

Blood Cell Analysis: Deep Learning Integration with Diffraction Phase Microscopy

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DESCRIPTION

Blood cell characterization plays an important role in diagnosing various diseases and monitoring patients' health conditions. Traditional methods for blood cell analysis often involve time-consuming manual processes and can lack accuracy and efficiency. In this article, we propose a novel approach for blood cell characterization by integrating deep learning techniques with Diffraction Phase Microscopy (DPM). DPM offers label-free, high-resolution imaging capabilities, making it suitable for studying biological samples like blood cells. Leveraging the power of deep learning, we develop a robust and accurate classification model capable of automatically identifying and classifying different blood cell types based on DPM images. We demonstrate the effectiveness of our proposed method through extensive experiments on real blood cell samples, achieving high classification accuracy and outperforming existing approaches. Our proposed framework has the potential to revolutionize blood cell analysis, enabling faster and more reliable diagnosis of various blood-related disorders.

Advancements in blood cell analysis

Blood cell analysis is a fundamental aspect of clinical diagnosis and medical research, providing valuable insights into an individual's health status. Various blood disorders, such as anemia, leukemia, and thrombocytopenia, can significantly impact a person's well-being and require accurate and timely diagnosis for appropriate treatment. Traditional methods for blood cell characterization often involve staining techniques followed by manual examination under a microscope. However, these methods are labor-intensive, time-consuming, and prone to human error. Moreover, the use of stains may alter the cellular structure, limiting the accuracy of analysis.

In recent years, advances in imaging technologies have paved the way for label-free and non-invasive approaches to blood cell analysis. Diffraction Phase Microscopy (DPM) is one such technique that offers high-resolution imaging capabilities

without the need for staining. DPM measures the phase shift of light passing through a sample, providing detailed information about cellular morphology and dynamics. By combining DPM with computational techniques, such as deep learning, it is possible to automate the process of blood cell characterization, offering advantages in terms of speed, accuracy, and scalability. In this article, we propose a novel framework for blood cell characterization based on deep learning and diffraction phase microscopy. The proposed methodology consists of the following steps:

Data acquisition: Blood cell samples are imaged using a diffraction phase microscope to obtain high-resolution phase contrast images.

Preprocessing: The acquired images are preprocessed to enhance contrast, remove noise, and normalize intensity levels, ensuring consistency across different samples.

Deep learning model: A Convolutional Neural Network (CNN) architecture is designed and trained using a large dataset of annotated blood cell images. The CNN learns to automatically extract relevant features from the DPM images and classify blood cells into different categories, such as red blood cells, white blood cells, and platelets.

Model evaluation: The trained model is evaluated using a separate test dataset to assess its performance in terms of accuracy, sensitivity, specificity, and other relevant metrics.

Integration: The trained model is integrated into a user-friendly software tool, allowing clinicians and researchers to perform automated blood cell analysis with ease.

Experimental results demonstrate the effectiveness of the proposed approach in accurately identifying and classifying different blood cell types based on DPM image. The deep learning model achieves high classification accuracy, outperforming existing methods for blood cell characterization. Moreover, the proposed framework exhibits robustness against

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variations in sample preparation and imaging conditions, making it suitable for real-world applications.

CONCLUSION

The article presents a novel approach for blood cell characterization using deep learning and diffraction phase microscopy. By leveraging the unique capabilities of DPM and

the computational power of deep learning, we have developed a robust and accurate classification model capable of automatically identifying and classifying different blood cell types. Our proposed framework offers several advantages over traditional methods, including speed, accuracy, and label-free imaging. Future work may involve further optimization of the deep learning model and integration with other imaging modalities for comprehensive blood cell analysis.