

# Analisis\_Sentimen

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```
library(e1071)
```

```
## Warning: package 'e1071' was built under R version 4.1.2
```

```
library(caret)
```

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

```
## Loading required package: ggplot2
```

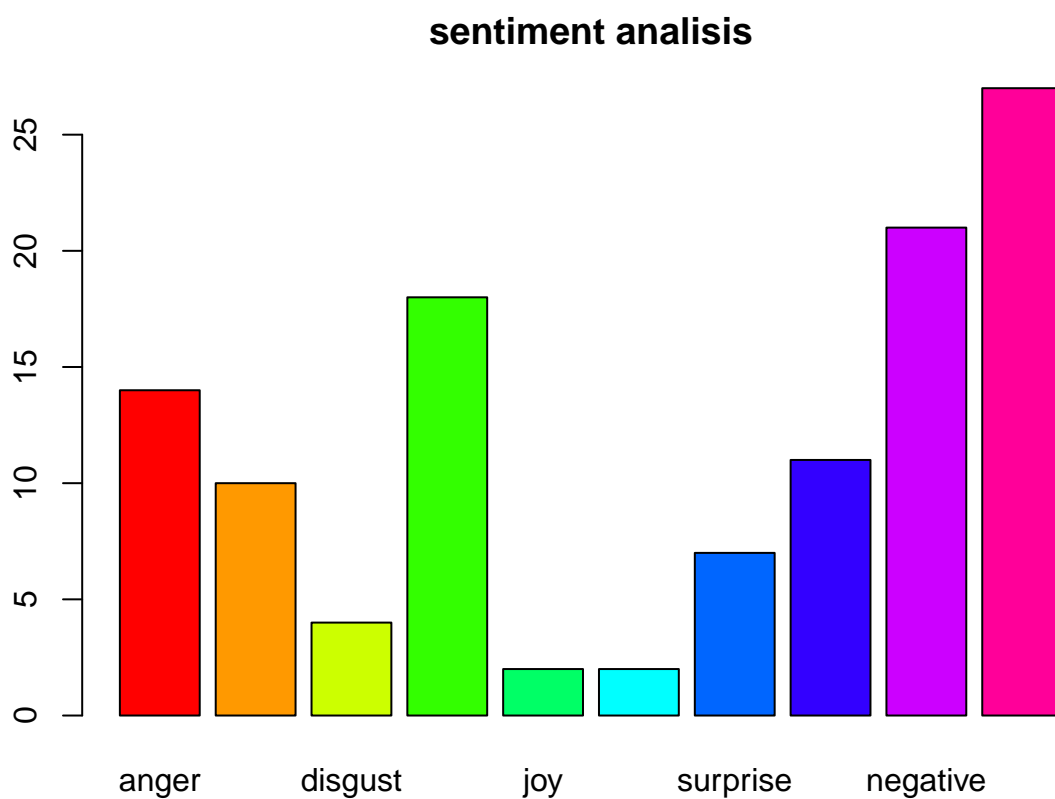
```
## Loading required package: lattice
```

```
library(syuzhet)
```

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

```
datavaksin<-read.csv("vaksin.csv", stringsAsFactors = FALSE)  
review <- as.character(datavaksin$text)  
s<-get_nrc_sentiment(review)
```

```
review_combine<-cbind(datavaksin$text, s)  
par(mar=rep(3,4))  
a<-barplot(colSums(s), col=rainbow(10), ylab='count', main='sentiment analisis')
```



```
brplt<-a
```

```
library(tm)
```

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

```
## Loading required package: NLP
```

```
##
```

```
## Attaching package: 'NLP'
```

```
## The following object is masked from 'package:ggplot2':
```

```
##
```

```
##   annotate
```

```
library(RTextTools)
```

```
## Loading required package: SparseM
```

```
##
```

```
## Attaching package: 'SparseM'
```

```
## The following object is masked from 'package:base':
##
##     backsolve
```

```
library(e1071)
library(dplyr)
```

```
##
## Attaching package: 'dplyr'
```

```
## The following objects are masked from 'package:stats':
##
##     filter, lag
```

```
## The following objects are masked from 'package:base':
##
##     intersect, setdiff, setequal, union
```

```
library(caret)
df<-read.csv("vaksin.csv", stringsAsFactors = FALSE)
glimpse(df)
```

```
## Registered S3 method overwritten by 'cli':
##   method      from
##   print.tree tree
```

```
## Rows: 1,000
## Columns: 2
## $ X      <int> 1, 2, 3, 4, 5, 6, 7, 8, 9, 10, 11, 12, 13, 14, 15, 16, 17, 18, 19~
## $ text <chr> "hi siapa warga astrazeneca u all dah dapat booster ke \n\nke aku~
```

```
set.seed(20)
df<-df[sample(nrow(df)),]
df<-df[sample(nrow(df)),]
glimpse(df)
```

```
## Rows: 1,000
## Columns: 2
## $ X      <int> 223, 831, 568, 122, 927, 958, 708, 680, 683, 867, 256, 338, 304, ~
## $ text <chr> "InHealthVC AstraZeneca Go Lavanya", "Adapun jenis vaksin yang di~
```

```
corpus<-Corpus(VectorSource(df$text))
corpus
```

```
## <<SimpleCorpus>>
## Metadata: corpus specific: 1, document level (indexed): 0
## Content: documents: 1000
```

```
inspect(corpus[1:10])
```

```
## <<SimpleCorpus>>
## Metadata: corpus specific: 1, document level (indexed): 0
## Content: documents: 10
##
## [1] InHealthVC AstraZeneca Go Lavanya
## [2] Adapun jenis vaksin yang disuntikan adalah Moderna AstraZeneca dan Coronavac
## [3] jenis vaksin dos penggalak disyorkan\nComirnaty Pfizer\nAstraZeneca\nCoronaVac Sinovac\n\nInfo
## [4] Warga Diminta Tak Ragu dengan Vaksin Pfizer dan AstraZeneca Menkes Pastikan Aman\nProkes Tangk
## [5] RT bernamadotcom TWG syor pemberian vaksin Comirnaty AstraZeneca CoronaVac sebagai dos penggalak
## [6] so dos penggalak ni aku boleh dapat paling awal bulan huhu \n\nsekurangkurangnya enam bulan se
## [7] RT TedHilbert Kematian akibat vaksin covid terbukti oleh otopsi StopPaksaVaksin UmatDukungReuni
## [8] Hallo jangan lupa ya daftar Vaksinasi dosis di Unikama pendaftaran mulai dari November \n\nDa
## [9] Emang sih Aku setelah vaksin astrazeneca malah berasa imun tubuh berkurang dikit dikit cape kur
## [10] RT restasukabumi Assalamualaikum Wr Wb\n\nSelamat siang warga Sukabumi sekitarnya\n\nGerai Vaks
```

```
corpus.clean<- corpus %>%
  tm_map(content_transformer(tolower))%>%
  tm_map(removePunctuation)%>%
  tm_map(removeNumbers)%>%
  tm_map(removeWords,stopwords(kind = "en"))%>%
  tm_map(stripWhitespace)
```

```
## Warning in tm_map.SimpleCorpus(., content_transformer(tolower)): transformation
## drops documents
```

```
## Warning in tm_map.SimpleCorpus(., removePunctuation): transformation drops
## documents
```

```
## Warning in tm_map.SimpleCorpus(., removeNumbers): transformation drops documents
```

```
## Warning in tm_map.SimpleCorpus(., removeWords, stopwords(kind = "en")):
## transformation drops documents
```

```
## Warning in tm_map.SimpleCorpus(., stripWhitespace): transformation drops
## documents
```

```
dtm<-DocumentTermMatrix(corpus.clean)
```

```
inspect(dtm[1:10,1:20])
```

```
## <<DocumentTermMatrix (documents: 10, terms: 20)>>
## Non-/sparse entries: 41/159
## Sparsity : 80%
## Maximal term length: 19
## Weighting : term frequency (tf)
## Sample :
## Terms
## Docs astrazeneca comirnaty coronavac dan dos inhealthvc jenis penggalak pfizer
```

```
##      1      1      0      0  0  0      1      0      0      0
##     10      0      0      0  0  0      0      0      0      0
##      2      1      0      1  1  0      0      1      0      0
##      3      1      1      1  0  1      0      1      1      1
##      4      1      0      0  1  0      0      0      0      1
##      5      1      1      1  0  1      0      0      1      0
##      6      0      0      0  0  2      0      0      1      0
##      7      0      0      0  0  0      0      0      0      0
##      8      0      0      0  0  0      0      0      0      0
##      9      1      0      0  0  0      0      0      0      0
##      Terms
## Docs vaksin
##      1      0
##     10      1
##      2      1
##      3      1
##      4      1
##      5      1
##      6      0
##      7      1
##      8      0
##      9      1
```

```
df.train<-df[1:50,]
df.test<-df[51:100,]

dtm.train<-dtm[1:50,]
dtm.test<-dtm[51:100,]

corpus.clean.train<-corpus.clean[1:50]
corpus.clean.test<-corpus.clean[51:100]

dim(dtm.train)
```

```
## [1] 50 2515
```

```
fivefreq<-findFreqTerms(dtm.train,5)
length(fivefreq)
```

```
## [1] 13
```

```
dtm.train.nb <-DocumentTermMatrix (corpus.clean.test,control = list(dictionary=fivefreq))
dim(dtm.train.nb)
```

```
## [1] 50 13
```

```
dtm.test.nb <-DocumentTermMatrix (corpus.clean.test,control = list(dictionary=fivefreq))
dim(dtm.test.nb)
```

```
## [1] 50 13
```

```

convert_count<-function(x){
  y <-ifelse(x>0,1,0)
  y <-factor(y,levels=c(0,1),labels=c ("no","yes"))
  y
}
trainNB<-apply(dtm.train.nb, 2, convert_count)
testNB<-apply(dtm.test.nb, 1, convert_count)
library(wordcloud)

```

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

```
## Loading required package: RColorBrewer
```

```
wordcloud(corpus.clean,min.freq = 4,max.words=100,random.order=F,colors=brewer.pal (8,"Dark2"))
```

```
## Warning in strwidth(words[i], cex = size[i], ...): conversion failure on
## 'pedazitosdemí' in 'mbcsToSbcs': dot substituted for <ed>
```

```
## Warning in text.default(x1, y1, words[i], cex = size[i], offset = 0, srt
## = rotWord * : conversion failure on 'pedazitosdemí' in 'mbcsToSbcs': dot
## substituted for <ed>
```

```
## Warning in strwidth(words[i], cex = size[i], ...): conversion failure on
## 'rosalíasotés' in 'mbcsToSbcs': dot substituted for <ed>
```

```
## Warning in strwidth(words[i], cex = size[i], ...): conversion failure on
## 'rosalíasotés' in 'mbcsToSbcs': dot substituted for <e9>
```

```
## Warning in text.default(x1, y1, words[i], cex = size[i], offset = 0, srt
## = rotWord * : conversion failure on 'rosalíasotés' in 'mbcsToSbcs': dot
## substituted for <ed>
```

```
## Warning in text.default(x1, y1, words[i], cex = size[i], offset = 0, srt
## = rotWord * : conversion failure on 'rosalíasotés' in 'mbcsToSbcs': dot
## substituted for <e9>
```

