

**CSE 535**

# **Project 3 : Evaluation of IR Models**

**Information Retrieval**

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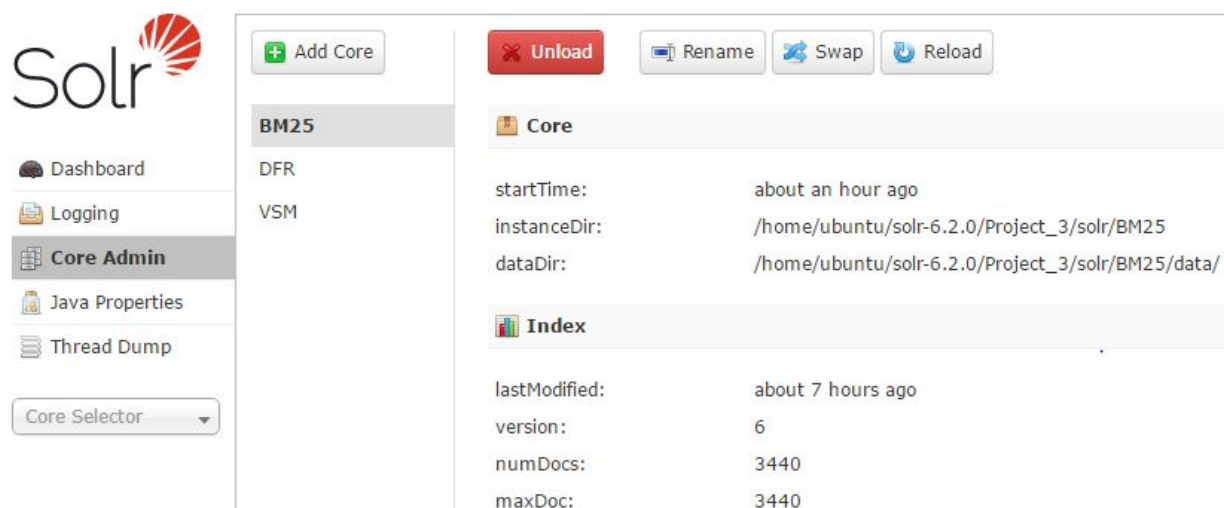
## OVERVIEW

The objective of this IR project is to implement the three IR models namely Vector Space Model, BM25 and Divergence from Randomness model using Solr, evaluate the IR system and improve the search result based on the understanding of the models.

- ❑ **BM25** : The Best Matching (BM25) algorithm is a probabilistic Information Retrieval (IR) model. Its often called Okapi Weighting, which is a ranking function in Information Retrieval used by search engines to rank matching documents according to their relevance to given search query.
- ❑ **Vector Space Model** : Also known as Term Vector Model, is an algebraic model used for representing documents and queries as vectors in the term space. It allows computing a continuous degree of similarity between queries and documents, ranking documents according to their possible relevance, partial matching.
- ❑ **Divergence from Randomness** : The Divergence from Randomness (DFR) paradigm is a generalisation of one of the very first models of Information Retrieval, Harter's 2-Poisson indexing-model. It is based on the following components : Randomness Model, First Normalization and Term Frequency Normalization.

We successfully implemented the 3 models and created 3 cores for the same. Changes were made as we made progress on improvisation of the model

Please see the snapshot of the Solr UI displaying the 3 core named according to their models.



The screenshot displays the Solr Admin interface. On the left is a sidebar with navigation links: Dashboard, Logging, Core Admin (selected), Java Properties, and Thread Dump. Below these is a 'Core Selector' dropdown. The main content area shows the configuration for the 'BM25' core. At the top, there are buttons for '+ Add Core', 'Unload', 'Rename', 'Swap', and 'Reload'. The 'Core' section lists: startTime: about an hour ago, instanceDir: /home/ubuntu/solr-6.2.0/Project\_3/solr/BM25, and dataDir: /home/ubuntu/solr-6.2.0/Project\_3/solr/BM25/data/. The 'Index' section lists: lastModified: about 7 hours ago, version: 6, numDocs: 3440, and maxDoc: 3440.

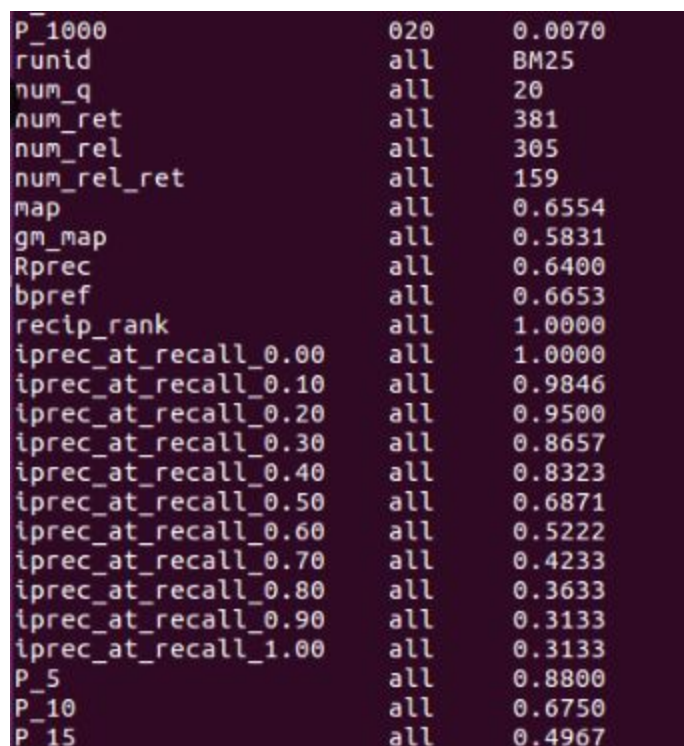
## BM25 Model

**Initial Implementation** : This is the default model. In order to implement the model changes were made in the schema.xml file by inserting the following statement.

```
<similarity class="solr.BM25SimilarityFactory">
```

On calculating the MAP value through trec\_eval the result received was **0.6554**.

Please find below the screenshot for the same:



```
P_1000      020      0.0070
runid       all      BM25
num_q       all      20
num_ret     all      381
num_rel     all      305
num_rel_ret all      159
map         all      0.6554
gm_map      all      0.5831
Rprec       all      0.6400
bpref       all      0.6653
recip_rank  all      1.0000
iprec_at_recall_0.00 all 1.0000
iprec_at_recall_0.10 all 0.9846
iprec_at_recall_0.20 all 0.9500
iprec_at_recall_0.30 all 0.8657
iprec_at_recall_0.40 all 0.8323
iprec_at_recall_0.50 all 0.6871
iprec_at_recall_0.60 all 0.5222
iprec_at_recall_0.70 all 0.4233
iprec_at_recall_0.80 all 0.3633
iprec_at_recall_0.90 all 0.3133
iprec_at_recall_1.00 all 0.3133
P_5         all      0.8800
P_10        all      0.6750
P_15        all      0.4967
```

**Improvement** : In order to get a better MAP value , we played around the values of the solr.BM25SimilarityFactory class in order to get the improved score.

**K<sub>1</sub>**: This parameter controls how quickly an increase in term frequency results in term-frequency saturation. **The default value is 1.2** Lower values result in quicker saturation, and higher values in slower saturation.

**b**: This parameter controls how much effect field-length normalization should have. **The default is 0.75**.

Some of the value pair used in order to end up to a final value

- $(k_1, b) = (1.0, 1.0) \Rightarrow \text{MAP} = 0.6363$
- $(k_1, b) = (1.0, 0.7) \Rightarrow \text{MAP} = 0.6563$
- $(k_1, b) = (1.25, 0.7) \Rightarrow \text{MAP} = 0.6573$

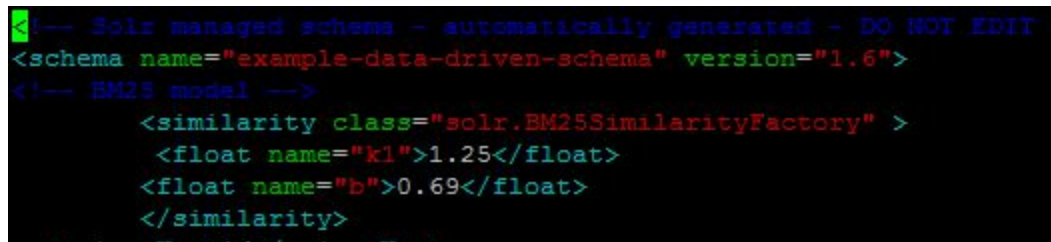
#### Final value:

$k_1 = 1.25$

$b = 0.69$

**MAP = 0.6871**

Below are the changes in the schema.xml file



```

<!-- Solr managed schema - automatically generated - DO NOT EDIT -->
<schema name="example-data-driven-schema" version="1.6">
  <!-- BM25 model -->
    <similarity class="solr.BM25SimilarityFactory" >
      <float name="k1">1.25</float>
      <float name="b">0.69</float>
    </similarity>

```

On making further changes in values of  $b$  and  $k_1$  there is negligible change in the MAP value so we decided to make changes in the **schema.xml** file by adding those filter classes that would enhance the performance and in turn the MAP value.

Please see the screenshot below :

```

<fieldType name="text_de" class="solr.TextField" positionIncrementGap="50">
  <analyzer>
    <tokenizer class="solr.StandardTokenizerFactory"/>
    <filter class="solr.LowerCaseFilterFactory"/>
    <filter class="solr.StopFilterFactory" format="snowball" words="lang/stopwords_de.txt" ignoreCase="true"/>
    <filter class="solr.GermanNormalizationFilterFactory"/>
    <filter class="solr.GermanLightStemFilterFactory"/>
    <filter class="solr.PorterStemFilterFactory"/><!-- Added to increase the MAP value -->
  </analyzer>
  <filter class="solr.BeiderMorseFilterFactory" nameType="GENERIC" ruleType="APPROX" concat="true" languageSet="auto"/><!-- A
  <filter class="solr.NGramFilterFactory" minGramSize="3" maxGramSize="15"/><!-- Added to increase the MAP value -->
  <filter class="solr.EnglishMinimalStemFilterFactory"/>
  </analyzer>
</fieldType>

```

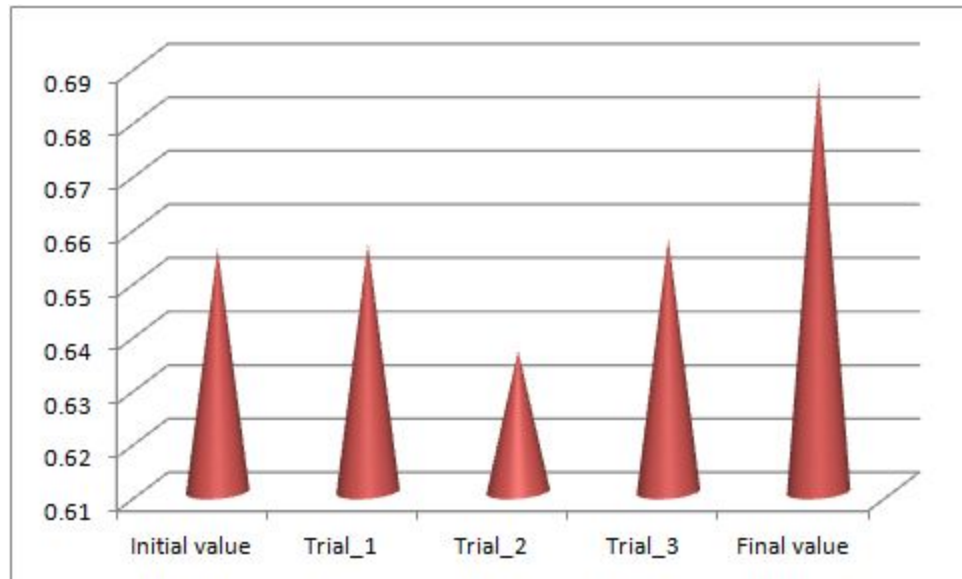
We have added following Filter classes in schema.xml:

- **Porter Stem Filter** - This filter applies Porter Stem Algorithm for English, it returns the root word while dropping the suffix(es). It has been benchmarked as four times faster than the English Snowball stemmer, so can provide a performance enhancement.
- **Beider Morse Filter** - This filter allows identification of similar names, even if they are spelled differently or in different languages.
- **N Gram Filter** - It generates n-gram tokens of sizes in the given range of minGramSize and maxGramSize.

After doing the above changes we successfully landed to an improved MAP value.

P_1000	all	0.0070
runid	all	BM25
num_q	all	20
num_ret	all	651
num_rel	all	305
num_rel ret	all	180
map	all	0.6871
gm_map	all	0.6133
Rprec	all	0.6565
bpref	all	0.7086
recip_rank	all	1.0000
iprec_at_recall_0.00	all	1.0000
iprec_at_recall_0.10	all	0.9846
iprec_at_recall_0.20	all	0.9500
iprec_at_recall_0.30	all	0.8657
iprec_at_recall_0.40	all	0.8315
iprec_at_recall_0.50	all	0.7190
iprec_at_recall_0.60	all	0.5928
iprec_at_recall_0.70	all	0.5125
iprec_at_recall_0.80	all	0.4844
iprec_at_recall_0.90	all	0.3312
iprec_at_recall_1.00	all	0.3312
P_5	all	0.8800
P_10	all	0.6750
P_15	all	0.5000
P_20	all	0.3975
P_30	all	0.2900
P_100	all	0.0900
P_200	all	0.0450
P_500	all	0.0180
P_1000	all	0.0080

The Graphical Analysis of BM25 model is :



## Divergence From Randomness (DFR model)

There are three components of DFR (strings).

**Initial Implementation:** To implement this model in the default mode , a similarity class was added to the schema.xml file as under with the default values with respect to the information given in the project guidelines.

```
<!-- DFR model -->
    <similarity class="solr.DFRSimilarityFactory">
        <str name="basicModel">G</str>
        <str name="afterEffect">B</str>
        <str name="normalization">H2</str>
    </similarity>
```

On calculating the MAP value through trec\_eval the result received was **0.6468**

**Improvement** : In order to increase the map score , there was need to change the values of the 3 strings.

We have to put in the right combination as much possible of **basicModel** ,**afterEffect** and **normalization** in order to improve the score.



We finally settled for the following :

- 1.basicModel:**Be**: Limiting form of Bose-Einstein
2. afterEffect:**B**: Ratio of two Bernoulli processes
- 3.normalization: **H2**: term frequency density inversely related to length

```

<!-- Solr managed schema - automatically generated - DO NOT EDIT -->
<schema name="example-data-driven-schema" version="1.6">
  <!-- DFR model -->
  <similarity class="solr.DFRSimilarityFactory">
    <str name="basicModel">Be</str>
    <str name="afterEffect">B</str>
    <str name="normalization">H2</str>
    <float name="c">1000</float>
  </similarity>

```

With Improved MAP value =**0.6965**

Some of the trial values used in order to end up to a final value were :

- (basicModel , afterEffect ,normalization) => (D: Divergence approximation of the Binomial,B: Ratio of two Bernoulli processes, H2: term frequency density inversely related to length) MAP =0.6822
- (basicModel , afterEffect ,normalization) =>(P: Poisson approximation of the Binomial, B: Ratio of two Bernoulli processes,H3:term frequency normalization provided by Dirichlet prior) MAP=0.6420
- (basicModel , afterEffect ,normalization)=> (Be: Limiting form of Bose-Einstein,L: Laplace's law of succession,H2: term frequency density inversely related to length) MAP=0.6894

Along with the additions in schema.xml ( similar to that for BM25 model) we also modified the query and added The DisMax query parser to improve the score. It is designed to process simple phrases entered by users and to search for individual terms across several fields using different weighting (boosts) based on the significance of each field.

<http://54.68.99.12:8983/solr/DFR/select?q=%27+qText+%27&fl=id%2Cscore&wt=json&indent=true&rows=20&defType=dismax>

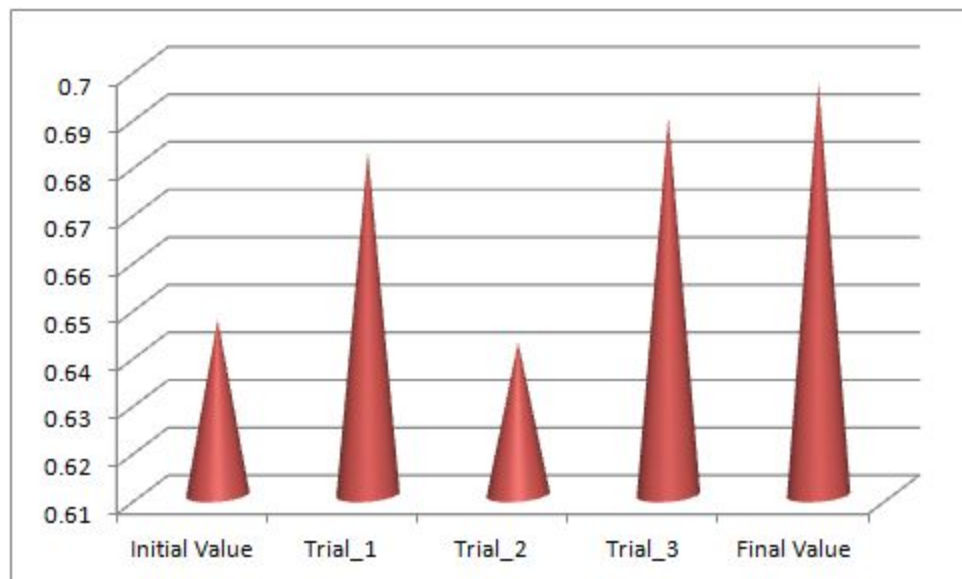
```

</fieldType>
<fieldType name="text_de" class="solr.TextField" positionIncrementGap="100">
  <analyzer>
    <tokenizer class="solr.StandardTokenizerFactory"/>
    <filter class="solr.LowerCaseFilterFactory"/>
    <filter class="solr.StopFilterFactory" format="snowball" words="lang/stopwords_de.txt" ignoreCase="true"/>
    <filter class="solr.GermanNormalizationFilterFactory"/>
    <filter class="solr.GermanLightStemFilterFactory"/>
    <filter class="solr.PorterStemFilterFactory"/><!-- Added to increase the MAP value -->
  <filter class="solr.BeiderMorseFilterFactory" nameType="GENERIC" ruleType="APPROX" concat="true" languageSet="auto"></filter><!-- A
    <filter class="solr.NGramFilterFactory" minGramSize="3" maxGramSize="15"/><!-- Added to increase the MAP value -->
  </analyzer>
</fieldType>

```

Similar changes were made to text\_en and text\_ru.

**The Graphical Analysis of DFR model is :**



## Vector Space Model(VSM)

The vector space model procedure can be divided into three stages.

The first stage is the document indexing where content bearing terms are extracted from the document text.

The second stage is the weighting of the indexed terms to enhance retrieval of document relevant to the user.

The last stage ranks the document with respect to the query according to a similarity measure.

Similarity Class for VSM : **ClassicSimilarityFactory** was added to schema.xml



On running the default settings for VSM the result obtained for MAP value was **0.6469**.

```

P_100      020      0.0700
P_200      020      0.0350
P_500      020      0.0140
P_1000     020      0.0070
ruid       all      VSM
num_q      all      20
num_ret    all      381
num_rel    all      305
num_rel_ret all      154
map        all      0.6469
gm_map     all      0.5739
Rprec      all      0.6573
bpref      all      0.6578
rectp_rank all      1.0000
iprec_at_recall_0.00 all 1.0000
iprec_at_recall_0.10 all 0.9846
iprec_at_recall_0.20 all 0.9464
iprec_at_recall_0.30 all 0.8561
iprec_at_recall_0.40 all 0.8373
iprec_at_recall_0.50 all 0.6477
iprec_at_recall_0.60 all 0.5020
iprec_at_recall_0.70 all 0.4064
iprec_at_recall_0.80 all 0.3714
iprec_at_recall_0.90 all 0.3231
iprec_at_recall_1.00 all 0.3231
P_5        all      0.8600
P_10       all      0.6600
P_15       all      0.4900
P_20       all      0.3850
P_30       all      0.2567
P_100      all      0.0770
P_200      all      0.0385
P_500      all      0.0154
P_1000     all      0.0077

```

In order to improve the performance of this model we added some more stopwords to **stopwords\_en.txt**, **stopwords\_de.txt**, **stopwords\_ru.txt** so as to get better precision and recall values. Also we added a list of synonyms of common words to **synonyms.txt** so as to get better MAP result . These two changes can be seen in the screenshot below:

```

<fieldType name="text_en" class="solr.TextField" positionIncrementGap="100">
  <analyzer type="index">
    <tokenizer class="solr.StandardTokenizerFactory"/>
    <filter class="solr.StopFilterFactory" words="lang/stopwords_en.txt" ignoreCase="true"/>
    <filter class="solr.LowerCaseFilterFactory"/>
    <filter class="solr.EnglishPossessiveFilterFactory"/>
    <filter class="solr.KeywordMarkerFilterFactory" protected="protwords.txt"/>
    <filter class="solr.SynonymFilterFactory" synonyms="synonyms.txt"/>
  <!-- Added to increase the MAP value -->
  <filter class="solr.BeiderMorseFilterFactory" nameType="GENERIC" ruleType="APPROX" concat="true" languageSet="auto"/></filter><!-- A
  </analyzer>
<analyzer type="index"><!-- Added to increase MAP value-->
  <tokenizer class="solr.WhitespaceTokenizerFactory"/>
  <filter class="solr.HyphenatedWordsFilterFactory"/>
</analyzer>

```

Following Filter classes have been added:

- ❑ **Synonym Filter** - Each token is looked up in the list of synonyms and if a match is found, then the synonym is emitted in place of the token. The position value of the new tokens are set such they all occur at the same position as the original token.

- ❑ **Hyphenated Words Filter** - This filter is used to use hyphenated terms as one single terms in order to provide better matching results at the time of query searching and matching.

Both these filters have been added to provide better MAP results in the VSM model and it is done so by performing better matching of terms in the document with the query terms.

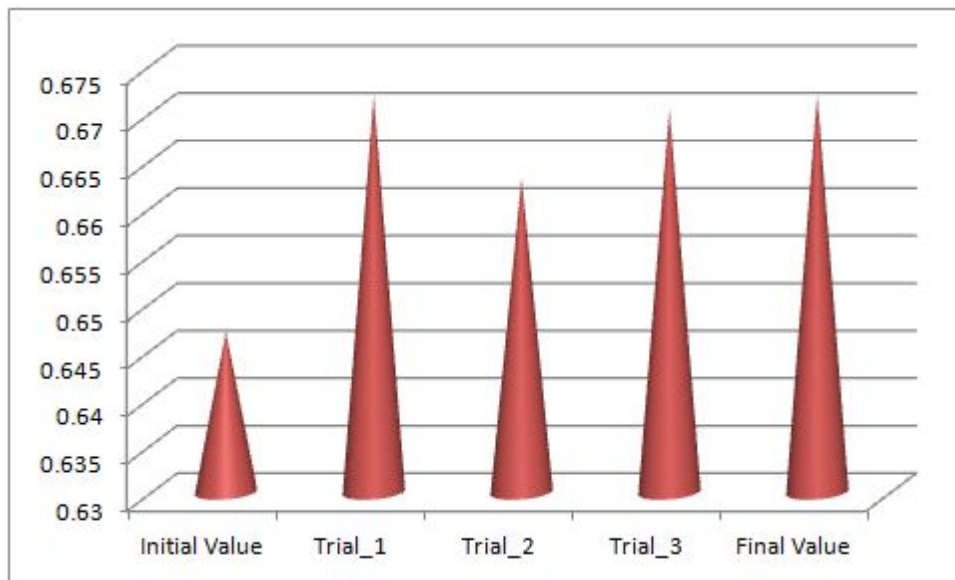
Before arriving to the final MAP value we had some intermediated values as

- MAP=0.6728
- MAP=0.6631
- MAP=0.6704

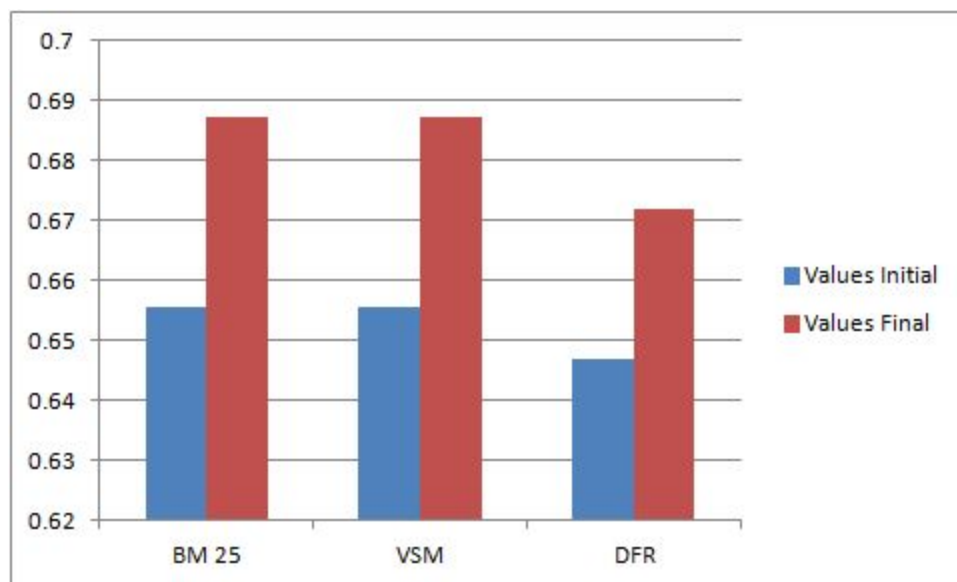
The final improved MAP value is **0.6719** which can be seen in the screenshot below:

num_ret	all	561
num_rel	all	305
num_rel_ret	all	170
map	all	0.6719
gm_map	all	0.5978
rprec	all	0.6738
bpref	all	0.6903
recip_rank	all	1.0000
iprec_at_recall_0.00	all	1.0000
iprec_at_recall_0.10	all	0.9846
iprec_at_recall_0.20	all	0.9464
iprec_at_recall_0.30	all	0.8578
iprec_at_recall_0.40	all	0.8391
iprec_at_recall_0.50	all	0.6682
iprec_at_recall_0.60	all	0.5570
iprec_at_recall_0.70	all	0.4926
iprec_at_recall_0.80	all	0.4445
iprec_at_recall_0.90	all	0.3435
iprec_at_recall_1.00	all	0.3435
P_5	all	0.8600
P_10	all	0.6600
P_15	all	0.4900
P_20	all	0.3925
P_30	all	0.2833
P_100	all	0.0850
P_200	all	0.0425
P_500	all	0.0170
P_1000	all	0.0085

The Graphical Analysis of VSM model is :



### Collective Graphical Analysis of the three models



## Conclusion

The three IR models have been implemented successfully and we tried to improve the performance of all the three models by using their properties and the features provided by Solr 6.2.0.

In BM25 model we settled for values of  $b$  and  $k_1$  as 0.69 and 1.25 respectively, as better results were achieved through these values and also by adding N gram filter, Beider Morse filter and Porter Stem Filter.

In DFR model with the help of the 3 components we improved the MAP value along with some changes in the schema.xml.

In VSM model we added Synonym Filter, Hyphenated Filter as well added more stopwords to the stopwords.txt file of all three language fields.

All these provided some improvements in the system but there is a lot more scope of improvement to do in future prospects.