

DAFTAR PUSTAKA

- Adhikari, S., Saha, S., Bandyopadhyay, T. K., & Ghosh, P. (2015). Efficiency of ISSR marker for characterization of *Cymbopogon* germplasms and their suitability in molecular barcoding. *Plant systematics and evolution*, 301, 439-450.
- Adelina, M., Harianto, S. P., & Nurcahyani, N. (2016). Keanekaragaman jenis burung di hutan rakyat pekon kelungu kecamatan kotaagung kabupaten tanggamus. *Jurnal Sylva Lestari*, 4(2), 51-60.
- Agyenim-Boateng KG, Lu J, Shi Y, Zhang D, Yin X. (2019). SRAP analysis of the genetic diversity of wild castor (*Ricinus communis* L.) in South China. *PLoS ONE*. 14(7): e0219667. <https://doi.org/10.1371/journal.pone.0219667>
- Ahmad, F. dan Y.S. Poerba. 2010. Penampilan random amplified polymorphic DNA pada *Azadirachta indica* A. Juss dari Taman Nasional Baluran. *Jurnal Teknik Lingkungan* 11(1) : 61-69.
- Alam, M.A., Juraimi, A.S., Rafii, M.Y. *et al.* (2015). Application of EST-SSR marker in detection of genetic variation among purslane (*Portulaca oleracea* L.) accessions. *Braz. J. Bot.* 38: 119–129. <https://doi.org/10.1007/s40415-014-0103-0>
- Aljumaili, S. J., Rafii, M. Y., Latif, M. A., Sakimin, S. Z., Arolu, I. W., & Miah, G. (2018). Genetic diversity of aromatic rice germplasm revealed by SSR markers. *BioMed research international*, 2018(1), 7658032
- Amirul Alam, M., Juraimi, A. S., Rafii, M. Y., Hamid, A. A., Kamal Uddin, M., Alam, M. Z., & Latif, M. A. (2014). Genetic improvement of Purslane (*Portulaca oleracea* L.) and its future prospects. *Molecular biology reports*, 41, 7395-7411.
- Amiryousefi, A., Hyvönen, J., & Poczai, P. (2018). iMEC: Online marker efficiency calculator. *Applications in plant sciences*, 6(6), e01159.
- Alqahtani, M. (2024). Biodiversity of *Artemisia* Species From Saudi Arabia Based On Morphological and Molecular Markers. *Applied Ecology & Environmental Research*, 22(3).
- Amiteye, S. (2021). Basic concepts and methodologies of DNA marker systems in plant molecular breeding. *Heliyon*, 7(10).
- Aziz, R.R., Tahir, N.AR. (2023). Genetic diversity and structure analysis of melon (*Cucumis melo* L.) genotypes using URP, SRAP, and CDDP markers. *Genet Resour Crop Evol.* 70: 799–813. <https://doi.org/10.1007/s10722-022-01462-y>
- Azmi, W. A., Puspita, P. J., Safira, U. M., Subositi, D., & Maruzy, A. (2022). Keragaman Genetik *Graptophyllum pictum* (L.) Griff dari Etnis Indonesia Timur Berbasis Sequence-Related Amplified Polymorphism. *Indonesian Journal of Agronomy/Jurnal Agronomi Indonesia*, 50(2).

- Azzam, C. R., Sultan, F. M., and Rizk, M. S. *et al.* (2023). SRAP and IRAP revealed molecular characterization and genetic relationships among cowpea (*Vigna unguiculata* L.) irradiated by gamma-ray. *Beni-Suef Univ J Basic Appl Sci* **12**, 109. <https://doi.org/10.1186/s43088-023-00448-8>
- Bargish, T. A., & Rahmani, F. (2016). SRAP markers based genetic analysis of *Silene* species. *Journal of Tropical Biology & Conservation (JTBC)*, *13*.
- Becker, J. M., Caldwell, G. A., & Zachgo, E. A. (Eds.). (1996). *Biotechnology: a laboratory course*. Academic Press.
- Bennett, P., Williamson, C., Sykes, L., MacIntyre, D. A., & Dixon, P. H. (2022). *Basic Science in Obstetrics and Gynaecology E-Book: Basic Science in Obstetrics and Gynaecology E-Book*. Elsevier Health Sciences.
- Bublyk, O., Andreev, I., Kalendar, R., Spiridonova, K., & Kunakh, V. (2013). Efficiency of different PCR-based marker systems for assessment of *Iris pumila* genetic diversity. *Biologia*, *68*(4), 613-620.
- Castanares, E., M. I. Dinolfo, A. Patriarca, and S. A. Stenglein. (2023). SRAP markers as an alternative tool for *Alternaria* classification. *Food Microbiology*. 116: 104370.
- Chen, J. Z., Li, T., Tian, X., Zheng, Y. H., Jin, Z. L., Cao, H. N., & Zong, C. W. (2022). Taxonomic status of intraspecific germplasm resources of *Vaccinium uliginosum* based on chloroplast matK gene and SRAP molecular markers. *Journal of Berry Research*, *12*(3), 315-327.
- Chňapek, M., Balážová, Ž., Špaleková, A. *et al.* (2024). Genetic diversity of maize resources revealed by different molecular markers. *Genet Resour Crop Evol.* <https://doi.org/10.1007/s10722-024-01908-5>
- Choe, Y. I., Song, S. R., Ho, U. H. *et al.* (2024). Sequence-related amplified polymorphism (SRAP) markers reveal genetic variation of rice strains obtained by millet DNA injection through coleoptile. *Genet Resour Crop Evol.* <https://doi.org/10.1007/s10722-024-02080-6>
- Danin, A., Buldrini, F., Bandini Mazzanti, M., Bosi, G., Caria, M. C., Dandria, D., ... Bagella, S. (2016). Diversification of *Portulaca oleracea* L. complex in the Italian peninsula and adjacent islands†. *Botany Letters*. *163*(3): 261–272. <https://doi-org.ezproxy.ugm.ac.id/10.1080/23818107.2016.1200482>
- Desta, M., A. Molla, and Z. Yusuf. (2020). Characterization of physico-chemical properties and antioxidant activity of oil from seed, leaf and stem of purslane (*Portulaca oleracea* L.). *Biotechnology Reports*. *27*(e00512).
- Dwivedi, S., Singh, S., Chauhan, U. K., & Tiwari, M. K. (2018). Inter and intraspecific genetic diversity (RAPD) among three most frequent species of macrofungi

(*Ganoderma lucidum*, *Leucoagricus* sp. and *Lentinus* sp.) of Tropical forest of Central India. *Journal of Genetic Engineering and Biotechnology*, 16(1), 133-141.

- Ekbiç, E., Okay, C.Ö. (2024). Assessment of elite pepper breeding lines using molecular markers. *Plant Biotechnol Rep.* 18: 515–524. <https://doi.org/10.1007/s11816-024-00911-7>
- Fatmawati, Y. 2022. Validasi genetik berdasarkan penanda retrotransposon dan evaluasi komponen hasil pada populasi f2 dan f3 hasil persilangan intergenerik [kacang hijau (*vigna radiata*) x buncis (*phaseolus vulgaris*)] (doctoral dissertation, universitas gadjah mada).
- Felix, F. C., Chagas, K. P. T. D., Ferrari, C. D. S., Vieira, F. D. A., & Pacheco, M. V. (2020). Applications of ISSR markers in studies of genetic diversity of *Pityrocarpa moniliformis*. *Revista Caatinga*, 33(4), 1017-1024.
- Finkeldey, R. dan H. Hattemer. 2007. Tropical Forestry: Tropical Forest Genetics. Institute of Forest Genetics and Forest Tree Breeding, Germany.
- Fosket, D. E. (1994). The size and complexity of plant genomes. *Plant Growth and Development: A Molecular Approach.*, 79-152.
- Ferraz, J. R., Rossetto, E. F. S., Hassemer G., and da S. Ribeiro J. E. L. (2024). Taxonomic notes for *Portulaca* (Portulacaceae) in South America II: synonymisation of *P. diegoi* under *P. hatschbachii* based on macro and micro characters. *Kew Bulletin*. 79:223–232.
- Giri, M.S., Jingade, P., Huded, A.K.C. *et al.* (2024). Genetic characterization of *Paramyothecium roridum* strains isolated from different host plants in coffee agroecosystem using SRAP, SCoT markers and ITS sequencing. *Indian Phytopathology*. <https://doi.org/10.1007/s42360-024-00753-x>
- Govindaraju, D. R. (1989). Variation in gene flow levels among predominantly self-pollinated plants. *Journal of Evolutionary Biology*, 2(3), 173-181.
- Grativol, C., da Fonseca Lira-Medeiros, C., Hemerly, A. S., & Ferreira, P. C. G. (2011). High efficiency and reliability of inter-simple sequence repeats (ISSR) markers for evaluation of genetic diversity in Brazilian cultivated *Jatropha curcas* L. accessions. *Molecular biology reports*, 38, 4245-4256.
- Hafizah, R. A., Adawiyah, R., Harahap, R. M., Hannum, S., & Santoso, P. J. (2018). Aplikasi marka SSR pada keanekaragaman genetik durian (*Durio zibethinus* Murr.) di Kabupaten Deli Serdang, Sumatra Utara. *Al-Kaunyah*, 11(1), 49-56.
- Hamrick, J. L., Godt, M. J. W., & Sherman-Broyles, S. L. (1992). Factors influencing levels of genetic diversity in woody plant species. In *Population Genetics of Forest Trees: Proceedings of the International Symposium on Population Genetics of*

Forest Trees Corvallis, Oregon, USA, July 31–August 2, 1990 (pp. 95-124). Springer Netherlands.

- Harahap, E.J., Rosmayati, dan D.S. Hanafiah. 2019. Uji polimorfik dan heterozigositas pada progeni F4 kedelai (*Glycine max* L.) Merrill) tahan salin dengan menggunakan marka SSR (Simple Sequence Repeat). *Jurnal Agrotek Lestari* 5(2) : 20-28.
- Hasnah, T.M. 2014. Keragaman genetik meranti (*Shorea leprosula* Miq.) asal Kalimantan dengan analisis isozim. *Jurnal penelitian dipterokarpa* 8(1) : 35-46.
- He, Y., Huang S., Xu G., Jiang O., Huang L., Sun C., Jin J., and Chen C. (2023). Structural characteristics and immunomodulation activity of a polysaccharide from purslane (*Portulaca oleracea*). *Journal of Functional Foods*. 109 (105781): 1-15.
- Hennink, S., & Zeven, A. C. (1990). The interpretation of Nei and Shannon-Weaver within population variation indices. *Euphytica*, 51, 235-240.
- Jia, S., Z. Yan, Y. Wang, Y. Wei, Z. Xie, and F. Zhang. (2017). Genetic diversity and relatedness among ornamental purslane (*Portulaca* L.) accessions unraveled by SRAP markers. *Biotech*. 7(241): 1-8. DOI 10.1007/s13205-017-0881-8.
- Kaundun, S. S., Zhyvoloup, A., & Park, Y. G. (2000). Evaluation of the genetic diversity among elite tea (*Camellia sinensis* var. *sinensis*) accessions using RAPD markers. *Euphytica*, 115, 7-16.
- Khade, Y.P., Sinhasane, S.R., Mainkar, P. *et al.* (2024). Exploring sequence-related amplified polymorphism (SRAP) markers for assessing genetic diversity in onion (*Allium cepa* L.) genotypes and their wild relatives. *Genet Resour Crop Evol.* <https://doi.org/10.1007/s10722-024-01978-5>
- Kinho, J., Na'iem, M., & Indrioko, S. (2016). Studi keragaman genetik *Diospyros rumphii* Bakh di Sulawesi Utara berdasarkan penanda isoenzim. *Jurnal Pemuliaan Tanaman Hutan*, 10(2), 95-109.
- Lamo, J.M., Rao, S.R. (2023). DNA markers in analysis of genetic diversity of *Curcuma longa* L. from Meghalaya. *Nucleus*. 66: 127–136. <https://doi.org/10.1007/s13237-023-00424-8>
- Lan, X., S. Guo, Y. Zhao, M. Zhang, D. Zhang, A. Leng, and X. Ying. (2023). A novel skeleton alkaloid from *Portulaca oleracea* L. and its bioactivities. *Fitoterapia*. 169(105608): 1-6.
- Li, C. M., Dong, H., Zhou, Q., & Goh, K. H. (2008). Biochips-fundamentals and applications. In *Electrochemical Sensors, Biosensors and their Biomedical Applications* (pp. 307-383).

- Li, G., & Quiros, C. F. (2001). Sequence-related amplified polymorphism (SRAP), a new marker system based on a simple PCR reaction: its application to mapping and gene tagging in Brassica. *Theoretical and applied genetics*, 103, 455-461.
- Li, J., Yang, X., & Mehri, S. (2021). Genetic diversity in *Stellaria* L.(Caryophyllaceae) using sequence related amplified polymorphism. *Genetika*, 53(3), 1369-1377.
- Li, S., Gan, X., Han, H., Zhang, X., & Tian, Z. (2018). Low within-population genetic diversity and high genetic differentiation among populations of the endangered plant *Tetracentron sinense* Oliver revealed by inter-simple sequence repeat analysis. *Annals of Forest Science*, 75, 1-11.
- Ma, M., Wang, T. & Lu, B. (2021). Assessment of genetic diversity in *Amomum tsao-ko* Crevost & Lemarié, an important medicine food homologous crop from Southwest China using SRAP and ISSR markers. *Genet Resour Crop Evol.* 68: 2655–2667. <https://doi.org/10.1007/s10722-021-01204-6>
- Majeed, H. O., Faraj, J. M., Rasul, K. S. *et al.* (2024). Evaluation of the genetic diversity and population structure of reticulated iris accessions in the Iraqi Kurdistan region using SCoT and SRAP markers. *Genet Resour Crop Evol.* <https://doi.org/10.1007/s10722-024-01884-w>
- Maloy, S., & Hughes, K. (Eds.). (2013). *Brenner's encyclopedia of genetics*. Academic Press.
- Manjunath, B. C., Chandrashekar, B. R., Mahesh, M., & Rani, R. V. (2011). DNA profiling and forensic dentistry—A review of the recent concepts and trends. *Journal of forensic and legal medicine*, 18(5), 191-197.
- Mansur, M., A.T.B.A. Mahmud, M.I.A. Dagong, L. Rahim, Rr S.R.A. Bugiwati, dan S. Baco. 2016. Keragaman genetik sapi bali di Kabupaten Barru berdasarkan karakteristik fenotipe dan DNA penciri mikrosatelit. *Jurnal Ilmu dan Teknologi Peternakan* 4(3) : 104-111.
- Marinho, M. A. O., G. Souza, L. P. Felix, abd R. de Carvalho. (2019). Comparative cytogenetics of the ACPT clade (Anacampserotaceae, Cactaceae, Portulacaceae, and Talinaceae): a very diverse group of the suborder Cactineae, Caryophyllales. *Protoplasma*. 256:805–814. <https://doi.org/10.1007/s00709-018-01334-2>.
- Misrianti, R., Sumantri, C., & Anggraeni, A. (2011). Keragaman gen hormon pertumbuhan reseptor (GHR) pada sapi perah Friesian Holstein. *Jity*, 16, 253-259.
- Mohammed, N.A., Afzal, M., Al-Faifi, S.A. *et al.* (2023). Effectiveness of sequence-related amplified polymorphism (SRAP) markers to assess the geographical origin and genetic diversity of collected lentil genotypes. *Plant Biotechnol Rep.* 17: 519–530. <https://doi.org/10.1007/s11816-023-00842-9>

- Mohamed, H. M., El-Assal, S. E. D. S. M., Gamal El-Din, A.. K. Y. *et al.* (2024). Analysis of genetic diversity and population structure in some Egyptian Berseem (*Trifolium alexandrinum*) accessions based on ISSR, SCoT and SRAP markers. *Genet Resour Crop Evol.* <https://doi.org/10.1007/s10722-024-01940-5>
- Monaghan, P. (2008). Early growth conditions, phenotypic development and environmental change. *Philosophical Transactions of the Royal Society B: Biological Sciences*, 363(1497), 1635-1645.
- Mo-Rong Xu, Meng-Shiou Lee, Bo-Cheng Yang, Hsiu-Chi Chang, Chao-Lin Kuo, Chia-Hsin Lin, Hsi-Jien Chen, Jai-Hong Cheng, Fang-Chun Sun. (2023). Development of a specific and sensitive diagnostic PCR for rapid molecular authentication of the medicinal plant *Portulaca oleracea*. *Molecular and Cellular Probes*. 67(101890): 1-8.
- Najaphy, A., Parchin, R. A., & Farshadfar, E. (2011). Evaluation of genetic diversity in wheat cultivars and breeding lines using inter simple sequence repeat markers. *Biotechnology & Biotechnological Equipment*, 25(4), 2634-2638.
- Obidike, N.. A., E. Victor, E. M. Obukohwo, O. P.. Ejiro, aand O. N. Blessing. (2024). *Portulaca oleraceae* leave extract enhanced renal function via enhancing urea and creatinine clearance in wistar rats. *European Journal of Medicineaal Chemistry Reports*. 11(200244).
- Obukohwo, O. M. (2024). Nutraceutical health benefit and safety utility of *Portulaca oleraceae*: A Review focus on Neuroooendocrine function. *Clinical Traditional Medicine and Pharmacology*. 200168.
- Ocampo, G. and J. T. Columbus. (2012). Molecular phylogenetics historical biogeography, and chromosome number evolution of *Portulaca* (Portulacaceae). *Molecular Phylogenetics and Evolution*. 63(1): 97-112.
- Panchariya, D. C., Dutta, P., Ananya, Mishra, A., Chawade, A., Nayee, N., ... & Kushwaha, S. K. (2024). Genetic marker: a genome mapping tool to decode genetic diversity of livestock animals. *Frontiers in Genetics*, 15, 1463474.
- Poczai, P., Varga, I., Laos, M., Cseh, A., Bell, N., Valkonen, J. P., & Hyvönen, J. (2013). Advances in plant gene-targeted and functional markers: a review. *Plant methods*, 9, 1-32.
- Prastowo, B. W., Rahardianti, R., Nur, E. M., & Taslihan, A. (2009). Profil Heterogenitas Genetik Induk Udang Windu (*Penaeus monodon*) Turunan F1 melalui Analisis DNA Mitokondria-RFLP dan RAPD. *Jurnal Perikanan Universitas Gadjah Mada*, 11(1), 25-30.
- Prevost, A., & Wilkinson, M. J. (1999). A new system of comparing PCR primers applied to ISSR fingerprinting of potato cultivars. *Theoretical and applied Genetics*, 98, 107-112.

- Priyono, B., & Abdullah, M. (2013). Keanekaragaman jenis kupu-kupu di Taman Kehati UNNES. *Biosaintifika: Journal of Biology & Biology Education*, 5(2), 100-105.
- Purnomo, E., and R.S. Ferniah. 2018. Polimorfisme cabai rawit dan cabai gendot dengan penanda RAPD (Random Amplified Polymorphic DNA) menggunakan primer OPA-8. *Berkala Bioteknologi*. 1(1) : 1-10.
- Putri, C. R., Kusumaningrum, H. P., & Kusdiyantini, E. (2016). Keragaman Genetik Jahe (*Zingiber officinale* Roscoe) menggunakan Teknik Penanda Molekuler Random Amplified Polymorphic DNA (RAPD). *Jurnal Akademika Biologi*, 5(2), 87-97.
- Raza, A., Farooq, A. B. U., Khan, W. A., Iqbal, A., Celik, S., Ali, M., & Khan, R. S. A. (2020). Polymorphic information and genetic diversity in Brassica species revealed by RAPD markers. *Biocell*, 44(4), 769.
- Restu, M., dan Pongtuluran, I. 2012. Seleksi primer untuk analisis keragaman genetik jenis bitti (*Vitex coffassus*). *Perennial*, 8(1): 25-29.
- Rianita, R., & Murni, P. (2023). Karakterisasi Morfologi dan Fenologi Pembungaan Krokot (*Portulaca oleracea* Linn.). *Biospecies*, 16(2), 54-62.
- Saini, M., Singh, S., Hussain, Z., & Yadav, A. (2010). RAPD analysis in mungbean [*Vigna radiata* (L.) Wilczek]: I. Assessment of genetic diversity.
- Sathyanarayana, S. H., & Wainman, L. M. (2024). Laboratory approaches in molecular pathology: the polymerase chain reaction. In *Diagnostic Molecular Pathology* (pp. 13-25). Academic Press.
- dos Santos, T. V. A., K. R. B. Leite, A. F. G. Trindade, *et al.* (2023). Inflorescence patterns within the genus *Portulaca* (Portulacaceae): analyses of anatomy and morphology, based on species occurring in Brazil. *Plant Syst Evol* **309**: 38. <https://doi.org/10.1007/s00606-023-01873-z>
- Sdouga, D., Kabtni, S., Hakim, L., Branca, F., Trifi-Farah, N., & Marghali, S. (2018). Integrated analysis for identifying *Portulaca oleracea* and its sub-species based on chloroplastic and nuclear DNA barcoding. *Plant Biosystems - An International Journal Dealing with All Aspects of Plant Biology*, 153(1), 25–31. <https://doi-org.ezproxy.ugm.ac.id/10.1080/11263504.2018.1435575>
- Serrote, C. M. L., Reiniger, L. R. S., Silva, K. B., dos Santos Rabaiolli, S. M., & Stefanel, C. M. (2020). Determining the polymorphism information content of a molecular marker. *Gene*, 726, 144175.
- Shamustakimova, A. O., Mavlyutov, Y. M., & Klimenko, I. A. (2021). Application of SRAP markers for DNA identification of Russian Alfalfa Cultivars. *Russian Journal of Genetics*, 57(5), 540-547.

- Sinaga, A., Putri, L. A. P., & Bangun, M. K. (2017). Analisis Pola Pita Andaliman (*Zanthoxylum Acanthopodium* DC) Berdasarkan Primer OPD 03, OPD 20, OPC 07, OPM 20, OPN 09. *Jurnal Agroekoteknologi Universitas Sumatera Utara*, 5(1), 109236.
- Solin, N.W.N.M., Sobir, dan N. Touran-Mathius. 2013. Keragaman genetik aksesori tetua saudara kandung (sibs) kelapa sawit Sura Deli berdasarkan penanda DNA mikrosatelit. *Buletin Palma* 14(2) : 100-108.
- Sopandi, A., & Herwanto, F. (2020). Eksplorasi dan karakterisasi morfologi tanaman kopi robusta (*Coffea robusta* L.) di dataran medium kecamatan lembah masurai kabupaten merangin. *Jurnal Sains Agro*, 5(2).
- Singh, N., Pal, A.K., Roy, R.K. *et al.* (2016) Assessment of genetic variation and population structure in Indian *Gladiolus* cultivars inferred from molecular markers. *Nucleus*.59: 235–244. <https://doi.org/10.1007/s13237-016-0181-4>
- Sivaramakrishna, P., & Yugandhar, P. (2020). A new species of the genus *Portulaca* L.(Portulacaceae) from the Eastern Ghats, India. *Journal of Asia-Pacific Biodiversity*, 13(4), 755-761.
- Stephenson, F. H. (2010). *Calculations for molecular biology and biotechnology: a guide to mathematics in the laboratory*. Academic press.
- Sukartini, S. (2007). Pengelompokan Aksesori Pisang Menggunakan Karakter Morfologi IPGRI. *Jurnal Hortikultura*, 17(1), 83642.
- Sulistyawati, P., & Widyatmoko, A. Y. P. B. C. (2017). Keragaman Genetik Aksesori Kayu Merah (*Pterocarpus Indicus* Willd) Menggunakan Penanda Random Amplified Polymorphism Dna. *Jurnal Pemuliaan Tanaman Hutan*, 11(1), 67-76.
- Sulistiyo, R. H., Soetopo, L., & Damanhuri, D. (2015). Eksplorasi dan identifikasi karakter morfologi porang (*Amorphophallus muelleri* B.) di Jawa Timur (Doctoral dissertation, Brawijaya University).
- Strong, W. L. (2016). Biased richness and evenness relationships within Shannon–Wiener index values. *Ecological indicators*, 67, 703-713.
- Terryana, R.T., K. Nugroho, H. Rijzaani, dan P. Lestari. 2018. Karakterisasi keragaman genetik 27 genotipe cabai berdasarkan marka SSR (Simple Sequence Repeat). *Jurnal Ilmu-Ilmu Hayati* 17(2) : 183-194.
- Tonk, F. A., Tosun, M., Ilker, E., Istipliler, D., & Tatar, O. (2014). Evaluation and comparison of ISSR and RAPD markers for assessment of genetic diversity in triticale genotypes. *Bulgarian Journal of Agricultural Science*, 20(6), 1413-1420.
- Tuomisto, H. (2012). An updated consumer's guide to evenness and related indices. *Oikos*, 121(8), 1203-1218.

- Vinodhini, R.T., Ashwin, N.M.R., Agisha, V.N. *et al.* (2023). Discovering The Molecular Variations Among Distinct *Sporisorium scitamineum* Isolates Using Sequence-Related Amplified Polymorphism (SRAP) Markers. *Sugar Tech* **25**, 1144–1151. <https://doi.org/10.1007/s12355-023-01276-x>
- Wang, G., Pan, J., Li, X., He, H., Wu, A., & Cai, R. (2005). Construction of a cucumber genetic linkage map with SRAP markers and location of the genes for lateral branch traits. *Science in China Series C: Life Sciences*, *48*, 213-220.
- Wei, W., Gao, Y., Wang, H., Cao, Y., & Li, J. (2024). Combined application of gasification filter cake and *Portulaca oleracea* to promote soil quality and tomato yields under irrigation with brackish water. *Horticultural Plant Journal*.
- Williams, J. G., Kubelik, A. R., Livak, K. J., Rafalski, J. A., & Tingey, S. V. (1990). DNA polymorphisms amplified by arbitrary primers are useful as genetic markers. *Nucleic acids research*, *18*(22), 6531-6535.
- Windarti, W. (2009). Keanekaragaman genetika pisang bergenom B berdasarkan penanda mikrosatelit. *Biosaintifika: Journal of Biology & Biology Education*, *1*(1).
- Zhang, Z., Yang, Q., Niu, Y. *et al.* (2021). Diversity analysis and establishment of core collection among *Akebia trifoliata* (Thunb.) Koidz. in Qinba mountain area of China using ISSR and SRAP markers. *Genet Resour Crop Evol.* **68**: 1085–1102 . <https://doi.org/10.1007/s10722-020-01051-x>
- Zheng, Y., Xu, S., Liu, J., Zhao, Y., & Liu, J. (2017). Genetic diversity and population structure of Chinese natural bermudagrass [*Cynodon dactylon* (L.) Pers.] germplasm based on SRAP markers. *PLoS One*, *12*(5), e0177508.
- Zhou, L., Yarra, R., Cao, H. *et al.* (2021). Sequence-Related Amplified Polymorphism (SRAP) Markers Based Genetic Diversity and Population Structure Analysis of Oil Palm (*Elaeis guineensis* Jacq.). *Tropical Plant Biol.* **14**: 63–71 . <https://doi.org/10.1007/s12042-020-09273-0>