



**The Mexican Health and Aging Study (MHAS/ENASEM):
Master Follow-up File 2001, 2003, 2012, 2015 and 2018**

Version 1

July 22, 2020

“The MHAS (Mexican Health and Aging Study) is partly sponsored by the National Institutes of Health/National Institute on Aging (grant number NIH R01AG018016) in the US and the *Instituto Nacional de Estadística y Geografía* (INEGI) in Mexico. Data files and documentation are public use and available at www.MHASweb.org.”

Suggested citation for this document:

MHAS (2020), “The Mexican Health and Aging Study (MHAS/ENASEM): Master File Follow-up 2001, 2003, 2012, 2015 and 2018. Version 1, July 2020.” Retrieved from www.MHASweb.org on (date).

Table of Contents

I. Introduction	1
II. Study Description	1
III. Structure of the MHAS Master Follow-up File	2
A. Identifiers	3
1. Unique Household ID (Variable name: CUNICAH, also called UNHHID)	3
2. Person Identifiers (Variable name: NP, CODENT01 in 2001, and CODENT03 in 2003).....	3
2001 Person Identification Code (Variable name: CODENT01)	4
2003 Person Identification Code (Variable name: CODENT03)	4
New Person identifier (Variable name: NP).....	4
Unique Person ID (Variable name: UNHHIDNP)	7
3. Sub-household ID (Variable name: ACTHOG and SUBHOG_XX)	7
Updated Household ID (Variable name: ACTHOG)	7
Sub-household ID (Variable name: SUBHOG_XX).....	7
B. Outcomes of the Interview, Sample and Study Status	9
1. Eligible (Variable name: ELEGIBLE_XX).....	9
2. Follow-New person (Variable name: NEW_SAMPLE_XX)	9
3. Type of non-interview (Variable name: TIPNE_XX)	10
4. Type of interview (Variable name: TIPENT_XX)	12
5. Reason for Proxy (Variable name: REASON_PROXY_XX)	13
6. Interview Date (Variable name: INT_DATE_XX).....	13
7. Wave Status – Alive or Deceased (Variable name: FALLECIDO_XX)	13
C. Other	14
1. Same dwelling (Variable name: MISMA_V_XX)	14
2. Interview Technique – Electronic versus Paper (Variable name: C_PAP_XX)	14
3. High Migration States (Variable name: EAM_XX).....	14
4. Age and Sex (Variable name: AGE_XX and SEX_XX).....	14
D. Weights	14
E. Sub-Samples and Ancillary Study	15
1. Selected for Anthropometric Measures 2001 & 2003 (Variable name: ANTRO_XX)	15
2. Result for Anthropometric Measures 2001 & 2003 (Variable name: RES_ANTRO_XX)	15
3. Sub-sample 2012 (Variable name: SUBSAMPLE_12).....	15
4. Result for Biomarkers Measures 2012 (Variable name: RES_BIOMARKERS_12)	15
5. Ancillary Study Variables: Cognitive Aging Study (Mex-Cog)	16
Mex-Cog 2016 Sub-sample (Variable name: SUBSAMPLE_16)	16
Mex-Cog 2016 Phase (Variable name: PHASE_MXCOG_16)	16
Mex-Cog 2016 Result for Each Component (Variables name: RES_MXCOG_16, RES_COGNITIVO_16, RES_INFORMANTE_16, RES_ANTRO_16)	16
Mex-Cog 2016 Other Variables (Variables name: RES_BIOMARKERS_16, EDTA_16, HBA1C_16)	17
C. Accessing the MHAS Data	17
D. Merging the MHAS Data	18
Appendix 1. Master Follow-up File Content	19
Appendix 2. Master Follow-up File Codebook	22

I. Introduction

This document describes the follow-up master file of the Mexican Health and Aging Study (MHAS/ENASEM) with information regarding the identifiers, the outcome of the interviews, weights, and related variables for the 2001, 2003, 2012, 2015, and 2018 waves of the study.

This descriptive file enhances but does not preclude the information provided in the previous “Follow-Up Master File For The Mexican Health And Aging Study (MHAS/ENASEM) 2001-2003” file (available [here](#)) and “Follow-Up Master File For The Mexican Health And Aging Study (MHAS/ENASEM) 2001-2012” (available [here](#)).

This file was designed to provide information at the individual level, starting from the individuals selected for the baseline survey in 2001 and including the new cohorts added in 2012 and 2018. The file includes the new samples added in the third and fifth waves. The file also includes a record for individuals that were selected but never interviewed, as well as those eligible and not eligible for re-interview in 2003, 2012, 2015 or 2018.

II. Study Description¹

The 2001 baseline survey of the Mexican Health and Aging Study (MHAS/ENASEM) is a national representative survey of individuals born prior to 1951—that is, the population aged 50 or older as of the year 2001. The study protocols and survey instruments are highly comparable to the U.S. Health and Retirement Study (HRS).

The MHAS is designed to examine the aging process and evaluate the impact of disease on health, function, and mortality of adults over the age of 50 living in Mexico. The baseline survey was conducted in the summer of 2001, and a follow-up visit to the same individuals was carried out in the summer of 2003. The MHAS baseline sample was selected from residents of both rural and urban areas, from the National Employment Survey (Encuesta Nacional de Empleo, ENE), carried out by the INEGI (Instituto Nacional de Estadística y Geografía) in Mexico. Households with at least one resident of ages 50 or older were eligible to be part of the MHAS baseline sample. If more than one person was age-eligible in the selected households, then one person was selected at random for the study. If the selected MHAS person was married or in a

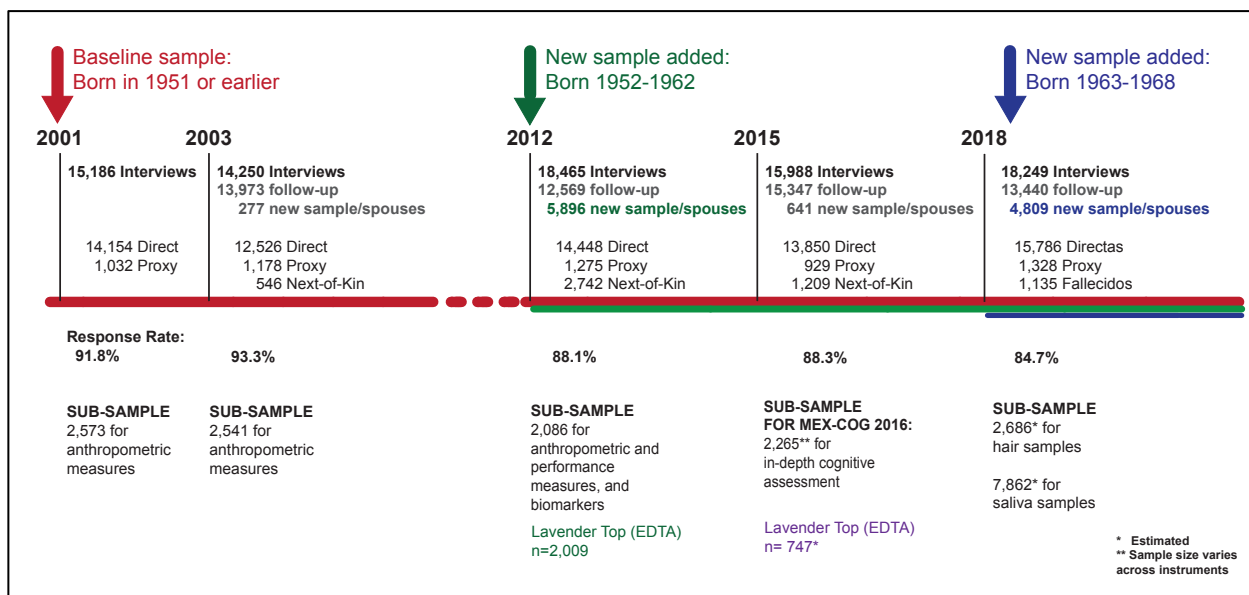
¹ Wong R, Michaels-Obregon A, Palloni A. Cohort Profile: The Mexican Health and Aging Study (MHAS). *Int. J. Epidemiol.* (2015). First published online: January 27, 2015. doi: 10.1093/ije/dyu263

consensual union, with the spouse residing in the same household, then the spouse or partner was also recruited to be part of the MHAS regardless of his/her age.

The 2012 survey was conducted in the fall of 2012, and a follow-up visit was carried out in the fall of 2015. In addition, the 2012 sample was refreshed by adding a representative sample of the population from the 1952-1962 birth cohorts, as well as their spouses/partners regardless of age. Similar to the baseline interview, the sampling frame for the new cohort sample was the Mexican National Employment and Occupation Survey (ENOE, previously named National Employment Survey, ENE) 2012.

The 2018 survey was completed in the fall of 2018, and a follow-up visit will be completed in 2021. Similar to the previous surveys, the 2018 sample was refreshed by adding a representative sample of the population from the 1963-1968 birth cohorts, as well as their spouses/partners regardless of age. The sampling frame for the new cohort sample was INEGI’s National Directory of Households.

Figure 1. MHAS 2001-2018 Timeline Sample Size and Response Rates Across Waves



III. Structure of the MHAS Master Follow-up File

The 2018 Master Follow-up File contains a record for every subject included in the MHAS sample (n=28,303), all the identifiers used in each wave and the cover variables. The sample includes

a record of selected individuals eligible for re-interview in 2003, 2012, 2015, or 2018, those not eligible for re-interview in 2003, 2012, or 2015, in addition to samples added in the third and fifth wave (2012 and 2018, respectively). The file contains all the identifiers used in each wave and survey variables including: outcome of the interviews, sampling weights, and other useful variables. Thus the 2018 Master Follow-up File serves as main file to merge any data within and across all the MHAS waves.

There are 106 variables in 2018 Master Follow-up File. The file contents and codebook are included in Appendix 1 and 2, respectively. Below we provide a brief description of the variables included in the data file.

A. Identifiers

The first set of variables in the Master Follow-up File are the household and person identifiers used in all the waves. The following is a short description of each household and individual identifier for each wave, and an explanation on how to use them to merge the data file within and across waves.

1. Unique Household ID (Variable name: CUNICAH, also called UNHHID)

In 2001, a random sequential number ranging from 1 to 11,000 was created to identify each household at baseline (CUNICAH also called UNHHID).

The new sample added in 2012 was assigned a value starting from 11,001 to 15,130 to identify each new household, while the new sample added in 2018 was assigned a value starting from 15,131 to 20,218 to identify each new household.

2. Person Identifiers (Variable name: NP, CODENT01 in 2001, and CODENT03 in 2003)

Within each household, in the initial interview, there are up to two persons under study (the selected person of eligible age, and spouse regardless of age). Once interviewed, the two individuals become part of the longitudinal study.

2001 Person Identification Code (Variable name: CODENT01)

In the baseline interview, the unique person ID (CODENT01) is assigned in the household as follows (this distinction is important, as it determines the sample weight for each person):

1 = Selected individual,

2 = Spouse.

2003 Person Identification Code (Variable name: CODENT03)

In 2003, these two values were dragged from 2001 with each respondent, and we added two possible values to create CODENT03:

3 = New spouse of the person who was = 1 in 2001, and

4 = New spouse of the person who was = 2 in 2001

In combination with the unique household ID in 2001, and the updated household ID in 2003, these codes uniquely identify individual subjects of study.

TIPS: How to use identifiers to merge 2001 and 2003 data files

In 2001, the variable CUNICAH serves as unique household identifier. This variable, in combination with the person identifier CODENT01 (also called PS3) serve as a unique person identifier for the first wave.

In 2003, the unique household identifier CUNICAH used in 2001 is supplemented with ACTHOG to form the unique household identifier. Also, these variables in combination with the person identifier for 2003 given by CODENT03 (also called ENT2) serve as a unique person identifier for the second wave.

New Person identifier (Variable name: NP)

Starting in 2012, the variable NP was created to identify each person in each original household, as follows:

NP=010 (selected person), NP=020 (spouse of selected person)

Figure 1. Assignment of NP values within the 010 family

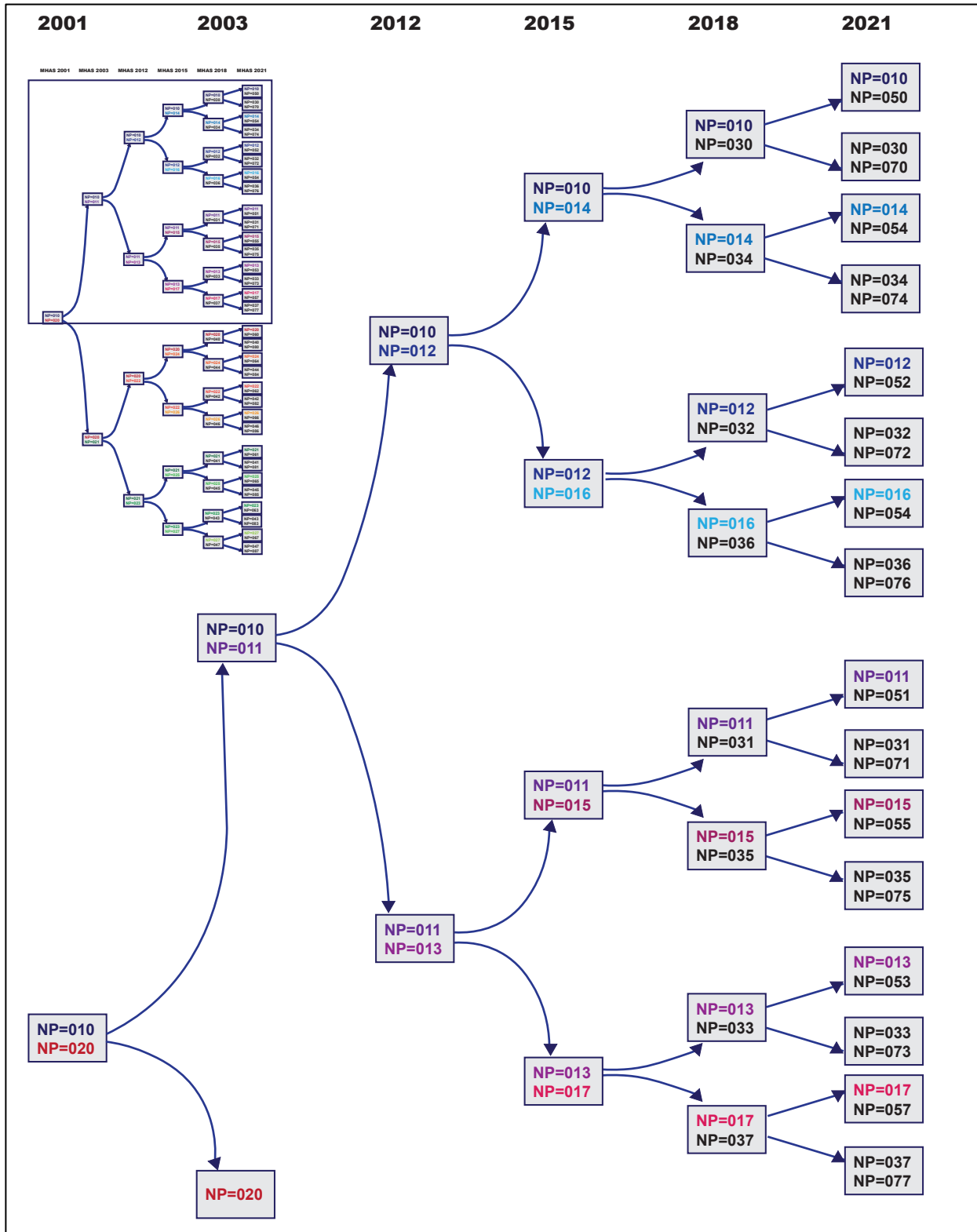
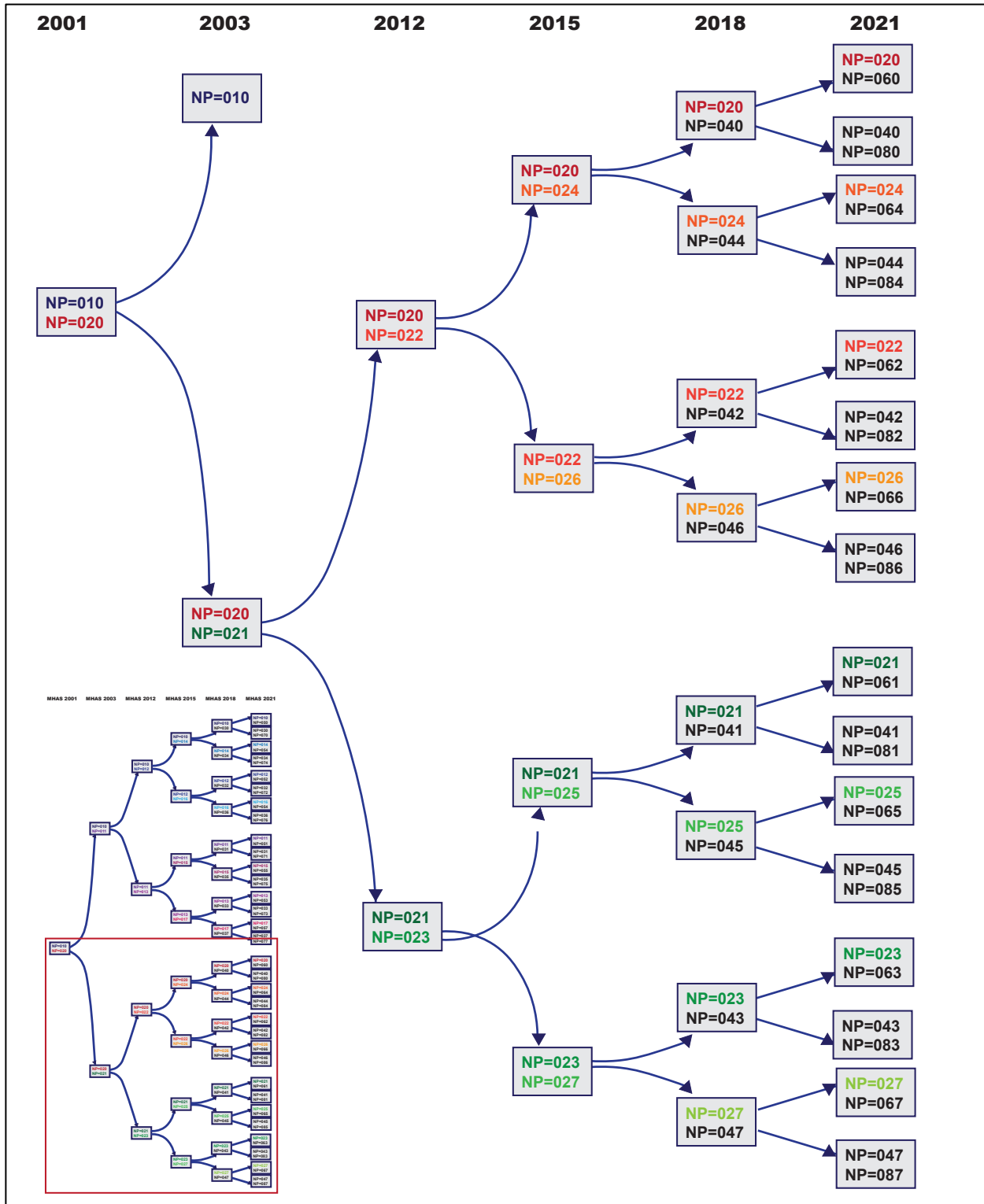


Figure 2. Assignment of NP values within the 020 family



From then on, if these subjects get new spouses, the spouses will be assigned codes that correspond to the person they are married to. If they are a spouse of the baseline subject $NP=010$, the new spouse is assigned 011 (for the first new spouse if there is one), 012 (for the second new spouse if there is one), etc. If they are spouses of selected person $NP=020$; the new spouse is assigned 021 (for the first new spouse if there is one), 022 (for the second new spouse if there is one), etc. The NP key assignment would be done in a fixed way. Figure 1 illustrates how NP is assigned in the “family” of the selected subject $NP=010$, while Figure 2 does it for the “family” of the selected subject $NP=020$.

In combination with the unique household ID ($CUNICAH$), this code uniquely identifies individual subjects of study.

Unique Person ID (Variable name: $UNHHIDNP$)

Starting in 2018, we added a new string variable that combines the unique household ID ($CUNICAH$) and the person ID (NP).

3. Sub-household ID (Variable name: $ACTHOG$ and $SUBHOG_XX$)

Updated Household ID (Variable name: $ACTHOG$)

In 2003, a household code was created to capture changes in the situation of the individual or couple interviewed in 2001. This is referred to as “updated household” ($ACTHOG$) and the codes reflect the type of change experienced, including divorce/separation, death, or new spouse. In the case of split couples as mentioned above, an interview was sought with both baseline respondents in their respective households and their new spouses if applicable. The updated-household codes also capture whether the household observed in 2003 contains the baseline *sampled* respondent, or the baseline *spouse* of the selected person.

Sub-household ID (Variable name: $SUBHOG_XX$)

A sub-household ID ($SUBHOG_XX$) was created to follow the modifications of the original household and new households that result from the changes occurred over time. These changes include: divorce/separation, death, or new spouse of the original subjects. The last two digits of the variable indicate the year of the respective survey. Thus, we constructed one variable for

each wave (SUBHOG01, SUBHOG03, SUBHOG12, SUBHOG15, and SUBHOG18), each of them reflecting the changes in the household recorded each wave. This new identifier was created to replace the “updated household” ID (ACTHOG) used in 2003.

Table 1. Description of SUBHOG

BASELINE HOUSEHOLD	
00	For all households that are new sample (in their baseline year)
NO CHANGES IN THE HOUSEHOLD	
01	The household contains an individual NP = 010 (first selected subject that entered the sample)
02	The household contains an individual NP=020 (subject in this household that entered the sample first)
03	The household contains an individual NP=011 (subject in this household that entered the sample first)
04	The household contains an individual NP=021 (subject in this household that entered the sample first)
<i>Note: Starting from 05 up to 10 we could assign new IDs in future waves.</i>	
CHANGES IN THE HOUSEHOLD	
<i>Note: The change can indicate a dissolution (due to the death of the spouse or separation – with or without a new union) or a new union. In case of a death in the household, this ID is assigned to the survivor.</i>	
11	The new household contains an individual NP=010 (subject in this new household that entered the sample first)
12	The new household contains an individual NP=020 (subject in this new household that entered the sample first)
13	The new household contains an individual NP=011 (subject in this new household that entered the sample first)
14	The new household contains an individual NP=021 (subject in this new household that entered the sample first)
15	The new household contains an individual NP=012 (subject in this new household that entered the sample first)
16	The new household contains an individual NP=022 (subject in this new household that entered the sample first)
17	The new household contains an individual NP=013 (subject in this new household that entered the sample first)
18	The new household contains an individual NP=023 (subject in this new household that entered the sample first)
19	The new household contains an individual NP=014 (subject in this new household that entered the sample first)
20	The new household contains an individual NP=024 (subject in this new household that entered the sample first)
<i>Note: Starting from 19 up to 30 we could assign new IDs in future waves.</i>	
DECEASED	
<i>Note: This ID is assigned to the deceased; to the survivor we assigned the ID from CHANGES IN THE HOUSEHOLD.</i>	
31	The deceased is NP=010
32	The deceased is NP=020
33	The deceased is NP=011
34	The deceased is NP=021
35	The deceased is NP=012
36	The deceased is NP=022
37	The deceased is NP=013
38	The deceased is NP=023
37	The deceased is NP=014
38	The deceased is NP=024
<i>Note: Starting from 39 up to 69 we could assign new IDs in future waves.</i>	
RE-UNION	
70	Two individuals (NP=010 ando NP=020) part of the study, they were together, separated, and got back together

- 71 Two individuals, from the family of 10, they were together, separated, and got back together
- 72 Two individuals, from the family of 20, they were together, separated, and got back together

NEW INDIVIDUAL

- 99 To the new individuals we assign 99 in the previous waves to the one when they enter the study

DECEASED

- 88 To the deceased we assign 88 in the posterior waves to the one when he/she was reported dead

NOT CONTACTED

- 77 To the individuals not contacted (whole household)

Source: MHAS (2013), The Mexican Health and Aging Study: “MHAS 2012 Data Files Description, Version1, September 2013.” Retrieved from www.MHASweb.org on (February 2015).

At baseline, the variable takes the value 00 to reflect no change in the household. Thus SUBHOG_01=00 for all households because they are all original households, new to the study in this wave. In the same way, for the new refresher sample of households added in 2012 and 2018, we assigned the same value (SUBHOG_12=00 and SUBHOG_18=00) to indicate that these are new to the study in this wave. For more information on how the values of SUBHOG_XX are assigned see “MHAS 2012 Data Files Description” available for download [here](#).

B. Outcomes of the Interview, Sample and Study Status

The file includes the following variables to indicate the outcome of the interviews at different levels:

1. Eligible (Variable name: ELEGIBLE_XX)

Starting in 2003, ELEGIBLE_XX indicates whether the individual is eligible for a follow-up interview in any given wave. As general rule, individuals were considered eligible for a re-interview if they completed an interview in the previous wave and they reported age was within the required age to be part of the MHAS sample.

2. Follow-New person (Variable name: NEW_SAMPLE_XX):

Starting in 2003, NEW_SAMPLE_XX indicates whether the interview is for a new person or a follow-up subject. In 2003, the new sample includes either a new spouse from a follow-up subject or a sample subject in 2001 who did not complete the interview in Wave 1.

In 2012, the new sample includes either a new spouse of a follow-up subject, or a sample subject

in 2001 who did not complete the interview in Wave 1. In addition, it includes the new sample of those born between 1952-1962.

Similar to 2003, in 2015 the new sample includes either a new spouse from a follow-up subject or a subject from the new 2012 sample who did not complete the interview in Wave 3.

In 2018, the new sample includes either a new spouse of a follow-up subject or the new sample added of those born between 1963-1968.

3. Type of non-interview (Variable name: TIPNE_XX)

Indicates whether a respondent was interviewed as well as the reason of non-interview. This is a key variable for users as it indicates the respondent's status in any given wave.

In 2001 and 2003, it is considered a household variable because if at least one member in the household completed the interview it is considered an “Interviewed” household. In 2001 and 2003 (TIPNE_01 and TIPNE_03) include the following options:

VALUE	LABEL	DESCRIPTION
0	Interviewed	At least one interview was completed in the household.
2	Nobody at time of interview	If nobody was present at the time of the visits (minimum of 5 visits).
3	Temporarily absent	A temporarily absent subject, is a person that at the time of the visit is not in the house, but intends to return during the collection period or before three months, regardless of the reason of the absence.
4	Refused to provide information	No interviews were completed in the household because the subject and the spouse (if any) refused to provide information and didn't allow others (proxy) to provide it either.
5	Other (occupied house)	If the interview was not completed for other reasons different from options 2 to 4.
6	Unoccupied, fit for living	If the house, with or without furniture, is unoccupied, but fit for living, for sale or rent.
7	Temporary housing	No permanent residents were found in the dwelling.
8	Unoccupied, inadequate to live in due to bad conditions or remodeling	If the house is unoccupied and not considered appropriate for living, that is, there is unacceptable structural conditions due either to its damage or other reasons like lack of windows and doors.
9	Unoccupied, temporarily used for other means different than housing	If the house is temporarily occupied by businesses, workshops, factories, etc.
10	Demolished	If the house has been destroyed or is in demolition process.

11	Tent, trailer, mobile home, etc.	If the tent, trailer, mobile home (etc.) has changed location by the time of the interview.
12	Business or permanent warehouse	If the house has been transformed or adapted from their normal living use to a commercial one, for business, or to store harvests, machinery, office supplies and other uses.
13	Other (specify)	If the house can't be classified in any of the previous options.
14	Selected (and spouse) moved and unable to reach or locate	If the selected subject was not located (or lost to follow-up).
15	Selected (and spouse) not fit for interview and no substitute informant	If the selected subject and the spouse (if any) could not complete the interview, for different reasons, and no substitute informant was available to complete the interview.
16	Selected deceased and no spouse	The selected subject was deceased without a spouse.
17	Selected deceased and no spouse within required age	The selected subject was deceased, and the spouse was not within the required age (50 years or older).
18	Selected (and spouse) temporarily absent and there is no adequate or available substitute informant	A household is considered "temporarily absent" if the selected subject and the spouse (if any) were never present at the time of the visits but they were identified living in the household. In this case, not substitute informant was available to complete the interview.
19	Other (specify)	No interviews were completed in the household for other reason not included in the previous options.

In 2012, TIPNE_12 includes the following options:

VALUE	DESCRIPTION
1	Interviewed
2	Incomplete Interview
3	Postponed Interview
4	Absent and there is no informant
5	Empty household or informant not adequate
6	Subject unable to complete interview without informant
7	Deceased without informant
8	Refused
9	Refused direct interview
10	Refused proxy interview
11	Refused next-of-kin interview
12	Changed location
13	Subject could not be located
14	Empty housing
15	Housing with temporary use
16	Housing with non-residential use, demolished or uninhabitable
17	Address not located
18	Area not safe
19	Other
20	Deceased subject from the new sample added in 2012

In 2015, TIPNE_15 includes the following options:

VALUE	DESCRIPTION
1	Complete Interview
2	Incomplete Interview
3	Postponed Interview
4	Absence of the subject or the proxy or next-of-kin informant
5	Absence of occupants or adequate informant
6	Subject not fit for interview and without proxy informant
7	Subject deceased without next-of-kin informant
8	Refusal
9	Refusal from the subject or the proxy or next-of-kin informant
10	Change of residence
11	Subject could not be located
12	Empty residence
13	Dwelling for temporary use
14	Dwelling with non-residential use, demolished or uninhabitable
15	Address not located
16	Area not safe
17	Other

In 2018, TIPNE_18 includes the following options:

VALUE	DESCRIPTION
1	Complete Interview
2	Incomplete Interview
3	Postponed Interview
4	Absence of the subject, or proxy or next-of-kin informant
5	Absence of occupants or adequate informant
6	Refusal
7	Subject not fit for interview and w/o proxy informant
8	Subject deceased without next-of-kin informant
9	Age out of the range for new sample
10	Change of residence
11	Subject could not be located
12	Residence could not be located
13	Other situation
14	Inhabited residence
15	Dwelling for temporary use
14	Dwelling with non-residential use, demolished or uninhabitable
16	Area not safe

4. Type of interview (Variable name: TIPENT_XX)

Indicates the type of individual interview. The MHAS makes every effort to obtain core interviews directly with respondents themselves. In cases where the respondent is unavailable or unable to participate, a proxy interview is sought with a knowledgeable informant. Starting in 2003, if the respondent was deceased an exit interview is sought with a next-of-kin. TIPENT_XX

indicates direct interviews or proxy interviews.

In 2001 and 2003, it also indicates the order of the interview in the household – first or second interview. The options included the following:

VALUE	DESCRIPTION	
	2001	2003
11	Direct, first interview	
12	Direct, second interview	
21	Proxy, first interview	
22	Proxy, second interview	
31		Next-of-kin interview, first interview
32		Next-of-kin interview, second interview
33		Next-of-kin interview, third interview
42		Non-response, second interview

Starting in 2012, TIPENT_XX uses the following options:

VALUE	DESCRIPTION
1	Direct, follow-up interview
2	Direct, new sample interview
3	Proxy, follow-up interview
4	Proxy, new sample interview
5	Next-of-kin

5. Reason for Proxy (Variable name: REASON_PROXY_XX)

In cases where the respondent is unable to complete a direct interview due to (1) health reason, (2) language reason, or (3) temporary absence a proxy interview is sought with a knowledgeable informant. The variable REASON_PROXY_XX indicates the reason a proxy interview was completed.

6. Interview Date (Variable name: INT_DATE_XX)

Indicates the date of the interview.

7. Wave Status – Alive or Deceased (Variable name: FALLECIDO_XX)

For all the eligible respondents in a given wave, the variable FALLECIDO_XX indicates whether a next-of-kin interview for a deceased subject was completed.

C. Other

1. Same dwelling (Variable name: MISMA_V_XX)

Indicates whether the household was located in the same dwelling or in a different dwelling or a group quarter (asylum, penitentiary, etc.) as in the previous wave.

2. Interview Technique – Electronic versus Paper (Variable name: C_PAP_XX)

Starting in 2012, the interviews were completed using a CAPI (Computer Assisted Personal Interview). However, in some cases due to security or technological issues the interview was completed using paper and pencil. C_PAP_XX indicates whether the interview was completed using the electronic questionnaire or paper and pencil.

3. High Migration States (Variable name: EAM_XX)

Starting in 2012, it indicates whether the subject is part of a high migration state that was oversampled.

4. Age and Sex (Variable name: AGE_XX and SEX_XX)

In each wave, the variable AGE_XX is calculated for the new sample using the self-reported birth date and the interview date. For the follow-up sample, the variable AGE_XX is the self-reported age unless the subject refuses to answer and doesn't know; in those cases the age is calculated similar to how it's done for the new sample.

The variable SEX_XX is the self-report given in each wave. In 2015, we observed differences in the reported sex across waves using the variables SEXO_01, SEXO_03, SEX_12, and SEX_15. These differences were carefully studied by the MHAS team and INEGI in Mexico and a new variable SEX_VALIDATED_15 was constructed with the validated sex variable. The user can use this variable instead, from 2001 to 2015.

D. Weights

The file also contains the sampling weights that were constructed for every wave of the study. MHAS has individual (FACTORI_XX), household (FACTORH_XX), anthropometric and

biomarkers (FACTORA_XX) factors. The individual weights are based on the birth cohort, household composition, and the place of residence (in urban and rural areas) and geographic areas. Also, the household weights are based on the sample design and sample selection criteria, and the calibration variables of all community dwelling households with at least one resident 50 years or older, based on the household composition, and the place of residence (urban and rural areas) and geographic areas. The sample selection involves 3 stages: selection of Primary Sampling Units (PSU), then the selection of households within each PSU, and finally the selection of subjects in those households. The sampling weights provided reflect this sample selection as well as post-stratification adjustment for non-response. They are used to expand the sample to the national population. For more information regarding the calculation of the weights please see the 2012 or the 2015 Methodological Document available in our website: http://mhasweb.org/Resources/DOCUMENTS/2012/Methodological_Document_2012%E2%80%93SEC.pdf and http://mhasweb.org/Resources/DOCUMENTS/2015/Methodological_Document_2015.pdf

E. Sub-Samples and Ancillary Study

1. Selected for Anthropometric Measures 2001 & 2003 (Variable name: ANTRO_XX)

In 2001 and 2003, ANTRO_XX indicates whether the subject was selected to be part of the sub-sample for anthropometric measures.

2. Result for Anthropometric Measures 2001 & 2003(Variable name: RES_ANTRO_XX)

In 2001 and 2003, RES_ANTRO_XX indicates whether the subject selected to be part of the sub-sample for anthropometrics measures participated in the collection of the measurements.

3. Sub-sample 2012 (Variable name: SUBSAMPLE_12)

SUBSAMPLE_12 indicates whether the subject was selected to be part of the 2012 sub-sample for anthropometric, biomarkers, and performance measures.

4. Result for Biomarkers Measures 2012 (Variable name: RES_BIOMARKERS_12)

RES_BIOMARKERS_12 indicates whether the subject was selected to be part of the 2012 sub-sample for anthropometric, biomarkers, and performance measures participated in the collection

of the measurements.

5. Ancillary Study Variables: Cognitive Aging Study (Mex-Cog)

Mex-Cog 2016 Sub-sample (Variable name: SUBSAMPLE_16)

In 2016, SUBSAMPLE_16 indicates whether the subject was selected to be part the Cognitive Aging Study (Mex-Cog) linked to the MHAS. The Mex-Cog 2016 sample was selected using the MHAS wave 4 (2015) as sampling frame. The criteria for eligibility for Mex-Cog were: first, aged 55 and older in MHAS 2015; and second, having completed a direct interview or a proxy interview for health reasons in the MHAS 2015. Only 8 of the 32 states were selected using stratified sampling procedures. The Mex-Cog contains three parts: Cognitive Assessment, Adequate Informant Survey, and Anthropometric and Performance Measures

Mex-Cog 2016 Phase (Variable name: PHASE_MXCOG_16)

PHASE_MXCOG_16 indicates whether the information was collected during Phase 1 in the spring (March-April) or Phase 2 in the fall (October-November) of 2016.

Mex-Cog 2016 Result for Each Component (Variables name: RES_MXCOG_16,

RES_COGNITIVO_16, RES_INFORMANTE_16, RES_ANTRO_16)

	Frequency	COMPONENTS			
		COG	INF	ANTRO	BIO
0000	983				
0010	2			✓	
0100	200		✓		
0101	2		✓		✓
0110	11		✓	✓	
0111	10		✓	✓	✓
1000	4	✓			
1010	127	✓		✓	
1011	62	✓		✓	✓
1100	9	✓	✓		
1101	1	✓	✓		✓
1110	1,162	✓	✓	✓	
1111	677	✓	✓	✓	✓
Total	3,250				

RES_MXCOG_16 is a string variable that indicates whether or not a component of the study was completed; from left to right each digit indicates each of following components respectively:

cognitive assessment (COG), adequate informant survey (INF), anthropometric and performance measures (ANTRO), and biomarkers (BIO). The table above illustrates the different combinations.

RES_COGNITIVO_16, RES_INFORMANTE_16, and RES_ANTRO_16 indicate whether the subject selected to be part of Mex-Cog 2016 completed the cognitive assessment, adequate informant survey, and/or anthropometric and performance measures portion, respectively. The variables also indicated the reason for not completing each component including refusal, loss to follow-up, deceased, or other.

Mex-Cog 2016 Other Variables (Variables name: RES_BIOMARKERS_16, EDTA_16, HBA1C_16)

RES_BIOMARKERS_16 indicates whether the subject completed the biomarkers portion of Mex-Cog 2016. While HBA1C_16 and EDTA_16 indicate if Glycosylated Hemoglobin was measured and if a Lavender Tube with EDTA was collected, respectively.

C. Accessing the MHAS Data

All study databases and documentation can be accessed free of charge from the MHAS study website after registration is completed. The platform is in English www.MHASweb.org and in Spanish www.ENASEM.org. The website also features a dynamic searchable database of publications using MHAS data and a discussion forum. **We ask all users to please inform us to info@mhasweb.com of any published work using the MHAS data.**

We also ask our users to include the following acknowledgement: “The MHAS (Mexican Health and Aging Study) is sponsored by the National Institutes of Health/National Institute on Aging (grant number NIH R01AG018016) and the Mexican National Institute of Statistics and Geography (*Instituto Nacional de Estadística y Geografía*, INEGI).

D. Merging the MHAS Data

Correctly merging data sets is critical to every research project. We always recommend starting every merge with the Master Follow-up File, in particular when merging data files across waves. As described above, the Master Follow-up File contains a record of every subject that has participated in the study and includes all the identifiers used in each wave and the result of the interview in each wave. In this process it is essential you identify the right variables to uniquely identify household and individual. Below we provide a short description on which identifiers to use in each wave.

How to use identifiers to merge data files

To merge data files at the household level you need to use the following variables:

- 1) UNHHID (also named CUNICAH) in 2001
- 2) CUNICAH+ACTHOG in 2003
- 3) CUNICAH+SUBHOG_XX in each wave from 2012 to 2018

Check that the identifiers use the same names both in the 'master' and the 'using dataset'. For example, the unique household ID CUNICAH is also named UNHHID in some data files.

To merge data files at the individual level you need to use the following variables:

- 1) UNHHID (also named CUNICAH)+CODENT01 (also named PS3) in 2001
- 2) CUNICAH+ACTHOG+CODENT03 (also named ENT2) in 2003
- 3) CUNICAH+NP in each wave from 2012 to 2018

Check that the identifiers use the same names both in the 'master' and the 'using dataset'. For example, the 2001 person ID CODENT01 is also named PS3 and the 2003 person ID CODENT03 is also named ENT2 in some data files.

Appendix 1. Master Follow-up File Content

Number of Observations		28,303	Version 2. July 2020 (includes weights)
Number of Variables		106	
Variable Name	Type	Variable Label	
unhhid	float	Unique Household identification code (=cunicah)	
UNHHID	str5	Unique Household ID (=CUNICAH) - STRING	
cunicah	float	Clave Unica del Hogar (=unhhid)	
CUNICAH	str5	Clave Unica del Hogar (=UNHHID) - STRING	
np	int	Person Number/ Numero de Persona	
NP	str3	Person Number/Numero de Persona - STRING	
UNHHIDNP	str8	Unique Person ID/Numero de Identificador de Persona Unico - STRING	
codent01	float	Person identification code 2001 (=ps3)	
subhog_01	int	2001 sub-household identifier	
acthog	byte	Update household code 2003	
codent03	byte	Person identification code 2003 (=ent2)	
subhog_03	long	2003 sub-household identifier	
subhog_12	byte	2012 sub-household identifier	
subhog_15	byte	2015 sub-household identifier	
subhog_18	byte	2018 sub-household identifier	
res_01	int	Outcome of the household interview 2001	
tipne_01	byte	Type of household non-interview 2001	
tipent_01	byte	Type of interview 2001	
reason_proxy_01	float	Reason for Proxy Interview 2001	
tamloc_01	byte	Locality size 2001	
factorh_01	long	Household weight 2001	
factori_01	long	Individual weight 2001	
factora_01	long	Biomarkers weight 2001	
edad_01	int	Age 2001	
sexo_01	float	Sex 2001 (Male=1)	
antro_01	byte	Selected for anthropometric measurement 2001	
res_antro_01	byte	Result for Anthropometric Measures 2003	
elegible_03	byte	Eligible for interview in 2003	
new_sample_03	byte	Follow-up/new person	
misma_v_03	byte	Same dwelling in 2003	
fallecido_03	byte	Died between 2001 and 2003	
res_03	int	Outcome for the household interview 2003	
res_ent_03	byte	Outcome for individual interview 2003	
tipne_03	byte	Type of non-interview 2003	
tipent_03	byte	Type of interview 2003	

reason_proxy_03	byte	Reason for Proxy Interview 2003
factori_03	long	Individual weight 2003
factorh_03	long	Household weight 2003
factora_03	long	Biomarkers weight 2003
antro_03	byte	Selected for anthropometric measurement 2003
res_antro_03	byte	Result for Anthropometric Measures 2003
edad_03	int	Age 2003
sexo_03	byte	Sex 2003 (Male=1)
mes_03	byte	Month of birth 2003
a_o_03	int	Year of birth 2003
elegible_12	float	Eligible for interview in 2012
new_sample_12	byte	Follow-up and new sample 2012
misma_v_12	byte	Same dwelling in 2012
tipne_12	byte	Type of non-interview 2012
tipent_12	float	Type of interview 2012
reason_proxy_12	byte	Reason for Proxy Interview 2012
int_date_12	str10	Interview date 2012
order_12	byte	Order of interview 2012 (in the household)
fallecido_12	float	Died between 2003 and 2012
tam_loc_12	float	Locality size 2012
eam_12	byte	High migration states 2012
factorh_12	long	Household weight 2012
factori_12	long	Individual weight 2012
factora_12	long	Biomarkers/Anthropometrics weight 2012
age_12	int	Age 2012
sex_12	byte	Sex 2012 (Male=1)
subsample_12	float	Selected subsample for Biomarkers/Anthropometrics 2012
res_biomarke~12	byte	Result for Biomarkers/Anthropometrics 2012
biomarkers_12	float	Sample with Biomarkers 2012
edta_12	float	2012 Completed EDTA-Lavender Tube
hbalc_12	float	2012 Completed HbA1c
elegible_15	float	Eligible for interview in 2015
new_sample_15	float	Follow-up and new sample/spouses 2015
misma_v_15	byte	Same dwelling as in 2012
tipne_15	byte	Type of non-interview 2015
tipent_15	float	Type of interview 2015
reason_proxy_15	byte	Reason for Proxy Interview 2015
int_date_15	str10	Interview date 2015
c_pap_15	byte	Interview technique - CAPI vs Paper
fallecido_15	float	Died between 2012 and 2015

tam_loc_15	byte	Locality size 2015
eam_15	float	Current Residence in High Migration States from 2012
factorh_15	long	Household weight 2015
factori_15	long	Individual weight 2015
age_15	float	Age 2015
sex_15	byte	Sex (Male=1)
sex_validate~15	float	Validated Sex 2015 (Male=1)
subsample_16	float	Selected subsample for Mex-Cog 2016
phase_mxcog_16	float	Mex-Cog 2016 Phase
res_mxcog_co~16	str4	Mex-Cog Result for Each Component (Character): COG/INF/ANTRO/BIOM
res_mxcog_16	float	Mex-Cog Result All Four Components
res_cognitiv~16	byte	Mex-Cog Result of Cognitive Assessment
res_informan~16	byte	Mex-Cog Result of Informant Interview
res_antro_16	byte	Mex-Cog Result for Anthropometrics 2016
biomarkers_16	float	Mex-Cog 2016 Sample with Biomarkers
edta_16	float	Mex-Cog 2016 Sample with EDTA-Lavender Tube
hbalc_16	float	Mex-Cog 2016 Sample with HbA1c
elegible_18	float	Eligible for interview in 2018
new_sample_18	float	Follow-up and new sample/spouses 2018
tipne_18	byte	Type of non-interview 2018
misma_v_18	byte	Same dwelling as in 2018
resul_hh_18	float	Result of Sections at Household Level 2018
tipent_18	float	Type of interview 2018
reason_proxy_18	byte	Reason for Proxy Interview 2018
int_date_18	str8	Interview date 2018
c_pap_18	byte	Interview technique - Electronic vs Paper
fallecido_18	float	Died between 2015 and 2018
tam_loc_18	byte	Locality size 2018
eam_18	byte	Current Residence in High Migration States from 2018
age_18	float	Age 2018
sex_18	float	Sex (Male=1)

Appendix 2. Master Follow-up File Codebook

 unhhid Unique Household identification code (=cunicah)

type: numeric (float)

range: [1,20218] units: 1
 unique values: 17,664 missing .: 0/28,303

Variable	Obs	Mean	Std. Dev.	Min	Max
unhhid	28,303	9794.42	5666.713	1	20218

 UNHHID Unique Household ID
 (=CUNICAH) - STRING

type: string (str5)

unique values: 17,664 missing "": 0/28,303

examples: "03926"
 "07891"
 "11727"
 "15182"

 cunicah Clave Unica del Hogar (=unhhid)

type: numeric (float)

range: [1,20218] units: 1
 unique values: 17,664 missing .: 0/28,303

Variable	Obs	Mean	Std. Dev.	Min	Max
cunicah	28,303	9794.42	5666.713	1	20218

 CUNICAH Clave Unica del Hogar (=UNHHID) - STRING

type: string (str5)

unique values: 17,664 missing "": 0/28,303

examples: "03926"
 "07891"
 "11727"
 "15182"

 np Person Number/ Numero de Persona

type: numeric (int)

range: [10,44] units: 1
 unique values: 13 missing .: 0/28,303

Person |
 Number/ |

Numero de Persona	Freq.	Percent	Cum.
10	17,621	62.26	62.26
11	236	0.83	63.09
12	4	0.01	63.11
13	5	0.02	63.12
14	110	0.39	63.51
15	2	0.01	63.52
20	10,234	36.16	99.68
21	56	0.20	99.88
22	1	0.00	99.88
24	30	0.11	99.99
34	2	0.01	99.99
40	1	0.00	100.00
44	1	0.00	100.00
Total	28,303	100.00	

NP Person Number/Numero de Persona - STRING

```

type: string (str3)
unique values: 13 missing "": 0/28,303
examples: "010"
           "010"
           "010"
           "020"

```

UNHHIDNP Unique Person ID/Numero de Identificador de Persona Unico - STRING

```

type: string (str8)
unique values: 28,303 missing "": 0/28,303
examples: "03926010"
           "07891020"
           "11727010"
           "15182010"

```

codent01 Person identification code 2001 (=ps3)

```

type: numeric (float)
label: codent01
range: [1,2] units: 1
unique values: 2 missing .: 12,901/28,303

```

Person identification code 2001 (=ps3)	Freq.	Percent	Cum.
1.Selected person	9,812	63.71	63.71
2.Spouse of selected person	5,590	36.29	100.00
Total	15,402	100.00	

subhog_01 2001 sub-household identifier

type: numeric (int)
label: clavesubhog

range: [0,99] units: 1
unique values: 3 missing .: 0/28,303

2001 sub-household identifier	Freq.	Percent	Cum.
00.Baseline HH	15,402	54.42	54.42
77.HH lost to follow-up/Not contacted	446	1.58	55.99
99.Not yet part of the study	12,455	44.01	100.00
Total	28,303	100.00	

acthog Update household code 2003

type: numeric (byte)
label: acthog

range: [0,32] units: 1
unique values: 9 missing .: 13,027/28,303

Update household code 2003	Freq.	Percent	Cum.
00.Neither separation nor death, no new	14,100	92.30	92.30
01.Neither separation nor death, new sp	198	1.30	93.60
10.Had a separation and contains sample	63	0.41	94.01
11.Had a separation and contains sample	8	0.05	94.06
20.Had a separation and contains spouse	61	0.40	94.46
21.Had a separation and contains spouse	12	0.08	94.54
30.One 2001 respondent died, no new spo	790	5.17	99.71
31.One 2001 respondent died, new spouse	16	0.10	99.82
32.Both 2001 respondents died	28	0.18	100.00
Total	15,276	100.00	

codent03 Person identification code 2003 (=ent2)

type: numeric (byte)
label: codent03

range: [1,4] units: 1
unique values: 4 missing .: 13,028/28,303

Person identification code 2003 (=ent2)	Freq.	Percent	Cum.
1.Selected person in 2001	9,675	63.34	63.34
2.Spouse of selected person in 2001	5,498	35.99	99.33
3.New spouse of selected person in 2001	95	0.62	99.95
4.New spouse-of-spouse of selected pers	7	0.05	100.00
Total	15,275	100.00	

subhog_03 2003 sub-household identifier

type: numeric (long)
label: clavesubhog

range: [0,99] units: 1
unique values: 11 missing .: 0/28,303

2003 sub-household identifier	Freq.	Percent	Cum.
00.Baseline HH	804	2.84	2.84
01.No change, HH contains NP=010	13,272	46.89	49.73
02.No change, HH contains NP=020	1	0.00	49.74
11.New HH contains NP=010	247	0.87	50.61
12.New HH contains NP=020	59	0.21	50.82
13.New HH contains NP=011	6	0.02	50.84
14.New HH contains NP=021	1	0.00	50.84
31.NP=010 Deceased	534	1.89	52.73
32.NP=020 Deceased	278	0.98	53.71
77.HH lost to follow-up/Not contacted	1,163	4.11	57.82
99.Not yet part of the study	11,938	42.18	100.00
Total	28,303	100.00	

subhog_12

2012 sub-household identifier

2012 sub-household identifier	Freq.	Percent	Cum.
00.Baseline HH	5,754	20.33	20.33
01.No change, HH contains NP=010	8,222	29.05	49.38
02.No change, HH contains NP=020	117	0.41	49.79
11.New HH contains NP=010	832	2.94	52.73
12.New HH contains NP=020	766	2.71	55.44
13.New HH contains NP=011	20	0.07	55.51
14.New HH contains NP=021	3	0.01	55.52
31.NP=010 Deceased	1,938	6.85	62.37
32.NP=020 Deceased	792	2.80	65.17
33.NP=011 Deceased	12	0.04	65.21
70.NP=010 & NP=020 separated, reunited	4	0.01	65.22
71.Subsequent NP=010 HH separated, reun	2	0.01	65.23
77.HH lost to follow-up/Not contacted	3,253	11.49	76.72
88.Deceased before current wave	546	1.93	78.65
99.Not yet part of the study	6,042	21.35	100.00
Total	28,303	100.00	

subhog_15

2015 sub-household identifier

type: numeric (byte)
label: clavesubhog

range: [0,99] units: 1
unique values: 15 missing .: 0/28,303

2015 sub-household identifier	Freq.	Percent	Cum.
00.Baseline HH	644	2.28	2.28
01.No change, HH contains NP=010	12,347	43.62	45.90
02.No change, HH contains NP=020	91	0.32	46.22
11.New HH contains NP=010	1,308	4.62	50.84
12.New HH contains NP=020	1,097	3.88	54.72
13.New HH contains NP=011	20	0.07	54.79
14.New HH contains NP=021	7	0.02	54.81
15.New HH contains NP=012	29	0.10	54.92
31.NP=010 Deceased	794	2.81	57.72
32.NP=020 Deceased	403	1.42	59.15
33.NP=011 Deceased	11	0.04	59.18
34.NP=021 Deceased	2	0.01	59.19
70.NP=010 & NP=020 separated, reunited	44	0.16	59.35
77.HH lost to follow-up/Not contacted	5,713	20.19	79.53

99.Not yet part of the study	5,793	20.47	100.00

Total	28,303	100.00	

subhog_18 2018 sub-household identifier

```

type: numeric (byte)
range: [0,77]
unique values: 18
units: 1
missing .: 6,764/28,303
mean: 6.82599
std. dev: 16.294

```

2018 sub-household identifier	Freq.	Percent	Cum.
00.Baseline HH	5,766	26.77	26.77
01.No change, HH contains NP=010	10,913	50.67	77.44
02.No change, HH contains NP=020	74	0.34	77.78
11.New HH contains NP=010	1,397	6.49	84.27
12.New HH contains NP=020	1,268	5.89	90.15
13.New HH contains NP=011	32	0.15	90.30
14.New HH contains NP=021	6	0.03	90.33
15.New HH contains NP=012	19	0.09	90.42
19. New HH contains NP=014	14	0.06	90.48
20.New HH contains NP=024	4	0.02	90.50
31.NP=010 Deceased	784	3.64	94.14
32.NP=020 Deceased	360	1.67	95.81
33.NP=011 Deceased	7	0.03	95.84
34.NP=021 Deceased	2	0.01	95.85
39.NP=014 Deceased	4	0.02	95.87
40.NP=024 Deceased	3	0.01	95.89
70.NP=010 & NP=020 separated, reunited	53	0.25	96.13
77.HH lost to follow-up/Not contacted	833	3.87	100.00

Total	21,539	100.00	

res_01 Outcome of the household interview 2001

```

type: numeric (int)
label: res01
range: [110,233]
unique values: 12
units: 1
missing .: 12,901/28,303

```

Outcome of the household interview 2001	Freq.	Percent	Cum.
110.Selected person, information provid	3,904	25.35	25.35
130.Selected person, information provid	362	2.35	27.70
202.Couple, Information on selected per	232	1.51	29.20
203.Couple, Information on selected per	4	0.03	29.23
210.Couple, Information on selected per	246	1.60	30.83
211.Couple, Information about both prov	422	2.74	33.57
212.Couple, Information provided by eac	9,460	61.42	94.99
213.Couple, Information on selected per	78	0.51	95.49
222.Couple, Information on both by spou	470	3.05	98.55
230.Couple, Information on selected per	6	0.04	98.58
232.Couple, Information on selected per	86	0.56	99.14
233.Couple, Information on both provide	132	0.86	100.00

Total	15,402	100.00	

tipne_01

Type of household non-interview 2001

type: numeric (byte)
label: tipne01

range: [0,0] units: 1
unique values: 1 missing .: 12,901/28,303

Type of household non-interview 2001	Freq.	Percent	Cum.
0.Interviewed	15,402	100.00	100.00
Total	15,402	100.00	

tipent_01

Type of interview 2001

type: numeric (byte)
label: tipent01

range: [11,22] units: 1
unique values: 4 missing .: 12,901/28,303

Type of interview 2001	Freq.	Percent	Cum.
11.Direct, first interview	9,424	61.19	61.19
12.Direct, second interview	4,946	32.11	93.30
21.Proxy, first interview	438	2.84	96.14
22.Proxy, second interview	594	3.86	100.00
Total	15,402	100.00	

reason_proxy_01

Reason for Proxy Interview 2001

type: numeric (float)
label: reason

range: [1,3] units: 1
unique values: 3 missing .: 27,271/28,303

Reason for Proxy Interview 2001	Freq.	Percent	Cum.
1.Health reasons	321	31.10	31.10
2.Language reasons	21	2.03	33.14
3.Temporary absence	690	66.86	100.00
Total	1,032	100.00	

tamloc_01

Locality size 2001

type: numeric (byte)
label: tamloc

range: [1,4] units: 1
unique values: 4 missing .: 12,776/28,303

Locality size 2001	Freq.	Percent	Cum.
1.100,000 - +	9,148	58.92	58.92
2. 15,000 - 99,999	2,325	14.97	73.89

3.	2,500 - 14,999		1,414	9.11	83.00
	4.< 2,500		2,640	17.00	100.00
-----+					
	Total		15,527	100.00	

factorh_01 Household weight 2001

```

type: numeric (long)
range: [16,35360]           units: 1
unique values: 1,840       missing .: 12,901/28,303
mean: 1014.07
std. dev: 1578.02

```

Variable	Obs	Mean	Std. Dev.	Min	Max
factorh_01	15,402	1014.068	1578.016	16	35360

factori_01 Individual weight 2001

```

type: numeric (long)
range: [0,44177]           units: 1
unique values: 2,125       missing .: 12,901/28,303
mean: 948.301
std. dev: 1686.75

```

Variable	Obs	Mean	Std. Dev.	Min	Max
factori_01	15,402	948.3007	1686.748	0	44177

factora_01 Biomarkers weight 2001

```

type: numeric (long)
range: [0,108420]          units: 1
unique values: 1,290       missing .: 12,901/28,303
mean: 948.301
std. dev: 4341.15

```

Variable	Obs	Mean	Std. Dev.	Min	Max
factora_01	15,402	948.3013	4341.146	0	108420

edad_01 Age 2001

```

type: numeric (int)
range: [18,105]           units: 1
unique values: 84         missing .: 13,117/28,303
mean: 60.2629
std. dev: 10.8379

```

Variable	Obs	Mean	Std. Dev.	Min	Max
----------	-----	------	-----------	-----	-----

```
-----+-----
edad_01 |      15,186      60.26294      10.83792      18      105
```

```
-----+-----
sexo_01                                     Sex 2001 (Male=1)
-----+-----
```

```
      type: numeric (float)
      label: sex
```

```
      range: [1,2]
unique values: 2
      units: 1
missing .: 12,901/28,303
```

Sex 2001 (Male=1)	Freq.	Percent	Cum.
1.Male	6,650	43.18	43.18
2.Female	8,752	56.82	100.00
Total	15,402	100.00	

```
-----+-----
antro_01                                     Selected for anthropometric measurement 2001
-----+-----
```

```
      type: numeric (byte)
      label: yesno
```

```
      range: [0,1]
unique values: 2
      units: 1
missing .: 12,901/28,303
```

Selected for anthropometric measurement 2001	Freq.	Percent	Cum.
0.No	12,458	80.89	80.89
1.Yes	2,944	19.11	100.00
Total	15,402	100.00	

```
-----+-----
res_antro_01                               Result for Anthropometric Measures 2003
-----+-----
```

```
      type: numeric (byte)
      label: yesno
```

```
      range: [0,1]
unique values: 2
      units: 1
missing .: 12,901/28,303
```

Result for Anthropometric Measures 2003	Freq.	Percent	Cum.
0.No	12,829	83.29	83.29
1.Yes	2,573	16.71	100.00
Total	15,402	100.00	

```
-----+-----
elegible_03                               Eligible for interview in 2003
-----+-----
```

```
      type: numeric (byte)
      label: elegible03
```

```
      range: [0,1]
      units: 1
```

unique values: 2 missing .: 12,901/28,303

Eligible for interview in 2003	Freq.	Percent	Cum.
0.Not eligible for 2003 interview	252	1.64	1.64
1.Eligible for 2003 interview	15,150	98.36	100.00
Total	15,402	100.00	

new_sample_03 Follow-up/new person

type: numeric (byte)
label: entrevista03

range: [1,2] units: 1
unique values: 2 missing .: 14,463/28,303

Follow-up/new person	Freq.	Percent	Cum.
1.Follow-up person	13,620	98.41	98.41
2.New person	220	1.59	100.00
Total	13,840	100.00	

misma_v_03 Same dwelling in 2003

type: numeric (byte)
label: misma_v03

range: [1,3] units: 1
unique values: 3 missing .: 13,028/28,303

Same dwelling in 2003	Freq.	Percent	Cum.
1.Same dwelling	14,475	94.76	94.76
2.Different dwelling	798	5.22	99.99
3.Group quarters (asylum, penitentiary)	2	0.01	100.00
Total	15,275	100.00	

fallecido_03 Died between 2001 and 2003

type: numeric (byte)
label: fallecido03

range: [0,1] units: 1
unique values: 2 missing .: 13,918/28,303

Died between 2001 and 2003	Freq.	Percent	Cum.
0.Alive in 2003	13,839	96.20	96.20
1.Died between 2001-2003	546	3.80	100.00
Total	14,385	100.00	

res_03 Outcome for the household interview 2003

type: numeric (int)
label: res03

range: [1100,3231] units: 1
unique values: 26 missing .: 13,028/28,303

Outcome for the household interview 2003	Freq.	Percent	Cum.
1100.One, direct 1st int.	3,428	22.44	22.44
1200.One, proxy 1st int.	331	2.17	24.61
1300.One, next of kin int.	257	1.68	26.29
1400.One, no 1st int.	279	1.83	28.12
2101.Two, direct 1st interview, direct	236	1.55	29.66
2102.Two, direct 1st int., proxy int. n	68	0.45	30.11
2104.Two, direct 1st int., no new perso	10	0.07	30.17
2110.Two, direct 1st int., direct 2nd i	7,880	51.59	81.76
2120.Two, direct 1st int., proxy 2nd in	1,156	7.57	89.33
2130.Two, direct 1st int., next of kin	456	2.99	92.31
2140.Two, direct 1st int., no 2nd int.	234	1.53	93.85
2201.Two, proxy 1st int., no 2nd int.,	16	0.10	93.95
2202.Two, proxy 1st int., proxy int. ne	12	0.08	94.03
2204.Two, proxy 1st int., no new person	2	0.01	94.04
2220.Two, proxy 1st int., proxy 2nd int	188	1.23	95.27
2230.Two, proxy 1st int., next of kin i	48	0.31	95.59
2240.Two, proxy 1st int., no 2nd int.	4	0.03	95.61
2301.Two, next of kin int., direct int.	2	0.01	95.63
2304.Two, next of kin int., no 2nd int.	2	0.01	95.64
2330.Two, next of kin int., next of kin	24	0.16	95.80
2340.Two, next of kin int., no 2nd int.	14	0.09	95.89
2401.Two, no 1st int., direct int. new	6	0.04	95.93
2404.Two, no 1st int., no 2nd int., no	6	0.04	95.97
2440.Two, no 1st int., no 2nd int.	604	3.95	99.92
3131.Three, direct 1st int.,next of kin	9	0.06	99.98
3231.Three, proxy 1st int., next of kin	3	0.02	100.00
Total	15,275	100.00	

res_ent_03 Outcome for individual interview 2003

type: numeric (byte)
label: res_ent03

range: [1,20] units: 1
unique values: 2 missing .: 14,053/28,303

Outcome for individual interview 2003	Freq.	Percent	Cum.
1.Complete interview	14,215	99.75	99.75
20.Incomplete interview	35	0.25	100.00
Total	14,250	100.00	

tipne_03 Type of non-interview 2003

type: numeric (byte)
label: tipne03

range: [0,29] units: 1
unique values: 9 missing .: 13,028/28,303

Type of non-interview 2003	Freq.	Percent	Cum.
0.Interviewed	14,386	94.18	94.18
21.Changed location and cannot be locat	317	2.08	96.26
22.Refused Core Interview	241	1.58	97.83
23.Refused Proxy Interview	5	0.03	97.87
24.Refused Next-of-Kin Interview	6	0.04	97.91
25.Unable without informant	24	0.16	98.06
26.Absent and there is no informant	213	1.39	99.46
27.Deceased without an informant	16	0.10	99.56
29.Other	67	0.44	100.00
Total	15,275	100.00	

tipent_03 Type of interview 2003

type: numeric (byte)
label: tipent03
range: [11,42] units: 1
unique values: 8 missing .: 13,917/28,303

Type of interview 2003	Freq.	Percent	Cum.
11.Direct, first interview	8,465	58.84	58.84
12.Direct, second interview	4,061	28.23	87.07
21.Proxy, first interview	457	3.18	90.25
22.Proxy, second interview	721	5.01	95.26
31.Next-of-kin interview, first intervi	277	1.93	97.18
32.Next-of-kin interview, second intervi	265	1.84	99.03
33.Next-of-kin interview, third intervi	4	0.03	99.05
42.Non-response, second interview	136	0.95	100.00
Total	14,386	100.00	

reason_proxy_03 Reason for Proxy Interview 2003

type: numeric (byte)
label: reason
range: [1,4] units: 1
unique values: 4 missing .: 27,125/28,303

Reason for Proxy Interview 2003	Freq.	Percent	Cum.
1.Health reasons	358	30.39	30.39
2.Language reasons	54	4.58	34.97
3.Temporary absence	763	64.77	99.75
4.Invalid code	3	0.25	100.00
Total	1,178	100.00	

factori_03 Individual weight 2003

type: numeric (long)
range: [0,46315] units: 1
unique values: 2,114 missing .: 12,776/28,303
mean: 907.109

std. dev: 1746.56

Variable	Obs	Mean	Std. Dev.	Min	Max
factori_03	15,527	907.1087	1746.563	0	46315

factorh_03 Household weight 2003

type: numeric (long)

range: [0,46315] units: 1
unique values: 2,063 missing .: 12,776/28,303
mean: 900.397
std. dev: 1731.9

Variable	Obs	Mean	Std. Dev.	Min	Max
factorh_03	15,527	900.3974	1731.898	0	46315

factora_03 Biomarkers weight 2003

type: numeric (long)

range: [0,167630] units: 1
unique values: 1,193 missing .: 12,776/28,303
mean: 900.398
std. dev: 4391.99

Variable	Obs	Mean	Std. Dev.	Min	Max
factora_03	15,527	900.3983	4391.992	0	167630

antro_03 Selected for anthropometric measurement 2003

type: numeric (byte)
label: yesno

range: [0,1] units: 1
unique values: 2 missing .: 13,028/28,303

Selected for			
anthropometric			
measurement 2003	Freq.	Percent	Cum.
0.No	12,370	80.98	80.98
1.Yes	2,905	19.02	100.00
Total	15,275	100.00	

res_antro_03 Result for Anthropometric Measures 2003

type: numeric (byte)
label: yesno

range: [0,1] units: 1
unique values: 2 missing .: 14,463/28,303

Result for Anthropometric Measures 2003	Freq.	Percent	Cum.
0.No	11,499	83.09	83.09
1.Yes	2,341	16.91	100.00
Total	13,840	100.00	

edad_03 Age 2003

type: numeric (int)
range: [21,999] units: 1
unique values: 85 missing .: 13,382/28,303
mean: 62.8848
std. dev: 27.5392

Variable	Obs	Mean	Std. Dev.	Min	Max
edad_03	14,910	62.19416	10.58009	21	107

sexo_03 Sex 2003 (Male=1)

type: numeric (byte)
label: sex
range: [1,2] units: 1
unique values: 2 missing .: 14,599/28,303

Sex 2003 (Male=1)	Freq.	Percent	Cum.
1.Male	5,827	42.52	42.52
2.Female	7,877	57.48	100.00
Total	13,704	100.00	

mes_03 Month of birth 2003

type: numeric (byte)
range: [1,99] units: 1
unique values: 13 missing .: 28,096/28,303
mean: 16.8309
std. dev: 29.3187

Variable	Obs	Mean	Std. Dev.	Min	Max
mes_03	184	6.559783	3.601261	1	12

a_o_03 Year of birth 2003

type: numeric (int)
range: [1910,9999] units: 1

unique values: 51 missing .: 28,096/28,303

mean: 2373.21
std. dev: 1810.98

Variable	Obs	Mean	Std. Dev.	Min	Max
a_o_03	196	1945.235	11.29357	1910	1974

elegible_12 Eligible for interview in 2012

type: numeric (float)
label: elegible

range: [0,2] units: 1
unique values: 3 missing .: 13,028/28,303

Eligible for interview in 2012	Freq.	Percent	Cum.
0.Not eligible for interview	415	2.72	2.72
1.Eligible for interview	14,291	93.56	96.27
2.Fallecido	569	3.73	100.00
Total	15,275	100.00	

new_sample_12 Follow-up and new sample 2012

type: numeric (byte)
label: sample

range: [1,2] units: 1
unique values: 2 missing .: 7,305/28,303

Follow-up and new sample 2012	Freq.	Percent	Cum.
1.Follow-up	14,286	68.04	68.04
2.New person	6,712	31.96	100.00
Total	20,998	100.00	

misma_v_12 Same dwelling in 2012

type: numeric (byte)
label: dwelling

range: [1,2] units: 1
unique values: 2 missing .: 9,013/28,303

Same dwelling in 2012	Freq.	Percent	Cum.
1.Same dwelling	18,788	97.40	97.40
2.Different dwelling	502	2.60	100.00
Total	19,290	100.00	

tipne_12

Type of non-interview 2012

type: numeric (byte)
label: nointerview

range: [1,20] units: 1
unique values: 20 missing .: 7,305/28,303

Type of non-interview 2012	Freq.	Percent	Cum.
1.Interviewed	18,465	87.94	87.94
2.Incomplete Interview	158	0.75	88.69
3.Postponed Interview	52	0.25	88.94
4.Absent and there is no informant	183	0.87	89.81
5.Empty household or informant not adeq	253	1.20	91.01
6.Subject unable to complete interview	12	0.06	91.07
7.Deceased without informant	120	0.57	91.64
8.Refused	403	1.92	93.56
9.Refused direct interview	236	1.12	94.69
10.Refused proxy interview	5	0.02	94.71
11.Refused next-of-kin interview	11	0.05	94.76
12.Changed location	113	0.54	95.30
13.Subject could not be located	690	3.29	98.59
14.Empty housing	95	0.45	99.04
15.Housing with temporary use	25	0.12	99.16
16.Housing with non-residential use, de	16	0.08	99.23
17.Address not located	89	0.42	99.66
18.Area not safe	11	0.05	99.71
19.Other	51	0.24	99.95
20.Deceased subject from the new sample	10	0.05	100.00
Total	20,998	100.00	

tipent_12

Type of interview 2012

type: numeric (float)
label: interview

range: [1,5] units: 1
unique values: 5 missing .: 9,838/28,303

Type of interview 2012	Freq.	Percent	Cum.
1.Direct, follow-up interview	8,868	48.03	48.03
2.Direct, new sample interview	5,580	30.22	78.25
3.Proxy, follow-up interview	959	5.19	83.44
4.Proxy, new sample interview	316	1.71	85.15
5.Next-of-kin	2,742	14.85	100.00
Total	18,465	100.00	

reason_proxy_12

Reason for Proxy Interview 2012

type: numeric (byte)
label: reason

range: [1,4] units: 1
unique values: 4 missing .: 27,028/28,303

Reason for Proxy |

Interview 2012	Freq.	Percent	Cum.
1.Health reasons	657	51.53	51.53
2.Language reasons	111	8.71	60.24
3.Temporary absence	506	39.69	99.92
4.Invalid code	1	0.08	100.00
Total	1,275	100.00	

int_date_12 Interview date 2012

type: string (str10)

unique values: 86 missing "": 9,633/28,303

examples: ""
"06/10/2012"
"14/11/2012"
"23/11/2012"

order_12 Order of interview 2012 (in the household)

type: numeric (byte)
label: o_interview

range: [1,3] units: 1
unique values: 3 missing .: 8,222/28,303

Order of interview 2012 (in the household)	Freq.	Percent	Cum.
1.First interview	14,571	72.56	72.56
2.Second interview	5,508	27.43	99.99
3.Third interview	2	0.01	100.00
Total	20,081	100.00	

fallecido_12 Died between 2003 and 2012

type: numeric (float)

range: [0,1] units: 1
unique values: 2 missing .: 9,838/28,303

Died between 2003 and 2012	Freq.	Percent	Cum.
0	15,723	85.15	85.15
1	2,742	14.85	100.00
Total	18,465	100.00	

tam_loc_12 Locality size 2012

type: numeric (float)
label: tam_loc

range: [1,4] units: 1
unique values: 4 missing .: 7,325/28,303

Locality size 2012	Freq.	Percent	Cum.
1.Population = 100,000+	12,673	60.41	60.41
2.Population = 15,000 - 99,999	2,282	10.88	71.29
3.Population = 2,500 - 14,999	2,250	10.73	82.01
4.Population <2,500	3,773	17.99	100.00
Total	20,978	100.00	

eam_12 High migration states 2012

type: numeric (byte)

range: [0,1] units: 1
unique values: 2 missing .: 9,838/28,303

High migration states 2012	Freq.	Percent	Cum.
0	12,486	67.62	67.62
1	5,979	32.38	100.00
Total	18,465	100.00	

factorh_12 Household weight 2012

type: numeric (long)

range: [0,72628] units: 1
unique values: 3,049 missing .: 7,305/28,303

mean: 1390.29
std. dev: 2837.92

Variable	Obs	Mean	Std. Dev.	Min	Max
factorh_12	20,998	1390.289	2837.92	0	72628

factori_12 Individual weight 2012

type: numeric (long)

range: [0,97546] units: 1
unique values: 3,209 missing .: 7,305/28,303

mean: 1365.85
std. dev: 3022.62

Variable	Obs	Mean	Std. Dev.	Min	Max
factori_12	20,998	1365.852	3022.617	0	97546

 factora_12 Biomarkers/Anthropometrics weight 2012

type: numeric (long)
 range: [0,133613] units: 1
 unique values: 848 missing .: 7,305/28,303
 mean: 360.481
 std. dev: 2127.92

Variable	Obs	Mean	Std. Dev.	Min	Max
factora_12	20,998	360.4812	2127.915	0	133613

 age_12 Age 2012

type: numeric (int)
 range: [21,999] units: 1
 unique values: 90 missing .: 9,445/28,303
 mean: 84.8225
 std. dev: 132.1

Variable	Obs	Mean	Std. Dev.	Min	Max
age_12	18,474	65.85049	11.51231	21	115

 sex_12 Sex 2012 (Male=1)

type: numeric (byte)
 range: [1,2] units: 1
 unique values: 2 missing .: 9,508/28,303

Sex 2012 (Male=1)	Freq.	Percent	Cum.
1	8,170	43.47	43.47
2	10,625	56.53	100.00
Total	18,795	100.00	

 subsample_12 Selected subsample for Biomarkers/Anthropometrics 2012

type: numeric (float)
 range: [0,1] units: 1
 unique values: 2 missing .: 9,835/28,303

Selected subsample for Biomarkers/ Anthropometrics 2012	Freq.	Percent	Cum.
0	16,160	87.50	87.50
1	2,308	12.50	100.00

-----+-----
 Total | 18,468 100.00

res_biomarkers_12 Result for Biomarkers/Anthropometrics 2012

type: numeric (byte)
 label: biomarkers_res

range: [1,13] units: 1
 unique values: 8 missing .: 25,999/28,303

Result for Biomarkers/Anthropometrics 2012	Freq.	Percent	Cum.
1.Complete Interview	2,086	90.54	90.54
3.Subject not fit for the interview	1	0.04	90.58
5.Absence of subject	68	2.95	93.53
6.Refusal	92	3.99	97.53
7.Subject Deceased	1	0.04	97.57
8.Change of residence	11	0.48	98.05
12.Other	28	1.22	99.26
13.Household refused to provide informa	17	0.74	100.00
Total	2,304	100.00	

biomarkers_12 Sample with Biomarkers 2012

type: numeric (float)

range: [0,1] units: 1
 unique values: 2 missing .: 25,995/28,303

Sample with Biomarkers 2012	Freq.	Percent	Cum.
0	222	9.62	9.62
1	2,086	90.38	100.00
Total	2,308	100.00	

edta_12 2012 Completed EDTA-Lavender
 Tube

type: numeric (float)

range: [0,1] units: 1
 unique values: 2 missing .: 25,982/28,303

2012 Completed EDTA- Lavender Tube	Freq.	Percent	Cum.
0	310	13.36	13.36
1	2,011	86.64	100.00
Total	2,321	100.00	

hbalc_12 2012 Completed HbA1c

```

type: numeric (float)
range: [0,1]
unique values: 2
units: 1
missing .: 25,995/28,303

```

2012 Completed HbA1c	Freq.	Percent	Cum.
0	272	11.79	11.79
1	2,036	88.21	100.00
Total	2,308	100.00	

```
-----
elegible_15                                     Eligible for interview in 2015
-----
```

```

type: numeric (float)
label: eligible

```

```

range: [0,2]
unique values: 3
units: 1
missing .: 7,305/28,303

```

Eligible for interview in 2015	Freq.	Percent	Cum.
0.Not eligible for interview	417	1.99	1.99
1.Eligible for interview	17,839	84.96	86.94
2.Fallecido	2,742	13.06	100.00
Total	20,998	100.00	

```
-----
new_sample_15                                 Follow-up and new sample/spouses 2015
-----
```

```

type: numeric (float)
label: sample15

```

```

range: [1,3]
unique values: 3
units: 1
missing .: 10,317/28,303

```

Follow-up and new sample/spouses 2015	Freq.	Percent	Cum.
1.Follow-up sample	16,983	94.42	94.42
2.New spouse from follow-up sample	306	1.70	96.12
3.New person from 2012 new sample	697	3.88	100.00
Total	17,986	100.00	

```
-----
misma_v_15                                     Same dwelling as in 2012
-----
```

```

type: numeric (byte)
label: yesno

```

```

range: [1,2]
unique values: 2
units: 1
missing .: 12,315/28,303

```

Same dwelling as in 2012	Freq.	Percent	Cum.
1.Yes	15,788	98.75	98.75
2.No	200	1.25	100.00
Total	15,988	100.00	

tipne_15 Type of non-interview 2015

type: numeric (byte)
label: nointerview15

range: [1,17] units: 1
unique values: 17 missing .: 10,317/28,303

Type of non-interview 2015	Freq.	Percent	Cum.
1.Complete Interview	15,884	88.31	88.31
2.Incomplete Interview	125	0.69	89.01
3.Postponed Interview	50	0.28	89.29
4.Absence of the subject or the proxy o	119	0.66	89.95
5.Absence of occupants or adequate info	157	0.87	90.82
6.Subject not fit for interview and wit	4	0.02	90.84
7.Subject deceased without next-of-kin	14	0.08	90.92
8.Refusal	274	1.52	92.44
9.Refusal from the subject or the proxy	278	1.55	93.99
10.Change of residence	35	0.19	94.18
11.Subject could not be located	786	4.37	98.55
12.Empty residence	68	0.38	98.93
13.Dwelling for temporary use	14	0.08	99.01
14.Dwelling with non-residential use, d	6	0.03	99.04
15.Address not located	69	0.38	99.43
16.Area not safe	10	0.06	99.48
17.Other	93	0.52	100.00
Total	17,986	100.00	

tipent_15 Type of interview 2015

type: numeric (float)
label: interview2

range: [1,5] units: 1
unique values: 5 missing .: 12,315/28,303

Type of interview 2015	Freq.	Percent	Cum.
1.Direct, follow-up interview	13,254	82.90	82.90
2.Direct, new spouse interview	596	3.73	86.63
3.Proxy, follow-up interview	884	5.53	92.16
4.Proxy, new spouse interview	45	0.28	92.44
5.Next-of-kin	1,209	7.56	100.00
Total	15,988	100.00	

reason_proxy_15 Reason for Proxy Interview 2015

type: numeric (byte)
label: reason

range: [1,3] units: 1
 unique values: 3 missing .: 27,374/28,303

Reason for Proxy Interview 2015	Freq.	Percent	Cum.
1.Health reasons	567	61.03	61.03
2.Language reasons	55	5.92	66.95
3.Temporary absence	307	33.05	100.00
Total	929	100.00	

 int_date_15 Interview date 2015

type: string (str10)
 unique values: 74 missing "": 10,629/28,303
 examples: ""
 "03/11/2015"
 "13/10/2015"
 "22/10/2015"

 c_pap_15 Interview technique - CAPI vs Paper

type: numeric (byte)
 label: capi
 range: [0,1] units: 1
 unique values: 2 missing .: 13,524/28,303

Interview technique - CAPI vs Paper	Freq.	Percent	Cum.
0.CAPI	14,576	98.63	98.63
1.Paper	203	1.37	100.00
Total	14,779	100.00	

 fallecido_15 Died between 2012 and 2015

type: numeric (float)
 range: [0,1] units: 1
 unique values: 2 missing .: 10,317/28,303

Died between 2012 and 2015	Freq.	Percent	Cum.
0	16,777	93.28	93.28
1	1,209	6.72	100.00
Total	17,986	100.00	

 tam_loc_15 Locality size 2015

type: numeric (byte)
label: tam_loc

range: [1,4] units: 1
unique values: 4 missing .: 10,317/28,303

Locality size 2015	Freq.	Percent	Cum.
1.Population = 100,000+	10,658	59.26	59.26
2.Population = 15,000 - 99,999	2,343	13.03	72.28
3.Population = 2,500 - 14,999	1,624	9.03	81.31
4.Population <2,500	3,361	18.69	100.00
Total	17,986	100.00	

eam_15 Current Residence in High Migration States from 2012

type: numeric (float)

range: [0,1] units: 1
unique values: 2 missing .: 10,317/28,303

Current Residence in High Migration States from 2012	Freq.	Percent	Cum.
0	11,126	61.86	61.86
1	6,860	38.14	100.00
Total	17,986	100.00	

factorh_15 Household weight 2015

type: numeric (long)

range: [0,51137] units: 1
unique values: 2,893 missing .: 10,317/28,303

mean: 1488.52
std. dev: 2760.26

Variable	Obs	Mean	Std. Dev.	Min	Max
factorh_15	17,986	1488.517	2760.262	0	51137

factori_15 Individual weight 2015

type: numeric (long)

range: [0,90984] units: 1
unique values: 3,186 missing .: 10,317/28,303

mean: 1254.83
std. dev: 2641.32

Variable	Obs	Mean	Std. Dev.	Min	Max
----------	-----	------	-----------	-----	-----

```
-----+-----
factori_15 |      17,986      1254.832      2641.321          0      90984
-----+-----
```

```
age_15                                                                                               Age 2015
-----+-----
```

```

      type: numeric (float)
      range: [22,999]
unique values: 85
      units: 1
      missing .: 13,510/28,303
      mean: 66.3865
      std. dev: 18.6787

```

```
-----+-----
Variable |      Obs      Mean      Std. Dev.      Min      Max
-----+-----
age_15 |    14,789    66.13422    10.66113      22     113
-----+-----
```

```
sex_15                                                                                               Sex (Male=1)
-----+-----
```

```

      type: numeric (byte)
      label: sex
      range: [1,2]
unique values: 2
      units: 1
      missing .: 10,342/28,303

```

```
-----+-----
Sex |
(Male=1) |      Freq.      Percent      Cum.
-----+-----
 1.Male |      7,666      42.68      42.68
 2.Female |     10,295      57.32     100.00
-----+-----
Total |     17,961     100.00
-----+-----
```

```
sex_validated_15                                                                                   Validated Sex 2015 (Male=1)
-----+-----
```

```

      type: numeric (float)
      label: sex
      range: [1,2]
unique values: 2
      units: 1
      missing .: 5,909/28,303

```

```
-----+-----
Validated |
Sex 2015 |
(Male=1) |      Freq.      Percent      Cum.
-----+-----
 1.Male |      9,860      44.03      44.03
 2.Female |     12,534      55.97     100.00
-----+-----
Total |     22,394     100.00
-----+-----
```

```
subsample_16                                                                                   Selected subsample for Mex-Cog 2016
-----+-----
```

```

      type: numeric (float)
      range: [0,1]
unique values: 2
      units: 1
      missing .: 13,524/28,303

```

```

Selected |
subsample |
for Mex-Cog |
  2016 |      Freq.      Percent      Cum.
-----+-----
    0 |    11,529      78.01      78.01
    1 |     3,250      21.99     100.00
-----+-----
  Total |    14,779     100.00

```

```

-----
phase_mxcog_16                                     Mex-Cog 2016 Phase
-----

```

```

      type: numeric (float)
      label: phase

      range: [1,2]                                units: 1
unique values: 2                                missing .: 25,053/28,303

```

```

  Mex-Cog |
  2016 Phase |      Freq.      Percent      Cum.
-----+-----
  1.Phase 1 |     1,821      56.03      56.03
  2.Phase 2 |     1,429      43.97     100.00
-----+-----
    Total |     3,250     100.00

```

```

-----
res_mxcog_comp_16                               Mex-Cog Result for Each Component (Character): COG/INF/ANTRO/BIOM
-----

```

```

      type: string (str4)

      unique values: 13                            missing "": 25,053/28,303

```

```

Mex-Cog Result for |
Each Component |
(Character): |
COG/INF/ANTRO/BIOM |      Freq.      Percent      Cum.
-----+-----
    0000 |         983      30.25      30.25
    0010 |          2         0.06      30.31
    0100 |         200         6.15      36.46
    0101 |          2         0.06      36.52
    0110 |         11         0.34      36.86
    0111 |         10         0.31      37.17
    1000 |          4         0.12      37.29
    1010 |        127         3.91      41.20
    1011 |         62         1.91      43.11
    1100 |          9         0.28      43.38
    1101 |          1         0.03      43.42
    1110 |       1,162      35.75      79.17
    1111 |         677      20.83     100.00
-----+-----
    Total |       3,250     100.00

```

```

-----
res_mxcog_16                                     Mex-Cog Result All Four Components
-----

```

```

      type: numeric (float)
      label: result

      range: [1,6]                                units: 1

```


unique values: 6

missing .: 25,053/28,303

Mex-Cog Result All Four Components	Freq.	Percent	Cum.
1.Complete	1,839	56.58	56.58
2.Incomplete	428	13.17	69.75
3.Refusal	267	8.22	77.97
4.Lost to Follow-up	647	19.91	97.88
5.Deceased	67	2.06	99.94
6.Subject not fit for interview & witho	2	0.06	100.00
Total	3,250	100.00	

res_cognitivo_16

Mex-Cog Result of Cognitive Assessment

type: numeric (byte)
label: result

range: [1,6]
unique values: 5

units: 1
missing .: 25,053/28,303

Mex-Cog Result of Cognitive Assessment	Freq.	Percent	Cum.
1.Complete	2,042	62.83	62.83
3.Refusal	306	9.42	72.25
4.Lost to Follow-up	807	24.83	97.08
5.Deceased	67	2.06	99.14
6.Subject not fit for interview & witho	28	0.86	100.00
Total	3,250	100.00	

res_informante_16

Mex-Cog Result of Informant Interview

type: numeric (byte)
label: result

range: [1,6]
unique values: 5

units: 1
missing .: 25,053/28,303

Mex-Cog Result of Informant Interview	Freq.	Percent	Cum.
1.Complete	2,072	63.75	63.75
3.Refusal	295	9.08	72.83
4.Lost to Follow-up	806	24.80	97.63
5.Deceased	67	2.06	99.69
6.Subject not fit for interview & witho	10	0.31	100.00
Total	3,250	100.00	

res_antro_16

Mex-Cog Result for Anthropometrics 2016

type: numeric (byte)
label: result

range: [1,6]
unique values: 5

units: 1
missing .: 25,053/28,303

Mex-Cog Result for Anthropometrics 2016	Freq.	Percent	Cum.
---	-------	---------	------

1.Complete	2,051	63.11	63.11
3.Refusal	303	9.32	72.43
4.Lost to Follow-up	804	24.74	97.17
5.Deceased	67	2.06	99.23
6.Subject not fit for interview & witho	25	0.77	100.00
Total	3,250	100.00	

biomarkers_16 Mex-Cog 2016 Sample with Biomarkers

type: numeric (float)

range: [0,1] units: 1
unique values: 2 missing .: 25,053/28,303

Mex-Cog			
2016 Sample			
with			
Biomarkers	Freq.	Percent	Cum.
0	2,502	76.98	76.98
1	748	23.02	100.00
Total	3,250	100.00	

edta_16 Mex-Cog 2016 Sample with EDTA-Lavender Tube

type: numeric (float)

range: [0,1] units: 1
unique values: 2 missing .: 25,053/28,303

Mex-Cog			
2016 Sample			
with EDTA-			
Lavender Tube	Freq.	Percent	Cum.
0	2,500	76.92	76.92
1	750	23.08	100.00
Total	3,250	100.00	

hbA1c_16 Mex-Cog 2016 Sample with HbA1c

type: numeric (float)
label: EDTA, but label does not exist

range: [0,1] units: 1
unique values: 2 missing .: 25,053/28,303

Mex-Cog			
2016 Sample			
with HbA1c	Freq.	Percent	Cum.
0	2,109	64.89	64.89
1	1,141	35.11	100.00
Total	3,250	100.00	

elegible_18 Eligible for interview in 2018

type: numeric (float)
label: elegendible

range: [0,2] units: 1
unique values: 3 missing .: 10,328/28,303

Eligible for interview in 2018	Freq.	Percent	Cum.
0.Not eligible for interview	1,054	5.86	5.86
1.Eligible for interview	15,698	87.33	93.20
2.Fallecido	1,223	6.80	100.00
Total	17,975	100.00	

new_sample_18 Follow-up and new sample/spouses 2018

type: numeric (float)
label: sample18

range: [1,4] units: 1
unique values: 4 missing .: 6,764/28,303

Follow-up and new sample/spouses 2018	Freq.	Percent	Cum.
1.R from follow-up sample	15,699	72.89	72.89
2.R from follow-up sample w/o informati	36	0.17	73.05
3.New spouse from follow-up sample	77	0.36	73.41
4.New person from 2018 new sample	5,727	26.59	100.00
Total	21,539	100.00	

tipne_18 Type of non-interview 2018

type: numeric (byte)
label: nointerview18

range: [1,16] units: 1
unique values: 15 missing .: 6,764/28,303

Type of non-interview 2018	Freq.	Percent	Cum.
1.Complete Interview	18,249	84.73	84.73
2.Incomplete Interview	7	0.03	84.76
3.Postponed Interview	180	0.84	85.59
4.Absence of the subject, or proxy or n	111	0.52	86.11
5.Absence of occupants or adequate info	722	3.35	89.46
6.Refusal	932	4.33	93.79
7.Subject not fit for interview and w/o	3	0.01	93.80
8.Subject deceased without next-of-kin	11	0.05	93.85
10.Change of residence	147	0.68	94.54
11.Subject could not be located	546	2.53	97.07
12.Residence could not be located	61	0.28	97.35
13.Other situation	425	1.97	99.33
14.Inhabited residence	96	0.45	99.77
15.Dwelling for temporary use	40	0.19	99.96
14.Dwelling with non-residential use, d	9	0.04	100.00
Total	21,539	100.00	

misma_v_18

Same dwelling as in 2018

```

type: numeric (byte)
label: yesno

range: [1,2]
unique values: 2

units: 1
missing .: 11,189/28,303

```

Same dwelling as in 2018	Freq.	Percent	Cum.
1.Yes	16,600	97.00	97.00
2.No	514	3.00	100.00
Total	17,114	100.00	

resul_hh_18

Result of Sections at Household Level 2018

```

type: numeric (float)
label: resul_h

range: [1,3]
unique values: 3

units: 1
missing .: 6,764/28,303

```

Result of Sections at Household Level 2018	Freq.	Percent	Cum.
1.Complete	17,806	82.67	82.67
2.Incomplete	873	4.05	86.72
3.Without Information	2,860	13.28	100.00
Total	21,539	100.00	

tipent_18

Type of interview 2018

```

type: numeric (float)
label: interview

range: [1,5]
unique values: 5

units: 1
missing .: 10,054/28,303

```

Type of interview 2018	Freq.	Percent	Cum.
1.Direct, follow-up interview	11,183	61.28	61.28
2.Direct, new sample interview	4,603	25.22	86.50
3.Proxy, follow-up interview	1,122	6.15	92.65
4.Proxy, new sample interview	206	1.13	93.78
5.Next-of-kin	1,135	6.22	100.00
Total	18,249	100.00	

reason_proxy_18

Reason for Proxy Interview 2018

type: numeric (byte)

label: reason
 range: [1,3] units: 1
 unique values: 3 missing .: 26,975/28,303

Reason for Proxy Interview 2018	Freq.	Percent	Cum.
1.Health reasons	709	53.39	53.39
2.Language reasons	45	3.39	56.78
3.Temporary absence	574	43.22	100.00
Total	1,328	100.00	

int_date_18 Interview date 2018

type: string (str8)
 unique values: 91 missing "": 11,160/28,303
 examples: ""
 "01/12/18"
 "12/12/18"

c_pap_18 Interview technique - Electronic vs Paper

type: numeric (byte)
 label: capi18
 range: [1,2] units: 1
 unique values: 2 missing .: 11,190/28,303

Interview technique - Electronic vs Paper	Freq.	Percent	Cum.
1.Electronic	8,969	52.41	52.41
2.Paper	8,144	47.59	100.00
Total	17,113	100.00	

fallecido_18 Died between 2015 and 2018

type: numeric (float)
 range: [0,1] units: 1
 unique values: 2 missing .: 6,764/28,303

Died between 2015 and 2018	Freq.	Percent	Cum.
0	20,404	94.73	94.73
1	1,135	5.27	100.00

Total | 21,539 100.00

tam_loc_18 Locality size 2018

type: numeric (byte)
label: tam_loc

range: [1,4] units: 1
unique values: 4 missing .: 6,764/28,303

Locality size 2018	Freq.	Percent	Cum.
1.Population = 100,000+	12,558	58.30	58.30
2.Population = 15,000 - 99,999	2,786	12.93	71.24
3.Population = 2,500 - 14,999	2,199	10.21	81.45
4.Population <2,500	3,996	18.55	100.00

Total	21,539	100.00	

eam_18 Current Residence in High Migration States from 2018

type: numeric (byte)

range: [0,1] units: 1
unique values: 2 missing .: 6,764/28,303

Current Residence in High Migration States from 2018	Freq.	Percent	Cum.
0	15,140	70.29	70.29
1	6,399	29.71	100.00

Total	21,539	100.00	

age_18 Age 2018

type: numeric (float)

range: [17,999] units: 1
unique values: 90 missing .: 10,028/28,303

mean: 67.2678
std. dev: 47.6274

Variable	Obs	Mean	Std. Dev.	Min	Max
age_18	18,229	64.95315	11.91534	17	115

sex_18 Sex (Male=1)

type: numeric (float)
label: sex

range: [1,2] units: 1
unique values: 2 missing .: 6,925/28,303

Sex (Male=1)	Freq.	Percent	Cum.
1.Male	9,165	42.87	42.87
2.Female	12,213	57.13	100.00
Total	21,378	100.00	