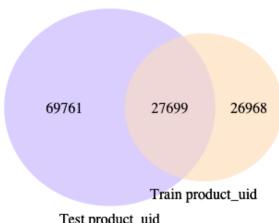
Data Analytics Final Project

Home Depot Competition

Group Members: Tong Wei (TOW6), Fengxi Liu (FEL34), Jinrong Liu (JIL181)

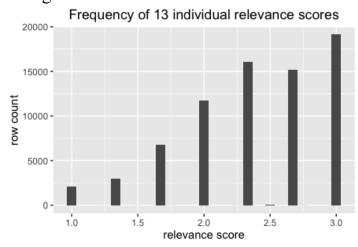
1. Data Exploration

a. Venn Diagram

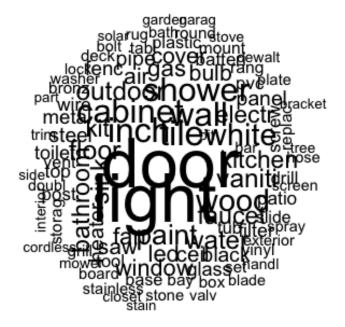


Test product_uid

b. Histogram



c. Words Cloud



d. Codes

```
#------
#install.packages("tm")
#install.packages("wordcloud")
#install.packages("SnowballC")
library(tm)
library(wordcloud)
library(SnowballC)

trainCorpus <- Corpus(VectorSource(train$search_term)) #create a corpus
trainCorpus <- tm_map(trainCorpus, PlainTextDocument) #convert the corpus to a plain text document
trainCorpus <- tm_map(trainCorpus, removePunctuation)
trainCorpus <- tm_map(trainCorpus, removeWords, stopwords('english'))
trainCorpus <- tm_map(trainCorpus, stemDocument)
wordcloud(trainCorpus, max.words = 100, random.order = FALSE)</pre>
```

2. Preparing Data

We do the data preparing following the steps on the guidance.

Step one:

Lowercase. Tolower function is used here.

```
train[] <- lapply(train, tolower)
Result:</pre>
```



Step two:

Remove stopwords and stem data:

We used the functions in QDAP package to achieve removing stopwords in a for loop. At the same time, we found we can do the stem in the loop. We choose porter stemmer in the SnowballC package, and stemming data in the second for loop inside the first loop.

```
ifor (i in 1:length(train$id)) {
    unit <- train$product_title[i]
    unit <- rm_stopwords(unit,Top25Words,unlist=TRUE)
    for (m in 1:length(unit)) {
        temp[m]=wordStem(unit[m],language = "porter")
        unit[m]<-temp[m]
    }
    p <- paste(unit,collapse =' ')
    product_title <- rbind(product_title,p)
}
Results:

briggs and stratton lawn mower

Before

After</pre>
```

Step three:

Correct misspelled words

This step is important because this will greatly affect our final result. We use which_misspelled function in QDAP package and look up into the QDAP dictionary. We can achieve all the correcting in a loop.

3. Model Selection and Model Evaluation

Model choosing: gbm model (Generalized Boosted Regression Models) R: install.packages('gbm')

```
gbm_model <- gbm.fit(train[,7:9],train$relevance,distribution = "gaussian",interaction.depth = 2,shrinkage=0.05,n.trees=500)
test_relevance <- predict(gbm_model,test[,6:8],n.trees=600)
test_relevance <- ifelse(test_relevance>3,3,test_relevance)
test_relevance <- ifelse(test_relevance<1,1,test_relevance)</pre>
```

Model evaluation: k-fold validation

K-fold cross validation is one way to improve over the holdout method. The data set is divided into k subsets, and the holdout method is repeated k times. Each time, one of the k subsets is used as the test set and the other k-l subsets are put together to form a training set. Then the average error across all k trials is computed.

RMSE result:

```
pred =ifelse(pred<0,0,pred)
obs = test2$count
error = obs-pred
me=mean(error)
rmse=sqrt(mean(error^2))
rmse</pre>
```

1551 new TongWei 0.50507