

# Diving for Treasure in a Sea of Scientific Literature

## Extracting Scientific Information from Free Text Articles

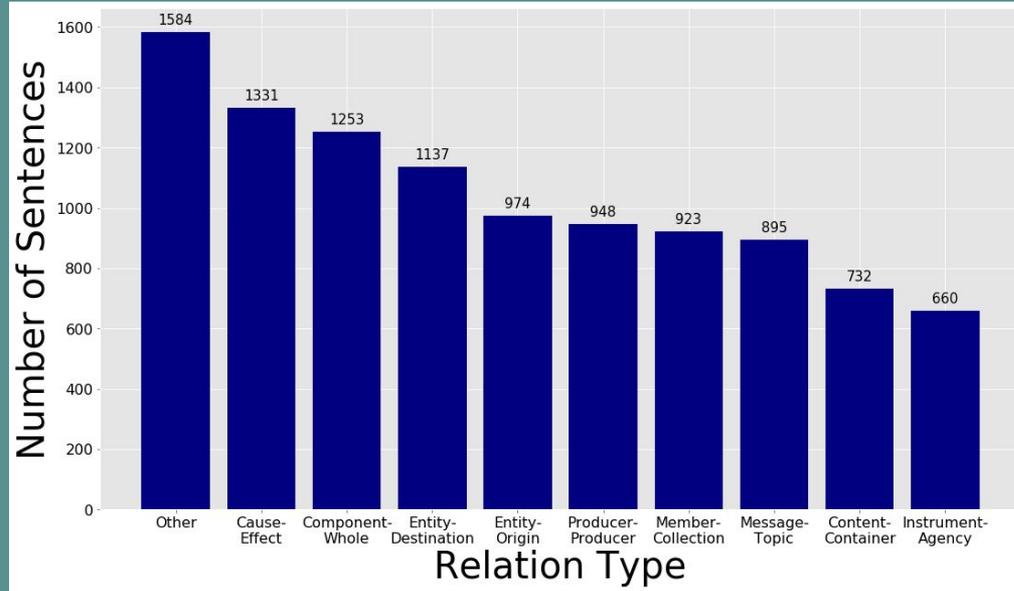
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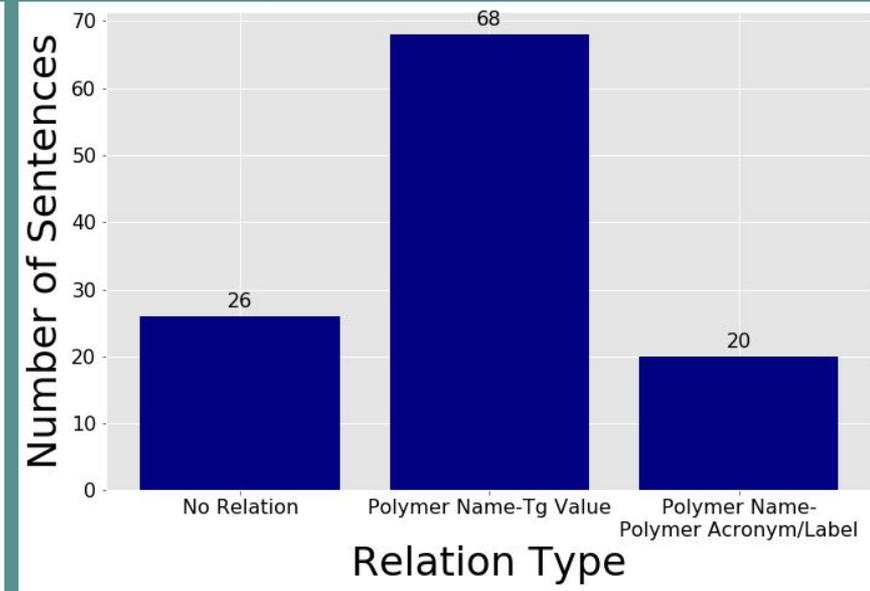
**Why?**

# Dataset Distribution

## Noun Dataset Distribution



## Polymer Dataset Distribution



# Dataset Distribution

## Noun Data

- The dataset was provided by SemEval Task 8
- 8000 training sentences, 2717 testing sentences, each labelled with entities, and the relation type.

Example: 'Wings' and 'Fly'  
Cause-Effect Relationship

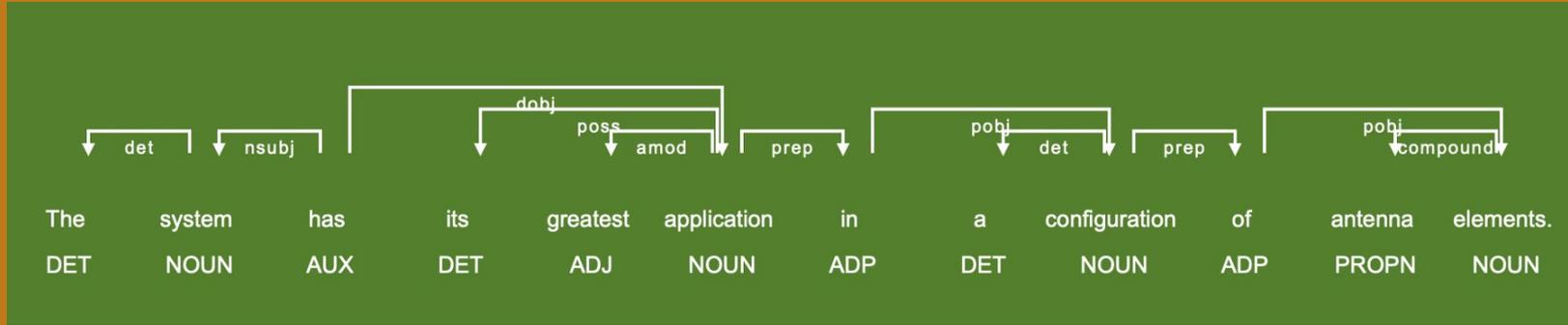
## Polymer Data

- The dataset was provided by *Macromolecules* journal
- 114 sentences, each labelled with entities, and the relation type.

Example: '5 7-dibutyl-1  
3-dehydroadamantane' and '3'  
Polymer Name and Label Relationship

# What is a Dependency Parser?

Extracts relationships between words based on parts of speech and semantic relationships

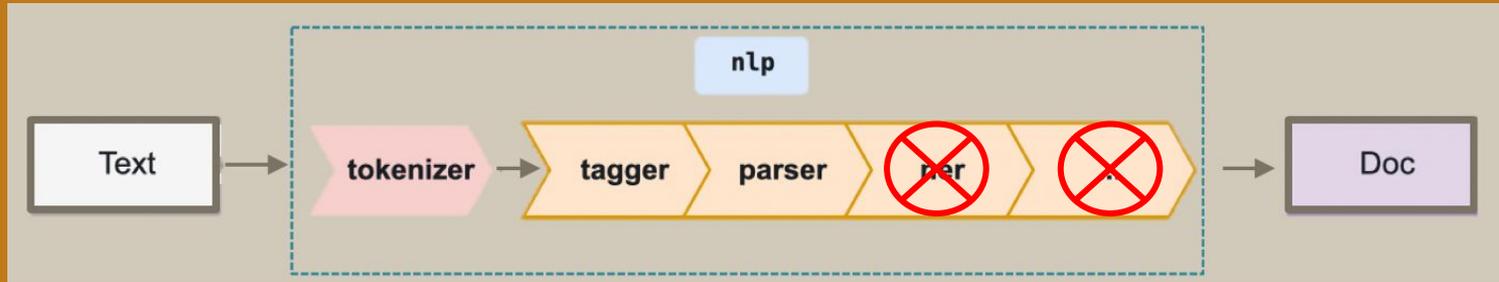
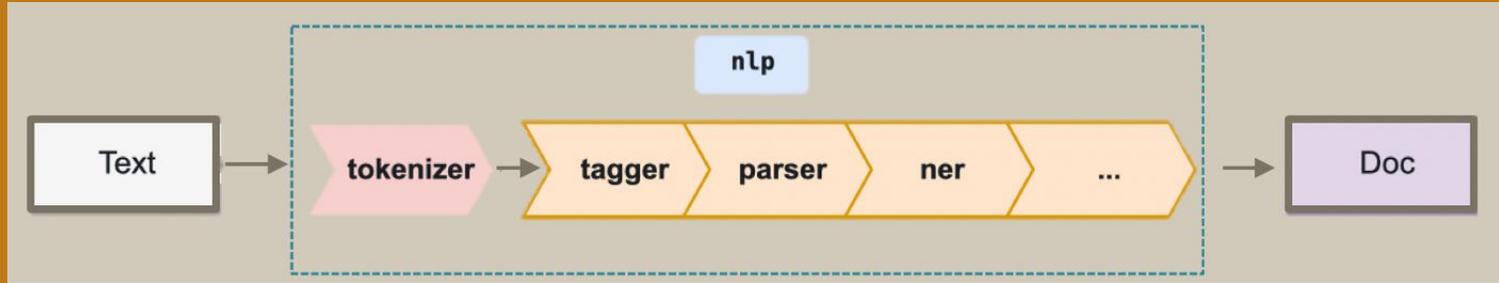


## Named Entity

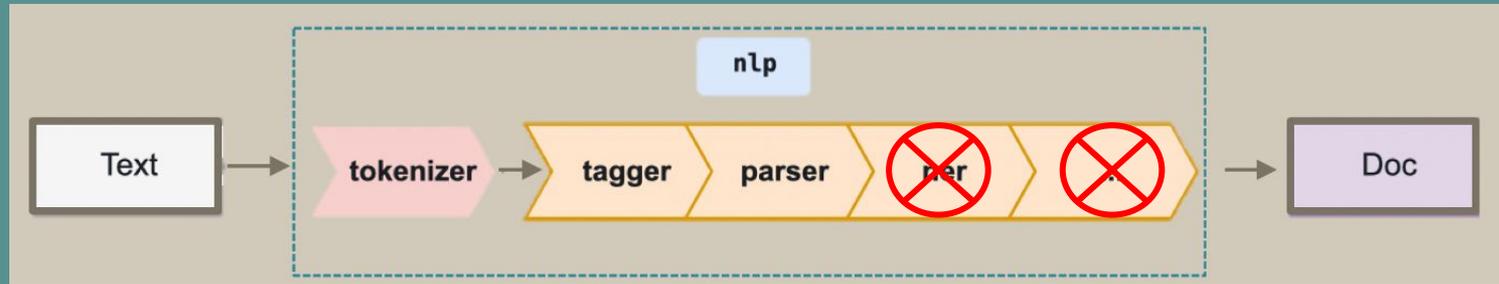
**Tagger:** Extracts relationships between words based on word type (person, place, location, etc)



# spaCy NLP Pipeline



**INPUT SENTENCE:** 'The system as described above has its greatest application in an arrayed configuration of antenna elements .'



**OUTPUT:** [('configuration', 'Whole', 12), ('elements', 'Component', 15)]

# Customizations - Tokenizer and Word Embedding

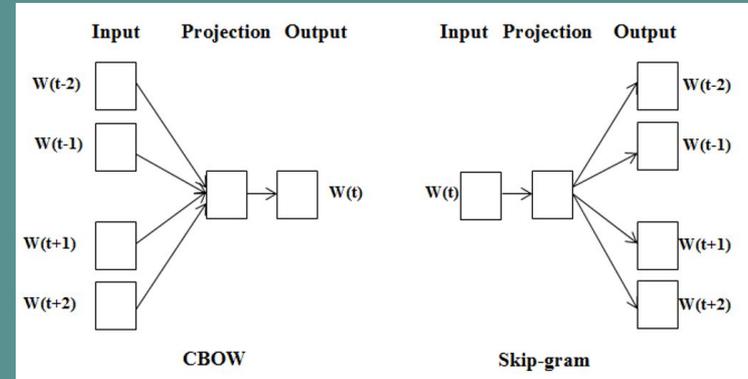
**Original Sentence:** The **glass-to-rubber** transition of poly (**BC-co-BS**) copolymers was investigated by DSC on melt quenched samples.

Default Tokenizer:

*(The, **glass**, -, **to**, -, **rubber**, transition, of, poly, (**BC**, -, **co**, -, **BS**), copolymers, was, investigated, by, DSC, on, melt, quenched, samples, .)*

Custom Tokenizer:

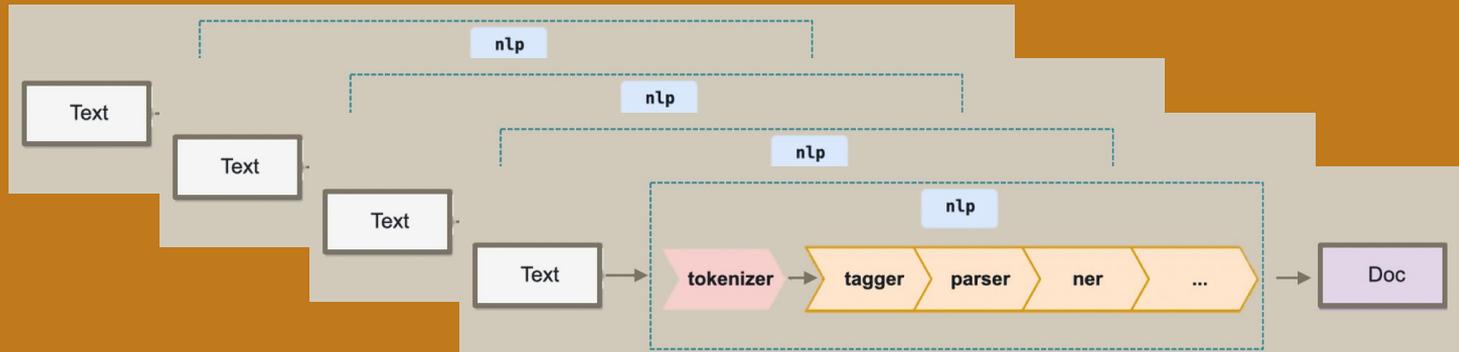
*(The, **glass-to-rubber**, transition, of, poly, (**BC-co-BS**), copolymers, was, investigated, by, DSC, on, melt, quenched, samples, .)*



# Scaling the pipeline

- Our models identify relations in fragments of text
- There are millions of papers to be processed

⇒ Scale by slicing the data and parallelizing across cores and nodes (with Parsl)



- RCC Midway Cluster

```
import parsl
import time
import os
from parsl.app.app import python_app, bash_app
from parsl.providers import SlurmProvider
from parsl.launchers import SrunLauncher
from parsl.addresses import address_by_hostname
from parsl.executors import HighThroughputExecutor
from parsl.config import Config

num_nodes = 1
num_cores_per_node = 32
config = Config(
    executors=[
        HighThroughputExecutor(
            label='Midway_HTEX',
            worker_debug=False,
            address=address_by_hostname(),
            cores_per_worker=1,
            max_workers=num_cores_per_node,
            provider=SlurmProvider(
                'gm4',
                launcher=SrunLauncher(),
                nodes_per_block=num_nodes,
                cores_per_node=num_cores_per_node,
                init_blocks=1,
                max_blocks=1,
                exclusive=False,
                scheduler_options='#SBATCH --qos=gm4-cpu',
                worker_init='source activate freetext',
                walltime='12:00:00'
            ),
        ),
    ],
)

parsl.load(config)
```

```
@python_app

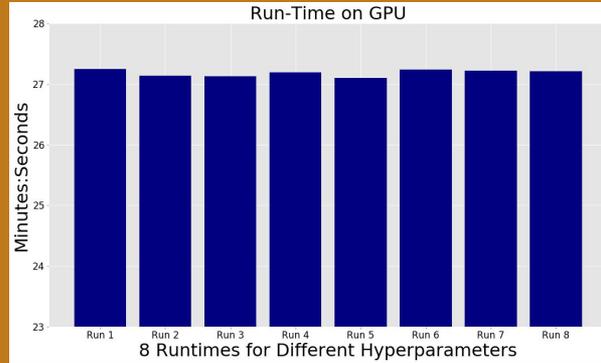
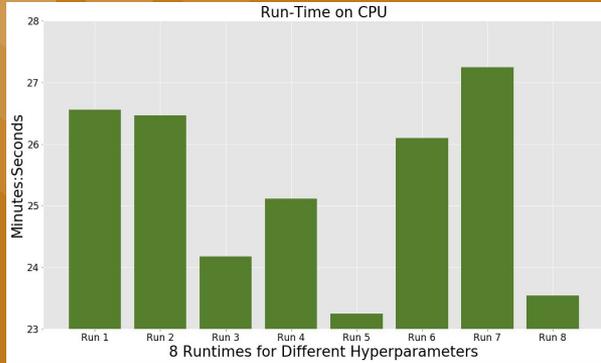
def generate(string):
    from datetime import datetime
    import spacy
    tic = datetime.now()
    nlp = spacy.load("./nlp_model")
    result = []
    docs = nlp(string)
    result.append([(t.text, t.dep_, t.idx) for t in docs if t.dep_ != "-" and t.dep_
!= "ROOT"])
    toc = datetime.now()
    return tic, toc, result

import pandas as pd
df = pd.read_table('./polymer.txt', delim_whitespace=False, names=('A')) #reading
training data
data = [_ for _ in df['A']]
untoken = []
for sent in data:
    delta = sent.split()
    string = ''
    for i in delta:
        string = string + i + " "
    untoken.append(string)

# Wait for all apps to finish and collect the results
out = list()
for i in untoken:
    out.append(generate(i))

tics, tocs, outputs = list(), list(), list()
for i in out:
    tic, toc, result = i.result()
    tics.append(tic)
    tocs.append(toc)
    outputs.append(result)

# Print results
with open("nodes1_results.txt", "w") as result:
    result.write(str(outputs))
    result.write(str(max(tocs) - max(tics)))
```



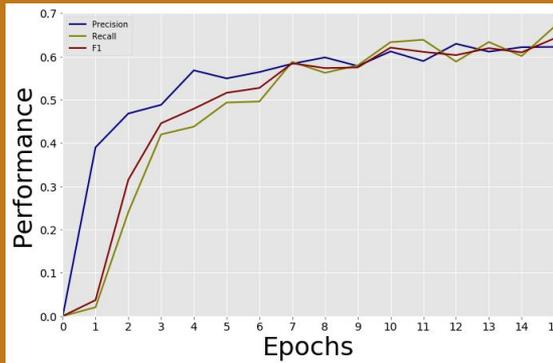
## CPU vs GPU

- Used Model 3 and default CBOW parameters
- spaCy not optimized for GPU acceleration
- Fastest CPU runtime for training is 23 minutes

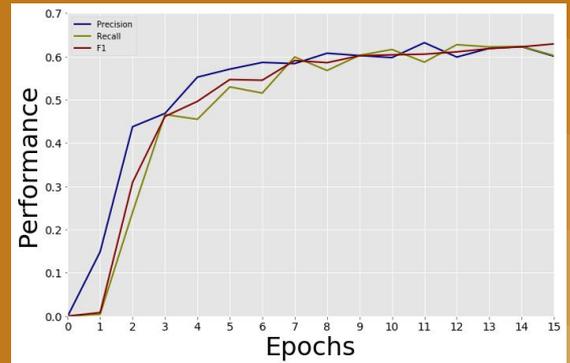
## Performance

- Performance over 15 epochs for Skipgram and CBOW with default hyperparameters for Model 3

Performance Over 15 Epochs for CBOW

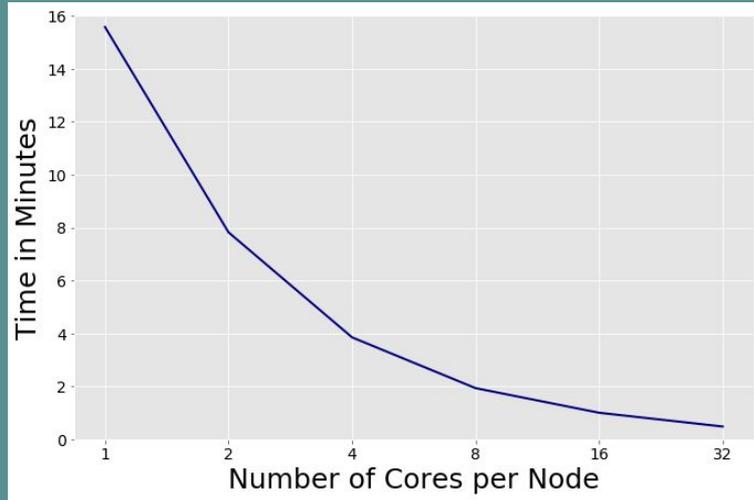


Performance Over 15 Epochs for Skip-gram

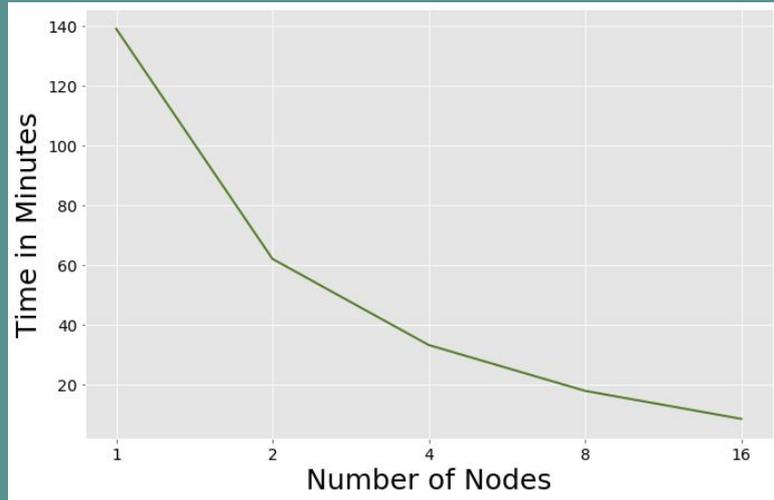


# Scaling

Per Core Scaling



Per Node Scaling



# Conclusion



spaCy



ChemDataExtractor

