Ensemble Classification of Grants using LDA-based Features

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Abstract

Classifying research grants into useful categories is a vital task for a funding body to give structure to the portfolio for analysis, informing strategic planning and decision-making. Automating this classification process would save time and effort, providing the accuracy of the classifications is maintained. We employ five classification models to classify a set of BBSRC-funded research grants in 21 research topics based on unigrams, technical terms and Latent Dirichlet Allocation models. To boost precision, we investigate methods for combining their predictions into five aggregate classifiers. Evaluation confirmed that ensemble classification models lead to higher precision. It was observed that there is not a single best-performing aggregate method for all research topics. Instead, the best-performing method for a research topic depends on the number of positive training instances available for this topic. Subject matter experts considered the predictions of aggregate models to correct erroneous or incomplete manual assignments.

Keywords: research grant classification, document classification, topic models

1. Introduction

On a worldwide scale, funding bodies fund thousands of research grants per year. In the United Kingdom, national funding bodies are mostly organised by discipline in research councils. For example, the Biotechnology and Biological Sciences Research Council (BBSRC) funds research in biotechnology and biology, the Medical Research Council (MRC) funds research in medicine, the Engineering and Physical Sciences Research Council (EPSRC0 funds research in engineering and physical sciences etc. BBSRC processes, on average, over 1600 research grant applications per year.

Classifications give structure to research portfolios, enabling timely, accurate portfolio analysis, to inform strategic planning and decision making, whilst also assisting in the process of peer review. Currently classification is a manual and subjective process taking considerable time and effort. Being able to assign classifications automatically will enable greater accuracy, consistency and timelines and has the potential to be used across a broad range of funding mechanisms. In addition, automatic classification offers the potential to amend the set of topics flexibly. The efficiency savings can free up capacity to perform increasingly sophisticated analyses. However, automatic assignment is only meaningful if the associated error rate is very limited, even if no assignment can be produced for some grants. Increasing classification accuracy requires that enough manually annotated data is available, to avoid error propagation and ensure that the resulting classification model will be able to generalise. The current study showed that automatic classifications were useful in correcting manual assignments.

research topics. We combine a variety of classifiers whose features encode the words and technical terms in the description of funded grants as well as Latent Dirichlet Allocation (LDA) topics. In particular, for each research topic we train five classification models based on Support Vector Machines (SVMs) that employ as features different combinations of words, techincal terms, LDA topics (Blei et al., 2003) and link LDA topics (Erosheva et al., 2004). LDA topics are used as a means of reducing sparsity in the space of words and technical terms.

Combining classifiers aims at boosting precision. The predictions of the five basic classification models are combined using five voting-based aggregate classifiers. We investigate whether avoiding to produce final predictions for instances on which the basic classification models disagree leads to more accurate final predictions.

As evaluation data we use a set of freely available descriptions of BBSRC funded research grants to be assigned to 21 research topics. Combining diverse classifiers proves to be beneficial towards increasing classification performance without leaving many instances unclassified. The precision achievable by combined classification models for a research topic depends on the number of positive training instances available for this topic.

The remaining of the paper is structured as follows: section 2. briefly reviews related work. In section 3., we present five basic SVM classifiers based on SVMs and LDA models. Section 5. discusses our experimentation with the basic classifiers and evaluation results. It presents five aggregate classifiers and the evaluation results of applying them to the dataset at hand. Finally, section 6. concludes the paper and summarises futore work dimensions.

In this study, we address the task of classifying grants into

field	value
reference number	BB/C000072/1
title	Understanding the role and regulation of cation homeostasis during citric acid stress in the spoilage yeast Saccharomyces cerevisiae
abstract	Yeasts are able to spoil foods and beverages because they have evolved mechanisms that allow them to adapt and grow under the extreme environmental conditions, such as acid pH, that are often used to preserve foods and drinks. Citric acid (E330) and its salts are used extensively in the food and have rege inductor to control pH and act as preservatives to prevent microhild
	growth, but little is known about how yeasts adapt and grow in the presence of the acid
objectives	We have identified three regulatory proteins, the protein kinases Hog1p, Sky1p and Ptk2p, whose function is required for optimal adaptation of spoilage yeast to the inhibitory effect of citric acid. Thus, the overall aim of this research will be to; a) identify the other component proteins of the signalling pathways used by Hog1p, Sky1p and Ptk2p; b) determine whether Hog1p, Sky1p and Ptk2p interact, or are components of the same, or different, signalling pathways; and, b) identify what proteins Hog1p, Sky1p and Ptk2p interact with, and thus regulate, to switch on the adaptive mechanisms that allow the yeast to grow in the presence of citric acid stress. To fulfil these aims we will
initiative	Proteomics and Cell Function (PCF)
holding institution	University of St. Andrews
holding department	Biology
principal investigator	Dr Peter Coote
research topics	MFS, MIC

Table 1: Example grant description

2. Related work

LDA models have been applied extensively for a variety of purposes, e.g. modelling dependencies in text (Griffiths et al., 2005), matrix factorisation (Agarwal and Chen, 2010), decomposing biodiversity data (Valle et al., 2014), identifying musical key-profiles (Hu and Saul, 2009) and cyber security (Aswani et al., 2015). Similarly to the present study, Torkkola (2003) used LDA models to reduce the dimensionality of token representation. Moreover, LDA topics have been used as classification features in Lee et al. (2015).

Semi-supervised LDA models have been used for classifying documents, rating movies from reviews, predicting the popularity of webpages and for image classification and annotation (Mcauliffe and Blei, 2008; Lacoste-Julien et al., 2009; Wang et al., 2009). Using semi-supervised LDA models for the current task is a matter of future work.

3. Classification Models

We build classification models for assigning a grant description to one or more predefined research topics. A grant description is structured in fields, as shown in Table 1. We hypothesise that field content similarities between grants correspond to thematic similarities and thus should be the basis for assigning grants to research topics. As our baseline classification model, we employ a Support Vector Machine (SVM) trained on features representing words in the textual fields of the grant descriptions, i.e. title, abstract and objectives.

Analysing the results and errors of the baseline experiment revealed that frequently grant descriptions in the same research topic do not share many words, due to the limited number of instances in the data set. The number of funded grants over the years is significant in terms of research but marginally sufficient to train a machine learner. To address

#	abbreviation	research topic
1	AGE	ageing
2	AH	animal health
3	AW	animal welfare
4	CS	crop science
5	DH	diet and health
6	EG	bioenergy
7	IB	industrial biotechnology
8	IMM	immunology
9	MFS	microbial food safety
10	MIC	microbiology
11	NS	neuroscience and behaviour
12	PHM	pharmaceuticals
13	PS	plant science
14	REG	regenerative biology
15	DDD	replacement, reduction and
15	KKK	refinement of animals in research
16	SB	systems biology
17	SC	stem cells
18	SS	soil science
19	STR	structural biology
20	SYN	synthetic biology
21	TD	technology development

Table 2: Research topics

sparsity, we employed LDA models (Blei et al., 2003; Griffiths and Steyvers, 2004; Torkkola, 2003) to assign probabilistically grant descriptions to a predefined number of topics in an unsupervised manner based on content similarity. Using LDA topic features to represent grants addresses sparsity, because the LDA topics are less than the distinct words.

A second analysis outcome is that for some grants it is important to consider fields other than the free text ones. For example, the principal investigator's name (table 1) can provide hints for classifying a grant, given that a researcher usually investigates a small number of areas and applies for follow-up grants. This extra Information can be accommodated in LDA models. The standard LDA model assumes the same distribution for all features. However, information of different fields is not equally important for the task. To cover this requirement, we use the link LDA model (Erosheva et al., 2004), which allows considering different views to describe a single grant description. We employ separate views for the free text fields, the award details and the principal investigator's name.

Thirdly, although free text fields often contain multiword technical terms, they are not parts of the feature representation. Technical terms, as well-known/common phrases (concepts), occur naturally in text and are often shared between grant descriptions. Complimentary to words, terms are used as features in the standard LDA model or as an extra view in the link LDA model. We synthesise the above components into the classifiers:

- **SVM:** A SVM trained on words in the free text fields, i.e. title abstract and objectives (table 1).
- **LDA**: A SVM trained on the topic distribution of an LDA that considers words in the free text fields.
- **LDA + terms:** A SVM trained on the topic distribution of an LDA that considers word and multiword terms in the free text fields.
- link LDA: A SVM trained on the topic distribution of a link LDA that collectively considers the following views: (a) free text fields, (b) award details, i.e. initiative, holding institution and department, and (c) principal investigator's name.
- **link LDA + terms:** A SVM trained on the topic distribution of a link LDA that considers the views above plus a view for multiword terms extracted from the free text fields.

4. Data

The dataset consists of 5,462 descriptions that have been awarded by the BBSRC between 2008 and 2013 or with annual spend between 2008/09 and 2013/14. For joint or transferred grants, only one grant per group was included in the dataset: the lead grant for joint grants or the original grant for transferred grants. 3,924 grants have been manually assigned to one or more research topics in table 2 by subject matter experts. These grants were used as training data while the remaining 1,538 descriptions comprise the test data. Manual annotations of test set descriptions were provided at a later stage for evaluation. The description of each grant in these two sets consists of the fields shown in table 1. The abstract, the objectives and the initiative under which the grant was funder may be empty. Table 3 shows the numbers and associated percentages of positive and negative instances per research topic in the training data.

5. Experiments

We evaluate five classification models, introduced in section 3., and we investigate prediction ensembles that can

research	positiv	e instances	negative instances				
topic	%	#	%	#			
MIC	28.9	444	71.1	1094			
PS	22.4	345	77.6	1193			
NS	14.6	225	85.4	1313			
TD	14.4	221	85.6	1317			
CS	13.7	211	86.3	1327			
STR	13.6	209	86.4	1329			
AH	12.5	193	87.5	1345			
IMM	11.1	170	88.9	1368			
IB	8.7	134	91.3	1404			
AGE	5.0	77	95.0	1461			
SB	4.8	74	95.2	1464			
PHM	4.1	63	95.9	1475			
DH	4.0	62	96.0	1476			
SYN	4.0	62	96.0	1476			
SC	3.4	53	96.6	1485			
EG	2.5	39	97.5	1499			
MFS	2.5	38	97.5	1500			
REG	2.2	34	97.8	1504			
SS	2.1	32	97.9	1506			
AW	1.2	19	98.8	1519			
RRR	1.0	16	99.0	1522			

Table 3: Positive and negative instance counts and percentages per research topic sorted in order of decreasing positive instances.

boost performance. In the pre-processing stage, the free text fields of grant descriptions, i.e. title abstract and objectives and the fields describing award details, i.e. initiative, holding institution and department (table 1), were sentence split and tokenised. Technical terms were extracted using TerMine¹ (Frantzi et al., 2000). For our experiments, the LibLinear SVM implementation² (Fan et al., 2008) was used, while the LDA model (Blei et al., 2003) and link LDA model (Erosheva et al., 2004) were implemented from scratch and parameters were set as follows: alpha = beta = 0.5. LDA models were trained for 2000 iterations to induce 500 topics. As evaluation measures, we employ precision (P), recall (R) and F-score (F₁):

$$P = \frac{TP}{TP + FP}, R = \frac{TP}{TP + FN}$$
$$F_1 = 2\left(P^{-1} + R^{-1}\right)^{-1}$$

TP, FP and FN stand for true positives, false positives and false negatives, respectively.

As this study is part of a collaboration of the National Centre for Text Mining (NaCTeM)³ and the BBSRC, to evaluate the classification models of section 3. we followed a procedure similar to the SemEval series⁴. Initially, the BB-SRC provided fully annotated training data and unannotated test data. Classification models were trained on the

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<sup>3</sup>nactem.ac.uk
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<sup>4</sup>alt.gcri.org/semeval2015
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¹nactem.ac.uk/software/termine

²In particular, L2-regularized L2-loss support vector classification (dual) solver was employed, with parameter settings: C = 1, eps = 0.01, p = 0.1.

	Classification model														
research		SVM			LDA		LL)A + ter	ms	l	ink LD	4	link l	L DA + 1	terms
topic	Р	R	F_1	Р	R	F_1	Р	R	F_1	Р	R	F_1	Р	R	F_1
MIC	91.9	74.6	82.3	87.9	72.1	79.2	86.5	73.9	79.7	85.6	73.9	79.3	89.1	73.4	80.5
PS	97.4	87.3	92.1	94.2	84.1	88.8	94.5	84.9	89.5	93.1	82.6	87.6	95.4	78.0	85.8
NS	99.0	85.8	91.9	98.0	86.7	92.0	97.0	87.6	92.1	98.0	85.3	91.2	96.6	88.4	92.3
TD	77.8	72.9	75.2	80.0	76.0	78.0	73.3	73.3	73.3	77.9	70.1	73.8	75.7	74.7	75.2
CS	88.7	77.7	82.8	89.3	75.7	81.8	85.4	74.9	79.8	85.3	71.6	77.8	88.2	74.4	80.7
STR	91.1	78.0	84.0	87.3	72.3	79.1	86.5	76.6	81.2	84.8	74.6	79.4	87.7	71.3	78.6
AH	93.1	77.2	84.4	92.3	80.8	86.2	94.1	82.4	87.9	92.0	77.7	84.3	89.0	79.3	83.8
IMM	94.5	71.2	81.2	90.8	69.4	78.7	90.9	76.5	83.1	84.2	68.8	75.7	87.3	72.9	79.5
IB	87.0	64.9	74.4	70.3	72.4	71.3	81.1	76.9	78.9	83.2	73.9	78.3	79.7	70.2	74.6
AGE	92.7	49.4	64.4	83.3	45.5	58.8	85.7	39.0	53.6	81.3	33.8	47.7	87.9	37.7	52.7
SB	81.9	48.7	61.0	74.4	43.2	54.7	81.3	52.7	63.9	76.2	43.2	55.2	76.2	43.2	55.2
PHM	87.5	22.2	35.4	87.5	33.3	<i>48.3</i>	74.4	46.0	56.9	76.7	36.5	49.5	77.8	33.3	46.7
DH	95.9	75.8	84.7	86.3	71.0	77.9	91.8	72.6	81.1	84.9	72.6	78.3	85.7	77.4	81.4
SYN	96.6	45.2	61.5	93.9	50.0	65.3	93.1	43.6	59.3	97.1	53.2	68.8	90.0	58.1	70.6
SC	100.0	58.5	73.8	86.7	73.6	79.6	92.3	67.9	78.3	84.6	62.3	71.7	79.6	66.0	72.2
EG	90.0	46.2	61.0	78.6	56.4	65.7	78.1	64.1	70.4	81.3	66.7	73.2	67.7	59.0	63.0
MFS	92.9	34.2	50.0	61.1	29.0	39.3	76.2	42.1	54.2	84.2	42.1	56.1	80.0	21.1	33.3
REG	92.3	35.3	51.1	77.8	41.2	53.9	70.0	41.2	51.9	82.4	41.2	54.9	83.3	29.4	43.5
SS	90.5	59.4	71.7	77.8	65.6	71.2	88.9	75.0	81.4	81.5	68.8	74.6	87.5	65.6	75.0
AW	80.0	21.1	33.3	75.0	15.8	26.1	85.7	31.6	46.2	66.7	21.1	32.0	85.7	31.6	46.2
RRR	0.0	0.0	0.0	0.0	0.0	0.0	0.0	0.0	0.0	0.0	0.0	0.0	0.0	0.0	0.0
average	86.7	56.5	66.5	79.6	57.8	65.5	81.3	61.1	68.7	80.0	58.1	66.2	80.5	57.4	65.3

Table 4: Results per classification model and research topic (%). Scores greater than 85% or less than 50% are printed in bold or italics, respectively. Research topics are sorted in order of decreasing positive instances (see Table 3).

training set to produce predictions for the grant descriptions in the test set. Each classification model was trained for each research topic. Then, each test grant description was presented to each of these models and produced five estimates for each research topic. Finally, the BBSRC disclosed the gold-standard assignments of test grant descriptions to research topics.

Table 4 presents evaluation results per classification model and research topic. The best performing models are SVMand LDA + terms. In some cases, the remaining 3 models achieve comparable or better results. For example, *link* LDA + terms achieved the maximum F-score for the *NS* topic. For some research topics, e.g. *AW*, *MFS* and *RRR*, all five models achieved very low results. The reason is sparsity of positive instances for these topics, as shown in table 3. For *RRR* the results are zero because none of the classifiers identified true positives.

Automatic assignments are useful for decision making only if they are very precise. Thus, we investigate ways of combining the predictions of the five classification models, to increase precision. Cross-verifying predictions of grant descriptions to research topics, lowers the possibility of accepting a prediction. However, accepted predictions are expected to be more accurate than any of the five models. We consider the following methods:

- **5 agreements:** we accept a positive prediction if all five classification models agree.
- **4 positives:** we accept a positive prediction if at least four models agree.
- **3 positives:** we accept a positive prediction if at least three models agree.
- **2 positives:** we accept a positive prediction if at least two

models agree.

1 positive: we accept a positive prediction if produced by one or more models.

In all methods, we accept a negative prediction if all five models agree. These methods do not produce any prediction for some grant descriptions, e.g. if the five models do not agree when using *5 agreements*.

Table 5 shows evaluation scores of the five ensemble methods for each of the 21 research topics. It can be observed that the stricter the aggregate algorithm is, the higher precision and the lower recall it achieves. However, there is not a particular aggregate model that achieves the best balance in this trade-off between precision and recall, i.e. the highest F-score, for all research topics.

As mentioned previously, all aggregate methods except for 1 positive are lossy, i.e. do not produce predictions for all test instances. Table 6 enlightens this issue, by showing the percentages of omitted instances over the total number of training instances for each aggregation method and for each research topic. The results in table 5 confirm the increase of precision at the expense of a small percentage of grant descriptions for which no prediction was produced (see column omitted instances). We observe that the stricter an aggregation method is, the more lossy it is. As table 6 is sorted in order of decreasing positive training instances available for a research topic, we observe that topics with more positive training instances tend to be more lossy. This is probably because for topics with small numbers of positive training instances the five basic classifiers tend to be very strict and as a result agree with each other more often. Table 5 shows that the methods 5 agreements and 4 positives are the best performing for many research topics but

		Classification ensemble model													
research	5 aş	5 agreements 4 positives			4 positives			positiv	es	2	positiv	es	1	positiv	'e
topic	Р	R	F_1	Р	R	F_1	Р	R	F_1	Р	R	F_1	Р	R	F_1
MIC	99.2	81.0	89.2	94.4	84.4	89.1	91.8	85.7	88.6	87.0	86.8	86.9	77.3	87.4	82.0
PS	100.0	90.8	95.2	97.9	91.6	94.7	95.7	92.1	93.9	94.2	92.5	93.3	89.1	92.8	90.9
NS	100.0	93.1	96.4	99.5	93.6	96.4	98.5	93.8	96.1	97.6	94.0	95.8	93.8	94.2	94.0
TD	93.1	83.6	88.1	86.5	86.0	86.2	81.4	87.4	84.3	74.3	88.2	80.6	63.8	89.1	74.3
CS	96.8	82.1	88.8	94.9	85.1	89.8	88.5	86.1	87.3	87.9	87.1	87.5	76.5	87.7	81.7
STR	98.3	88.5	93.1	92.7	90.3	91.5	89.7	91.2	90.4	84.1	92.1	87.9	80.2	92.8	86.0
AH	97.7	89.0	93.1	96.5	89.6	92.9	94.6	90.8	92.6	91.7	91.2	91.5	83.9	91.7	87.6
IMM	99.0	81.9	89.6	93.3	84.1	88.5	92.4	85.2	88.6	88.7	86.5	87.6	80.5	87.7	83.9
IB	94.0	85.1	89.4	87.5	88.4	88.0	84.8	89.6	87.2	79.9	91.3	85.2	67.2	91.8	77.6
AGE	100.0	35.0	51.9	100.0	46.9	63.9	88.6	54.4	67.4	88.6	60.0	71.6	76.1	66.2	70.8
SB	100.0	50.0	66.7	93.6	52.7	67.4	85.7	53.6	65.9	71.7	59.4	65.0	64.9	64.9	64.9
PHM	100.0	19.4	32.4	100.0	35.9	52.8	91.3	45.7	60.9	85.3	53.7	65.9	64.4	60.3	62.3
DH	97.1	82.9	89.5	93.3	85.7	89.4	92.2	87.0	89.5	86.4	87.9	87.2	80.9	88.7	84.6
SYN	100.0	54.3	70.4	100.0	59.0	74.2	96.8	65.2	77.9	94.9	69.8	80.4	86.8	74.2	80.0
SC	100.0	74.3	85.3	96.9	77.5	86.1	92.1	79.6	85.4	84.4	80.9	82.6	77.2	83.0	80.0
EG	93.8	65.2	76.9	94.7	69.2	80.0	88.5	74.2	80.7	75.0	77.1	76.1	63.3	79.5	70.5
MFS	100.0	16.7	28.6	100.0	34.8	51.6	92.9	46.4	61.9	94.4	53.1	68.0	59.0	60.5	59.7
RE	100.0	33.3	50.0	100.0	39.1	56.3	87.5	50.0	63.6	77.8	50.0	60.9	66.7	58.8	62.5
SS	94.4	77.3	85.0	95.3	80.0	87.0	87.5	80.8	84.0	81.5	81.5	81.5	75.0	84.4	79.4
AW	75.0	21.4	33.3	80.0	26.7	40.0	80.0	26.7	40.0	80.0	26.7	40.0	80.0	42.1	55.2
RRR	0.0	0.0	0.0	0.0	0.0	0.0	0.0	0.0	0.0	0.0	0.0	0.0	0.0	0.0	0.0
average	98.1	79.4	87.8	94.6	82.1	87.9	91.2	83.6	87.2	87.0	84.8	85.9	78.4	86.0	82.0

Table 5: Results per classification ensemble model and research topic (%). Scores greater than 85% or less than 50% are printed in bold or italics, respectively. Research topics are sorted in order of decreasing positive instances (see Table 3).

	Classification ensemble model								
topic	5 agreements	4 Dositives	3 positives	2 positives	I positive				
MIC	17.0	11.8	8.9	5.1	0.0				
PS	7.4	5.1	3.6	2.1	0.0				
NS	3.3	2.3	1.8	1.0	0.0				
TD	11.6	9.0	6.8	4.4	0.0				
CS	7.7	5.5	3.9	2.8	0.0				
STR	8.1	6.0	4.4	2.2	0.0				
AH	5.1	4.4	2.9	2.0	0.0				
IMM	5.8	4.3	3.5	2.2	0.0				
IB	7.5	5.7	4.6	2.5	0.0				
AGE	3.5	2.9	2.1	1.5	0.0				
SB	3.1	2.8	2.5	1.4	0.0				
PHM	3.5	2.9	2.3	1.6	0.0				
DH	2.2	1.5	1.1	0.6	0.0				
SYN	2.2	2.0	1.4	0.9	0.0				
SC	2.0	1.6	1.2	0.8	0.0				
EG	2.2	2.0	1.5	0.9	0.0				
MFS	2.3	2.0	1.6	1.4	0.0				
RE	1.5	1.4	0.9	0.8	0.0				
SS	1.2	1.0	0.8	0.6	0.0				
AW	0.4	0.3	0.3	0.3	0.0				
RRR	0.0	0.0	0.0	0.0	0.0				
average	4.6	3.6	2.7	1.7	0.0				

Table 6: Percentage of omitted instances per classification ensemble model and research topic (%). Research topics are sorted in order of decreasing positive instances (see Table 3).

not for all. To investigate this further, we computed separate average scores for research topics where the aggregate methods perform well or not so well. In particular, as shown in table 8 we computed average scores for research topics for which the 5 agreements method achieved an F-score higher than 75%, i.e. AH, CS, DH, EG, IB, IMM, MIC, NS, PS, SB, SC, SS, STR and TD. Average scores for the remaining research topics are illustrated in table 8. We observe that the best performing aggregate method for the high-performance research topics is 5 agreements, while the best performing method for the low-performance research topics is 2 positives. Due to the sparsity of positive training instances for the low-performance research topics (table 3) the classification models rarely produce positive predictions. Thus, requiring a positive prediction from three, four or five models is a very strict criterion, affecting performance negatively.

Disagreements between manual assignments and automatic predictions were studied by subject matter experts. Analysis revealed that some manual assignments were wrong or incomplete, decreasing the error rate from 2.04% to 1.38%, when using the *5 agreements* combination method and averaging over all research topics.

6. Conclusion & Future Work

We investigated the task of classifying funded grants descriptions into research topics. The task is currently done manually. A precise automatic solution would speed up the process of informing decision makers, while minimising the cost. We investigated five classification models based on feature representations that consist of words, terms and LDA-induced topics. To boost precision, we combined predictions using five different methods. We concluded that

Method	P R		\mathbf{F}_1	omitted instances
5 agreements	98.1	84.4	90.7	6.0
4 positives	94.4	86.5	90.2	4.5
3 positives	91.2	87.5	89.3	3.4
2 positives	86.9	88.4	87.6	2.0
1 positive	78.4	89.1	83.4	0.0

Table 7: Results per prediction combination method for the research topics AH, CS, DH, EG, IB, IMM, MIC,NS, PS,SB, SC, SS, STR, TD (%).

Method	P R		\mathbf{F}_1	omitted instances
5 agreements	98.7	34.4	51.0	2.1
4 positives	97.3	42.5	59.1	1.8
3 positives	89.9	49.0	63.4	1.4
2 positives	84.4	54.4	66.2	1.0
1 positive	70.5	61.1	65.5	0.0

Table 8: Results per prediction combination method for the research topics AGE, AW, MFS, PHM, REG, RRR, SB, SYN (%).

accepting a prediction only when five classification models agree achieved high precision for research topics for which adequate positive instances are available. BBSRC experts used the results to improve manual assignments.

In the future, we will pursue further precision increase, especially for research topics associated with very few positive instances. We plant to consider methods for adapting Support Vector Machines to highly imbalanced datasets (Akbani et al., 2004; Batuwita and Palade, 2013; Wang and Japkowicz, 2010; Tang et al., 2009). The relevant literature suggests various methods for smoothing the effect of the small number of positives, including oversampling and undersampling (Akbani et al., 2004; Batuwita and Palade, 2013; Wang and Japkowicz, 2010; Tang et al., 2009; Chawla, 2005; Wang et al., 2014; Ganganwar, 2012). It is worth applying these methods to the particular problem at hand, in an attempt to improve classification performance for topic labels of very high sparsity of positive training instances.

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