Statistics/Data analysis

```
      name:  <unnamed>
       log:  R:\RIECE_RELEASE V2-2016\Combine_baseline_resurvey2016\codebook_sc/b10.sc
> ml
  log type:  smcl
 opened on:  12 Feb 2026, 14:02:32

1 . codebookr _all,all

                Dataset:  \RIECE_RELEASE V2-2016\Combine_baseline_resurvey2016\stata\sc
> ramble/b10_run.dta
             Last saved:  12 Feb 2026 14:02

                  Label:  [none]
   Number of variables:  79
Number of observations:  1,666
                  Size:  3,886,778 bytes ignoring labels, etc.
          Unique Values:  A list of all of the possible non-missing values
                          for the variable and the description of the values.
 Unique Missing Values:  There are four types of missing values


              .a or -6 or RF: The subject explicitly refused to answer the question
                        when he or she should have.
              .b or -7 or NA: The subject was never asked the question
                        for one reason or another. Usually this results
                        from "skip patterns" that occur.
              .c or -8 or DK: The subject was unable to answer the question
                        either because he or she had no opinion or because
                        the required information was not available.
              .d or -9 or MI: Items should be filled out but have no data entry found.
                        This is enumerator's own mistake. The circumstances
                        can be interviewers failing to ask a question
                        or forgetting to record a response


      Numeric Missing*:   .a or -6                    String Missing*:   RF
                          .b or -7                                       NA
                          .c or -8                                       DK
                          .d or -9                                       MI

-------------------------------------------------------------------------------
pid                                                                 personal id
-------------------------------------------------------------------------------

                  type:  string (str18)

         unique values:  1,666                    missing "":  0/1,666

              examples:  "201591160601205H1"
                         "201691130216999H1"
                         "201691160104180H1"
                         "201691161706142H6"

-------------------------------------------------------------------------------
iyear                                                                      year
-------------------------------------------------------------------------------

                  type:  string (str4)

         unique values:  2                        missing "":  0/1,666

            tabulation:  Freq.  Value
                           626  "2015"
                         1,040  "2016"

-------------------------------------------------------------------------------
hhid                                                             household id
-------------------------------------------------------------------------------

                  type:  string (str15)
```

unique values: **1,413** missing "": **0/1,666**

examples: **"201591160601205"**
"201691130216999"
"201691160104180"
"201691161706142"

---

prov **province**

---

type: string (**str2**)

unique values: **2** missing "": **0/1,666**

tabulation:

Freq.	Value
1,501	**"91"**
165	**"93"**

---

amp **amphoe**

---

type: string (**str2**)

unique values: **7** missing "": **0/1,666**

tabulation:

Freq.	Value
165	**"12"**
308	**"13"**
134	**"14"**
164	**"15"**
608	**"16"**
45	**"17"**
242	**"18"**

---

tam **tambon**

---

type: string (**str2**)

unique values: **15** missing "": **0/1,666**

tabulation:

Freq.	Value
71	**"01"**
259	**"02"**
141	**"04"**
65	**"05"**
60	**"06"**
69	**"07"**
67	**"08"**
109	**"09"**
155	**"10"**
96	**"11"**
167	**"13"**
52	**"14"**
169	**"15"**
116	**"17"**
70	**"19"**

---

moo **moo**

---

type: string (**str2**)

unique values: **21** missing "": **0/1,666**

tabulation:	Freq.	Value
	168	"01"
	75	"02"
	166	"03"
	172	"04"
	151	"05"
	175	"06"
	86	"07"
	173	"08"
	108	"09"
	84	"10"
	61	"11"
	51	"12"
	46	"13"
	14	"14"
	12	"15"
	41	"16"
	10	"17"
	14	"18"
	31	"19"
	21	"22"
	7	"24"

---

strucid — **structure ID**

---

type: string (**str3**)

unique values: **182** missing "": **0/1,666**

examples: **"011"**
"034"
"071"
"157"

---

cid — **children id**

---

type: string (**str18**)

unique values: **1,666** missing "": **0/1,666**

examples: **"201591160601205H1"**
"201691130216999H1"
"201691160104180H1"
"201691161706142H6"

---

ch10 — **Chicken Pox**

---

type: numeric (**byte**)
label: **ch10**

range: [**1,5**] units: **1**
unique values: **3** missing .: **0/1,666**
unique missing codes: **1** missing *: **3/1,666**

tabulation:	Freq.	Numeric	Label
	116	1	yes
	1,493	3	no
	54	5	not sure
	3	.b	

---

ch11 — **Influenza vaccine (in past 12 months)**

---

type: numeric (**byte**)
label: **ch11**

range: [**1,5**] units: **1**
unique values: **3** missing .: **0/1,666**
unique missing codes: **1** missing *: **3/1,666**

tabulation:	Freq.	Numeric	Label
	117	**1**	yes
	1,483	**3**	no
	63	**5**	not sure
	3	**.b**	

---

ch12 **HIB (Vaccine against meningitis)**

---

type: numeric (**byte**)
label: **ch12**

range: [**1,5**] units: **1**
unique values: **3** missing .: **0/1,666**
unique missing codes: **1** missing *: **3/1,666**

tabulation:	Freq.	Numeric	Label
	116	**1**	yes
	1,479	**3**	no
	68	**5**	not sure
	3	**.b**	

---

ch38 **maternal and child health book (pink book) was recorded**

---

type: numeric (**byte**)
label: **ch38**

range: [**1,3**] units: **1**
unique values: **2** missing .: **1,111/1,666**

tabulation:	Freq.	Numeric	Label
	441	**1**	yes
	114	**3**	no
	1,111	**.**	

---

nvac_1 **BCG (Tuberculosis vaccine) (not display)**

---

type: string (**str197**), but longest is str0

unique values: **0** missing "": **1,666/1,666**

tabulation:	Freq.	Value
	1,666	**""**

---

ch3_1 **Has the child received vaccine in the folloing lists? 1**

---

type: numeric (**byte**)
label: **ch3**

range: [**1,5**] units: **1**
unique values: **3** missing .: **0/1,666**
unique missing codes: **1** missing *: **5/1,666**

tabulation:	Freq.	Numeric	Label
	1,622	**1**	yes
	8	**3**	no
	31	**5**	not sure
	5	**.d**	

---

ch3b_1 **Where does that data come from? 1**

---

type: numeric (**byte**)
label: **ch3b**

range: [**1,3**] units: **1**
unique values: **2** missing .: **0/1,666**
unique missing codes: **1** missing *: **6/1,666**

tabulation:	Freq.	Numeric	Label
	1,415	**1**	from child's health book (pink book)
	245	**3**	primary caregiver's answer
	6	**.d**	

---

nvac_2 **HBV (Hepatitis-B vaccine) (not display)**

---

type: string (**str197**), but longest is str0

unique values: **0** missing "": **1,666/1,666**

tabulation:	Freq.	Value
	1,666	""

---

ch3_2 **Has the child received the vaccine in the following lists? 2**

---

type: numeric (**byte**)
label: **ch3**

range: [**1,5**] units: **1**
unique values: **3** missing .: **0/1,666**
unique missing codes: **1** missing *: **5/1,666**

tabulation:	Freq.	Numeric	Label
	1,628	**1**	yes
	3	**3**	no
	30	**5**	not sure
	5	**.d**	

---

ch3b_2 **Where does that data come from? 2**

---

type: numeric (**byte**)
label: **ch3b**

range: [**1,3**] units: **1**
unique values: **2** missing .: **0/1,666**
unique missing codes: **1** missing *: **6/1,666**

tabulation:	Freq.	Numeric	Label
	1,416	**1**	from child's health book (pink book)
	244	**3**	primary caregiver's answer
	6	**.d**	

---

nvac_3 **OPV (Polio vaccine) (not display)**

---

type: string (**str197**), but longest is str0

unique values: **0** missing "": **1,666/1,666**

tabulation: Freq. Value
1,666 ""

---

ch3_3 **Has the child received vaccine in the folloing lists? 3**

---

type: numeric (**byte**)
label: **ch3**

range: [**1,5**] units: **1**
unique values: **3** missing .: **0/1,666**
unique missing codes: **1** missing *: **5/1,666**

tabulation: Freq. Numeric Label
1,617 **1** yes
14 **3** no
30 **5** not sure
5 **.d**

---

ch3b_3 **Where does that data come from? 3**

---

type: numeric (**byte**)
label: **ch3b**

range: [**1,3**] units: **1**
unique values: **2** missing .: **0/1,666**
unique missing codes: **1** missing *: **6/1,666**

tabulation: Freq. Numeric Label
1,415 **1** from child's health book (pink book)
245 **3** primary caregiver's answer
6 **.d**

---

nvac_4 **DTP,DPT (vaccine against diphtheria- tetanus- whooping cough) (not display)**

---

type: string (**str197**), but longest is str0

unique values: **0** missing "": **1,666/1,666**

tabulation: Freq. Value
1,666 ""

---

ch3_4 **Has the child received vaccine in the folloing lists? 4**

---

type: numeric (**byte**)
label: **ch3**

range: [**1,5**] units: **1**
unique values: **3** missing .: **0/1,666**
unique missing codes: **1** missing *: **5/1,666**

tabulation: Freq. Numeric Label
1,606 **1** yes
26 **3** no
29 **5** not sure
5 **.d**

ch3b_4 **Where does that data come from? 4**

type:	numeric (**byte**)		
label:	**ch3b**		
range:	[**1**,**3**]	units:	**1**
unique values:	**2**	missing .:	**0**/**1,666**
unique missing codes:	**1**	missing *:	**6**/**1,666**

tabulation:	Freq.	Numeric	Label
	1,415	**1**	from child's health book (pink book)
	245	**3**	primary caregiver's answer
	6	**.d**	

nvac_5 **M or MR or MMR (vaccine against measles - mumps - rubella) (not display)**

type:	string (**str197**), but longest is str0		
unique values:	**0**	missing "":	**1,666**/**1,666**

tabulation:	Freq.	Value
	1,666	""

ch3_5 **Has the child received vaccine in the folloing lists? 5**

type:	numeric (**byte**)		
label:	**ch3**		
range:	[**1**,**5**]	units:	**1**
unique values:	**3**	missing .:	**0**/**1,666**
unique missing codes:	**1**	missing *:	**5**/**1,666**

tabulation:	Freq.	Numeric	Label
	1,617	**1**	yes
	14	**3**	no
	30	**5**	not sure
	5	**.d**	

ch3b_5 **Where does that data come from? 5**

type:	numeric (**byte**)		
label:	**ch3b**		
range:	[**1**,**3**]	units:	**1**
unique values:	**2**	missing .:	**0**/**1,666**
unique missing codes:	**1**	missing *:	**6**/**1,666**

tabulation:	Freq.	Numeric	Label
	1,414	**1**	from child's health book (pink book)
	246	**3**	primary caregiver's answer
	6	**.d**	

nvac_6 **JE (Encephalitis vaccine) (not display)**

type:	string (**str197**), but longest is str0		
unique values:	**0**	missing "":	**1,666**/**1,666**

tabulation: Freq. Value
1,666 ""

---

ch3_6 **Has the child received vaccine in the folloing lists? 6**

---

type: numeric (**byte**)
label: **ch3**

range: [**1,5**] units: **1**
unique values: **3** missing .: **0/1,666**
unique missing codes: **1** missing *: **5/1,666**

tabulation: Freq. Numeric Label
1,595 **1** yes
37 **3** no
29 **5** not sure
5 **.d**

---

ch3b_6 **Where does that data come from? 6**

---

type: numeric (**byte**)
label: **ch3b**

range: [**1,3**] units: **1**
unique values: **2** missing .: **0/1,666**
unique missing codes: **1** missing *: **6/1,666**

tabulation: Freq. Numeric Label
1,414 **1** from child's health book (pink book)
246 **3** primary caregiver's answer
6 **.d**

---

ch13_hospital **Birth place on hospital?**

---

type: numeric (**float**)
label: **ch13_hospital**

range: [**0,1**] units: **1**
unique values: **2** missing .: **252/1,666**

tabulation: Freq. Numeric Label
67 **0** no
1,347 **1** yes
252 **.**

---

ch13 **Birth place (unvailable)**

---

type: string (**str98**), but longest is str0

unique values: **0** missing "": **1,666/1,666**

tabulation: Freq. Value
1,666 ""

---

ch14 **By whom**

---

type: numeric (**byte**)
label: **ch14**

range:	[1,5]	units:	1
unique values:	3	missing .:	0/1,666
unique missing codes:	1	missing *:	277/1,666

tabulation:	Freq.	Numeric	Label
	750	1	doctor
	635	3	nurse
	4	5	others (specify)
	277	.b	

---

ch14_des — **Other (specify) : By whom (not display)**

---

type:	string (**str42**), but longest is str0		
unique values:	0	missing "":	1,666/1,666

tabulation:	Freq.	Value
	1,666	""

---

ch15_w — **Week : Age of pregnancy**

---

type:	numeric (**byte**)		
range:	[28,44]	units:	1
unique values:	16	missing .:	0/1,666
unique missing codes:	2	missing *:	298/1,666

tabulation:	Freq.	Value
	1	28
	1	29
	5	30
	5	31
	4	32
	10	33
	16	34
	28	35
	65	36
	188	37
	402	38
	358	39
	226	40
	52	41
	6	42
	1	44
	293	.b
	5	.c
mean:	38.2646	
std. dev:	1.68087	

percentiles:	10%	25%	50%	75%	90%
	37	38	38	39	40

---

ch15_d — **Day : Age of pregnancy**

---

type:	numeric (**byte**)		
range:	[0,15]	units:	1
unique values:	12	missing .:	0/1,666
unique missing codes:	2	missing *:	298/1,666

tabulation:	Freq.	Value
	558	0
	169	1
	146	2
	111	3
	144	4
	118	5
	115	6
	1	7
	3	8
	1	9
	1	11
	1	15
	293	.b
	5	.c

mean: 1.98538
std. dev: 2.16741

percentiles:	10%	25%	50%	75%	90%
	0	0	1	4	5

---

ch16 How did you give birth (Caesarean section or normal labor) (not display)

---

type: string (**str96**), but longest is str0

unique values: **0** missing "": **1,666/1,666**

tabulation:	Freq.	Value
	1,666	""

---

ch17 Complications during delivery

---

type: numeric (**byte**)
label: **ch17**

range: [**1,3**] units: **1**
unique values: **2** missing .: **0/1,666**
unique missing codes: **1** missing *: **505/1,666**

tabulation:	Freq.	Numeric	Label
	1,112	1	no
	49	3	yes (specify)
	505	.b	

---

ch17_des yes (specify) : Complications during delivery (not display)

---

type: string (**str66**), but longest is str0

unique values: **0** missing "": **1,666/1,666**

tabulation:	Freq.	Value
	1,666	""

---

ch18 Complications after birth

---

type: numeric (**byte**)
label: **ch18**

range: [**1,3**] units: **1**
unique values: **2** missing .: **0/1,666**
unique missing codes: **1** missing *: **735/1,666**

tabulation:	Freq.	Numeric	Label
	916	**1**	no
	15	**3**	yes (specify)
	735	**.b**	

---

ch18_des **yes (specify) : Complications after birth (not display)**

---

type: string (**str63**), but longest is str0

unique values: **0** missing "": **1,666/1,666**

tabulation:	Freq.	Value
	1,666	**""**

---

ch19 **Stressed**

---

type: numeric (**byte**)
label: **ch19**

range: [**3,3**] units: **1**
unique values: **1** missing .: **0/1,666**
unique missing codes: **1** missing *: **1,658/1,666**

tabulation:	Freq.	Numeric	Label
	8	**3**	no
	1,658	**.b**	

---

ch20 **Depressed**

---

type: numeric (**byte**)
label: **ch20**

range: [**3,3**] units: **1**
unique values: **1** missing .: **0/1,666**
unique missing codes: **1** missing *: **1,658/1,666**

tabulation:	Freq.	Numeric	Label
	8	**3**	no
	1,658	**.b**	

---

ch21 **Drinking alcohol**

---

type: numeric (**byte**)
label: **ch21**

range: [**3,3**] units: **1**
unique values: **1** missing .: **0/1,666**
unique missing codes: **1** missing *: **1,658/1,666**

tabulation:	Freq.	Numeric	Label
	8	**3**	no
	1,658	**.b**	

---

ch22 **Date of Birth (unvailable)**

---

type: string (**str20**), but longest is str0

unique values: **0** missing "": **1,666/1,666**

tabulation:	Freq.	Value
	1,666	**""**

ch22_d — **Day of Birth (unvailable)**

type:	numeric (**byte**)		
range:	[.,.]	units:	.
unique values:	**0**	missing .:	**1,666/1,666**

tabulation:

Freq.	Value
1,666	.

mean: .
std. dev: .

percentiles:	10%	25%	50%	75%	90%
	.	.	.	.	.

ch22_m — **Month of Birth (unvailable)**

type:	numeric (**byte**)		
label:	**ch22_m**		
range:	[.,.]	units:	.
unique values:	**0**	missing .:	**1,666/1,666**

tabulation:

Freq.	Numeric	Label
1,666	.	

ch22_y — **Year of Birth (unvailable)**

type:	numeric (**int**)		
range:	[.,.]	units:	.
unique values:	**0**	missing .:	**1,666/1,666**

tabulation:

Freq.	Value
1,666	.

mean: .
std. dev: .

percentiles:	10%	25%	50%	75%	90%
	.	.	.	.	.

ch23 — **Gender**

type:	numeric (**byte**)		
label:	**ch23**		
range:	[**1,3**]	units:	**1**
unique values:	**2**	missing .:	**0/1,666**

tabulation:

Freq.	Numeric	Label
860	**1**	male
806	**3**	female

ch24 — **Weight at birth**

type:	numeric (**double**)		
range:	[**1,5.21**]	units:	**.0001**
unique values:	**345**	missing .:	**0/1,666**
unique missing codes:	**1**	missing *:	**38/1,666**

mean:	**3.04945**
std. dev:	**.469257**

percentiles:	10%	25%	50%	75%	90%
	2.48	**2.78**	**3.0575**	**3.33**	**3.59**

---

ch25 **Length at birth**

---

type:	numeric (**float**)		
label:	**ch25**, but **32** nonmissing values are not labeled		
range:	[**27,63**]	units:	**.1**
unique values:	**32**	missing .:	**0**/**1,666**
unique missing codes:	**2**	missing *:	**114**/**1,666**

tabulation:	Freq.	Numeric	Label
	1	**27**	
	1	**28**	
	3	**31**	
	1	**32**	
	3	**33**	
	2	**37**	
	1	**39**	
	3	**41**	
	4	**42**	
	8	**43**	
	2	**44**	
	15	**45**	
	23	**46**	
	48	**47**	
	1	**47.5**	
	89	**48**	
	133	**49**	
	1	**49.5**	
	197	**50**	
	205	**51**	
	1	**51.5**	
	242	**52**	
	197	**53**	
	165	**54**	
	91	**55**	
	54	**56**	
	37	**57**	
	17	**58**	
	4	**59**	
	1	**61**	
	1	**62**	
	1	**63**	
	113	**.c**	
	1	**.d**	

---

ch26 **Head circumference**

---

type:	numeric (**double**)		
label:	**ch26**, but **25** nonmissing values are not labeled		
range:	[**25,54**]	units:	**.1**
unique values:	**25**	missing .:	**0**/**1,666**
unique missing codes:	**2**	missing *:	**193**/**1,666**

tabulation:	Freq.	Numeric	Label
	1	25	
	3	27	
	8	28	
	16	29	
	88	30	
	1	30.5	
	189	31	
	8	31.5	
	327	32	
	12	32.5	
	351	33	
	5	33.5	
	253	34	
	5	34.5	
	139	35	
	1	35.5	
	43	36	
	2	36.5	
	6	37	
	7	38	
	1	40	
	3	48	
	1	49	
	2	52	
	1	54	
	191	.b	
	2	.d	

---

ch27 **Apgar Score(1 minute)**

---

type: numeric (**byte**)
label: **ch27**, but **9** nonmissing values are not labeled

range: [**1,10**] units: **1**
unique values: **9** missing .: **0**/**1,666**
unique missing codes: **2** missing *: **194**/**1,666**

tabulation:	Freq.	Numeric	Label
	1	1	
	1	3	
	1	4	
	7	5	
	12	6	
	24	7	
	146	8	
	1,175	9	
	105	10	
	192	.b	
	2	.d	

---

ch28 **Apgar Score(5 minute)**

---

type: numeric (**byte**)
label: **ch28**, but **5** nonmissing values are not labeled

range: [**6,10**] units: **1**
unique values: **5** missing .: **0**/**1,666**
unique missing codes: **2** missing *: **202**/**1,666**

tabulation:	Freq.	Numeric	Label
	5	6	
	4	7	
	23	8	
	125	9	
	1,307	10	
	197	.b	
	5	.d	

---

ch29 **Congenital disorder**

---

type: numeric (**byte**)
label: **ch29**

range: [**1,3**] units: **1**
unique values: **2** missing .: **0/1,666**
unique missing codes: **1** missing *: **383/1,666**

tabulation:	Freq.	Numeric	Label
	21	1	yes
	1,262	3	no
	383	.b	

---

ch29_des **yes (specify) : Congenital disorder (not display)**

---

type: string (**str2**), but longest is str0

unique values: **0** missing "": **1,666/1,666**

tabulation:	Freq.	Value
	1,666	""

---

ch30 **Newborn health**

---

type: numeric (**byte**)
label: **ch30**

range: [**1,3**] units: **1**
unique values: **2** missing .: **0/1,666**
unique missing codes: **1** missing *: **595/1,666**

tabulation:	Freq.	Numeric	Label
	1,035	1	good
	36	3	abnormal (specify)
	595	.b	

---

ch30_des **abnormal (specify) (not display)**

---

type: string (**str124**), but longest is str0

unique values: **0** missing "": **1,666/1,666**

tabulation:	Freq.	Value
	1,666	""

---

ch31 **Discharged date (unvailable)**

---

type: string (**str27**), but longest is str0

unique values: **0** missing "": **1,666/1,666**

tabulation: Freq. Value
1,666 ""

---

ch31_d — **Day of discharge (unvailable)**

---

type: numeric (**byte**)

range: [.,.] units: .
unique values: **0** missing .: **1,666/1,666**

tabulation: Freq. Value
1,666 .
mean: .
std. dev: .

percentiles: 10% 25% 50% 75% 90%
. . . . .

---

ch31_m — **Month of discharge (unvailable)**

---

type: numeric (**byte**)
label: **ch31_m**

range: [.,.] units: .
unique values: **0** missing .: **1,666/1,666**

tabulation: Freq. Numeric Label
1,666 .

---

ch31_y — **Year of discharge (unvailable)**

---

type: numeric (**int**)

range: [.,.] units: .
unique values: **0** missing .: **1,666/1,666**

tabulation: Freq. Value
1,666 .
mean: .
std. dev: .

percentiles: 10% 25% 50% 75% 90%
. . . . .

---

ch32 — **Weight at discharge date**

---

type: numeric (**int**)

range: [**1720,4450**] units: **1**
unique values: **99** missing .: **0/1,666**
unique missing codes: **2** missing *: **1,502/1,666**

tabulation: Freq. Value
1 1720
1 1930
1 2020
1 2130
1 2200
1 2240
1 2260
1 2270
1 2300
1 2310
1 2350

3	2360
1	2370
1	2400
1	2430
1	2435
2	2450
2	2460
1	2465
1	2480
1	2502
1	2525
1	2526
1	2540
2	2550
4	2600
2	2610
3	2620
1	2625
1	2630
2	2650
3	2660
2	2670
1	2690
4	2700
1	2710
1	2726
3	2750
3	2755
1	2770
1	2788
10	2800
1	2828
1	2831
1	2840
1	2850
1	2860
1	2880
1	2890
2	2900
3	2910
4	2920
4	2930
1	2940
1	2950
1	2960
1	2965
1	2990
5	3000
1	3035
1	3040
1	3050
1	3055
1	3070
1	3075
1	3088
9	3100
2	3110
1	3150
2	3160
1	3180
1	3210
1	3220
1	3240
3	3250
3	3300
1	3315
1	3320
1	3325
1	3330
1	3345
1	3350
1	3360

```
                      1  3370
                      1  3380
                      1  3390
                      4  3400
                      1  3430
                      2  3440
                      3  3450
                      1  3460
                      1  3470
                      2  3500
                      1  3600
                      1  3650
                      1  3700
                      1  3710
                      1  3960
                      1  4450
                  1,490  .b
                     12  .c
            mean:   2906.82
       std. dev:   401.509

     percentiles:        10%       25%       50%       75%       90%
                        2430      2640      2905      3160      3400
```

---

ch33 **Vitamin k**

```
            type:  numeric (byte)
           label:  ch33

           range:  [1,3]                        units:  1
     unique values:  2                       missing .:  0/1,666
unique missing codes:  1                     missing *:  329/1,666

     tabulation:  Freq.   Numeric  Label
                  1,331         1  inject
                      6         3  not spray
                    329        .b
```

---

ch34 **Newborn Screening (unavailable)**

```
            type:  string (str29), but longest is str0

     unique values:  0                       missing "":  1,666/1,666

     tabulation:  Freq.  Value
                  1,666  ""
```

---

ch34_d **Month of newborn screening (unavailable)**

```
            type:  numeric (byte)

           range:  [.,.]                        units:  .
     unique values:  0                       missing .:  1,666/1,666

     tabulation:  Freq.  Value
                  1,666  .
            mean:        .
       std. dev:        .

     percentiles:       10%       25%       50%       75%       90%
                          .         .         .         .         .
```

---

ch34_m **Year of newborn screening (unavailable)**

---

type: numeric (**byte**)
label: **ch34_m**

range: [.,.] units: .
unique values: **0** missing .: **1,666/1,666**

tabulation: Freq. Numeric Label
1,666 .

---

ch34_y **Year of newborn screening (unvailable)**

---

type: numeric (**int**)

range: [.,.] units: .
unique values: **0** missing .: **1,666/1,666**

tabulation: Freq. Value
1,666 .
mean: .
std. dev: .

percentiles: 10% 25% 50% 75% 90%
. . . . .

---

ch35 **Congenital Hypothyroidism Screening**

---

type: numeric (**byte**)
label: **ch35**

range: [**1,1**] units: **1**
unique values: **1** missing .: **0/1,666**
unique missing codes: **1** missing *: **1,621/1,666**

tabulation: Freq. Numeric Label
45 **1** normal
1,621 **.b**

---

ch36 **PKU Screening**

---

type: numeric (**byte**)
label: **ch36**

range: [**1,1**] units: **1**
unique values: **1** missing .: **0/1,666**
unique missing codes: **1** missing *: **1,657/1,666**

tabulation: Freq. Numeric Label
9 **1** normal
1,657 **.b**

---

ch37 **Interviewer note (unvailable)**

---

type: string (**str328**), but longest is str0

unique values: **0** missing "": **1,666/1,666**

tabulation: Freq. Value
1,666 ""

---

note_cleaner **Data cleaner note (not display)**

---

type: string (**str76**), but longest is str0

unique values: **0** missing "": **1,666/1,666**

tabulation: Freq. Value
1,666 ""

---

year_survey **year survey**

---

type: numeric (**float**)

range: [**2016,2016**] units: **1**
unique values: **1** missing .: **0/1,666**

tabulation: Freq. Value
1,666 **2016**
mean: **2016**
std. dev: **0**

percentiles: 10% 25% 50% 75% 90%
2016 **2016** **2016** **2016** **2016**

---

ch_added **children added**

---

type: numeric (**float**)
label: **ch_add**

range: [**0,1**] units: **1**
unique values: **2** missing .: **0/1,666**

tabulation: Freq. Numeric Label
1,595 **0** no
71 **1** yes

---

survey_name **survey round**

---

type: string (**str12**)

unique values: **2** missing "": **0/1,666**

tabulation: Freq. Value
1,040 **"BASELINE2016"**
626 **"RESURVEY2016"**

---

ch22_age **Child's Age**

---

type: numeric (**double**)

range: [**.08,6.41**] units: **.001**
unique values: **75** missing .: **0/1,666**

tabulation: Freq. Value
2 **.08**
2 **.25**
1 **.33**
2 **.42**
1 **.5**
4 **.58**
2 **.666**
4 **.75**
4 **.833**
6 **.92**
12 **1**
19 **1.08**

20	1.16
22	1.25
30	1.33
29	1.41
25	1.5
19	1.58
17	1.66
31	1.75
32	1.83
26	1.91
36	2
31	2.08
27	2.16
34	2.25
27	2.33
19	2.41
28	2.5
25	2.58
36	2.66
23	2.75
43	2.83
36	2.91
41	3
16	3.08
28	3.16
35	3.25
32	3.33
24	3.41
29	3.5
43	3.58
38	3.66
44	3.75
53	3.83
47	3.91
43	4
29	4.08
24	4.16
27	4.25
22	4.33
25	4.41
21	4.5
30	4.58
26	4.66
1	4.72
28	4.75
30	4.83
17	4.91
42	5
23	5.08
19	5.16
17	5.25
29	5.33
20	5.41
20	5.5
24	5.58
13	5.66
11	5.75
6	5.83
4	5.91
5	6
1	6.25
1	6.33
3	6.41

mean:	3.33545
std. dev:	1.33329

percentiles:	10%	25%	50%	75%	90%
	1.5	2.25	3.41	4.33	5.16

ch31_month — **Time between birthday and discharge day**

type: numeric (**double**)

range: [**0,2.06**] units: **.01**
unique values: **20** missing .: **1,483/1,666**

tabulation:	Freq.	Value
	1	0
	1	.04
	54	.06
	50	.07
	45	.1
	14	.13
	2	.16
	1	.18
	1	.19
	1	.2
	4	.23
	1	.24
	1	.26
	1	.35
	1	.37
	1	.45
	1	1.06
	1	1.12
	1	1.41
	1	2.06
	1,483	.

mean: .121475
std. dev: .209595

percentiles:	10%	25%	50%	75%	90%
	.06	.06	.07	.1	.13

ch34_month — **Time between birthday and newborn screening day**

type: numeric (**double**)

range: [**0,2.06**] units: **.01**
unique values: **16** missing .: **974/1,666**

tabulation:	Freq.	Value
	4	0
	3	.03
	1	.04
	324	.06
	210	.07
	114	.1
	7	.11
	16	.13
	2	.16
	1	.18
	2	.19
	1	.29
	1	.43
	1	.47
	3	1.06
	2	2.06
	974	.

mean: .083656
std. dev: .128194

percentiles:	10%	25%	50%	75%	90%
	.06	.06	.07	.07	.1

```
2 . log close
      name:  <unnamed>
       log:  R:\RIECE_RELEASE V2-2016\Combine_baseline_resurvey2016\codebook_sc/b10.sc
> ml
  log type:  smcl
 closed on:  12 Feb 2026, 14:02:45
```