Biomedical Dataset Recommendation

Xu Wang, Frank van Harmelen and Zhisheng Huang Vrije University Amsterdam, De Boelelaan 1105, 1081 HV Amsterdam, The Netherlands

Keywords: Dataset Recommendation, Scientific Datasets.

Abstract: Dataset search is a special application of information retrieval, which aims to help scientists with finding the datasets they want. Current dataset search engines are query-driven, which implies that the results are limited by the ability of the user to formulate the appropriate query. In this paper we aim to solve this limitation by framing dataset search as a recommendation task: given a dataset by the user, the search engine recommends similar datasets. We solve this dataset recommendation task using a similarity approach. We provide a simple benchmark task to evaluate different approaches for this dataset recommendation task. We also evaluate the recommendation task with several similarity approaches in the biomedical domain. We benchmark 8 different similarity metrics between datasets, including both ontology-based techniques and techniques from machine learning. Our results show that the task of recommending scientific datasets based on meta-data as it occurs in realistic dataset collections is a hard task. None of the ontology-based methods manage to perform well on this task, and are outscored by the majority of the machine-learning methods. Of these ML methods only one of the approaches performs reasonably well, and even then only reaches 70% accuracy.

1 INTRODUCTION

Dataset search is an application of dataset retrieval, which is specialization of information retrieval (Kunze and Auer, 2013). Dataset search aims to help people to find datasets they want. Scientists, journalists and decision makers would be typical users of such a dataset search service. A dataset is "a set of related observations which are organized and formatted for a specific purpose" (Chapman et al., 2020). A dataset can be of widely different formats: tables, files, images, structured objects or others (Google Developers, 2021).

As one particular application case of dataset search, a dataset search engine returns datasets based on a user query. A number of such dataset search engines are now operating in practice, for example Google Dataset Search¹ or the Mendeley Data search engine² operated by Elsevier. Such querydriven search engines come with the limitation that the search engine is highly reliant on an appropriate search query presented by the user.

In (Wang et al., 2020), we provided a recommendation paradigm for dataset search, which can be paraphrased as "if you like this dataset, you'll also like that one". This recommendation paradigm gives us the motivation for this paper, with bringing a new paradigm "if user like a query or a dataset, the dataset search engine could recommend a similar dataset to the user". This paradigm could be applied to a dataset search engine in two scenarios: a) when users can provide a search query with an appropriate description, the engine could recommend datasets similar to the search query; b) when user cannot provide that query, the user can provide a dataset as a query, and the search engine will recommend a similar dataset.

In this paper we are aiming to answer following research questions: 1) How to precisely define the task of recommending similar datasets from queries or from other datasets? 2) How to evaluate different approaches to this recommendation task? 3) Can we set up experiments to test this evaluation task in a specific scientific domain?

There are many related works on dataset recommendation. In (Leme et al., 2013), they provide a dataset recommendation task by using the link between dataset and bibliographic domain (e.g. DBLP). In (Ellefi et al., 2016), dataset recommendation works by considering schema overlap between datasets. In (Patra et al., 2020), they provide a dataset recommendation approach with help of publications and interests of researchers. In (Singhal et al., 2013), the au-

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Wang, X., van Harmelen, F. and Huang, Z. Biomedical Dataset Recommendation. DOI: 10.5220/0010521801920199 In Proceedings of the 10th International Conference on Data Science, Technology and Applications (DATA 2021), pages 192-199

IN Proceedings of the Four International Conference of Data Science, rectinology and Applications (DATA 2021), pages 192-1 ISBN: 978-989-758-521-0

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¹https://datasetsearch.research.google.com/

²https://data.mendeley.com/



Figure 1: Overview of main works in this paper.

thors used the context of user's interest to find interesting research datasets. All these approaches differ from the approach we take in this paper, namely providing dataset recommendations based on only the meta-data descriptions of datasets as provided by the authors of the datasets.

We given an overview of our paper in Figure 1, which shows the two parts of our work: the recommendation task and the evaluation of different approaches for this task. The recommendation task will try to recommend similar datasets chosen from candidate datasets for a given target (which could be a dataset or a query). The evaluation will evaluate the recommendation work by checking if the recommended dataset belongs to the same scientific domain as the target. This choice of success-criterion (do we manage to recommend a dataset from the same domain as the target) may sound like a lowbarrier. However, our experimental results in section 5 will actually show that even this seemingly simple success-criterion is far from trivial to achieve.

The main contributions of this paper are: 1) We provide a novel dataset recommendation task. This task recommends similar datasets from a set of candidate datasets given a target that consists of another dataset or a query. We also provide several approaches for this similar dataset recommendation task, by applying a number of similarity approaches. 2) We provide a procedure for evaluating the different approaches to the similar dataset recommendation task. Different from expert judgement, our (simpler) evaluation task is to compare the research domain between the given dataset/query (the target) and the recommended dataset (the output). As argued above, this seemingly easy success-criterion will turn out to be already very hard to achieve by many of the approaches we test. 3) We run experiments to test several approaches to the similar dataset recommendation task for targets in the biomedical domain. The results show that single approach (BM25) outperforms others in our experiments.

2 PRELIMINARIES

In this section we will introduce the different similarity definitions we will use in this paper.

Wu-Palmer. Wu-Palmer is an edge-based semantic similarity metric to calculate the similarity between concepts in hierarchical structure or ontology (Wu and Palmer, 1994). The Wu-Palmer similarity between two concept C_A and C_B is defined as

$$Sim_{WuP}(C_A, C_B) = \frac{2 * Path(R, LCS(C_A, C_B))}{Path(R, C_A) + Path(R, C_B)}$$

where $LCS(C_A, C_B)$ is the least common subsumer of C_A and C_B ; Path(R, n) is the length of the path between the ROOT node *R* and node *n* in the ontology structure.

Resnik. Resnik is a content-based semantic similarity metric to calculate the similarity between concepts with the help of their information content (Resnik, 1995). The Resnik similarity between two concepts C_A and C_B is defined as:

$$Sim_{Res}(C_A, C_B) = -\log p(LCS(C_A, C_B))$$
$$= -\log \frac{\sum_{n \in words(LCS(C_A, C_B))} count(n)}{N}$$

where $LCS(C_A, C_B)$ is the least common subsumer of C_A and C_B ; $p(LCS(C_A, C_B))$ is the probability of $LCS(C_A, C_B)$ in the definition of information content; $words(LCS(C_A, C_B))$ is the set of concepts which are subsumed by $LCS(C_A, C_B)$ in the ontology structure; N is the total number of concepts in the ontology structure.

LDA. LDA (latent Dirichlet allocation) is a generative probabilistic model for a corpus, which could used a set of topics to represent a document (Blei et al., 2003). The LDA similarity between two content fields is calculated by first extracting topics from each content field and then to check the coverage between two sets of topics.

Doc2Vec. Doc2vec is a method to create a numeric representation of a document (Le and Mikolov, 2014). The Doc2vec similarity metric between two content fields is calculated by first transferring each of content fields into pretrained a vector space and then calculate the distance between the two content fields in this vector space.

BM25. BM25 (Best Match 25) is a ranking function that scores documents based on their relevance to a given text query (Robertson et al., 1995). The BM25 approach is used by search engines to rank matching documents according to their relevance to a search query. In this paper, we consider textual content of our target as "query" and use BM25 to find relevant content from candidate datasets.

Bert. Bert (Bidirectional Encoder Representations from Transformers) is a multi-layer neural language representation model (Devlin et al., 2019). We use Bert to compute the similarity between two contents fields by first transferring each content to a vector using Bert's encoder. Then we will calculate the similarity between the two vectors with cosine similarity. This similarity score will be used as the similarity between two content fields.

3 DATASET RECOMMENDATION

In this section we will introduce the dataset recommendation task and a number of different approaches to this task based on similarity metrics between datasets.

3.1 Concept-based Similarity between Datasets

Based on the similarity metrics we introduced above, here we will introduce the similarity approach between datasets. Our concept-based similarity approach between datasets only focuses on the ontology concepts extracted from the textual content of the meta-data fields of datasets. These extracted ontology concepts are the concepts which appear both in the ontology and in the textual content of the meta-data.

Restricting the meta-data fields to the concepts from ontologies will enable the use of ontology-based similarity approaches. In particular, if the ontology captures the terminology of a research domain, the concept extraction of a dataset will contain the concepts from that domain, and will ignore the "noise" terms in the meta-data fields of datasets, which are useless for similarity approach. For example, if we use the UMLS ontology³ for concept extraction, the extracted concepts will always belong to the biomedical research domain, and concepts from other domains (e.g. Computer Science) will be ignored. This will then benefit us when we generate recommendations for biomedical datasets. After the introduction of our concept extraction approach, we will now introduce concept-based similarity between datasets based on the Wu-Palmer and Resnik metrics.

Def. 1 (Concept-based Similarity between Datasets). Given two datasets D_A and D_B , and given a similarity approach between concepts, denoted as sim. Concept-based similarity $ConSim(D_A, D_B)$ between D_A and D_B is defined as follows:

$$ConSim(D_A, D_B) = \frac{\sum \{sim(c_A, c_B) | c_A \in D_A, c_B \in D_B\}}{|D_A| * |D_B|}$$

where c_A is a concept from textual meta-data content of dataset D_A and similarly c_B ; $sim(c_A, c_B)$ is the similarity between c_A and c_B , calculated by similarity metric sim; $|D_A|$ is the number of all concepts in the meta-data of D_A and similarly $|D_B|$.

We can instantiate this generic definition with specific similarity metrics by replacing *sim* with Wu-Palmer or Resnik.

We also define concept-based similarity between a query and a dataset. In this definition, we extract concepts from the query and then apply the same concept-based similarity calculation as above. The definition of concept-based similarity between a query Q and a dataset D is then as follows:

$$ConSim(Q,D) = \frac{\sum \{sim(c_Q, c_D) | c_Q \in Q, c_D \in D\}}{|Q| * |D|}$$
(1)

where c_Q is a concept from content query Q and |Q| is the number of all concepts occurring in Q.

3.2 Content-based Similarity between Datasets

Here we will introduce the content-based similarity between datasets which could be used with ML-based and IR-based approaches. The content we consider here is still the textual meta-data content of datasets, which are title and description.

Def. 2 (Content-based Similarity between Datasets). Given two datasets D_A and D_B . Given a similarity approach between contents, denoted as sim. Content similarity ContSim (D_A, D_B) between D_A and D_B is the similarity between C_A and C_B , where C_A is the textual content of D_A and C_B is the textual content of D_B .

We can then instantiate this definition of contentbased similarity with content metrics such as LDA, Doc2vec, BM25 and Bert. For LDA, BM25 and Bert, we can directly apply them to content-based similarity because all of them are similarity metrics between

³https://www.nlm.nih.gov/research/umls/index.html

contents. For Doc2vec, we consider the whole textual meta-data content of a dataset as a "document" and then calculate the similarity between such "documents" of datasets.

Similar to concept-based similarity between query and dataset, we also have to define content-based similarity between query and dataset. We define contentbased similarity between a query Q and a dataset D as followa:

$$ContSim(Q,D) = sim(C_O,C_D)$$
(2)

where C_Q is the content of query Q; C_D is the textual meta-data content of dataset D; $sim(C_Q, C_D)$ is a similarity metric between contents, which could be LDA, Doc2vec, BM25 and Bert.

3.3 The Similar Dataset Recommendation Task

After introducing the two types of approaches for the similarity between datasets (concept-based and content-based), we will now introduce the task of recommending a similar dataset.

Def. 3 (Similar Dataset Recommendation Task *SDR* for Dataset). Given a target dataset D_T and a list of candidate datasets CL_D , the similar dataset recommendation *SDR* task is to return a sorted list of candidate datasets $SCL_D = \{D_1, D_2, ..., D_n\}$ so that $Sim(D_T, D_1) \ge Sim(D_T, D_2) \ge \ge Sim(D_T, D_n) > 0$, where $D_1, D_2, ..., D_n \in CL_D$. Then $D_1 \in SCL_D$ is the dataset recommended by this task for D_T , denoted as $RD_{D_T}^{SDR}$.

We also define the similar dataset recommendation task for a query.

Def. 4 (Similar Dataset Recommendation Task *SDR* for a Query). Given a target query Q and a list of candidate datasets CL_D , the similar dataset recommendation task *SDR* is to return a sorted list of candidate datasets $SCL_D = \{D_1, D_2,, D_n\}$ so that $Sim(Q, D_1) \ge Sim(Q, D_2) \ge \ge Sim(Q, D_n) > 0$, where $D_1, D_2,, D_n \in CL_D$. Then D_1 in SCL_D is the dataset recommended by this task for Q, denoted as RD_D^{SDR} .

After introducing the similar dataset recommendation task for both dataset and query, we can now turn to a variety of similar dataset recommendation approaches by applying different similarity mentrics to this task.

Also, we identify three types of recommendation approaches, Concept-based and Content-based, and Hybrid one. The Concept-based similar dataset recommendation approach applies concept-based similarity metrics (Wu-Palmer or Resnik) to the dataset recommendation task. The Content-based approach on the other hand applies content-based similarity metrics (LDA, Doc2vec, BM25 or Bert) to the dataset recommendation task. Finally, the Hybrid approach combines concept-based and LDA metrics.

4 EVALUATION FOR DATASET RECOMMENDATION

In this section we will introduce both our evaluation method and the gold standard we will use in the evaluation.

We deliberately choose an evaluation task that is at first sign very easy to achieve: we count a dataset recommendation as correct, if the recommendation is in the same scientific domain as the original target. As we will see, even this apparently rather easy evaluation task will turn out to be difficult to achieve by almost all of the methods that we test in our experiment. A benefit of this choice of evaluation task is that a clear gold standard is available: In many online opensource dataset repositories, such as figshare⁴ or Harvard Dataverse⁵, there are categories or subjects that show the subject-area or research domain and topics of datasets. Two examples are shown in Figure 2 and Figure 3.

Since in our evaluation procedure we will take all our targets from biomedical datasets or queries, the evaluation task comes down to predicting which of the candidate datasets (some 200.000 datasets from all kinds of scientific domains) belong to the biomedical domain.

We can now give a more precise definition of our evaluation task:

Def. 5 (Simple Evaluation Task on Similar Dataset Recommendation for Dataset). Given a list of target datasets L_{D_T} ; a list of datasets L_{D_C} as recommendation candidate datasets; a list of similar dataset recommendation approaches L_{SDR} ; and the gold standard scientific domains Standard_{D'_T} for each D'_T in L_{D_T} . The evaluation task for similarity dataset recommendation is to find the approach SDR' from L_{SDR} so that for every SDR'' $\in L_{SDR}$ (SDR'' \neq SDR'), $|\{RD_{D'_T}^{SDR'} \subseteq_{Domain}$ Standard_{$D'_T}}| \geq |\{RD_{D'_T}^{SDR''} \subseteq_{Domain}$ Standard_{$D'_T}}|, where <math>\subseteq_{Domain}$ means subset relationship on domain area; $RD_{D'_T}^{SDR'} \subseteq_{Domain}$ Standard_{D'_T} $means the domain of dataset <math>RD_{D'_T}^{SDR'}$ is the subset of Standard_{D'_T}.</sub></sub></sub>

⁴https://figshare.com/

⁵https://dataverse.harvard.edu/



Figure 2: Gold Standard Example 1 (figshare).

and similarly for evaluating recommendations based on a target that is a query:

Def. 6 (Simple Evaluation Task on Similar Dataset Recommendation for Query). Given a list of target datasets L_Q ; a list of datasets L_D as recommendation candidate datasets; a list of similar dataset recommendation approaches L_{SDR} , and the gold standard scientific domains Standard_{Q'} for each Q' in L_Q . The valuation task for similarity dataset recommendation is to find the approach SDR' from L_{SDR} so that for every SDR'' $\in L_{SDR}$ (SDR'' \neq SDR'), $|\{RD_{Q'}^{SDR'} \subseteq_{Domain} Standard_{Q'}\}| \geq |\{RD_{Q'}^{SDR''} \subseteq_{Domain} Standard_{Q'}\}|$, where \subseteq_{Domain} means subset relationship on domain area; $RD_{Q'}^{SDR'} \subseteq_{Domain} Standard_{Q'}$ means the domain of dataset $RD_{Q'}^{SDR'}$ is the subset of Standard_{Q'}.

In all our evaluation matching between two domains, we also consider synonyms of the domain. This means that we do not only do exact matching, but also match one domain to the synonym of that domain.

5 EXPERIMENTS

In this section we will introduce our experiments.

5.1 Preparation

First off, we will introduce the datasets and the gold standard used in our experiments.

Target Datasets. Thanks to our colleagues from Elsevier, all the datasets we used are from the Elsevier data repository⁶. All the target datasets in our experiments are 190 datasets derived from Elsevier

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Figure 3: Gold Standard Example 2 (Harvard Dataverse).

DataSearch⁷ based on 19 biomedical queries (shown in Table 1), with the top10 datasets per query. These queries all come from a carefully validated set of test queries for Elsevier's Mendeley data search engine. As a result, we can be certain that all of these 190 datasets are indeed from the biomedical domain, as they are the top10 results on 19 validated biomedical queries.

Candidate Datasets. We use 209.998 datasets from Mendeley Data⁸. These datasets are randomly selected from about 18 million datasets in the Elsevier data repository. Recommendation datasets in our experiments must be selected from these \approx 210k candidate datasets. There are about 600 domains for these candidate datasets without considering synonym of domain.

These datasets contain textual meta-data (title and description of the dataset), topic meta-data (scientific domain of the dataset) and so on. In the recommendation approach, we only consider the textual metadata (title and description) of datasets. We also use the "topic" meta-data, (the "subjectArea" in the metadata schema of the datasets) only in the evaluation. This is because of the particularity of our evaluation approach. Our evaluation approach is to evaluate the scientific research domain. In the metadata of the dataset, the "topic" meta-data is the scientific research domain. Therefore, we only consider topic in the evaluation approach, not in the recommendation approach.

Training Datasets. For training the ML-based distance metrics described above, we used ≈ 21 M datasets (21.110.372 to be precise) as our training corpus. We obviously took care that this training set is disjoint from the 210k candidate datasets, so as to avoid data leakage from the training phase in our experiment.

⁶https://data.mendeley.com/research-data/ ?repositoryType=NON_ARTICLE_BASED_ REPOSITORY

⁷https://datasearch.elsevier.com/

⁸https://data.mendeley.com/

Query	Content	Query	Content
E2	Protein Degradation mechanisms	E54	glutamate alcohol interaction
E7	oxidative stress ischemic stroke	E66	calcium signalling in stem cells
E8	middle cerebral artery occlusion mice	E67	phylogeny cryptosporidium
E17	Risk factors for combat PTSD	E68	HPV vaccine efficacy and safety
E26	mab melting temperature	E78	c elegans neuron degeneration
E28	mutational analysis cervical cancer	E79	mri liver fibrosis
E31	metformin pharmacokinetics	E80	Yersinia ruckeri enteric red mouth disease
E35	prostate cancer DNA methylation	E89	Electrocardiogram variability OR ECG variability
E50	EZH2 in breast cancer	E94	pinealectomy circadian rhythm
E88	Flavonoids cardiotoxicity		

Table 1: The corpus of 19 biomedical queries.

Gold Standard. According to what we discussed in last section, we consider the categories or subjects of datasets as gold standard for evaluation task. In the \approx 210k candidate datasets, there are 60.484 datasets which are labeled by "None" subject or "Uncategorized".

Evaluation. We use check if domain of recommended dataset is same or similar to the one of target datasets by hand. This is doable and easy to do because it's not hard to check if two domains are same or similar for human (not even be a domain expert).

5.2 Experimental Set-up

In this part we will introduce the set-up of our two evaluation experiments. In total we use eight dataset recommendation approaches for these experiments:

- Two concept-based recommendation (with Wu-Palmer and Resnik)
- Two hybrid recommendation (with LDA+Wu-Palmer and LDA+Resnik)
- Four content-based recommendation (with LDA, Doc2vec, BM25 and Bert_Base)

Exp1: Evaluation of Dataset Recommendation Approaches from Dataset Targets. In this experiment, we will evaluate the dataset recommendation approaches when given a dataset as the input target. Given a list of recommendation approaches, the pipeline of Exp1 is:

- 1. Using 19 biomedical queries to get 190 target candidates: We send each query to the Elsevier Datasearch API, and we select the top 10 returned datasets as target candidates.
- 2. For each recommendation approach *SDR* in the list of recommendation approaches given above:
 - (a) For each target dataset, using recommendation approach *SDR* to find the Top1, Top5 and Top10 similar datasets from \approx 210k candidate datasets.

- (b) We consider a dataset recommended by *SDR* as a success-count when the domain of this dataset meets the domain of the target dataset.
- 3. The recommendation approach with the best performance is the one with the highest successcount over all 190 target candidates.

Exp2: Evaluation of Dataset Recommendation Approaches from Query Targets. Different from Exp1, we will here recommend similar datasets from a given query as target. Given a list of recommendation approaches, the pipeline of Exp2 is:

- 1. Using the 19 biomedical queries as target queries.
- 2. For each recommendation approach *SDR* from the list of approaches given above:
 - (a) For each target query, we use *SDR* to find the Top1, Top5 and Top10 similar datasets from \approx 210k candidate datasets.
 - (b) We consider a dataset recommended by *SDR* as a success-count when the domain of this similar dataset meets the domain of the target query.
- 3. The recommendation approach with best performance is the one with highest success-count over 19 target queries.

5.3 Results and Analysis

In this section we will show the results of the two experimental scenarios that we set up above. In Table 2 and Table 3, we show the results of two experimental scenarios for both Exp1 and Exp2:

- Scenario 1: consider all TopN datasets, whether or not they contained subject-area metadata.
- Scenario 2: consider only TopN datasets with subject-area metadata.

Scenario 1. In the first experimental scenario (shown in Table 2), we consider all the TopN similar datasets for each target dataset or query, even if some similar datasets do not contain the subject-area in their meta-data. In such a case, we take the conservative position, and consider it a failure if the recommendation approach recommends a dataset that has

		Тор	10 Judgen	nent	Тор	5 Judgen	nent	Top1 Judgement		
Experiment	Measure	success	All		success	All		success	All	
			total	fraction		total	fraction		total	fraction
	Wu-Palmer	772	1660	0.47	464	830	0.56	92	166	0.55
	Resnik	859	1660	0.52	514	830	0.62	96	166	0.58
	LDA+Wup	674	1700	0.4	506	850	0.6	41	170	0.24
Evn1 (Dataset nair)	LDA+Res	429	1700	0.25	359	850	0.42	19	170	0.11
Expr (Dataset pair)	BM25	1351	1900	0.71	744	950	0.78	143	190	0.75
	Bert_Base	1465	1900	0.77	800	950	0.84	159	190	0.84
	LDA	1360	1900	0.72	767	950	0.81	154	190	0.81
	Doc2vec	941	1900	0.5	582	950	0.61	144	190	0.76
	Wu-Palmer	83	190	0.44	44	95	0.46	6	19	0.32
	Bert_Base	118	190	0.62	66	95	0.69	11	19	0.58
Even? (Quary dataset nair)	BM25	151	190	0.79	83	95	0.87	17	19	0.89
Exp2 (Query-uataset pair)	Resnik	64	190	0.34	39	95	0.41	6	19	0.32
	LDA	60	190	0.32	34	95	0.36	8	19	0.42
	Doc2vec	123	190	0.65	76	95	0.8	16	19	0.84

Table 2: Scenario 1: Top10, Top5 and Top1 results.

Table 3: Scenario 2: Top10, Top5 and Top1 results.(Only consider datasets with subject-area).

		Top10 Judgement			Top5 Judgement			Top1 Judgement			
Experiment	Measure	success	cess Only-SubjectArea		success	Only-SubjectArea		success	ess Only-SubjectArea		
			total	fraction		total	fraction		total	fraction	
	Wu-Palmer	772	1253	0.62	464	760	0.61	92	163	0.56	
	Resnik	859	1306	0.66	514	795	0.65	96	164	0.59	
	LDA+Wup	674	1264	0.53	506	839	0.6	41	170	0.24	
Evn1 (Dataset nair)	LDA+Res	429	1126	0.38	359	817	0.44	19	170	0.11	
Exp1 (Dataset pair)	BM25	1351	1715	0.79	744	948	0.78	143	190	0.75	
	Bert_Base	1465	1736	0.84	800	950	0.84	159	190	0.84	
	LDA	1360	1694	0.8	767	950	0.81	154	190	0.81	
	Doc2vec	941	1539	0.61	582	940	0.62	144	190	0.76	
	Wu-Palmer	83	155	0.54	44	93	0.47	6	19	0.32	
	Bert_Base	118	167	0.71	66	95	0.69	11	19	0.58	
Evn2 (Query-dataset nair)	BM25	151	169	0.89	83	95	0.87	17	19	0.89	
Exp2 (Query-uataset pair)	Resnik	64	154	0.42	39	95	0.41	6	19	0.32	
	LDA	60	159	0.38	34	95	0.36	8	19	0.42	
	Doc2vec	123	150	0.82	76	95	0.8	16	19	0.84	

no subject-area in their meta-data. In Table 2, there is a difference between the "total" number of judgements performed for each approach. This is because in the concept-based approaches, no concept could be extracted for some datasets or queries. We have removed these cases from the counts in the table, since no meaningful similarity score can be computed. Table 2 shows that the Bert-based approach outperforms all others in Exp1, reaching 80%, while LDA-based approach and BM25-based approaches also perform well by reaching about 80% accuracy. The table also shows that the concept-based approaches all seriously underperform on the task when using datasets as the original target.

The table also shows the reason why we test they Hybrid approach by combining LDA with conceptbased approaches. As Table 2 shows, the LDA approach performs very well on Exp1 (third best performance), which means that LDA can extract relatively correct topics in the majority of cases. This prompted the hypothesis: extracted topics of datasets by LDA could be used as concepts to help the concept-based approach. However, the data in Table 2 shows that this hypothesis was false: the hybrid approach of LDA plus either Wu-Palmer or Resnik performs no better (and sometimes even substantially worse) than Wu-Palmer or Resnik on their own.

In Exp2 in Table 2, we test 6 approaches without considering the Hybrid approach. Because the LDA approach doesn't perform well in Exp2, we do not expect it to boost the concept-based approaches. In Exp2, the BM25-based approach outperforms others by reaching 85% accuracy. The Bert-based and Doc2vec-based approaches also perform well. Again, we see a substantial underperformance of either of the two concept-based approaches.

Scenario 2. Scenario 2 is a revised version of scenario 1, in which we only consider TopN datasets that mention the subject-area in their meta-data, so that we can be certain about the correctness of the recommendation. For Exp1 of this scenario, the results of Table 3 show that (again) both the Bert-based and the LDA-based approach perform well, reaching 80% accuracy, with the Bert-based approach outperforming others. Also, the BM25-based approach performs well by reaching 75%. For Exp2, the BM25-based approach even outperforms all others by reaching about 89% accuracy. The Doc2vec-based approach also performs well with reaching about 80% accuracy.

Overall, we can conclude as the final result that BM25-based approach performs best in both Exp1 and Exp2 among all the approaches we tested.

6 CONCLUSION

In this paper, we provided a novel task for recommending scientific datasets. This task recommends similar dataset based on a target that is either a query or another dataset. Based on this task, we introduced several approaches, using some popular similarity methods. Also, we executed experiments to evaluate these approaches on biomedical datasets.

There are a number of lessons that we can draw from our experiments: 1) it is notable that the task of recommending similar datasets based on only the meta-data from these datasets, and possibly a query, is much harder than one might expect, with even the best performing methods rarely scoring higher than 80%. 2) we see that the semantic, ontology-based methods are not capable of solving this task, and that the statistical methods from machine learning or information retrieval far outperform the semantic methods. Even boosting the ontology-based methods with a machine learning method did not give an acceptable result. 3) our results show that the BM25-based approach performs well on the task of dataset recommendation from both a query target and a dataset target, reaching 70% accuracy in our experiments.

In this paper, we have used only textual meta-data for dataset recommendation. Given the high variability in the syntax and semantics of the content of datasets (ranging from gene sequences to geographical maps to spreadsheets with financial data), it is near impossible to use this dataset contents. Nevertheless, there are other signals that could be considered for similar dataset recommendation. Authors of datasets would be one of these signals: the co-author network could be used for dataset recommendation if we could match dataset's author into this network. The Open Academic Graph (OAG) is a very popular and large knowledge graph that unifies two separate billion-scale academic graphs MAG and AMiner (Microsoft, 2021; Sinha et al., 2015; Tang et al., 2008). We will investigate in future work if this resource can be exploited to improve the task of recommending similar datasets that we defined in this paper.

ACKNOWLEDGEMENT

This work has been funded by the Netherlands Science Foundation NWO grant nr. 652.001.002 which is also partially funded by Elsevier. The first author is funded by by the China Scholarship Council (CSC) under grant number 201807730060.

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