# Auxiliary Variables help Improving Group Robustness Through Bias Amplification

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## Abstract

Neural networks produced by standard training tend to suffer from poor accuracy on rare subgroups despite achieving high accuracy on average due to the *spurious correlation*. Previously we proposed BAM, a novel two-stage training algorithm, comprising a *bias amplification* stage with the learnable *auxiliary variable* and rebalanced training stage using upweighted error samples, which empirically improves the worst-group accuracy in various benchmarks with state-of-the-art performance. To investigate the roles of different elements in BAM, including the *auxiliary variables* and *Class Difference* stopping criterion, in this paper, we adopt two new datasets and conduct extensive experiments with carefully controlled parameters. With these experiments, we not only justify the assumption on the effectiveness of BAM but also shows the ubiquity of ClassDiff, for which we provide detailed discussions.

## 1 Introduction

This is a followup work based on our prior work  $[22]$ <sup>[1](#page-0-0)</sup>.

The *spurious correlation* is a phenomenon in machine learning where a model tends to "learn" certain decision rules based on spurious features such as backgrounds, thus likely to have unintended behaviors for subgroups from a data distribution. It prevails in various fields, including computer vision [\[1\]](#page-6-0), natural language processing [\[9\]](#page-6-1), and reinforcement learning [\[20\]](#page-7-1), and due to the ubiquity and gravity of this problem, extensive efforts have been made to address it.

Prior works mainly include methods that require group annotations for the whole training set or only the validation set. For the latter, they usually train two models, where the first is used to identify minority samples, based on which a second model is trained focusing on improving worst-group classification performance [\[23,](#page-7-2) [26,](#page-7-3) [27,](#page-7-4) [40\]](#page-8-0). In contrast, in our prior work, we proposed the BAM algorithm, which only requires training one model. BAM also contributes some interesting insights, such as the two-stage training process comprising bias amplification with squared loss and rebalanced training. With experiments on some commonly used benchmarks [\[35,](#page-7-5) [30,](#page-7-6) [24,](#page-7-7) [30,](#page-7-6) [36,](#page-7-8) [30,](#page-7-6) [2,](#page-6-2) [18\]](#page-6-3), we observed that BAM leads to consistent improvement in worst-group accuracy and achieves state-ofthe-art performance. However, to better understand why this procedure works well and how this approach can be applied to more generalized settings, we need to look deeper into different elements in BAM and perform experiments on more carefully constructed datasets.

In this work, we adopt two additional datasets: Controlled-Waterbirds and Colored-MNIST. By carefully controlling their specifications and conducting experiments with various settings, such as

<span id="page-0-0"></span> $1$ <sup>W</sup>e only provide abridged version of Related Works and Methodology Sections to avoid too much repetition. Please refer to the original paper for more details.

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auxiliary variables, upweight factors, stopping epochs, and losses, we gain deeper insights into each of them and validate the robustness of BAM.

## 2 Related Works

A variety of recent work discussed different realms of robustness, for instance, class imbalance [\[10,](#page-6-4) [12,](#page-6-5) [15,](#page-6-6) [14,](#page-6-7) [33\]](#page-7-9), and robustness in distribution shift, where the target data distribution is different from the source data distribution [\[5,](#page-6-8) [39,](#page-8-1) [25,](#page-7-10) [19,](#page-6-9) [38\]](#page-8-2). In this paper, we mainly focus on improving group robustness. Categorized by the amount of information we have for training and validation, we discuss three directions below:

Improving Group Robustness with Training Group Annotations. Multiple works have used training group annotations to improve worst-group accuracy [\[3,](#page-6-10) [16,](#page-6-11) [8,](#page-6-12) [4,](#page-6-13) [31\]](#page-7-11). Other works include minimizing the worst-group training loss using distributionally robust optimization (Group-DRO) [\[30\]](#page-7-6), simple training data balancing (SUBG) [\[13\]](#page-6-14), and retraining the last layer of the model on the group-balanced dataset (DFR) [\[17\]](#page-6-15). These methods achieve state-of-the-art performance on all benchmark datasets. However, acquiring spurious attributes of the entire training set is costly and unrealistic in real-world datasets.

Improving Group Robustness with Validation Group Annotations Only. Acknowledging the cost of obtaining group annotations, many recent works focus on the setting where training group annotations are not available [\[7,](#page-6-16) [28,](#page-7-12) [21,](#page-7-13) [29\]](#page-7-14). Approaches closely related to our method usually use the first model to identify minority samples and then train a separate model based on the results predicted by the first model [\[37,](#page-7-15) [34\]](#page-7-16). For example, JTT [\[23\]](#page-7-2) first trains an ERM model to identify minority groups in the training set (similar to EIIL [\[6\]](#page-6-17)), and then trains a second ERM model with these selected samples to be upweighted. However, these approaches all focus on the robust training of the second model and fail to consider the potential of accumulating biased knowledge from the first model.

Improving Group Robustness without any Group Annotations. Relatively little work has been done under the condition that no group information is provided for both training and validation. [\[13,](#page-6-14) [23\]](#page-7-2) observe a significant drop ( $10\%$  -  $25\%$ ) in worst-group test accuracy if using the highest *average* validation accuracy as the stopping criterion without any group annotations. A recent work, GEORGE [\[32\]](#page-7-17), tries to separate unlabeled classes in deep model feature spaces and then use the generated pseudo labels to train the model via the distributionally robust optimization objective. However, there is a considerable performance gap between GEORGE and the supervised methods.

## <span id="page-1-0"></span>3 Methodology

In the previous work, we designed the BAM training algorithm consisting of a two stages.

In Stage 1, we train a bias-amplified model to bias toward majority group samples unintentionally. Inspired by the work done by [\[11\]](#page-6-18), we introduce trainable auxiliary variables  $b_i$  for each data sample and add it to the network's output. The objective function is formally defined as:

$$
R_1(\theta, B) = \frac{1}{n} \sum_{i=1}^{n} \ell(f_{\theta}(x_i) + \lambda b_i, y_i).
$$
 (1)

where  $f_{\theta}: \mathcal{X} \to \mathbb{R}^C$  is a neural network with parameters  $\theta$  and the class number  $C = |\mathcal{Y}|$ , and  $\lambda$  is a hyperparameter that controls the strength of the auxiliary variables. We also further adopt squared  $\lim_{y \to \infty} \ell(z, y) = \|z - e_y\|_2^2$  where  $e_y \in \mathbb{R}^C$  is the one-hot encoding for the label y.

In stage 2, we continue to train the (same) model using the error set we obtained in Stage 1, i.e., data points the model misclassifies. We adopt the rebalanced loss that upweights the examples in the error set E:

$$
R_2(\theta) = \mu \sum_{(x,y)\in E_1} \ell_{CE}(f_{\theta}(x), y) + \sum_{(x,y)\in D\setminus E} \ell_{CE}(f_{\theta}(x), y),
$$
\n(2)

where  $\ell_{CE}$  is the cross-entropy  $\{x,y\} \in E$   $\mu$  is a hyperparamter (upweight factor).

In particular, we have a special stopping criterion for Stage 2 training. Note that the group annotations may or may not be available in our setting. We will stop at the highest worst-group validation accuracy if it is available. However, if there are no group annotations presented, we will instead use the *minimum class difference*, i.e., stop when the sum of pairwise validation accuracy differences between classes is at the minimum. The class difference can be formally defined as

ClassDiff = 
$$
\sum_{i,j=1}^{C} |Acc(\text{class } i) - Acc(\text{class } j)|.
$$
 (3)

In this work, we use the same method while focusing more on the experiment side.

## 4 Experiments

#### 4.1 Datasets

The previous method we proposed, BAM, has been tested on datasets including Waterbirds, CelebA, CivilComments-WILDS, and MultiNLI (see Table [3](#page-9-0) for specifications) and outperformed state of the art on nearly all of them. Some ablation studies about loss functions and auxiliary variables have also been done to prove that all the model components are useful. However, as we see in Table [3,](#page-9-0) the sizes of different classes and groups are not balanced on all four datasets. As the training process becomes undetermined and unpredictable because of the nature of these datasets, we could not perform a fine-grained analysis of the role of each model component. In addition, Waterbirds and CelebA have significant distribution shifts in the training and test data, and two NLP datasets have too much noise that impedes deep learning models from learning well on the classification task, even without spurious correlation.

To further investigate how auxiliary variables, upweight factors, stopping epochs in Stage 1, and losses play their roles in BAM and address the problems above, we consider two new CV datasets: Controlled-Waterbirds and Colored-MNIST, and manually control their class and group counts. In Colored-MNIST, we set class 0 for numbers from 0 to 4 and class 1 for numbers from 5 to 9. We make the class labels spuriously correlated with colors and randomly flip the color of a small subset to generate minority groups. Figure [1](#page-2-0) shows what their majority and minority groups are. Table [1](#page-2-1) shows some parameters of the two datasets.

<span id="page-2-0"></span>

Figure 1: Based on their group sizes, for Controlled-Waterbirds on the left, WW and LL are majority groups. Similarly, for Colored-MNIST on the right, Class 0 in red and Class 1 in Green are majority groups.

<span id="page-2-1"></span>Table 1: Important parameters to construct Controlled-Waterbirds and Colored-MNIST.

<b>Dataset</b>	Parameter	Description	
Controlled-Waterbirds		group sizes of WW, WL, LW, LL in order	
	S	total dataset size	
Colored-MNIST		minority group ratios in class $0 < 1$	
	cr	class ratios of class $0 & 1$	
	S	total dataset size	

#### 4.2 The Role of Auxiliary Variables

To validate that auxiliary variables help improve the model performance in a non-trivial way, we did additional ablation studies on new datasets. Section [4.2](#page-3-0)

Figure [2](#page-3-0) shows the change of model performance with different auxiliary variables. Note that  $\lambda = 0$ will disable the auxiliary variables to fit training samples. With a moderately large  $\lambda \cdot b$ , it is easy to observe that the model achieves higher robust worst-group test accuracy than without  $b$  on both datasets. It indicates the important role of auxiliary variables.

<span id="page-3-0"></span>

Figure 2: Worst-group accuracy with different lambdas on Controlled-Waterbirds (left) and Colored-MNIST (right). For Controlled-Waterbirds, we fix dataset parameters  $d = \{1800, 200, 200, 1800\}$ and hyperparameters  $\mu = 100, T = 100$ . For Colored-MNIST, we fix dataset parameters  $fr =$  ${0.1, 0.1}$ ,  $cr = 1.0$ ,  $s = 50000$  and hyperparameters  $\mu = 10$ ,  $T = 50$ . For simplicity, we choose the same hyperparameters of Controlled-Waterbirds as in Waterbirds in the previous work.

#### 4.3 Negative Correlations of ClassDiff and Worst-Group Accuracy

Section [3](#page-1-0) It explains the definition of ClassDiff, which is mainly used to evaluate when to stop Stage 2 without the help of validation group annotations. In Figure 3 of the previous work BAM, we show the negative correlation between absolute validation class difference and worst-group accuracy on all four benchmarks (Waterbirds, CelebA, CivilComments-WILDS, MultiNLI). In this project, we did abundant experiments to verify its success on new datasets by varying the dataset sizes, class size ratios, and group size ratios over a large range. In every single setting we tested, we observed similar negative correlations. These findings suggest that ClassDiff could be useful in general when no group annotation is available, as it is robust across different datasets and parameters. Here we show one example per dataset each in Figure [3.](#page-3-1) Appendix [A.3](#page-9-1) displays a larger and random subset of our experiments without cherry picking, and explains the subset generation details.

<span id="page-3-1"></span>

Figure 3: Visualization on relations between ClassDiff and worst-group validation accuracy. Left: Controlled-Waterbirds.  $d = \{1400, 200, 600, 4200\}, \lambda = 50, T = 100, \mu = 100$ . Right: Colored-MNIST.  $s = 50000$ ,  $fr = \{0.2, 0.2\}$ ,  $cr = 1.0$ ,  $\lambda = 20$ ,  $T = 11$ ,  $\mu = 5$ .

## 5 Discussions

Different Dataset Sizes Since the total size of Waterbirds is limited, varying the total size could cause a very small size of minority groups. Figure [4](#page-4-0) shows the performance comparison of  $\lambda = 0$  vs.  $\lambda = 50$ . On Colored-MNIST, we see a tiny but persistent difference of different  $\lambda$ 's on robust worstgroup test accuracy from the left plot. The relation between ClassDiff and  $\lambda$  seems not observable, while ClassDiff achieves good performance on all the total dataset size s listed above. The robust test accuracy is comparably robust to s.

<span id="page-4-0"></span>

Figure 4: Robust worst-group accuracy and minimum ClassDiff accuracy for different dataset sizes on Colored-MNIST. We consider  $s \in \{1000, 2000, 5000, 20000, 50000\}$ , fix  $fr = \{0.1, 0.1\}$ ,  $cr = 1.0$ , and tune  $\mu \in \{5, 10, 20, 50\}, T \in \{20, 50\}.$ 

Different Class Imbalance Ratios Figure [5](#page-4-1) shows the robustness of BAM w.r.t. class imbalance ratios.

<span id="page-4-1"></span>

Figure 5: Robust worst-group accuracy for different class imbalance ratios. For Controlled-Waterbirds (left), we choose WW = 1400, WL = 200,  $\lambda = 50$ ,  $T = 100$ . For Colored-MNIST (right), we choose  $s = 20000$ ,  $fr = \{0.1, 0.1\}$ ,  $\lambda = 50$ ,  $T = 20$ . Each point is averaged over 3 random experiments.

Different Group Imbalance Ratios Figure [6](#page-4-2) shows the robustness of BAM w.r.t. group imbalance ratios.

<span id="page-4-2"></span>

Figure 6: Robust worst-group accuracy for different class imbalance ratios. For Controlled-Waterbirds (left), we choose WW = 1800, LL = 1800,  $\lambda = 50$ , T = 100. For Colored-MNIST (right), we choose  $s = 20000$ ,  $cr = 1.0$ ,  $\lambda = 50$ ,  $T = 50$ . Each point is averaged over 3 random experiments.

**Different Upweight Factors** Since Colored-MNIST is a relatively simple dataset for addressing spurious correlation problems, the change of worst-group performance is fairly tiny by varying the upweight factor  $\mu \in \{2, 5, 10, 20, 50\}$ . Appendix [A.4](#page-9-2) shows one example of the corresponding accuracy with all the  $\mu$ 's in the above set. On Controlled-Waterbirds, the trend is more obvious. In general, as  $\mu$  increases, the worst-group test accuracy first rises and then drops. However, over the set  $\mu \in \{10, 50, 70, 100, 120, 150\}$  that we tested, it is robust to  $\mu$  within a large range from 50 to 150. Figure [5,](#page-4-1) Figure [6](#page-4-2) and Table [2](#page-5-0) show such robustness.

<span id="page-5-0"></span>Table 2: Experiments on Controlled-Waterbirds dataset. Parameters and hyperparameters are  $d =$  $\{1800, 200, 200, 1800\}, \lambda = 50.$ 

	Upweight factor $\mu$						
	10	50	70	100	120	150	
10	74.4	84.6	86.1	84.6	84.6	84.6	
20	72.2	84.6	87.2	86.1	87.2	84.6	
50	63.9	86.1	86.1	88.9	86.1	86.1	

**Different Stage 1 Epochs** In Figure [10](#page-12-0) and Figure [11,](#page-13-0) KDE-plot shows how the values of auxiliary variables changes as Stage 1 proceeds. Carefully controlling all other parameters and hyperparameters, we observe some clear patterns below on these two datasets: First, as  $T$  grows, the logits become larger and increasingly influence the model prediction outputs. Second, minority and majority groups are easier to be separated from each other, resulting in better error sets.

In summary, a larger T will generally lead to a better error set, while the model prediction will be wrong on majority groups if  $\overline{T}$  is too large.



Figure 7: (a)(b) Epoch 1 and 100 in Stage 1 on Waterbirds. Logit 0 corresponds to the prediction on the waterbird class, and logit 1 corresponds to landbird. The group sizes are 1800, 200, 200, 1800 in order. (c)(d) Epoch 1 and 4 in stage 1 on Colored-MNIST. Logit 0 corresponds to the prediction on class 0, and logit 1 corresponds to class 1.

## 6 Conclusion

In this paper, we tested and analyzed BAM on two new datasets and verified the robustness of BAM varying a large range of parameters and hyperparameters (e.g., the total data size, class imbalance ratio, and group imbalance ratio,  $\mu$ , T). In addition, we demonstrate the effectiveness of auxiliary variables by varying values of  $\lambda$ . We also validate the negative correlation between ClassDiff and worst-group validation accuracy, which indicates its potential to supplant validation group annotations for less burden.

For future work, we will

- Look deeper into the effects of different losses in Stage 1.
- Validate BAM on more complex datasets such as CIFAR10-MNSIT, and test ClassDiff on multi-classification datasets.
- Aalternate the second stage of BAM with the method proposed by [\[40\]](#page-8-0), and analyze whether this could improve the performance of the second stage.

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## A Appendix

#### A.1 Team contributions



#### A.2 Prior Dataset Specifications

<span id="page-9-0"></span>



### <span id="page-9-1"></span>A.3 ClassDiff Visualization

We generate a random subset of experiments with size 10 and visualize their relations between ClassDiff and worst-group validation accuracy. First, we sort all the experiment logs on Greatlakes by their starting training time. Then, fix 42 as the random seed of Numpy and generate a random indices list with size 10. We select the experiments by these indices and plot the ClassDiff relations to avoid cherry picking.

#### <span id="page-9-2"></span>A.4 Upweight Factors Visualization

Table [4](#page-9-3) shows that Colored-MNIST does not have clear differences when vary upweight factors.

<span id="page-9-3"></span>Table 4: Experiments on Colored-MNIST dataset. Parameters and hyperparameters are  $s =$  $50000, fr = \{0.1, 0.1\}, cr = 1.0, \lambda = 50.$ 



A.5 Different Epochs visualization



Figure 8: 10 random experiments of the relations between ClassDiff and worst-group validation accuracy on Controlled-Waterbirds.



Figure 9: 10 random experiments of the relations between ClassDiff and worst-group validation accuracy on Colored-MNIST.

<span id="page-12-0"></span>

Figure 10: Different epochs in Stage 1 on Waterbirds. Logit 0 corresponds to the prediction on the waterbird class, and logit 1 corresponds to landbird. The group sizes are 1800, 200, 200, 1800 in order.

<span id="page-13-0"></span>

Figure 11: Different epochs in Stage 1 on Colored-MNIST. Logit 0 corresponds to the prediction on class 0, and logit 1 corresponds to class 1.