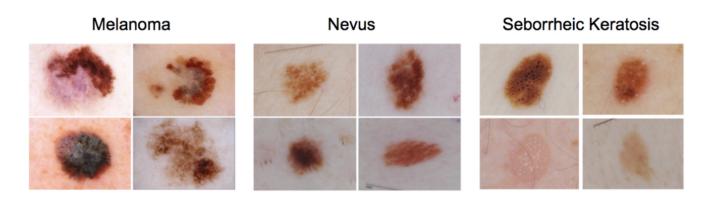
Mini Project: Dermatologist Al

Introduction

In this mini project, you will design an algorithm that can visually diagnose **melanoma**, the deadliest form of skin cancer. In particular, your algorithm will distinguish this malignant skin tumor from two types of benign lesions (**nevi** and **seborrheic keratoses**).

The data and objective are pulled from the 2017 ISIC Challenge on Skin Lesion Analysis Towards Melanoma Detection. As part of the challenge, participants were tasked to design an algorithm to diagnose skin lesion images as one of three different skin diseases (melanoma, nevus, or seborrheic keratosis). In this project, you will create a model to generate your own predictions.



Getting Started

1. Clone the **repository** and create a data/ folder to hold the dataset of skin images.

```
git clone https://github.com/udacity/dermatologist-ai.git
mkdir data; cd data
```

2. Create folders to hold the training, validation, and test images.

```
mkdir train; mkdir valid; mkdir test
```



- 5. Download and unzip the test data (5.1 GB).
- 6. Place the training, validation, and test images in the data/ folder, at data/train/, data/valid/, and data/test/, respectively. Each folder should contain three subfolders (melanoma/, nevus/, seborrheic_keratosis/), each containing representative images from one of the three image classes.

You are free to use any coding environment of your choice to solve this mini project! In order to rank your results, you need only use a pipeline that culminates in a CSV file containing your test predictions.

Create a Model

Use the training and validation data to train a model that can distinguish between the three different image classes. (After training, you will use the test images to gauge the performance of your model.)

If you would like to read more about some of the algorithms that were successful in this competition, please read **this article** that discusses some of the best approaches. A few of the corresponding research papers appear below.

- Matsunaga K, Hamada A, Minagawa A, Koga H. "Image Classification of Melanoma, Nevus and Seborrheic Keratosis by Deep Neural Network Ensemble".
 International Skin Imaging Collaboration (ISIC) 2017 Challenge at the International Symposium on Biomedical Imaging (ISBI).
- Daz IG. "Incorporating the Knowledge of Dermatologists to Convolutional Neural Networks for the Diagnosis of Skin Lesions". International Skin Imaging
 Collaboration (ISIC) 2017 Challenge at the International Symposium on Biomedical Imaging (ISBI). (github)
- Menegola A, Tavares J, Fornaciali M, Li LT, Avila S, Valle E. "RECOD Titans at ISIC
 Challenge 2017". International Skin Imaging Collaboration (ISIC) 2017 Challenge at the International Symposium on Biomedical Imaging (ISBI). (github)

While the original challenge provided additional data (such as the gender and age of the patients), we only provide the image data to you. If you would like to download this additional patient data, you may do so at the competition website.



begin with the ISIC Archive.

Evaluation

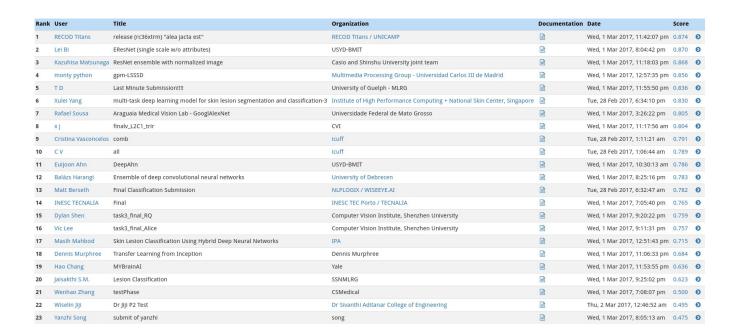
Inspired by the ISIC challenge, your algorithm will be ranked according to three separate categories.

Category 1: ROC AUC for Melanoma Classification

In the first category, we will gauge the ability of your CNN to distinguish between malignant melanoma and the benign skin lesions (nevus, seborrheic keratosis) by calculating the area under the receiver operating characteristic curve (ROC AUC) corresponding to this binary classification task.

If you are unfamiliar with ROC (Receiver Operating Characteristic) curves and would like to learn more, you can check out the documentation in **scikit-learn** or read **this Wikipedia article**.

The top scores (from the ISIC competition) in this category can be found in the image below.



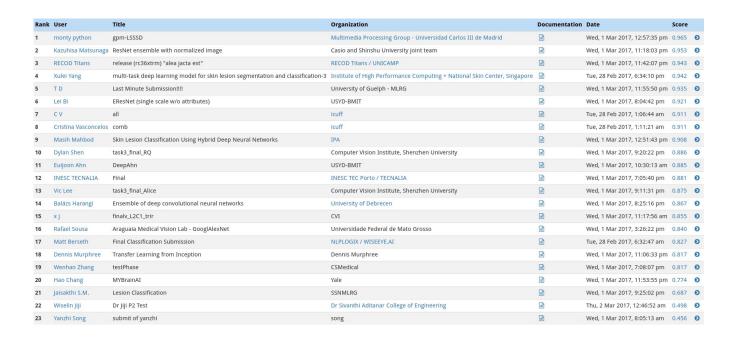
Category 2: ROC AUC for Melanocytic Classification

All of the skin lesions that we will examine are caused by abnormal growth of either melanocytes or keratinocytes, which are two different types of epidermal skin cells.



In the second category, we will test the ability of your CNN to distinuish between melanocytic and keratinocytic skin lesions by calculating the area under the receiver operating characteristic curve (ROC AUC) corresponding to this binary classification task.

The top scores in this category (from the ISIC competition) can be found in the image below.



Category 3: Mean ROC AUC

In the third category, we will take the average of the ROC AUC values from the first two categories.

The top scores in this category (from the ISIC competition) can be found in the image below.



4	роріеуі .	ERESNET (Single scale w/o attributes)	O2AD-RWI1		wed, 1 Mar 2017, 7:04:42 pm	0.896	O
5	Xulei Yang	multi-task deep learning model for skin lesion segmentation and classification-3	Institute of High Performance Computing + National Skin Center, Singapore	a	Tue, 28 Feb 2017, 5:34:10 pm	0.886	0
6	T D	Last Minute Submission!!!!	University of Guelph - MLRG		Wed, 1 Mar 2017, 10:55:50 pm	0.886	0
7	Cristina Vasconcelos	comb	icuff		Tue, 28 Feb 2017, 12:11:21 am	0.851	0
8	Cristina Vasconcelos	all	icuff	a	Tue, 28 Feb 2017, 12:06:44 am	0.850	Ð
9	Euijoon Ahn	DeepAhn	USYD-BMIT		Wed, 1 Mar 2017, 9:30:13 am	0.836	0
10	хj	finalv_L2C1_trir	CVI		Wed, 1 Mar 2017, 10:17:56 am	0.829	Ð
11	Balázs Harangi	Ensemble of deep convolutional neural networks	University of Debrecen		Wed, 1 Mar 2017, 7:25:16 pm	0.825	0
12	INESC TECNALIA	Final	INESC TEC Porto / TECNALIA		Wed, 1 Mar 2017, 6:05:40 pm	0.823	0
13	Rafael Sousa	Araguaia Medical Vision Lab - GooglAlexNet	Universidade Federal de Mato Grosso		Wed, 1 Mar 2017, 2:26:22 pm	0.823	Ð
14	Dylan Shen	task3_final_RQ	Computer Vision Institute, Shenzhen University		Wed, 1 Mar 2017, 8:20:22 pm	0.823	0
15	Vic Lee	task3_final_Alice	Computer Vision Institute, Shenzhen University		Wed, 1 Mar 2017, 8:11:31 pm	0.816	0
16	Masih Mahbod	Skin Lesion Classification Using Hybrid Deep Neural Networks	IPA		Wed, 1 Mar 2017, 11:51:43 am	0.811	0
17	Matt Berseth	Final Classification Submission	NLPLOGIX / WISEEYE.AI		Tue, 28 Feb 2017, 5:32:47 am	0.804	0
18	Dennis Murphree	Transfer Learning from Inception	Dennis Murphree		Wed, 1 Mar 2017, 10:06:33 pm	0.750	0
19	Hao Chang	MYBrainAI	Yale		Wed, 1 Mar 2017, 10:53:55 pm	0.705	0
20	Wenhao Zhang	testPhase	CSMedical		Wed, 1 Mar 2017, 6:08:07 pm	0.658	0
21	Jaisakthi S.M.	Lesion Classification	SSNMLRG		Wed, 1 Mar 2017, 8:25:02 pm	0.655	0
22	Wiselin Jiji	Dr Jiji P2 Test	Dr Sivanthi Aditanar College of Engineering		Wed, 1 Mar 2017, 11:46:52 pm	0.497	0
23	Yanzhi Song	submit of yanzhi	song		Wed, 1 Mar 2017, 7:05:13 am	0.465	0

Getting your Results

Once you have trained your model, create a CSV file to store your test predictions. Your file should have exactly 600 rows, each corresponding to a different test image, **plus** a header row. You can find an example submission file (sample_submission.csv) in the repository.

Your file should have exactly 3 columns:

- Id the file names of the test images (in the same order as the sample submission file)
- task_1 the model's predicted probability that the image (at the path in Id) depicts melanoma
- task_2 the model's predicted probability that the image (at the path in Id) depicts seborrheic keratosis

Once the CSV file is obtained, you will use the get_results.py file to score your submission. To set up the environment to run this file, you need to create (and activate) an environment with Python 3.5 and a few pip-installable packages:

```
conda create --name derm-ai python=3.5
source activate derm-ai
pip install -r requirements.txt
```

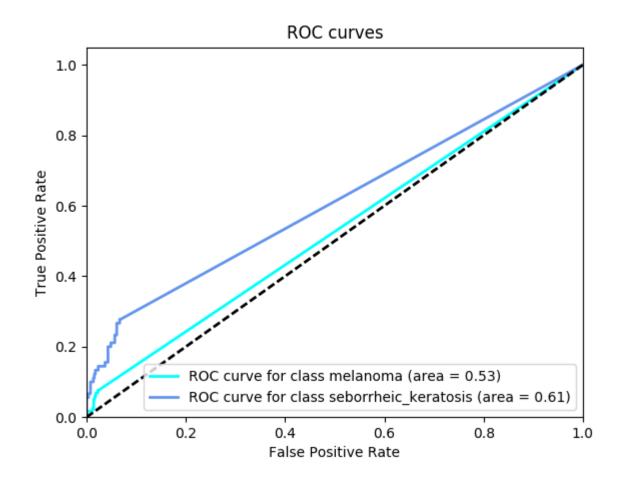


python get_results.py sample_predictions.csv

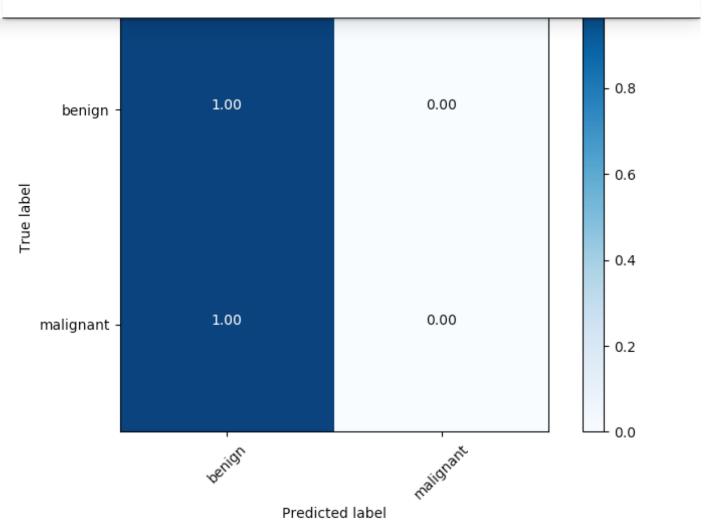
Check the terminal output for the scores obtained in the three categories:

Category 1 Score: 0.526 Category 2 Score: 0.606 Category 3 Score: 0.566

The corresponding **ROC curves** appear in a pop-up window, along with the **confusion matrix** corresponding to melanoma classification.







As you can see from the confusion matrix, the sample submission currently predicts that most of the images in the test dataset correspond to benign lesions. Let's see if your model can improve these results, towards better detecting cancer!

The code for generating the confusion matrix assumes that the threshold for classifying melanoma is set to 0.5. To change this threshold, you need only supply an additional command-line argument when calling the get_results.py file. For instance, to set the threshold at 0.4, you need only run:

```
python get_results.py sample_predictions.csv 0.4
```

To test **your own** submission, change the code to instead include the path to **your** CSV file.

