Classifying Breast Cytological Images using Deep Learning Architectures

Hasnae Zerouaoui¹¹^a and Ali Idri^{1,2}^b

¹Modeling, Simulation and Data Analysis, Mohammed VI Polytechnic University, Benguerir, Morocco ²Software Project Management Research Team, ENSIAS, Mohammed V University in Rabat, Morocco

- Keywords: Computer-aided Diagnosis, Breast Cancer, Classification, Deep Convolutional Neural Networks, Image Processing, Histological Images.
- Abstract: Breast cancer (BC) is a leading cause of death among women worldwide. It remains a critical challenge, causing over 10 million deaths globally in 2020. Medical images analysis is the most promising research area since it provides facilities for diagnosing several diseases such as breast cancer. The present paper carries out an empirical evaluation of recent deep Convolutional Neural Network (CNN) architectures for a binary classification of breast cytological images based fined tuned versions of seven deep learning techniques: VGG16, VGG19, DenseNet201, InceptionResNetV2, InceptionV3, ResNet50 and MobileNetV2. The empirical evaluations used: (1) four classification performance criteria (accuracy, recall, precision and F1-score), (2) Scott Knott (SK) statistical test to select the best cluster of the outperforming architectures, and (3) borda count voting system to rank the best performing architectures. All the evaluations were over the FNAC dataset which contain 212 images. Results showed the potential of deep learning techniques to classify breast cancer in malignant and benign, therefor the findings of this study recommend the use of MobileNetV2 for the classification of the breast cancer cytological images since it gave the best results with an accuracy of 98.54%.

1 INTRODUCTION

Breast cancer (BC) is still the leading cause of death among women worldwide (Metelko et al., 1995). It remains a global challenge, causing over 1 million deaths globally in 2019 (Bish et al., 2005). As the number of patients infected by this disease increases, it turns out to be increasingly hard for radiologists to accurately deal with the diagnosis process in the constrained accessible time (Zhang et al., 2011). Medical images analysis is one of the most promising research areas, it provides facilities for diagnosis and making decisions of several diseases such as breast cancer. Recently, more attention are paid to imaging modalities and Deep Learning (DL) in BC (Mendelson and Eb, 2019). Therefore, interpretation of these images requires expertise and consequently several algorithms have been developed and evaluated to improve and help oncologist's diagnosis.

In general, DL showed better performance in breast cancer detection, and provided high accurate classifications compared with classical Machine Learning (ML) techniques (Saha, Mukherjee and Chakraborty, 2016) (Sadoughi et al., 2018). For instance, the study (Saha, Mukherjee and Chakraborty, 2016) showed that the use of deep CNN architectures is very powerful and efficient in the domain of DL since it tested the InceptionRecurrent Residual CNN in the dataset BreakHis and it gave better results compared to existing techniques such as CNN and SVM. The study (Xie et al., 2019) used AlexNet and LeNet for the binary classification of the BreakHis dataset and showed an improvement of the accuracy compared to the traditional ML techniques. However, the present study develops and evaluates the performances measured in terms of accuracy, sensitivity, recall, precision and F1-score of seven of the most recent DL techniques in BC classification

Zerouaoui, H. and Idri, A.

Classifying Breast Cytological Images using Deep Learning Architectures

DOI: 10.5220/0010850000003123

In Proceedings of the 15th International Joint Conference on Biomedical Engineering Systems and Technologies (BIOSTEC 2022) - Volume 5: HEALTHINF, pages 557-564 ISBN: 978-989-758-552-4; ISSN: 2184-4305

^a https://orcid.org/0000-0001-7268-8404

^b https://orcid.org/0000-0002-4586-4158

Copyright © 2022 by SCITEPRESS - Science and Technology Publications, Lda. All rights reserved

over the FNAC dataset. To the best of our knowledge, this study is the first to evaluate and compare seven DL techniques (VGG16, VGG19, DenseNet201, InceptionResNetV2, InceptionV3, ResNet50 and MobileNetV2) using the Scott Knott (SK) statistical test and the borda count voting method in BC binary classification. Note that the SK test has been widely used to comparing, clustering and ranking multiple machine learning models for parameters tunning (Idri, Hosni and Abran, 2016; Ottoni et al., 2020) in different fields such as software engineering (Ottoni et al., 2020) and breast cancer (Idri et al., 2020). Therefore, we use the SK test since: (1) it shows high performance compared to other statistical tests such as Jollife (Jolliffe, Allen and Christie, 1989), Calinski and Corsten (Calinski and Corsten, 1985), and Cox and Spjotvoll (Worsley, 1986) and (2) its ability to select the best non-overlapping groups of machine learning techniques. Moreover, we use the Borda Count voting method (García-Lapresta and Martínez-Panero, 2002; Emerson, 2013) to rank the best SK selected techniques based on the four performance criterias.

The present study discusses two research questions (RQs):

- (RQ1): What is the overall performance of DL techniques in BC classification?
- (RQ2): Is there any DL techniques which distinctly outperform the others?

The main contributions of this empirical study are the following: (1) Designing seven DL architectures: VGG16, VGG19, DenseNet201, InceptionResNetV2, InceptionV3, ResNet50 and MobileNetV2 in BC classification, (2) Avoiding overfitting by using weight decay and L2 regularizers, (3) Comparing the performances of the seven architectures using SK clustering test and borda count voting method.

The remainder of this paper is organized as follow. Section 2 describes the related work. In Section 3, we present the configuration and parametrization of the seven DL techniques, the empirical methodology followed throughout the research, the data preparation which includes data acquisition and image processing and the abbreviations. Section 4 presents and discusses the empirical results. Section 5 outlines conclusions and future works.

2 RELATED WORKS

This section presents the results and the main findings of the related work as shown in Table 1, the results are summarized as follow:

- Accuracy is the most frequently criterion used to evaluate the performance of the DL techniques in BC when using balanced datasets (Zerouaoui and Idri, 2021).
- Most of the studies only compared two to three DL techniques. Although the DL architectures used in the selected studies were different, it is worth notable that the most investigated techniques were InceptionResNet, VGG16, VGG19 and ResNet50 (Alom *et al.*, 2019; Spanhol *et al.*, 2016; Xie *et al.*, 2019)
- Some studies (Kassani *et al.*, 2019; (Nahid, Mehrabi and Kong, 2018; Zhu *et al.*, 2019) combined more than two DL techniques in order to have better

Table 1: Summary of the literature review of the use of DL techniques in BC classification.

Authors	Findings and results
Alom et	Conduct a study on the use of Inception
al.	Recurrent Residual Convolutional Neural
(Alom et	Network (IRRCNN) which is a hybrid
al.,	DCNN architecture based on RCNN,
2019)	Residual Network and Inception tested on
	two datasets: BreakHis and BC
	classification challenge 2015.
LOGY	The performance was evaluated on image
	level, patch level, image based and patch-
	based analysis. The results show an
	improvement of 3,67% and 2.14% of
	accuracy on the BreakHis dataset compared
	to scientific results since 2016
Fabio et	Investigate a deep learning approach, using
al.	two DCNN architecture which are AlexNet
(Spanhol	and LeNet, in order to avoid hand crafted
et al.,	features. The results of the experiments
2016)	demonstrated an improved accuracy
	compared to the experiments that used
	traditional feature extractor techniques.
Xie et al.	Evaluate Inception_V3 and Inception_
(Xie et	ResNet_V2 for classification of Brekhis
al.,	using two types of learning: the supervised
2019)	and the unsupervised learning. The authors
	used the transfer learning by pre training the
	model on ImageNet, applying Data
	augmentation and Fine tuning using the
	BreakHis dataset. The experiment shows
	that the results of the augmented dataset is
	much better than the normal dataset, that
	Inception_ResNet_V2 has better results for
	the feature extraction and the supervised
	learning has a better accuracy than
	clustering

Table 1: Summary of the literature review of the use of DL techniques in BC classification (cont.).

Authors	Findings and results
Al Nahid et al. (Nahid, Mehrabi and Kong, 2018)	Use three models: CNN techniques, LSTM structure and the combination of CNN and LSTM on the BreakHis Dataset. As results 91% of accuracy was obtained using 200x images dataset, 96 of precision with 40x images dataset, and the best F-Measure was obtained using 40x and 100x datasets.
Kassani et al. (Kassani <i>et al.</i> , 2019)	Use of a combination of CNNs deep learning architectures such as VGG19, MobileNet and Densenet for feature extraction on 4 histopathological dataset images: ICIAR, BreakHis, Densenet and MobilNet. Then they compared the results of the classification on traditional single models and machine learning techniques namely Decision tree, Random Forest, XGBoost, AdaBoost and bagging classifier. The main finding is that the proposed ensemble method gives better results thant the solo methods with an accuracy of: 98.13%,
Chuang et al. (Zhu <i>et</i> <i>al.</i> , 2019)	Propose a hybrid architecture where they assembled multiple CNN architectures (Inception module, Residual Net and Batch Normalization techniques) and tested it on two datasets (BreakHis and BACH). The proposed model shows a comparable and better performance.
Jiang et al. (Jiang <i>et al.</i> , 2019)	Design a new CNN that includes a convolutional layer, a small SE-ResNet module and a fully connected layer, they tested their architecture on the BreakHis dataset and the results achieved 98,87% and 99,34% for the binary class and 90.66% and 93.81% for the multi class.

prediction results, since the ensemble methods in general outperformed their single techniques.

3 EXPERIMENT CONFIGURATION

This section presents the parameter tuning of the DL models, the empirical design, the data preparation followed and finally the abbreviations.

3.1 Experiment Configuration

Toward an automatic binary BC classification based on publicly available image FNAC dataset, the different DL architectures have been implemented using several parameters tuning experiments. All the images of the FNAC dataset were resized to 224x224 pixels except those of InceptionV3 and InceptionResNetV2 models that were resized to 299x299 since it is the default input size in their architectures. To train the models, we used the transfer learning technique where we downloaded the seven DL techniques pre-trained in the ImageNet dataset (Fei-Fei, Deng and Li, 2010). For the parameter tunning, we set the batch size to 32 and the number of epochs to 300. As for the optimization, we used Adam (adaptive moment estimation) (Kingma and Ba, 2015) with β 1=0.9, β 2=0.999, and an initial learning rate set to 0.0001 and decrease exponentially to 0.000001. Moreover, we used weight decay and L2- regularizers to reduce the overfitting for different models. A fully connected layer was trained with the ReLU, followed by a dropout layer with a probability of 0.5. We updated the last dense layer in all models to output two classes corresponding to benign and malignant instead of 1000 classes as was used for ImageNet.

3.2 Empirical Design

Figure 1 shows the methodology followed to carry out all the empirical evaluations. It consists of three steps we describe hereafter. Note that similar methodologies were used in (Worsley, 2009)(SHARMA *et al.*, 2003)(Azzeh, Nassif and Minku, 2015)(Idri, Abnane and Abran, 2018)(Zerouaoui *et al.*, 2021)(Idri and Abnane, 2017).



Figure 1: Experimental process.

3.3 Data Preparation

This section presents the data preparation process followed for the FNAC as described in Figure 2, which consists of Data pre-processing by using intensity normalization and Contrast Limited Adaptive Histogram Equalization (CLAHE) and Data augmentation. The Images of the FNAC dataset were captured by us using Leica ICC50 HD microscope using 400 resolution and 24 bits color depth and with 5 megapixels camera associated with the microscope(Saikia et al., 2019). Digitized images captured were then reviewed by experienced certified cyto-pathologists and selected a total of 212 images (113 Malignant and 99 Benign). The database can be downloaded from the link in (Saikia et al., 2019).



Figure 2: Data preparation process.

Data Processing: The next stage is to pre-process input images using intensity normalization and Contrast Limited Adaptive Histogram Equalization (CLAHE). Intensity normalization is a pre-processing step in image processing applications (Kassani *et al.*, 2019). We normalized input images to the standard normal distribution using min-max normalization of Equation 1. Furthermore, before feeding input images into the proposed models, CLAHE is a necessary step to improve the contrast in images as shown in Figure 3 (Makandar and Halalli, 2015; Kharel *et al.*, 2017).



Figure 3: Original and transformed images.

$$X_{norm} = \frac{x - x_{\min}}{x_{\max} - x_{\min}} \tag{1}$$

Data Augmentation: was used for the training process after dataset pre-processing and has the goal to avoid the risk of overfitting (Perez and Wang, 2017). Moreover, the strategies we used include

geometric transforms such as rescaling, rotations, shifts, shears, zooms and flips.

3.4 Abbreviations

To assist the reader and shorten the names of the DL techniques, we use the following naming rules in the rest of this paper. We abbreviate the name of each variant of DL techniques as shown in Table 2.

Table 2: Abbreviations used for the DL techniques for the FNAC dataset.

D.L techniques with the image magnification factor	Abbreviation
Noc 1(MOOIL
VGG 16	VGG16
VGG 19	VGG19
ResNet 50	Res50
Inception V3	INV3
Inception ResNet V2	INRES
DensNet201	DENS
MobilNetV2	MOB

4 **RESULTS**

This section presents and discusses the results of the empirical evaluations of seven DL techniques: ResNet50. InceptionV3, VGG16, VGG19, InceptionResNetV2, DenseNet201, and MobileNetV2, over the FNAC dataset. The performances of the DL techniques were evaluated using 5-fold cross validation and four criteria's: accuracy, recall, precision and F1-score. First the performance is compared in terms of accuracy of each DL technique (RQ1). Thereafter, we use the SK statistical test to cluster the selected DL techniques, and borda count to rank the DL techniques belonging to the best SK cluster (RQ2).

4.1 Accuracy Evaluation and Comparison of the Seven DL Techniques

This section compares the accuracy values of the seven DL techniques to each other over the FNAC dataset. Note that the training and testing of the DL techniques are implemented in Python using Keras and Tensorflow DL frameworks and run on a TPU processing unit of 8 cores with 35 GB in RAM and Linux-based OS, provided by google in Colab Notebook.

Figure 4 and Table 3 show the accuracy values of the baseline VGG16, VGG19, DenseNet201, InceptionResNetV2, InceptionV3, ResNet50 and

MobileNetV2 vs the number of epochs over the FNAC dataset. We observe that the best accuracy values were achieved with the number of epochs 25 using VGG16, VGG19, InceptionV3, DenseNet201, InceptionResNetV2 and MobileNetV2; and with the number of epochs 100 when using ResNet 50 since. The best accuracy value was achieved by MobileNet V2 (98.54%) followed by DenseNet201, VGG16, VGG19, InceptionV3 and InceptionResNetV2 with an accuracy value of 98.07%, 97.65%, 95.72%, 93.51% and 91.34% respectively. The worst accuracy value was achieved using ResNe 50 with a value of 91.34%



Figure 4: Accuracy values vs number of epochs of the seven deep learning architectures over the FNAC dataset (The abbreviation used in Figure 3 for the DL techniques for the FNAC dataset are defined in Table 4 of section 6.5).

Table 3: Accuracy values of the seven deep learning architectures over the FNAC dataset (The abbreviation used in Table 3 for the DL techniques for the FNAC dataset are defined in Table 1 of section 3.4).

DL technique	Accuracy (%)
VGG16	97.65
VGG19	95.72
INV3	95.06
Res50	91.34
INRES	93.51
DENS	98.07
MOB	98.54

4.2 Clustering DL Techniques using SK Test and Ranking Them using Borda Count

This step uses the SK statistical test to evaluate the predictive capabilities of the DL techniques evaluated in step 4.1 and discusses the ranking results when applying the borda count voting method based on accuracy, recall, precision and F1-score on the best SK clusters. Table 4 shows the values of the four performance measures of all the DL techniques over

FNAC dataset. Note that the SK test consists of grouping DL techniques with no significant difference between their accuracy values. Since the SK test requires that its inputs should be normally distributed we verified the normality of the data by the Kolmogorov-Smirnov test, and since the data is normally distributed we didn't use the Box Cox transformation (Sakia, 2012). Afterwards, we performed the SK test to cluster the selected DL techniques into overlapping free groups and identified the best group based on accuracy. The DL techniques belonging to the same group have similar predictive capability and the best group contains the DL techniques that have the highest value of accuracy.

Table 4: Best performance values of the DL techniques over the FNAC Dataset (The abbreviation used in Table 4 for the DL techniques for the FNAC dataset are defined in Table 1 of section 3.4).

DL	Accur acy (%)	Recall (%)	Precis ion (%)	F1 score (%)
VGG16	97.65	98.15	97.49	97.8
VGG19	95.72	94.98	96.95	95.95
INV3	95.06	94.19	96.5	95.32
Res50	91.34	91.9	91.98	91.86
INRES	93.51	94.28	93.79	93.94
DENS	98.07	98.42	98.86	98.63
MOB —	98.54	98.15	98.24	98.2

From the results of the SK test shown in Figure 5, it is noticeable that the SK test results gives 4 clusters which implies that the accuracy performances are highly influenced by the DL model used for the classification. The figure shows that the best SK cluster contains 3 DL models including DenseNet201, MobileNetV2 and VGG16 and that last SK cluster contains the DL model ReseNet50.



Figure 5: Results of SK test for the DL techniques over the FNAC dataset.

Table 5 shows the borda count ranking of the architectures belonging to the best SK clusters for the

FNAC dataset. The DL architecture MobileNetV2 is ranked first followed by DenseNet201 and finally VGG16.

Table	5:	Best	performance	values.
1 4010	۰.	2000	Perrormanee	

Borda Count Ranking	Deep Learning model
1	MOB
2	DENS
3	VGG16

As a summary, when using the cytological FNAC dataset, it is recommended to use the DL architecture MobileNetV2 since it was ranked first when using the borda count voting method based on accuracy, precision, recall and F1-score and achieved an accuracy of 98.54%. On top of that MobileNetV2 remains the perfect DL architecture since it is light weighted in terms of architecture and is designed for mobile and web application. Therefore, we highly recommend the use of MobileNetV2 for a binary cytological classification.

5 CONCLUSIONS

The present paper presented and discussed the results of an empirical comparative study of seven recent DL techniques (VGG16, VGG19, DenseNet201, InceptionResnetV2, InceptionV3, ResNet50 and MobileNetV2) for BC binary imaging classification. All the empirical evaluations used four performance criteria's, SK statistical test, and borda Count to assess and rank these seven DL techniques over the FNAC dataset. The findings of this study are:

(RQ1): What is the Overall Performance of DL Techniques in BC Classification?

The accuracy results of the seven DL techniques were highly influenced by the characteristics of the dataset. Nevertheless, we observed that MobileNetV2, DenseNet201, VGG16, VGG19 and InceptionV3 gave the best results. However, ReseNet50 underperformed compared to the others.

(RQ2): Is There Any DL Techniques, Which Distinctly Outperform the Others?

MobileNetV2 technique gave the best results since it belonged to the best SK clusters for the FNAC dataset and was ranked first using the borda count voting test based on accuracy, recall, precision and F1-score. As results we recommend the use of MobileNet V2 to develop DL computer assister diagnosis systems since it gives good results when using cytological images for binary BC classification.

Ongoing works investigate homogenous and heterogeneous ensembles whose members are deep learning techniques with different meta-learning techniques such as bagging, boosting and stacking for breast cancer imaging classification.

ACKNOWLEDGEMENTS

This work was conducted under the research project "Machine Learning based Breast Cancer Diagnosis and Treatment", 2020-2023. The authors would like to thank the Moroccan Ministry of Higher Education and Scientific Research, Digital Development Agency (ADD), CNRST, and UM6P for their support.

This study was funded by Mohammed VI polytechnic university at Ben Guerir Morocco.

Compliance with ethical standards.

Conflicts of interest/competing interests Not applicable.

Code availability not applicable.

REFERENCES

- Alom, M. Z. et al. (2019) 'Breast Cancer Classification from Histopathological Images with Inception Recurrent Residual Convolutional Neural Network', *Journal of Digital Imaging*. Journal of Digital Imaging. doi: 10.1007/s10278-019-00182-7.
- Azzeh, M., Nassif, A. B. and Minku, L. L. (2015) 'An empirical evaluation of ensemble adjustment methods for analogy-based effort estimation', *Journal of Systems and Software*. Elsevier Ltd., 103, pp. 36–52. doi: 10.1016/j.jss.2015.01.028.
- Bish, A. *et al.* (2005) 'Understanding why women delay in seeking help for breast cancer symptoms B', 58, pp. 321–326. doi: 10.1016/j.jpsychores.2004.10.007.
- Bony, S. *et al.* (2001) 'The relationship between mycotoxin synthesis and isolate morphology in fungal endophytes of Lolium perenne', *New Phytologist*, 152(1), pp. 125–137. doi: 10.1046/j.0028-646X.2001.00231.x.
- Calinski, T. and Corsten, L. C. A. (1985) 'Clustering Means in ANOVA by Simultaneous Testing', *Biometrics*, 41(1), p. 39. doi: 10.2307/2530641.
- Emerson, P. (2013) 'The original Borda count and partial voting', Social Choice and Welfare, 40(2), pp. 353– 358. doi: 10.1007/s00355-011-0603-9.
- Fei-Fei, L., Deng, J. and Li, K. (2010) 'ImageNet: Constructing a large-scale image database', *Journal of Vision*, 9(8), pp. 1037–1037. doi: 10.1167/9.8.1037.
- García-Lapresta, J. L. and Martínez-Panero, M. (2002) 'Borda count versus approval voting: A fuzzy

approach', Public Choice, 112(1), pp. 167-184. doi: 10.1023/A:1015609200117.

- Hamza, M. and Larocque, D. (2005) 'An empirical comparison of ensemble methods based on classification trees', *Journal of Statistical Computation* and Simulation, 75(8), pp. 629–643. doi: 10.1080/ 00949650410001729472.
- He, K. and Sun, J. (2016) 'Deep Residual Learning for Image Recognition'. doi: 10.1109/CVPR.2016.90.
- Hosni, M. et al. (2019) 'Reviewing ensemble classification methods in breast cancer', Computer Methods and Programs in Biomedicine, 177, pp. 89–112. doi: 10.1016/j.cmpb.2019.05.019.
- Huang, G. et al. (2017) 'Densely Connected Convolutional Networks'. doi: 10.1109/CVPR.2017.243.
- Idri, A. *et al.* (2020) 'Assessing the impact of parameters tuning in ensemble based breast Cancer classification', *Health and Technology*. Health and Technology, 10(5), pp. 1239–1255. doi: 10.1007/s12553-020-00453-2.
- Idri, A. and Abnane, I. (2017) 'Fuzzy Analogy Based Effort Estimation: An Empirical Comparative Study', *IEEE* CIT 2017 - 17th IEEE International Conference on Computer and Information Technology, (MI), pp. 114– 121. doi: 10.1109/CIT.2017.29.
- Idri, A., Abnane, I. and Abran, A. (2018) 'Evaluating Pred(p) and standardized accuracy criteria in software development effort estimation', *Journal of Software: Evolution and Process*, 30(4), pp. 1–15. doi: 10.1002/ smr.1925.
- Idri, A., Hosni, M. and Abran, A. (2016) 'Improved estimation of software development effort using Classical and Fuzzy Analogy ensembles', *Applied Soft Computing Journal*. Elsevier B.V., 49, pp. 990–1019. doi: 10.1016/j.asoc.2016.08.012.
- Jiang, Y. et al. (2019) 'Breast cancer histopathological image classification using convolutional neural networks with small SE-ResNet module', *PLoS ONE*, 14(3), pp. 1–21. doi: 10.1371/journal.pone.0214587.
- Jolliffe, I. T., Allen, O. B. and Christie, B. R. (1989) 'COMPARISON OF VARIETY MEANS USING advantage of this approach is that the divisions into groups can be done at more', 25, pp. 259–269.
- K, H. T. (2013) 'c r v i h o e f c r v i h o e f', 4(2), pp. 627– 635.
- Kassani, S. H. et al. (2019) 'Classification of Histopathological Biopsy Images Using Ensemble of Deep Learning Networks'. Available at: http:// arxiv.org/abs/1909.11870.
- Kharel, N. et al. (2017) 'Early diagnosis of breast cancer using contrast limited adaptive histogram equalization (CLAHE) and Morphology methods', 2017 8th International Conference on Information and Communication Systems, ICICS 2017, pp. 120–124. doi: 10.1109/IACS.2017.7921957.
- Kingma, D. P. and Ba, J. L. (2015) 'Adam: A method for stochastic optimization', 3rd International Conference on Learning Representations, ICLR 2015 - Conference Track Proceedings, pp. 1–15.
- Makandar, A. and Halalli, B. (2015) 'Breast Cancer Image Enhancement using Median Filter and CLAHE',

International Journal of Scientific & Engineering Research, 6(4), pp. 462–465. Available at: http:// www.ijser.org.

- Mendelson, E. B. and Eb, M. (2019) 'Imaging : Potentials and Limitations', *American Journal of Roentgenology*, (February), pp. 1–7. doi: 10.2214/AJR.18.20532.
- Metelko, Z. *et al.* (1995) 'Pergamon THE WORLD HEALTH ORGANIZATION QUALITY OF LIFE ASSESSMENT (WHOQOL): POSITION PAPER FROM THE WORLD HEALTH ORGANIZATION', 41(10).
- Mittas, N. and Angelis, L. (2013) 'Ranking and clustering software cost estimation models through a multiple comparisons algorithm', *IEEE Transactions on Software Engineering*, 39(4), pp. 537–551. doi: 10.1109/TSE.2012.45.
- Nahid, A. Al, Mehrabi, M. A. and Kong, Y. (2018) 'Histopathological breast cancer image classification by deep neural network techniques guided by local clustering', *BioMed Research International*, 2018. doi: 10.1155/2018/2362108.
- Ottoni, A. L. C. *et al.* (2020) 'Tuning of reinforcement learning parameters applied to SOP using the Scott– Knott method', *Soft Computing*. Springer Berlin Heidelberg, 24(6), pp. 4441–4453. doi: 10.1007/ s00500-019-04206-w.
- Perez, L. and Wang, J. (2017) 'The Effectiveness of Data Augmentation in Image Classification using Deep Learning'. Available at: http://arxiv.org/abs/ 1712.04621.
- Razzak, M. I., Naz, S. and Zaib, A. (no date) 'Deep Learning for Medical Image Processing: Overview , Challenges and the Future'.
- Sadoughi, F. et al. (2018) 'Artificial intelligence methods for the diagnosis of breast cancer by image processing: A review', Breast Cancer: Targets and Therapy, 10, pp. 219–230. doi: 10.2147/BCTT.S175311.
- Sagi, O. and Rokach, L. (2018) 'Ensemble learning: A survey', Wiley Interdisciplinary Reviews: Data Mining and Knowledge Discovery, 8(4), pp. 1–18. doi: 10.1002/widm.1249.
- Saha, M., Mukherjee, R. and Chakraborty, C. (2016) 'Computer-aided diagnosis of breast cancer using cytological images: A systematic review', *Tissue and Cell*. Elsevier Ltd, 48(5), pp. 461–474. doi: 10.1016/j.tice.2016.07.006.
- Saikia, A. R. *et al.* (2019) 'Comparative assessment of CNN architectures for classification of breast FNAC images', *Tissue and Cell.* Elsevier Ltd, 57, pp. 8–14. doi: 10.1016/j.tice.2019.02.001.
- Sakia, A. R. M. (2012) 'The Box-Cox transformation technique : a review', 41(2), pp. 169–178.
- Sandler, M. et al. (2018) 'MobileNetV2: Inverted Residuals and Linear Bottlenecks', Proceedings of the IEEE Computer Society Conference on Computer Vision and Pattern Recognition. IEEE, pp. 4510–4520. doi: 10.1109/CVPR.2018.00474.
- SHARMA, J. et al. (2003) 'Symbiotic Seed Germination and Mycorrhizae of Federally Threatened Platanthera praeclara (Orchidaceae)', The American Midland

HEALTHINF 2022 - 15th International Conference on Health Informatics

Naturalist, 149(1), pp. 104–120. doi: 10.1674/0003-0031(2003)149[0104:ssgamo]2.0.co;2.

- Simonyan, K. and Zisserman, A. (2015) 'Very deep convolutional networks for large-scale image recognition', 3rd International Conference on Learning Representations, ICLR 2015 - Conference Track Proceedings, pp. 1–14.
- Spanhol, F. A. et al. (2016) 'A Dataset for Breast Cancer Histopathological Image Classification', IEEE Transactions on Biomedical Engineering, 63(7), pp. 1455–1462. doi: 10.1109/TBME.2015.2496264.
- Szegedy, C. *et al.* (2014) 'Rethinking the Inception Architecture for Computer Vision'.
- Szegedy, C. *et al.* (no date) 'the Impact of Residual Connections on Learning', pp. 4278–4284.
- Worsley, A. K. J. (2009) 'A Non-Parametric Extension of a Cluster Analysis Method by Scott and Knott Published by: International Biometric Society Stable URL: http:// www.jstor.org/stable/2529369', 33(3), pp. 532–535.
- Worsley, K. J. (1986) 'Confidence regions and tests for a change-point in a sequence of exponential family random variables', *Biometrika*, 73(1), pp. 91–104. doi: 10.1093/biomet/73.1.91.
- Xie, J. et al. (2019) 'Deep learning based analysis of histopathological images of breast cancer', Frontiers in Genetics, 10(FEB), pp. 1–19. doi: 10.3389/ fgene.2019.00080.
- Zerouaoui, H. et al. (2021) 'Breast Fine Needle Cytological Classification Using Deep Hybrid Architectures BT -Computational Science and Its Applications – ICCSA 2021', in Gervasi, O. et al. (eds). Cham: Springer International Publishing, pp. 186–202.
- Zerouaoui, H. and Idri, A. (2021) 'Reviewing Machine Learning and Image Processing Based Decision-Making Systems for Breast Cancer Imaging'. Journal of Medical Systems.
- Zhang, G. et al. (2011) 'A review of breast tissue classification in mammograms', Proceedings of the 2011 ACM Research in Applied Computation Symposium, RACS 2011, pp. 232–237. doi: 10.1145/ 2103380.2103426.
- Zhu, C. et al. (2019) 'Breast cancer histopathology image classification through assembling multiple compact CNNs', BMC Medical Informatics and Decision Making, 19(1), pp. 1–17. doi: 10.1186/s12911-019-0913-x.