Domain-Specific NLP

Introduction

Pretrained (Large) Language Models are trained on content crawled over the internet, books, reports and news papers and are, hence **are open-domain**.

A textual domain is the distribution over language characterizing a given topic or genre [1].

• You are more likely to see the word "integer" in computer science than in news papers.

Contents

1. Domain-Specific Models

- a. Don't stop pre-training
- b. Specialized models (BioBERT, SciBERT, Galactica)

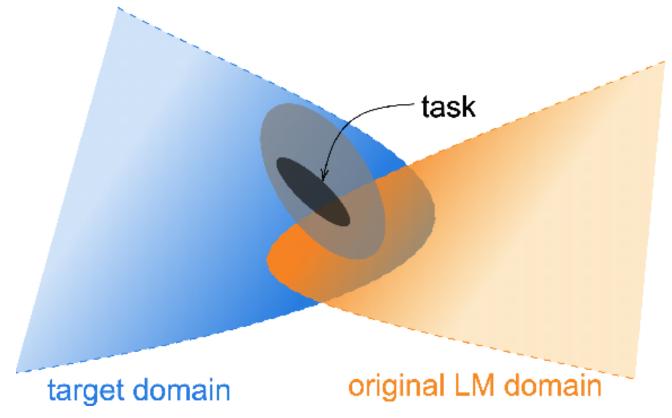
2. Unsupervised Classification Models

- a. Représentations out-of-the-box: limitations
- b. SimCSE, E5, GTE...

3. Learning Long-Range Dependencies

- a. Long-range attention models
- b. State-space models: \$4

Don't Stop Pretraining: Adapt Language Models to Domains and Tasks. [1]



Domain	Pretraining Corpus	# Tokens	Size	$\mathcal{L}_{RoB.}$	$\mathcal{L}_{ exttt{DAPT}}$
BIOMED	2.68M full-text papers from S2ORC (Lo et al., 2020)	7.55B	47GB	1.32	0.99
CS	2.22M full-text papers from S2ORC (Lo et al., 2020)	8.10B	48GB	1.63	1.34
News	11.90M articles from REALNEWS (Zellers et al., 2019)	6.66B	39GB	1.08	1.16
REVIEWS	24.75M AMAZON reviews (He and McAuley, 2016)	2.11B	11GB	2.10	1.93
RoBERTA (baseline)	see Appendix §A.1	N/A	160GB	[‡] 1.19	-

Table 1: List of the domain-specific unlabeled datasets. In columns 5 and 6, we report ROBERTA's masked LM loss on 50K randomly sampled held-out documents from each domain before $(\mathcal{L}_{ROB.})$ and after (\mathcal{L}_{DAPT}) DAPT (lower implies a better fit on the sample). \ddagger indicates that the masked LM loss is estimated on data sampled from sources similar to ROBERTA's pretraining corpus.



Figure 2: Vocabulary overlap (%) between domains. PT denotes a sample from sources similar to ROBERTA's pretraining corpus. Vocabularies for each domain are created by considering the top 10K most frequent words (excluding stopwords) in documents sampled from each domain.

			Additional Pretraining Phases		
Domain	Task	ROBERTA	DAPT	TAPT	DAPT + TAPT
BIOMED	СнемРкот †RCT	81.9 _{1.0} 87.2 _{0.1}	84.2 _{0.2} 87.6 _{0.1}	82.6 _{0.4} 87.7 _{0.1}	84.4 _{0.4} 87.8 _{0.1}
CS	ACL-ARC SCIERC	$63.0_{5.8} \\ 77.3_{1.9}$	75.4 _{2.5} 80.8 _{1.5}	67.4 _{1.8} 79.3 _{1.5}	75.6 _{3.8} 81.3 _{1.8}
News	HyperPartisan †AGNews	$86.6_{0.9}$ $93.9_{0.2}$	88.2 _{5.9} 93.9 _{0.2}	90.4 _{5.2} 94.5 _{0.1}	90.0 _{6.6} 94.6 _{0.1}
REVIEWS	[†] HELPFULNESS [†] IMDB	$65.1_{3.4} \\ 95.0_{0.2}$	66.5 _{1.4} 95.4 _{0.1}	68.5 _{1.9} 95.5 _{0.1}	68.7 _{1.8} 95.6 _{0.1}

Table 5: Results on different phases of adaptive pretraining compared to the baseline RoBERTa (col. 1). Our approaches are DAPT (col. 2, §3), TAPT (col. 3, §4), and a combination of both (col. 4).

"[..] the word distributions of general and biomedical corpora are quite different, which can often be a problem for biomedical text mining models." [2]

: 1. List of text corpora used for BioBERT

Corpus	# of words (B)	Domain
English Wikipedia BooksCorpus	2.5B 0.8B	General General
PubMed Abstracts PMC Full-text articles	4.5B 13.5B	Biomedical Biomedical

"We showed that **pre-training BERT on biomedical corpora is crucial in applying it to the biomedical domain**. Requiring minimal task-specific architectural modification, **BioBERT outperforms previous models on biomedical text mining tasks** such as NER, RE and QA."

Task	Dataset	BIOBERT	SCIBERT
	BC5CDR	88.85	90.01
NER	JNLPBA	77.59	77.28
	NCBI-disease	89.36	88.57
REL	ChemProt	76.68	83.64

Table 2: Comparing SciBERT with the reported BioBERT results on biomedical datasets.

NB: SciBERT was trained on curated textual data; not trained on code or script for example---at leat not trained directly and purposefully on this kind of data

"Unlike search engines, language models can potentially store, combine and reason about scientific knowledge." [4]

- Specialized models (BioBERT, SciBERT, Galactica) were trained on a rather small highly curated dataset.
- The data was standardized in markdown format.

14

Modality	Entity	Sequence	
Text	Abell 370	Abell 370 is a cluster	
IATEX	Schwarzschild radius	$r_{s} = \frac{2GM}{c^2}$	$r_s = rac{2GM}{c^2}$
Code	Transformer	<pre>class Transformer(nn.Module)</pre>	200000
SMILES	Glycine	C(C(=0)0)N	H'O N'H
AA Sequence	Collagen $lpha$ -1(II) chain	MIRLGAPQTL	0 ⁰⁰⁰ 00000000000000000000000000000000
DNA Sequence	Human genome	CGGTACCCTC	C G G T A C C C T G C C A T G G G A

Table 1: Tokenizing Nature. Galactica trains on text sequences that represent scientific phenomena.

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- 1. **Citations:** wrapped with special reference tokens [START_REF] and [END_REF].
- 2. **Step-by-Step Reasoning:** wrapped with a working memory token <work, mimicking an internal working memory context.
- 3. **Mathematics:** for mathematical content, with or without LaTeX, ASCII operations are splitted into individual characters. Parentheses are treated like digits. The rest of the operations allow for unsplit repetitions. Operation characters are !"#\$%&'*+,-./:;<=>?^_'| and parentheses are ()[]{}.

- 4. **Numbers:** splitted into individual tokens. For example 737612.62 > 7,3,7,6,1,2,.,6,2.
- 5. **SMILES formula:** wrapped with [START_SMILES] and [END_SMILES] and tokenized absed on characters. Similarly [START_I_SMILES] and [END_I_SMILES] is usedwhere isomeric SMILES is denoted.
- 6. **Amino acid sequences:** wrapped with [START_AMINO] and [END_AMINO] and apply character-based tokenization, treating each amino acid character as a single token. For example, MIRLGAPQTL -> M,I,R,L,G,A,P,Q,T,L.

1. **DNA sequences:** tokenized based on characters and wrapped inside [START_DNA] and [END_DNA]. For example, CGGTACCCTC -> C, G, G, T, A, C, C, C, T, C.

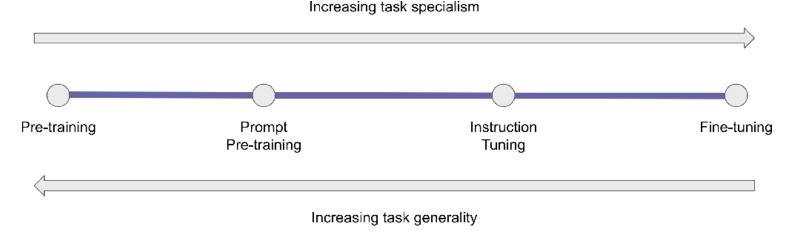


Figure 5: Prompt Pre-training. Pre-training weighs all tokens equally as part of the self-supervised loss. This leads to a weak relative signal for tasks of interest, meaning model scale has to be large to work. Instruction tuning boosts performance post hoc, and can generalize to unseen tasks of interest, but it risks performance in tasks that are distant from instruction set tasks. Prompt pre-training has a weaker task of interest bias than instruction tuning but less risk of degrading overall task generality.

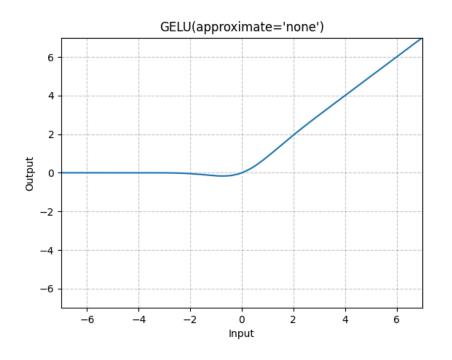
- GeLU Activation GeLU activations for all model sizes.
- Context Window a 2048 length context window.
- **No Biases** following PaLM, no biase in any of the dense kernels or layer norms.
- Learned Positional Embeddings learned positional embeddings for the model.
- **Vocabulary** vocabulary of 50k tokens using BPE. The vocabulary was generated from a randomly selected 2% subset of the training data.

Gaussian Error Linear Units function (GeLu)

$$GELU(x) = x * \Phi(x)$$

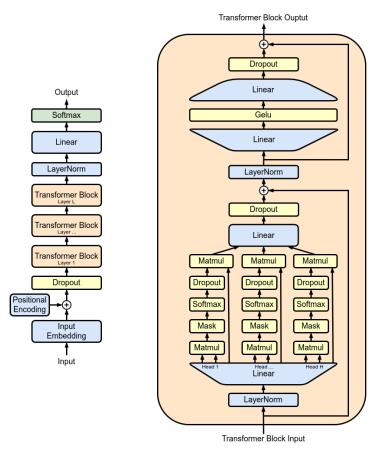
Where $\Phi(x)$ is the Gaussian function.

$$GELU(x) pprox x*rac{1}{2}(1+Tanh(rac{2}{\pi}*(x+0.044715*x^3)))$$



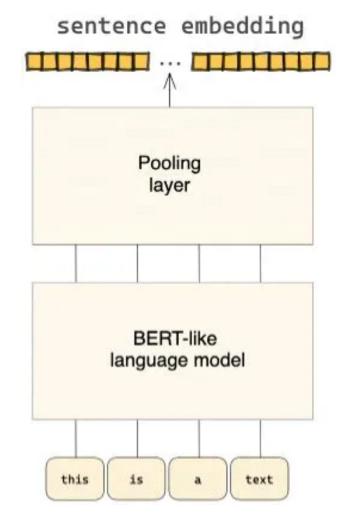
- Allows small negative values when x < 0.
- Avoids the dying ReLU problem.

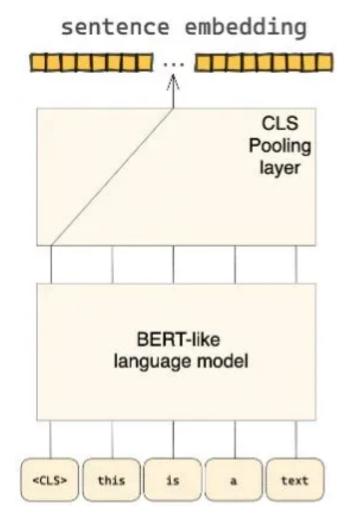
Why no biases?

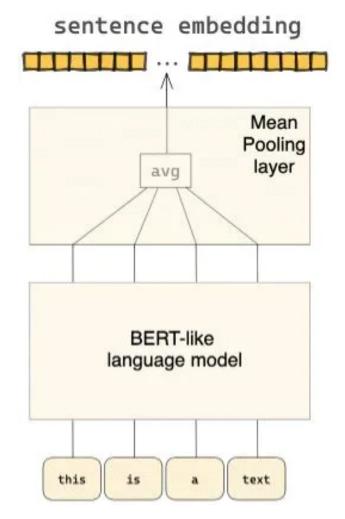


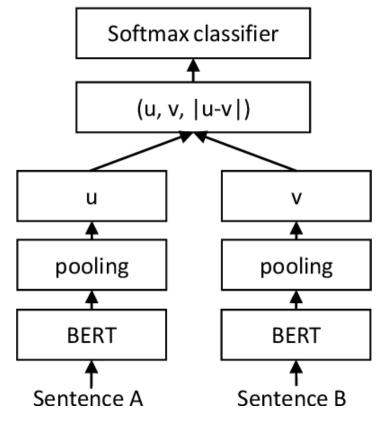
Unsupervised Classification Models

Embedding pooling is the process of combining token embeddings from an encoder model into a single vector representing the entire input sequence. Common methods include averaging (mean pooling), taking the maximum (max pooling), or using a special token like [CLS] or <s>.









[6]

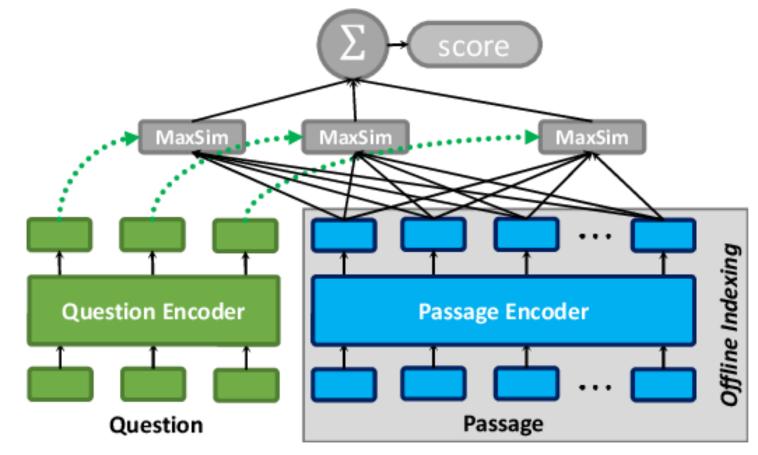


Figure 1. The late interaction architecture given

The data is being compressed mutliple time -> challeging document can be hard to embed.

Can we do better?

Contrastive learning uses similar data point and opposite ones in order for the model build close representations for the first ones and and more separated ones for the latter. [7]

- Unsupervised SimCSE: standard dropout as data augmentation
- Supervised SimCSE: use pairs in NLI datasets

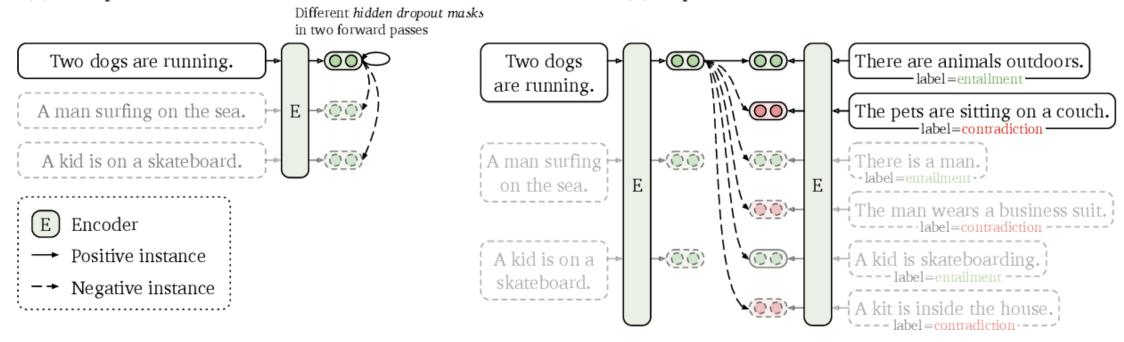
$$\mathcal{L}_{uns} = -lograc{exp(rac{sim(extbf{h}_i, extbf{h}_i^+)}{ au})}{\sum_{j=1}^{N} exp(rac{sim(extbf{h}_i, extbf{h}_i^+)}{ au})}$$

$$\mathcal{L}_{sup} = -lograc{exp(rac{sim(old h_i,old h_i^+)}{ au})}{\sum_{j=1}^N exp(rac{sim(old h_i,old h_j^+)}{ au}) + exp(rac{sim(old h_i,old h_j^-)}{ au})}$$

[8]

(a) Unsupervised SimCSE

(b) Supervised SimCSE

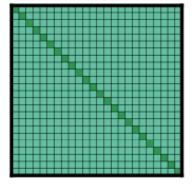


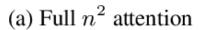
Contrastive learning mitigates anisotropy in language models by encouraging embeddings to be more uniformly distributed in the representation space. It pulls similar embeddings closer and pushes dissimilar ones apart, preventing over-clustering and ensuring better geometric properties for downstream tasks.

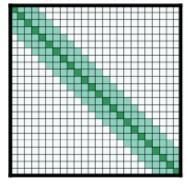
Learning Long-Range Dependencies

Long-range attention models

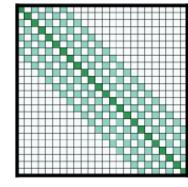
Sliding window attention: Longformer [11]



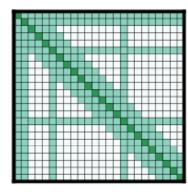




(b) Sliding window attention



(c) Dilated sliding window



(d) Global+sliding window

Long-range attention models

Sliding window attention: Mistral 7B [12]

2 Architectural details

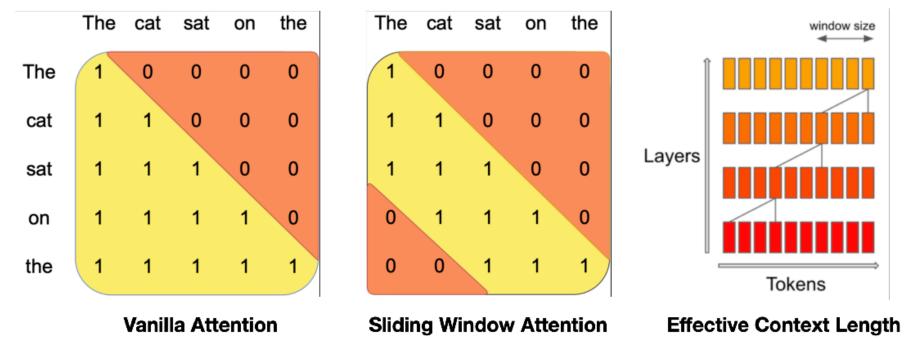
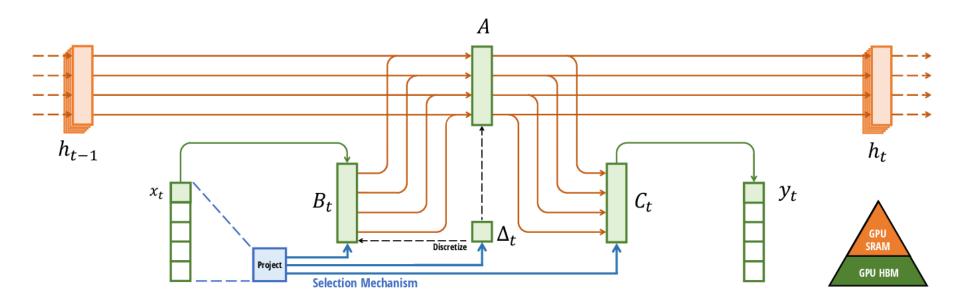


Figure 1. Cliding Window Attention. The number of executions in vanille attention is sundentic in the accurance

State-space models: Mamba



Questions?

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