

## Classify AI

**Classify AI is a software package designed to train, optimise and evaluate machine learning classifiers for automated reading of HIV Rapid Diagnostic Tests (RDTs). It compares the performance of 4 different approaches (MobileNet\_v2, MobileNet\_v3, ResNet50 and SVM), using a dataset of images captured in the field in rural South Africa.**

### Product Specification

Classify AI is a series of interdependent scripts written in Python 3.6.

### Description

This repository contains the code used to produce the results presented in the following publication:

Deep learning of HIV field-based rapid tests DOI:10.1038/s41591-021-01384-9

The work presented is the result of a collaboration between i-sense (<https://www.i-sense.org.uk/>) and the Africa Health Research Institute (AHRI, <https://www.ahri.org/>).

The datasets are available upon request, please see details in the publication.

The repository is divided in 2 subfolders corresponding to Figures 3 and 4 of the publication, respectively.

In Figure 3, the others used a dataset of 11, 374 images of HIV RDTs captured in the field, around AHRI. They trained and optimised classifiers, performed cross-validation to check for reproducibility. They produced 2 optimal classifiers, one for each type of HIV RDT used in the study.

In Figure 4, the authors conducted a pilot study to assess 1) the level of agreement between participants when interpreting HIV RDTs by eye; 2) the performance of their optimal classifiers (one for each type of HIV RDT used in the study) compared to traditional visual interpretation by humans.

### References

1. V Turbé, C Herbst, T Mngomezulu, S Meshkinfamfard, N Dlamini, T Mhlongo, T Smit, V Cherepanova, K Shimada, J Budd, N Arsenov, S Gray, D Pillay, K Herbst, M Shahmanesh & R A. McKendry (17 June 2021), <https://doi.org/10.1038/s41591-021-01384-9>, <https://www.nature.com/articles/s41591-021-01384-9>

### Category

Software/Bioinformatics

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