

MeSHup: A Corpus for Full Text Biomedical Document Indexing

Xindi Wang^{1,3}, Robert E. Mercer^{1,3}, and Frank Rudzicz^{2,3,4}

¹Department of Computer Science, University of Western Ontario, London, Ontario, Canada

²Department of Computer Science, University of Toronto, Toronto, Ontario, Canada

³Vector Institute for Artificial Intelligence, Toronto, Ontario, Canada

⁴Unity Health Toronto, Toronto, Ontario, Canada



BACKGROUND

MEDLINE

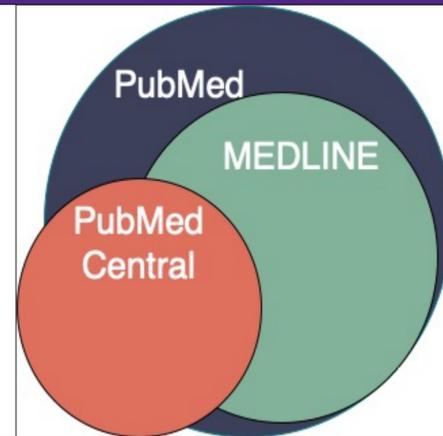
- Core database, contains more than 28 million references to a specific set of journals in biomedical science.

PubMed

- A free access search engine for abstracting and indexing biomedical citations.
- Comprises more than 33 million citations for biomedical literature from MEDLINE (as of Apr. 2022).
- Links to articles from publisher's websites and PubMed Central.

PubMed Central (PMC)

- Full-text archive for biomedical and life sciences journal articles.
- 7.9 million articles are archived in PMC



MOTIVATIONS

- Existing corpora only provide the title and abstract, while human annotators review the full text articles.
- Previous work focused on text information but are less concerned with metadata.

DATA SAMPLE

```

{"articles": [
  {"PMID": "27976717",
   "TITLE": "Temporal pairwise spike
             correlations fully capture
             single-neuron information",
   "ABSTRACT": "To crack the neural
                code and read out the
                information neural spikes
                convey, [...] ",
   "INTRO": "Throughout the central
              nervous system of a mammalian
              brain [...] ",
   "METHODS": "Deriving the correlation
                theory of neural information [
                ...] ",
   "RESULTS": "We are interested in the
                information contained in a
                spike train r(t) about a
                stimulus s(t)[...]",
   "DISCUSS": "The list of spike timing
                features that have been
                implicated in neural coding
                includes [...] ",
   "FIG_CAPTIONS": "Dimensionality of
                    neural information coding [...]",
   "TABLE_CAPTIONS": "Parameter sets
                      across neuron models. [...]",
   "JOURNAL": "Nature communications",
   "YEAR": "2016",
   "DOI": "10.1038/ncomms13805",
   "AUTHORS": [
     "Amadeus, Dettner",
     "Sabrina, Munzberg",
     "Tatjana, Tchumatchenko" ],
   "MeSH": {
     "D000200": "Action Potentials",
     "D008959": "Models, Neurological",
     "D009474": "Neurons",
     "D059010": "Single-Cell Analysis"
   },
   "CHEMICALS": "None",
   "SUPPLMeSH": "None"
 },
  ...
],
  ...
}

```

DATASET CONSTRUCTION

Data resources

- PubMed Central Open Access in BioC format (BioC-PMC)
- MEDLINE / PubMed Annual Baseline Repository (MBR)

Constrains

- Articles indexed by human annotators only
- English articles only

Information extracted from BioC-PMC

- Eight BioC sections are selected to construct the new corpus: title, abstract, introduction, methods, results, discussion, figure captions, and table captions.

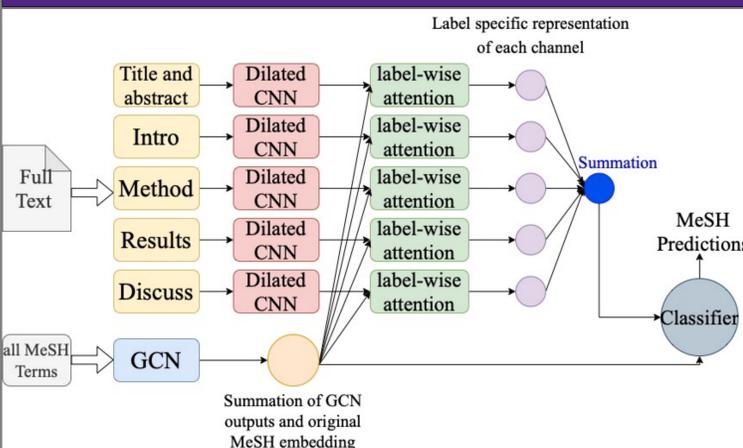
Information extracted from MBR

- Metadata: PMID, authors, journal name, publication year, DOI, MeSH terms, supply MeSH, and chemical list.

MeSHup CORPUS

- Contains a set of 1, 342, 667 biomedical documents.
- Each article has full textual information and metadata associated with it.

BASELINE MODEL AND RESULTS



Bipartition evaluation	Methods		
	Titles and Abstracts	Full Texts	
Example based	EBF	0.183	0.259
	EBP	0.503	0.588
	EBR	0.112	0.166
Micro-averaged	MiF	0.177	0.259
	MiP	0.473	0.604
	MiR	0.110	0.164
Macro-averaged	MaF	0.362	0.367
	MaP	0.798	0.810
	MaR	0.234	0.237

Table 3: Comparison using only titles and abstracts and full texts across bipartition evaluation. Bold: best scores in each row.

CONTRIBUTIONS

- We release a large-scale annotated MeSH indexing corpus, MeSHup.
- We train an end-to-end multichannel model that incorporates different sections of the full text article to show that full texts are more informative in the MeSH indexing tasks compared to the titles and abstracts only