

Dissertation Critique: Exploring Machine Learning Techniques Using Patient Interactions In Online Health Forums to Classify Drug Safety

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Abstract

Patient generated health data represents an area of active research interest for its potential applications in improving the public health. The study of Pharmacovigilance is one such area, focused on monitoring drugs once they have been released to market. Dr. Brant Chee's 2011 dissertation applying machine learning techniques to patient messages in on-line health forums explores how watch list drugs from the United States Food and Drug Administration can be detected via these forum messages, ultimately with the intent to alert consumers to drug safety concerns.

Keywords: Drug Safety, Pharmacovigilance, NLP

1. Summary of Research

Dr. Brant Chee's 2011 dissertation *Exploring Machine Learning Techniques Using Patient Interactions in Online Health Forums to Classify Drug Safety* describes Chee's research in applying natural language processing (NLP) techniques in conjunction with Naive Bayes and Support Vector Machine classifiers to identify candidate *watch list* drugs from online patient forums. Watch list drugs are those drugs identified by the United States Food and Drug Administration (FDA) as presenting a significant health or safety risk to drug consumers, thereby prompting regulatory action to better inform the consumer or directly protect the consumer by removing the drug from market or reducing its accessibility. Chee's dissertation seeks to answer the specific questions:

- Can Machine Learning classification methods using text features extracted from online health forums be used to identify FDA watch list drugs?
- Is the sentiment of the forum message useful in identifying these drugs?
- Similarly, are the drug effect entities useful in identifying watchlist drugs?

This research is accomplished through an empirical study using a corpus from the Yahoo! public health forums, against which Chee applies various NLP techniques to define and distill a feature space for classification using Naive Bayes and Support Vector machines for

detecting watchlist drugs. Drugs detected are evaluated against watchlist drugs found via the FDA Adverse Event Reporting System (AERS) to determine the utility of the approach and its applicability in Pharmacovigilance.

1.1 Background on Pharmacovigilance, AERS and Social Media

The dissertation begins with an extensive background discussion on adverse drug reactions and current surveillance techniques. Adverse drug reactions are defined by the FDA and World Health Organization (WHO) as "A response to a drug which is noxious and unintended and which occurs at doses normally used in man for prophylaxis, diagnosis, or therapy of disease or for modification of physiological function."?. Chee continues by introducing Pharmacovigilance as "the study of drugs once released to market" Chee, and the important regulatory agencies practicing it are mentioned - the World Health Organization (WHO) and United States Food and FDA. The FDA Adverse Event Reporting system (AERS) is discussed as comparison with it is central to the work. AERS was constructed to house mandatory drug safety reports from drug manufacturers, distributors and health care facilities, as well as voluntary reports submitted by consumers (patients), physicians and other healthcare providers. Reports are evaluated by the Center for Drug Evaluation and Research (CDER) and Center for Biologics Evaluation and Research (CBER) within the FDA for drug safety signals, which may then be elevated for further review by clinicians, epidemiologists and other expertise to determine the next steps, up to and including the removal of a drug from the market.

Chee identifies a major limitation in AERS and other *spontaneous reporting systems* in that they are known to have high underreporting rates (?), due to the likelihood of a patient reporting an event only if they feel their healthcare provider has not paid attention to the adverse drug reaction observed (?). This deficiency is presented as motivation for Chee's work exploring social media as a data source. Social media provides a venue for patients to share their health information in anonymous setting as patients are not always transparent nor truthful with their physicians. Online health forums create an environment where patients can find those having similar backgrounds, conditions and challenges, which in turn prompt rich social interactions where patient disclose their opinions and observations about their current drug regimen effectiveness and perceived adverse events. Chee feels these forums represent an untapped means to crowdsource data for the pharmacovigilance task.

1.2 Experimental Data

The data selected for the dissertation's experimentation is a Yahoo! corpus containing 12.5 million messages from various Yahoo Health group forums. As the data is a raw export containing a combination of message metadata, raw text and HTML, it must first be studied to better understand its composition and what NLP techniques should be applied to better prepare it for experimentation.

1.2.1 TOKENIZATION STUDY OF DATA

Chee conducts an initial study of the Yahoo! corpus by selecting at random 500 messages, stripping them of html tags, numerical and punctuation only tokens (\$, %, :), :(, etc), then tokenizing them on white spaces with trailing punctuation. The tokens are then evaluated

in several rounds of classification using lexicons obtained or constructed by Chee to aid in understanding message composition. Lexicons for English and foreign language were drawn from the OpenOffice project (?). Drug names were taken from the Drugs@FDA website. A medical and disease terminology lexicon was constructed from terms on MedicineNet, Wikipedia, and the MedDRA lexicon from FDA AERS. The names lexicon was constructed using names extracted from email addresses in message headers in the corpus, popular baby names from the United States (US) Social Security Administration, and popular common names from the US 1990 Census.

The classification process was iterative. If a token did not initially classify as English, web (a lexicon of web slang), medical, or drug name it was manually inspected and classified into *error types*: Foreign Language, Names (augmenting name lexicon), Spelling Errors, Compound Words, Slang, Abbreviations, Web, Unknown Words, Numbers and Garbage. Chee’s analysis produced some interesting average metrics for the messages:

Average # of Tokens per Messsage:	172.21
Average # of Drug Name Tokens:	.29
Average # of Error Type Tokens:	7.09
Average # of Name Toknes:	5.34
Average # of Medical Tokens:	.81

Of the error tokens, over 54% of them were found to be foreign language tokens - primarily Indonesian and Spanish - motivating Chee to incorporate Foreign Language lexicons from OpenOffice to speed up classification. A primary concern for Chee was the presence of spelling errors, but the classification results show only a .8% error rate, which Chee uses to rationalize sticking with dictionary based approaches for word classification for their high precision. Finally, it is acknowledged that Named Entity Recognition (NER) is challenging in this context. FDA approved drugs represent a closed class of nominals, but foreign drugs, herbs and other chemicals are not available in a comprehensive list. Dictionary approaches to classifying drug outcomes are challenged by the use of slang terms.

1.2.2 A VOCABULARY FOR EXPERIMENTATION

Chee describes performance concerns training SVMs for classification using all the words in the message, given the $O(kn)$ training time for n training instances using k features (words). Additionally, multiple words together in order can convey a different meaning than separate single words, such as "vitamin a" compared to "vitamin" and "a".

These constraints motivate the use of word-grams - unigrams, bigrams and trigrams specifically - as a way to capture more accurate meaning. Chee references (??)’s work proposing the most informative words in a message would be the mid-frequently occurring ones, electing to take the top $k - n$ most frequently occurring terms in a message as the most important terms, where k is the top number of terms, minus n accounting for simple function words like *a*, *or* and *the*.

Specialized lexicons are developed as a means to ensure the classifiers that will be trained do not overfit to only those drugs in the drug lexicon, preventing the identification of previously unseen (unlabeled) watchlist drugs. The lexicons selected to use in the classification are:

- drugs - a drug list from drugs.com
- medical - medical terminology extracted from MedicineNet
- sentiment - a sentiment lexicon from the combination of SentiWordNet and Linguistic Inquiry and Word Count (LIWC)
- medra - the MedDRA terms for drug adverse events/outcomes from FDA AERS
- disease - a disease list from Wikipedia

These five lexicons allow for twenty-nine different datasets of features to be constructed from the Yahoo! corpus messages for the watchlist drug classification experiments. The feature vector used in classification is then the intersection of those terms found in *all* of the lexicons used in that particular test. For example, if the sentiment and drug lexicons are used together, the feature vector has only those terms that occur in both lexicons.

1.3 Language Identification for Messages

The previous study identified a significant number of foreign language messages in the corpus. While these text processing techniques are language agnostic, removing the foreign language messages will reduce the feature vector length for training, as well as acknowledge the audience for this study is English speaking.

Messages containing non-romanized text are removed first using Unicode language detection. Since non-English languages in romanized text are harder to discern, Chee compares and contrasts character n-gram approaches by (??) with dictionary approaches from (?) for foreign language classification. Dictionaries are opted for given the simple, binary nature of the problem: Is a message English or not? Dictionaries for foreign language words are taken from the OpenOffice project and used in conjunction with the medical, drug, disease and name lexicons mentioned earlier are used to evaluate a linear inequality for each message to determine if it will be kept or not.

First the messages are stripped of tokens containing web addresses, punctuation only (emoticons), as well as short words, email addresses or tokens that are already on an ignore list. Remaining tokens in each message are counted up using the following algorithm:

- if (word in ignore) OR (word length ≤ 2) OR (word contains "@") ++ignore
- else if (word in English) ++english
- else if (word in Drug) ++drug
- else if (word in medical) ++medical
- else if (word in name) ++name
- else if (word in foreign) ++foreign
- else unknown++

Table 1:

Once the counts are obtained, the following linear inequality is evaluated:

$$4 * \textit{foreign} + \textit{unknown} + \textit{ignore} > \textit{english} + \textit{drugs} + \textit{medical}$$

The weighting of the foreign words is selected to ensure messages contain less than 25% foreign words to be considered english. Foreign messages are removed and English messages retained. This resulted in a reduction from 12,520,438 messages to 10,178,710 messages, and a reduction in the number of unique terms per message from 2.5 to 2.

1.4 Experimentation and Results

The goal of the dissertation is to develop a classification system for drugs based on how people are talking about them in online message forums. Processed forum messages are first organized by drug, divided into test and training sets, converted into feature vectors, and then run through Support Vector Machine and Naive Bayes classification algorithms. Chee conducts a multitude of these experiments using two versions of the feature vector structure, multiple combinations of the lexicons described earlier, usage of top $n - k$ terms and BNS for term selection to feed feature identification using the lexicons, as well as cost-weighted and unweighted variants of Naive Bayes and SVMs.

1.4.1 CLASS SEPARABILITY CONFIRMED VIA KULLBACK-LEIBLER DIVERGENCE

Chee begins with an initial experiment to determine how separable conversations regarding watchlist and non-watchlist drugs will be in order to validate the classification approach. Kullback-Leibler divergence (KL divergence) is used to measure the difference in word frequency distribution between messages in each category, with a smoothing technique applied to make sure each term is represented - albeit with very low frequency - in each distribution to prevent infinite divergence. A series of comparisons are done between the watchlist and non-watchlist frequencies, as well as term frequencies in the Google Web 1T 5-gram corpus and Reuters Corpus to provide perspective.

- $D_{jk}(\textit{Watchlist}||\textit{Non}) = .1684$
- $D_{jk}(\textit{Non}||\textit{Watchlist}) = .1778$
- $D_{jk}(\textit{Watchlist}||\textit{Google}) = 1.4178$
- $D_{jk}(\textit{Watchlist}||\textit{Reuters}) = 1.3279$
- $D_{jk}(\textit{Non}||\textit{Google}) = 1.1804$
- $D_{jk}(\textit{Non}||\textit{Reuters}) = 1.0815$

Clearly Watchlist Nonwatchlist diverge, and show a different degree of divergence from the Google and Reuters corpi, indicating some separability.

Interesting term frequency differences:

Table 2:

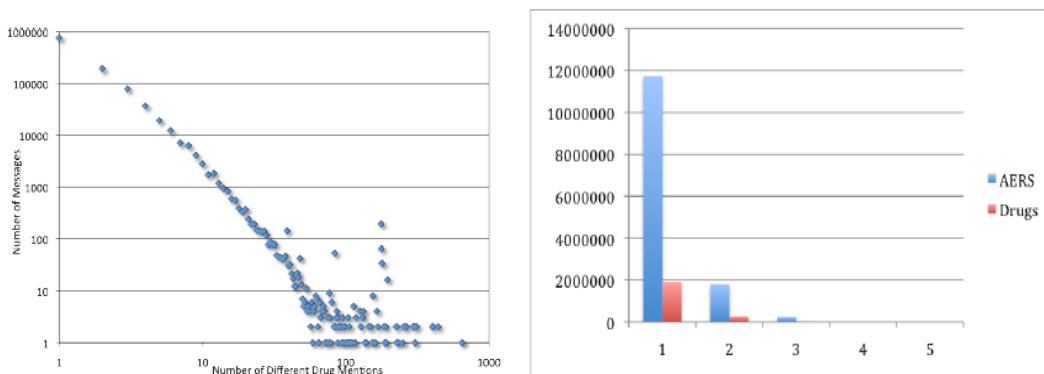
- i $D_{kl}(Watchlist||Non) = .007875$
- my $D_{kl}(Watchlist||Non) = .003018$
- me $D_{kl}(Watchlist||Non) = .0002256$
- you $D_{kl}(Watchlist||Non) = .001712$
- i'm $D_{kl}(Watchlist||Non) = 9.64E - 04$

The above terms are overexpressed in the watchlist messages compared to non-watchlist, and are considered indicative of emotional writing - just the kind of writing that might occur when discussing a drug with an adverse effect. Chee concludes these results support separability of the two classes of message based on their word features.

1.4.2 NAMED ENTITY RECOGNITION IN MESSAGES

Another challenge presented by Chee is how to identify the drug(s) and adverse event(s) themselves within a message - a problem of Named Entity Recognition (NER). A dictionary based approach using a drug lexicon compiled from FDA and Drugs.com, and adverse event lexicon using the Medical Dictionary for Regulatory Activities (MedDRA) is used to query a Lucene index built atop the processed Yahoo! corpus messages. The index construction applied a lowercase filter with stemming, presenting a problem with common words that are also part of drug names. For example, the drug name *Commit* is indistinguishable from the verb *commit*. This was addressed by replacing drug names with common words by their generic (chemical) name in the lexicon to favor precision of query results over recall.

A series of phrase searches were issued against the index to determine prevalence of MedDRA terms (adverse events) and drug names. Two important conclusions are drawn thanks to the resultant plots:



The first plot demonstrates a zipfian distribution for drug name mentions in the messages, with more than 96% of messages mentioning at most 5 drugs, adding confidence to the hypothesis that messages can be associated with at most a handful of drugs. Chee elects to eliminate messages having more than 5 unique drugs as these frequently constitute lists of drugs posted or SPAM messages. The second plot shows that adverse events are mentioned

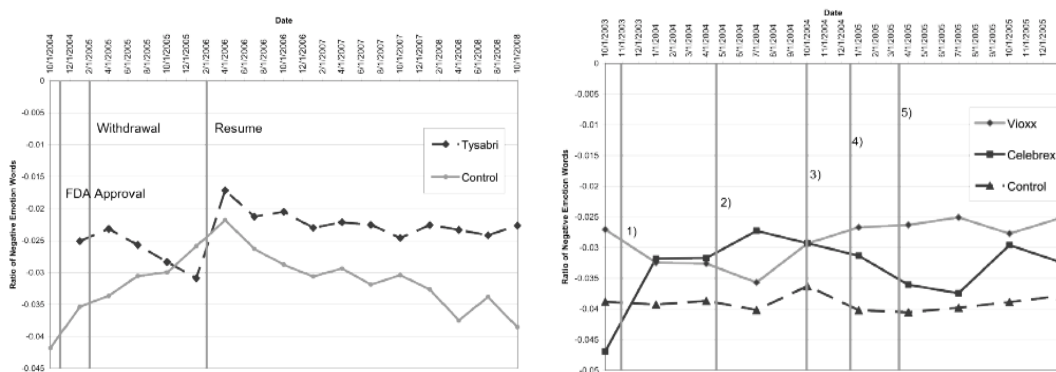
far more often than drug names, suggesting that a preceding post containing a drug name eliminates referencing it in subsequent posts - much like replies on a forum. Investigating this is an area for future research. Note that the numbers on the X-axis for the 2nd plot refer to the number of terms (1, 2, ...) that comprise the drug name or adverse event - a fact not really relevant for this analysis.

1.4.3 HANDLING MULTIPLE DRUG NAMES

If multiple drugs are mentioned in a single message, it becomes difficult to discern which drug the message should apply to. Another study is conducted to evaluate the typical distance (number of characters) between the top 25 co-occurring drugs in messages to determine if the messages could be segmented into relevant portions for each drug. The analysis shows a Zipfian distribution in the character separation, indicating that most drugs are talked about together within a single sentence, or in adjacent sentences, leading Chee to conclude separation is not possible and that any adverse events mentioned in a message should just be attributed to all drugs mentioned within the message.

1.4.4 SENTIMENT FEATURE

A final experiment preceeding the main classification work is done to validate how message sentiment will be determined. Chee hypothesizes that the positive or negative valence in a message represents drug satisfaction, making it an interesting feature to incorporate into the classification experiments. A lexicon using the positive emotion, negative emotion, anxiety, anger and sadness terms from LIWC is constructed, augmented to include several emoticons (:), :(, ..) and acronyms (LOL, ROFL, ..). Two case studies are then executed using messages sampled from specific groups in the corpus, and the change in drug sentiment is analyzed over the drug's pre-recall, recall, and post-recall timeframes. This sentiment is compared with a control sentiment derived from those messages in the samples not containing the drug to look for a statistically significant difference.



The left-most plot shows the sentiment change for Tysabri pre-recall, recall (withdrawal) and post-recall (resume) over the control. It shows a reasonably intuitive change in negative valence to Tysabri having been introduced (more positive), withdrawn (negative), reintroduced (hopeful therefore positive), then stabilizing. The right plot shows sentiment change for a pair of commonly used pain relievers - Vioxx and Celebrex - over the course of several public announcements and a withdrawal of Vioxx (sections 1 through 3), then Celebrex (4

and 5). ANOVA is applied to both case studies to determine statistical significance for each drug in each segment against the control. Both were found to be statistically significant with $p < .001$.

1.4.5 FEATURE SELECTION, TRAINING AND TEST DATA SIZE

The introduction to the main body of classification experiments is preceeded by a brief discussion on the features vectors used, as well as how training, test and validation sets are constructed to support the use of 10-fold Cross Validation for classifier evaluation.

Two types of feature vector are decided upon and then leveraged in the experiments. The first feature vector type is generated over general vocabulary terms in the messages, selected based on frequency cutoffs. This vector is then augmented with counts from the various specialized lexicons mentioned earlier: medical, diseases, drugs, sentiment and reactions (MedRA). The second feature vector uses only the specialized lexicon.

The richness of the training data is of foremost concern to Chee. There are only 435 drugs having 500 or more unique messages, and only 575 drugs having more than 250 messages, with 63 and 77 watchlist drugs mentioned in each respectively. Therefore approximately 90% of message instances reference non-watchlist drugs, creating a data scarcity problem when attempting to classify watchlist drugs. Chee decides upon a minimum cutoff of 250 messages per drug for that drug's messages to be included in training and testing. An experiment is constructed to evaluate techniques to address the scarcity, including scaling features, selecting different ratios of negative to positive training examples such as 1 to 1 and 2 to 1, and experimenting with different split ratios in cross-validation - 90/10 and 80/20. These experiments are dubbed inconclusive and not elaborated on further in the dissertation.

1.4.6 CLASSIFIER AND LEXICON SELECTION

The next goal of the dissertation's experimentation is to discover the best performing combinations of classifier, lexicons and feature selection as a means to inform the construction of a *meta classifier* to be used in watchlist drug prediction. The inconclusive nature of the data set sizing experiments prompts settling on the following methodology for classifier training and evaluation:

- test and training sets are sampled with the same distribution as the original data. No use of 1 to 1, 2 to 1 or similar negative to positive sample ratios.
- Data is divided into a 90/10 split, where 90% of the samples are used to train and 10% used to validate
- The splits themselves are sampled per the original distribution.

The classifier evaluation experiments evaluate combinations of classifier type (Support Vector Machines or Naive Bayes), the optional use of normalization and cost-weighting, and various combinations of the lexicons described earlier. Cost-weighting applies a greater penalty to incorrectly classifying a positive example than a negative one during training, as a means to offset the low prevalence (13%) of watchlist drugs in the training data. The

experimental results and analysis make heavy use of acronyms to understand the combinations evaluated:

Acronym	Description
UNB	Un-normalized Naive Bayes
UNBC	Un-normalized Naive Bayes with Cost Weighting
NNB	Normalized Naive Bayes
NNBC	Normalized Naive Bayes with Cost Weighting
SVM	Un-normalized SVM
SVMC	Un-normalized SMV with Cost Weighting
NSVM	Normalized SVM
dis	disease lexicon
react	reactions lexicon
drugs	drugs lexicon
sent	sentiment lexicon
med	medications lexicon

For example, *drugs_dis_sent_react_NSVMC* would equate to an experiment using the Normalized SVM with Cost Weighting, incorporating the drugs, disease, sentiment and reaction lexicons.

A series of 240 experiments were run using the combinations above to ascertain the best combination of lexicons to use when evaluated according to accuracy, F1 score, and area under the ROC curve (AUC). Accuracy is reviewed by Chee first, and the top 3 classifier configurations for each accuracy test are presented for context:

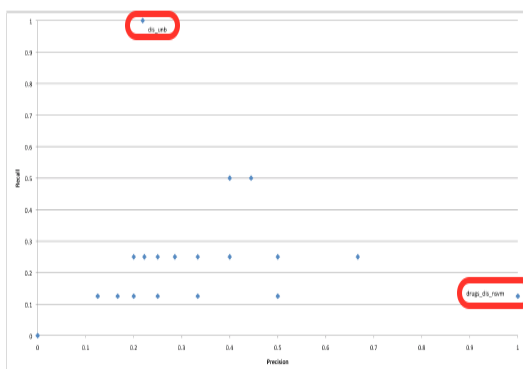
Experiment Type	Configuration	Accuracy	Lower Bound	Upper Bound
CV	dis_react_NSVM	0.903288201	0.793206235	0.957882075
CV	drugs_dis_sent_react_NSVM	0.901353965	0.790801818	0.956684149
CV	drugs_sent_react_NSVM	0.901353965	0.790801818	0.956684149
Test	drugs_dis_sent_react_UNB	0.879310345	0.771204077	0.940291098
Test	drugs_sent_react_UNB	0.879310345	0.771204077	0.940291098
Test	dis_sent_react_UNB	0.879310345	0.771204077	0.940291098
Test(CW)	drugs_dis_NSVMC	0.879310345	0.771204077	0.940291098
Test(CW)	drugs_sent_SVMC	0.862068966	0.750738501	0.928415984
Test(CW)	drugs_sent_NSVMC	0.862068966	0.750738501	0.928415984

The accuracy of a naive baseline classifier that labels all instances as negative would be 86.7% per concentrations of non-watchlist and watchlist drugs in the training data. While several configurations exceed this, when considering 95% confidence interval defined by the lower and upper bounds there is uncertainty that any classifier is more accurate than the naive baseline. The cost weighting shows no accuracy benefits either.

Discussed next are the F1 and area under the ROC curve (AUC) scores. Analysis of the F1 score also prompts inspection regarding the recall and precision tradoffs therein, as the F1 score represents a harmonic mean of the two:

Experiment Type	Configuration	F1 Score
CV	drugs_dis_react_NSVM	0.476190476
CV	drugs_dis_react_UNB	0.472527473
CV	drugs_UNB	0.4698795518
Test	drugs_UNB	0.470588235
Test	drugs_dis_UNB	0.444444444
Test	drugs_dis_sent_react_UNB	0.363636364
Test(CW)	drugs_dis_SVMC	0.444444444
Test(CW)	drugs_UNBC	0.444444444
Test(CW)	drugs_react_SVMC	0.384615385

Experiment Type	Configuration	AUC
CV	drugs_UNB	0.7592
CV	drugs_dis_UNB	0.7564
CV	med_drugs_NNB	0.7545
Test	dis_UNB	0.7514
Test	drugs_dis_UNB	0.6850
Test	dis_sent_react_UNB	0.6675
Test (CW)	drugs_UNBC	0.7825
Test (CW)	drugs_dis_UNBC	0.7075
Test (CW)	drugs_dis_SVMC	0.6900



The F1 scores are lower where cost-weighting is applied, comparable to accuracy. The image above shows how recall (Y axis) and precision (X) access vary across classifier configurations in the test instance, and a comparable image was seen with cost-weighting: low numbers of high recall or precision outliers, and most of the mass concentrated at low recall and precision. AUC does contain higher scores for cost weighted classifiers compared to those non-weighted.

As the goal of this experiment was to identify the value of the lexicons, their prevalence in the top 10 best performing weighted and non-weighted classifiers for each metric is reviewed:

Drugs	33	Weighted Classifiers heightDrugs	20
Disease	32	Disease	18
Sentiment	28	Sentiment	15
Reactions	27	Reactions	10
Medical	7	Medical	5

The ranking favors the drugs, disease and sentiment lexicons, which are then selected for use in the prediction problem. Unfortunately this leads to overfitting concerns because a classifier will learn on the drug names and diseases associated with watchlist drugs. Chee notes disappointment that the reactions lexicon does not rank higher, indicating it may not really capture how people speak about adverse events.

1.4.7 EVALUATING THE BNS LEXICON

Another area of experimental exploration is how the initial word n-gram features are selected to populate the feature vectors. As previously discussed, the top $n - k$ technique is preferred to prompt selection of those word n-grams that are not necessarily the most frequent, but in theory the most informative.

Bi-Normal Separation (BNS) is an alternative technique explored by Chee for discovering those word n-grams. BNS identifies those n-grams that are differentially expressed between two classes: watchlist and non-watchlist messages. BNS lexicons are constructed using the test subset of data for the top 15,000, 10,000 and 5,000 word n-grams, then used in combination with the drugs, diseases and sentiment lexicons, as well as a set of "special features" consisting of numerical counts of:

- disease mentions
- drug mentions
- medical terminology
- sentiment containing terms
- AERS terminology

Experiments were run again using Naive Bayes and SVM classifiers, with optional normalization, optional cost-weighting, and the optional inclusion of the numerical features. The top classifier configuration for each combination of BNS n-grams and numerical features is shown below with its accuracy and confidence bounds:

Configuration	Numericals	BSN	Accuracy	Lower Bound	Upper Bound
bns_drug_dis_sent_NNB	Yes	All(5k,10k,15k)	0.8762	0.7602	0.9405
bns_drugs_dis_sent_NNB	No	All	0.8762	0.7602	0.9405
bns_drugs_dis_sent_NNB	Yes	(10k, 15k)	0.8762	0.7602	0.9405
bns_drugs_dis_sent_NNB	No	(10k, 15k)	0.8762	0.7602	0.9405
bns_drugs_dis_sent_NNB	Yes	15k	0.8762	0.7602	0.9405
bns_drugs_dis_sent_NNB	No	15k	0.8762	0.7602	0.9405

There was no clear advantage to a specific number of BNS features, nor the inclusion of the additional numerical features. Furthermore, the F1 and AUC scores for this test showed no advantage to a specific BNS or numerical feature combination, though un-normalized Naive Bayes did perform better in F1 and AUC than normalized Naive Bayes as seen above. Regardless, none of the tests showed better accuracy, F1 or AUC than the previous tests using specialized lexicons only. Chee hypothesizes this is in part due to the limited number of messages in the test set BNS could select n-grams from.

1.4.8 PREDICTION WATCHLIST DRUGS

The previous experiments were used to assemble a meta-classifier using the best classifiers based on accuracy, F1 and AUC scores. The following were selected:

- Normalized SVM using disease, reaction (AERS) lexicons, having 90.33% Accuracy
- Normalized SVM using drugs, disease and reaction lexicons, having an F1 score of 0.4762
- Un-normalized Naive Bayes classifier with an AUC score of 0.7592

Two meta-classification experiments are constructed to examine the False Positives produced as an indicator for a possible *future* watchlist drugs if used in an applied setting. The first experiment uses the classifier configurations stated above, but the training data is modified to denote drugs *withdrawn* from the market as non-watchlist. It is important to distinguish that a withdrawn drug would have at one point been a watchlist drug, as drugs are placed on the watchlist - even if briefly - before being removed from the market. By looking for these withdrawn drugs marked non-watchlist in the set of False Positives from the training results, we evaluate the ability to discern new watchlist drugs. This experiment was executed via training runs to build 100 classifiers of each configuration. A scoring methodology is applied using the following linear combination:

$$\frac{\# \text{ of False Positives}}{\# \text{ of Occurrences}} * (\# \text{ of False Positives}) * (\# \text{ of Classifier Types})$$

Each occurrence equates to an instance of the withdrawn drug actually being classified in the training. The table below shows the top 5 scores, as well as those withdrawn drugs identified as false positives.

Drug	Positives	Occurrences	Classifiers	Score
<i>clozapine</i> , Clozaril, FazaClo	31	64	3	45.047
<i>fludarabine</i> , Fludara, Oforta	29	61	3	41.361
<i>methylphenidate</i> , Concerta, Daytrana Metadate CD, Metadate ER, Methylin Ritalin, Ritalin LA, Ritalin-SR	25	50	3	37.500
<i>morphine</i> , Astramorph PF Avinza, Duramorph, Infumorph Kadian, MS Contin, MSIR Morphine IR, Oramorph SR, RMS, Roxanol	15	38	3	15.474
<i>meloxicam</i> , Mobic	15	50	3	13.500
<i>thalidomide</i> , Thalomid	10	36	1	2.778
<i>temazepam</i> , Restoril	11	50	1	2.420
<i>hydromorphone</i> , Diluadid, Dilaudid-HP Exalgo, Palladone	9	42	1	1.929
<i>trovafloxacin</i> , Trovan	9	46	1	1.761
<i>rofecoxib</i> , Vioxx	9	50	1	1.620
<i>sibutramine</i> , Meridia	5	27	1	0.926
<i>cerivastatin</i> , Baycol	1	33	1	0.030

Additionally, the generic Sibutramine (Meridia) is of particular interest to this study, as it is under review but not yet an official watchlist drug. FDA has issued safety communications as of November of 2009, and the European Union has removed it from the market.

The second experiment entails removing the withdrawn drugs entirely from training, classifying them after a classifier is built for each fold during cross-validation. This approach is presumed to identify the withdrawn drugs with greater confidence as they are not present in the training data:

Drug	Positives	Occurrences	Classifiers	Score
<i>methylphenidate</i> , Concerta Daytran, metadate(CD, ER) Methylin(ER), Ritalin(LA,SR)	30	34	3	79.412
<i>morphine</i> , Astramorph(PF) Avinza, Duramorph, Infumorph Kadian, MS Contin, MSIR Oramorph SR, RMS, Roxanol	13	38	3	13.342
<i>quetiapine</i> , Seroquel, Seroquel XR	14	31	2	12.645
<i>indomethacin</i> , Indocin Indocin IV, Indocin SR	19	37	1	8.500
<i>sibutramine</i> , Meridia	17	34	1	8.500

The second experiment wholly removes the withdrawn drugs from the training set in each cross validation run, using them only in the test set applied to each classifier built from each fold. This approaches

in cross validation, then introduce

2. Discussion of Contributions

Chee asserts that his research develops a technique for discerning drug Safety events from public data sources, and that he is developing a "crowd sourced" means of Pharmacovigilance. Is this the case? Specific aspects to discuss are:

1) Exploration of classification techniques to discern FDA watch list drugs from free text 2) Exploration of the Yahoo! public Health message forums as potential data source for adverse drug event mining 3) Approach generalizes to other social mediums?

3. Research Critique

A discussion of the methodology - were there any gaps? - good parts/bad parts?

4. Literature Review

What else is out there that is relevant in this space? Other studies that have used public data for medical purposes?

5. Application Areas

The most likely area is drug safety surveillance in uncontrolled settings.

- could we also use this in discovering underlying conditions? - drug combinations? - confounding factor discovery?? Write up 1 to 2 pages here.

6. Concluding Remarks

Conclude the critique with a few endcap statements about what I learned from it, where it could go, how it could motivate future research, etc.

7. Paper Criteria (Grading)

The critique should include a summary of the research reported, a discussion of the major contributions claimed, and an assessment of the significance of those contributions and of the research itself. The critique should also include a brief literature review of the topic related to the thesis, discussion of relevant algorithms, and application areas for the research reported.

Where appropriate, the critique should include a comparison with other issues discussed in class. Students are encouraged to select a dissertation that is related to their course projects. The evaluation criteria for the critique are as follows: Overview of the research reported (20) Review of the related literature (15) Major contributions of the thesis (20) Understanding of techniques and algorithms (20) Application areas (15) Proper construction and readability of paper (10)

7.1 Sentiment, Named Entities and Classification

Of specific interest to the dissertation is applying sentiment analysis and named entity recognition to the messages in these forums. Sentiment analysis is presented as challenging because the domain dependent nature (??) can make it difficult to differentiate between positive and negative sentiment on words and phrases alone. Chee's approach is to calculate the probability of a specific word given a positive or negative class: $P(word|negativeorpositive)$. The hand crafted lexicons Linguistic Inquiry Word Count (LIWC) and SentiWordNet are leveraged to generate sentiment scores on words. Support Vector Machines (SVMs) trained on words as features can also be used to separate positive phrases of text from negative.

Named entity recognition (NER) is necessary for identifying drug names and effects, such as headaches or vomiting. The challenge posed by doing so on forum data is the relaxed structure and oft-present grammatical errors make leveraging existing NLP tools trained on grammatically correct text difficult. Chee draws upon the work of Hearst (?) for automatically acquiring hyponyms from text. Hyponyms are words having more specific meaning than general or subordinate terms, thereby providing strong indication the discovered words are drugs or drug effects.

Classification techniques are employed by Chee to solve the problems of NER, sentiment analysis and assigning class labels to the message forum text. Specifically, Support Vector Machines (SVM) and Naive Bayes classifiers are used.

7.2 Support Vector Machines

SVMs map features into a high-dimensional space using a kernel function (?). A hyperplane is constructed that defines the decision boundary between two classes in this decision space, with those new observations being classified based on which side of the hyperplane they fall on. Chee quotes studies by Forman Joachims stating SVM's strengths in text classification, justifying its use in comparison to Naive Bayes. - LibSVM was used with a radial basis function (RBF) kernel - SVM solves the following optimization problem

$$\min_{u,b,\xi} \frac{1}{2} w^T w + C \sum_{i=1}^l \xi_i$$

$$y_i(w^T \phi(x_i) + b) - 1 - \xi_i$$

$$\xi_i \geq 0$$

- RBF's are non-linear in nature which gave some accuracy advantages over linear - RBF's are trained on two parameters C and γ - Grid search method using cross-validation is employed to look for C and γ because it parallelizes well - C is the penalty parameter for the error term - RBF kernel is defined as $K(x_i, x_j) = \exp(-\gamma \|x_i - x_j\|^2)$, $\gamma > 0$

7.3 Naive Bayes Classification

The dissertation uses Naive Bayes classification to address the NLP problems faced by Chee. Their use was somewhat counterintuitive because Naive Bayes Classifiers assume independence of features (words), whereas we know in real world settings that if a word like

”aspirin” were present, there is a greater probability of the the words ”headache” or ”pain” being present than ”lemonade”. However, in applied settings they still do reasonably well (??). NB has done well in SPAM detection (?) and make sense as a first step for their simplicity (no hyperparameters).

- given word grams w in messages about a drug D - $p(w_i|C)$ probability the i - th word is from class C , C is watchlist or non-watchlist drugs. - $p(D|C) = \prod p(w_i|C)$ - probability of a given drug given the class - W = watchlist, so $P(D|W) = \prod p(w_i|W)$. - Bayes rule writes this as

$$p(W|D) = \frac{p(W)}{p(D)} \prod p(w_i|W)$$

$$p(\neg W|D) = \frac{p(\neg W)}{p(D)} \prod p(w_i|\neg W)$$

Chee combines these two probability modles with the maximum a posteriori (MAP) decision rule to pick the most likely hypothesis. ;discuss maximum a posteriori method; - The method of MAP then estimates θ as the mode of the posterior distribution of this random variable

7.4 Feature Selection

- BNS (Bi-Normal Separation) cited by (?) outperforms other methods for rating ranking feature importance for Classification

- IG (Information Gain) Best practice suggested words occuring less than 3 times in a data set should be removed

7.5 Evaluation Metrics Used

- watchlist drugs are the positive examples - non-watchlist drugs are the negative examples
 - watchlist drugs that are false positive are the interesting ones from Classification - had to work with a 90/10 split where 90% of instances are one class (non-watchlist) and 10% are another (watchlist) it is difficult to outperform a naive classifier that just marks everything as non-watchlist - Receiver Operating Characterisits (ROC) curves are used with their Area Under the Curve (AUC) evaluated. - ROC curves are the true positive

References