

DAFTAR PUSTAKA

- [1] R. E. Litz, *The Mango: Botany, Production and Uses*, 2nd ed. Wallingford, UK: CABI, 2009.
- [2] M. A. P. Suwantra, K. N. Ramadhani, and F. Sthevanie, "Leaf species identification using multi texton histogram and support vector machine," in *Proceedings of the 2020 3rd International Conference on Image and Graphics Processing*, 2020, pp. 41–45.
- [3] U. S. Rao, R. Swathi, V. Sanjana, L. Arpitha, K. Chandrasekhar, P. K. Naik, et al., "Deep learning precision farming: Grapes and mango leaf disease detection by transfer learning," *Global Transitions Proceedings*, vol. 2, no. 2, pp. 535–544, November 2021.
- [4] C. Vijay and K. Pushpalatha, "Dv-pso-net: A novel deep mutual learning model with heuristic search using particle swarm optimization for mango leaf disease detection," *Journal of Integrated Science and Technology*, vol. 12, no. 5, pp. 804–804, 2024.
- [5] Z. Liu, H. Mao, C.-Y. Wu, C. Feichtenhofer, T. Darrell, and S. Xie, "A convnet for the 2020s," in *Proceedings of the IEEE/CVF Conference on Computer Vision and Pattern Recognition*, 2022, pp. 11976–11986.
- [6] X. Wang, Y. Wang, J. Zhao, and J. Niu, "Eca-convnext: A rice leaf disease identification model based on convnext," in *Proceedings of the IEEE/CVF Conference on Computer Vision and Pattern Recognition*, 2023, pp. 6234–6242.
- [7] H. Li, M. Qi, B. Du, Q. Li, H. Gao, J. Yu, C. Bi, H. Yu, M. Liang, G. Ye, et al., "Maize disease classification system design based on improved convnext," *Sustainability*, vol. 15, no. 20, p. 14858, October 2023.
- [8] Z. Liu, Y. Lin, Y. Cao, H. Hu, Y. Wei, Z. Zhang, S. Lin, and B. Guo, "Swin transformer: Hierarchical vision transformer using shifted windows," in

Proceedings of the IEEE/CVF International Conference on Computer Vision, 2021, pp. 10012–10022.

- [9] H.-J. Yu and C.-H. Son, “Leaf spot attention network for apple leaf disease identification,” in 2020 IEEE/CVF Conference on Computer Vision and Pattern Recognition Workshops (CVPRW), 2020, pp. 52–53.
- [10] K. He, X. Zhang, S. Ren, and J. Sun, “Deep residual learning for image recognition,” Available online, pp. 770–778, 2016.
- [11] K. He, X. Zhang, S. Ren, and J. Sun, “Identity mappings in deep residual networks,” in Computer Vision—ECCV 2016: 14th European Conference, Amsterdam, The Netherlands, October 11–14, 2016, Proceedings, Part IV, vol. 14. Springer, 2016, pp. 630–645.
- [12] S. Zagoruyko and N. Komodakis, “Wide residual networks,” CoRR, vol. abs/1605.07146, 2016.
- [13] S. Woo, S. Debnath, R. Hu, X. Chen, Z. Liu, I. S. Kweon, and S. Xie, “Convnext v2: Co-designing and scaling convnets with masked autoencoders,” Available online, pp. 16133–16142, 2023.
- [14] S. Ali, M. Ibrahim, S. I. Ahmed, M. Nadim, M. R. Mizanur, M. M. Shejunti, and T. Jabid, “Mangoleafbd dataset,” Mendeley Data, V1, doi: 10.17632/hxsnvwty3r.1, 2022.
- [15] M. M. Rahman, K. A. Kowser, M. Islam, and A. S. S. Md Nazmul Arefin, “Mango leaf disease identification dataset: (MLDID),” Mendeley Data, V1, doi: 10.17632/jpwtpv2c4s.1, 2024.
- [16] M. A. S. Nirob, P. Bishshash, A. K. M. F. K. Siam, S. Mia, T. Khatun, and M. S. Uddin, “Mango dataset: A comprehensive resource for agricultural research and disease detection,” Mendeley Data, V1, doi: 10.17632/fn8dgm4hb5.1, 2024.

- [17] M. M. H. Shakib, S. Mustofa, and M. T. Ahad, “MLD24: An image dataset for mango leaf disease detection,” Mendeley Data, V1, doi: 10.17632/6dvpywm2m2.1, 2024.
- [18] I. Loshchilov and F. Hutter, “Decoupled weight decay regularization,” Available online, 2019.
- [19] R. Mohammed, J. Rawashdeh, and M. Abdullah, “Machine learning with oversampling and undersampling techniques: Overview study and experimental results,” in Proceedings of the 2020 International Conference on Intelligent Computing and Intelligent Systems (ICICS). IEEE, 2020, pp. 243–248.
- [20] D. Berrar, “Cross-validation,” in Encyclopedia of Bioinformatics and Computational Biology, S. Ranganathan, M. Gribskov, K. Nakai, and C. Schönbach, Eds. Oxford: Academic Press, 2019, pp. 542–545.
- [21] J. M. Gorriz, F. Segovia, J. Ramirez, A. Ortiz, and J. Suckling, “Is k-fold cross validation the best model selection method for machine learning?” arXiv preprint, arXiv:2401.16407, 2024.