

# Obtaining population data

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## Population data

This file downloads detailed US population data by year state and age group from:

<http://seer.cancer.gov/popdata/download.html>

This is a large dataset which is then processed to obtain relevant summarised datasets:

- population of each state by year

The raw data is obtained from the url if it doesn't already exist:

```
filename<-"us.1969_2014.19ages.adjusted.txt.gz"
data_url <-"http://seer.cancer.gov/popdata/yr1969_2014.19ages/us.1969_2014.19ages.adjusted.txt.gz"

if (!file.exists(filename)){
  download.file(data_url, filename)
}

# Load data and rename data column to something more friendly
#
raw_data<-read.table(filename, stringsAsFactors = F)
names(raw_data)[1]<-"raw"
head(raw_data)
```

```
##              raw
## 1 1969AL01001991910000000159
## 2 1969AL01001991910100000657
## 3 1969AL01001991910200001137
## 4 1969AL01001991910300000956
## 5 1969AL01001991910400000721
## 6 1969AL01001991910500000424
```

The adjusted dataset is used which takes into account the population shifts in certain states following hurricane Katrina.

From time to time the data sets are updated - check and change the data\_url above to get the latest data

It can be seen that the raw data is compacted into just one column. The next section parses and summarises this column.

## Data format

The format of this data is described here:

<http://seer.cancer.gov/popdata/popdic.html>

The key elements needed for this module are:

field	start column	end column	Data type
Year	1	4	numeric
State postal abbreviation	5	6	

|character | |Population |19 |26 |numeric |

Field	Start Col	End Col	Data Type
Year	1	4	numeric
State	5	6	char
Population	19	26	numeric

The following code extracts the relevant columns and caches the results.

```
raw_data$year<-as.numeric(substr(raw_data$raw, 1,4))
raw_data$state<-as.character(substr(raw_data$raw, 5,6))
raw_data$population<-as.numeric(substr(raw_data$raw, 19,26))
```

## Create a new dataset

In this case we just want population per state per year:

```
summary <- raw_data %>% group_by(year, state) %>% summarise(population=sum(population))

# write out as a csv
write.table(summary, "states.csv", sep="," , row.names = F)
```