 | Technical Documentation: See https://github.com/NORC-UChicago/CODI-PQ for more information and full details on data sources and methodologies. |  |  |  |  |  |  |  |  |
| 16 | Query Date: Friday, August 27, 2021 1:49:38 PM |  |  |  |  |  |  |  |  |
| 17 | Suggested Citation: Tanenbaum, E., Campbell, S., Chelluri, D., Zalsha, S., Boim, J., Paddock, S., Copeland, K. (2021). Clinical and Community Data Initiative Prevalence Query (CODI-PQ) SAS programs (version 2015-2019) |  |  |  |  |  |  |  |  |
| 18 | Caveats |  |  |  |  |  |  |  |  |
| 19 | Patients with either missing or invalid age, sex, height, weight, or geography are not included in counts and prevalence results. |  |  |  |  |  |  |  |  |
| 20 | The method used to calculate the standard errors are documented in the technical documentation. |  |  |  |  |  |  |  |  |
| 21 | The population is based on age-race-sex-location specific counts from the 2014-2018 American Community Survey Five-year Estimates released by the Census Bureau on December 19, 2019. |  |  |  |  |  |  |  |  |
| 22 | The method used to calculate age-adjusted prevalence is documented in the technical documentation. |  |  |  |  |  |  |  |  |

Possible Result Errors

There are several reasons that CODI-PQ may not produce some or all results as described in the table that follows.

Table 33: CODI-PQ Results Error Codes

| Error | Description |
| --- | --- |
| One or more demographic or geographic category has no groups selected. One or more group must be selected in each category. Please ensure each demographic and geographic category has one or more groups selected (e.g., age group, select an age range for inclusion). | One or more categories are not selected. For example, a minimum of one year, sex, age group, geography, and racial group must be selected (Y). See Step 2.4.4 for more details. |
| Years are out of scope. CODI-PQ was developed between 2019 and 2021, see Implementation Guide for more details. | Starting year cannot be before 2000, ending year cannot be after 2030. CODI-PQ may be inappropriate to implement on medical encounters outside of 2015 through 2021. Please review the methodology in full to determine whether CODI-PQ is appropriate for your needs. |
| Geographic level (GEO\_GROUP) has been left blank or has been set to an unacceptable value. To remedy issue, please update the GEO\_GROUP variable to either STATE, ZCTA-3, or county. | The observed (or unadjusted, or crude) count of youth and teens in the study population. |
| State and/or GEO3 is incorrectly specified. Review the lists and ensure each value is: Surrounded by quotations, comma delimited, and/or the correct length (e.g., "08001", "08002", "08003", etc.). | Ensure the GEO\_LIST is set to the correct format. 1. State is a FIPS number, not a state abbreviation, 2. All numbers must be in single quotes, 3. There is a space and a comma whenever selecting multiple locations, and 4. The text is within the function %STR( );  Examples:  If GEO\_GROUP = STATE; /\*@Note: Level of geography (ACCEPTED VALUES: STATE/ZCTA3, or STATE/COUNTY) \*\*\*/  /\*\*\*/ %LET GEO\_LIST = %STR('08', ‘10’);  If GEO\_GROUP = STATE/ZCTA3; /\*@Note: Level of geography (ACCEPTED VALUES: STATE/ZCTA3, or STATE/COUNTY) \*\*\*/  /\*\*\*/ %LET GEO\_LIST = %STR('51221', ‘51224’);  If GEO\_GROUP = STATE/COUNTY; /\*@Note: Level of geography (ACCEPTED VALUES: STATE/ZCTA3, or STATE/COUNTY) \*\*\*/  /\*\*\*/ %LET GEO\_LIST = %STR('08001', ‘08002’); |
| Current selections return an insufficient number of patients and do not meet minimum threshold to estimate sample weights. Consider including additional demographic categories (e.g., races, age groups, sex), geographies, or years. | To determine the demographic group(s) with insufficient sample size, consider running the optional sample check (Sample\_Check = Y). |
| Iterative proportional fitting weighting routine has failed to converge. Please revise selection criteria and rerun algorithm. | Weighting is not possible using iterative proportionate fitting under certain circumstances. For example, according to a SAS SUGI paper, (Izrael, 2004)  “Oh and Scheuren [4] note that the available convergence proofs make strong assumptions about the cell counts in the cross-classification of the raking variables – that no cells are empty or that some particular combination of nonempty cells is present. They recommend setting up the raking problem in a “sensible” manner to avoid: 1) imposing too many marginal constraints on the sample, 2) defining marginal categories that contain a small percentage of the sample, and 3) imposing contradictory constraints on the sample.  …  Convergence may be slow if 1) any categories contain fewer than 5% of the sample cases, 2) the size of the difference between each control total and the weighted sample margin prior to raking. If some differences are large, the number of iterations will typically be higher.” |
| A SAS error has occurred within the algorithm. Review the SAS log or contact a system administrator for further assistance. | SAS errors occur when syntax is not properly specified. Common reasons for SAS errors include missing semi-colons, single or double quotes, mismatched quotes, deleting the “/\*” that is before a comment or “\*/” after a comment, etc., etc. In addition to reviewing your SAS code and log, consider contacting SAS technical support, and/or make a new copy of the software from Github. |

Additional messages may be displayed but are not indicative of an error. For example, the percentage of persons with imputed race that are included in the prevalence estimates.

Table 34: CODI-PQ Results Error Codes

|  |  |
| --- | --- |
| Comment | Description |
| RACE Imputed: (Error) of race values were imputed. Please be advised, prevalence may incur additional bias with imputed race values. Extreme caution is encouraged when the proportion of imputed race values exceeds 40%. | If the user allows records with imputed race to be included in the analysis, then the percentage of records (crude) with imputed race is reported in the results. |
| Weighting cells were consolidated for: | Statistical weighting is conducted by age group, race, sex, and geography. If the sample size is insufficient in an age group or geography, weighting cells may be collapsed (combined). Race and sex do not allow for consolidation of weighting cells. |
| CODI PQ was developed between 2019 and 2021 and tested with EHR from 2015 through 2019. Please review the Implementation Guide in full to determine whether CODI-PQ methodology is appropriate for your use case when used outside of these date ranges. | Users may choose to employ CODI-PQ outside of the testing period. It is recommended that the user carefully review all methods prior to doing so. |

Example Sample Checker Results

CODI-APQ generates optional sample size checker results in the SAS output or results window. Table 38 provides an example results table. Factors include age categories, race, and sex. The values include all demographics selected by the user.

For example, the results below were generated based on the user’s selections of only including persons ages 2 – 4, race either of Black or Asian, and both male and female. The user may choose to exclude males and rerun the analysis if partial information is of interest. Why? Results show see that male has an insufficient number of patients as well as Asian. CODI-APQ automatically removes racial categories if the sample size is insufficient (see A.6 Statistical Weights) but does not remove male or female automatically.

Table 35: CODI-PQ Sample Size Checker Results

|  |  |  |
| --- | --- | --- |
| **Checker\_Results** | **Factor** | **Value** |
| Sample Size Is Sufficient | Age Categories | 2 - 4 |
| Sample Size Is Sufficient | Race | Black |
| Sample Size Is Insufficient | Race | Asian |
| Sample Size Is Sufficient | Sex | Female |
| Sample Size Is Insufficient | Sex | Male |

###### State FIPS codes

Note: for a list of all state and county codes, visit USDA’s website: <https://www.nrcs.usda.gov/wps/portal/nrcs/detail/national/home/?cid=nrcs143_013697>

Table 36. State FIPS Codes

| Name | Postal Code | FIPS |
| --- | --- | --- |
| Alabama | AL | 01 |
| Alaska | AK | 02 |
| Arizona | AZ | 04 |
| Arkansas | AR | 05 |
| California | CA | 06 |
| Colorado | CO | 08 |
| Connecticut | CT | 09 |
| Delaware | DE | 10 |
| District of Columbia | DC | 11 |
| Florida | FL | 12 |
| Georgia | GA | 13 |
| Hawaii | HI | 15 |
| Idaho | ID | 16 |
| Illinois | IL | 17 |
| Indiana | IN | 18 |
| Iowa | IA | 19 |
| Kansas | KS | 20 |
| Kentucky | KY | 21 |
| Louisiana | LA | 22 |
| Maine | ME | 23 |
| Maryland | MD | 24 |
| Massachusetts | MA | 25 |
| Michigan | MI | 26 |
| Minnesota | MN | 27 |
| Mississippi | MS | 28 |
| Missouri | MO | 29 |
| Montana | MT | 30 |
| Nebraska | NE | 31 |
| Nevada | NV | 32 |
| New Hampshire | NH | 33 |
| New Jersey | NJ | 34 |
| New Mexico | NM | 35 |
| New York | NY | 36 |
| North Carolina | NC | 37 |
| North Dakota | ND | 38 |
| Ohio | OH | 39 |
| Oklahoma | OK | 40 |
| Oregon | OR | 41 |
| Pennsylvania | PA | 42 |
| Rhode Island | RI | 44 |
| South Carolina | SC | 45 |
| South Dakota | SD | 46 |
| Tennessee | TN | 47 |
| Texas | TX | 48 |
| Utah | UT | 49 |
| Vermont | VT | 50 |
| Virginia | VA | 51 |
| Washington | WA | 53 |
| West Virginia | WV | 54 |
| Wisconsin | WI | 55 |
| Wyoming | WY | 56 |

###### Glossary

**ACS** – American Community Survey. CODI-PQ relies on ACS population counts for statistical weighting.

**AEMR** – Ambulatory Electronic Medical Record. Used to test CODI-PQ.

**AEMR-US** – Ambulatory United States Electronic Medical Record data

Source: AEMR-US version 5 OMOP 5 (Aug 2019 release) accessed through the E360TM Software-as-a-Service (SaaS) Platform.

**Age-Adjusted Prevalence** – Is a prevalence that controls for the effects of differences in population age distributions. When comparing across geographic areas, age adjusting is typically used to control for the influence that different population age distributions might have on health encounter prevalences. Age-adjustment (or age standardization) is the same as calculating a weighted average. It weights the age-specific prevalence observed in a population of interest by the proportion of each age group in a standard population. The standard population are published by the CDC and represent the U.S. 2000 population in each age group.

**Age Groups** – Age groups include ages 2 to 4, 5 to 9, 10 to 14, 15 and 17, and 18 to 19 years of age.

**BMI** – Body Mass Index. Used to categorize a person’s height and weight into various categories (e.g., underweight, overweight, etc.).

**BMI Percentile** – Categorization of a youth or teen’s height, weight, age, and sex into one of five categories: underweight, healthy weight, overweight, obesity, and/or severe obesity.

**CDC** – Centers for Disease Control and Prevention

**CDM** – Common Data Model

**Census Tract** – Small, relatively permanent statistical subdivisions of a county or equivalent entity that are updated prior to each decennial census. The primary purpose of census tracts is to provide a stable set of geographic units for the presentation of statistical data. Census tracts generally have a population size between 1,200 and 8,000 people.

**CODI** – Previously the “Childhood Obesity Data Initiative” currently the “The Clinical and Community Data Initiative.” CODI 1.0 and 2.0 are projects led by the Centers for Disease Control and Prevention originally designed to enhance data capacity for users interested in exploring the efficacy of weight-related intervention and prevention strategies.

**CODI-PQ** – CODI prevalence query (CODI\_PQ in SAS programs)

**CODI-PQ-GEO3** – CODI PQ applied on EHR with state and a three digit geographic identifier

**Converge** – Property (exhibited by the statistical weighting function) of approaching a limit more and more closely as an argument (variable) of the function increases or decreases or as the number of terms of the series increases. Crude Prevalence of BMI percentile – is the total number of people within a particular BMI percentile (e.g., severe obesity) in a specified geographic area (state, county, ZCTA-3, etc.) for a specified group of people (age, race, or all people) divided by the total population for the same geographic area and same specified group for a specific time period (e.g., 2016) and multiplied by 100.

**COUNTY Data –** When referenced in all capital letters, it refers to EHRs linked to a patient’s state and county FIPS code.

**CSV** – Comma Separated Value. All input files should be in CSV.

**DHDN** – Distributed Health Data Network

**EHR** – Electronic Health Records. Digital records of patient health information. An EHR contains the patient's records from multiple providers and provides a more holistic, long-term view of a patient's health.

**EMR** – Electronic Medical Records. Digital records of patient health information. A digital version of a patient's chart.

**Execute** – In SAS software is the process by which a computer or virtual machine executes the instructions of a computer program. The term run is used synonymously in SAS. A related definition refers to the specific action of a user starting, launching, or invoking a program.

**FFRDC** – Federally Funded Research and Development Center

**FIPS Codes** – Numbers which uniquely identify geographic areas. The number of digits in FIPS codes vary depending on the level of geography. State-level FIPS codes have two digits, county-level FIPS codes have five digits of which the first two digits are the FIPS code of the state to which the county belongs followed by three digits which represent a county within the state.

**Geographic Area** – Geographic area is defined based on either 1) the state and county or 2) the state and ZCTA-3.

**GEO3** – Geographic area identified by three numbers. GEO3 is defined based on either the state and 1) county or 2) ZCTA-3.

**Growthcleanr** – An open-source R package for assessing height and weight record data from EHR systems, focused on categorizing the plausibility of individual record based on longitudinal analysis of each patient subject.

**Health FFRDC** – Centers for Medicare & Medicaid Services Alliance to Modernize Healthcare federally funded research and development center.

**Healthy Weight** – Body Mass Index value 5th percentile to less than the 85th percentile

**Informed Presence** – The belief that patients do not randomly go to the provider’s office and thus are not randomly included in EHRs.

**Imputation** – Estimating a value for a specific data item (e.g., race) where the response is missing or unusable.

**Iterative Proportional Fitting** – (IPF or raking) is an iterative algorithm for proportionally adjusting a matrix or contingency table of non-negative elements to produce a new 'similar' table with specified positive marginal totals in at least two dimensions.

**MSE** – Mean Squared Error

**NCHS** – National Center for Health Statistics

**NHANES** – National Health and Nutrition Examination Survey, a probability-based survey that might be more representative of the general population.

**Obesity** – 95th percentile to less than 120 percent of the body mass index (BMI) value for the 95th percentile.

**Overweight** – Body Mass Index value 85th to less than the 95th percentile

**Open-Access Program** – a program made freely available to libraries and end users

**Open-Source Program** – a program made freely available to libraries and end users, written in software that is free of charge.

**PCORnet** – Patient Centered Outcomes Research Network

**Pre-processing CODI-PQ** – a set of SAS programs that are executed once and only once per AEMR data file. It is also known as the data inputs and link population data.

**Prevalence** – proportion of a particular population found to be affected by a medical condition at a specific time.

**PUF** – Public Use File

**Quickstart** – a SAS program which requires user input. Only the Quickstart programs are needed along with user specifications to run the pre-processing and/or the PQ.

**Race Imputation** – Imputing missing race data, see also imputation. Setting race imputation to yes allows the programs to include all available EHRs for youth and teens even if the medical record did not include a known race. See Imputation for further clarification.

**Random Sample** - a method of selecting a sample from a population in such a way that every possible sample that could be selected has a predetermined probability of being selected.

**RDM** – CODI Research Data Model

**RLDM** – CODI Record Linkage Data Model

**Run** – in SAS software is the process by which a computer or virtual machine executes the instructions of a computer program. The term execute is used synonymously. A related definition refers to the specific action of a user starting, launching, or invoking a program.

**SAS –** SAS is a statistical software suite

**Sample** – The observed (or unadjusted, or crude) count of youth and teens in the study population.

**SDOH** – Social Determinants of Health

**Severe Obesity** – 120 percent or greater of the BMI value for the 95th percentile

**Statistical Weights** – A statistical weight is an amount given to increase or decrease the importance of an item. Weights are commonly given for people when a sample and not a census is taken. The value of the weight can be thought of as denoting the number of youth and teens in the population represented by that sample person in EHR, accounting for differences between the distribution of the sample and total populations.

Note: the use of statistical weights is encouraged for all analyses because the data comes from a nonprobability sample with no known probabilities of selection. Failure to use statistical weights may yield biased results and overstated significance levels.

**Suppression/Presentation Guidelines for Proportions** – Guidelines used by all of HHS which provide criteria for presenting or suppressing proportions. The multistep NCHS Data Presentation Standards for Proportions are based on a minimum denominator sample size and on criteria based on the absolute and relative widths of a CI calculated using the Clopper-Pearson method.

**Synthea** – An open-source, synthetic patient generator that models the medical history of synthetic patients.

**Teen –** A person aged 13 to 19

**Underweight** – Body Mass Index value less than 5th percentile

**Variance** – A measure of how far a set of numbers is spread out from their average value

**Weight Category** – Categorization of a person’s height, weight, age, and sex (BMI percentile) into one of five categories: underweight, healthy weight, overweight, obesity, and/or severe obesity.

**Weights** – See Statistical Weights or Weight Category

**Weighted Prevalence** – Prevalence based on weighted counts where are equal to crude prevalence with statistical weights applied.

**Youth** – A person aged 2 to 12

**ZCTA-3** – The first three digits of a ZIP code tabulation area (ZCTA3 in SAS)

**ZCTA-3 data** – Refers to a data file of ER linked to a patient’s ZIP-3 and thus ZCTA-3

###### Abbreviations and Acronyms

| ACRONYM | DEFINITION |
| --- | --- |
| ACS | American Community Survey |
| ADHD | Attention Deficit Hyperactivity Disorder |
| AEMR | Ambulatory Electronic Medical Record |
| BMI | Body Mass Index |
| CDC | Centers for Disease Control and Prevention |
| CI | Confidence Interval |
| CODI | Clinical and Community Data Initiative |
| CODI-PQ | Clinical and Community Data Initiative Prevalence Query |
| CSV | Comma Separated Value |
| DHDN | Distributed Health Data Network |
| EHR | Electronic Health Record |
| EMR | Electronic Medical Record |
| FFRDC | Federally Funded Research and Development Center |
| HHS | U.S. Department of Health and Human Services |
| IG | Implementation Guide |
| IPW | Inverse-Probability Weighting |
| MSE | Mean Square Error |
| NCHS | National Center for Health Statistics |
| NHANES | National Health and Nutrition Examination Survey |
| PUF | Public Use File |
| SAS | A Statistical Software Suite |
| SDOH | Social Determinants of Health |
| SFTP | Secured File Transfer Protocol |
| ZCTA | ZIP Code Tabulation Area |

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1. DHDNs are traditionally networks of medical organizations that map their EHRs concepts to a common data model so that users can query similar information across organizations. [↑](#footnote-ref-2)
2. [About Child & Teen BMI | Healthy Weight, Nutrition, and Physical Activity | CDC](https://www.cdc.gov/healthyweight/assessing/bmi/childrens_bmi/about_childrens_bmi.html) [↑](#footnote-ref-3)
3. CODI data partners are organizations and institutions which facilitate CODI data exchange by contributing and hosting data that can be accessed through the CODI infrastructure for queries and other research or programmatic uses of the data. [↑](#footnote-ref-4)
4. IQVIA's Ambulatory Electronic Medical Record (AEMR-US) database contains de-identified medical records and encounter from 44,000 physicians and 315 networks in the U.S. covering the period from January 2006 through May 2019. These data include provider medical specialty, patient variables such as examination data, year of birth, gender, race and ethnicity, and medical variables such as diagnoses, procedures, medication prescriptions records, and patient and family history captured during a patient encounter. Contributing practices consist of medium to large physician offices, outpatient clinics, and physician groups. Because examination date and year of birth, but not age, were available, age was calculated from the examination date and the midpoint of the birth year (July 2). [↑](#footnote-ref-5)
5. <https://synthetichealth.github.io/synthea/> [↑](#footnote-ref-6)
6. Non-probability sample is a group of individuals based on a sampling method in which not all members of the population have an equal chance of being a part of the sample. In probability sampling, each member of the population has a known chance of being selected. Thus, probability sampling is more stringent than non-probability sampling. [↑](#footnote-ref-7)
7. ACS 2019 file for use with CODI-PQ is available for download from https://sft.mitre.org/#/folder/6281923. The 2019 ACS data was used for model calibration. Use of other years of ACS data requires recalibration of the model due to changes in population counts. [↑](#footnote-ref-8)
8. SAS outputs a dot (.) instead of a numeric value when results are suppressed. Suppression occurs by row and may include one or more than one row of results. [↑](#footnote-ref-9)
9. Note: Pre-Processing is labeled Pre\_Processing\_GEO3 within the folder. Prevalence query is labeled CODI\_PQ\_GEO3 within the folder. [↑](#footnote-ref-10)
10. Note: The sickle cell disease indicator is not date sensitive since it is used for imputing race. Thus, the value can be calculated across all available data years and reconciled across years. Example: patient is identified with sickle cell disease in 2016. All records, regardless of year, for this patient would be identified as having sickle cell disease. [↑](#footnote-ref-11)
11. Note: If more than one year is selected, the first record of each SUBJID is kept with all subsequent records excluded from prevalence results to meet statistical weighting assumptions. [↑](#footnote-ref-12)
12. Klein & Schoenborn, 2001. [↑](#footnote-ref-13)
13. ACS 2019 file for use with CODI-PQ is available for download from https://sft.mitre.org/#/folder/6281923. The 2019 ACS data was used for model calibration. Use of other years of ACS data requires recalibration of the model due to changes in population counts. [↑](#footnote-ref-14)
14. CDC provided Ambulator Electronic Medical Record data under a Data Use Agreement with the Health FFRDC. [↑](#footnote-ref-15)
15. The Synthea package is based on Walonoski, et al., 2017 and is available at: <https://synthetichealth.github.io/synthea/>. [↑](#footnote-ref-16)
16. ZCTAs are areal representations of ZIP code service areas created by the Census Bureau. Approximately 99% of ZIP-3’s with a population greater than zero are equal to ZCTA-3 and thus ZCTA-3 and ZIP-3 are used interchangeably within the analysis. [↑](#footnote-ref-17)
17. Annual Estimates of the Resident Population by Sex, Age, Race, and Hispanic Origin for the United States: April 1, 2010, to July 1, 2019 (NC-EST2019-ASR6H); Source: U.S. Census Bureau, Population Division; Release Date: June 2020. [↑](#footnote-ref-18)
18. Little, 1993. [↑](#footnote-ref-19)
19. Wolter, 2007. [↑](#footnote-ref-20)
20. Parker et al., 2017. [↑](#footnote-ref-21)
21. The complement of a proportion p is (1 – p). The complement of the number of encounters in the numerator for p is the number of encounters in the numerator for (1 – p). [↑](#footnote-ref-22)
22. Wolter, 2007. [↑](#footnote-ref-23)
23. Note: prevalence of obesity will include two categories: those that are category 4 and 4a. [↑](#footnote-ref-24)
24. <https://www.census.gov/programs-surveys/geography/guidance/geo-areas/zctas.html> [↑](#footnote-ref-25)
25. Best & Shepherd, 2020. [↑](#footnote-ref-26)
26. Parker JD, Talih M, Malec DJ, et al, 2017. [↑](#footnote-ref-27)
27. Within this report, the term measurement means the process by which we describe and ascribe meaning to the key concepts we are investigating. At its core, measurement is about defining one’s terms in as clear and precise a way as possible. [↑](#footnote-ref-28)
28. The CODI SDOH Prioritization report is available at <https://github.com/NORC-UChicago/CODI-PQ>. [↑](#footnote-ref-29)
29. The variables are needed at a minimum for the CODI-PQ results. Actual ACS file may include additional variables. [↑](#footnote-ref-30)
30. One record per patient per year, thus a patient may be included multiple times in the EHR if multiple years are on the file. [↑](#footnote-ref-31)
31. The variables are needed at a minimum for the CODI-PQ results. Actual file may include additional variables. [↑](#footnote-ref-32)
32. Note: borders and shading are for demonstration purposes only. CSV exports columns separated with a comma. The results can be imported into Excel. Results based on synthetic data. Exact wording in Order 3 through 19 may vary from those shown in the example. [↑](#footnote-ref-33)
33. Note: borders and shading are for demonstration purposes only. CSV exports columns separated with a comma. The results can be imported into Excel. Results based on synthetic data. Exact wording in Order 3 through 22 may vary from those shown in the example. [↑](#footnote-ref-34)