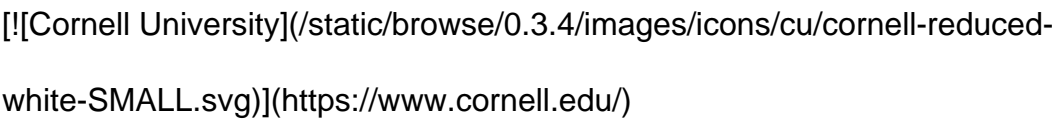


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> generalizability. Many studies have focused on specific features without
> exploring more comprehensive approaches. In our case, we formulate a deep
> learning technique for malaria-infected cell classification using
> traditional CNNs and transfer learning models notably VGG19, InceptionV3,
> and Xception. The models were trained using NIH datasets and tested using
> different performance metrics such as accuracy, precision, recall, and
> F1-score. The test results showed that deep CNNs achieved the highest
> accuracy -- 97%, followed by Xception with an accuracy of 95%. A machine
> learning model SVM achieved an accuracy of 83%, while an Inception-V3
> achieved an accuracy of 94%. Furthermore, the system can be accessed through
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