

# The Quest for Reliable Melanoma Detection

7 Years at the Vanguard of Skin Lesion Analysis

#### Sandra Avila

Institute of Computing University of Campinas, Brazil

ISIC Workshop, CVPR 2021



# Skin Cancer

Why do we care?



# RECOD Lab.

melanoma research 7 years 2014–2021









# Metrópole

Inteligência artificial pode diagnosticar tumor de pele que até médico não vê





# Classification

Segmentation

Systematic Evaluation

**Syn**thesis

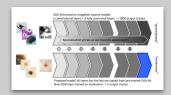
**Deb**iasing

**Data Augmentation** 

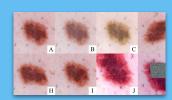




Statistical learning approach for robust melanoma screening

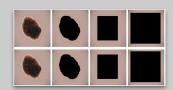


Towards automated melanoma screening: Proper computer vision & reliable results



Data augmentation for skin lesion analysis

Skin lesion synthesis with generative adversarial networks



Debiasing skin lesion datasets and models? Not so fast

Less is more: Sample selection and label conditioning improve skin lesion segmentation

2014

2015

2016

2017

2018

2019

2020

2021

Fixed Weights

Rew Model

Input

Original Model

Towards robust melanoma screening: A case for enhanced mid-level features

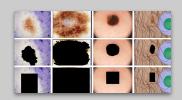
Transfer schemes for deep learning in image classification



Knowledge transfer for melanoma screening with deep learning

RECOD Titans at ISIC challenge 2017

Data, depth, and design: Learning reliable models for skin lesion analysis



(De)Constructing bias on skin lesion datasets

Solo or ensemble? Choosing a CNN architecture for melanoma classification



GAN-based data augmentation and anonymization for skin-lesion analysis: A critical review

An evaluation of self-supervised pre-training for skin-lesion analysis

# Towards Automated Melanoma Screening: Proper Computer Vision & Reliable Results

Michel Fornaciali, Micael Carvalho, Flávia Vasques Bittencourt, Sandra Avila, Eduardo Valle

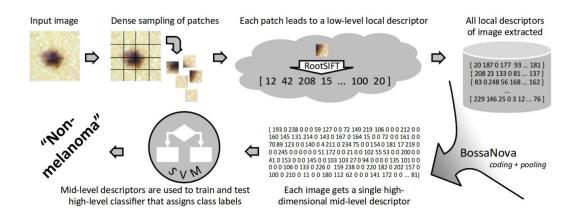
Abstract—In this paper we survey, analyze and criticize current art on automated melanoma screening, reimplementing a baseline technique, and proposing two novel ones. Melanoma, although highly curable when detected early, ends as one of the most dangerous types of cancer, due to delayed diagnosis and treatment. Its incidence is soaring, much faster than the number of trained professionals able to diagnose it. Automated screening appears as an alternative to make the most of those professionals, focusing their time on the patients at risk while safely discharging the other patients. However, the potential of automated malanama diagnosis is summently unfulfilled due to



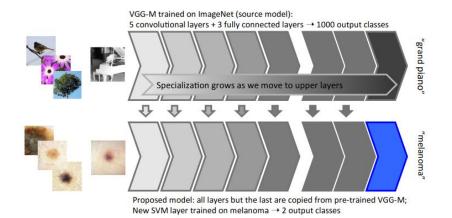




Paper, Code & Data: <a href="https://github.com/learningtitans/melanoma-screening">https://github.com/learningtitans/melanoma-screening</a>



Bag-of-Visual-Words pipeline



Deep Transfer Learning pipeline

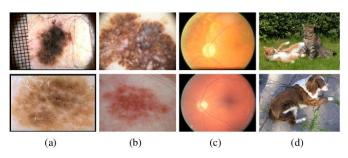
#### **Knowledge Transfer for Melanoma Screening with Deep Learning**

Afonso Menegola<sup>†‡</sup>, Michel Fornaciali<sup>†‡</sup>, Ramon Pires°, Flávia Vasques Bittencourt•, Sandra Avila<sup>†</sup>, Eduardo Valle<sup>†\*</sup>

†RECOD Lab, DCA, FEEC, University of Campinas (Unicamp), Brazil °RECOD Lab, IC, University of Campinas (Unicamp), Brazil •School of Medicine, Federal University of Minas Gerais (UFMG), Brazil

#### ABSTRACT

Knowledge transfer impacts the performance of deep learning — the state of the art for image classification tasks, including automated melanoma screening. Deep learning's greed for large amounts of training data poses a challenge for medical tasks, which we can alleviate by recycling knowledge from models trained on different tasks, in a scheme called *transfer learning*. Although much of the best art on automated melanoma screening employs some form of transfer learning, a systematic evaluation was missing. Here we investigate the presence of transfer, from which task the transfer is sourced, and





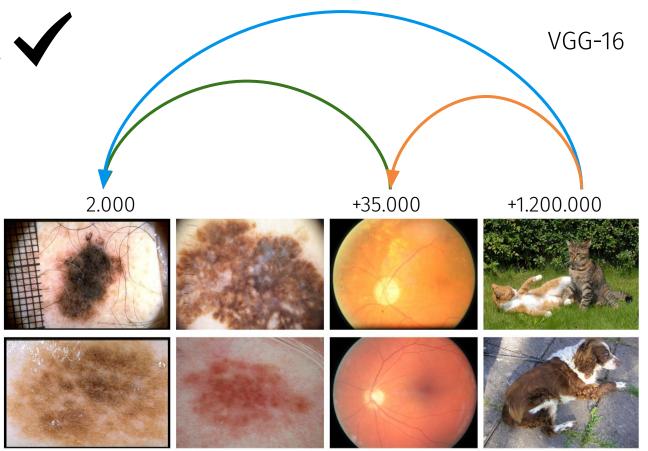
Paper, Code & Data: <a href="https://github.com/learningtitans/melanoma-transfer">https://github.com/learningtitans/melanoma-transfer</a>

ImageNet -> Melanoma

Double Transfer:

ImageNet -> Retina &

Retina -> Melanoma



## RECOD Titans at ISIC Challenge 2017

Afonso Menegola<sup>†</sup>, Julia Tavares<sup>†</sup>, Michel Fornaciali, Lin Tzy Li, Sandra Avila, Eduardo Valle\*

#### HISTORY

Our team has worked on melanoma classification since early 2014 [1], and has employed deep learning with transfer learning for that task since 2015 [2]. Recently, the community has started to move from traditional techniques towards deep learning, following the general trend of computer vision [3]. Deep learning poses a challenge for medical applications, due to the need of very large training sets. Thus, transfer learning becomes crucial for success in those applications, motivating our paper for ISBI 2017 [4].

Our team participated in Parts 1 and 3 of the ISIC Challenge 2017, described below in that order. Although our team has a long experience with skin-lesion classification (Part 3), this Challenge was the very first time we worked on skin-lesion segmentation (Part 1).

The code needed to reproduce our results is at our code repository<sup>4</sup>.

#### C. Data Augmentation

We used online image augmentation, with up to 10% horizontal and vertical shifts, up to 20% zoom, and up to  $270^\circ$  degrees rotation. Images were first resized — we tried  $256\times256$  and  $128\times128$ , ultimately keeping the latter, which was faster and resulted in similar performance. Transforming the images before resizing them was slower and did not improve the results.

#### D. Experiments

Our first attempt was a model based on the VGG network [6]. The first part of the model consisted of the VGG-16 layers











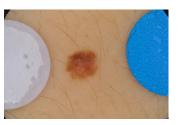


Melanoma Classification

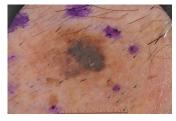


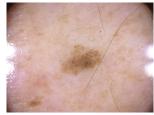
Paper, Code & Data: <a href="https://arxiv.org/abs/1703.04819">https://arxiv.org/abs/1703.04819</a>

## ISIC Challenge 2017









Training data 2000 images

95.1%

(internal validation)

Validation data 150 images

90.8%

Test data 600 images

87.4%

## RECOD Titans at ISIC Challenge 2017

Afonso Menegola<sup>†</sup>, Julia Tavares<sup>†</sup>, Michel Fornaciali, Lin Tzy Li, Sandra Avila, Eduardo Valle\*

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Melanoma Classification



Paper, Code & Data: <a href="https://arxiv.org/abs/1703.04819">https://arxiv.org/abs/1703.04819</a>

## ISIC Challenge 2017

Models + data



Image resolution

Class/sample-weighting schemes

Curriculum learning

SVM decision layer

Training and test augmentation



Patient data



Per-image normalization







### $2^9$ factors × 5 datasets = 2560 experiments

Neurocomputing 383 (2020) 303-313



Contents lists available at ScienceDirect

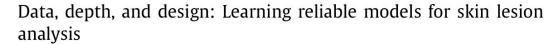
#### Neurocomputing

journal homepage: www.elsevier.com/locate/neucom









Eduardo Valle<sup>a,\*</sup>, Michel Fornaciali<sup>a</sup>, Afonso Menegola<sup>a</sup>, Julia Tavares<sup>a</sup>, Flávia Vasques Bittencourt<sup>b</sup>, Lin Tzy Li<sup>c,d</sup>, Sandra Avila<sup>c</sup>

- <sup>a</sup> School of Electrical and Computing Engineering, University of Campinas (UNICAMP), Av. Albert Einstein 400, Campinas, SP 13083-852, Brazil
- <sup>b</sup> School of Medicine, Federal University of Minas Gerais (UFMG), Alamedalvaro Celso 55, Belo Horizonte, MG 30150-260, Brazil
- c Institute of Computing, University of Campinas (UNICAMP), Av. Albert Einstein 1251, Campinas, SP 13083-852, Brazil
- d Samsung R&D Institute Brazil (SRBR), Campinas, SP, Brazil







"Amount of train data has disproportionate influence, explaining **46%** of the variation in performance." "Other than data, the most important factor was the **use of data augmentation on test**."

**Table 3** Selected lines from the 176-line ANOVA table; most of the omitted lines (126) had p-values  $\geq 0.05$ . Absolute explanation based on  $\eta^2$ -measure, relative explanation ignores residuals and choice of test dataset (j).

	Factor	<i>p</i> -value	Explanation (%)		Best AUC (%)		Worst AUC (%)	
			Abs.	Rel.	Treatment	Mean	Treatment	Mean
a	Model architecture	< 0.001	0	1	resnet	84	inception	83
b	Train dataset	< 0.001	5	46	full	85	challenge	81
c	Input resolution	< 0.001	1	5	598	84	299-305	82
d	Data augmentation	0.17	0	0	default	83	custom	83
e	Input normalization	0.001	0	0	default	83	erase mean	83
f	Use of segmentation	< 0.001	0	2	no	84	yes	83
g	Duration of training	0.003	0	0	full	83	half	83
h	SVM layer	< 0.001	0	4	no	84	yes	83
i	Augmentation on test	< 0.001	1	12	yes	84	no	82
j	Test dataset	< 0.001	75		full.split	96	edra.clinical	66

Paper, Code & Data: <a href="https://github.com/learningtitans/data-depth-design">https://github.com/learningtitans/data-depth-design</a>

#### Data Augmentation for Skin Lesion Analysis

Fábio Perez<sup>1</sup>, Cristina Vasconcelos<sup>2</sup>, Sandra Avila<sup>3</sup>, and Eduardo Valle<sup>1</sup>

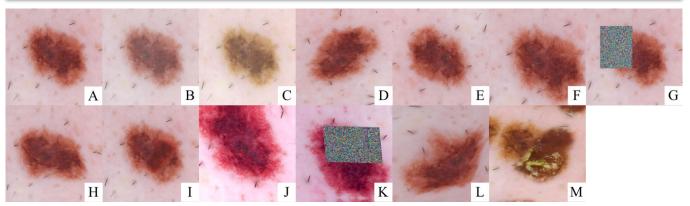
<sup>1</sup>RECOD Lab, DCA, FEEC, University of Campinas (Unicamp), Brazil <sup>2</sup>Computer Science Department, IC, Federal Fluminense University (UFF), Brazil <sup>3</sup>RECOD Lab, IC, University of Campinas (Unicamp), Brazil





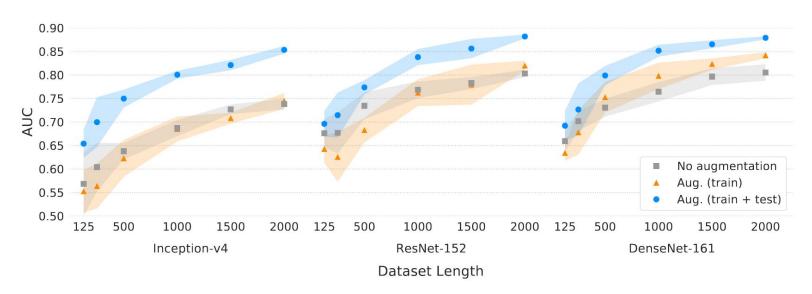






Paper, Code & Data: <a href="https://github.com/fabioperez/skin-data-augmentation">https://github.com/fabioperez/skin-data-augmentation</a>

#### Augmentation on Training & Testing



Paper, Code & Data: <a href="https://github.com/fabioperez/skin-data-augmentation">https://github.com/fabioperez/skin-data-augmentation</a>

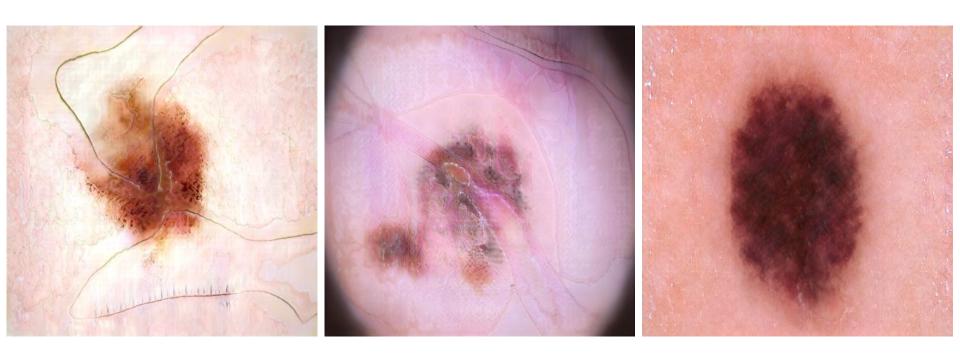
# Let's generate data!

Generative Adversarial Networks (GANs)

# I SEE BAD DATA



Spoiler alert! The film: "The Sixth Sense", the line: "I see dead people."



... using Progressive Growing GANs (PGAN)

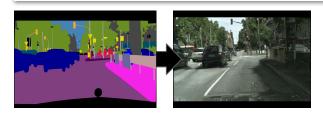
#### Skin Lesion Synthesis with Generative Adversarial Networks

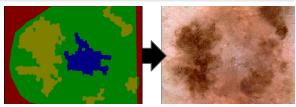
Alceu Bissoto<sup>1</sup>, Fábio Perez<sup>2</sup>, Eduardo Valle<sup>2</sup>, and Sandra Avila<sup>1</sup>

<sup>1</sup>RECOD Lab, IC, University of Campinas (Unicamp), Brazil

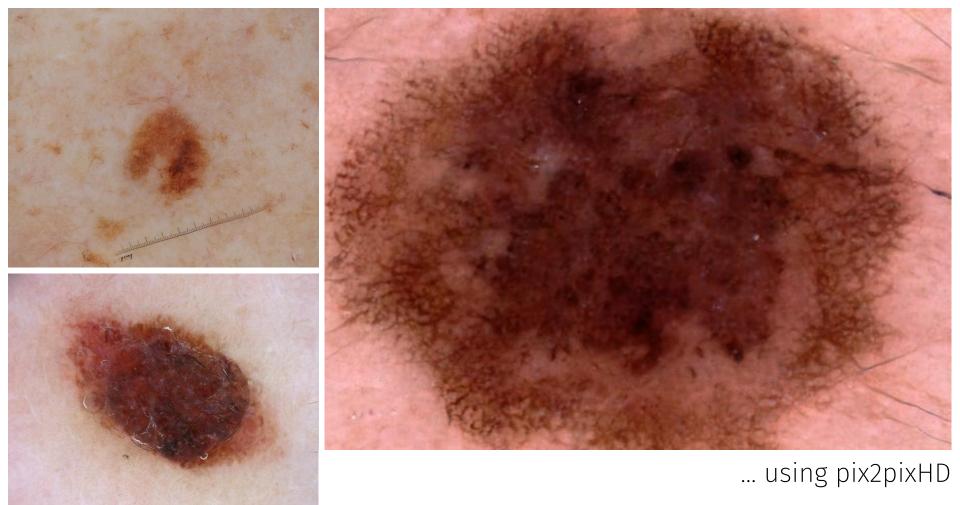
<sup>2</sup>RECOD Lab, DCA, FEEC, University of Campinas (Unicamp), Brazil



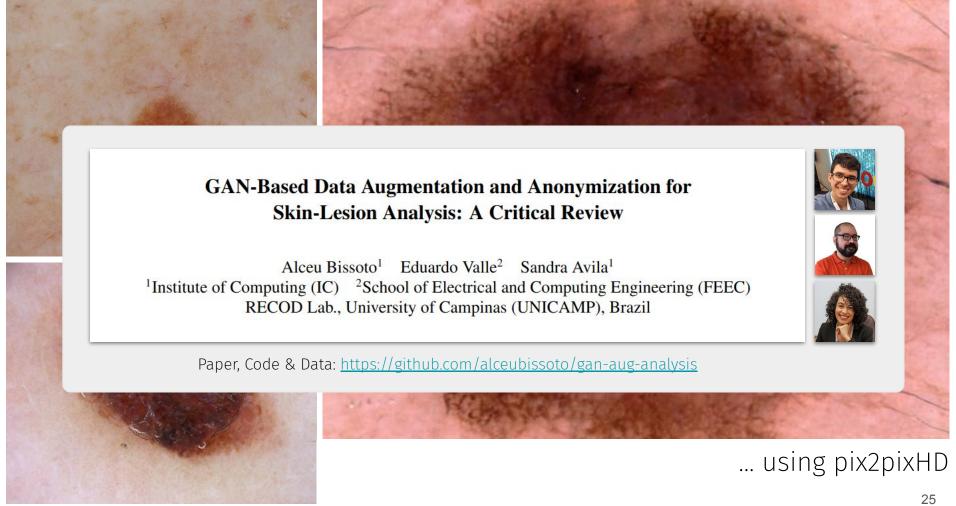




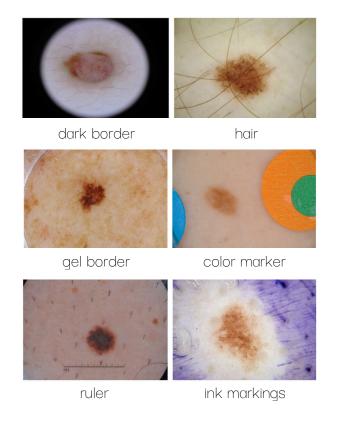
Paper, Code & Data: <a href="https://github.com/alceubissoto/gan-skin-lesion">https://github.com/alceubissoto/gan-skin-lesion</a>



A. Bissoto, F. Perez, E. Valle, S. Avila, "Skin lesion synthesis with generative adversarial networks", ISIC Workshop, MICCAI, 2018.



Dataset **biases** may **inflate** the performance of models!



#### (De)Constructing Bias on Skin Lesion Datasets

Alceu Bissoto<sup>1</sup> Michel Fornaciali<sup>2</sup> Eduardo Valle<sup>2</sup> Sandra Avila<sup>1</sup>

<sup>1</sup>Institute of Computing (IC) <sup>2</sup>School of Electrical and Computing Engineering (FEEC)

RECOD Lab., University of Campinas (UNICAMP), Brazil

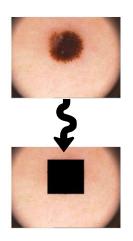




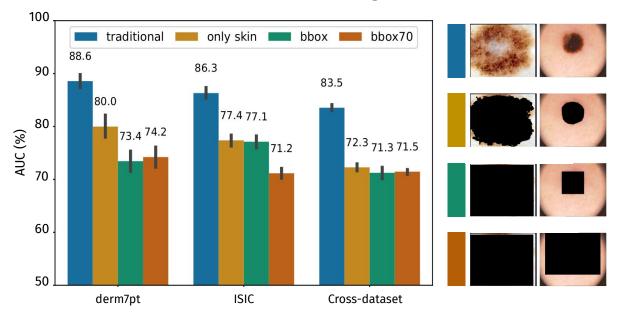
Paper, Code & Data: <a href="https://github.com/alceubissoto/deconstructing-bias-skin-lesion">https://github.com/alceubissoto/deconstructing-bias-skin-lesion</a>

# Is it possible to be completely bias free?

But, if we destroy the "important" information in the data?



Performance<sup>1</sup> of **157** dermatologists on ISIC: **67% AUC** 



#### **Debiasing Skin Lesion Datasets and Models? Not So Fast**

Alceu Bissoto<sup>1</sup> Eduardo Valle<sup>2</sup> Sandra Avila<sup>1</sup>

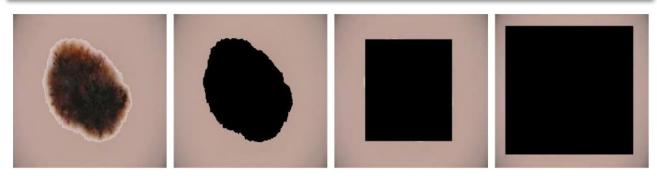
<sup>1</sup>Institute of Computing (IC) <sup>2</sup>School of Electrical and Computing Engineering (FEEC)

RECOD Lab., University of Campinas (UNICAMP), Brazil



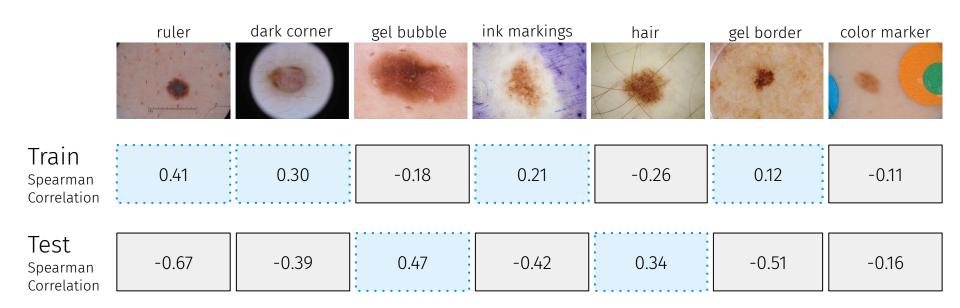






Paper, Code & Data: <a href="https://github.com/alceubissoto/debiasing-skin">https://github.com/alceubissoto/debiasing-skin</a>

# Trap Sets



# AN EVALUATION OF SELF-SUPERVISED PRE-TRAINING FOR SKIN-LESION ANALYSIS

#### A PREPRINT

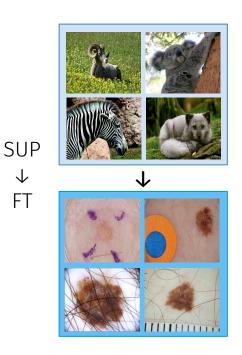
Levy Chaves<sup>1</sup> Alceu Bissoto<sup>1</sup> Eduardo Valle<sup>2</sup> Sandra Avila<sup>1</sup>

Institute of Computing (IC) <sup>2</sup>School of Electrical and Computing Engineering (FEEC)

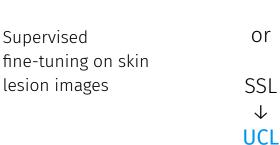
RECOD Lab., University of Campinas (UNICAMP), Brazil



Paper, Code & Data: <a href="https://github.com/VirtualSpaceman/ssl-skin-lesions">https://github.com/VirtualSpaceman/ssl-skin-lesions</a>



Supervised training on labeled natural images





SSL

FT

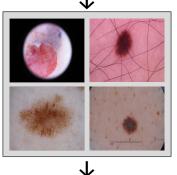
or

SCL

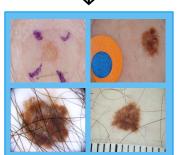
 $\downarrow$ 

FT

Self-supervised pre-training on unlabeled natural images



Self-supervised learning on unlabeled or labeled skin lesion images



Supervised fine-tuning on skin lesion images

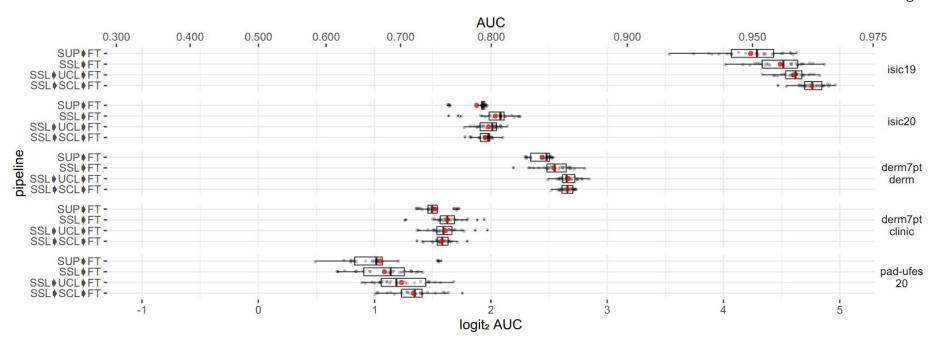
SSL: Self-Supervised Learning

UCL/SCL: Unsupervised/Supervised Contrastive Learning

FT: Fine-tuning

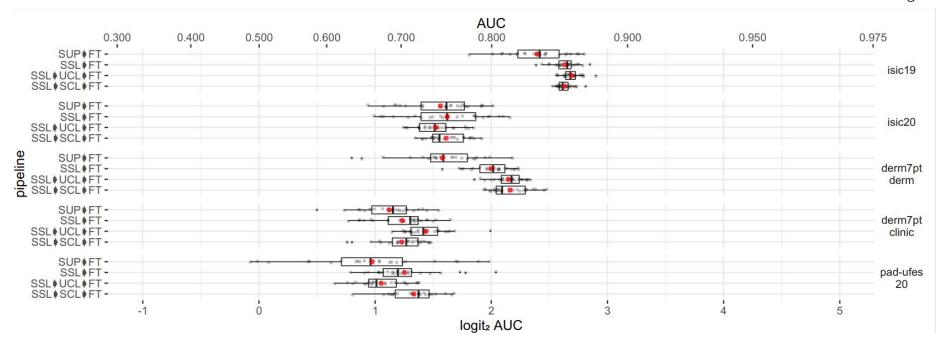
# 100% of labeled images

### Training on the ISIC 2019 dataset (~15,000 images)



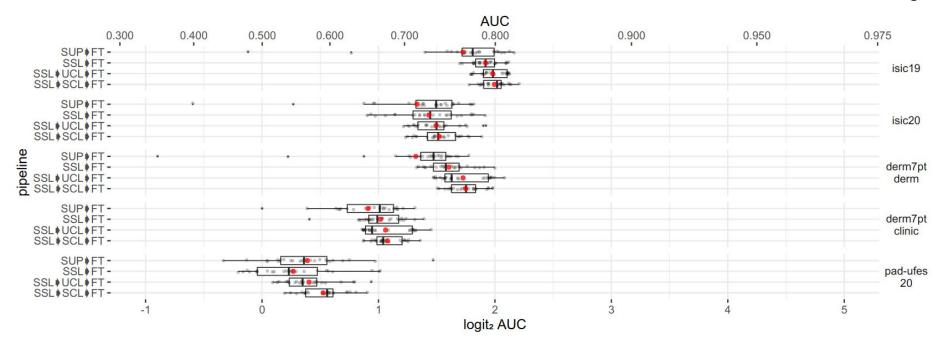
# 10% of labeled images

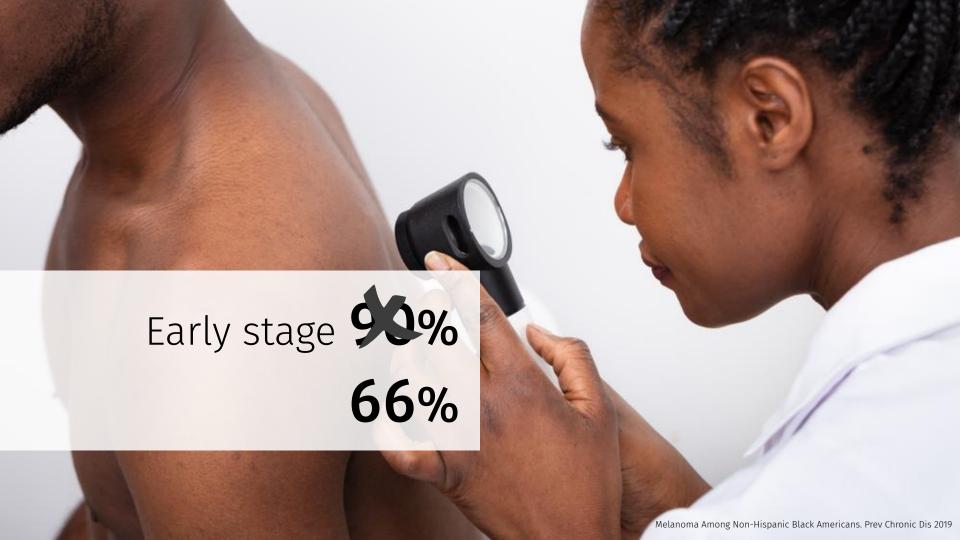
### Training on the ISIC 2019 dataset (~1500 images)



# 1% of labeled images

### Training on the ISIC 2019 dataset (~150 images)





#### The New York Times

### Dermatology Has a Problem With Skin Color

Common conditions often manifest differently on dark skin. Yet physicians are trained mostly to diagnose them on white skin.



By Roni Caryn Rabin

Aug. 30, 2020



Dr. Jenna Lester, director of the Skin of Color Program at University of California, San Francisco.

# Fairness of Classifiers Across Skin Tones in Dermatology

Newton M. Kinyanjui<sup>1,4</sup>, Timothy Odonga<sup>1,4</sup>, Celia Cintas<sup>1</sup>, Noel C. F. Codella<sup>2</sup>, Rameswar Panda<sup>3</sup>, Prasanna Sattigeri<sup>2</sup>, and Kush R. Varshney<sup>1,2</sup>

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<sup>2</sup> IBM Research – T. J. Watson Research Center, Yorktown Heights, NY 10598, USA
 <sup>3</sup> IBM Research – Cambridge, Cambridge, MA 02142, USA
 <sup>4</sup> Carnegie Mellon University Africa, Kigali, Rwanda

MICCAI 2020

