



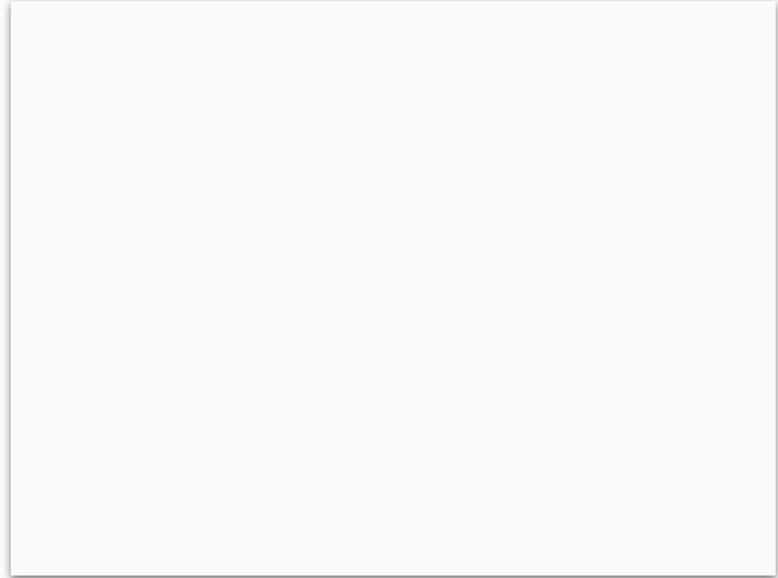
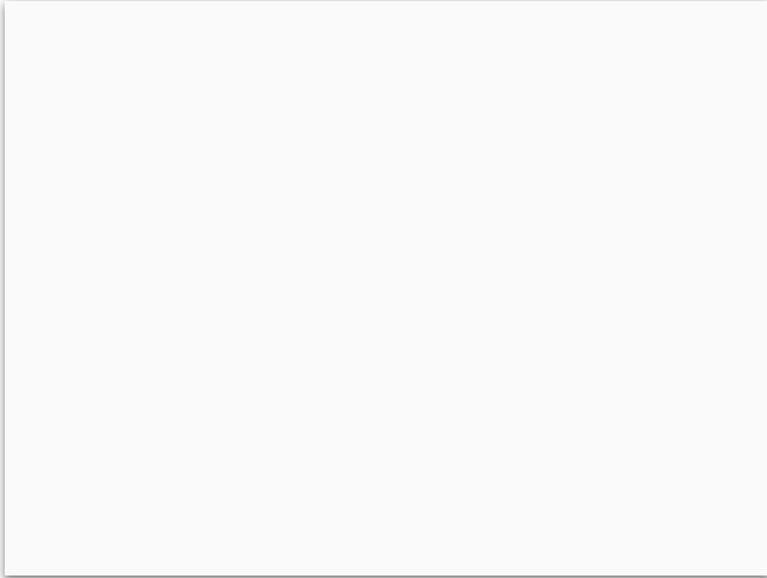
Countering Bias in AI Methods for Text Data

Nicolas Audinet de Pieuchon

6th of February 2026



Bias in AI Methods for Text Data



Bias in AI Methods for Text Data

Many types of bias: statistical bias,
model bias, representational bias,
social biases of all kinds

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Bias in AI methods leads to **real harm!**

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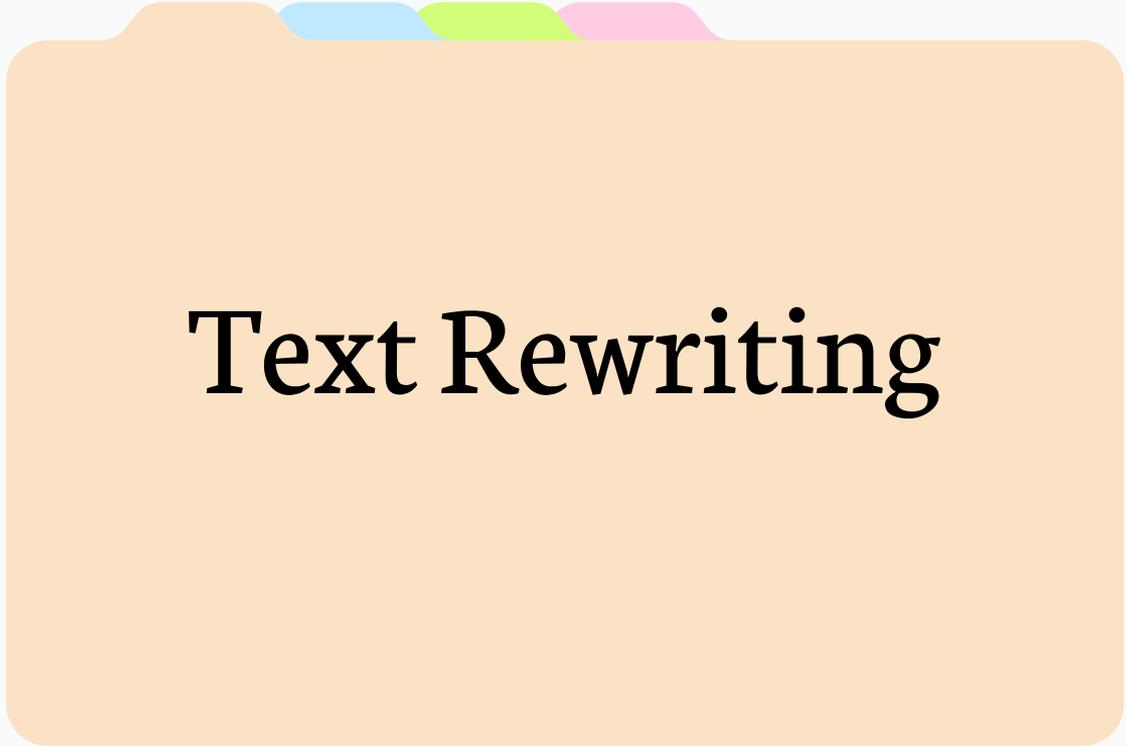
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Why?

- 1) To protect people
- 2) Applications to causal inference
- 3) To see if we can!



Text Rewriting

Product Review:

I bought this album because I loved the title song. It's such a great song, how bad can the rest of the album be, right? Well, the rest of the songs are just filler and aren't worth the money I paid for this it's either shameless bubblegum or over sentimentalized depressing tripe ...

Product Review:

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Sentiment:
negative

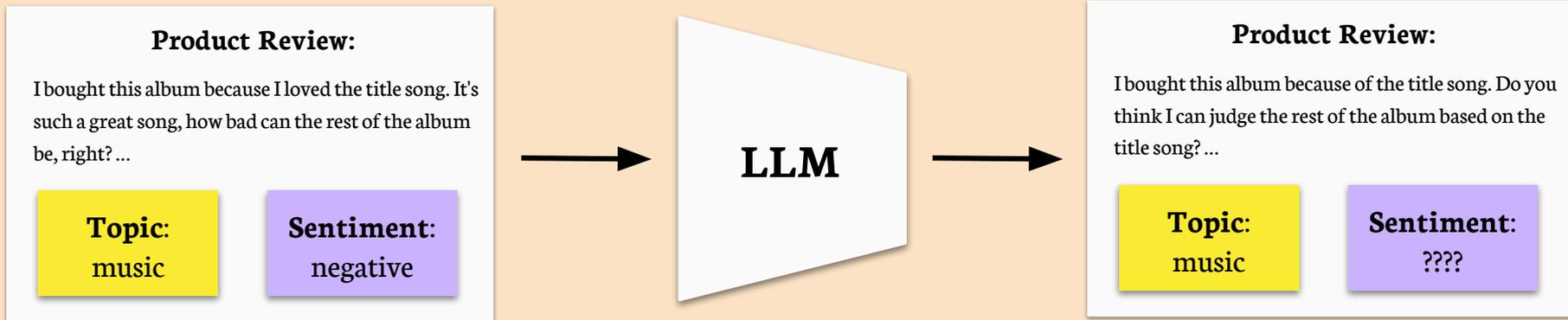
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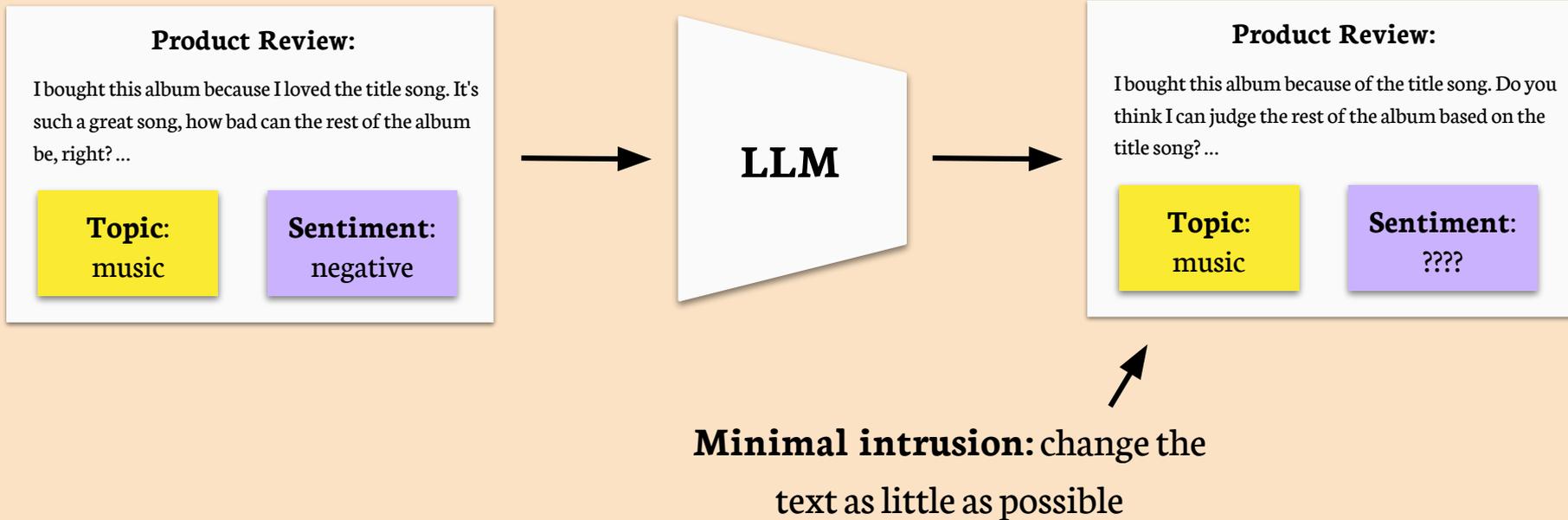
Sentiment:
negative

Topic:
music

Can we use an LLM to rewrite the review to make it neutral?



Can we use an LLM to rewrite the review to make it neutral?



Text Rewriting

Downstream
Debiasing

Representation
Blinding

Future Work

**Product
Reviews**

Text Rewriting

Downstream
Debiasing

Representation
Blinding

Future Work

**Product
Reviews**



```
graph TD; A[Product Reviews] --> B[Classifier]; A --> C[Classifier]; B --> D[Topic]; C --> E[Sentiment];
```

Classifier

Classifier

Topic

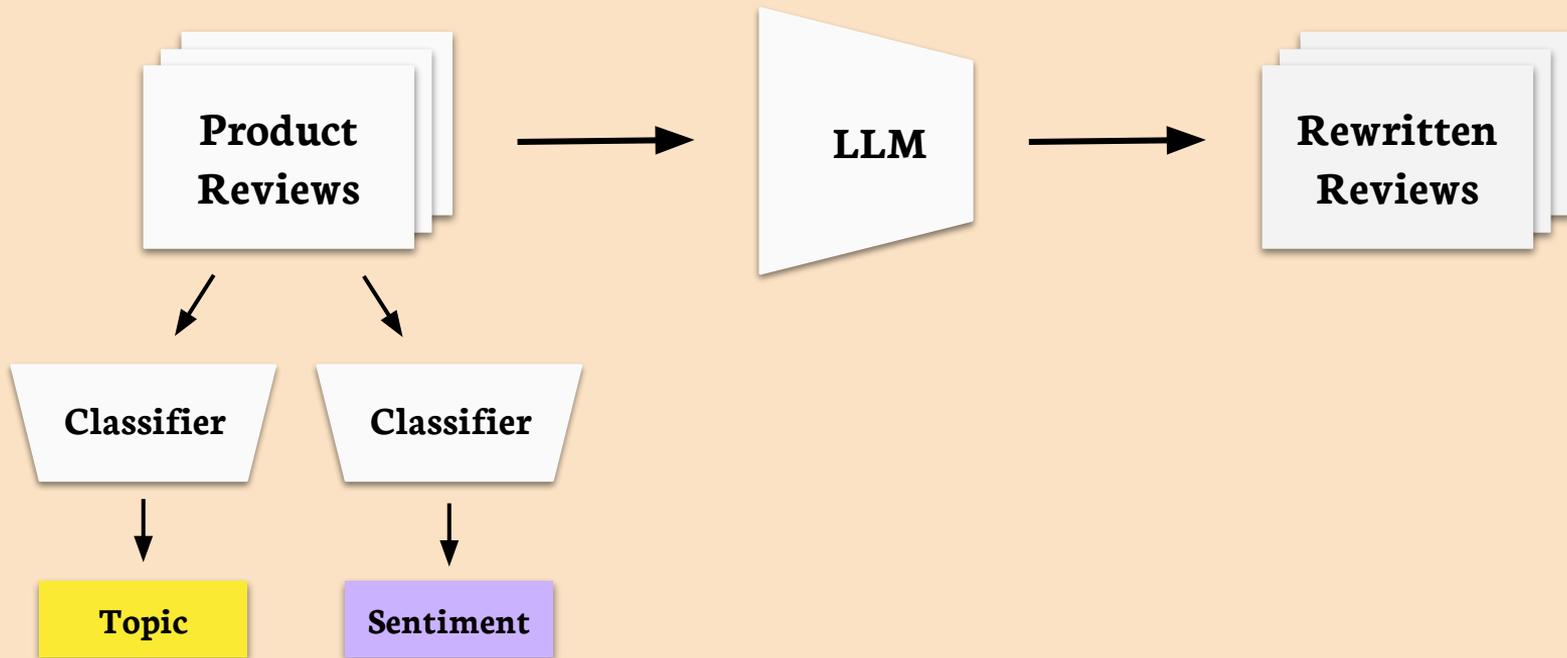
Sentiment

Text Rewriting

Downstream
Debiasing

Representation
Blinding

Future Work

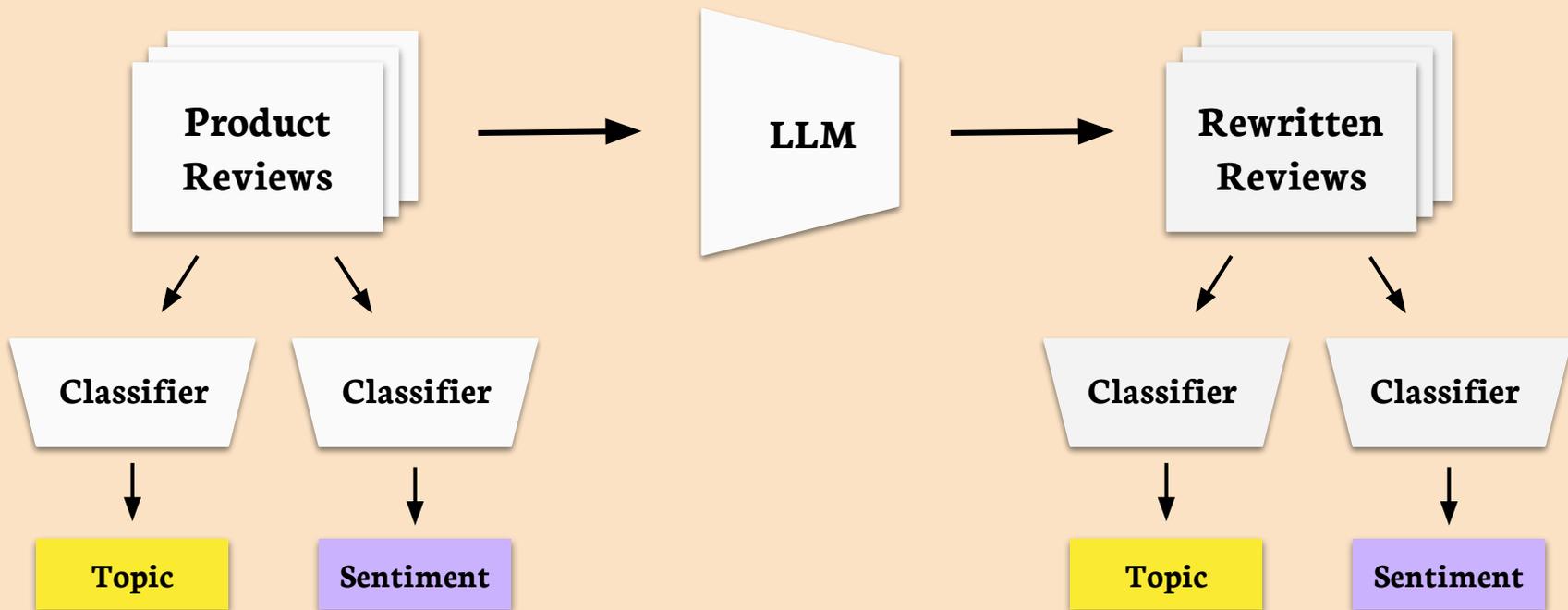


Text Rewriting

Downstream
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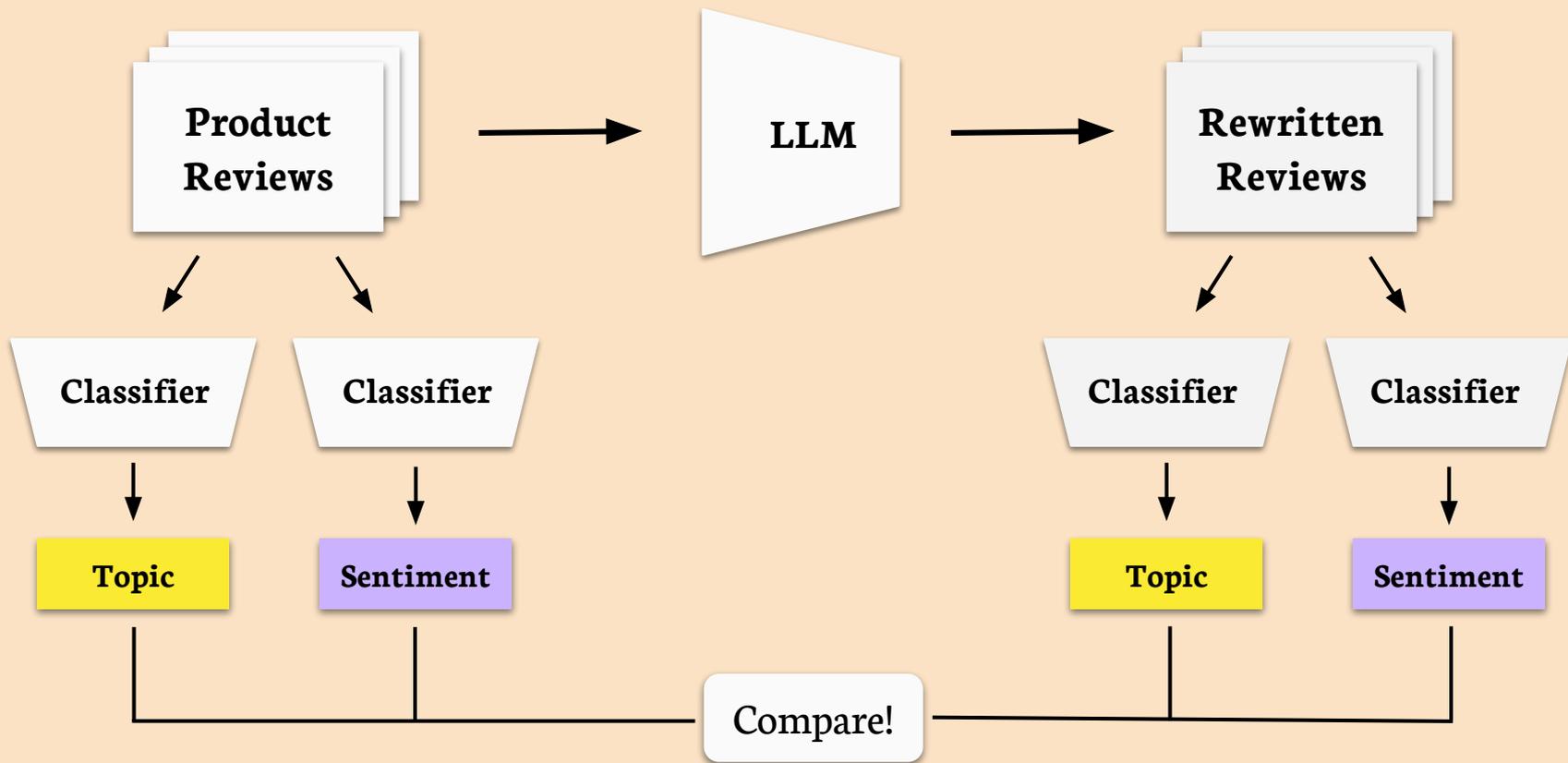


Text Rewriting

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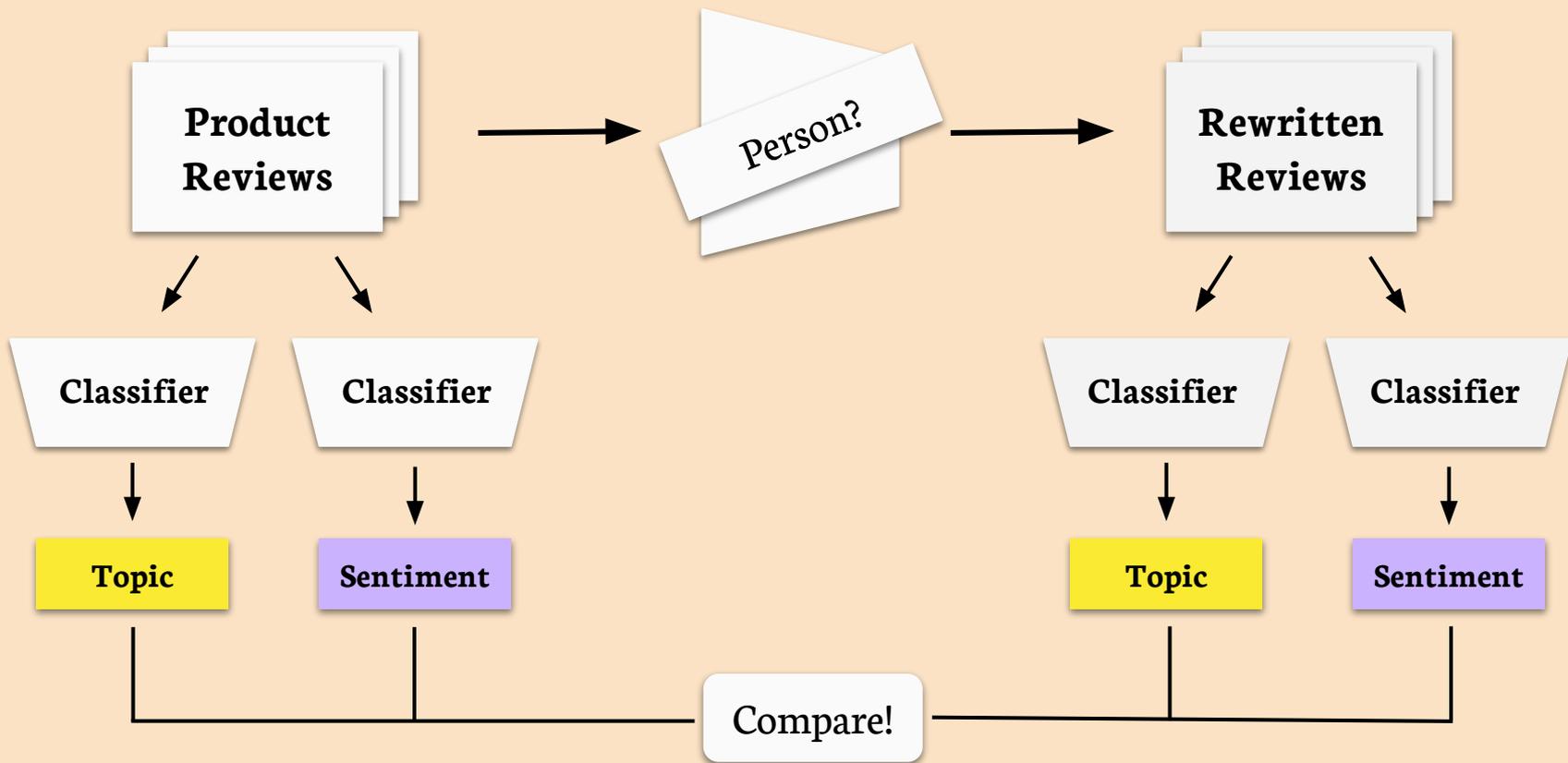


Text Rewriting

Downstream
Debiasing

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Future Work



Text Rewriting

Downstream
Debiasing

Representation
Blinding

Future Work

Setting	Prompt	Sentiment Accuracy ↓	Topic Accuracy ↑
No distillation		0.885 ± 0.035	0.946 ± 0.026

Text Rewriting

Downstream
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Future Work

Setting	Prompt	Sentiment Accuracy ↓	Topic Accuracy ↑
No distillation		0.885 ± 0.035	0.946 ± 0.026
Mean projection		0.524 ± 0.054	0.946 ± 0.026

Text Rewriting

Downstream
Debiasing

Representation
Blinding

Future Work

Setting	Prompt	Sentiment Accuracy ↓	Topic Accuracy ↑
No distillation		0.885 ± 0.035	0.946 ± 0.026
Mean projection		0.524 ± 0.054	0.946 ± 0.026
Human*	Prompt chaining	0.800 ± 0.145	0.842 ± 0.165

Setting	Prompt	Sentiment Accuracy ↓	Topic Accuracy ↑
No distillation		0.885 ± 0.035	0.946 ± 0.026
Mean projection		0.524 ± 0.054	0.946 ± 0.026
Human*	Prompt chaining	0.800 ± 0.145	0.842 ± 0.165
Mistral 7B	Paraphrase	0.891 ± 0.037	0.951 ± 0.024
	Few-shot	0.877 ± 0.023	0.951 ± 0.015
	Prompt chaining	0.841 ± 0.039	0.953 ± 0.023
GPT 4	Paraphrase	0.899 ± 0.034	0.951 ± 0.024
	Few-shot	0.824 ± 0.045	0.955 ± 0.024
	Prompt chaining	0.757 ± 0.044	0.945 ± 0.023

Setting	Prompt	Sentiment Accuracy ↓	Topic Accuracy ↑
No distillation		0.885 ± 0.035	0.946 ± 0.026
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Can LLMs Disentangle Text?

1.

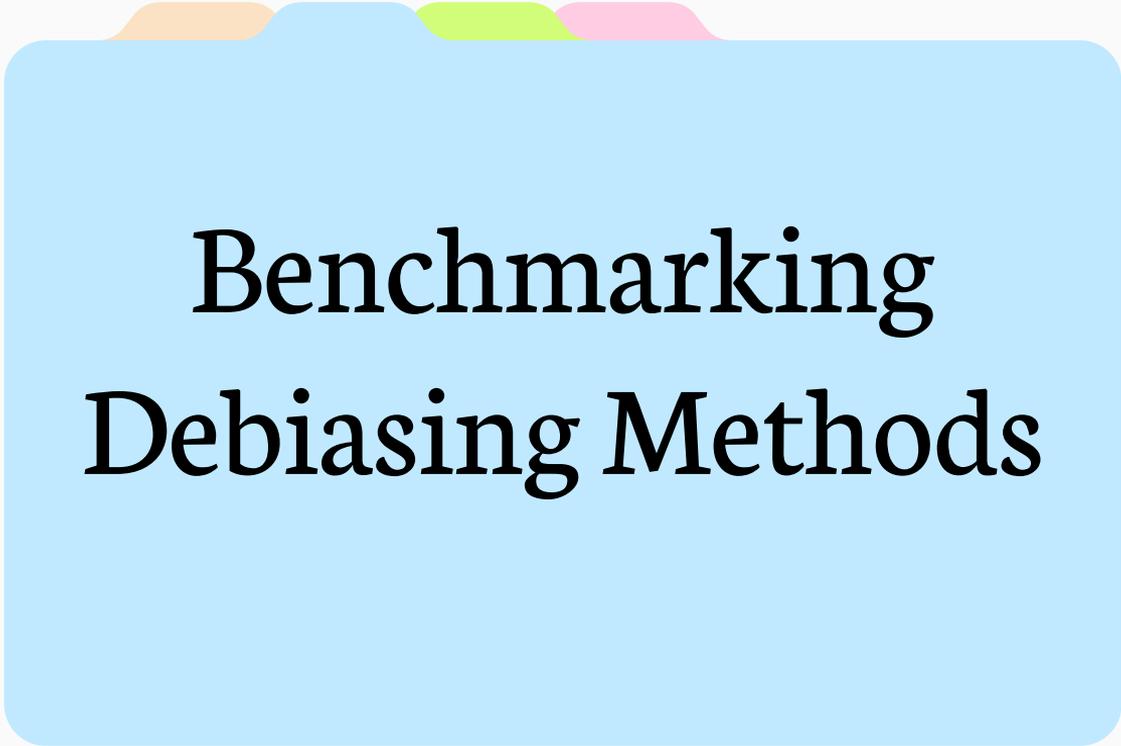
Some success, but
nowhere near baselines

2.

Difficult task for both
humans and LLMs

3.

Performance might be
task-dependent



Benchmarking Debiasing Methods

Advances in post-hoc debiasing methods ...

Using Imperfect Surrogates for Downstream Inference: Design-based Supervised Learning for Social Science Applications of Large Language Models

Naoki Egami^{*1}, Musashi Hinck², Brandon M. Stewart^{*2}, Hanying Wei¹
¹Columbia University, ²Princeton University

Abstract

In computational social science (CSS), researchers analyze documents to explain social and political phenomena. In most scenarios, CSS researchers first obtain labels for documents and then explain labels using interpretable regression analyses in the second step. One increasingly common way to annotate documents cheaply at scale is through large language models (LLMs). However, like other scalable ways of producing annotations, such surrogate labels are often imperfect and biased. We present a new algorithm for using imperfect annotation surrogates for downstream statistical analyses while guaranteeing statistical properties—like asymptotic unbiasedness and proper uncertainty quantification—which are *fundamental* to CSS research. We show that direct use of surrogate labels in downstream statistical analyses leads to substantial bias and invalid confidence intervals, even with high surrogate accuracy of 80–90%. To address this, we build on debiased machine learning to propose the *design-based supervised learning* (DSL) estimator. DSL employs a doubly-robust procedure to combine surrogate labels with a smaller number of high-quality, gold-standard labels. Our approach guarantees valid inference for downstream statistical analyses, even when surrogates are arbitrarily biased and without requiring stringent assumptions, by controlling the probability of sampling documents for gold-standard labeling. Both our theoretical analysis and experimental results show that DSL provides valid statistical inference while achieving root mean squared errors comparable to existing alternatives that focus only on prediction without inferential guarantees.

RESEARCH

MACHINE LEARNING

Prediction-powered inference

Anastasios N. Angelopoulos¹, Stephen Bates², Clara Fanrjang³, Michael I. Jordan⁴, Tijana Zrnica⁵

Prediction-powered inference is a framework for performing valid statistical inference when an experimental dataset is supplemented with predictions from a machine-learning system. The framework yields simple algorithms for computing provably valid confidence intervals for quantities such as means, quantiles, and linear and logistic regression coefficients without making any assumptions about the machine-learning algorithm that supplies the predictions. Furthermore, more accurate predictions translate to smaller confidence intervals. Prediction-powered inference could enable researchers to draw valid and more data-efficient conclusions using machine learning. The benefits of prediction-powered inference were demonstrated with datasets from proteomics, astronomy, genomics, remote sensing, census analysis, and ecology.

Imagine a scientist has a machine-learning system that can supply accurate predictions about a phenomenon far more cheaply than any gold-standard experimental technique. The scientist may wish to use these predictions as evidence in drawing scientific conclusions. For example, accurate predictions of three-dimensional structures have been made for a vast catalog of known protein sequences (1, 2) and are now being used in proteomics studies (3, 4). Such machine-learning systems are increasingly common in modern scientific inquiry, in domains ranging from cancer prognosis to microclimate modeling. Predictions are not perfect, however, and this may lead to incorrect conclusions. Moreover, as predictions best other predictions, the cumulative effect can amplify the imperfections. How can modern science leverage machine-learning predictions in a statistically principled way?

One way to use predictions is to follow the imputation approach: Proceed as if they are gold-standard measurements. Although this lets the scientist draw conclusions cheaply and quickly owing to the high-throughput nature of the machine-learning system, the conclusions may be invalid because the predictions may have biases.

Another possibility is to apply the classical approach: Ignore the machine-learning predictions and only use the available gold-standard measurements, which are typically far less abundant than the machine-learning predictions. This approach is also problematic because the machine-learning predictions are often more abundant than the gold-standard measurements. Prediction-powered inference provides a protocol for combining predictions, which are abundant but not always trustworthy, with gold-standard data, which are trusted but scarce, to compute confidence intervals and P values. The resulting confidence intervals and P values are statistically valid, as in the classical approach, but also leverage the information contained in the predictions, as in the imputation approach, to make the confidence intervals smaller and the P values more powerful.

Prediction-powered inference applies to any machine-learning system; as such, it absolves the need for case-by-case analyses dependent on the machine-learning algorithm on hand. The proposed protocol thereby could enable researchers to report on and assess the evidence for their conclusions in a fully standardized way.

Protocol for prediction-powered inference

The protocol for prediction-powered inference proceeds as follows. The scientist wishes to construct a confidence interval for a quantity θ^* , such as the mean outcome or a regression coefficient quantifying the statistical association between the outcome and a feature. Toward this goal, they have access to a small gold-standard dataset of features paired with outcomes, $(X, Y) = ((X_1, Y_1), \dots, (X_n, Y_n))$, as

Prediction-powered inference uses the gold-standard dataset to quantify and correct for the errors made by the machine-learning algorithm on the unlabeled dataset, thereby enabling researchers to reliably incorporate predictions when constructing confidence intervals. The three-step protocol is outlined below and visualized in Fig. 1.

1) Estimation. The first step is to select an estimand θ^* . The estimand is the quantity the scientist is interested in knowing—for example, the mean outcome $E[Y]$, median outcome $\text{median}(Y)$, a linear regression coefficient obtained by regressing Y onto X , etc.

2) Measure of fit and rectifier. The key step is to identify the right measure of fit m_θ and rectifier A_θ for the selected estimand. For every candidate value of the estimand θ , the measure of fit m_θ is computed on the unlabeled dataset imputed with predictions, (X', Y') and quantifies how likely θ^* is to be equal to θ on the basis of the imputed data. The closer m_θ is to zero, the more plausible it is for θ^* to be equal to θ . The rectifier A_θ is a notion of prediction error that is relevant for the estimand of interest. It is defined as the difference of the measure of fit m_θ computed on the labeled data, (X, Y) , and the labeled data when the true outcomes are replaced with predicted ones, (X, Y') . If the predictions are perfect, the rectifier is equal to zero.

Table 1 states the appropriate measure of fit and rectifier for common estimands of interest: the mean outcome, median outcome, q -quantile of the outcome, and linear and logistic regression coefficients when regressing Y onto X . A general recipe for deriving the right measure of fit and corresponding rectifier for a broad class of other estimands is provided in the SM.

3) Prediction-powered confidence interval. Finally, the measure of fit and rectifier are carefully combined to form a prediction-powered confidence interval for θ^* . This process is called rectifying the confidence interval. The prediction-powered confidence interval is constructed as $C^{PP} = \{\theta \text{ such that } |m_\theta + A_\theta| \leq \omega(\alpha)\}$ and is guaranteed to contain the esti-

X1	X2	...	Y
1	T	...	By now, we all know that House Intelligence Committee ...
5	F	...	Donald Trump won't tell you this on his Twitter feed. The ...
2	F	...	His accusation that President Obama wiretapped him has ...
...
4	T	...	The rainbow flag has become a symbol for LGBT rights ...

X1	X2	...	Y
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Annotate by hand

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Annotate by hand

Generate annotations
with an LLM

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Debiasing methods let
you use both!

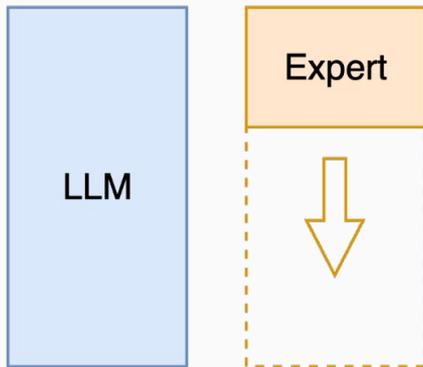
Annotate by hand

Generate annotations
with an LLM

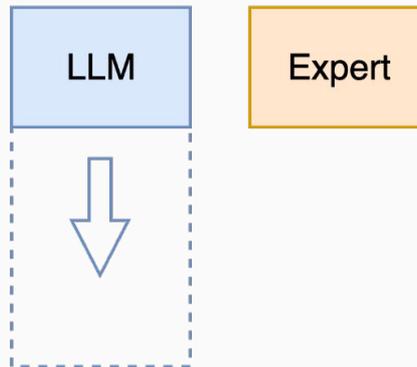
RQ1: When is it preferable to use debiasing methods over just the expert annotations?

RQ2: What are the performance differences between debiasing methods?

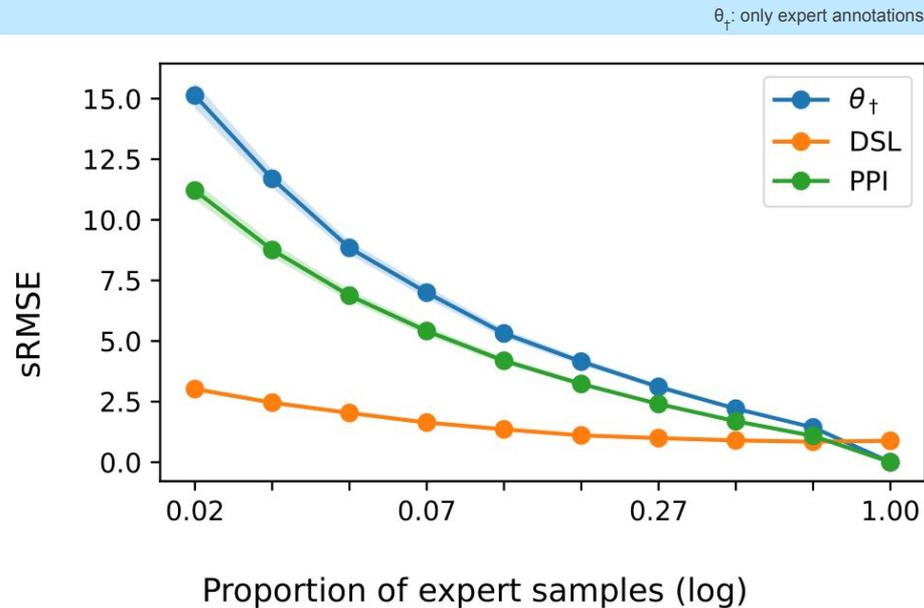
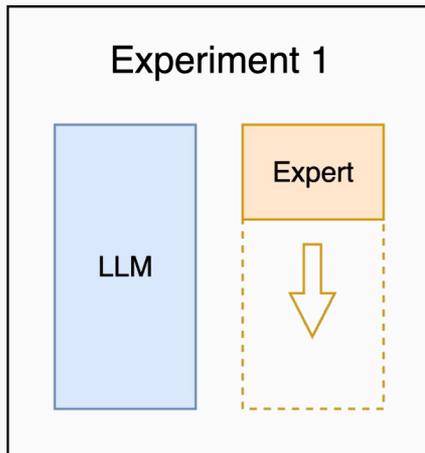
Experiment 1



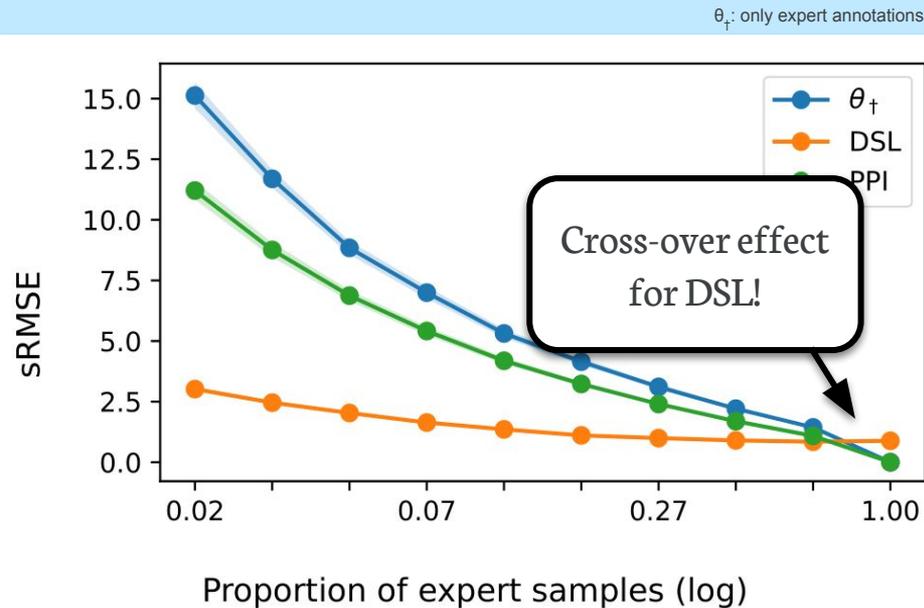
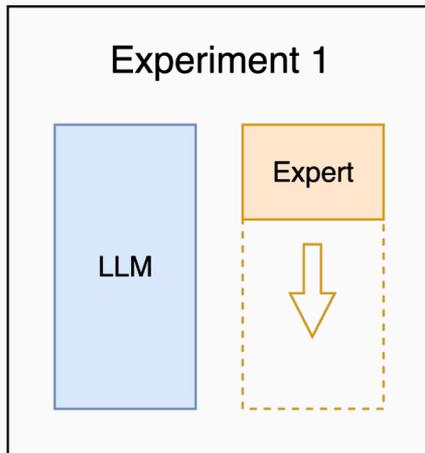
Experiment 2

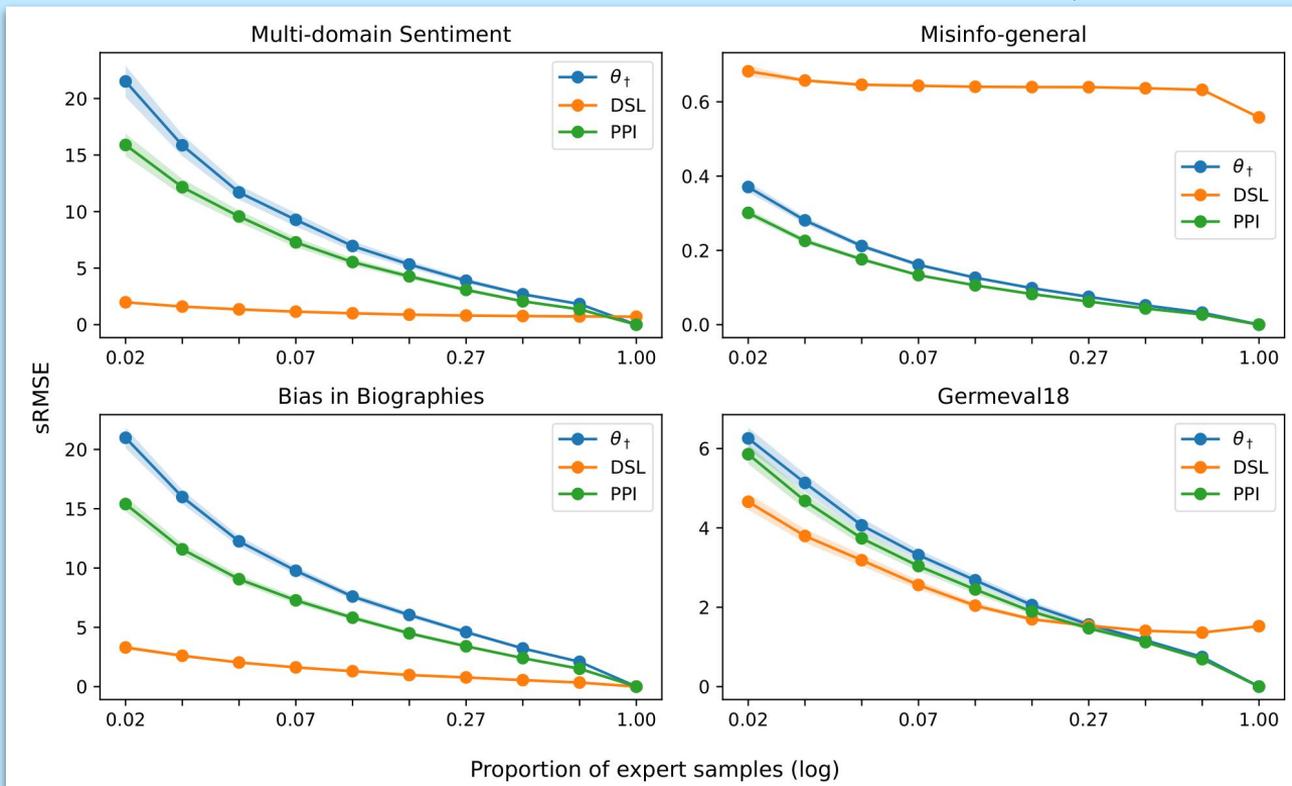


Experiment 1

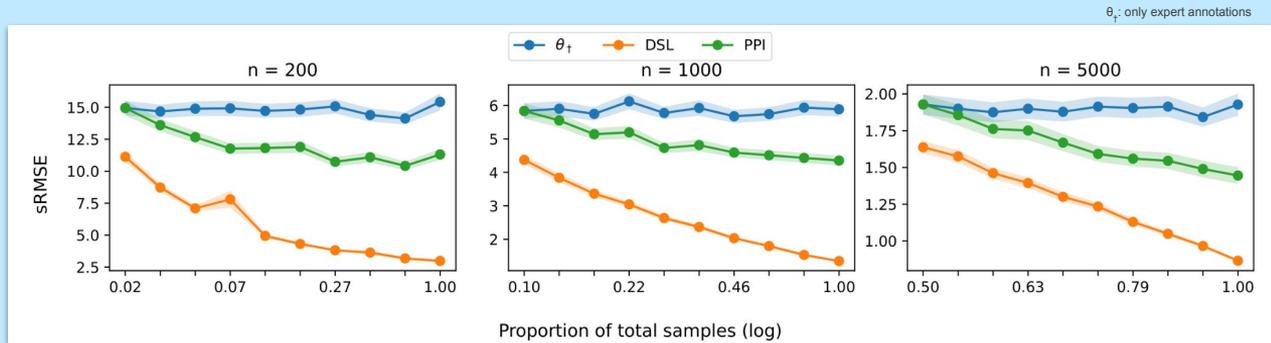
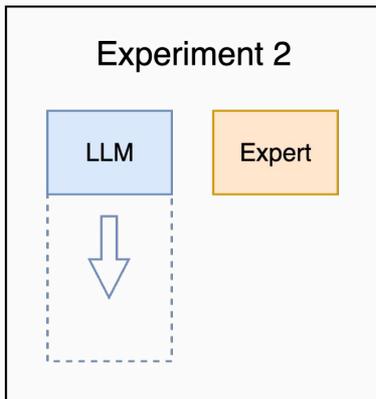


Experiment 1



θ_{\dagger} : only expert annotations

Experiment 2

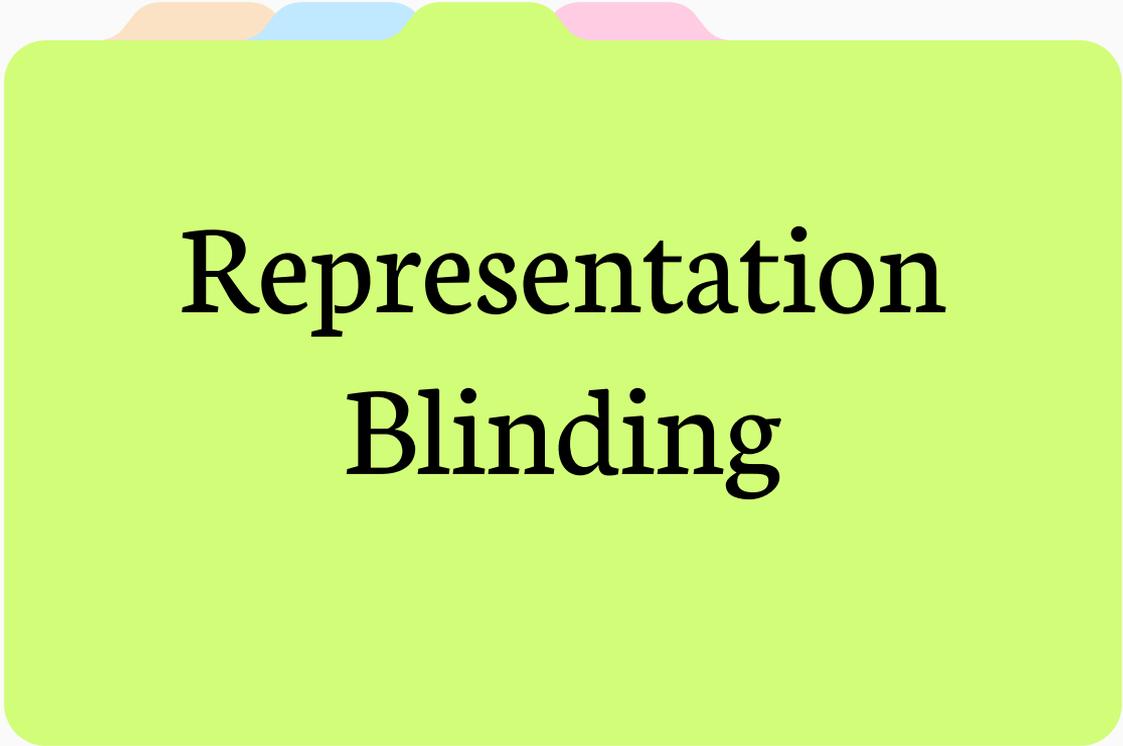


RQ1: When is it preferable to use debiasing methods over just the expert annotations?

Both DSL or PPI is more efficient than using only expert annotations.

RQ2: What are the performance differences between debiasing methods?

DSL tends to outperform PPI, but performance is more dataset-dependant.



Representation Blinding

Advances in mechanistic interpretability ...

Transcoders Find Interpretable

Jacob Dunefsky*
Yale University
New Haven, CT 06511
jacob.dunefsky@yale.edu

SPARSE FEATURE CIRCUITS: DISCOVERING AND EDITING INTERPRETABLE CIRCUITS IN LANGUAGE MODELS

Samuel Marks*
Northeastern University

Yonatan Belinkov
Technion – IIT

Can Feng
Independent

David Bau
Northeastern University

Need

Ab

A key goal in mechanistic interpretability is to understand the internal workings of language models. However, the complexity of modern models makes this task difficult. In particular, interpreting the behavior of sparse autoencoders (SAEs)—a type of neuron that is active in a sparse manner—is often intractable. We address this by introducing a novel method for circuit analysis through MLP sublayers. Our method is input-dependent and input-invariant, and we demonstrate that it performs at least on par with SAEs in terms of interpretability. Finally, we apply this method to GPT2-small. Our results suggest that tracing model computations involving MLP sublayers is possible. Available at <https://github.com/jacob>

1 Introduction

1 INTRODUCTION

The key challenge of interpretability research is to discover the internal workings of neural networks (NNs). Much recent work explores model components, for example by implicating certain neurons (e.g., Gou et al., 2022) or MLP modules in factual recall

A Survey on Sparse Interpreting the Internal Mechanisms

Dong Shu^{1,†}, Xuansheng Wu^{2,†}, Haoyang
Ziyu Yao⁴, Ninghao Liu¹,
¹Northwestern University ²UC
³New Jersey Institute of Technology
dongshu2024@northwestern.edu, {xwu,
hzs54, mengnan.du}@njit.edu, {d

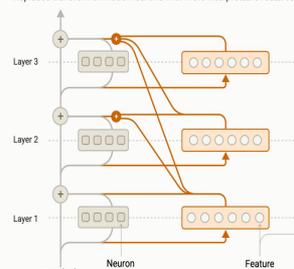
Abstract

Large Language Models (LLMs) have transformed natural language processing, yet their internal mechanisms remain largely opaque. Recently, mechanistic interpretability has attracted significant attention from the research community as a means to understand the inner workings of LLMs. Among various mechanistic interpretability approaches, Sparse Autoencoders (SAEs) have emerged as a promising method due to their ability to disentangle the complex, superimposed features within LLMs into more interpretable components. This paper presents a comprehensive survey of SAEs for interpreting and understanding the internal workings of LLMs. Our major contributions include: (1) exploring the technical framework of SAEs, covering basic architecture, design improvements, and effective training strategies; (2) examining different approaches to explaining SAE features, categorized into input-based and output-based explanation methods; (3) dis-

Circuit Tracing: Revealing Computational Graphs in Language Models

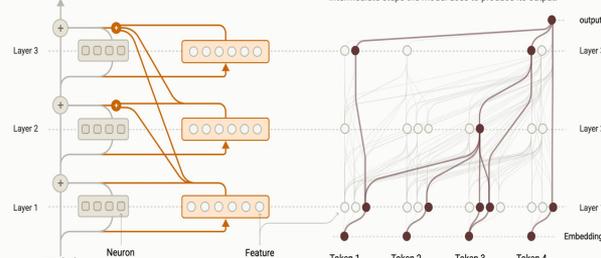
Replacement Model

Replaces transformer model neurons with more interpretable features.



Attribution Graph

Depicts influence of features on one another, allowing us to trace intermediate steps the model uses to produce its output.



We introduce a method to uncover mechanisms underlying behaviors of language models. We produce graph descriptions of the model's computation on prompts of interest by tracing individual computational steps in a "replacement model". This replacement model substitutes a more interpretable component (here, a "cross-layer transcoder") for parts of the underlying model (here, the multi-layer perceptrons) that it is trained to approximate. We develop a suite of visualization and validation tools we use to investigate these "attribution graphs" supporting simple behaviors of an 18-layer language model, and lay the groundwork for a [companion paper](#) applying these methods to a frontier model, Claude 3.5 Haiku.

Text Rewriting

Downstream
Debiasing

Representation
Blinding

Future Work

Ambiguous Dataset

Male Professor

Female Nurse

Text Rewriting

Downstream
Debiasing

Representation
Blinding

Future Work

Ambiguous Dataset

Male Professor

Female Nurse

train

LLM

Classifier
Head

Text Rewriting

Downstream
Debiasing

Representation
Blinding

Future Work

Ambiguous Dataset

Male Professor

Female Nurse

train

LLM

Classifier
Head

Balanced Dataset

Male
Professor

Female
Professor

Male
Nurse

Female
Nurse

test

Can we remove male/female information
from the text representation?

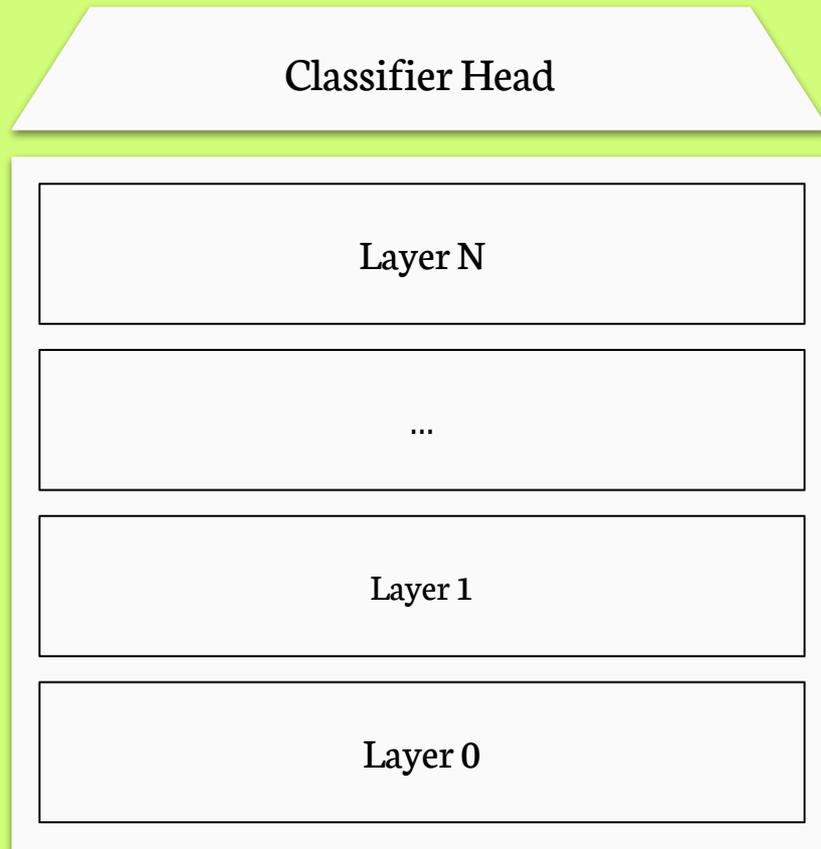
Can we remove male/female information
from the text representation?

SHIFT method:

Can we remove male/female information
from the text representation?

SHIFT method:

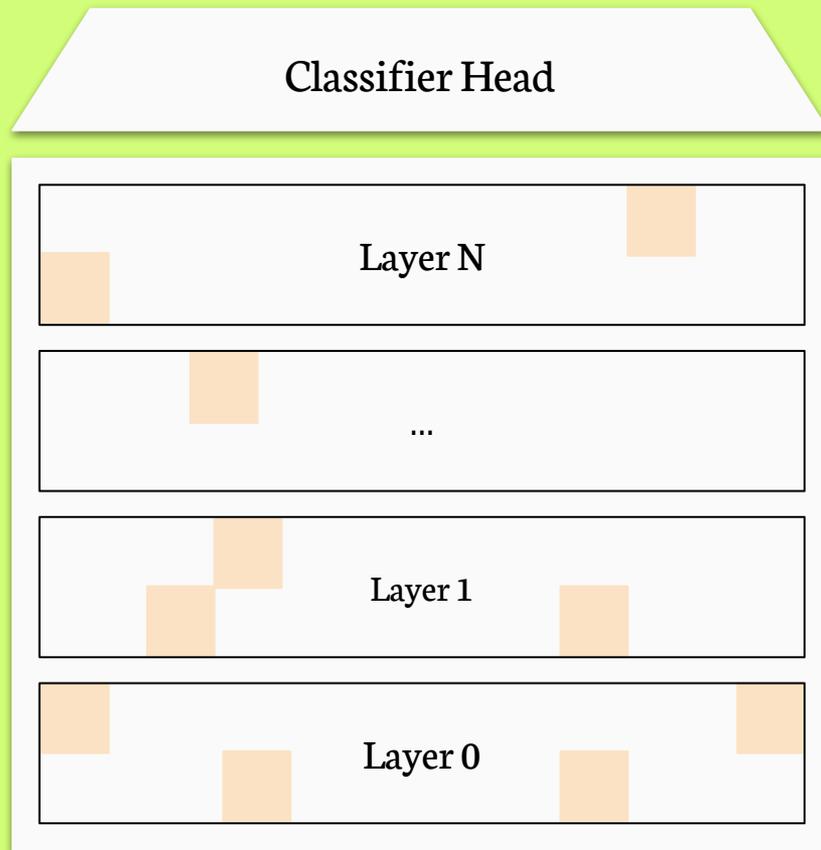
- 1) Train the classifier



Can we remove male/female information from the text representation?

SHIFT method:

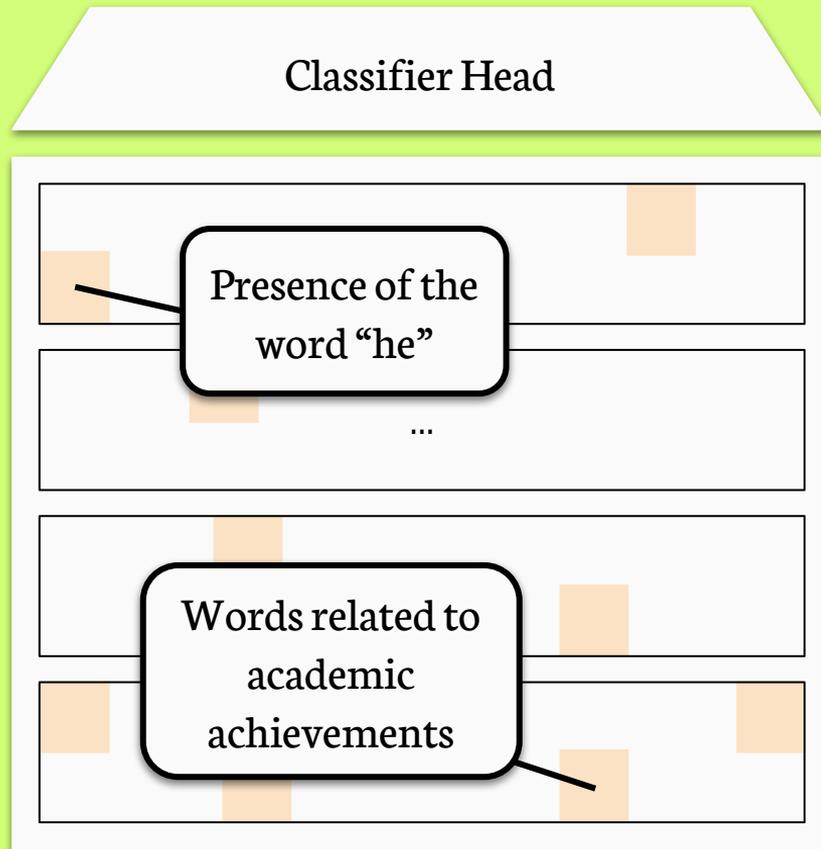
- 1) Train the classifier
- 2) Find relevant features



Can we remove male/female information from the text representation?

SHIFT method:

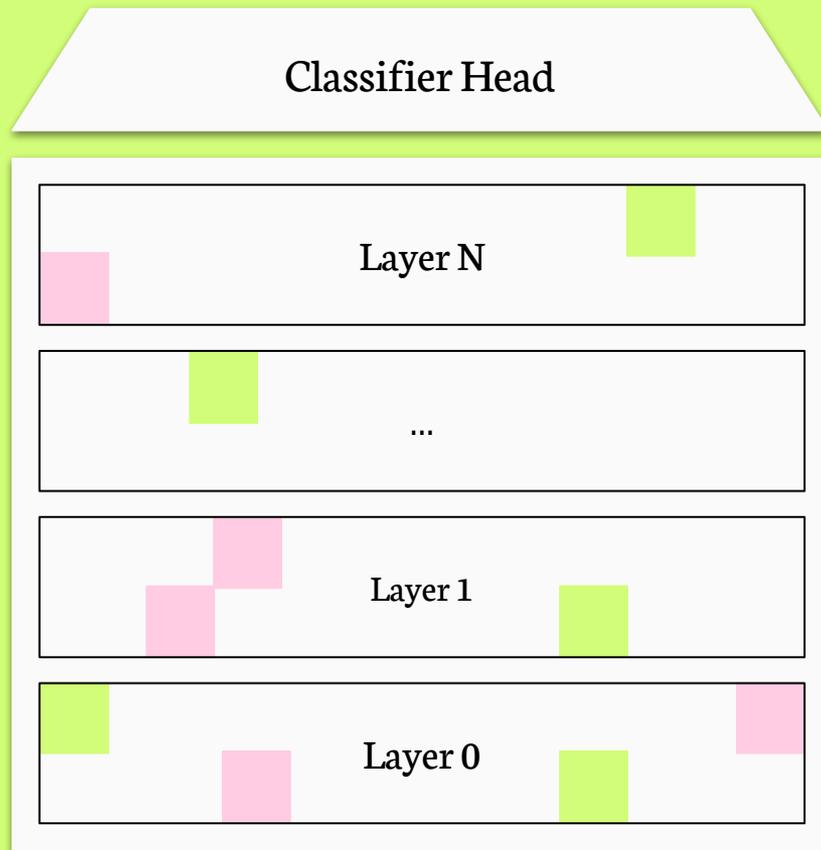
- 1) Train the classifier
- 2) Find relevant features
- 3) Interpret features



Can we remove male/female information from the text representation?

SHIFT method:

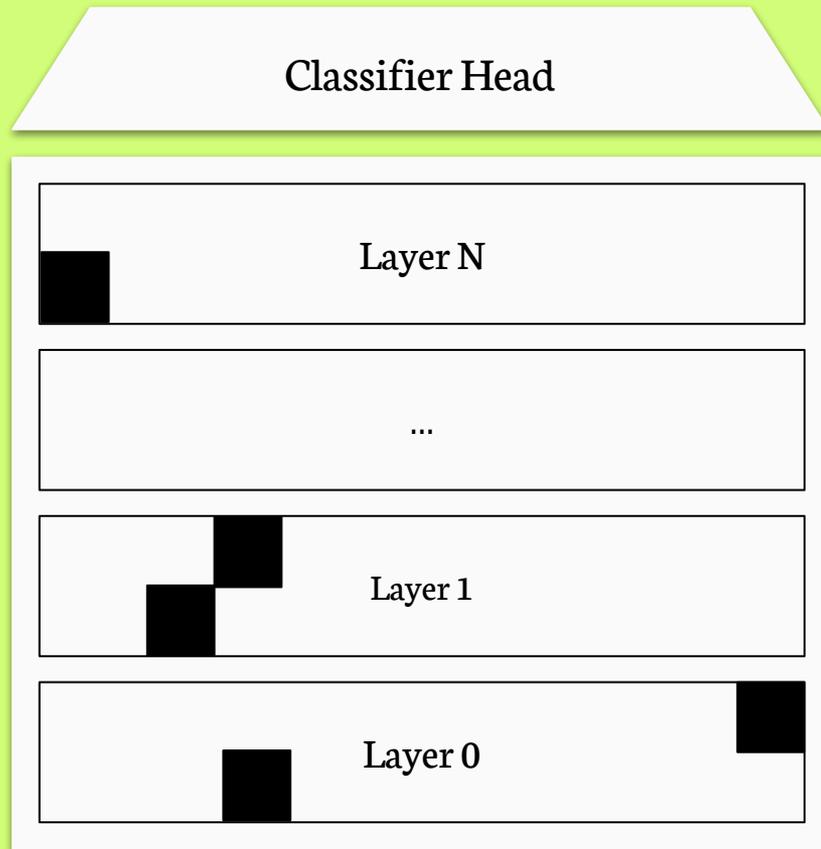
- 1) Train the classifier
- 2) Find relevant features
- 3) Interpret features
- 4) Select features



Can we remove male/female information from the text representation?

SHIFT method:

- 1) Train the classifier
- 2) Find relevant features
- 3) Interpret features
- 4) Select features
- 5) Remove and re-train

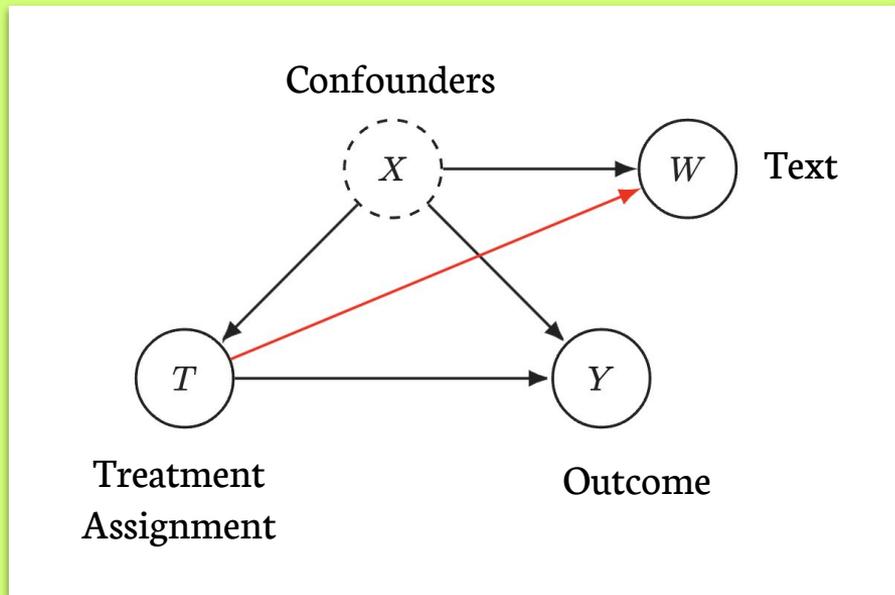


Pythia-70M				
Method	↑ Profession	↓ Gender	↑ Worst group	
Original	61.9	87.4	24.4	
CBP	83.3	60.1	67.7	
Random	61.8	87.5	24.4	
SHIFT	88.5	54.0	76.0	
SHIFT + retrain	93.1	52.0	89.0	
Neuron skyline	75.5	73.2	41.5	
Feature skyline	88.5	54.3	62.9	
Oracle	93.0	49.4	91.9	

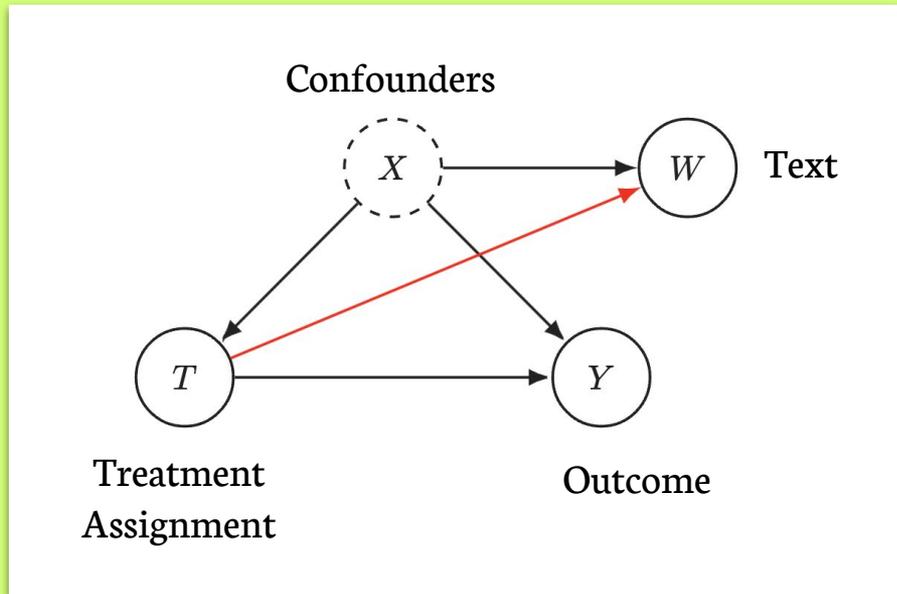
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Pythia-70M			
Method	↑Profession	↓Gender	↑Worst group
Original	61.9	87.4	24.4
CBP	83.3	60.1	67.7
Random	61.8	87.5	24.4
SHIFT	88.5	54.0	76.0
SHIFT + retrain	93.1	52.0	89.0
Neuron skyline	75.5	73.2	41.5
Feature skyline	88.5	54.3	62.9
Oracle	93.0	49.4	91.9

Can we apply this to treatment leakage?



Can we apply this to treatment leakage?



... we'll see!

Right now the results look encouraging but are not there yet.

I'd like to work together!

If you found any of it useful or
interesting, come have a chat :)



Thank you for listening!

References

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