# **STATS202 Final Project**

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### **Summary**

#### Selection

I conduct visualizations first. Some initial data exploration shows:

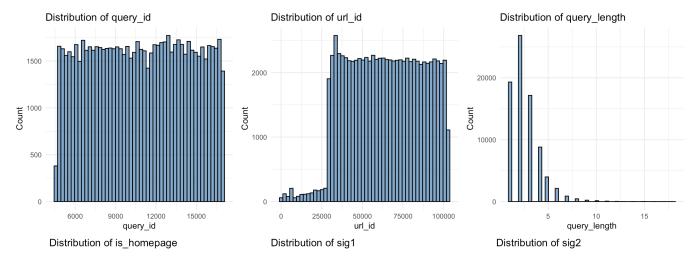
- The dependent variable is not imbalanced ( $\hat{p}=.4371$ )
- id and query\_id are unique identifiers.
- url\_id is not clearly a unique identifier because the distribution of histogram buckets is not uniform.
- is\_homepage is a binary variable.
- query\_length is a count.
- sig3, sig4, sig5, sig6 are all highly skewed.

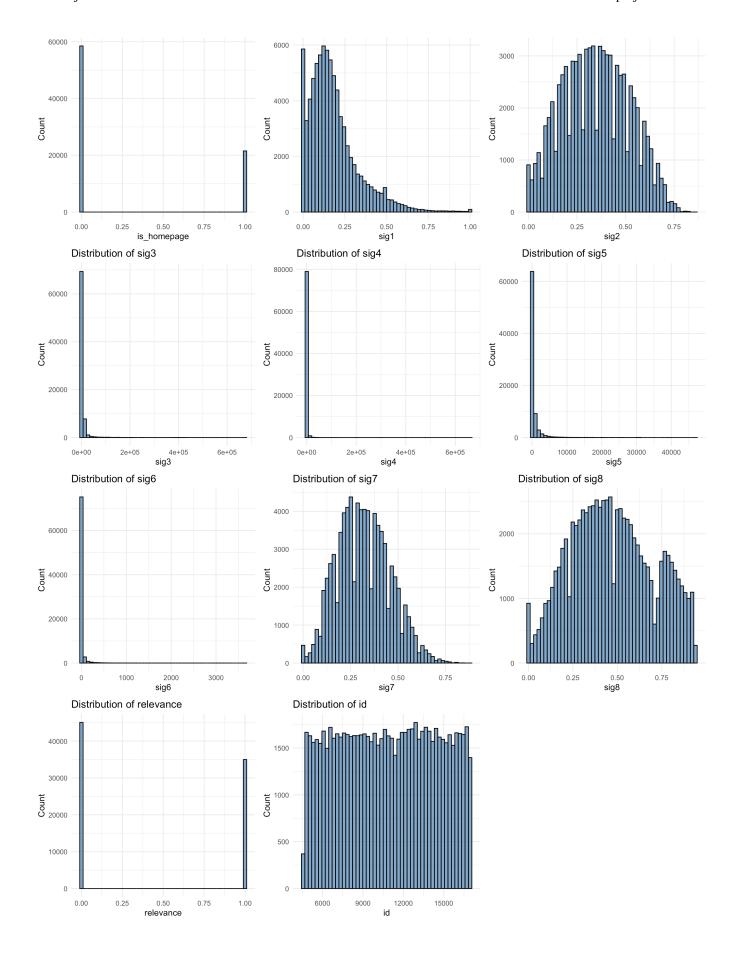
I visualize the density of each predictor grouped by dependent variable. The overlay of the plots helps visualize whether the variable of interest is systematically higher for certain values of the predictors. To make the contrast more conspicuous, I log-transform the skewed variables. I also group predictor values into deciles and condition on the dependent variable. sig2 appears conspicuously useful as a separator of the dependent variable.

For completeness I investigate multicollinearity but find no significant (|r| > .8) correlations. Variance inflation is also more germane to inference than to prediction; the point estimates with multicollinearity do not change even while standard errors increase.

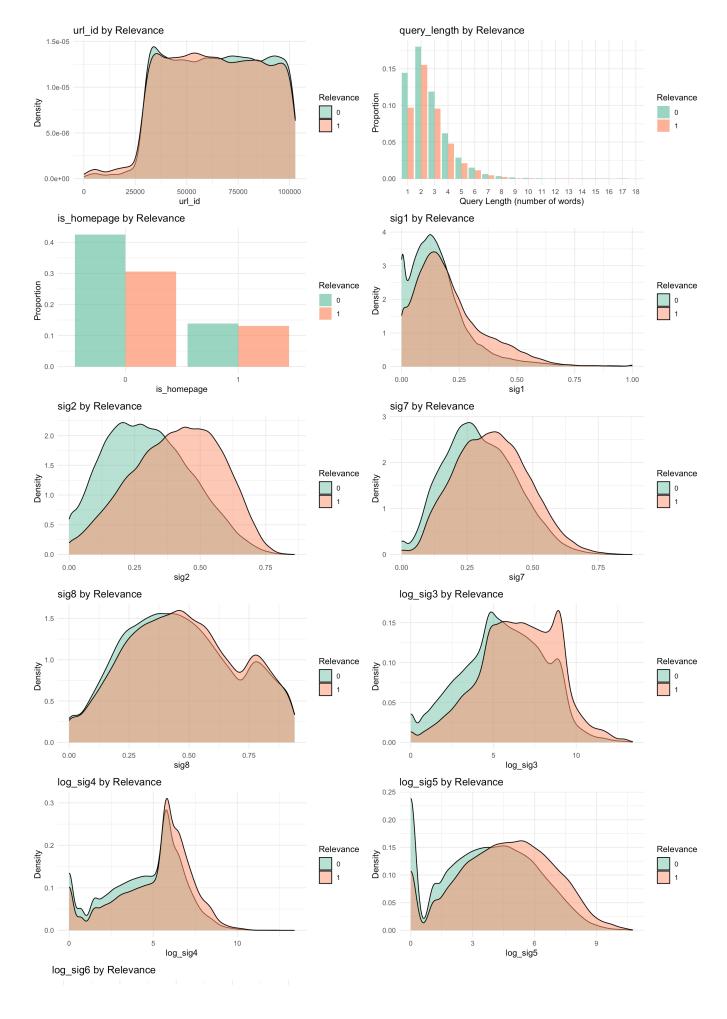
### Visualizations

#### Distributions of Variables

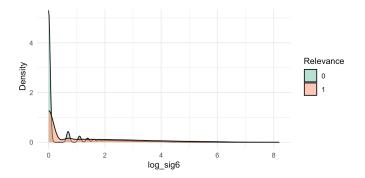




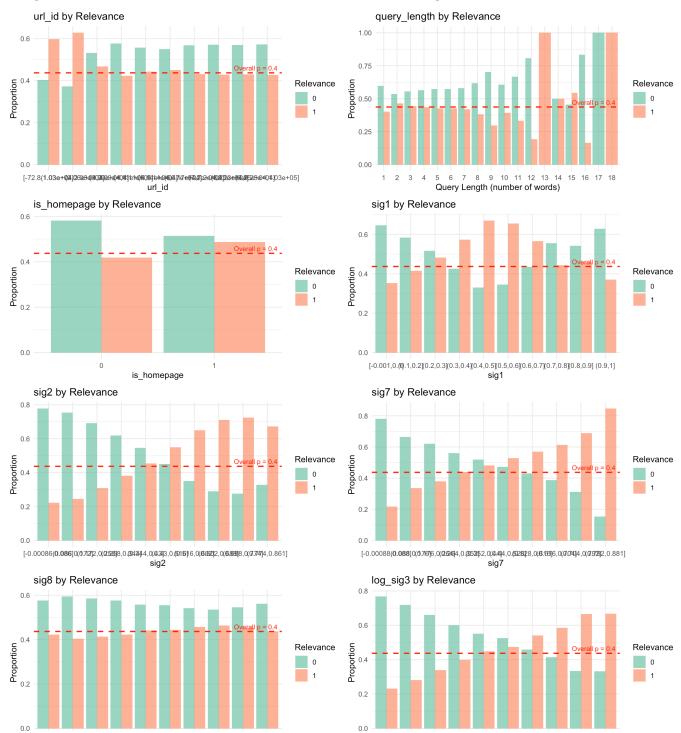
Histogram of Select Predictors Conditioned on Dependent Variable

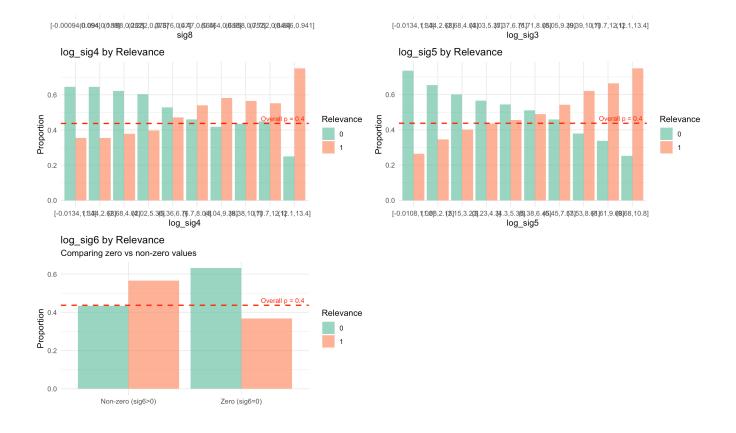


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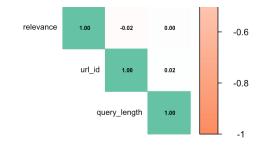
## Proportions of Select Predictors Conditioned on Dependent Variables Deciles





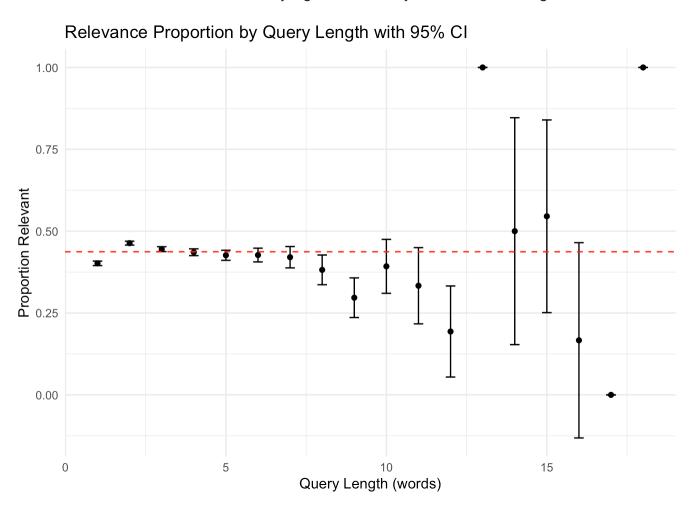
#### Correlation Plot





### Query Length

Longer query lengths have a very high proportion of relevant entries. I calculate standard errors to see whether these trends are statistically significant or may result in over-fitting.

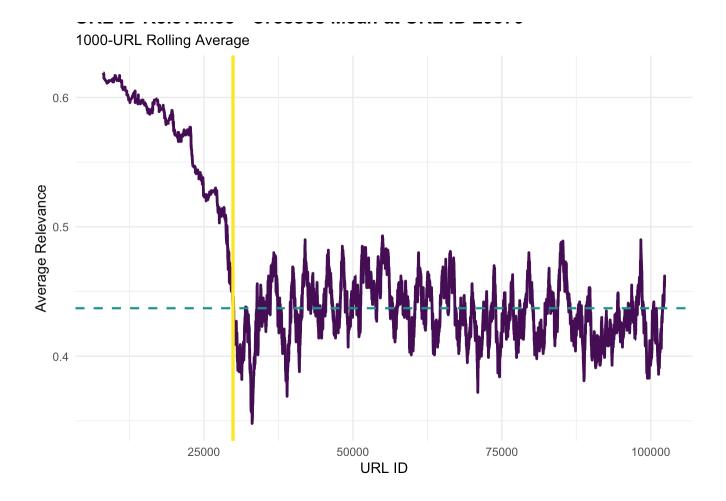


They are not statistically significant.

#### **URL ID**

Crossing point at URL ID: 29879

**URL ID Relevance - Crosses Mean at URL ID 29879** 



### **Pre-processing**

I code is\_homepage as an unordered factor.

I remove query\_length because the most predictive lengths are also the most rare. It is sensible to represent query\_length as a factor variable, because the proportion of relevance does not appear to increase linearly, but factors introduce various challenges with prediction. If some factor is not present in the future data, then our trained model will not work. A compromise is necessary.

url\_id will need to be handled differently. The numeric scale is not sensible but neither are factors, which would lead to over-fitting. Ultimately, it seems risky to include data that may be idiosyncratic, so I exclude it for conservatism.

sig2, sig7, and sig8 appear to have regular drop-offs in the histograms. This is an artifact of the measurement scale, which records values in increments of 0.01. Some histogram bins therefore align with the allowed values and receive higher counts, while others fall between these values and receive few or none, producing the visible pattern of regular dips. This is not a problem for analysis.

There are no missing values.

#### **Transformation**

I apply the following transformations:

- Log-transform skewed variables (sig3, sig4, sig5) for linear models.
  - This linearizes for LASSO, and monotone transformations are unlikely to hurt more flexible modeling strategies.
  - I use log1p from base R, which adds a constant of 1 to each value to avoid issues with zeros. The loss of the usual interpretability - where natural logs approximate percentage changes - is not important here.
- Two-part transformation for sig6: create a dummy variable indicating zero values, then log-transform the positive values. This preserves the information in zeros (which show a higher proportion of relevance) while making the positive values less skewed.
- I add binary variables for query\_length 1 and 2, which are statistically significant according to my earlier visualization.
- I add a factor variable for url\_id below 30000, the cutoff at which mean relevance was highest.
- After transforming, I remove url\_id and other variables.

Note that I do not standardize at this stage. Since I will later use cross-validation, scaling of the variables should happen within each fold where appropriate.

# **Data Mining**

I will use 10-fold cross-validation to estimate generalization error for each modeling strategy. In all cases, I will evaluate model accuracy, picking the tuning parameters and then the strategy which maximizes accuracy.

My modeling strategies are:

- 1. LASSO logistic regression: Suitable for correlated predictors and sparse data to reduce variance. Factor variables are one-hot encoded. The tuning variable is  $\lambda$ , imposing a progressively higher penalty that shrinks coefficients towards zero. I use a "kitchen-sink" approach with all interactions.
- 2. KNN: A simple, intuitive method that makes minimal assumptions about the data's functional form. Factor variables are one-hot encoded; the curse of dimensionality is not salient given the relatively low dimensions. The tuning variable is k, or the number of nearest neighbors.
- 3. Random Forests: Well-suited to low-dimensional tabular data and capable of capturing complex, nonlinear interactions automatically without needing the earlier transformations; especially attractive given apparent splines in some variables (sig1). Factor variables are not one-hot encoded because one-hot encoding increases the number of predictors, resulting in over-representation of factor variables by random forests. The tuning parameters are below:

```
rf_grid <- expand.grid(
  mtry = c(2, 4, 6, default_mtry),
  colitrule = "gini"</pre>
```

```
min.node.size = c(1, 5, 10)
)
```

4. I also use boosting with XGBoost. The tuning parameters are below:

```
params <- list(
  objective = "binary:logistic",
  eval_metric = "error",
  max_depth = 6,
  eta = 0.1,
  subsample = 0.8,
  colsample_bytree = 0.8
)</pre>
```

### Interpretation/Evaluation

Boosting with XGBoost performs the best.

LASSO keeps many variables. The coefficients represent changes in the log odds ratio but analysts are discouraged from making inferences about them; it is a known feature of penalized regressions that correlated covariates may be included in the final model without having an underlying causal effect on the dependent variable.

The random forests model uses a generic 500 trees. sig2 is the most important feature. When sig2 is permuted, the model's accuracy drops the most compared to any other predictor, so the forest relies heavily on it for splitting and prediction.

These results generally concur with the earlier visualizations. sig2 appeared to uniquely partition dependent and independent variables.

Below is a table summarizing results, followed by visualizations of each modeling strategy's variable selection methods and performance.

#### Model Performance Comparison

CV Accuracy	Optimal Parameters
0.66605	$\lambda = 1e-04$
0.66600	k = 149
0.66614	mtry = 3
0.66737	iterations = 46
	0.66605 0.66600 0.66614

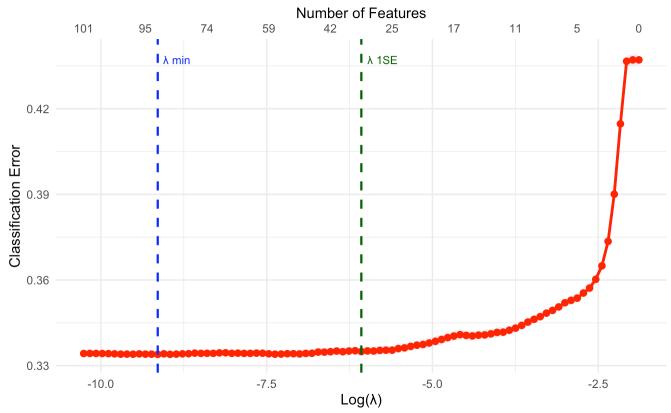
#### LASSO

Head of LASSO Selected Features (91 variables + intercept)

Feature	Coefficient
sig2	2.8803
is_homepage1:sig7	-1.7308
(Intercept)	-1.5162
sig1:sig7	-1.4295
sig7:sig8	-1.3312
is_homepage1:sig1	1.3215

#### **LASSO Cross-Validation**

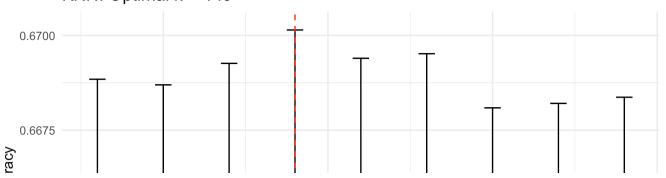
Min  $\lambda$  = 1e-04 | 1SE  $\lambda$  = 0.0023

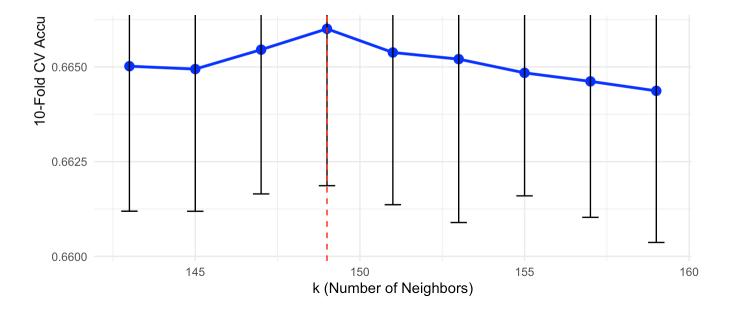


Features at min: 93 | Features at 1SE: 30

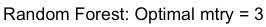
# KNN

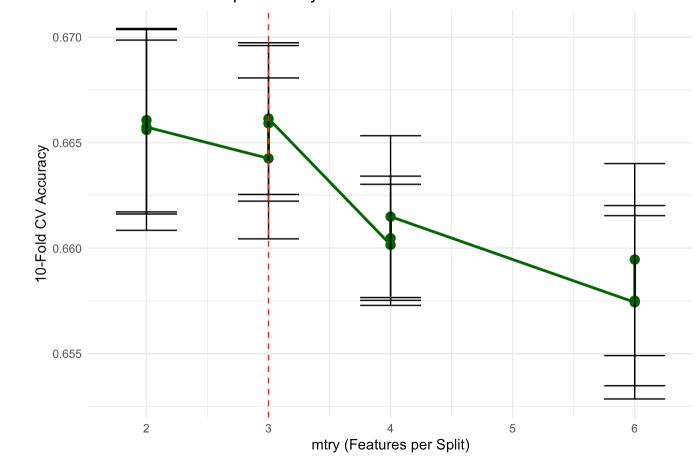
KNN: Optimal k = 149





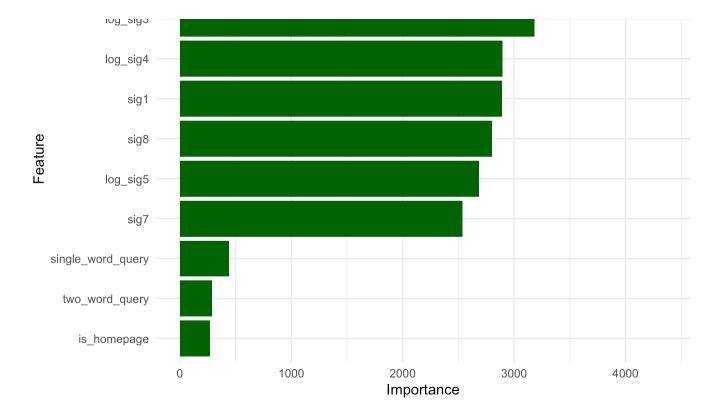
# **Random Forests**



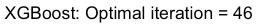


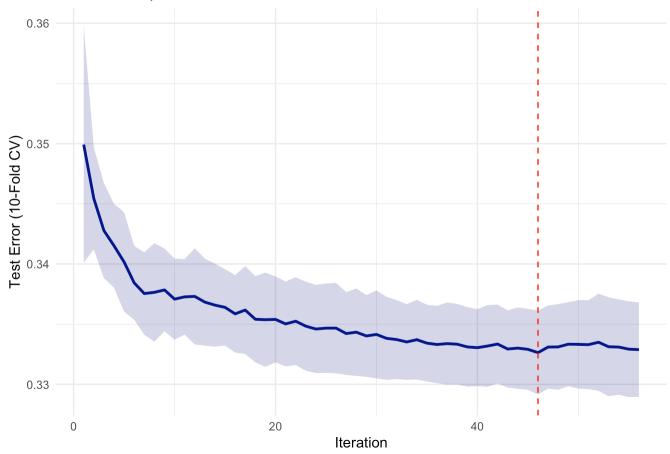
Top 10 Important Features

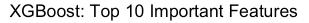


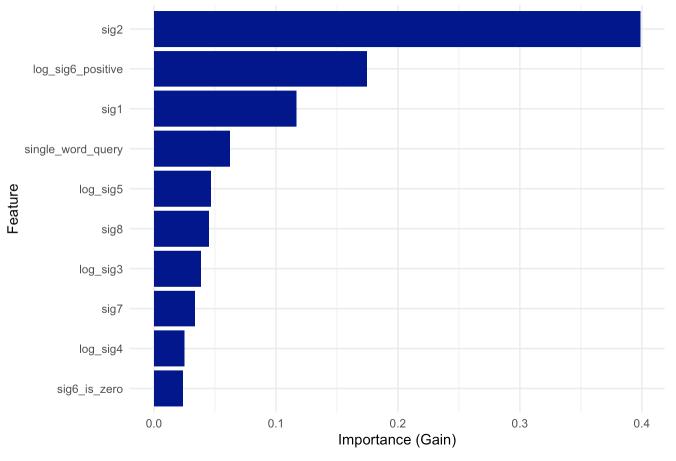


# XGBoost









XGBoost CV Accuracy: 0.6674

# **Testing**

A script for creating the test submissions is included in the GitHub Repository: https://github.com/24-bee-supply/stats-202-project.