

DS5011 Project Description

DS5011 Project: Behavioral Risk Factor Surveillance System Dashboard

Introduction

BRFSS, or the Behavioral Risk Factor Surveillance System, is the world's largest and longest-running telephone survey system for collecting health-related information from adults in the U.S.. It is a collaborative project between the Centers for Disease Control and Prevention (CDC) and all U.S. states, the District of Columbia, and U.S. territories. The survey gathers data on health behaviors, chronic conditions, and preventive care to monitor public health and inform policy.

What BRFSS collects

- **Health-related risk behaviors:** Such as tobacco use, alcohol consumption, and physical activity.
- **Chronic health conditions:** Data on conditions like heart disease, diabetes, and cancer.
- **Preventive care:** Information on access to and use of health services, including vaccinations and screenings.
- **Demographics and other factors:** Includes information on health care access, sleep patterns, and other demographic data.

How BRFSS is conducted

- **Surveys:** Telephone surveys are conducted monthly using landlines and cellular phones.
- **Data collection:** Each state conducts its own survey with the assistance of the CDC, and the data is aggregated at the state level.
- **Population:** The target population is the civilian noninstitutionalized adult population aged 18 and older.

- **Scale:** The survey system completes over 400,000 adult interviews each year across all participating states and territories.

Why BRFSS is important

- **Monitoring health:** It provides state-specific data that helps monitor public health trends and assess progress toward national health objectives.
- **Informing policy:** The data is used to design and evaluate public health programs and policies aimed at improving the health of the population.
- **Research:** It serves as a crucial data source for researchers studying health-related issues across the country.

About the Dataset

The BRFSS data repository is enormous and not available to the public. However, summary statistics are publicly available [here](#) in *.csv format at the relatively modest size of 1GB. Please download this summary data into your IDE as tibble `df` to explore it with this markdown file.

The *.csv file has 2763102 rows and 27 columns.

For each question asked in column `Question`, summary statistics include *percentage confidence intervals* for each possible response value in column `Response`. The left (right) endpoint of this interval is in `Confidence_limit_Low` (`Confidence_limit_High`). Most importantly for your project goals, these percentages are computed on grouped data. Grouping is done always using columns `Locationabbr` (state/territory) and `Year`. For this broadest grouping, columns `Break_Out` and `Break_Out_Category` take value “Overall”. Further grouping is indicated in those columns, including by: age, household income, sex, race, and highest education level.

Note that the use of ID columns in large table provide *keys* to the values in other columns. Key matching is much more efficient than value matching. Here is the key:value list between `BreakOutCategoryID` and `Break_Out_Category`, for example

```
# A tibble: 6 x 3
# Groups:   BreakOutCategoryID [6]
  BreakOutCategoryID Break_Out_Category  count
  <chr>              <chr>            <int>
1 CAT1               Overall           116860
2 CAT2               Sex              221825
3 CAT3               Age Group        621025
4 CAT4               Race/Ethnicity  768868
5 CAT5               Education Attained 439384
6 CAT6               Household Income  595140
```

Survey questions are assigned both a `Topic` and a `Class`, which may help in organizing the questions into layers. For example, here is a possible layering scenario.

```
layerQ <- df |>
  select(Class, Topic, Question) |>
  distinct()
layerQ |>
  group_by(Class, Topic) |>
  summarize(numberQuestions=n()) |>
  print(n=75)
```

	Class	Topic	numberQuestions
	<chr>	<chr>	<int>
1	Alcohol Consumption	Alcohol Consumption	1
2	Alcohol Consumption	Binge Drinking	1
3	Alcohol Consumption	Heavy Drinking	2
4	Cholesterol Awareness	Cholesterol Checked	2
5	Cholesterol Awareness	Cholesterol High	1
6	Chronic Health Indicators	Arthritis	4
7	Chronic Health Indicators	Asthma	2
8	Chronic Health Indicators	COPD	1
9	Chronic Health Indicators	Cardiovascular Disease	4
10	Chronic Health Indicators	Depression	1
11	Chronic Health Indicators	Diabetes	1
12	Chronic Health Indicators	Kidney	1
13	Chronic Health Indicators	Other Cancer	1
14	Chronic Health Indicators	Skin Cancer	1
15	Chronic Health Indicators	Vision	1
16	Colorectal Cancer Screening	Blood Stool Test	4
17	Colorectal Cancer Screening	Colonoscopy	1
18	Colorectal Cancer Screening	Sigmoidoscopy	2
19	Colorectal Cancer Screening	USPSTF Recommendations	2
20	Days of Poor Health	Healthy Days	2
21	Demographics	Age	1
22	Demographics	Disability status	7
23	Demographics	Education	1
24	Demographics	Employment	1
25	Demographics	Hearing	1
26	Demographics	Income	1
27	Demographics	Internet	1

28 Demographics	Marital Status	1
29 Demographics	Number of Children	1
30 Demographics	Race	1
31 Demographics	Rent/Own Home	1
32 Demographics	Sex	1
33 Demographics	Veteran Status	1
34 E-Cigarette Use	E-Cigarette Use	5
35 Fruits and Vegetables	Fruit Consumption	1
36 Fruits and Vegetables	Vegetable Consumption	1
37 HIV-AIDS	HIV Test	1
38 Health Care Access/Coverage	Health Care Cost	2
39 Health Care Access/Coverage	Health Care Coverage	2
40 Health Care Access/Coverage	Last Checkup	1
41 Health Care Access/Coverage	Personal Care Provider	2
42 Health Care Access/Coverage	Under 65 Coverage	1
43 Health Status	Fair or Poor Health	1
44 Health Status	Overall Health	1
45 Hypertension Awareness	High Blood Pressure	1
46 Immunization	Flu Shot	1
47 Immunization	Pneumonia Vaccination	1
48 Immunization	Shingle Vaccination	1
49 Immunization	Tetanus Shot	1
50 Injury	Drink and Drive	1
51 Injury	Seatbelt Use	2
52 Lung Cancer Screening	Had CAT/CT Chest Scan	1
53 Oral Health	All Teeth Removed	1
54 Oral Health	Dental Visit	1
55 Oral Health	Teeth Removed	1
56 Overweight and Obesity (BMI)	BMI Categories	1
57 Physical Activity	Aerobic Activity	1
58 Physical Activity	Exercise	1
59 Physical Activity	Physical Activity Index	1
60 Physical Activity	Strength Activity	2
61 Prostate Cancer	PSA Test	1
62 Tobacco Use	Current Smoker Status	1
63 Tobacco Use	Smokeless Tobacco	1
64 Tobacco Use	Smoker Status	1
65 Women's Health	Mammogram	3
66 Women's Health	Pap Test	2

Finally, the survey questions themselves have IDs (sometimes several).

```
# A tibble: 228 x 4
```

TopicId	ClassId	QuestionID	Question	
<chr>	<chr>	<chr>	<chr>	
1	TOPIC01	CLASS15	_PAINDEX3	Participated in 150 minutes or more of Aerobic Ph~
2	TOPIC01	CLASS15	_PAINDEX2	Participated in 150 minutes or more of Aerobic Ph~
3	TOPIC01	CLASS15	_PAINDEX1	Participated in 150 minutes or more of Aerobic Ph~
4	TOPIC01	CLASS15	_PAINDEX	Participated in 150 minutes or more of Aerobic Ph~
5	TOPIC02	CLASS05	AGE	What is your age?
6	TOPIC03	CLASS01	DRNKANY6	Adults who have had at least one drink of alcohol~
7	TOPIC03	CLASS01	DRNKANY5	Adults who have had at least one drink of alcohol~
8	TOPIC04	CLASS13	_ALTETH3	Adults aged 65+ who have had all their natural te~
9	TOPIC04	CLASS13	_ALTETH2	Adults aged 65+ who have had all their natural te~
10	TOPIC05	CLASS03	_DRDXAR2	Adults who have been told they have arthritis (va~
11	TOPIC05	CLASS03	_DRDXAR3	Adults who have been told they have arthritis (va~
12	TOPIC05	CLASS03	_DRDXAR1	Adults who have been told they have arthritis (va~
13	TOPIC05	CLASS03	_LMTACT3	Are you now limited in any way in any of your usu~
14	TOPIC05	CLASS03	_LMTACT2	Are you now limited in any way in any of your usu~
15	TOPIC05	CLASS03	_LMTACT1	Are you now limited in any way in any of your usu~
16	TOPIC05	CLASS03	_LMTWRK3	Do arthritis or joint symptoms now affect whether~
17	TOPIC05	CLASS03	_LMTWRK2	Do arthritis or joint symptoms now affect whether~
18	TOPIC05	CLASS03	_LMTWRK1	Do arthritis or joint symptoms now affect whether~
19	TOPIC05	CLASS03	_LMTSCL1	Does arthritis or joint symptoms interfere with y~
20	TOPIC06	CLASS03	_CASTHM1	Adults who have been told they currently have ast~
# i 208 more rows				

The output below shows that there are approximately 100 questions and some of the questions have multiple values of QuestionID.

# A tibble: 99 x 2	Question	countQIDs
	<chr>	<int>
1	About how long has it been since you last visited a doctor for a r~	1
2	Adults aged 18-64 who have any kind of health care coverage (varia~	3
3	Adults aged 45-75 who have had a blood stool test within the past ~	1
4	Adults aged 50+ who have ever had a sigmoidoscopy or colonoscopy (~	1
5	Adults aged 50+ who have had a blood stool test within the past tw~	1
6	Adults aged 50-75 who have had a blood stool test within the past ~	2
7	Adults aged 65+ who have ever had a pneumonia vaccination (variabl~	2
8	Adults aged 65+ who have had a flu shot within the past year (vari~	3
9	Adults aged 65+ who have had all their natural teeth extracted (va~	2
10	Adults that have had any permanent teeth extracted (variable calculu~	2
11	Adults who are current e-cigarette users (variable calculated from~	1
12	Adults who are current e-cigarette users (variable calculated from~	1

```

13 Adults who are current smokers (variable calculated from one or mo~ 1
14 Adults who are limited in any activities because of physical, ment~ 1
15 Adults who had some form of health insurance (variable calculated ~ 2
# i 84 more rows

```

Why is there a need for multiple values of `QuestionID` for the same question? Let's investigate this further for the `Question` matching "(Heavy drinkers).*(male).*(14 drinks)". Here is the count of each income breakout in the dataset for this question.

```

# A tibble: 12 x 3
# Groups:   BreakoutID [12]
  BreakoutID Break_Out      countBreakIDs
  <chr>       <chr>           <int>
1 INCOME01    Less than $15,000     6
2 INCOME02    $15,000-$24,999     6
3 INCOME03    $25,000-$34,999     6
4 INCOME04    $35,000-$49,999     6
5 INCOME05    $50,000-$99,999     6
6 INCOME06    $100,000-$199,999    6
7 INCOME07    $200,000+          6
8 INCOME1     Less than $15,000     12
9 INCOME2     $15,000-$24,999     12
10 INCOME3    $25,000-$34,999     12
11 INCOME4    $35,000-$49,999     12
12 INCOME5    $50,000+          12

```

A new income-level breakout was introduced in the last 3 years of the dataset, providing 7 instead of 5 categories. As a result of the new division, a new value of `QuestionID` was assigned. Care must be taken when combining results when a question has multiple values of `QuestionID`.

Project Goal

In this effort you are to produce a dashboard interface to the BRFSS dataset, allowing the user to select the survey question and the dashboard displays the various breakouts of the survey results for the question. Each subpanel of the dashboard should subset the results by only one criteria (age, for example). There should be a subpanel devoted to choosing the question. In addition, for each possible response, a:

- **confidence interval** overall in an **Overall** subpanel
- **confidence interval** versus gender in a **By_Gender** subpanel

- confidence interval versus age in a **By_Age_Group** subpanel
- confidence interval versus state/territory in a **By_Location** subpanel
- confidence interval versus level of education in a **By_Education** subpanel
- confidence interval versus income level in a **By_Income** subpanel
- confidence interval versus year in a **Temporal** subpanel

Grading metrics and design considerations

Function over fashion. The dashboard does not need to be fancy, and it must function accurately and repeatedly. Emphasize function, not fashion, in your work product. One more time: wrangle Dash Plotly or RShiny last, or not at all, but wrangle the workflow accuracy. Lastly, a pretty and inaccurate dashboard will receive a pretty low score. Most of the credit for this project will be given to accurate aggregation of the data, not the display of the aggregation. Do not overtweak the user interface! Below are some suggestions to consider when deploying your user interface.

Question selection. There are 100 questions, so the selection of a question should be layered. Consider a top-down layering of Class/Topic/Question, for example. Do not present the user with keys, only the value of the keys.

Subpanel Visualization. Please consider this important problem carefully. Subset the results *only* by one variable in each subpanel. For example, subset by year only in one subpanel, geography in another subpanel, age in a third subpanel, etc. This means that all other breakouts must be aggregated! More on this later.

State/territory subpanel. There are more than 50 regions to consider, so use a compact way to represent the geographical results, such as a map, or aggregating states into national regions, or showing highest, lowest and average geographical regions.

User adaptable subpanel complexity. Consider having, for each subpanel, a “more” feature and a “less” feature. This enables the user to adjust the granularity of each subpanel. **And**, provide only a few (5?) levels of granularity for each subpanel. Otherwise, you risk underwhelming or overwhelming your audience.

Proposed workflow

- Consider a parquet-formatted dataset, based on observed speedup of information retrieval.
- Design the question selection tool.
- Design the aggregation workflow for each subpanel, beginning with the Overall subpanel.
- Design the Overall subpanel without “more” or “less” options.

- Augment the Overall subpanel with “more” or “less” options.
- Design the other subpanels without “more” or “less” options.
- Modify these subpanels to include “more” or “less” options.

Aggregation Strategy for “Overall” Breakout

Most of the questions have a breakout group called “Overall”. As the name suggests, this is the broadest aggregation so far in the dataset. However, percentages labeled “Overall” are still per-territory, per-year summary metrics and not true overall results! Here, we consider a method to aggregate and produce confidence limits for response percentages.

Note that the sample size is provided in column `Sample_Size` along with a percentage estimate in column `Data_value`. The percentage estimate and the sample size allow you to translate per-breakout percentage estimates to per-breakout person estimates prior to aggregation. Upon aggregation, the aggregate sample size can be used to determine an aggregate percentage confidence interval. If the possible responses to a questions do not vary over time or territory, then aggregation is straightforward. Let’s consider that case first.

The chunk below illustrates how one could aggregate the per-territory per-year “Overall” value for `Break_Out` (“BO1” value for `BreakOutID`). However, if we “Run Cell” repeatedly, we see some problems with our aggregation. Try it!

```
save_rows <- c(2150434,830046,131103)
qidx<- sample(save_rows,size=1)
df |>
  select(Question,Break_Out,Response,ResponseID,
  Sample_Size,Data_value) |>
  filter(Question==Question[qidx]) |>
  filter(Break_Out=="Overall") |>
  na.omit() |>
  rename(persons=Sample_Size) |>
  mutate(Sample_Size=persons*100/Data_value) |>
  group_by(ResponseID,Response) |>
  summarize(agg_ss=sum(Sample_Size),
  agg_persons=sum(persons),
  agg_percent=agg_persons*100/agg_ss,
  agg_percent_sdev=sqrt(agg_percent*(100-agg_percent)/agg_ss),
  agg_low_ci_limit=agg_percent - 2*agg_percent_sdev,
  agg_high_ci_limit=agg_percent + 2*agg_percent_sdev) |>
  print(n=20)
```

```

# A tibble: 11 x 8
# Groups:   ResponseID [11]
  ResponseID Response      agg_ss agg_persons agg_percent agg_percent_sdev
  <chr>       <chr>        <dbl>     <dbl>       <dbl>            <dbl>
1 RESP025    Employd     829310.    403874     48.7           0.0549
2 RESP026    Self employd 952084.    79307      8.33           0.0283
3 RESP027    No work< yr 4345159.   137203     3.16           0.00839
4 RESP028    No work >yr  5004899.   138615     2.77           0.00734
5 RESP029    Homemkr     1025198.   67172      6.55           0.0244
6 RESP030    Student      2854783.   150163     5.26           0.0132
7 RESP031    Retired      9426205.   1750896    18.6            0.0127
8 RESP032    Unable to work 6332814.  411954     6.51           0.00980
9 RESP137    Employed     4116750.   2015613    49.0            0.0246
10 RESP141   Homemaker    4500671.   251868     5.60           0.0108
11 RESP172   Self-employed 4674301.   424403     9.08           0.0133
# i 2 more variables: agg_low_ci_limit <dbl>, agg_high_ci_limit <dbl>

```

By a few trials of the above chunk, it may be seen that the response options changed over the course of this survey for some questions. Often, the changes are trivial, for example, “Employd” vs. “Employed”, or “Homemkr” vs “Homemaker”. In other cases, the response options have different refinement, for example, “\$50,000+”, vs the finer options “\$50,000-99,999”, “\$100,000-199,999”, and “\$200,000+”. In all these cases, note that the value of `ResponseID` changes when the response option changes.

In order to aggregate properly, we need to merge these responses. Without this merge, “Homemkr” and “Homemaker” responses will not be pooled. Fortunately, this may be done by a simple mapping of every value of `ResponseID` to a new one, reflecting the merge. I have drafted merge rules for `responseID` values in the function below. Please start with that function and modify it as necessary.

```

merge_ResponseID <- function(char_vec) {
  char_vec <- str_replace(char_vec, "RESP025", "RESP137")
  char_vec <- str_replace(char_vec, "RESP026", "RESP172")
  char_vec <- str_replace(char_vec, "RESP029", "RESP141")
  char_vec <- str_replace(char_vec, "RESP230", "RESP020")
  char_vec <- str_replace(char_vec, "RESP231", "RESP020")
  char_vec <- str_replace(char_vec, "RESP232", "RESP020")
  char_vec <- str_replace(char_vec, "RESP196", "RESP199")
  char_vec <- str_replace(char_vec, "RESP197", "RESP199")
  char_vec <- str_replace(char_vec, "RESP198", "RESP199")
  char_vec <- str_replace(char_vec, "RESP199", "RESP199")
  char_vec <- str_replace(char_vec, "RESP200", "RESP008")
  char_vec <- str_replace(char_vec, "RESP194", "RESP005")

```

```

    char_vec <- str_replace(char_vec,"RESP195","RESP006")
    return(char_vec)
}

```

After the `ResponseID` column is merged, the following function changes the `Response` column. Both functions should be updated in tandem.

```

merge_Response <- function(ResponseID,Response) {
  idx <- str_detect(ResponseID,"RESP137")
  Response[idx] <- "Employed"
  idx <- str_detect(ResponseID,"RESP172")
  Response[idx] <- "Self-employed"
  idx <- str_detect(ResponseID,"RESP141")
  Response[idx] <- "Homemaker"
  idx <- str_detect(ResponseID,"RESP020")
  Response[idx] <- "$50,000+"
  idx <- str_detect(ResponseID,"RESP199")
  Response[idx] <- "A/A Native, Asian,Other"
  idx <- str_detect(ResponseID,"RESP008")
  Response[idx] <- "Multiracial"
  idx <- str_detect(ResponseID,"RESP005")
  Response[idx] <- "White"
  idx <- str_detect(ResponseID,"RESP006")
  Response[idx] <- "Black"
  # make all responses lower case
  Response <- tolower(Response)
  return(Response)
}

```

One of your tasks will be to map each one of these values of `ResponseID` to a (possible) new one, reflecting how the responses are merged, starting with the draft functions above. There will only be a small number of changes in this process, as most of the response options do not need to be merged.

Here the draft functions are applied to the same questions. Are there improvements?

```

  save_rows <- c(2150434,830046,131103)
  qidx<- sample(save_rows,size=1)
  lDf <- df |>
    select(Question,Break_Out,Response,ResponseID,
    Sample_Size,Data_value) |>
    filter(Question==Question[qidx]) |>

```

```

filter(Break_Out=="Overall")

lDf$ResponseID <- merge_ResponseID(lDf$ResponseID)
lDf$Response <- merge_Response(unlist(lDf$ResponseID),
unlist(lDf$Response))

lDf |>
na.omit() |>
rename(persons=Sample_Size) |>
mutate(Sample_Size=persons*100/Data_value) |>
group_by(ResponseID,Response) |>
summarize(agg_ss=sum(Sample_Size),
agg_persons=sum(persons),
agg_percent=agg_persons*100/agg_ss,
agg_percent_sdev=sqrt(agg_percent*(100-agg_percent)/agg_ss),
agg_low_ci_limit=agg_percent - 2*agg_percent_sdev,
agg_high_ci_limit=agg_percent + 2*agg_percent_sdev) |>
print(n=20)

```

```

# A tibble: 6 x 8
# Groups:   ResponseID [6]
  ResponseID Response      agg_ss agg_persons agg_percent agg_percent_sdev
  <chr>      <chr>       <dbl>     <dbl>        <dbl>          <dbl>
1 RESP010    18-24 years  2646706.    333789       12.6         0.0204
2 RESP011    25-34 years  3556121.    607216       17.1         0.0200
3 RESP012    35-44 years  4456831.    727847       16.3         0.0175
4 RESP013    45-54 years  5686151.    945513       16.6         0.0156
5 RESP014    55-64 years  7465206.    1236977      16.6         0.0136
6 RESP015    65+ years   9930209.    2048338      20.6         0.0128
# i 2 more variables: agg_low_ci_limit <dbl>, agg_high_ci_limit <dbl>

```

General Aggregation Approach

To aggregate “Overall” breakouts, it is important to merge the response options, as shown above. Aggregating of “Overall” breakouts is relatively straightforward, however, because there can be no change in the **breakout categories** over year or territory. “Overall” doesn’t change throughout this survey. This is not the case for other breakout categories, however. Let’s see an example of a question, whose responses are merged, below. Note the difference in the number of breakouts in the “Household Income” breakout category.

```

#my_Question <- "What is your race/ethnicity?"
#my_Question <- df$Question[sample(dim(df)[1],1)]
my_Question <- "Ever told you that you have a form of depression?"
lDf <- df |>
  filter(Question==my_Question) |>
  select(Question, QuestionID, Response,
  ResponseID, Break_Out, Break_Out_Category)

lDf$ResponseID <- merge_ResponseID(lDf$ResponseID)
lDf$Response <- merge_Response(unlist(lDf$ResponseID),
  unlist(lDf$Response))

lDf |>
  filter(Response=="yes") |>
  select(Question, QuestionID, Response,
  Break_Out, Break_Out_Category) |>
  distinct() |>
  group_by(Question, QuestionID, Break_Out_Category) |>
  summarize(countBreakOuts=n()) |>
  arrange(Break_Out_Category)

```

```

# A tibble: 12 x 4
# Groups:   Question, QuestionID [2]
  Question           QuestionID Break_Out_Category countBreakOuts
  <chr>              <chr>     <chr>                  <int>
1 Ever told you that you have a f~ ADDEPEV2    Age Group            6
2 Ever told you that you have a f~ ADDEPEV3    Age Group            6
3 Ever told you that you have a f~ ADDEPEV2    Education Attained      4
4 Ever told you that you have a f~ ADDEPEV3    Education Attained      4
5 Ever told you that you have a f~ ADDEPEV2    Household Income        5
6 Ever told you that you have a f~ ADDEPEV3    Household Income        8
7 Ever told you that you have a f~ ADDEPEV2    Overall                 1
8 Ever told you that you have a f~ ADDEPEV3    Overall                 1
9 Ever told you that you have a f~ ADDEPEV2    Race/Ethnicity          8
10 Ever told you that you have a f~ ADDEPEV3   Race/Ethnicity          8
11 Ever told you that you have a f~ ADDEPEV2    Sex                      2
12 Ever told you that you have a f~ ADDEPEV3   Sex                      2

```

To get started, here is a list of the breakouts, with BreakoutID and BreakOutCategoryID included.

```
# A tibble: 58 x 4
```

# Groups: BreakOutCategoryID, BreakoutID [58]				
	BreakOutCategoryID	BreakoutID	Break_Out	count
	<chr>	<chr>	<chr>	<int>
1	CAT1	B01	Overall	116860
2	CAT2	SEX1	Male	110490
3	CAT2	SEX2	Female	111335
4	CAT3	AGE01	18-24	99515
5	CAT3	AGE02	25-34	99773
6	CAT3	AGE03	35-44	99848
7	CAT3	AGE04	45-54	100186
8	CAT3	AGE05	55-64	100164
9	CAT3	AGE06	40-49	1172
10	CAT3	AGE07	50-59	4262
11	CAT3	AGE08	60-64	2020
12	CAT3	AGE09	65+	100435
13	CAT3	AGE10	65-74	3396
14	CAT3	AGE11	75+	3396
15	CAT3	AGE12	21-30	304
16	CAT3	AGE13	31-40	304
17	CAT3	AGE14	41-50	304
18	CAT3	AGE15	51-60	304
19	CAT3	AGE16	61-65	304
20	CAT3	AGE17	60-69	2242
21	CAT3	AGE18	70-74	534
22	CAT3	AGE19	70-75	1600
23	CAT3	AGE20	21-25	106
24	CAT3	AGE21	26-35	106
25	CAT3	AGE22	36-45	106
26	CAT3	AGE23	46-55	106
27	CAT3	AGE24	56-65	106
28	CAT3	AGE25	65-75	324
29	CAT3	AGE26	70-80	108
30	CAT4	RACE01	White, non-Hispanic	77207
31	CAT4	RACE02	Black, non-Hispanic	77207
32	CAT4	RACE03	American Indian or Alaskan Native, non-~	77207
33	CAT4	RACE04	Asian, non-Hispanic	77207
34	CAT4	RACE05	Native Hawaiian or other Pacific Island~	77207
35	CAT4	RACE06	Other, non-Hispanic	77207
36	CAT4	RACE07	Multiracial, non-Hispanic	77207
37	CAT4	RACE08	Hispanic	77207
38	CAT4	RACE1	White, non-Hispanic	30504
39	CAT4	RACE2	Black, non-Hispanic	29950
40	CAT4	RACE3	Hispanic	30366

41	CAT4	RACE4	Other, non-Hispanic	30364
42	CAT4	RACE5	Multiracial, non-Hispanic	30028
43	CAT5	EDUCA1	Less than H.S.	109834
44	CAT5	EDUCA2	H.S. or G.E.D.	109850
45	CAT5	EDUCA3	Some post-H.S.	109848
46	CAT5	EDUCA4	College graduate	109852
47	CAT6	INCOME01	Less than \$15,000	25494
48	CAT6	INCOME02	\$15,000-\$24,999	25494
49	CAT6	INCOME03	\$25,000-\$34,999	25494
50	CAT6	INCOME04	\$35,000-\$49,999	25494
51	CAT6	INCOME05	\$50,000-\$99,999	25494
52	CAT6	INCOME06	\$100,000-\$199,999	25494
53	CAT6	INCOME07	\$200,000+	25494
54	CAT6	INCOME1	Less than \$15,000	83342
55	CAT6	INCOME2	\$15,000-\$24,999	83346
56	CAT6	INCOME3	\$25,000-\$34,999	83329
57	CAT6	INCOME4	\$35,000-\$49,999	83323
58	CAT6	INCOME5	\$50,000+	83342

Let's begin with "CAT6", or income breakout. Clearly, "INCOME05", "INCOME06", and "INCOME07" are a refinement of "INCOME5". For "CAT4", or race, we also see a refinement, ("RACE03", "RACE04", "RACE05", and "RACE06" being a refinement of "RACE4"). Finally, for "CAT3", or age, we see two breakout groups (AGE01 through AGE11, and AGE12 through AGE26). Some of these values of `BreakoutID` have full or partial overlap, while others reflect a refinement.

One possible approach to merging the breakout categories mimics that for response options: first change the `BreakoutID` values, then change the `Break_Out` values. The following are draft functions which attempt this.

```
merge_BreakoutID <- function(char_vec) {
  char_vec <- str_replace(char_vec, "INCOME01", "INCOME1")
  char_vec <- str_replace(char_vec, "INCOME02", "INCOME2")
  char_vec <- str_replace(char_vec, "INCOME03", "INCOME3")
  char_vec <- str_replace(char_vec, "INCOME04", "INCOME4")
  char_vec <- str_replace(char_vec, "INCOME05", "INCOME5")
  char_vec <- str_replace(char_vec, "INCOME06", "INCOME5")
  char_vec <- str_replace(char_vec, "INCOME07", "INCOME5")
  char_vec <- str_replace(char_vec, "RACE01", "RACE1")
  char_vec <- str_replace(char_vec, "RACE02", "RACE2")
  char_vec <- str_replace(char_vec, "RACE08", "RACE3")
  char_vec <- str_replace(char_vec, "RACE04", "RACE4")
  char_vec <- str_replace(char_vec, "RACE05", "RACE4")
```

```

char_vec <- str_replace(char_vec,"RACE06","RACE4")
char_vec <- str_replace(char_vec,"RACE03","RACE4")
char_vec <- str_replace(char_vec,"RACE07","RACE5")
return(char_vec)
}

```

```

merge_Break_Out <- function(BreakoutID,Break_Out) {
  idx <- str_detect(BreakoutID,"INCOME5")
  Break_Out[idx] <- "$50,000+"
  idx <- str_detect(BreakoutID,"RACE1")
  Break_Out[idx] <- "White"
  idx <- str_detect(BreakoutID,"RACE2")
  Break_Out[idx] <- "Black"
  idx <- str_detect(BreakoutID,"RACE3")
  Break_Out[idx] <- "Hispanic"
  idx <- str_detect(BreakoutID,"RACE4")
  Break_Out[idx] <- "A/A Native, Asian,Other"
  idx <- str_detect(BreakoutID,"RACE5")
  Break_Out[idx] <- "Multiracial"
  return(Break_Out)
}

```

Let's apply these functions to merge values in `BreakoutID` and `Break_Out` for the same question.

```

#my_Question <- "What is your race/ethnicity?"
#my_Question <- df$Question[sample(dim(df)[1],1)]
my_Question <- "Ever told you that you have a form of depression?"
lDf <- df |>
  filter(Question==my_Question) |>
  select(Question,QuestionID,Response,
  ResponseID,Break_Out, BreakoutID,Break_Out_Category)

lDf$ResponseID <- merge_ResponseID(lDf$ResponseID)
  lDf$Response <- merge_Response(unlist(lDf$ResponseID),
  unlist(lDf$Response))
lDf$BreakoutID <- merge_BreakoutID(lDf$BreakoutID)
  lDf$Break_Out <- merge_Break_Out(unlist(lDf$BreakoutID),
  unlist(lDf$Break_Out))

lDf |>
  filter(Response=="yes") |>

```

```

  select(Question,QuestionID,Response,
Break_Out, Break_Out_Category) |>
  distinct() |>
  group_by(Question,QuestionID,Break_Out_Category) |>
  summarize(countBreakOuts=n()) |>
  arrange(Break_Out_Category)

```

```

# A tibble: 12 x 4
# Groups:   Question, QuestionID [2]
  Question           QuestionID Break_Out_Category countBreakOuts
  <chr>              <chr>     <chr>                <int>
1 Ever told you that you have a f~ ADDEPEV2    Age Group            6
2 Ever told you that you have a f~ ADDEPEV3    Age Group            6
3 Ever told you that you have a f~ ADDEPEV2    Education Attained    4
4 Ever told you that you have a f~ ADDEPEV3    Education Attained    4
5 Ever told you that you have a f~ ADDEPEV2    Household Income       5
6 Ever told you that you have a f~ ADDEPEV3    Household Income       5
7 Ever told you that you have a f~ ADDEPEV2    Overall                 1
8 Ever told you that you have a f~ ADDEPEV3    Overall                 1
9 Ever told you that you have a f~ ADDEPEV2    Race/Ethnicity          5
10 Ever told you that you have a f~ ADDEPEV3   Race/Ethnicity          5
11 Ever told you that you have a f~ ADDEPEV2    Sex                      2
12 Ever told you that you have a f~ ADDEPEV3   Sex                      2

```

As with response options, a complete aggregation will require that a mapping of BreakoutID values, reflected above, to a merged BreakoutID. **One of your responsibilities is to merge values of BreakoutID, using your best judgement.** The draft functions above serve as starting points. Such a mapping will allow aggregation of responses to the same questions across different versions of breakout category.

Pre-Beta Prototype Dashboard

In this section we consider a workflow, from choosing a question, to summarizing the results.

```

# prototype_workflow_choose_question
# interactive ONLY - prompt user for class
#option <- unique(layerQ$Class)
#idx <- utils::menu(option,title="Which class?")
#my_class <- option[idx]
#slayerQ <- layerQ |>

```

```

#     filter(Class==my_class)
## prompt user for topic
#option <- unique(slayerQ$Topic)
#idx <- utils::menu(option,title="Which topic?")
#my_topic <- option[idx]
#vslayerQ <- slayerQ |>
#     filter(Topic==my_topic)
## prompt user for question
#option <- unique(vslayerQ$Question)
#idx <- utils::menu(option,title="Which question?")
#my_q <- option[idx]
# from RDS file - question stored there
my_q <- readRDS("my_q.RDS")

qDf <- df |>
  filter(Question==my_q) |>
  filter(!(Locationabbr == "US")) |>
  filter(!(Locationabbr == "UW"))

# merges
qDf$ResponseID <- merge_ResponseID(qDf$ResponseID)
  qDf$Response <- merge_Response(unlist(qDf$ResponseID),
  unlist(qDf$Response))
qDf$BreakoutID <- merge_BreakoutID(qDf$BreakoutID)
  qDf$Break_Out <- merge_Break_Out(unlist(qDf$BreakoutID),
  unlist(qDf$Break_Out))

# comment out if next chunk is used
(my_q <- str_extract(my_q,"[^()]+"))

```

[1] "Days when physical health status not good "

```

plotDf <- qDf |>
  filter(BreakOutCategoryID=="CAT1") |>
  select(Response,Sample_Size) |>
  na.omit() |>
  rename(persons=Sample_Size) |>
  group_by(Response) |>
  summarize(agg_persons=sum(persons))

plotDf <- plotDf |>
  mutate(agg_ss= sum(agg_persons)) |>

```

```

    mutate(agg_percent=agg_persons*100/agg_ss) |>
    mutate( agg_percent_sdev=sqrt(
      agg_percent*(100-agg_percent)/agg_ss)) |>
    mutate(agg_low_ci_limit=
      agg_percent - 2*agg_percent_sdev) |>
    mutate(agg_high_ci_limit=
      agg_percent + 2*agg_percent_sdev) |>
    select(-c(agg_persons,agg_ss,agg_percent_sdev))
plotDf |>
  select(-agg_percent) |>
  print(n=20)

```

```

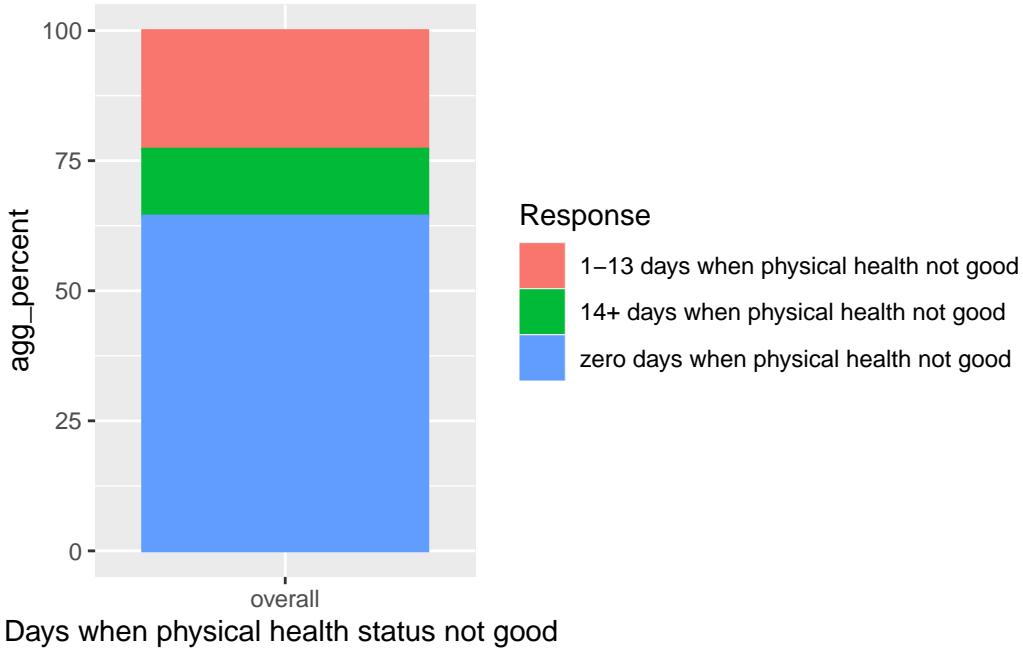
# A tibble: 3 x 3
Response          agg_low_ci_limit agg_high_ci_limit
<chr>                <dbl>             <dbl>
1 1-13 days when physical health not good     22.7            22.8
2 14+ days when physical health not good      12.8            12.9
3 zero days when physical health not good      64.3            64.4

```

```

plotDf |>
  ggplot( aes(x="overall",y=agg_percent,
  fill=Response,color=Response,position="fill")) +
  geom_col() + xlab(my_q)

```



```

## summary by overall
#(my_q <- str_extract(my_q, "[^()]+"))
#plotDf <- qDf |>
#  filter(BreakOutCategoryID=="CAT1") |>
#  select(Response, Sample_Size, Data_value) |>
#  na.omit() |>
#  rename(persons=Sample_Size) |>
#  mutate(Sample_Size=persons/Data_value*100) |>
##ssSum <- plotDf |>
##  summarize(sum(persons))
##ssSum <- unlist(ssSum)
##numResponseOptions <- length(unique(plotDf$Response))
##ssVec <- rep(ssSum, each=numResponseOptions)
##plotDf |>
#  group_by(Response) |>
#  summarize(agg_ss=sum(Sample_Size),
#  agg_persons=sum(persons),
#  agg_percent=agg_persons*100/agg_ss,
#  agg_percent_sdev=sqrt(agg_percent*(100-agg_percent)/agg_ss),
#  agg_low_ci_limit=agg_percent - 2*agg_percent_sdev,
#  agg_high_ci_limit=agg_percent + 2*agg_percent_sdev) |>
#  select(-c(agg_persons,agg_percent_sdev))
#

```

```

#   plotDf |>
#     select(-agg_percent) |>
#     print(n=20)
#   plotDf |>
#     ggplot( aes(x="overall",y=agg_percent,
#     fill=Response,color=Response,position="fill")) +
#     geom_col() + xlab(my_q)

# comment out if next chunk is used
# summary by territory
(my_q)

```

[1] "Days when physical health status not good "

```

plotDf <- qDf |>
  filter(BreakOutCategoryID=="CAT1") |>
  select(Locationabbr,Response,Sample_Size) |>
  na.omit() |>
  rename(persons=Sample_Size) |>
  group_by(Locationabbr) |>
  summarize(Response=Response,
  persons=persons,
  agg_ss=sum(persons))

```

Warning: Returning more (or less) than 1 row per `summarise()` group was deprecated in dplyr 1.1.0.
 i Please use `reframe()` instead.
 i When switching from `summarise()` to `reframe()`, remember that `reframe()` always returns an ungrouped data frame and adjust accordingly.

```

plotDf <- plotDf |>
  group_by(Locationabbr,Response) |>
  summarize(agg_ss=agg_ss,
  agg_persons=sum(persons),
  agg_percent=agg_persons*100/agg_ss,
  agg_percent_sdev=
  sqrt(agg_percent*(100-agg_percent)/agg_ss),
  agg_low_ci_limit=
  agg_percent - 2*agg_percent_sdev,
  agg_high_ci_limit=

```

```

agg_percent + 2*agg_percent_sdev) |>
distinct() |>
select(-c(agg_persons,agg_percent_sdev,agg_ss))

```

Warning: Returning more (or less) than 1 row per `summarise()` group was deprecated in dplyr 1.1.0.
 i Please use `reframe()` instead.
 i When switching from `summarise()` to `reframe()`, remember that `reframe()` always returns an ungrouped data frame and adjust accordingly.

```

plotDf |>
  select(-agg_percent) |>
  print(n=20)

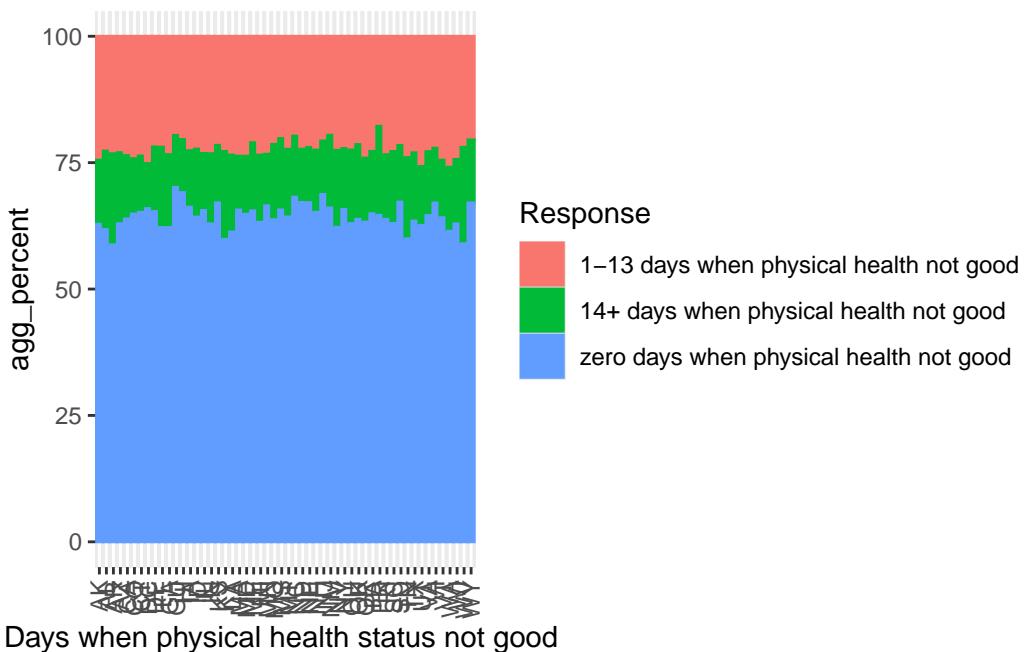
# A tibble: 162 x 4
# Groups:   Locationabbr, Response [162]
  Locationabbr Response      agg_low_ci_limit agg_high_ci_limit
  <chr>        <chr>            <dbl>             <dbl>
1 AK          1-13 days when physical heal~    23.9              25.1
2 AK          14+ days when physical healt~    12.2              13.1
3 AK          zero days when physical heal~    62.2              63.5
4 AL          1-13 days when physical heal~    22.2              23.2
5 AL          14+ days when physical healt~    15.1              16.0
6 AL          zero days when physical heal~    61.2              62.4
7 AR          1-13 days when physical heal~    22.7              23.8
8 AR          14+ days when physical healt~    17.6              18.5
9 AR          zero days when physical heal~    58.1              59.3
10 AZ         1-13 days when physical heal~    22.6              23.4
11 AZ         14+ days when physical healt~    13.7              14.4
12 AZ         zero days when physical heal~    62.5              63.4
13 CA         1-13 days when physical heal~    23.2              24.0
14 CA         14+ days when physical healt~    12.2              12.8
15 CA         zero days when physical heal~    63.4              64.3
16 CO         1-13 days when physical heal~    23.8              24.6
17 CO         14+ days when physical healt~    10.7              11.2
18 CO         zero days when physical heal~    64.4              65.3
19 CT         1-13 days when physical heal~    23.3              24.1
20 CT         14+ days when physical healt~    10.8              11.4
# i 142 more rows

```

```

plotDf |>
  ggplot( aes(x=Locationabbr,
y=agg_percent,
fill=Response,color=Response,position="fill")) +
  geom_col() + xlab(my_q) +
  theme(axis.text.x = element_text(angle = 90, vjust = 0.5, hjust=1))

```



```

## summary by territory
#(my_q)
#plotDf <- qDf |>
#  filter(BreakOutCategoryID=="CAT1") |>
#  select(Locationabbr,Response,Sample_Size,Data_value) |>
#  na.omit() |>
#  rename(persons=Sample_Size) |>
#  mutate(Sample_Size=persons*100/Data_value) |>
#  group_by(Locationabbr,Response) |>
#  summarize(agg_ss=sum(Sample_Size),
#  agg_persons=sum(persons),
#  agg_percent=agg_persons*100/agg_ss,
#  agg_percent_sdev=sqrt(agg_percent*(100-agg_percent)/agg_ss),
#  agg_low_ci_limit=agg_percent - 2*agg_percent_sdev,
#  agg_high_ci_limit=agg_percent + 2*agg_percent_sdev) |>

```

```

#      select(-c(agg_ss,agg_persons,agg_percent_sdev))
# plotDf |>
#      select(-agg_percent) |>
#      print(n=10)
# plotDf |>
#      ggplot( aes(x=Locationabbr,
#                  y=agg_percent,
#                  fill=Response,color=Response,position="fill")) +
#                  geom_col() + xlab(my_q) +
#                  theme(axis.text.x = element_text(angle = 90, vjust = 0.5, hjust=1))
#(my_q)

```

```

# comment out if next chunk is used
# summary by year
(my_q)

```

[1] "Days when physical health status not good "

```

plotDf <- qDf |>
  filter(BreakOutCategoryID=="CAT1") |>
  select(Year,Response,Sample_Size) |>
  na.omit() |>
  rename(persons=Sample_Size) |>
  group_by(Year) |>
  summarize(Response=Response,
            persons=persons,
            agg_ss=sum(persons))

```

Warning: Returning more (or less) than 1 row per `summarise()` group was deprecated in dplyr 1.1.0.

i Please use `reframe()` instead.
i When switching from `summarise()` to `reframe()`, remember that `reframe()` always returns an ungrouped data frame and adjust accordingly.

```

plotDf <- plotDf |>
  group_by(Year,Response) |>
  summarize(agg_ss=agg_ss,
            agg_persons=sum(persons),
            agg_percent=agg_persons*100/agg_ss,
            agg_percent_sdev=

```

```

sqrt(agg_percent*(100-agg_percent)/agg_ss),
agg_low_ci_limit=
agg_percent - 2*agg_percent_sdev,
agg_high_ci_limit=
agg_percent + 2*agg_percent_sdev) |>
distinct() |>
select(-c(agg_persons,agg_percent_sdev,agg_ss))

```

Warning: Returning more (or less) than 1 row per `summarise()` group was deprecated in dplyr 1.1.0.

- i Please use `reframe()` instead.
- i When switching from `summarise()` to `reframe()`, remember that `reframe()` always returns an ungrouped data frame and adjust accordingly.

```

plotDf |>
  select(-agg_percent) |>
  print(n=100)

```

	Year Response	agg_low_ci_limit	agg_high_ci_limit
	<dbl> <chr>	<dbl>	<dbl>
1	2019 1-13 days when physical health not ~	23.8	24.0
2	2019 14+ days when physical health not g~	14.1	14.3
3	2019 zero days when physical health not ~	61.8	62.1
4	2020 1-13 days when physical health not ~	18.1	18.3
5	2020 14+ days when physical health not g~	10.8	11.0
6	2020 zero days when physical health not ~	70.7	71.0
7	2021 1-13 days when physical health not ~	20.9	21.2
8	2021 14+ days when physical health not g~	11.8	12.0
9	2021 zero days when physical health not ~	66.9	67.2
10	2022 1-13 days when physical health not ~	24.8	25.1
11	2022 14+ days when physical health not g~	13.3	13.5
12	2022 zero days when physical health not ~	61.5	61.8
13	2023 1-13 days when physical health not ~	25.3	25.6
14	2023 14+ days when physical health not g~	13.7	13.9
15	2023 zero days when physical health not ~	60.6	60.9

```

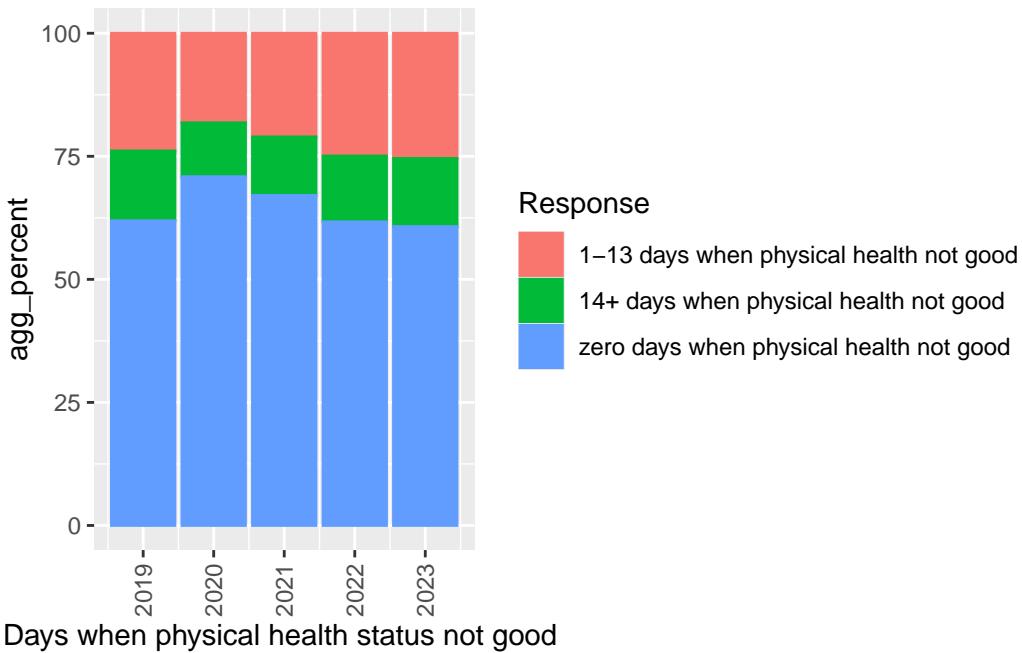
plotDf |>
  ggplot( aes(x=Year,

```

```

y=agg_percent,
fill=Response,color=Response,position="fill")) +
  geom_col() + xlab(my_q) +
  theme(axis.text.x = element_text(angle = 90, vjust = 0.5, hjust=1))

```



```

##summary by year
#(my_q)
#plotDf <- qDf |>
#  filter(BreakOutCategoryID=="CAT1") |>
#  select(Year,Response,Sample_Size,Data_value) |>
#  na.omit() |>
#  rename(persons=Sample_Size) |>
#  mutate(Sample_Size=persons*100/Data_value) |>
#  group_by(Year,Response) |>
#  summarize(agg_ss=sum(Sample_Size),
#            agg_persons=sum(persons),
#            agg_percent=agg_persons*100/agg_ss,
#            agg_percent_sdev=sqrt(agg_percent*(100-agg_percent)/agg_ss),
#            agg_low_ci_limit=agg_percent - 2*agg_percent_sdev,
#            agg_high_ci_limit=agg_percent + 2*agg_percent_sdev) |>
#            select(-c(agg_ss,agg_persons,agg_percent_sdev))
#  plotDf |>

```

```

#       select(-agg_percent) |>
#       print(n=20)
#   plotDf |>
#       ggplot( aes(x=Year,y=agg_percent,
#       fill=Response,color=Response,position="fill")) +
#           geom_col() + xlab(my_q)

# comment out if next chunk is used
# summary by gender
(my_q)

```

[1] "Days when physical health status not good "

```

plotDf <- qDf |>
  filter(BreakOutCategoryID=="CAT2") |>
  select(Break_Out,Response,Sample_Size) |>
  na.omit() |>
  rename(persons=Sample_Size) |>
  group_by(Break_Out) |>
  summarize(Response=Response,
  persons=persons,
  agg_ss=sum(persons))

```

Warning: Returning more (or less) than 1 row per `summarise()` group was deprecated in dplyr 1.1.0.

- i Please use `reframe()` instead.
- i When switching from `summarise()` to `reframe()`, remember that `reframe()` always returns an ungrouped data frame and adjust accordingly.

```

plotDf <- plotDf |>
  group_by(Break_Out,Response) |>
  summarize(agg_ss=agg_ss,
  agg_persons=sum(persons),
  agg_percent=agg_persons*100/agg_ss,
  agg_percent_sdev=
    sqrt(agg_percent*(100-agg_percent)/agg_ss),
  agg_low_ci_limit=
    agg_percent - 2*agg_percent_sdev,
  agg_high_ci_limit=
    agg_percent + 2*agg_percent_sdev) |>

```

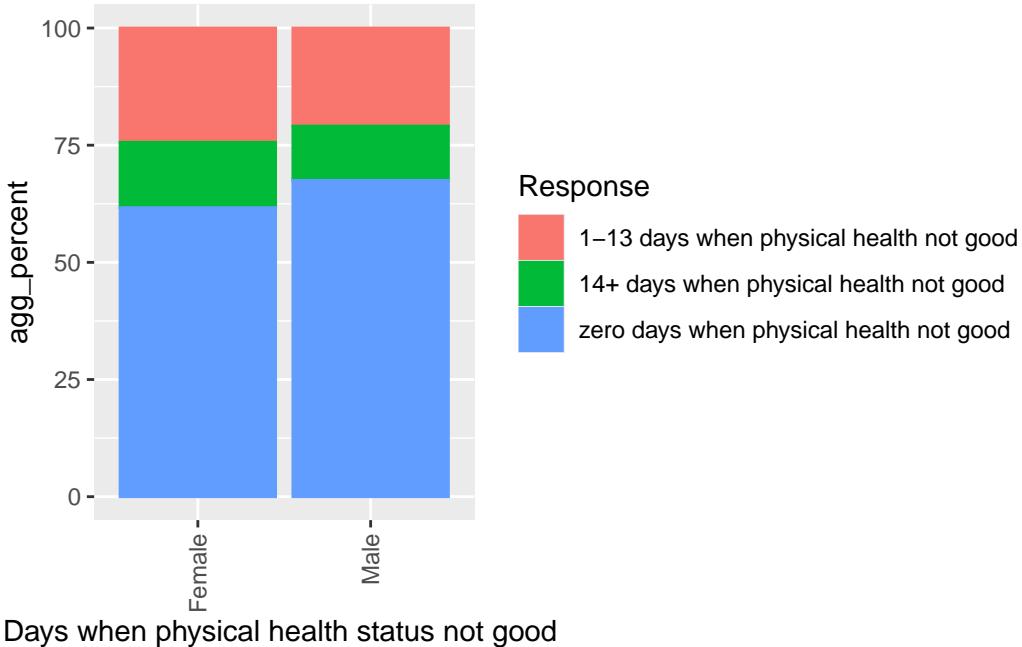
```
distinct() |>  
select(-c(agg_persons,agg_percent_sdev,agg_ss))
```

Warning: Returning more (or less) than 1 row per `summarise()` group was deprecated in dplyr 1.1.0.
i Please use `reframe()` instead.
i When switching from `summarise()` to `reframe()`, remember that `reframe()` always returns an ungrouped data frame and adjust accordingly.

```
plotDf |>  
  select(-agg_percent) |>  
  print(n=100)
```

```
# A tibble: 6 x 4  
# Groups: Break_Out, Response [6]  
Break_Out Response      agg_low_ci_limit agg_high_ci_limit  
<chr>    <chr>          <dbl>           <dbl>  
1 Female   1-13 days when physical health n~ 24.3 24.5  
2 Female   14+ days when physical health no~ 13.9 14.0  
3 Female   zero days when physical health n~ 61.6 61.8  
4 Male     1-13 days when physical health n~ 20.8 21.0  
5 Male     14+ days when physical health no~ 11.5 11.7  
6 Male     zero days when physical health n~ 67.4 67.6
```

```
plotDf |>  
  ggplot( aes(x=Break_Out,  
y=agg_percent,  
fill=Response,color=Response,position="fill")) +  
  geom_col() + xlab(my_q) +  
  theme(axis.text.x = element_text(angle = 90, vjust = 0.5, hjust=1))
```



```
## summary by gender
#(my_q)
#plotDf <- qDf |>
#  filter(BreakOutCategoryID=="CAT2") |>
#  select(Break_Out,Response,Sample_Size,Data_value) |>
#  na.omit() |>
#  rename(persons=Sample_Size) |>
#  mutate(Sample_Size=persons*100/Data_value) |>
#  group_by(Break_Out,Response) |>
#  summarize(agg_ss=sum(Sample_Size),
#            agg_persons=sum(persons),
#            agg_percent=agg_persons*100/agg_ss,
#            agg_percent_sdev=sqrt(agg_percent*(100-agg_percent)/agg_ss),
#            agg_low_ci_limit=agg_percent - 2*agg_percent_sdev,
#            agg_high_ci_limit=agg_percent + 2*agg_percent_sdev) |>
#            select(-c(agg_ss,agg_persons,agg_percent_sdev))
#  plotDf |>
#    select(-agg_percent) |>
#    print(n=20)
#
#  plotDf |>
#    ggplot( aes(x=Break_Out,
#                y=agg_percent,
```

```
#       fill=Response,color=Response,position="fill")) +  
#       geom_col() + xlab(my_q)
```

```
# comment out if next chunk is used  
# summary by age  
(my_q)
```

```
[1] "Days when physical health status not good "
```

```
plotDf <- qDf |>  
  filter(BreakOutCategoryID=="CAT3") |>  
  select(Break_Out,Response,Sample_Size) |>  
  na.omit() |>  
  rename(persons=Sample_Size) |>  
  group_by(Break_Out) |>  
  summarize(Response=Response,  
            persons=persons,  
            agg_ss=sum(persons))
```

Warning: Returning more (or less) than 1 row per `summarise()` group was deprecated in dplyr 1.1.0.

i Please use `reframe()` instead.
i When switching from `summarise()` to `reframe()`, remember that `reframe()` always returns an ungrouped data frame and adjust accordingly.

```
plotDf <- plotDf |>  
  group_by(Break_Out,Response) |>  
  summarize(agg_ss=agg_ss,  
            agg_persons=sum(persons),  
            agg_percent=agg_persons*100/agg_ss,  
            agg_percent_sdev=  
            sqrt(agg_percent*(100-agg_percent)/agg_ss),  
            agg_low_ci_limit=  
            agg_percent - 2*agg_percent_sdev,  
            agg_high_ci_limit=  
            agg_percent + 2*agg_percent_sdev) |>  
  distinct() |>  
  select(-c(agg_persons,agg_percent_sdev,agg_ss))
```

```

Warning: Returning more (or less) than 1 row per `summarise()` group was deprecated in
dplyr 1.1.0.
i Please use `reframe()` instead.
i When switching from `summarise()` to `reframe()`, remember that `reframe()` 
  always returns an ungrouped data frame and adjust accordingly.

```

```

plotDf |>
  select(-agg_percent) |>
  print(n=100)

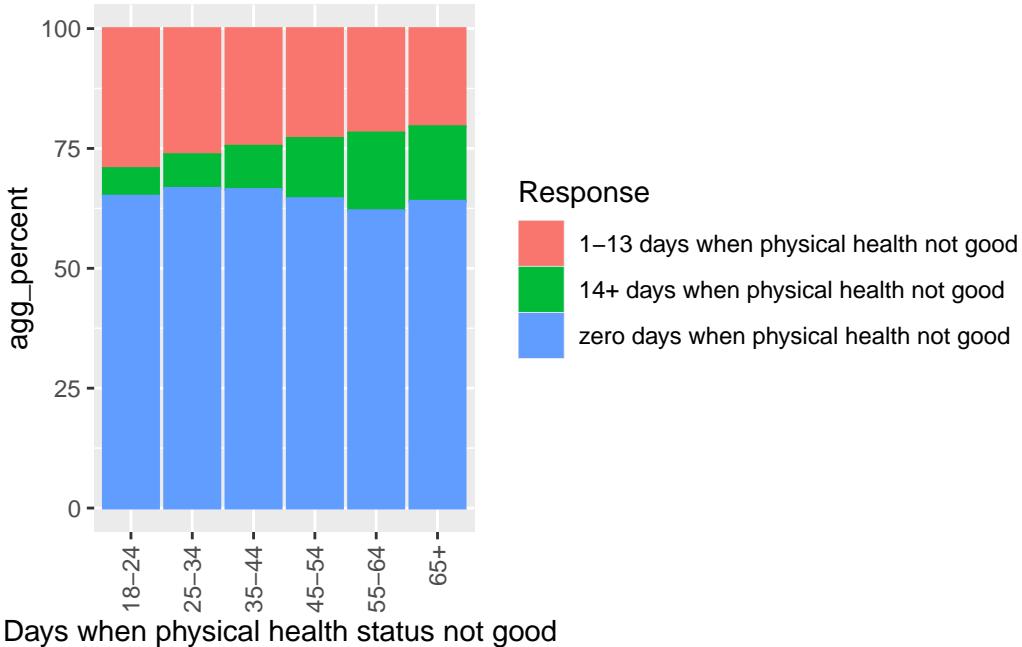
```

	Break_Out	Response	agg_low_ci_limit	agg_high_ci_limit
	<chr>	<chr>	<dbl>	<dbl>
1	18-24	1-13 days when physical health ~	28.9	29.5
2	18-24	14+ days when physical health n~	5.62	5.88
3	18-24	zero days when physical health ~	64.8	65.3
4	25-34	1-13 days when physical health ~	26.1	26.5
5	25-34	14+ days when physical health n~	6.92	7.13
6	25-34	zero days when physical health ~	66.5	66.9
7	35-44	1-13 days when physical health ~	24.4	24.7
8	35-44	14+ days when physical health n~	8.91	9.13
9	35-44	zero days when physical health ~	66.3	66.6
10	45-54	1-13 days when physical health ~	22.8	23.1
11	45-54	14+ days when physical health n~	12.4	12.7
12	45-54	zero days when physical health ~	64.4	64.7
13	55-64	1-13 days when physical health ~	21.6	21.9
14	55-64	14+ days when physical health n~	16.1	16.3
15	55-64	zero days when physical health ~	61.9	62.2
16	65+	1-13 days when physical health ~	20.4	20.6
17	65+	14+ days when physical health n~	15.5	15.6
18	65+	zero days when physical health ~	63.9	64.1

```

plotDf |>
  ggplot( aes(x=Break_Out,
  y=agg_percent,
  fill=Response,color=Response,position="fill")) +
    geom_col() + xlab(my_q) +
    theme(axis.text.x = element_text(angle = 90, vjust = 0.5, hjust=1))

```



```

## summary by age group
#(my_q)
#plotDf <- qDf |>
#  filter(BreakOutCategoryID=="CAT3") |>
#  select(Break_Out,Response,Sample_Size,Data_value) |>
#  na.omit() |>
#  rename(persons=Sample_Size) |>
#  mutate(Sample_Size=persons*100/Data_value) |>
#  group_by(Break_Out,Response) |>
#  summarize(agg_ss=sum(Sample_Size),
#  agg_persons=sum(persons),
#  agg_percent=agg_persons*100/agg_ss,
#  agg_percent_sdev=sqrt(agg_percent*(100-agg_percent)/agg_ss),
#  agg_low_ci_limit=agg_percent - 2*agg_percent_sdev,
#  agg_high_ci_limit=agg_percent + 2*agg_percent_sdev) |>
#  select(-c(agg_ss,agg_persons,agg_percent_sdev))
# plotDf |>
#  select(-agg_percent) |>
#  print(n=20)
# plotDf |>
#  ggplot( aes(x=Break_Out,
#  y=agg_percent,
#  fill=Response,color=Response,position="fill")) +

```

```

#           geom_col() + xlab(my_q)

# comment out if next chunk is used
# summary by race
(my_q)

[1] "Days when physical health status not good "

plotDf <- qDf |>
  filter(BreakOutCategoryID=="CAT4") |>
  select(Break_Out,Response,Sample_Size) |>
  na.omit() |>
  rename(persons=Sample_Size) |>
  group_by(Break_Out) |>
  summarise(Response=Response,
            persons=persons,
            agg_ss=sum(persons))

```

Warning: Returning more (or less) than 1 row per `summarise()`` group was deprecated in dplyr 1.1.0.

- i Please use `reframe()`` instead.
- i When switching from `summarise()`` to `reframe()`` , remember that `reframe()`` always returns an ungrouped data frame and adjust accordingly.

```

plotDf <- plotDf |>
  group_by(Break_Out,Response) |>
  summarise(agg_ss=agg_ss,
            agg_persons=sum(persons),
            agg_percent=agg_persons*100/agg_ss,
            agg_percent_sdev=
              sqrt(agg_percent*(100-agg_percent)/agg_ss),
            agg_low_ci_limit=
              agg_percent - 2*agg_percent_sdev,
            agg_high_ci_limit=
              agg_percent + 2*agg_percent_sdev) |>
  distinct() |>
  select(-c(agg_persons,agg_percent_sdev,agg_ss))

```

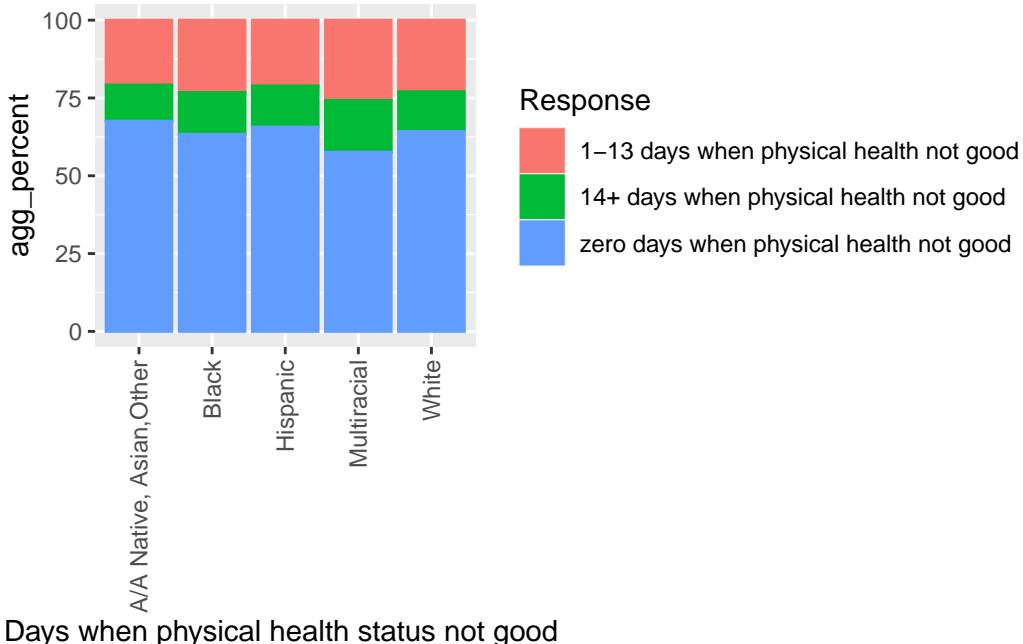
Warning: Returning more (or less) than 1 row per `summarise()`` group was deprecated in

```
dplyr 1.1.0.
i Please use `reframe()` instead.
i When switching from `summarise()` to `reframe()`, remember that `reframe()` always returns an ungrouped data frame and adjust accordingly.
```

```
plotDf |>
  select(-agg_percent) |>
  print(n=100)
```

	Break_Out	Response	agg_low_ci_limit	agg_high_ci_limit
	<chr>	<chr>	<dbl>	<dbl>
1	A/A Native, Asian,Other	1-13 days when ph~	20.6	21.1
2	A/A Native, Asian,Other	14+ days when phy~	11.4	11.8
3	A/A Native, Asian,Other	zero days when ph~	67.3	67.8
4	Black	1-13 days when ph~	23.0	23.4
5	Black	14+ days when phy~	13.3	13.6
6	Black	zero days when ph~	63.1	63.6
7	Hispanic	1-13 days when ph~	20.9	21.3
8	Hispanic	14+ days when phy~	13.1	13.4
9	Hispanic	zero days when ph~	65.4	65.9
10	Multiracial	1-13 days when ph~	25.4	26.2
11	Multiracial	14+ days when phy~	16.3	17.0
12	Multiracial	zero days when ph~	57.1	58.1
13	White	1-13 days when ph~	22.9	23.1
14	White	14+ days when phy~	12.7	12.8
15	White	zero days when ph~	64.2	64.3

```
plotDf |>
  ggplot( aes(x=Break_Out,
  y=agg_percent,
  fill=Response,color=Response,position="fill")) +
    geom_col() + xlab(my_q) +
    theme(axis.text.x = element_text(angle = 90, vjust = 0.5, hjust=1))
```



Days when physical health status not good

```
## summary by race
#(my_q)
#plotDf <- qDf |>
#  filter(BreakOutCategoryID=="CAT4") |>
#  select(Break_Out,Response,Sample_Size,Data_value) |>
#  na.omit() |>
#  rename(persons=Sample_Size) |>
#  mutate(Sample_Size=persons*100/Data_value) |>
#  group_by(Break_Out,Response) |>
#  summarize(agg_ss=sum(Sample_Size),
#            agg_persons=sum(persons),
#            agg_percent=agg_persons*100/agg_ss,
#            agg_percent_sdev=sqrt(agg_percent*(100-agg_percent)/agg_ss),
#            agg_low_ci_limit=agg_percent - 2*agg_percent_sdev,
#            agg_high_ci_limit=agg_percent + 2*agg_percent_sdev) |>
#            select(-c(agg_ss,agg_persons,agg_percent_sdev))
#plotDf |>
#  select(-agg_percent) |>
#  print(n=20)
#
#plotDf |>
#  ggplot( aes(x=Break_Out,y=agg_percent,
#             fill=Response,color=Response,position="fill")) +
#  geom_bar(stat="identity")
```

```

#           geom_col() + xlab(my_q)

# comment out if next chunk is used
# summary by education level
(my_q)

[1] "Days when physical health status not good"

plotDf <- qDf |>
  filter(BreakOutCategoryID=="CAT5") |>
  select(Break_Out,Response,Sample_Size) |>
  na.omit() |>
  rename(persons=Sample_Size) |>
  group_by(Break_Out) |>
  summarise(Response=Response,
            persons=persons,
            agg_ss=sum(persons))

```

Warning: Returning more (or less) than 1 row per `summarise()`` group was deprecated in dplyr 1.1.0.

- i Please use `reframe()`` instead.
- i When switching from `summarise()`` to `reframe()`` , remember that `reframe()`` always returns an ungrouped data frame and adjust accordingly.

```

plotDf <- plotDf |>
  group_by(Break_Out,Response) |>
  summarise(agg_ss=agg_ss,
            agg_persons=sum(persons),
            agg_percent=agg_persons*100/agg_ss,
            agg_percent_sdev=
              sqrt(agg_percent*(100-agg_percent)/agg_ss),
            agg_low_ci_limit=
              agg_percent - 2*agg_percent_sdev,
            agg_high_ci_limit=
              agg_percent + 2*agg_percent_sdev) |>
  distinct() |>
  select(-c(agg_persons,agg_percent_sdev,agg_ss))

```

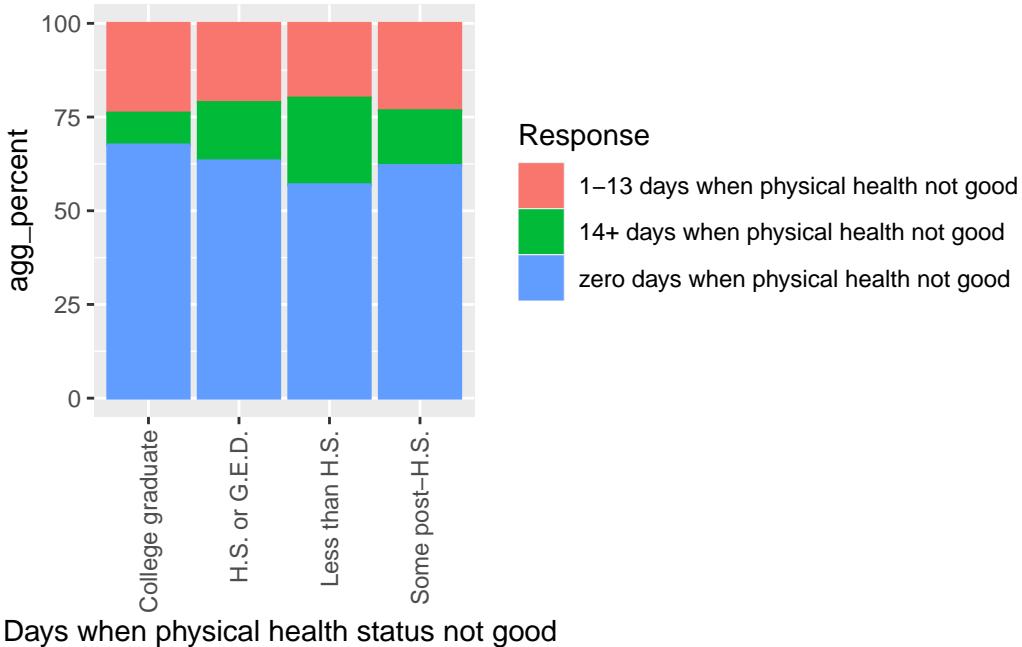
Warning: Returning more (or less) than 1 row per `summarise()`` group was deprecated in

```
dplyr 1.1.0.  
i Please use `reframe()` instead.  
i When switching from `summarise()` to `reframe()`, remember that `reframe()`  
  always returns an ungrouped data frame and adjust accordingly.
```

```
plotDf |>  
  select(-agg_percent) |>  
  print(n=100)
```

```
# A tibble: 12 x 4  
# Groups: Break_Out, Response [12]  
  Break_Out      Response      agg_low_ci_limit agg_high_ci_limit  
  <chr>        <chr>            <dbl>             <dbl>  
1 College graduate 1-13 days when physical ~          23.8           24.0  
2 College graduate 14+ days when physical h~          8.44            8.56  
3 College graduate zero days when physical ~         67.5            67.7  
4 H.S. or G.E.D.  1-13 days when physical ~         21.0            21.2  
5 H.S. or G.E.D.  14+ days when physical h~         15.5            15.7  
6 H.S. or G.E.D.  zero days when physical ~        63.2            63.5  
7 Less than H.S.  1-13 days when physical ~         19.7            20.1  
8 Less than H.S.  14+ days when physical h~         22.9            23.4  
9 Less than H.S.  zero days when physical ~        56.7            57.2  
10 Some post-H.S. 1-13 days when physical ~        23.2            23.4  
11 Some post-H.S. 14+ days when physical h~         14.5            14.7  
12 Some post-H.S. zero days when physical ~        62.0            62.2
```

```
plotDf |>  
  ggplot( aes(x=Break_Out,  
              y=agg_percent,  
              fill=Response,color=Response,position="fill")) +  
    geom_col() + xlab(my_q) +  
    theme(axis.text.x = element_text(angle = 90, vjust = 0.5, hjust=1))
```



```

## summary by education level
#(my_q)
#plotDf <- qDf |>
#  filter(BreakOutCategoryID=="CAT5") |>
#  select(Break_Out,Response,Sample_Size,Data_value) |>
#  na.omit() |>
#  rename(persons=Sample_Size) |>
#  mutate(Sample_Size=persons*100/Data_value) |>
#  group_by(Break_Out,Response) |>
#  summarize(agg_ss=sum(Sample_Size),
#  agg_persons=sum(persons),
#  agg_percent=agg_persons*100/agg_ss,
#  agg_percent_sdev=sqrt(agg_percent*(100-agg_percent)/agg_ss),
#  agg_low_ci_limit=agg_percent - 2*agg_percent_sdev,
#  agg_high_ci_limit=agg_percent + 2*agg_percent_sdev) |>
#  select(-c(agg_ss,agg_persons,agg_percent_sdev))
#plotDf |>
#  select(-agg_percent) |>
#  print(n=20)
#plotDf |>
#  ggplot( aes(x=Break_Out,
#  y=agg_percent,
#  fill=Response,color=Response,position="fill")) +

```

```

#           geom_col() + xlab(my_q)

# comment out if next chunk is used
# summary by income
(my_q)

[1] "Days when physical health status not good"

plotDf <- qDf |>
  filter(BreakOutCategoryID=="CAT6") |>
  select(Break_Out,Response,Sample_Size) |>
  na.omit() |>
  rename(persons=Sample_Size) |>
  group_by(Break_Out) |>
  summarize(Response=Response,
            persons=persons,
            agg_ss=sum(persons))

```

Warning: Returning more (or less) than 1 row per `summarise()`` group was deprecated in dplyr 1.1.0.

- i Please use `reframe()`` instead.
- i When switching from `summarise()`` to `reframe()`` , remember that `reframe()`` always returns an ungrouped data frame and adjust accordingly.

```

plotDf <- plotDf |>
  group_by(Break_Out,Response) |>
  summarize(agg_ss=agg_ss,
            agg_persons=sum(persons),
            agg_percent=agg_persons*100/agg_ss,
            agg_percent_sdev=
              sqrt(agg_percent*(100-agg_percent)/agg_ss),
            agg_low_ci_limit=
              agg_percent - 2*agg_percent_sdev,
            agg_high_ci_limit=
              agg_percent + 2*agg_percent_sdev) |>
  distinct() |>
  select(-c(agg_persons,agg_percent_sdev,agg_ss))

```

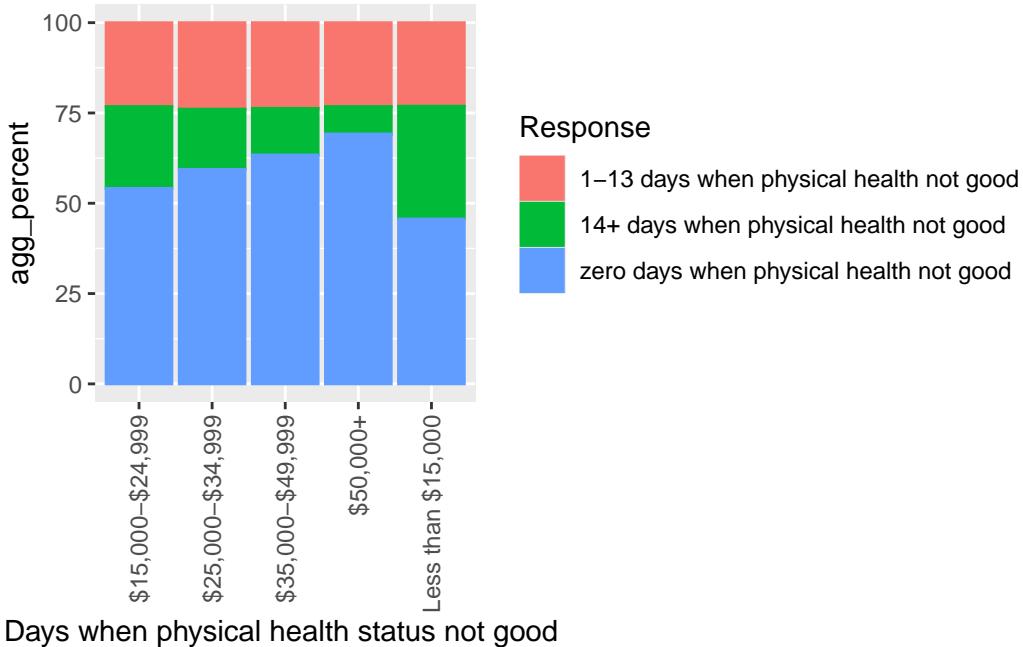
Warning: Returning more (or less) than 1 row per `summarise()`` group was deprecated in

```
dplyr 1.1.0.
i Please use `reframe()` instead.
i When switching from `summarise()` to `reframe()`, remember that `reframe()` always returns an ungrouped data frame and adjust accordingly.
```

```
plotDf |>
  select(-agg_percent) |>
  print(n=100)
```

	Break_Out	Response	agg_low_ci_limit	agg_high_ci_limit
	<chr>	<chr>	<dbl>	<dbl>
1	\$15,000-\$24,999	1-13 days when physical~	23.0	23.4
2	\$15,000-\$24,999	14+ days when physical ~	22.4	22.8
3	\$15,000-\$24,999	zero days when physical~	53.9	54.4
4	\$25,000-\$34,999	1-13 days when physical~	23.8	24.2
5	\$25,000-\$34,999	14+ days when physical ~	16.5	16.8
6	\$25,000-\$34,999	zero days when physical~	59.2	59.6
7	\$35,000-\$49,999	1-13 days when physical~	23.6	23.9
8	\$35,000-\$49,999	14+ days when physical ~	12.7	13.0
9	\$35,000-\$49,999	zero days when physical~	63.2	63.6
10	\$50,000+	1-13 days when physical~	23.1	23.3
11	\$50,000+	14+ days when physical ~	7.53	7.64
12	\$50,000+	zero days when physical~	69.1	69.3
13	Less than \$15,000	1-13 days when physical~	22.9	23.4
14	Less than \$15,000	14+ days when physical ~	31.0	31.5
15	Less than \$15,000	zero days when physical~	45.4	45.9

```
plotDf |>
  ggplot( aes(x=Break_Out,
  y=agg_percent,
  fill=Response,color=Response,position="fill")) +
    geom_col() + xlab(my_q) +
    theme(axis.text.x = element_text(angle = 90, vjust = 0.5, hjust=1))
```



```

## summary by income
#(my_q)
#plotDf <- qDf |>
#  filter(BreakOutCategoryID=="CAT6") |>
#  select(Break_Out,Response,Sample_Size,Data_value) |>
#  na.omit() |>
#  rename(persons=Sample_Size) |>
#  mutate(Sample_Size=persons*100/Data_value) |>
#  group_by(Break_Out,Response) |>
#  summarize(agg_ss=sum(Sample_Size),
#  agg_persons=sum(persons),
#  agg_percent=agg_persons*100/agg_ss,
#  agg_percent_sdev=sqrt(agg_percent*(100-agg_percent)/agg_ss),
#  agg_low_ci_limit=agg_percent - 2*agg_percent_sdev,
#  agg_high_ci_limit=agg_percent + 2*agg_percent_sdev) |>
#  select(-c(agg_ss,agg_persons,agg_percent_sdev))
#plotDf |>
#  select(-agg_percent) |>
#  print(n=20)
#
#plotDf |>
#  ggplot( aes(x=Break_Out,
#  y=agg_percent,

```

```
#     fill=Response,color=Response,position="fill")) +
#             geom_col() + xlab(my_q)
```

Project Guidelines

- **IDE/Language** Any. Free. Publicly available.
- **team size** 1. 2. or 3. period.
- **deliverables**
 - team members listing (due 11/10/2025)
 - link-shared 10-minute recorded presentation (due 12/3/2025). *Walk-through dashboard demo. Equal time for each team member.*