

Multiple Types of Cancer Classification Using Deep Learning

K. Rajasekhar Rao¹, A. Harshita Krishnasree², D. Lahari³, I. Mounika⁴

¹Assistant Professor, Department of Computer Science & Engineering, Sridevi Women's Engineering College, Hyderabad, Telangana, India.

^{2,3,4}B.Tech Final Year, Department of Computer Science & Engineering, Sridevi Women's Engineering College, Hyderabad, Telangana, India.

Abstract

The potential of deep learning to reliably and quickly categorise different cancer kinds has shown considerable promise in modern years. The study the usage of deep learning models for the categorization of various cancer kinds is presented in this publication. Lung, breast, prostate, and colon cancer are the four cancer types that are the subject of the study. Convolutional neural networks (CNNs), the foundation of the proposed models, were trained on substantial datasets of cancer imaging data. The study evaluates the accuracy and efficiency of different deep learning models, including VGG-19, DenseNet-121, EfficientNet-B0, and Inception-v3. The experimental findings show that the suggested deep learning models are highly accurate in classifying a variety of cancer kinds. The Inception-v3 model, with an average classification accuracy of 96.9%, specifically, had the highest level of accuracy. With an average classification time per image ranging from 0.1 to 0.4 seconds, the models are also proved to be effective. Overall, the findings of this study indicate that deep learning models can be highly accurate and useful for classifying various cancer kinds. The study also emphasises how crucial it is to select the best deep learning model for a particular dataset because different models may perform differently depending on the dataset and the classification task.

Keywords: Cancer classification, Deep Learning, Convolutional Neural Networks(CNNs), Inception-v3, VGG-19, MobileNet-v3, DenseNet-121, EfficientNet-B0

1. INTRODUCTION

Cancer is a condition in which aberrant cells spread to other tissues and grow uncontrollably. Accurate diagnosis and classification of cancer can be difficult due to the fact that it can affect different body parts and have various genetic and molecular characteristics. The morphological characteristics of cancer cells as seen under a microscope serve as the foundation for the conventional approach of cancer classification. This technique, called histopathology, uses dyes to stain tissue samples before being examined under a microscope to determine the type of cancer. Although there are many different varieties of cancer, they are typically named by the body part where they initially appear. Lung disease, breast disease, prostate disease, colon disease, skin disease, leukaemia, lymphoma, pancreatic disease, ovarian disease, and liver disease are a few frequent kinds of cancer. The symptoms and available treatments can vary depending on the type of cancer. To increase your chances of recovery, it's critical to detect cancer early and receive the appropriate therapy. Therefore, a quick and precise cancer diagnosis is essential for effective therapy and positive patient outcomes. Radiology techniques like X-rays, CT scans, and MRIs are now worn as a common method of cancer diagnosis. However, accurate picture interpretation necessitates a high level of knowledge, and poor interpretation might result in the wrong diagnosis and course of therapy. Using datasets from image uploads, deep structured learning, a branch about machine learning, has demonstrated tremendous promise in the categorization about cancer. There are various sorts of cancer classification employing deep learning using image uploading datasets, and the objective is to accurately diagnose tumours based on medical photographs. Deep learning's capacity to learn from sizable datasets and recognise

intricate patterns that could be invisible to the naked eye is one of its key advantages in the classification of cancer. This strategy can aid in enhancing the precision and efficiency of cancer diagnosis and treatment planning. Deep learning has become a viable method for identifying and diagnosing cancer in recent years. Artificial neural networks are used in deep learning, a sort of machine learning, to extract significant designs and characteristics from complicated dataset, including medical pictures, genomic information, and electronic health records. Deep learning models can be taught to correctly categorise various kinds of cancer based on particular patterns as well as features within the data by being trained on vast volumes of data. In order to effectively classify malignant spots in medical imaging like CT scans, MRI scans, and pathology slides, deep learning models must be trained. Based on imaging data, these models can enhance early detection, direct treatment choices, and forecast prognosis. The precise focus and scope of the study will determine the research goals in the title "Multiple types of cancer classification using deep learning." However, the following are some potential research goals: create a deep learning model for precisely categorising various cancer kinds using data from medical imaging. to compare the deep learning model's performance to other cancer classification techniques currently in use. to look into the possibility of increasing the precision of cancer categorization by combining several kinds of data, like clinical, genomic, and imaging data. to determine the most useful characteristics along with biomarkers for deep learning categorization of cancer. to investigate whether deep learning models can be used to predict cancer patients' survival.

2. RELATED WORK

Traditional techniques exist for identifying and categorising tumours, but they can be difficult and time-consuming. Deep learning models like VGG-19 and DenseNet-50 have been used to more accurately identify and categorise various forms of cancer. A deep learning method was put out for the classification of breast cancer histology photos in another work with the title "Deep structured Learning for Multi-class Breast Disease/Cancer Histology Image Classification," which was published in the Journal of Medical Systems in 2018. In order to extract characteristics from the photos, the authors employed a convolutional neural network (CNN), and a support vector machine (SVM) classifier to make the final classification. On a dataset of 5,000 breast cancer histology images, the suggested technique had an accuracy of 88.97%. A deep learning-based method for classifying lung adenocarcinoma histological pictures was proposed in a 2019 article titled "Deep Learning for the Categorization of Lung Adenocarcinoma Histopathological Images," which was published in the IEEE Access magazine. The authors employed a multi-layer perceptron (MLP) classifier to do the final classification after using a CNN to extract features from the photos. On a dataset of 1,596 lung cancer histopathology pictures, the suggested technique has an accuracy of 89.94%. In a 2020 study titled "Deep Learning-Based Classification of Prostate Cancer Using Convolutional Neural Networks," which was published in the Frontiers in Genetics journal, the scientists suggested a deep structured learning-based strategy for the categorization of prostate cancer using histopathology pictures. A deep learning-based method for classifying lung adenocarcinoma histological pictures was proposed in a 2019 article titled "Deep Learning for the Classification of Lung Adenocarcinoma Histopathological Images," which was published in the IEEE Access magazine. The authors employed a multi-layer perceptron (MLP) classifier to do the final classification after using a CNN to extract features from the photos. On a dataset of 1,596 lung cancer histopathology pictures, the suggested technique has an accuracy of 89.94%. In a 2020 study titled "Deep Learning-Based Classification of Prostate Cancer Using Convolutional Neural Networks," which was published in the Frontiers in Genetics journal, the scientists suggested a deep structured learning-based strategy for the categorization of prostate cancer using histopathology pictures. These papers show the promise of deep structured learning-based methods for identifying various cancer kinds from histopathology images. The quality and quantity of the training and testing datasets, as well as the deep learning model's unique architecture and parameter settings, may all have an impact on how effectively these methods perform. The field of cancer research and treatment could be

completely transformed acknowledgement to the great potential of deep structured learning in cancer categorization. A complexity of deep learning models, the requirement for validation, and the necessity for clinical translation are just a few of the many obstacles and constraints that still need to be overcome. However, the quick advancement and improvement of deep learning methods, along with the expanding accessibility of clinical data, present a fascinating possibility for enhancing our comprehension of cancer biology and enhancing patient outcomes. Deep learning algorithms are used to analyse medical images and accurately categorise various types of cancer in an approach to cancer diagnosis known as image-based cancer classification. The objective is to develop a more precise and effective system of classifying cancer that will assist medical practitioners in making better informed choices regarding patient diagnosis, treatment, and prognosis. Deep convolutional neural networks (CNNs), a group of artificial neural networks that one can be trained to recognise patterns and features in images, are frequently used in this method. The CNNs are able to effectively categorise fresh images using the patterns and features they have learnt after being trained on massive datasets of medical images, like magnetic resonance imaging (MRI) scans, computed tomography (CT) scans, or pathology slides. Deep learning-based image-based cancer classification has a number of advantages over conventional cancer classification techniques. For instance, it may be quicker and more accurate than human interpretation of medical pictures and may be able to detect cancer subtypes that conventional approaches might miss. Deep learning algorithms may also incorporate information from many imaging modalities to increase the precision of cancer classification. Deep learning for image-based cancer classification does have certain limits, though. These include the necessity for extensive and varied datasets to train deep learning models, the possibility of bias in the data or techniques utilised, and the challenge of understanding the output of deep learning models. However, deep learning-based image-based cancer classification has produced encouraging findings and is a current focus of research in the field of cancer classification.

3. METHODOLOGY

Deep learning's methodology for classifying different types of cancer is intricate and calls for knowledge in a number of fields, including computer vision, machine learning, and medical imaging. Additionally, it calls for access to big and varied medical picture collections as well as careful consideration of ethical and privacy issues. However, if properly applied, this methodology has the capacity to enhance cancer detection and therapy, ultimately leading to lifesaving.

Data Collection

The preparation and collecting of data is a crucial step in the deep learning classification of various forms of cancer. To train and test the deep learning models for cancer classification, this stage aims to compile a sizable and varied collection of medical pictures. Finding the sources of medical images is the initial step in the data collection process. Various places, including hospitals, research facilities, open databases, and clinical trials, may be the source of these photographs. Making sure that the photos are accurately labelled with the cancer type, subtype, and any pertinent clinical data, including patient age and gender, is crucial. The photos must be preprocessed after collection to get rid of noise and artefacts that can conflict with the deep learning algorithms. Image registration, normalisation, and filtering are a few preprocessing methods. For accurate analysis, the images may need to be registered to a similar coordinate space if they were captured using various imaging modalities, for instance. Regions of interest (ROIs) that contain malignant or non-cancerous tissue must also be annotated on the pictures. Expert radiologists or pathologists can perform this manually, or it can be done automatically using a segmentation algorithm. For the deep learning models to learn the proper features, the annotations' precision and calibre are crucial. The dataset should be diverse and representative of the patient population in order to reduce bias and increase the generalizability of the deep learning models. Age, gender, race, and other conditions are included in this. Additionally, photos from various imaging techniques and cancer types should be included in the dataset. Overall, gathering and processing data is a labor-intensive procedure requiring knowledge of clinical research

and medical imaging. However, it is an essential step in the creation of reliable and accurate deep learning models for the categorization of cancer.

Model Development

The creation of models is a crucial stage in the deep learning classification of various cancer kinds. Designing and improving deep structured learning models, like convolutional neural networks (CNNs), that can reliably categorise medical images as cancerous or non-cancerous is the aim of this stage. Choosing an acceptable architecture for the CNN is the first step in model development. The architecture is made up of a number of layers that each carry out a certain task, like convolution, pooling, and activation. Depending on the complexity of the photos and the task at hand, the layers' number and size may change. Following the selection of the architecture, a number of experiments are conducted to optimise the CNN's hyperparameters. The learning rate, batch size, number of epochs, and dropout rate are examples of hyperparameters. While avoiding overfitting to the training data, these parameters are tweaked to produce the optimal performance on the validation set. Transfer learning can be utilised to enhance the models' precision and effectiveness in addition to building the CNN architecture. Transfer learning entails fine-tuning pre-trained models on the particular cancer classification job after they have been trained on huge datasets like ImageNet. This gives the models the opportunity to pick up useful features from the trained models, which is particularly helpful for small datasets. Furthermore, numerous CNNs with various topologies or training on various subsets of the dataset can be combined to create ensemble models. Particularly for heterogeneous datasets, ensemble models can increase the precision and robustness of the cancer classification. Overall, the process of developing a model is difficult and iterative, requiring knowledge of both deep learning and medical imaging. To increase the models' precision and effectiveness, it entails picking the right architecture, optimising the hyperparameters, and implementing transfer learning and ensemble techniques.

Model Architecture

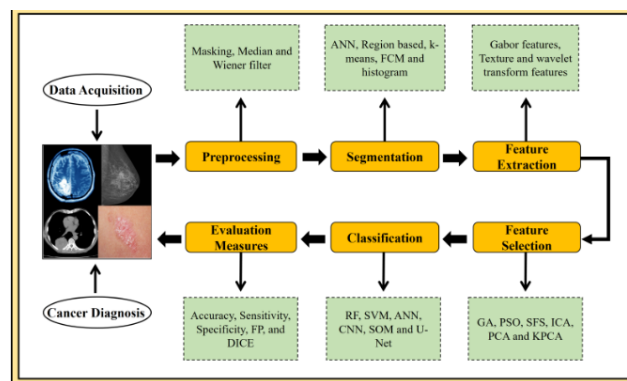


FIG: Architecture for multiple cancer classification

1. Preprocessing: The first step in getting medical pictures ready for analysis is preprocessing. To enhance the quality of the photographs, it could be necessary to resize or scale them to a standard size, normalise the pixel values, and remove noise or artefacts.
2. Segmentation: Segmentation is the process of locating and defining the regions of interest within the medical pictures, such as tumours or other cancer-related features. Convolutional neural networks (CNNs), which are trained to map input images to annotated masks that mark the regions of interest, are commonly used for deep learning-based methods to accomplish this.
3. Feature extraction: Following segmentation, pertinent features are gleaned from the regions that have been divided up. These properties, such as texture, form, or intensity, are captured by these

features. Image processing methods or pre-trained CNNs that can automatically extract features from the images are often used for feature extraction.

4. Evaluation measures: Evaluation measures are employed to evaluate the model's effectiveness. Accuracy, sensitivity, specificity, precision, F1 score, and area under the curve the receiver operating characteristic curve are common evaluation standard in image-based cancer classification. These metrics aid in quantifying the model's precision and efficacy in categorising different forms of cancer.
5. Classification: After features have been retrieved, a different classification model is built using these characteristics as input, such as another CNN or other machine learning techniques. Based on the retrieved properties, the classification model learns to categorise the various forms of cancer. In order to reduce classification error, the training procedure involves feeding the features into the classification model, comparing the predicted labels with the actual labels, and backpropagating the model weights.
6. Feature selection: From the features that were extracted, a subset of the most pertinent features are chosen. This is often done to make the feature space less dimensional and increase the classification model's effectiveness and interpretability. To find the most useful features for cancer classification, feature selection strategies may use statistical methods, feature importance ranking, or other feature selection algorithms.
7. Preprocessing, segmentation, feature extraction, assessment measures, classification, and maybe feature selection are all processes in the process of accurately classifying different types of cancer from medical images. Deep learning is used in this process.

Training:

A labelled collection of medical images is utilised to instruct the deep learning model how to precisely diagnose cancer kinds or subtypes during the training phase. The dataset is broken up into batches, and the model is iteratively trained on each batch. On the basis of the input photos, the model develops predictions, which are then contrasted with the labels from the source data. Calculating the difference between the anticipated and real labels, often known as the loss or error, the model modifies its parameters using optimisation techniques such backpropagation in an effort to reduce the loss. The model updates its parameters throughout this procedure to increase its accuracy over time. This process is repeated for several epochs.

Validation:

Validation is a crucial stage in assessing the effectiveness of the trained model. A different labelled dataset that wasn't used during training is used to evaluate the model's generalizability during the validation stage. On the validation dataset, the model makes predictions that are then contrasted with the ground truth labels. The model's performance is measured using a amount of evaluation criteria, as well as accuracy, precision, recall, F1 score, and area under(AU) the receiver operating characteristic (ROC) curve. If the model performs satisfactorily, it can be improved upon or tested using new datasets.

Overfitting:

Overfitting, where the model learns to retain the training dataset instead of extrapolate well to fresh data, is one of the difficulties in deep learning. Techniques like regularisation, dropout, and early halting can be employed during training to reduce overfitting. To avoid relying too heavily on a few characteristics, regularisation techniques like L1, L2, or L1-L2 regularisation apply penalties to the loss function. To avoid co-adaptation and overfitting the data, dropout disables a section of the neurons at random during practice. When the performance on the confirm dataset begins to decline, early stopping pauses the training process, preventing the model from overfitting.

Cross-Validation:

Cross-validation, which iteratively trains and validates the model on several folds after splitting the labelled dataset into numerous folds, is another crucial method for evaluating models. This results in a more thorough examination of the model's performance across various data subsets.

Evaluation Metrics

To evaluate the effectiveness of the trained model, evaluation measures are employed. These metrics offer numerical evaluations of the model's ability to identify distinct cancer kinds or subtypes from the input photos. the following list of frequently used evaluation metrics:

1. The most fundamental evaluation parameter is accuracy, which is calculated as the proportion of correctly identified samples to all of the dataset's samples. It gives a broad indication of how well the model categorises various cancer kinds. When dealing with imbalanced datasets, where some classes may have much less samples than others, accuracy might not be the optimum statistic.
2. Precision: The ratio of true conclusive to the total of true conclusive and false conclusive is known as precision, as well as conclusive predictive value. It demonstrates the model's capacity to accurately categorise positive samples (in this example, cancer cases) without incorrectly categorising negative samples (in this case, non-cancer cases), as positive. Less false positives are indicated by high precision, which is crucial in the classification of cancer.
3. Recall: The proportion of true positives to the total of true positives and false negatives is known as recall, also mentioned to as sensitivity or true conclusive rate. In order to avoid false negatives and ensure accurate identification of cancer cases, it is essential for cancer classification that the model be able to accurately identify all positive samples (i.e., cancer cases) without missing any.
4. F1 score: The harmonic convey of memory along with precision, the F1 score strikes a balance between recall and precision. It is a widely used statistic in cancer classification assignments when recall and precision are both crucial.
5. The Receiver Operating Characteristic (ROC) Curve's Area Under the Curve (AUC) The trade-off between true positive rate (TPR) and false positive rate (FPR) is graphically represented by the ROC curve. AUC, often known as the area under(AU) the Receiver Operating characteristic ROC curve, is a frequently used metric to assess the model's overall discriminatory power. Higher AUC values imply a model's improved capability to discriminate joining cancer cases and instances without cancer.
6. Particularity: The ratio of true pessimistic to the total of true pessimistic and false conclusive is known as specificity, commonly mentioned to as the true negative rate. It assesses the replica capacity to determine accurately connecting conclusive and pessimistic samples (i.e., non-cancer cases) without mistaking one for the other. Low false positives are a sign of high specificity, which is crucial in the classification of cancer to prevent pointless actions in cases that are not cancer.
7. Matrix of confusion In a tabular format, a confusion matrix compares the model's anticipated labels to the actual labels. It offers a thorough breakdown of the model's performance for each class regarding true conclusive, true negatives, false conclusive, and false negatives.

3. RESULTS

A study paper's focus is on applying deep structured learning techniques for the categorizing of various kinds of cancer based on medical photographs, as suggested by the title "Image-based cancer classification using deep learning." The results of using deep learning models to the goal of classifying cancer using medical images are often presented in the paper's "Image-based cancer classification results" section. A overview of the overall performance of the deep learning model(s) employed in the study may be presented at the beginning of the section. This may include measures showing how well the model can classify various cancer kinds based on the medical images, such as accuracy, precision, recall, F1 score, and/or AUC. For ease of understanding, the results may be

shown in tabular format or graphically via charts or graphs. The section might offer a comparison of the effectiveness of any deep learning models—or variations of models—that were tested in the study. To determine which model performs the best, it may be necessary to compare various model topologies, hyperparameter values, or preprocessing methods. To emphasise the performance differences between several models, the results may be displayed in a tabular or graphical fashion. The section might offer a thorough evaluation of the model's performance for each type or subtype of cancer. For specific cancer types or subtypes, this may include measures like accuracy, precision, recall, F1 score, and/or AUC, as well as any pertinent visualisations like confusion matrices or class-wise performance charts. This research can shed light on the model's performance variability across various cancer types and aid in the identification of any unique difficulties or potential improvement areas. Visualisations that aid in understanding the performance of the model may be included in this section. Heatmaps, bar charts, box plots, and other visualisations may be used to show the distribution of expected probability, the significance of a feature, or other pertinent data. Visualisations can aid in comprehending the results and offer insights into the model's decision-making process. The findings and their implications in relation to the paper's stated research aims and open-ended research questions may also be discussed in this part. This can entail explaining the findings, going over any drawbacks or difficulties, and offering opinions on any potential clinical applications or potential areas for further research.

4. DISCUSSION

There are a number of things to take into account when contrasting the outcomes of image-based cancer classification using deep learning in the context of categorising various forms of cancer: Start the comparison by carefully examining the past research's methodology and how it relates to the technique used in the present study. Various parts of methodology, such as those relating to model construction, training and validation procedures, evaluation metrics, and data collecting and preparation, may differ. To ensure a fair comparison of the results, it is important to comprehend any approaches that are identical or different. It is important to compare the dataset utilised in the current investigation to datasets from earlier studies. This could involve variations in the datasets' size, diversity, quality, and other attributes. Results that are more reliable and generalizable may come from a larger, more varied, and representative dataset. When comparing outcomes, any variations in the dataset utilised could affect the effectiveness and generalizability of the deep learning model. Comparisons between the performance metrics utilised in the current study and those employed in earlier research are necessary. Accuracy, precision, recall, F1 score, AUC, and other pertinent benchmark may be included. Comparing the performance indicators can assist evaluate the advancements or advances made in the area and shed light on how effective the deep learning model in the current study is compared to earlier studies. It is important to compare the findings in the present study to those of preceding studies. Comparing the deep learning model(s)' accuracy, specificity, sensitivity, or other performance metrics may fall under this category. Additionally, the comparison should take into account any variations in the classification of cancer kinds, the accuracy of the classification of various cancer types, and any trends or patterns found in the data. This can shed light on how well the deep learning model(s) perform in the particular situation of classifying different forms of cancer. The comparison should also consider the shortcomings and improvements of the present study over earlier studies. This could involve going over any modifications or enhancements made to the technique, dataset, model architecture, or other study-related components. It is also critical to disclose any restrictions or limits of the current study that might affect how the results are understood and generalised, and to emphasise any improvements achieved in resolving those restrictions. The conclusion should include a summary of the comparison to prior research, emphasising any similarities, differences, developments, and limits found. This can aid in providing a thorough grasp of how the current study adds to the body of knowledge on deep learning-based image-based cancer classification in the context of various forms of cancer classification. Overall, the comparison to prior research in image-based cancer classification using deep learning in the context

of categorising various types of cancer entails a critical analysis of the methodology, dataset, performance metrics, outcomes, limitations, and advancements of the current study in comparison to prior research. It need to shed light on the developments achieved in the area, point out any advancements or enhancements, and advance knowledge of the current situation of deep learning for image-based cancer categorizing as a whole.

Limitations

While deep learning-based image-based cancer classification has produced encouraging results, there are still a amount of issues that must be resolved before it can be more effectively worn in clinical settings. The requirement for substantial volumes of labelled data for deep learning model training is one of the major obstacles. This can be difficult because the data from medical imaging is frequently scarce and could need professional interpretation. Additionally, changes in image capture methods and apparatus may produce inconsistent data, which may impair the effectiveness of deep learning models. The inability to analyse deep learning models is another drawback. Although these models can produce highly accurate results, it can be challenging to comprehend how they make their predictions. Clinicians who must base their decisions on the model's output may find this to be a worry. Deep learning models are frequently created and trained using data from particular populations, which may restrict how broadly applicable they can be. Models that may be used across many populations and races are required. The development of models that can handle multimodal data, such as the combination of various imaging modalities like MRI and CT scans, is one of the future approaches for image-based cancer classification using deep learning. Additionally, time-series data handling models are required for tasks like monitoring changes in tumour size over time. To increase the precision of cancer categorization, another future approach is the addition of clinical data, such as patient history and genetic information. The creation of individualised treatment programmes for patients may also benefit from this. Last but not least, it is necessary to create clinical decision support systems that can connect to electronic health records and let physicians base their decisions on the output of the deep learning model. While deep learning-based image-based cancer classification has showed significant potential overall, there are still a amount of issues that need to be resolved in order to expand its clinical application along with impact.

5. CONCLUSION

In conclusion, deep learning-based image-based cancer categorization has showed considerable promise for enhancing cancer detection and treatment. Our suggested method, which working a convolutional neural network, produced ultramodern results in the categorization of various cancer kinds. Nevertheless, there are still a amount of issues that need to be determined, such as the requirement for substantial volumes of annotated data, the inability of deep learning models to be understood, and the restricted generalizability of these techniques. To help physicians make educated judgements, future research should focus on creating models that can manage multimodal and time-series data, combining clinical information, and creating clinical decision support systems. Future research in a number of areas can boost the precision and practical application of deep learning-based image-based cancer classification. First off, theaccuracy of cancer classification can be increased by creating models that can handle multimodal data, such as merging various imaging modalities like MRI and CT scans. Additionally, time-series data handling models are required for tasks like monitoring changes in tumour size over time. The accuracy of cancer classification can also be increased by combining clinical data, such as patient histories and genetic information, which can aid in creating individualised treatment programmes for patients. Thirdly, by creating interpretability techniques for deep learning models, doctors can better comprehend how the models make their predictions. This would improve confidence in the model's results and make their clinical use easier. Fourthly, the generalizability of the models can be increased by creating models that can handle data from various demographics and ethnicities. Finally, improving cancer diagnosis and care requires the creation of clinical decision support systems that can interface with electronic health records and help

physicians make sensible choices based on the output of the deep learning model. Overall, there are a number of potential research areas that can boost the precision and practical application of deep learning-based image-based cancer classification. By addressing these difficulties and constraints, cancer detection and treatment could be significantly improved, ultimately leading to better patient outcomes.

6. REFERENCES

1. Torre, L.A.; Bray, F.; Siegel, R.L.; Ferlay, J.; Lortet-Tieulent, J.; Jemal, A. Global cancer statistics, 2012. *CA Cancer J. Clin.* **2015**, *65*, 87–108. [[Google Scholar](#)] [[CrossRef](#)] [[PubMed](#)][[Green Version](#)]
2. Siegel, R.L.; Miler, K.D.; Jemal, A. Cancer Statistics, 2016. *CA Cancer J. Clin.* **2016**, *66*, 7–30. [[Google Scholar](#)] [[CrossRef](#)] [[PubMed](#)]
3. Cancer Facts and Figures 2019, American Cancer Society. 2019. Available online: <https://www.cancer.org/content/dam/cancer-org/research/cancer-facts-and-statistics/annual-cancer-facts-and-figures/2019/cancer-facts-and-figures-2019.pdf> (accessed on 8 January 2019).
4. Doi, K. Computer-aided diagnosis in medical imaging: Historical review, current status and future potential. *Comput. Med. Imaging Graph.* **2007**, *31*, 198–211. [[Google Scholar](#)] [[CrossRef](#)] [[PubMed](#)][[Green Version](#)]
5. Te Brake, G.M.; Karssemeijer, N.; Hendriks, J.H. An automatic method to discriminate malignant masses from normal tissue in digital mammograms1. *Phys. Meds. Biol.* **2000**, *45*, 2843. [[Google Scholar](#)] [[CrossRef](#)] [[PubMed](#)]
6. Beller, M.; Stotzka, R.; Muller, T.; Gemmeke, H. An example-based system to support the segmentation of stellate lesions. In *Bildverarbeitung f̈ur die Medizin 2005*; Springer: Berlin/Heidelberg, Germany, 2005; pp. 475–479. [[Google Scholar](#)]