

# Covid19VaccineAnalysisInR

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## R Markdown

This is an R Markdown document. Markdown is a simple formatting syntax for authoring HTML, PDF, and MS Word documents. For more details on using R Markdown see <http://rmarkdown.rstudio.com>.

When you click the **Knit** button a document will be generated that includes both content as well as the output of any embedded R code chunks within the document. You can embed an R code chunk like this:

```
library(ggthemes)
```

```
## Warning: package 'ggthemes' was built under R version 4.1.2
```

```
library(tidyverse)
```

```
## Warning: package 'tidyverse' was built under R version 4.1.2
```

```
## -- Attaching packages ----- tidyverse 1.3.1 --
```

```
## v ggplot2 3.3.5      v purrr   0.3.4
## v tibble  3.1.4      v dplyr  1.0.7
## v tidyr   1.1.4      v stringr 1.4.0
## v readr   2.1.1      v forcats 0.5.1
```

```
## Warning: package 'tidyr' was built under R version 4.1.2
```

```
## Warning: package 'readr' was built under R version 4.1.2
```

```
## Warning: package 'purrr' was built under R version 4.1.2
```

```
## Warning: package 'dplyr' was built under R version 4.1.2
```

```
## Warning: package 'forcats' was built under R version 4.1.2
```

```
## -- Conflicts ----- tidyverse_conflicts() --
## x dplyr::filter() masks stats::filter()
## x dplyr::lag()     masks stats::lag()
```

```
library(colorspace)
library(scales)
```

```
##
## Attaching package: 'scales'

## The following object is masked from 'package:purrr':
##
##     discard

## The following object is masked from 'package:readr':
##
##     col_factor
```

```
library(ggplot2)
library(dplyr)
library(DT)
```

```
## Warning: package 'DT' was built under R version 4.1.2
```

```
library(tidyr)
library(data.table)
```

```
## Warning: package 'data.table' was built under R version 4.1.2
```

```
##
## Attaching package: 'data.table'

## The following objects are masked from 'package:dplyr':
##
##     between, first, last

## The following object is masked from 'package:purrr':
##
##     transpose
```

```
library(stringr)
library(summarytools)
```

```
## Warning: package 'summarytools' was built under R version 4.1.2
```

```
##
## Attaching package: 'summarytools'

## The following object is masked from 'package:tibble':
##
##     view
```

```
library(lubridate)
```

```
## Warning: package 'lubridate' was built under R version 4.1.2
```

```
##
```

```
## Attaching package: 'lubridate'
```

```
## The following objects are masked from 'package:data.table':
```

```
##
```

```
##     hour, isoweek, mday, minute, month, quarter, second, wday, week,  
##     yday, year
```

```
## The following objects are masked from 'package:base':
```

```
##
```

```
##     date, intersect, setdiff, union
```

## Including Plots

You can also embed plots, for example:

```
info_vac<-fread(file = "country_vaccinations.csv")  
head(info_vac)
```

```
##      country iso_code      date total_vaccinations people_vaccinated  
## 1: Afghanistan   AFG 2021-02-22              0              0  
## 2: Afghanistan   AFG 2021-02-23             NA             NA  
## 3: Afghanistan   AFG 2021-02-24             NA             NA  
## 4: Afghanistan   AFG 2021-02-25             NA             NA  
## 5: Afghanistan   AFG 2021-02-26             NA             NA  
## 6: Afghanistan   AFG 2021-02-27             NA             NA  
##      people_fully_vaccinated daily_vaccinations_raw daily_vaccinations  
## 1:                NA                NA                NA  
## 2:                NA                NA             1367  
## 3:                NA                NA             1367  
## 4:                NA                NA             1367  
## 5:                NA                NA             1367  
## 6:                NA                NA             1367  
##      total_vaccinations_per_hundred people_vaccinated_per_hundred  
## 1:                0                0  
## 2:                NA                NA  
## 3:                NA                NA  
## 4:                NA                NA  
## 5:                NA                NA  
## 6:                NA                NA  
##      people_fully_vaccinated_per_hundred daily_vaccinations_per_million  
## 1:                NA                NA  
## 2:                NA                35  
## 3:                NA                35  
## 4:                NA                35  
## 5:                NA                35  
## 6:                NA                35
```

```
##
## 1: Johnson&Johnson, Oxford/AstraZeneca, Pfizer/BioNTech, Sinopharm/Beijing
## 2: Johnson&Johnson, Oxford/AstraZeneca, Pfizer/BioNTech, Sinopharm/Beijing
## 3: Johnson&Johnson, Oxford/AstraZeneca, Pfizer/BioNTech, Sinopharm/Beijing
## 4: Johnson&Johnson, Oxford/AstraZeneca, Pfizer/BioNTech, Sinopharm/Beijing
## 5: Johnson&Johnson, Oxford/AstraZeneca, Pfizer/BioNTech, Sinopharm/Beijing
## 6: Johnson&Johnson, Oxford/AstraZeneca, Pfizer/BioNTech, Sinopharm/Beijing
##
##      source_name      source_website
## 1: World Health Organization https://covid19.who.int/
## 2: World Health Organization https://covid19.who.int/
## 3: World Health Organization https://covid19.who.int/
## 4: World Health Organization https://covid19.who.int/
## 5: World Health Organization https://covid19.who.int/
## 6: World Health Organization https://covid19.who.int/
```

```
str(info_vac)
```

```
## Classes 'data.table' and 'data.frame':  31240 obs. of  15 variables:
## $ country      : chr  "Afghanistan" "Afghanistan" "Afghanistan" "Afghanistan"
## $ iso_code     : chr  "AFG" "AFG" "AFG" "AFG" ...
## $ date         : IDate, format: "2021-02-22" "2021-02-23" ...
## $ total_vaccinations : num  0 NA NA NA NA NA 8200 NA NA NA ...
## $ people_vaccinated : num  0 NA NA NA NA NA 8200 NA NA NA ...
## $ people_fully_vaccinated : num  NA NA NA NA NA NA NA NA NA NA ...
## $ daily_vaccinations_raw : num  NA NA NA NA NA NA NA NA NA NA ...
## $ daily_vaccinations : num  NA 1367 1367 1367 1367 ...
## $ total_vaccinations_per_hundred : num  0 NA NA NA NA NA 0.02 NA NA NA ...
## $ people_vaccinated_per_hundred : num  0 NA NA NA NA NA 0.02 NA NA NA ...
## $ people_fully_vaccinated_per_hundred: num  NA NA NA NA NA NA NA NA NA NA ...
## $ daily_vaccinations_per_million : num  NA 35 35 35 35 35 35 41 46 52 ...
## $ vaccines     : chr  "Johnson&Johnson, Oxford/AstraZeneca, Pfizer/BioNTech, S
## $ source_name   : chr  "World Health Organization" "World Health Organization"
## $ source_website : chr  "https://covid19.who.int/" "https://covid19.who.int/" "
## - attr(*, ".internal.selfref")=<externalptr>
```

```
dim (info_vac)
```

```
## [1] 31240    15
```

```
info_temp_vac <- info_vac[,1:13]
colnames(info_temp_vac)
```

```
## [1] "country"      "iso_code"
## [3] "date"         "total_vaccinations"
## [5] "people_vaccinated" "people_fully_vaccinated"
## [7] "daily_vaccinations_raw" "daily_vaccinations"
## [9] "total_vaccinations_per_hundred" "people_vaccinated_per_hundred"
## [11] "people_fully_vaccinated_per_hundred" "daily_vaccinations_per_million"
## [13] "vaccines"
```

```
data.frame("Total_NA" = colSums(is.na(info_temp_vac))) %>%
  mutate ("Percentage_of_NA" = (colSums(is.na(info_temp_vac))/dim(info_temp_vac)[1]) %>%
    round (3) * 100)
```

##	Total_NA	Percentage_of_NA
## country	0	0.0
## iso_code	0	0.0
## date	0	0.0
## total_vaccinations	13789	44.1
## people_vaccinated	14686	47.0
## people_fully_vaccinated	17445	55.8
## daily_vaccinations_raw	16819	53.8
## daily_vaccinations	292	0.9
## total_vaccinations_per_hundred	13789	44.1
## people_vaccinated_per_hundred	14686	47.0
## people_fully_vaccinated_per_hundred	17445	55.8
## daily_vaccinations_per_million	292	0.9
## vaccines	0	0.0

```
info_temp_vac[is.na(info_temp_vac)] = 0
neglect = c('Faeroe Islands', 'Saint Lucia', 'Saint Vincent and the Grenadines', 'England', 'Northern Ireland')
info_temp_vac <- info_temp_vac %>%
  filter (!country %in% neglect)
unique(info_temp_vac$country)
```

## [1]	"Afghanistan"	"Albania"
## [3]	"Algeria"	"Andorra"
## [5]	"Angola"	"Anguilla"
## [7]	"Antigua and Barbuda"	"Argentina"
## [9]	"Armenia"	"Aruba"
## [11]	"Australia"	"Austria"
## [13]	"Azerbaijan"	"Bahamas"
## [15]	"Bahrain"	"Bangladesh"
## [17]	"Barbados"	"Belarus"
## [19]	"Belgium"	"Belize"
## [21]	"Benin"	"Bermuda"
## [23]	"Bhutan"	"Bolivia"
## [25]	"Bonaire Sint Eustatius and Saba"	"Bosnia and Herzegovina"
## [27]	"Botswana"	"Brazil"
## [29]	"British Virgin Islands"	"Brunei"
## [31]	"Bulgaria"	"Burkina Faso"
## [33]	"Cambodia"	"Cameroon"
## [35]	"Canada"	"Cape Verde"
## [37]	"Cayman Islands"	"Central African Republic"
## [39]	"Chad"	"Chile"
## [41]	"China"	"Colombia"
## [43]	"Comoros"	"Congo"
## [45]	"Cook Islands"	"Costa Rica"
## [47]	"Cote d'Ivoire"	"Croatia"
## [49]	"Cuba"	"Curacao"
## [51]	"Cyprus"	"Czechia"
## [53]	"Democratic Republic of Congo"	"Denmark"
## [55]	"Djibouti"	"Dominica"

## [57]	"Dominican Republic"	"Ecuador"
## [59]	"Egypt"	"El Salvador"
## [61]	"Equatorial Guinea"	"Estonia"
## [63]	"Eswatini"	"Ethiopia"
## [65]	"Fiji"	"Finland"
## [67]	"France"	"French Polynesia"
## [69]	"Gabon"	"Gambia"
## [71]	"Georgia"	"Germany"
## [73]	"Ghana"	"Gibraltar"
## [75]	"Greece"	"Greenland"
## [77]	"Grenada"	"Guatemala"
## [79]	"Guernsey"	"Guinea"
## [81]	"Guinea-Bissau"	"Guyana"
## [83]	"Honduras"	"Hong Kong"
## [85]	"Hungary"	"Iceland"
## [87]	"India"	"Indonesia"
## [89]	"Iran"	"Iraq"
## [91]	"Ireland"	"Isle of Man"
## [93]	"Israel"	"Italy"
## [95]	"Jamaica"	"Japan"
## [97]	"Jersey"	"Jordan"
## [99]	"Kazakhstan"	"Kenya"
## [101]	"Kosovo"	"Kuwait"
## [103]	"Kyrgyzstan"	"Laos"
## [105]	"Latvia"	"Lebanon"
## [107]	"Lesotho"	"Liberia"
## [109]	"Libya"	"Liechtenstein"
## [111]	"Lithuania"	"Luxembourg"
## [113]	"Macao"	"Madagascar"
## [115]	"Malawi"	"Malaysia"
## [117]	"Maldives"	"Mali"
## [119]	"Malta"	"Mauritania"
## [121]	"Mauritius"	"Mexico"
## [123]	"Moldova"	"Monaco"
## [125]	"Mongolia"	"Montenegro"
## [127]	"Montserrat"	"Morocco"
## [129]	"Mozambique"	"Myanmar"
## [131]	"Namibia"	"Nauru"
## [133]	"Nepal"	"Netherlands"
## [135]	"New Caledonia"	"New Zealand"
## [137]	"Nicaragua"	"Niger"
## [139]	"Nigeria"	"Niue"
## [141]	"North Macedonia"	"Northern Cyprus"
## [143]	"Norway"	"Oman"
## [145]	"Pakistan"	"Palestine"
## [147]	"Panama"	"Papua New Guinea"
## [149]	"Paraguay"	"Peru"
## [151]	"Philippines"	"Pitcairn"
## [153]	"Poland"	"Portugal"
## [155]	"Qatar"	"Romania"
## [157]	"Russia"	"Rwanda"
## [159]	"Saint Helena"	"Samoa"
## [161]	"San Marino"	"Sao Tome and Principe"
## [163]	"Saudi Arabia"	"Senegal"

```
## [165] "Serbia" "Seychelles"
## [167] "Sierra Leone" "Singapore"
## [169] "Sint Maarten (Dutch part)" "Slovakia"
## [171] "Slovenia" "Solomon Islands"
## [173] "Somalia" "South Africa"
## [175] "South Korea" "South Sudan"
## [177] "Spain" "Sri Lanka"
## [179] "Sudan" "Suriname"
## [181] "Sweden" "Switzerland"
## [183] "Syria" "Taiwan"
## [185] "Tajikistan" "Thailand"
## [187] "Timor" "Togo"
## [189] "Tonga" "Trinidad and Tobago"
## [191] "Tunisia" "Turkey"
## [193] "Turkmenistan" "Turks and Caicos Islands"
## [195] "Tuvalu" "Uganda"
## [197] "Ukraine" "United Arab Emirates"
## [199] "United Kingdom" "United States"
## [201] "Uruguay" "Uzbekistan"
## [203] "Vanuatu" "Venezuela"
## [205] "Vietnam" "Wallis and Futuna"
## [207] "Yemen" "Zambia"
## [209] "Zimbabwe"
```

```
info_temp_vac$vaccines <- str_replace_all(info_temp_vac$vaccines, " ", "")
usage_vac<- unique(info_temp_vac$vaccines)
vaccine<- vector()
for (i in usage_vac){
  for (j in strsplit(i, ",")){
    vaccine<- c(vaccine, j)
  }
}
used_vcc<- unique(vaccine)
used_vcc
```

```
## [1] "Johnson&Johnson" "Oxford/AstraZeneca" "Pfizer/BioNTech"
## [4] "Sinopharm/Beijing" "Sinovac" "SputnikV"
## [7] "Moderna" "Covaxin" "CanSino"
## [10] "Sinopharm/Wuhan" "Abdala" "Soberana02"
## [13] "QazVac" "Sinopharm/HayatVax" "EpiVacCorona"
## [16] "RBD-Dimer"
```

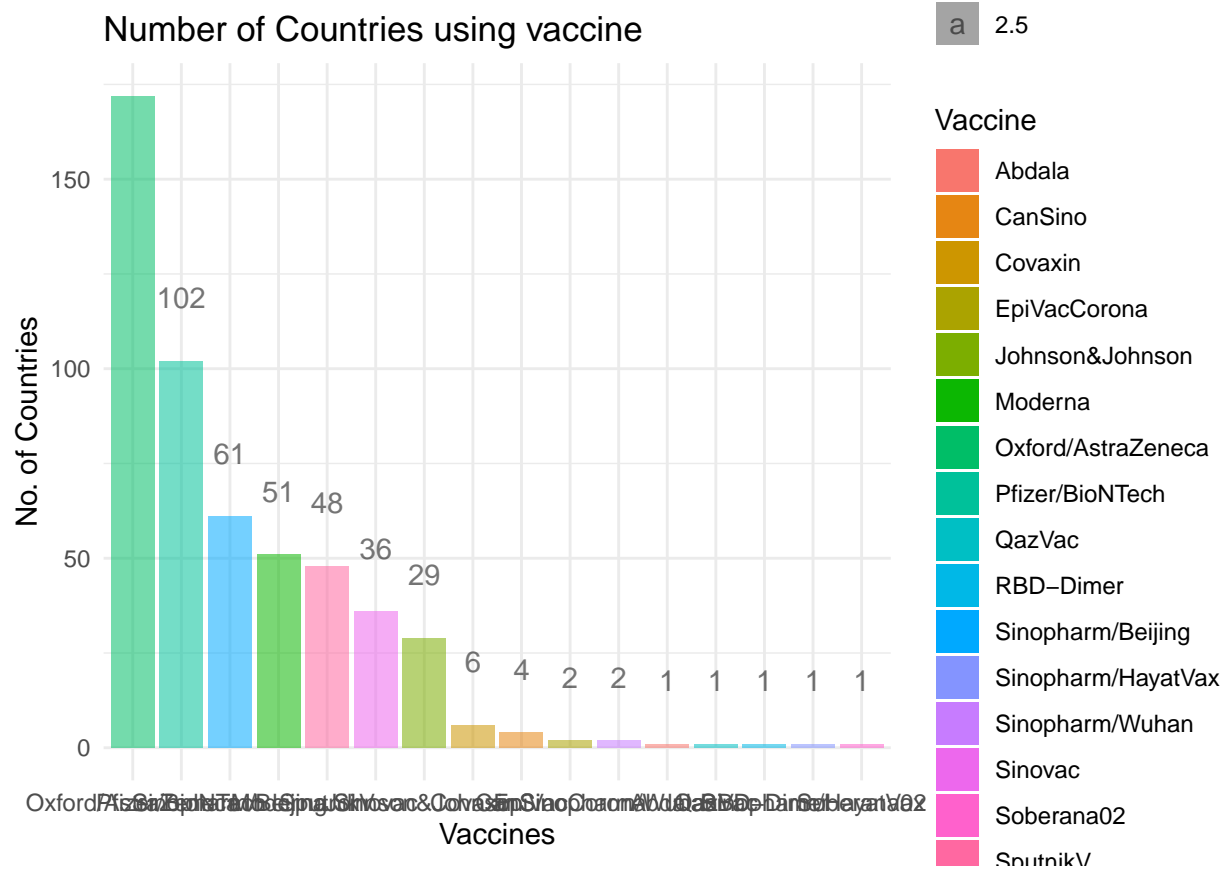
```
info_vac_val <- data.frame(matrix(ncol = length(used_vcc), nrow = 0))
for (i in info_temp_vac$vaccines){
  info_vac_val<- rbind(info_vac_val, Vectorize(grepl, USE.NAMES = TRUE)(used_vcc, str_replace_all(i, " "
}))
info_vac_val[info_vac_val == TRUE] = 1
info_vac_val[info_vac_val == FALSE] =0
colnames(info_vac_val) <- paste0(unique(vaccine))

countrywise_vacc<- info_vac_val %>%
mutate(country = info_temp_vac$country)%>%
group_by(country)%>%
```

```

summarise_all(sum)
data <- data.frame("No_of_countries"= apply(countrywise_vacc[-1],2, function(c)sum(c!=0)))
cbind("Vaccine"=row.names(data),data) %>%
ggplot(mapping=aes(x=reorder(Vaccine, -No_of_countries), y=No_of_countries, fill = Vaccine, alpha=2.5))+
geom_col() +
labs(x = "Vaccines", y = "No. of Countries", title = "Number of Countries using vaccine")+
geom_text(aes(label = No_of_countries), vjust=-2.5)+
theme_minimal()

```



Note that the `echo = FALSE` parameter was added to the code chunk to prevent printing of the R code that generated the plot.