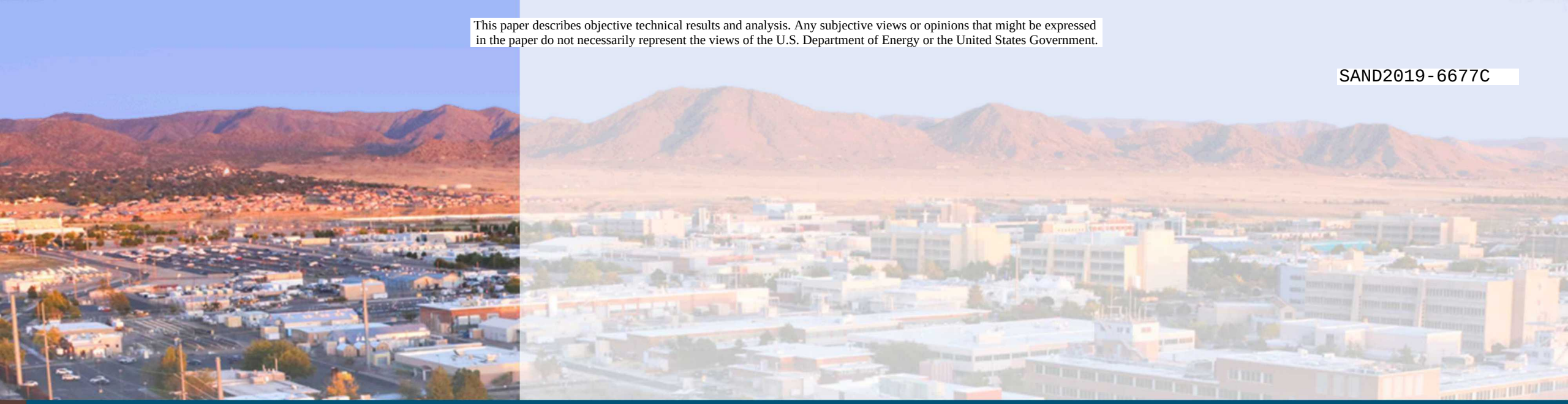
SAND2019-6677C



Automated Recognition of Dual Use Publications

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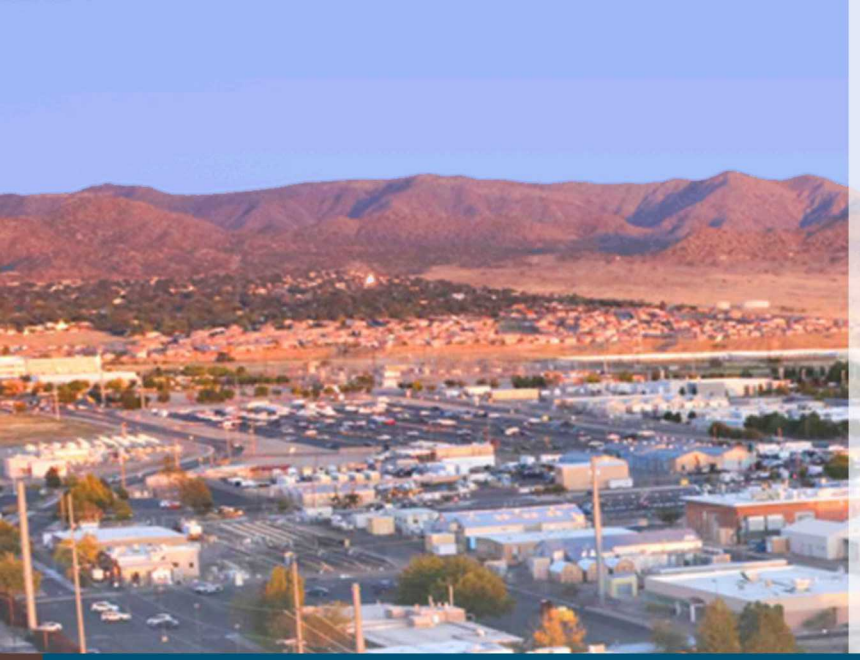
Motivation

- Research in biological fields has great potential to do good as well as harm.
- Unlike other fields with potential for Dual-Use Research Concerns (DURC) (e.g., nuclear engineering) this line is not as well understood.
- Journals and grants now rely on manual content review to identify material that could be misused.
- The community would benefit from automated recommendation methods to expedite the review process.



Data and Methods

Data

- UIUC has provided 272 PubMed abstracts manually labeled by a Subject Matter Expert (SME) according to the following levels of DURC:
  - Not DURC
  - Potential DURC
  - DURC
- Labels are binarized into:
  - Not DURC
  - At least potentially DURC
- Data format example:

```
      Label Text
0        y0  (Title-Abstract from paper 0)...
1        y1  (Title-Abstract from paper 1)...
2        y1  (Title-Abstract from paper 2)...
3        y0  (Title-Abstract from paper 3)...
...
```

- Also provided: list of DURC-related keywords used by SME to identify DURC publications.

Data preprocessing

- We preprocess the titles and abstracts accordingly:
  - Abstracts and titles are all set to lower case.
  - Abstracts and titles are tokenized.
  - Stop words are removed.
  - Words with document frequency of >0.7 are removed.
  - Words that appear in fewer than 1 document are removed.
- We compare this baseline to two additional preprocessing methods:
  1. Remove all tokens that are not keywords identified by a SME.
  2. Remove all tokens that are keywords identified by a SME.
- These methods are used as control to identify which tokens are most significant to the classifier's decision boundary.

Representation of data

- For analysis, the Pubmed articles' titles and abstracts are converted into a numerical vector representing normalized word occurrences.
- Each entry in the vector is a word's term-frequency inverse document frequency (TFIDF)
- TFIDF highlights words that are most signature to a given document.

$$\text{tfidf}(t, d, D) = \text{tf}(t, d) * \text{idf}(t, D)$$

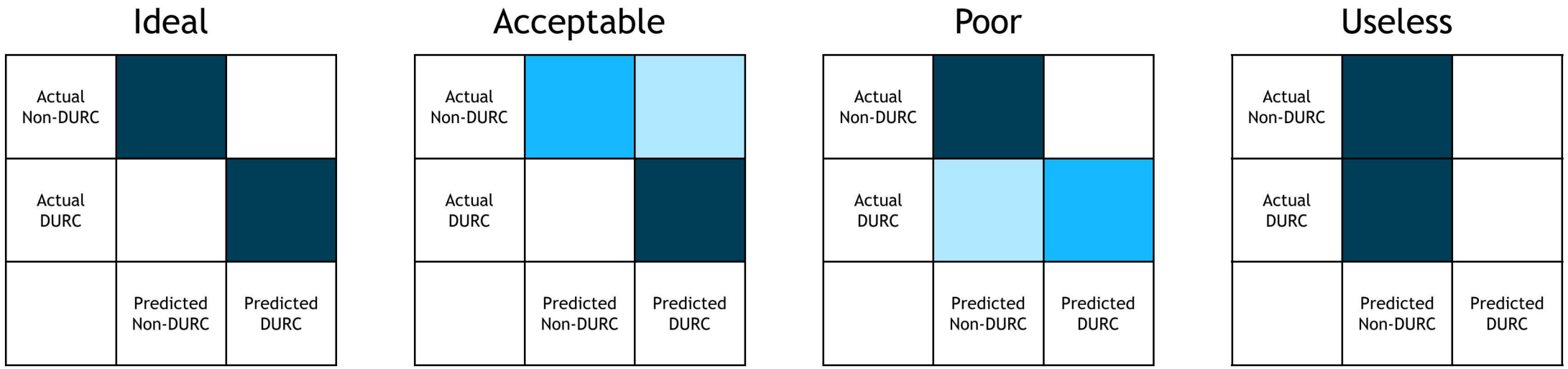
- Where:
  - **tf** corresponds to the frequency that term ***t*** appears in document ***d***.
  - **idf** offsets the term frequency according to how many documents in the corpus ***D*** contain the term ***t***.

Methods

- For classification, we inspect several possible classifiers
  - Linear SVC: C=0.5
  - K-Nearest Neighbors (KNN): K=15
  - Gaussian Naïve Bayes
  - Random Forest (RF): n_estimators=100, max_depth=2
- Additionally, we also inspect the performance of these classifiers 2D representations of the data using:
  - PCA
  - SVD
  - Spectral Embedding with RBF kernel
  - TSNE

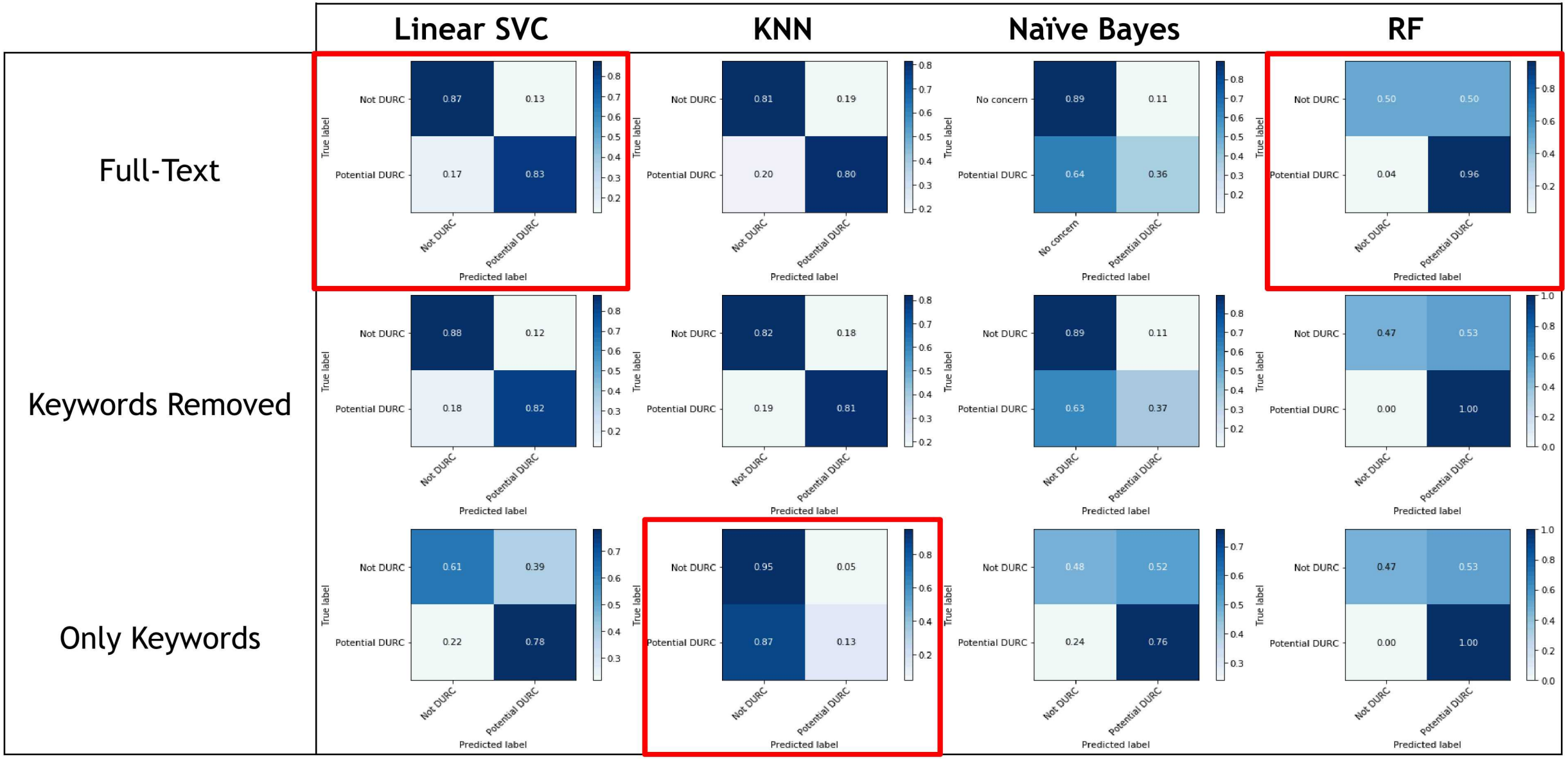
Performance Measurement

- Classifier performance is averaged over stratified 5-fold cross validation.
- Performance is summarized using a Confusion Matrix.
- We wish to find as many DURC papers as possible, constrained by also reducing the number of false positives a manual reviewer would have to read.
- Ideally, each class accuracy should be maximized.
- Pragmatically, we favor DURC class accuracy over (and empirically, at the expense of) non-DURC class accuracy.
- DURC labelled data were weighed more heavily than non-DURC for Linear SVC and Random Forest to emphasize recall.



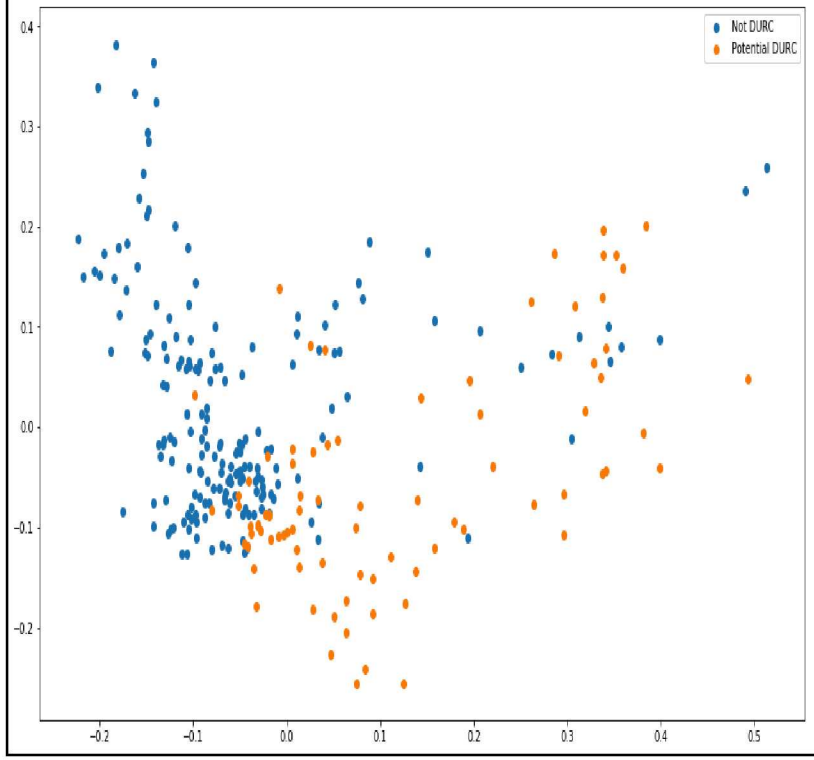
Results

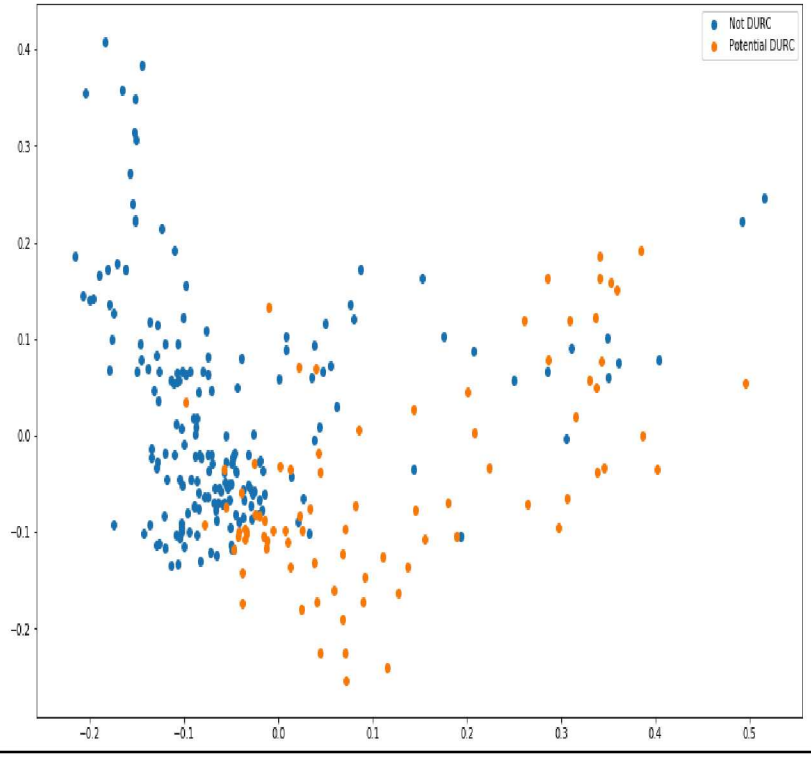
Classification Performance on Full-Dimensionality Data

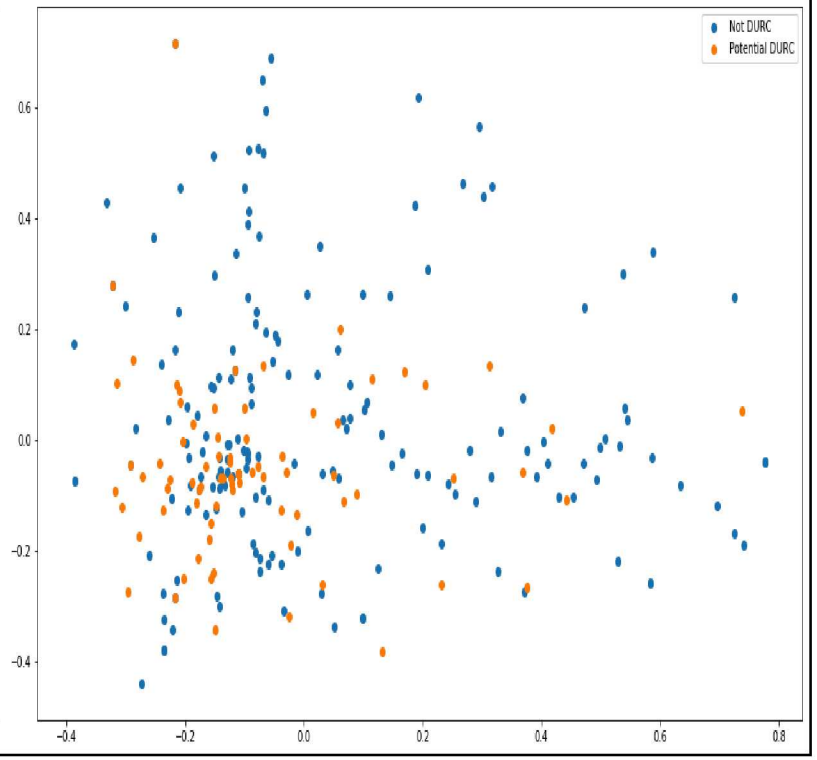


PCA Dimensionality Reduction

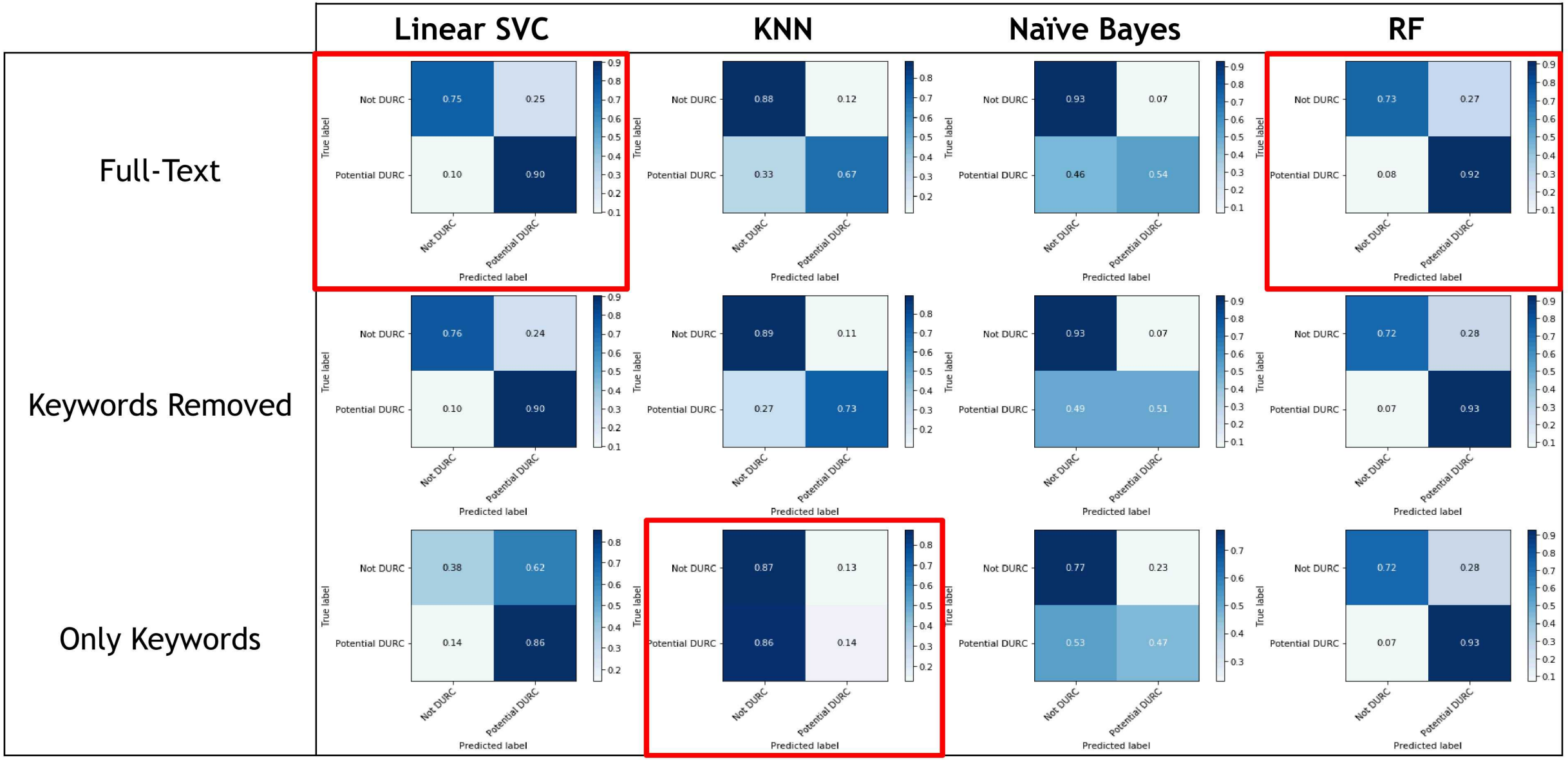
Full-Text	Keywords Removed	Only Keywords



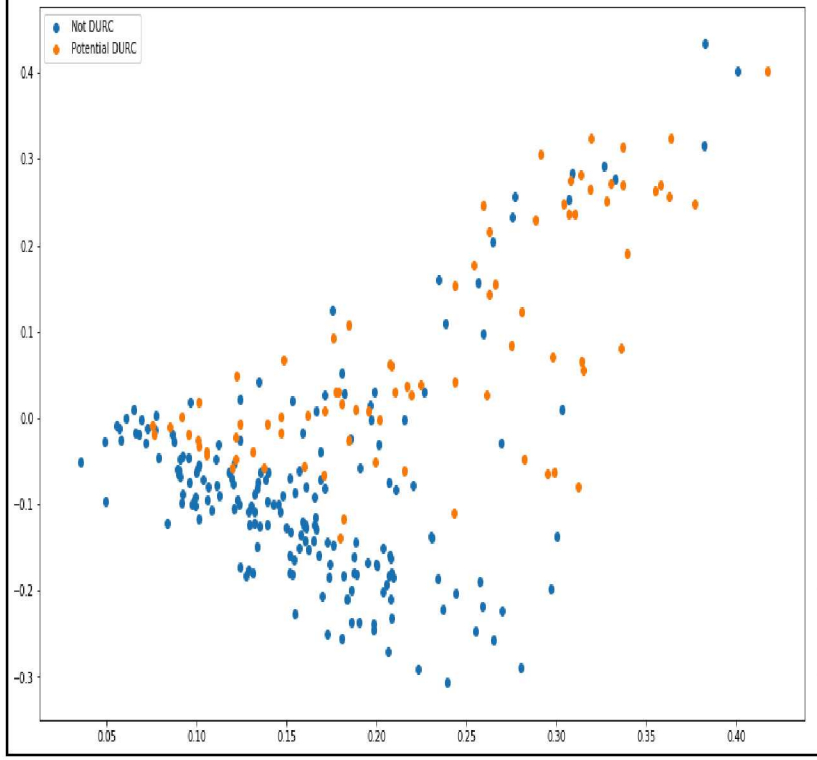
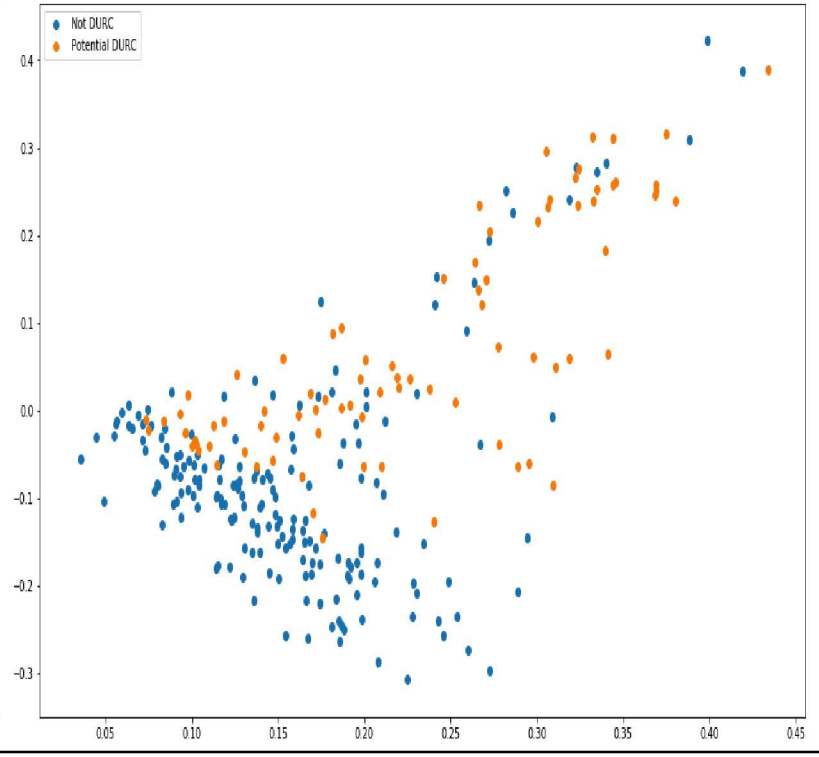
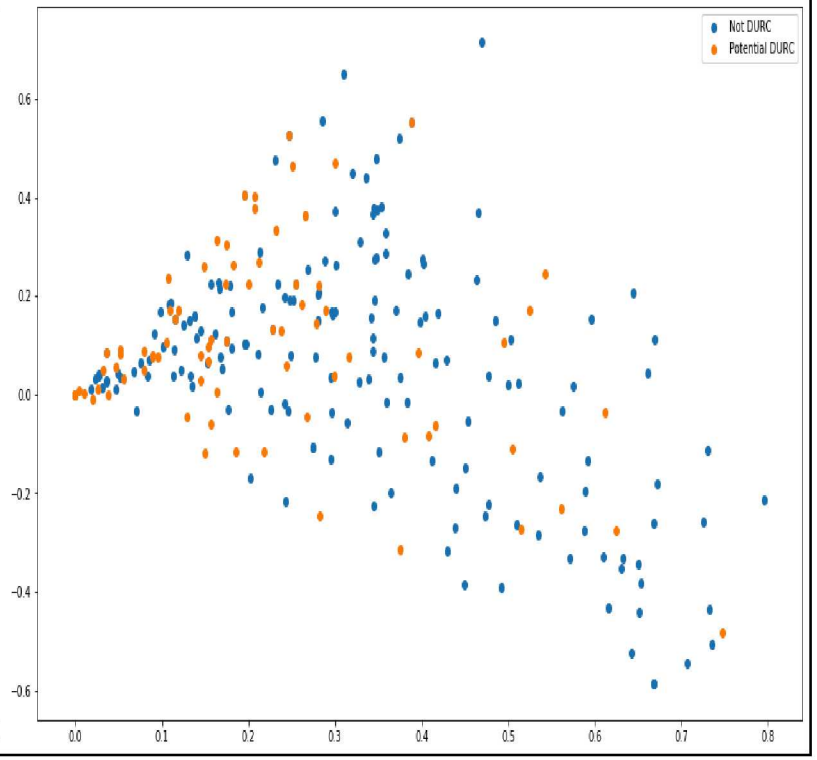




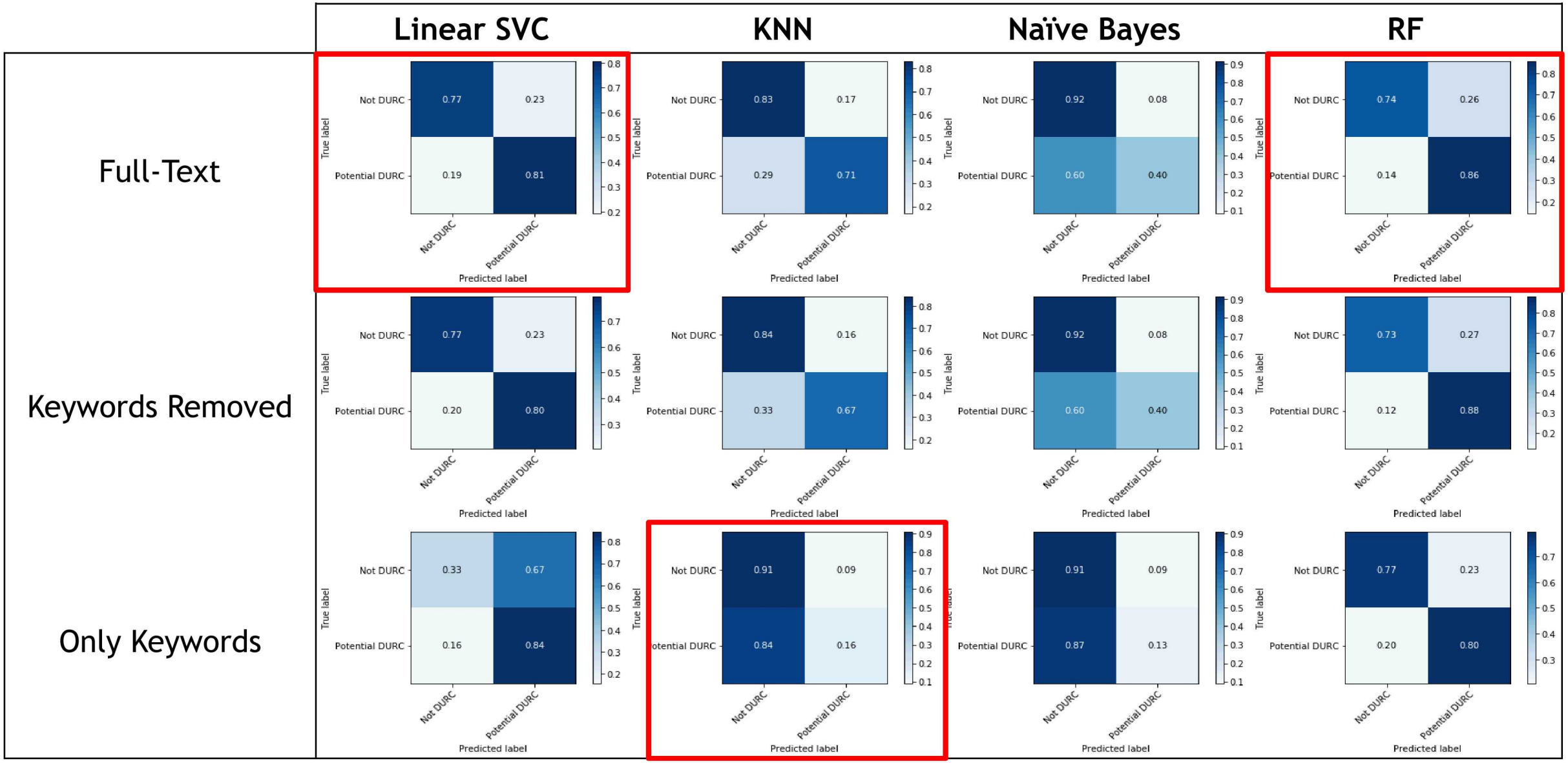
Classification Performance on PCA-Reduced Data



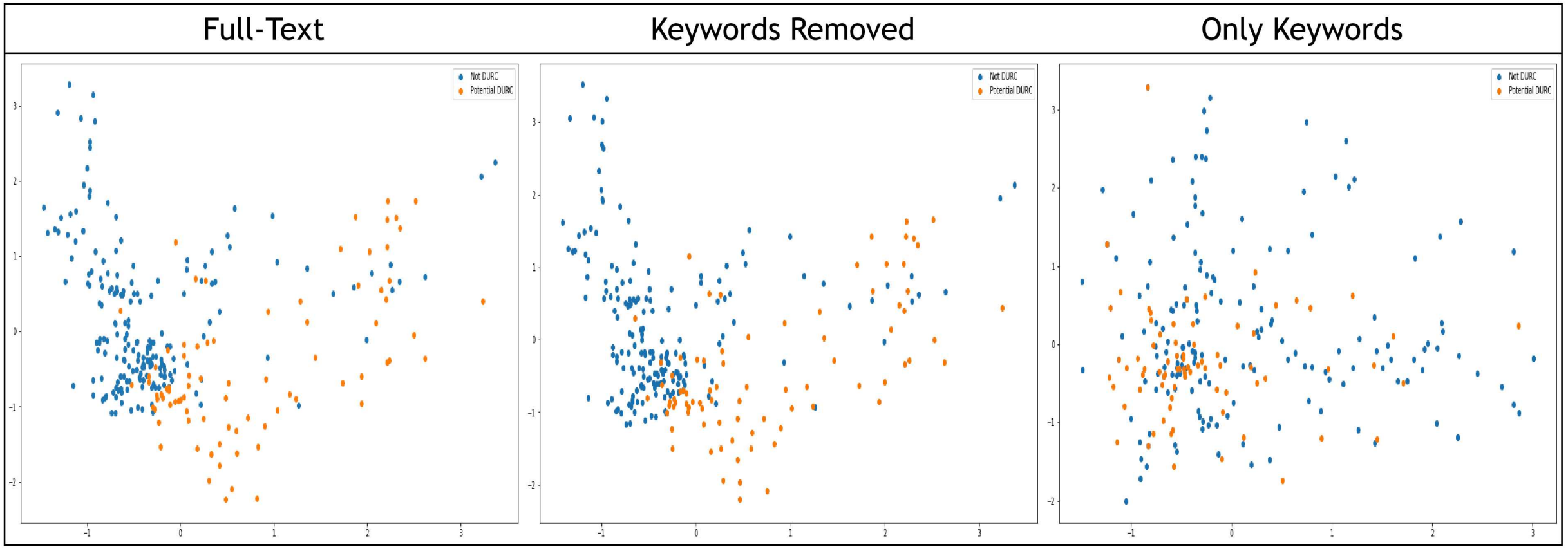
SVD Dimensionality Reduction

Full-Text	Keywords Removed	Only Keywords
		

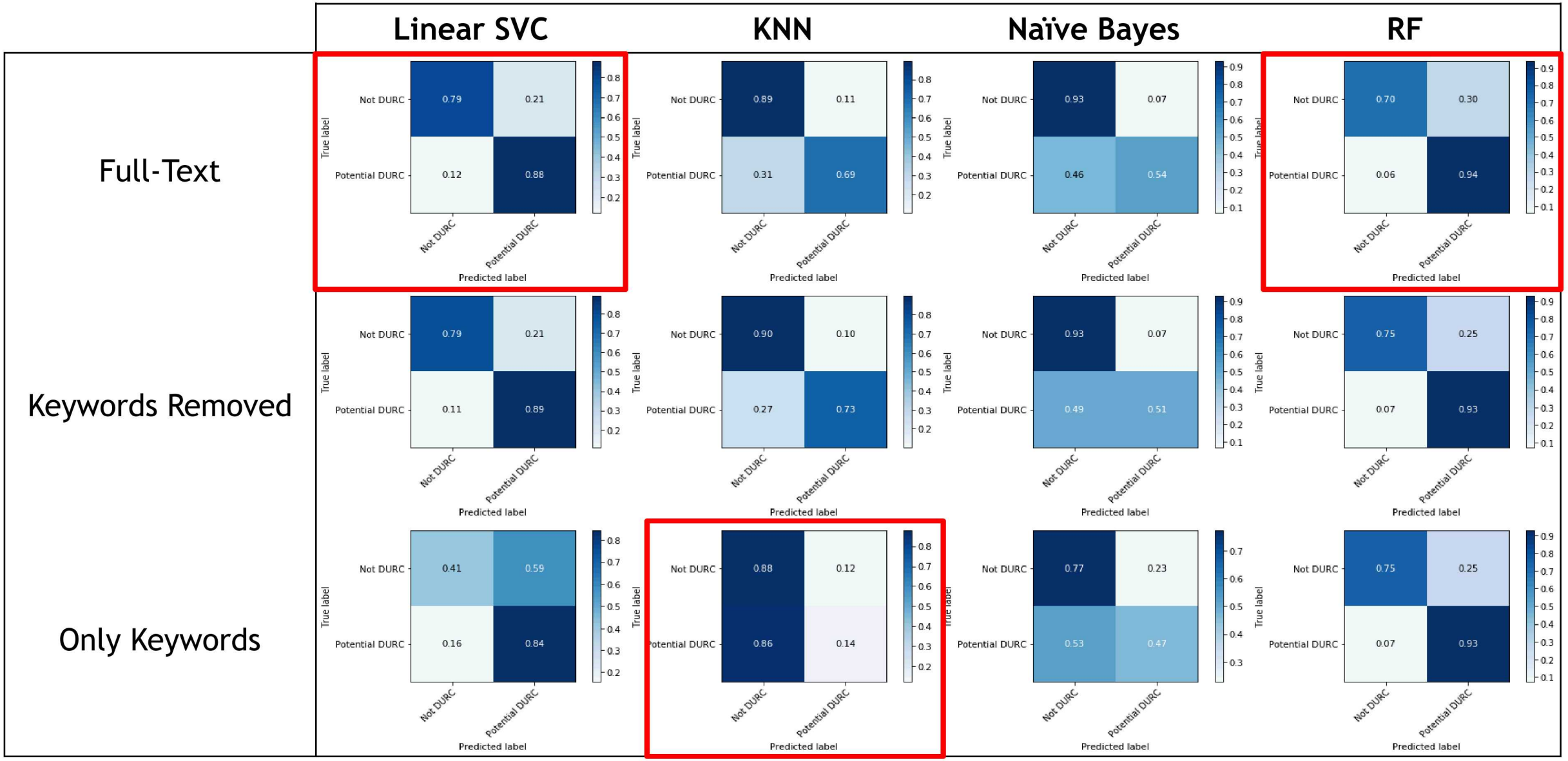
Classification Performance on SVD-Reduced Data



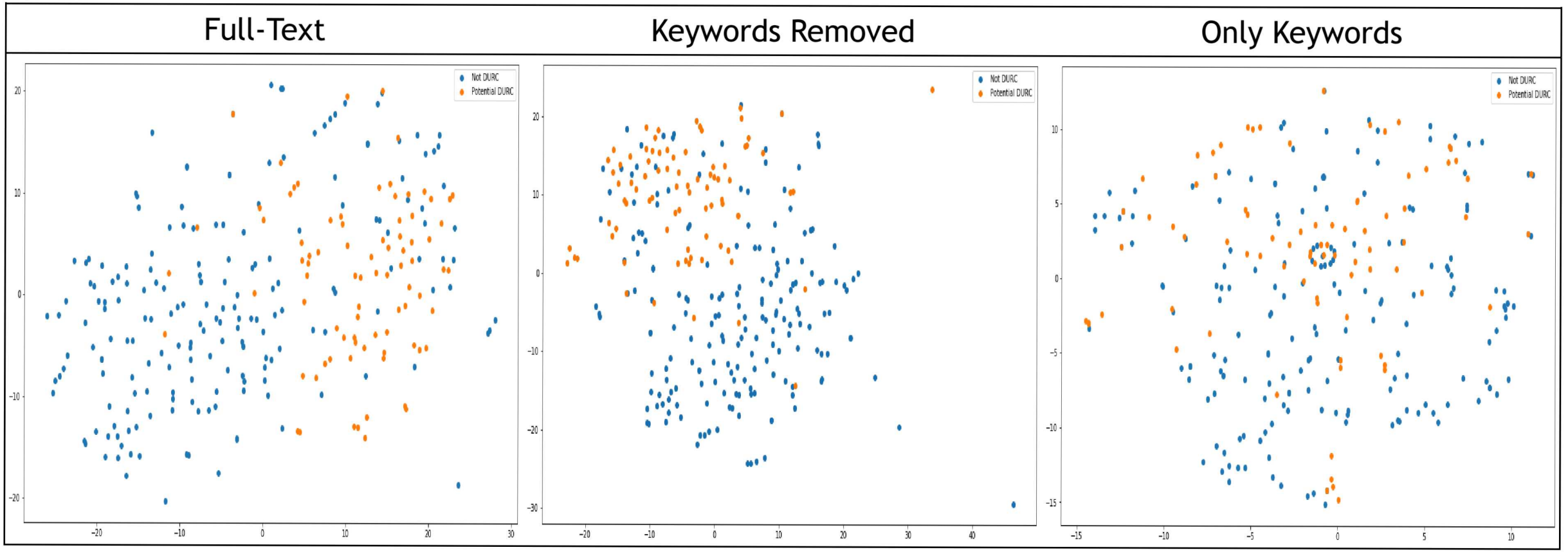
Spectral Embedding Dimensionality Reduction



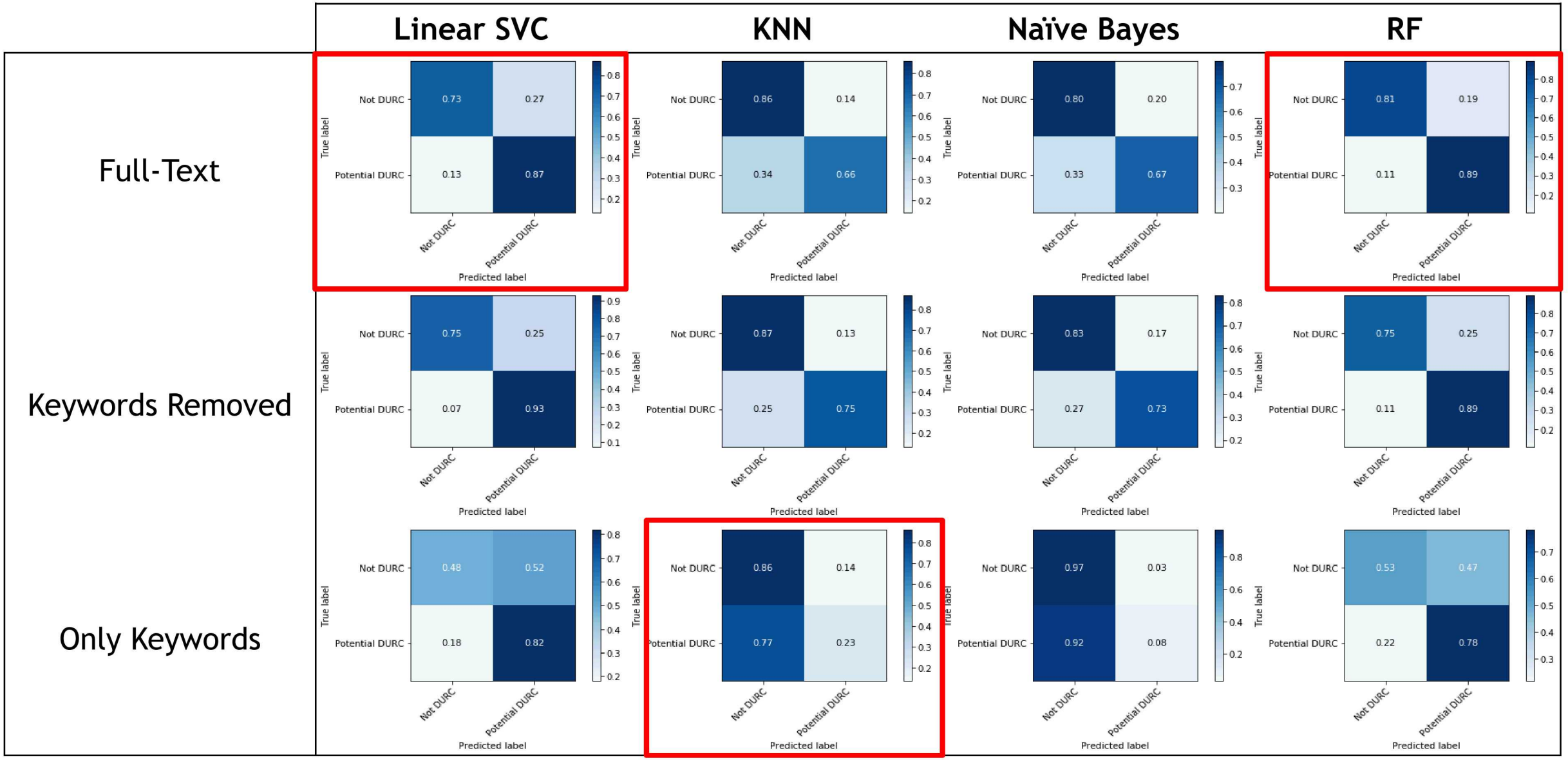
Classification Performance on Spectral Embedding-Reduced Data



TSNE Dimensionality Reduction



Classification Performance on TSNE-Reduced Data



Analysis

Principal Conclusions

- Both Linear SVC and Random Forest were able to achieve ~90% detection rates of DURC in our data set.
- KNN performed best on full dimensionality data, but suffered when only using keywords.
- Given that DURC and non-DURC papers were not completely separable when projected, the experiments that had the highest detection rate tended to have a higher false-positive rate.
- Dimensionality reduction improved the performance of Random Forest considerably, significantly reducing the rate of false positives.
- After dimensionality reduction, false positive rates were less than 30%, significantly down-selecting the number of documents required for manual curation.
- Contextual words (Non-keywords) were critical for classification performance.

Future Work

- Our work was done on a dataset of limited size. Thus, additional work will be needed to ensure the performance generalizes to larger and more varied datasets.
- More work must be done to better explain what factors contributed to the ease of classification in our dataset.

Questions?

Provided Keywords

highly
specificity
optimize
wide
better
irreversible
cytopathogenicity
screen
mosaic
translocate
strategic
pathogenicity
cytotoxicity
increase
functional
different
immunogenicity
regulate
pathogen
versatile
combine
penetrate

optimal
amplify
robust
robustness
efficiency
functionality
catalytic activity
deliver
combinatorial
potentiate
superior
improve
effective
bioagent
multivalent
solubility
more
induce
elevated
selectivity
switch
elevate

refine
diversity
introduce
novel
toxicity
virulence factor
resistance
potentiation
genetic
cytotoxic
biological activity
enhance
recombine
exogenous
host specificity
toxic
virulent
efficacy
virulence
chimeric
upregulate
stable

efficient
stability
new
biosynthesize
change
maximize
genetically
lethality
exchange
engineer
broad
secrete
edit
modular
design
swap
diversify
synthesize
resilient
alter
bacteria
selective

cytomodulin
modulate
virion
titer
build
lethal
recombinant
reversible
encapsulate
pathogenesis
reconstruct
pure
bind
coexpress
modularity
toxin
traceless
endocytose
cellular activity
conjugate
redesign
potent

antibiotic resistance
replace
fuse
virus
potency
expand
synthetic
mutate
vary
enzymatic activity
modify
specific
construct
heterologous
versatility
infectivity
maximal
rapid
purer
higher
accelerate
broaden