

Mitigating Racial Bias in Clinical Prediction Models: A D3M-Inspired Data Selection Approach

DBDS Student Talks Presentation

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Agenda

1. Research Question Formulation
2. Experimental Outline
3. Background
4. Results
5. Analysis
6. Future Directions
7. Acknowledgments

Research Question Formulation

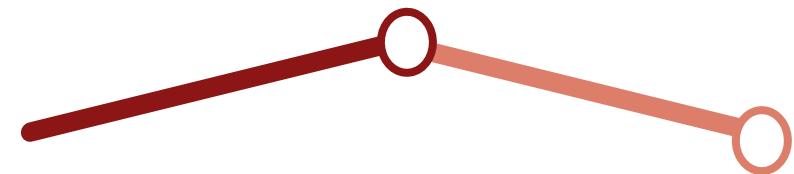
What are computational strategies to mitigate race-based outputs
in LLMs?

- Machine learning models often reflect biases present in training data.
- Models inadvertently learn and amplify these biases, leading to unfair or inaccurate predictions.
- The sheer size of datasets makes it impossible to manually check and correct every data point.
- This necessitates **automated methods for debiasing datasets.**

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Experimental Outline

- **Objective:** Improve worst subgroup performance for racial, protected, and underresourced groups using debiased fine-tuning.

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- **Method:** Apply data selection method (D3M-inspired) to balance training datasets.
- **Expected Outcome:** Boost accuracy in underperforming subgroups and achieve fairer, more robust models.

Background

How do we quantify data's impact in training?

- **Data attribution:** the task of predicting model outputs/behavior at test-time as a function of the input training data.
- In other words: *What would happen if I trained the model on a given subset of my training set?*

Background

- **TRAK:** data attribution method giving us coefficients (scores) to help identify examples that exacerbate discrepancies in group performance
- **D3M:** allows us to actually remove the examples and retrain on a dataset without the harmful examples

$$A_i = \frac{\sum_{g \in \mathcal{G}} \exp(\beta \ell_g) \cdot \tau(g)_i}{\sum_{g' \in \mathcal{G}} \exp(\beta \ell_{g'})}$$

we set $\beta = 1$ (hyperparameter controlling the smoothness of the maximum)

loss of a base classifier $\theta(S)$ on group g (evaluated on the validation set)

group alignment score: the impact of training sample i on the overall worst-group performance

the i -th coefficient for group g

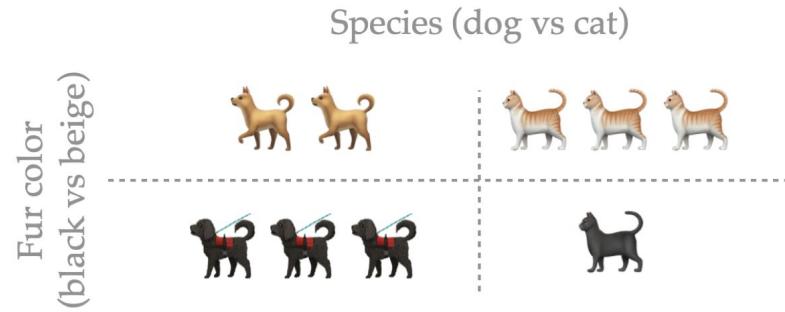
TRAK: Tracing with the Randomly-projected After Kernel

Algorithm 1 TRAK for multi-class classifiers (as implemented)

```
1: Input: Learning algorithm  $\mathcal{A}$ , dataset  $S$  of size  $n$ , sampling fraction  $\alpha \in (0, 1]$ , correct-class likelihood function  $p(z; \theta)$ , projection dimension  $k \in \mathbb{N}$ 
2: Output: Matrix of attribution scores  $\mathbf{T} \in \mathbb{R}^{n \times n}$ 
3:  $f(z; \theta) := \log\left(\frac{p(z; \theta)}{1-p(z; \theta)}\right)$  ▷ Margin function  $f_\theta$ 
4: for  $m \in \{1, \dots, M\}$  do
5:   Sample random  $S' \subset S$  of size  $\alpha \cdot n$ 
6:    $\theta_m^* \leftarrow \mathcal{A}(S')$  ▷ Train a model on  $S'$ 
7:    $\mathbf{P} \sim \mathcal{N}(0, 1)^{p \times k}$  ▷ Sample projection matrix
8:    $\mathbf{Q}^{(m)} \leftarrow \mathbf{0}_{n \times n}$ 
9:   for  $i \in \{1, \dots, n\}$  do
10:     $\phi_i \leftarrow \mathbf{P}^\top \nabla_{\theta} f(z_i; \theta_m^*)$  ▷ Compute gradient at  $\theta_m^*$  and project to  $k$  dimensions
11:     $\mathbf{Q}_{ii}^{(m)} \leftarrow 1 - p(z_i; \theta^*)$  ▷ Compute weighting term
12:   end for
13:    $\Phi_m \leftarrow [\phi_1; \dots; \phi_n]^\top$ 
14: end for
15:  $\mathbf{T} \leftarrow \left[ \frac{1}{m} \sum_{m=1}^M \Phi_m (\Phi_m^\top \Phi_m)^{-1} \Phi_m^\top \right] \left[ \frac{1}{m} \sum_{m=1}^M \mathbf{Q}^{(m)} \right]$ 
16: return SOFT-THRESHOLD( $\mathbf{T}$ )
```

D3M: Data Debiasing via Datamodeling

Training data correlation between **class (species)** and **extra feature (color)** leads to disparate performance.



Goal: “debias” dataset to improve *worst-group accuracy* (WGA):

$$\text{WGA} = \min_{\text{group } \in \{\text{tan dog, black dog, tan cat, black cat}\}} \text{Acc}(\text{group})$$

D3M: Data Debiasing via Datamodeling

Our approach: Data Debiasing via Datamodeling (D3M)

Compute impact of each training sample on WGA by predicting WGA as a function of dataset selection.

Worst-group accuracy on a (small) validation set

$$\text{WGA} \approx \sum D_i \cdot A_i$$

Binary; whether we select the i -th training sample

"Group alignment score"
Learned coefficient; impact
of point i on WGA

Find selection that maximizes worst-group accuracy by removing most harmful examples.



- + Only changes data
- + Less accuracy gap
- + Competitive accuracy
- + Only needs test set labels

D3M: Data Debiasing via Datamodeling

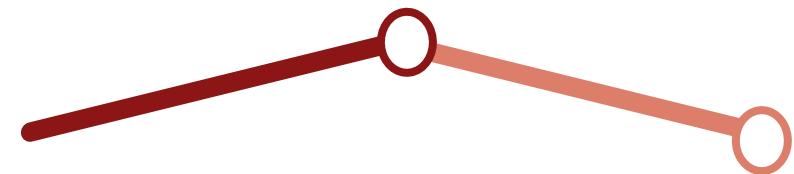
- Given a training dataset S_{train} and a validation dataset S_{val} , produce a classifier f to minimize **worst case loss** over predefined subpopulations

$$\max_{g' \in \mathcal{G}} \mathbb{E}_{(x,y,g) \sim \mathcal{D}} [\overline{\ell(f(x), y)} | g = g']$$

Annotations explaining the components:

- data distribution \mathcal{D} over triplets (x_i, y_i, g_i)
- loss function that quantifies the error of the classifier f in predicting the label y for input x
- set of distinct subpopulations in the data \mathcal{G}
- input $x_i \in X$
label $y_i \in Y$
subgroup label $g_i \in \mathcal{G}$
- (loss function is being evaluated conditional on the data belonging to subgroup g')

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Experimental Plan

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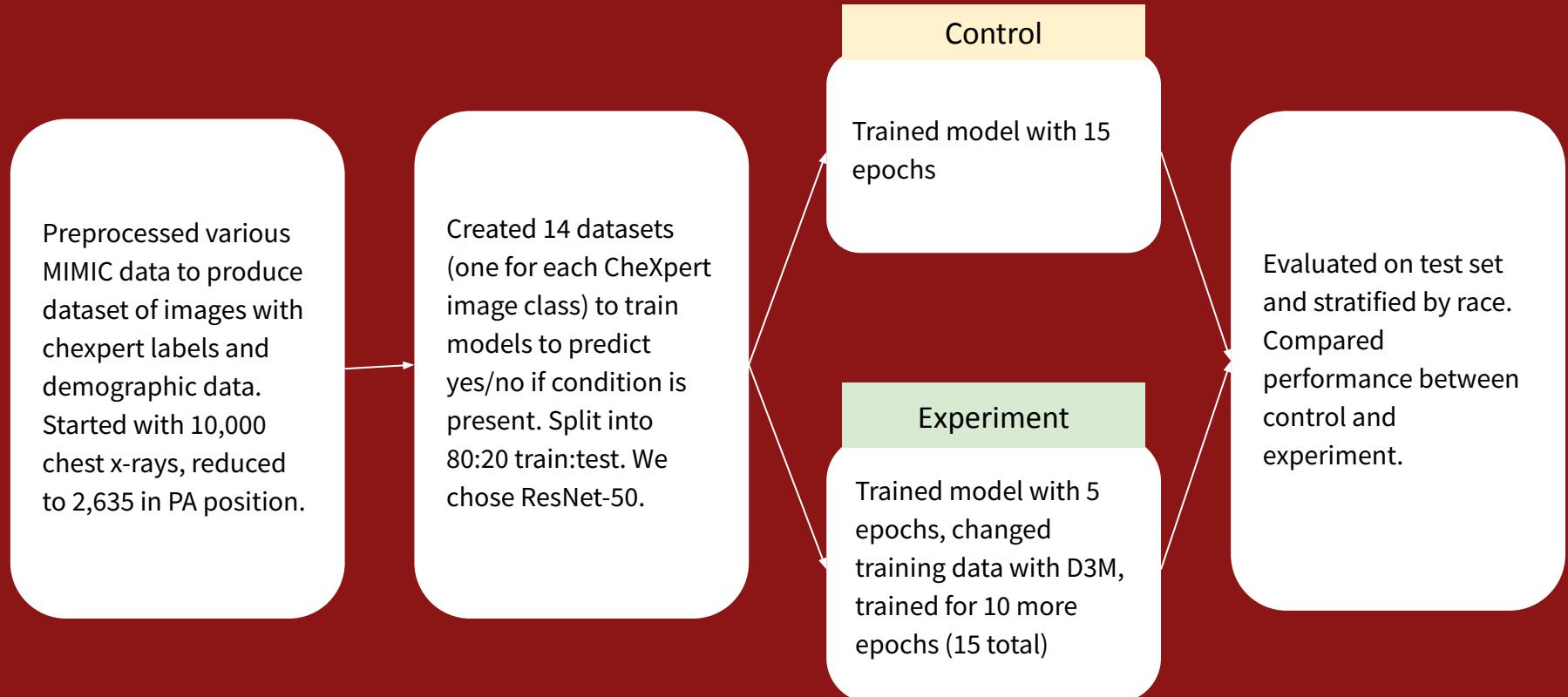
- **Datasets:** MIMIC-CXR (metadata) + MIMIC-CXR-JPG (images and CheXpert labels) + MIMIC-IV (demographic/racial data)
- **Models:** ResNet-9, ResNet-50
- **Procedure:** Break down 14 classes into binary datasets. Find the classes with the most discrepancies between groups. Train model - after 5 epochs training, apply 10 epochs of D3M + separately 10 epochs of normal training (15 each).

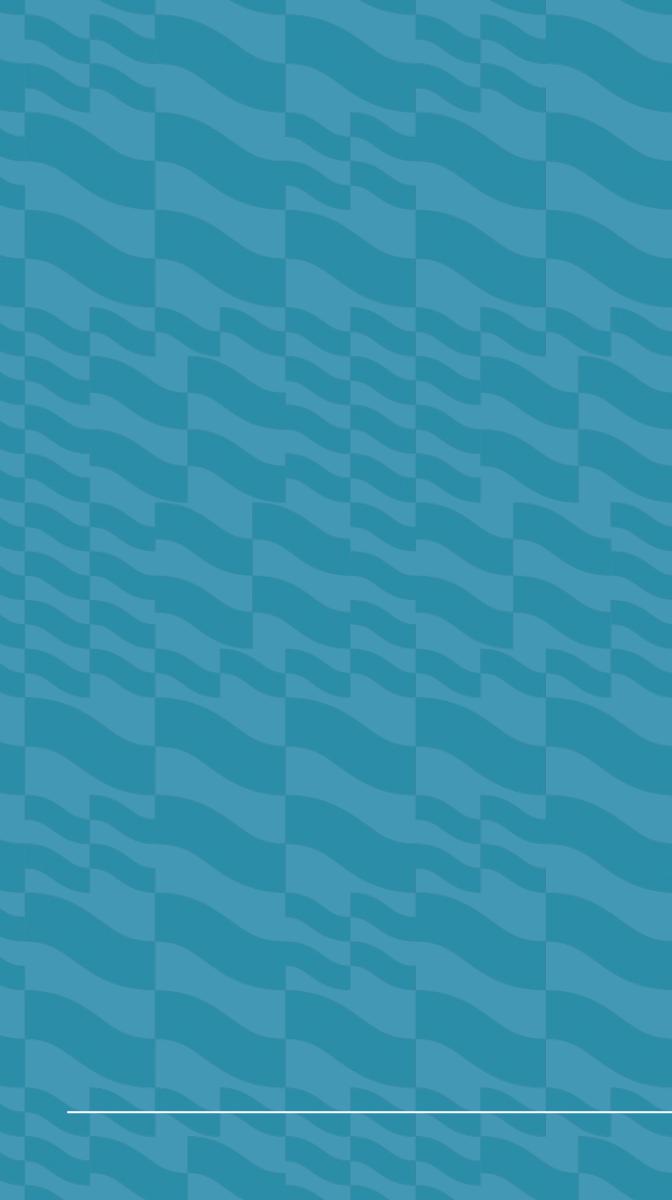
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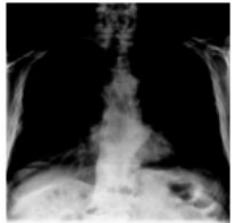


Results

TRAK

Top scoring TRAK images from the train set

Target: No Finding



No Finding



No Finding



No Finding



No Finding



No Finding

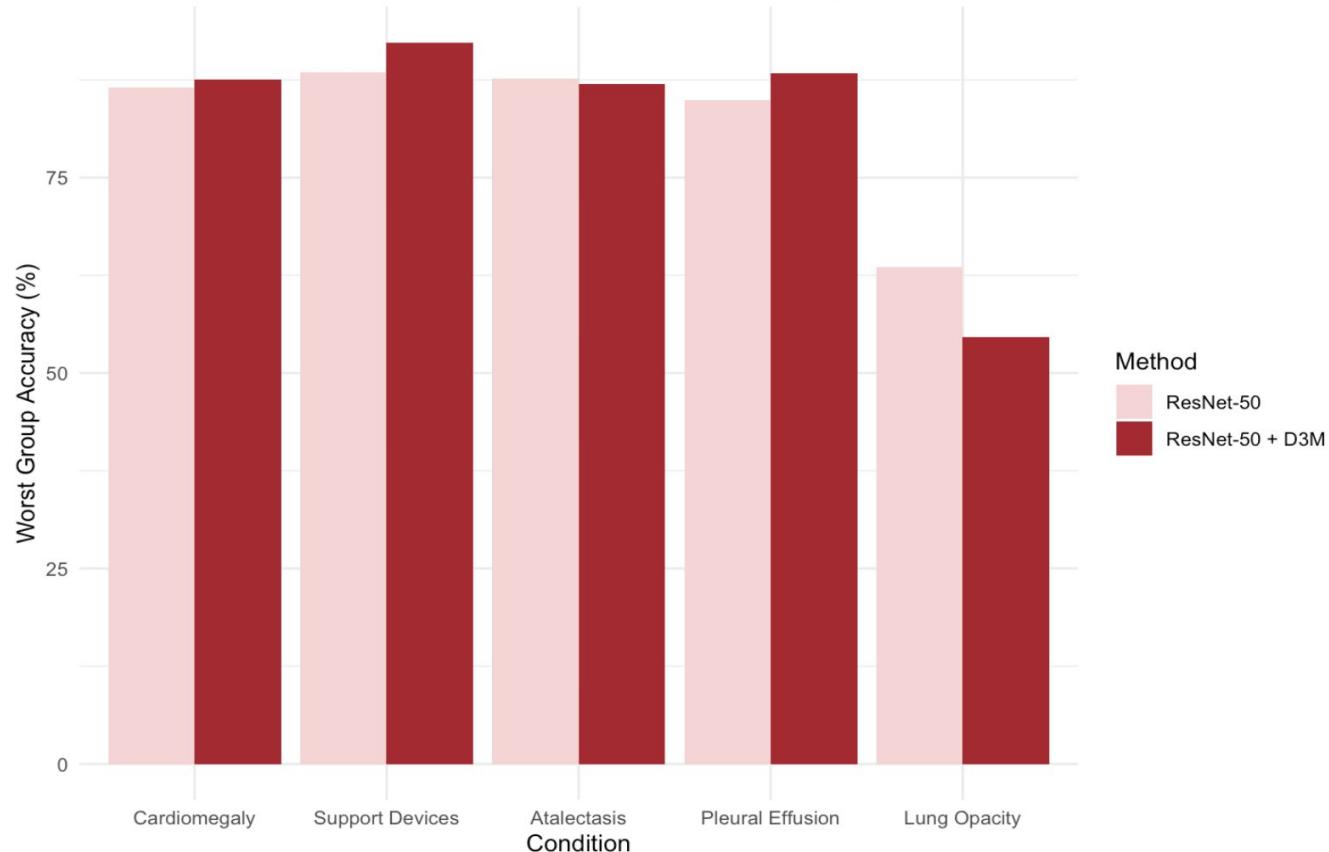


Worst Group Accuracies

Method	Worst Group Accuracy (%)				
	Cardiomegaly	Atalectasis	Support Devices	Pleural Effusion	Lung Opacity
ResNet-50	86.5%	87.7%	88.5%	84.9%	63.6%
ResNet-50 + D3M	87.6%	87.0%	92.3%	88.4%	54.6%

Worst Group Accuracies

Worst Group Accuracy Per Condition Across Racial Groups



Cardiomegaly

Race	Accuracy (%)			Δ in n
	ResNet-50 (n)	ResNet-50 + D3M (n)	Δ	
Asian	100.0 (65)	100.0 (51)	0.0	-14
Black or African American	86.5 (336)	87.6 (291)	+1.1	-45
Hispanic/Latino	92.1 (148)	92.1 (132)	0.0	-16
Other	95.2 (60)	95.2 (52)	0.0	-8
Unknown	93.9 (425)	93.9 (393)	0.0	-32
White	89.4 (1072)	88.3 (976)	-1.1	-96
Overall	90.5 (2108)	90.1 (1897)	0.0	-211

Support Devices

Race	Accuracy (%)			Δ in n
	ResNet-50 (n)	ResNet-50 + D3M (n)	Δ	
Asian	100.0 (65)	100.0 (48)	0.0	-17
Black or African American	98.9 (336)	98.9 (293)	0.0	-43
Hispanic/Latino	88.5 (148)	92.3 (139)	+3.8	-9
Other	100.0 (60)	100.0 (64)	0.0	4
Unknown	97.2 (425)	97.4 (377)	+0.2	-48
White	88.9 (1072)	89.3 (976)	+0.4	-96
Overall	93.0 (2108)	93.4 (1897)	0.4	-211

Atalectasis

Race	Accuracy (%)			Δ in n
	ResNet-50 (n)	ResNet-50 + D3M (n)	Δ	
Asian	93.8 (65)	93.8 (52)	0.0	-13
Black or African American	92.8 (336)	92.8 (299)	0.0	-37
Hispanic/Latino	91.4 (148)	91.4 (128)	0.0	-20
Other	91.7 (60)	95.8 (50)	+4.1	-10
Unknown	90.8 (425)	89.1 (373)	-1.7	-52
White	87.7 (1072)	87.0 (971)	-0.7	-101
Overall	89.8 (2108)	89.2 (1897)	-0.6	-211



Pleural Effusion

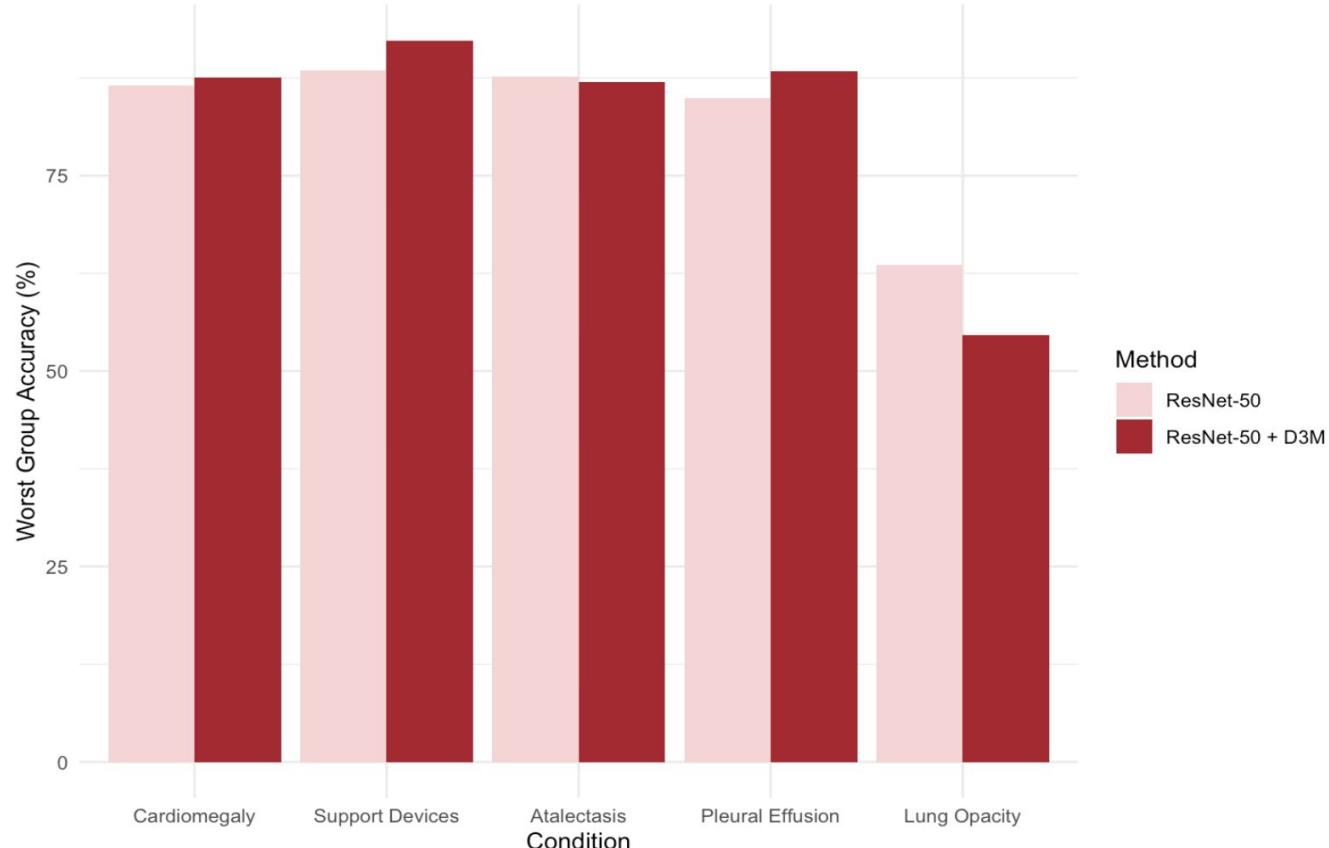
Race	Accuracy (%)			Δ in n
	ResNet-50 (n)	ResNet-50 + D3M (n)	Δ	
Asian	100.0 (65)	100.0 (59)	0.0	-6
Black or African American	84.9 (336)	88.4 (303)	+3.5	-33
Hispanic/Latino	91.9 (148)	86.5 (133)	-5.4	-15
Other	100.0 (60)	100.0 (54)	0.0	-6
Unknown	95.9 (425)	94.9 (355)	-1.0	-70
White	88.4 (1072)	89.1 (971)	+0.7	-101
Overall	90.1 (2108)	90.5 (1897)	0.4	-211

Lung Opacity

Race	Accuracy (%)			Δ in n
	ResNet-50 (n)	ResNet-50 + D3M (n)	Δ	
Asian	63.6 (65)	54.6 (60)	-9.1	-5
Black or African American	87.3 (336)	82.3 (299)	-5.0	-37
Hispanic/Latino	87.6 (148)	87.5 (132)	-0.1	-16
Other	89.5 (60)	89.5 (52)	0.0	-8
Unknown	92.6 (425)	92.3 (379)	-0.3	-46
White	85.2 (1072)	85.2 (975)	0.0	-97
Overall	86.7 (2108)	85.8 (1897)	-0.9	-211

Worst Group Accuracies

Worst Group Accuracy Per Condition Across Racial Groups

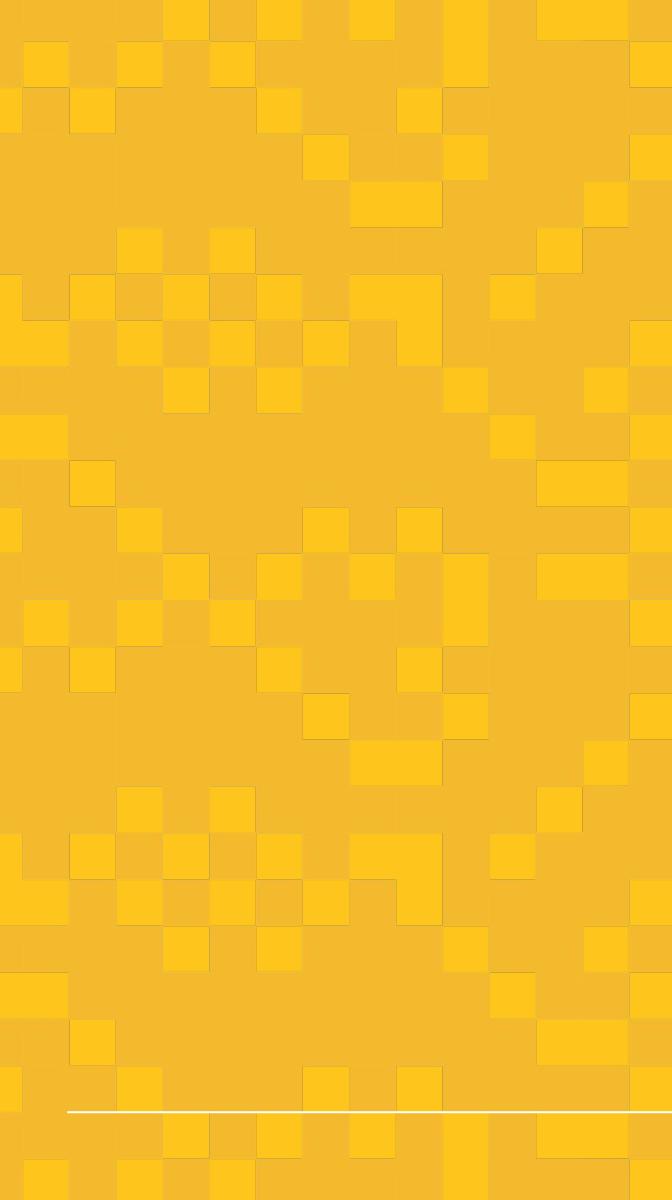


Quick note on D3M examples removed

```
def get_debiased_train_indices(
    self, group_alignment_scores, use_heuristic=True, num_to_discard=None
):
    """
    If use_heuristic is True, training examples with negative score will be discarded,
    and the parameter num_to_discard will be ignored
    Otherwise, the num_to_discard training examples with lowest scores will be discarded.
    """
    if use_heuristic:
        return [i for i, score in enumerate(group_alignment_scores) if score >= 0]

    if num_to_discard is None:
        raise ValueError("num_to_discard must be specified if not using heuristic.")

    sorted_indices = sorted(
        range(len(group_alignment_scores)),
        key=lambda i: group_alignment_scores[i],
    )
    return sorted_indices[num_to_discard:]
```



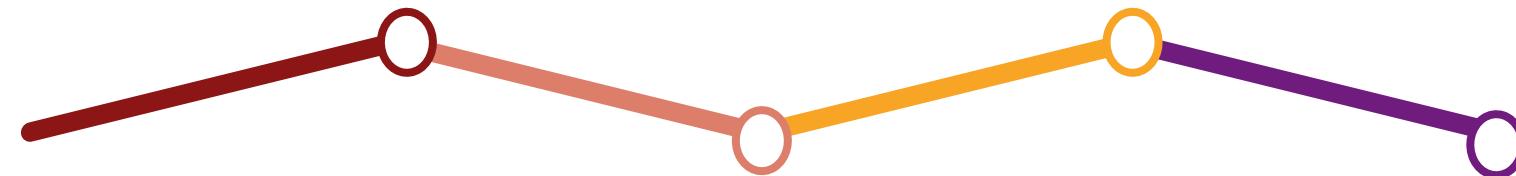
Analysis

Analysis

- **Did it work?** D3M increased accuracy of some subgroups but decreased accuracy of others.
- **Considerations:** Dataset size, domain, type of classification.
- **Purpose:** The intended purpose of D3M in our context was specifically to improve accuracy of minority groups.

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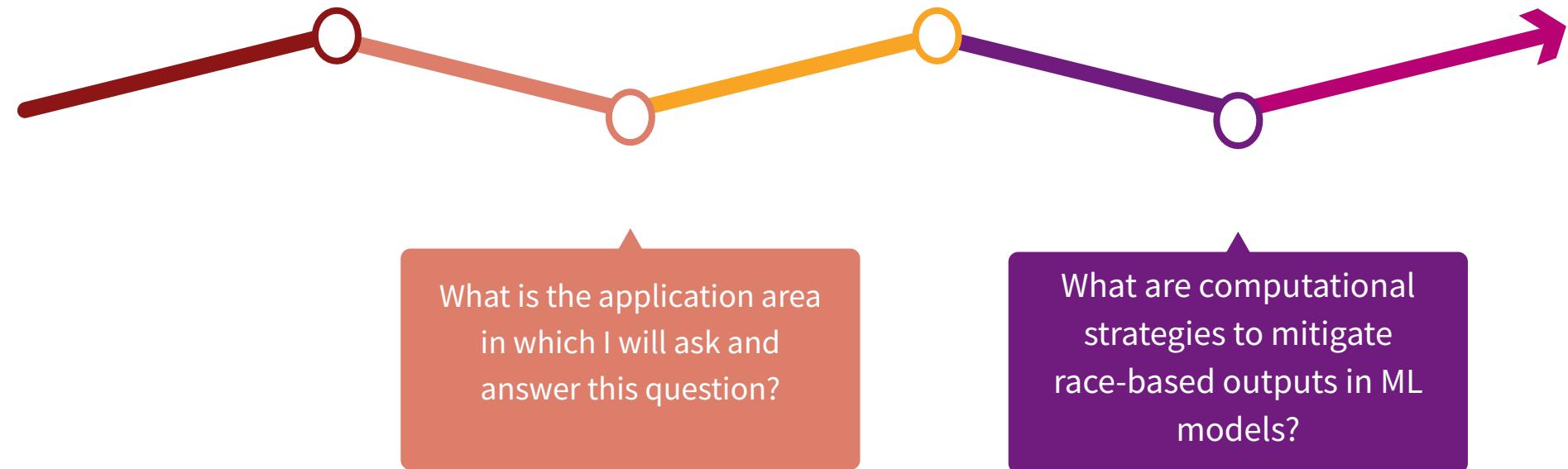


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Future Directions

- More data!
- Error analysis - what examples contributed to misclassification, specifically for which race?
- Other tasks beyond chest x-rays

A vertical decorative border on the left side of the slide features a repeating pattern of interlocking red shapes, resembling stylized leaves or petals, against a dark red background.

Thank You!

Huge thank you to Nigam!

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Alyssa Unell

Citations

Park, S. M., Georgiev, K., Ilyas, A., Leclerc, G., & Madry, A. (2023). TRAK: Attributing Model Behavior at Scale. arXiv preprint arXiv:2303.14186.

Jain, S., Hamidieh, K., Georgiev, K., Ilyas, A., Ghassemi, M., & Madry, A. (2024). Data Debiasing with Datamodels (D3M): Improving Subgroup Robustness via Data Selection. arXiv preprint arXiv:2406.16846.

I learned SO MUCH this rotation!

- Cuda
- Semaphore
- Not to use the login node
- Research Park
- Srun
- Importance of word choice

Background

- **TRAK:** data attribution method giving us coefficients (scores) to help identify examples that exacerbate discrepancies in group performance
- **D3M:** allows us to actually remove the examples and retrain on a dataset without the harmful examples