Towards Fairness & Robustness in Machine Learning for Dermatology

Skin-tone representation disparities in dermatology datasets for machine learning applications

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The team

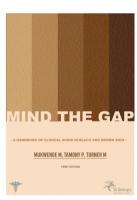




Disparities in Dermatology

- In African American population, melanoma is often diagnosed at an advanced stage with deeper tumors [MSL⁺17, WEK⁺11].
- 5 year survival rates for acral lentiginous melanoma (ALM) is 82.6% in caucasian population, but only 77.2% in african american patients. [MCH15].
- The paucity of images of skin manifestations of COVID-19 in patients with darker skin is a problem, because it may make identification of COVID-19 presenting with cutaneous manifestations more difficult for both dermatologists and the public. [LJZ⁺20]
- Dermatologists started an international registry to catalog examples of skin manifestations of Covid-19. The registry compiled more than 700 cases, but only 34 of disorders in Hispanic and 13 in Black patients were submitted. [Rab20]





The cover of the "Mind the Gap" handbook, written by Malone Mukwende, with two of his lecturers, Peter Tamony and Margot Turner.

How these disparities are reflected in Healthcare Machine Learning models?

- Are standard dermatology image datasets used in ML tasks biased with respect to skin tone? Can we quantify this?
- ² Are ML models **robust against changes** in the clinical setting or unknown diseases samples?

Machine Learning & Dermatology

Skin disease diagnosis using machine learning

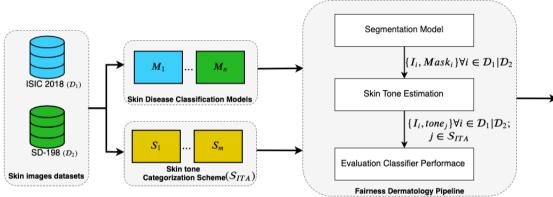
- Benchmark model for melanoma diagnosis outperforms trained dermatologists [CNP⁺16]
- 2 ISIC challenges (https://www.isic-archive.com/)

 Predictive inequity in computer vision with respect to skin type

- 1 Automated face image analysis for gender classification [BG18]
- 2 Pedestrian detection systems [WHM19]
- Out-of-distribution detection in dermatology [AYAG19, GNS⁺19, ZZL19, CHP⁺ss, PAT19, PST⁺20]



Overview : Proposed Framework



Kinyanjui, et al. "Estimating skin tone and effects on classification performance in dermatology datasets."MICCAI 2020.



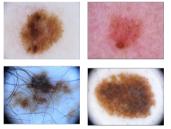
Skin Disease Graphical Content Warning

Note that we will show **skin disease examples** that could be **sensitive** or **triggering** to some viewers. We notice this, so viewers can prepare themselves to adequately engage or, if necessary, disengage for their own well-being.

Datasets

ISIC 2018

- 10015 dermoscopic images
- 7 disease classes
- 2594 images with ground truth segmentation masks for diseased area



SD-198

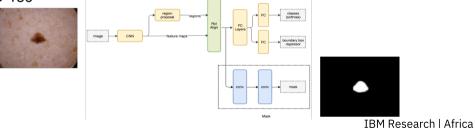
- 6548 clinical images
- 198 disease classes
- No segmentation data



Segmentation to Obtain Non-Diseased Region

Finetune Mask R-CNN model ([HGDG17])

- Adjust pretrained classifier with a FastRCNNPredictor with 2 classes (background and mask)
- Adjust mask predictor with new MaskRCNNPredictor with 2 classes and 512 hidden neurons
- Further apply thresholding techniques on predicted grayscale mask including contour extraction for ISIC2018 and grid search for optimal binary thresholding for SD-136

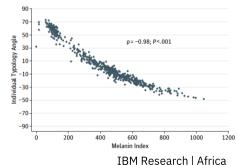


Skin Tone Metric of Non-Diseased Region

- 1 Given non-diseased pixels, characterize them with a skin tone metric
 - 1 Use individual typology angle (ITA) [WWdPR15], Highly correlated with melanin index
 - 2 ITA = $\tan^{-1} \left(\frac{L-50}{b}\right) \times \frac{180}{\pi}$ Where *L* is luminance and *b* quantifies amount of yellow.
 - 3 Use pixels with *L* and *b* values within 1 standard deviation to deal with outliers.
- 2 Bin into categories [CSD+15]

ITA Range	Skin Tone Category	Abbreviation
$ITA>55^\circ$	Very Light	very_lt
$48^\circ < ITA \le 55^\circ$	Light 2	lt2
$41^\circ < ITA \le 48^\circ$	Light 1	lt1
$34.5^\circ < ITA \le 41^\circ$	Intermediate 2	int2
28° < ITA \leq 34.5 $^\circ$	Intermediate 1	int1
$19^\circ < ITA \le 28^\circ$	Tanned 2	tan2
$10^\circ<{ m ITA}\le19^\circ$	Tanned 1	tan1
$ITA \leq 10^\circ$	Dark	dark

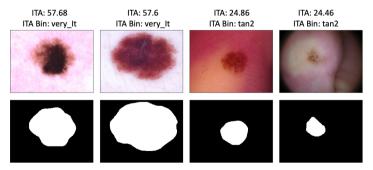
Figure from [WWdPR15].



Results

Metrics for segmentation on ISIC 2018

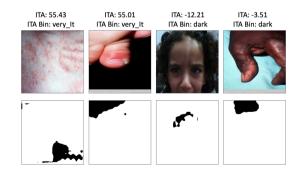
The Mask R-CNN model yields an accuracy of 0.956, a false negative rate of 0.024, and a mean absolute error in ITA computation of 0.428 degrees. [KOC⁺19]



Results (Cont.)

Metrics for segmentation on SD-136

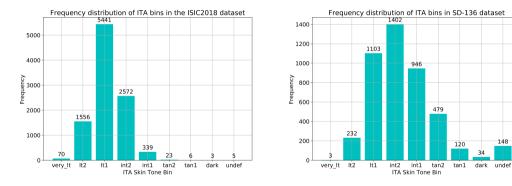
The segmentation model on the SD-136 dataset yield an accuracy of 0.802, a false negative rate of 0.076, and a mean absolute error in ITA computation of 3.572 degrees. [KOC⁺19]



Results (Cont.)

Skin Tone Distribution

There is underrepresentation of darker skin tones in both datasets



How these disparities are reflected in Healthcare Machine Learning models?

- Are standard dermatology image datasets used in ML tasks biased with respect to skin tone? Can we quantify this?
- 2 Are ML models robust against changes in the clinical setting or unknown diseases samples?

OOD for Skin Disease Classifiers

Recent advances in deep learning have led to breakthroughs in the development of automated skin disease classification. As we observe an **increasing interest** in these models in the **dermatology space**, it is crucial to address aspects such as the **robustness** and fairness of these solutions.

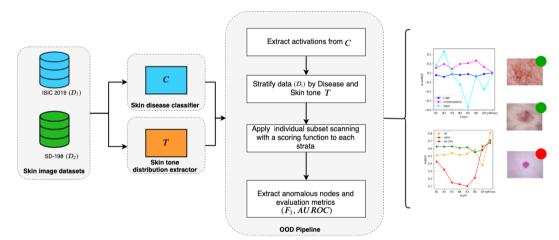
We validated our approach in two use cases:

- 1 Different clinical settings.
- 2 Unknown disease classes.



Example images from unknown disease case (top) and clinical setting changes (bottom). IBM Research | Africa

Overview: Proposed Approach



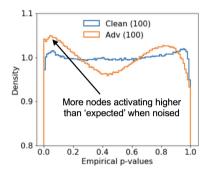
Out-of-Distribution Detection in Dermatology using Input Perturbation and Subset Scanning [KTC⁺21] IBM Research | Africa

Treat Neural Networks as data-generating systems and apply anomalous pattern detection methods to activation data.

Subset Scanning efficiently searches over a large combinatorial space in order to find groups of records that differ the most from 'expected' behavior.

Some goodies about this type of approach:

- 1 We can provide detection improvements at run time.
- 2 We can abstract from domains and focus only on the deep representation of the input.
- No need to re-train or have labeled examples of the anomalies.
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Assumption

Activations from abnormal images have a different distribution of p-values than normal samples.

p-value is the proportion of background activations (H_0), drawn from the same node for several clean samples, greater than the activation from a test sample.

$$\max_{\alpha} \varphi(\alpha, N_{\alpha}, N) = \frac{N_{\alpha} - N\alpha}{\sqrt{N}} \quad (1)$$

Where N_{α} is the number of p-values less than α N is the number of p-values α is the level of significance φ is a scoring function

How we score a test sample?

Scoring functions operate on a test sample in order to measure how much the p-values deviate from uniform.

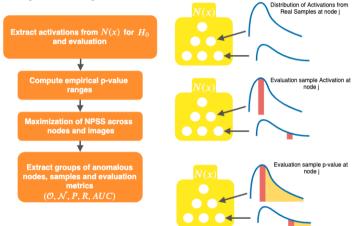
NPSS maximization

Scoring functions may be viewed as set functions that operate on subsets of nodes. We search for the highest scoring subset of nodes that maximize the deviance from uniform.

$$F(S) = \max_{\alpha} F_{\alpha}(S) = \max_{\alpha} \varphi(\alpha, N_{\alpha}(S), N(S))$$
(2)

Group vs. Individual Scanning

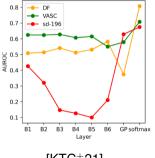
For group-based scanning our search space is: $S = X_{\hat{S}} \times O_{\hat{S}}$, where $X_{\hat{S}}$ is a subset of test samples and $O_{\hat{S}}$ is a subset of nodes' activations. For individual scanning we work with only one X_i .



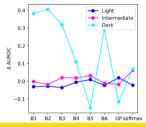
Preliminary results

Results across settings

The layers for detecting new class are different from the ones for OOD



[KTC+21]



Fairness of OOD detectors

We see varying performances for samples of Dark skin tones. This instability of performance for samples of Dark skin tones may be partially because network is trained on the ISIC 2019 dataset that **heavily lacks** samples of **Dark skin tones**.

Conclusions and future work

- The two skin disease datasets are biased towards lighter skin with majority of the samples between ITA values [34.5°, 48°].
- We can provide a single OOD detection for multiple scenarios (clinical setting change or unknown disease)
- Implementation of better segmentation models for clinical images for all skin tones.
- **4** Experiments around stratification of skin tone by disease.
- 5 How a fair distribution looks like in this case?

Other interesting work at the Kenya Lab

1 Subset Scanning

- Cintas, C., Speakman, S., Akinwande, V., Ogallo, W., Weldemariam, K., Sridharan, S. and McFowland, E. Detecting Adversarial Attacks via Subset Scanning of Autoencoder Activations and Reconstruction Error. International Joint Conference on Artificial Intelligence (IJCAI) 2020.
- Cintas, C., Das, P., Quanz, B., Speakman, S., Akinwande, V. and Chen, P.Y., 2021.
 Towards creativity characterization of generative models via group-based subset scanning. In Synthetic Data Generation Workshop at ICLR 2021.
- 2 ML in Healthcare
 - Tadesse et al. Unsupervised Discovery of Subgroups with Anomalous Maternal and Neonatal Outcomes with WHO's Safe Childbirth Checklist as Intervention. NeurIPS Workshop on Machine Learning for Public Health (Best Paper Award), December 2020.
 - Speakman et al. Automatic Stratification of Tabular Health Data. American Medical Informatics Association Annual Symposium (AMIA) 2021.

Asante, Thanks, Gracias!



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- Hannah Kim, Girmaw Abebe Tadesse, Celia Cintas, Skyler Speakman, and Kush Varshney, *Out-of-distribution detection in dermatology using input perturbation and subset scanning*, arXiv preprint arXiv:2105.11160 (2021).
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