

DAFTAR PUSTAKA

- Ali, A., M. U. Rehman, S. M. Ahmad, T. Mehraj, I. Hussain, A. Nadeem, M. U. R. Mir, and S. A. Ganie. 2022. In Silico Tools for Analysis of Single-Nucleotide Polymorphisms in the Bovine Transferrin Gene. *Animals*. 12(6): 1-16. <https://doi.org/10.3390/ani12060693>.
- Andrés, J., K. Maisey, J. M. Yáñez, H. Valenzuela, P. A. Cea, L. Tort, A. M. Sandino, and M. Imarai. 2020. Single-Nucleotide Polymorphisms (SNPs) Mining and Their Effect on the Tridimensional Protein Structure Prediction in a Set of Immunity-Related Expressed Sequence Tags (EST) in Atlantic Salmon (*Salmo salar*). *Frontiers in Genetics*. 10. <https://doi.org/10.3389/fgene.2019.01406>.
- Aoyama, T., T. Kobayashi, M. Takahashi, S. Nagasaka, K. Usuda, Y. Kakei, Y. Ishimaru, H. Nakanishi, S. Mori, and N. K. Nishizawa. 2009. OsYSL18 is a rice iron (III)–deoxymugineic acid transporter specifically expressed in reproductive organs and phloem of lamina joints. *Plant molecular biology*. 70 (6): 681–692. <https://doi.org/10.1007/s11103-009-9500-3>.
- Azis, A., M. Izzati, S. Haryanti. 2015. Aktivitas Antioksidan dan Nilai Gizi dari Beberapa Jenis Beras dan Millet sebagai Bahan Pangan Fungsional Indonesia. *Jurnal Biologi*. 4(1): 45-61.
- Balk, J., and T. A. Schaedler. 2014. Iron cofactor assembly in plants. *Annual review of plant biology*. 65: 125–153. <https://doi.org/10.1146/annurev-arplant-050213-035759>.
- Bashir, K., H. Inoue, S. Nagasaka, M. Takahashi, H. Nakanishi, S. Mori, and N. K. Nishizawa. 2006. Cloning and characterization of deoxymugineic acid synthase genes from graminaceous plants. *The Journal of biological chemistry*. 281(43):32395–32402. <https://doi.org/10.1074/jbc.M604133200>
- Bashir, K., Y. Ishimaru, and N.K. Nishizawa. 2010. Iron Uptake and Loading into Rice Grains. *Rice*. 3: 122–130. <https://doi-org.ezproxy.ugm.ac.id/10.1007/s12284-010-9042-y>.
- Budiman, H., F. S. H. Krismastuti, and Nuryatini. 2009. Estimation of Measurement Uncertainty in The Determination of Fe Content in Powdered Tonic Food Drink Using Graphite Furnace Atomic Absorption Spectrometry. *Indo. J. Chem*. 9 (2): 247–253.
- Butler, J.M. 2012. *Advanced Topics in Forensic DNA Typing: Methodology*. Academic Press. Maryland. p. 347. <https://doi.org/10.1016/B978-0-12-374513-2.00012-9>.
- Carrillo, J. T. and D. Borthakur. 2021. Methods for metal chelation in plant homeostasis: Review. *Plant Physiology and Biochemistry*. 163: 95–107. <https://doi.org/10.1016/J.PLAPHY.2021.03.045>.
- Chen, G., J. Li, H. Han, R. Du, and X. Wang. 2022. Physiological and Molecular Mechanisms of Plant Responses to Copper Stress. *International Journal of Molecular Sciences*. 23(21):12950. <https://doi.org/10.3390/ijms232112950>.
- Chu, H. H., J. Chiecko, T. Punshon, A. Lanzirrotti, B. Lahner, D. E. Salt, and E. L. Walker. 2010. Successful reproduction requires the function of *Arabidopsis* *Yellow Stripe-Like1* and *Yellow Stripe-Like3* metal-nicotianamine

- transporters in both vegetative and reproductive structures. *Plant physiology*. 154(1): 197–210. <https://doi.org/10.1104/pp.110.159103>.
- Conte, S. S., H. H. Chu, D. C. Rodriguez, T. Punshon, , K. A. Vasques, D. E. Salt, and E. L. Walker. 2013. *Arabidopsis thaliana Yellow Stripe1-Like4 and Yellow Stripe1-Like6* localize to internal cellular membranes and are involved in metal ion homeostasis. *Frontiers in plant science*. 4(283): 1-15. <https://doi.org/10.3389/fpls.2013.00283>.
- Dayanti, F. Ghina, A. Djuminar, A. Dermawan, dan A. Tantan. 2019. Perbandingan Nilai Pengukuran Kuantitatif Hasil Ekstraksi DNA *Salmonella typhi* Menggunakan Metode Boiling, NaOH, Kit Komersial. *Jurnal Riset Kesehatan Poltekkes Depkes Bandung*. 11(1). 350–357.
- Dewanata, P. A., dan M. Mushlih. 2021. Differences in DNA Purity Test Using UV-Vis Spectrophotometer and Nanodrop Spectrophotometer in Type 2 Diabetes Mellitus Patients. *Indonesian Journal of Innovation Studies*. 15:1-10. <https://doi.org/10.21070/ijins.v15i.553>.
- Dey, S., P. Regon, S. Kar, and S. K. Panda. 2020. Chelators of iron and their role in plant's iron management. *Physiology and Molecular Biology of Plants*. 26(8): 1541–1549. <https://doi.org/10.1007/s12298-020-00841-y>.
- DiDonato, R. J., Jr, L. A. Roberts, T. Sanderson, R. B. Eisley, & E. L. Walker. 2004. *Arabidopsis Yellow Stripe-Like2 (YSL2)*: a metal-regulated gene encoding a plasma membrane transporter of nicotianamine-metal complexes. *The Plant journal: for cell and molecular biology*. 39(3): 403–414. <https://doi.org/10.1111/j.1365-313X.2004.02128.x>
- Fahma, R. 2022. Deteksi *Single Nucleotide Polymorphisms (SNPs)* Gen *OsFER2* Pada Lima Kultivar Padi Putih Dan Berpigmen (*Oryza sativa* L.). Skripsi. Universitas Gadjah Mada, Yogyakarta.
- Gomes, A. and B. Korf. 2018. Genetic Testing Techniques. *Pediatric Cancer Genetics*. 47–64. <https://doi.org/10.1016/B978-0-323-48555-5.00005-3>.
- Grimm, B., Y. A. Purwestri, T. R. Nuringtyas, F. A. Susanto, T. Fan, and L. Roling, 2019. Basic Methods in Plant Molecular Biology and Plant Physiology. Universitas Gadjah Mada Yogyakarta-Humboldt University Berlin.
- Handiwirawan, E., R. R. Noor, C. Sumantri, Subandriyo, dan I. Inounu. 2012. Identifikasi *Single Nucleotide Polymorphism (SNP)* pada Gen *Mono Amine Oxidase A (MAO-A)* sebagai Penanda Genetik untuk Sifat Agresif pada Domba. *JITV*. 17 (4): 258-275.
- Herman, Y. P. Ardila, F. Nugraha, dan D. I. Roslim. 2014. Analisis Sekuen Intron 1 sampai Sebagian Ekson 4 dari Gen *Feritin2* pada Tiga Genotipe Padi (*Oryza sativa* L.) Lokal Indragiri Hilir, Riau. XXIX: 21–26. <https://journal.uir.ac.id/index.php/dinamikapertanian/article/view/856>.
- Hernawan, E. dan V. Meylani. 2016. Analisis Karakteristik Fisikokimia Beras Putih, Beras Merah, Dan Beras Hitam (*Oryza sativa* L., *Oryza nivara* dan *Oryza sativa* L. *indica*). *Jurnal Kesehatan Bakti Tunas Husada*. 15(1): 79-91.
- Horiike, T. 2016. An Introduction to Molecular Phylogenetic Analysis. *Reviews in Agricultural Science*. 4: 36-45. Doi: 10.7831/ras.4.36
- Inoue, H., T. Kobayashi, T. Nozoye, M. Takahashi, Y. Kakei, K. Suzuki, M. Nakazono, H. Nakanishi, S. Mori, and N. K. Nishizawa. 2009. Rice *OsYSL15* is an iron-regulated iron (III)-deoxymugineic acid transporter expressed in the roots and is essential for iron uptake in early growth of the seedlings. *The*

- Journal of biological chemistry*. 284(6): 3470–3479.
<https://doi.org/10.1074/jbc.M806042200>.
- Ishimaru, Y., M. Suzuki, T. Tsukamoto, K. Suzuki, M. Nakazono, T. Kobayashi, Y. Wada, S. Watanabe, S. Matsuhashi, M. Takahashi, H. Nakanishi, S. Mori, and N. K. Nishizawa. 2006. Rice plants take up iron as an Fe³⁺-phytosiderophore and as Fe²⁺. *The Plant journal: for cell and molecular biology*. 45(3). 335–346. <https://doi.org/10.1111/j.1365-313X.2005.02624.x>.
- Ishimaru Y, Masuda H, Bashir K, Inoue H, Tsukamoto T, Takahashi M, Nakanishi H, Aoki N, Hirose T, Ohsugi R, Nishizawa, N.K. 2010. Rice metal-nicotianamine transporter, *OsYSL2*, is required for the long-distance transport of iron and manganese. *Plant J*. 62:379–390.
- Jin, Y., S. Liu, Z. Yuan, Y. Yang, S. Tan, and Z. Liu. 2016. *Genomics in Aquaculture*. Academic Press. Alabama. p.73. <https://doi.org/10.1016/B978-0-12-801418-9.00004-4>.
- Juliano, B.O. and A.P.P. Tuano. 2019. *Rice: Gross structure and composition of the rice grain*. Elsevier. Los Baños. p:31. DOI: 10.1016/B978-0-12-811508-4.00002-2.
- Kobayashi, T. and N. K. Nishizawa. 2012. Iron uptake, translocation, and regulation in higher plants. *Annual review of plant biology*. 63: 131–152. <https://doi.org/10.1146/annurev-arplant-042811-105522>.
- Koike, S., Inoue, H., Mizuno, D., Takahashi, M., Nakanishi, H., Mori, S., and Nishizawa, N.K. 2004. *OsYSL2* is a rice metal-nicotianamine transporter that is regulated by iron and expressed in the phloem. *Plant J*. 39:415-424.
- Kumari, P., P. Kumar, and VK. Sharma. 2018. Candidate gene markers based preliminary genetic analysis in relation to grain zinc and iron accumulation in rice varieties. *Journal of Pharmacognosy and Phytochemistry*. 7(4): 1300-1305. 10.13140/RG.2.2.15829.32484.
- Kurnianingsih, N. R., R. Susandarini, F. A. Susanto, T. R. Nuringtyas, G. Jenkins, and Y. A. Purwestri. 2019. Characterization of Indonesian pigmented rice (*Oryza sativa*) based on morphology and single nucleotide polymorphism. *Biodiversitas*. 20(4):1208-1214.
- Kristamtini, Taryono, P. Basunanda, dan R. H. Murti. 2016. Keragaman Genetik Kultivar Padi Beras Hitam Lokal Berdasarkan Penanda Mikrosatelit. *Jurnal AgroBiogen*. 10(2):69-76.
- Liu, P.P. 2022. Talking Glossary of Genomic and Genetic Terms. <https://www.genome.gov/genetics-glossary>. Diakses tanggal 04 Januari 2023.
- Liu, T., J. Zeng, K. Xia, T. Fan, Y. Li, Y. Wang, X. Xu, and M. Zhang. 2012. Evolutionary expansion and functional diversification of oligopeptide transporter gene family in rice. *Rice*. 5(12):1–14. <https://doi.org/10.1186/1939-8433-5-12>.
- Lorenz T. C. 2012. Polymerase chain reaction: basic protocol plus troubleshooting and optimization strategies. *Journal of visualized experiments : JoVE*. 63: e3998. <https://doi.org/10.3791/3998>.
- Mahendrakar, M. D., M. Parveda, P. B. K. Kishor, and R. K. Srivastava. 2020. Discovery and validation of candidate genes for grain iron and zinc metabolism in pearl millet [*Pennisetum glaucum* (L.) R. Br.]. *Scientific reports*. 10(1). 1-16. <https://doi.org/10.1038/s41598-020-73241-7>.

- Meng, F., Y. Wei, and X. Yang. 2005. Iron content and bioavailability in rice. *Journal of Trace Elements in Medicine and Biology*. 18: 333–338.
- Mouza, Z. 2020. Deteksi Gen *Ferritin OsFER1* dan *OsFER2* Pada Padi Hitam (*Oryza sativa* L. ‘Cempo Ireng’). Skripsi. Universitas Gadjah Mada, Yogyakarta.
- Mowidu, I., H.S. Bambang, H.P. Benito, dan N.H.U. Sri. 2015. Kadar Fe Total Pada Tanah Sawah Rawa Lebak di Kabupaten Poso. *Jurnal AgroPet*. 12(1):7-11.
- Nozoye, T., H. Inoue, M. Takahashi, Y. Ishimaru, H. Nakanishi, S. Mori, and N. K. Nishizawa. 2007. The expression of iron homeostasis-related genes during rice germination. *Plant molecular biology*. 64(1-2): 35–47. <https://doi.org/10.1007/s11103-007-9132-4>.
- Palmgren, M. G., S. Clemens, L. E. Williams, U. Krämer, S. Borg, J. K. Schjørring, and D. Sanders. 2008. Zinc biofortification of cereals: problems and solutions. *Trends in Plant Science*. 13(9): 464–473. <https://doi.org/10.1016/j.tplants.2008.06.005>.
- Panda, B., S. Sharma, P. Mohapatra, A. and Das. 2014. Iron Nutrition *vis-à-vis* Aconitase Activity and Ferritin Accumulation in Tropical *Indica* Rice Cultivars Differing in Grain Iron Concentration. *American Journal of Plant Sciences*. 5: 2829-2841. doi: [10.4236/ajps.2014.518299](https://doi.org/10.4236/ajps.2014.518299).
- Pratiwi, A. 2022. Deteksi *Single Nucleotide Polymorphisms (SNPs)* Gen *OsFER1* Pada Lima Kultivar Padi Putih Dan Berpigmen (*Oryza sativa* L.). Skripsi. Universitas Gadjah Mada, Yogyakarta.
- Pratiwi, R. and Y. A. Purwestri. 2017. Black Rice as a Functional Food in Indonesia. *Functional Foods in Health and Disease*. 7(3): 182-194. DOI: 10.31989/ffhd.v7i3.310.
- Putri, A. dan S. Wathon. 2018. Aplikasi *Single Nucleotide Polymorphism (SNP)* dalam Studi Farmakogenomik untuk Pengembangan Obat. *BioTrends*. 9(2): 69-74.
- Puig, S., L.R. Alonso, A.M. Romero, and M.T. Martínez-Pastor. 2017. The elemental role of iron in DNA synthesis and repair. *Metallomics*. 9(11):1483–1500. <https://doi.org/10.1039/c7mt00116a>
- Rajalakshmi, S. 2017. International Journal of Pharmaceutical, Chemical and Biological Sciences Different Types of PCR Techniques and Its Applications. *IJPCBS*. 7(3): 285–292. www.ijpcbs.com.
- Reuter, M., Y. Küpper, A. Schmitz, J. P. Breuer, U. Wend, and J. Hennig. 2005. Detection of new single nucleotide polymorphisms by means of real time PCR. *Journal of Genetics*. 84 (3): 341-345.
- Robert, F. and J. Pelletier. 2018. Exploring the Impact of Single-Nucleotide Polymorphisms on Translation. *Front. Genet*. 9:1-11. doi:10.3389/fgene.2018.00507.
- Rout, G. R., and S. Sahoo. 2015. Role of iron in plant growth and metabolism. *Rev. Agric. Sci*. 3: 1–24. doi: 10.7831/ras.3.1.
- Sahu, P. K., S. Mondal, D. Sharma, G. Vishwakarma, V. Kumar, and B. K. Das. 2017. InDel marker based genetic differentiation and genetic diversity in traditional rice (*Oryza sativa* L.) landraces of Chhattisgarh, India. *PLoS ONE*. 12(11). <https://doi.org/10.1371/journal.pone.0188864>.

- Sotomayor-Vivas, C., E. Hernández-Lemus, and R. Dorantes-Gilardi. 2022. Linking protein structural and functional change to mutation using amino acid networks. *PloS one*. 17(1):1-23. doi.org/10.1371/journal.pone.0261829.
- Sudiarta, W., O. Ratnayani, dan A.K. Veliyana. 2019. Analisis Kadar Logam Besi dalam Susu Bubuk Formula Kehamilan secara Spektrofotometri Serapan Atom. *J. Media Sains*. 3 (1): 1-6.
- Sulartini, N.W.S., G.R. Sadimantara, T. Wijayanto, dan Muhidin. 2011. Pengujian kadar antosianin padi gogo beras merah hasil koleksi plasma nutfah sulawesi tenggara. *J Crop Agro*. 4(2): 43-48.
- Tasma, I. M. 2015. Pemanfaatan Teknologi Sekuensing Genom untuk Mempercepat Program Pemuliaan Tanaman. *J. Litbang Pert*. 32(2). 159–168.
- Takahashi, M., Y. Terada, I. Nakai, H. Nakanishi, E. Yoshimura, S. Mori, and N. K. Nishizawa. 2003. Role of nicotianamine in the intracellular delivery of metals and plant reproductive development. *The Plant cell*. 15(6): 1263–1280. <https://doi.org/10.1105/tpc.010256>.
- Thomson, M.J., E.M. Septingsih, F. Suwardjo, T.I. Santoso, T.S. Sintenga, and S.R. McCouch. 2007. Genetic Diversity Analysis of Traditional and Improved Indonesian Rice (*Oryza sativa* L.) Germplasm Using Microsatellite Markers. *Theor Appl Genet*. 114(3): 559-568.
- Tripathi, K. K., O. P. Govila., R. Warriar and V. Ahuja. 2011. *Biology of Oryza sativa* L. (*rice*). Department of Biotechnology, Ministry of Sciences & Technology, Government of India. New Delhi.
- Utama, Z.H. 2015. *Budidaya Padi pada Lahan Marjinal: Kiat Meningkatkan Produksi Padi*. Penerbit ANDI. Yogyakarta. pp.10-12.
- Wang, M., W. Gruissem, and N. K. Bhullar. 2013. Nicotianamine synthase overexpression positively modulates iron homeostasis-related genes in high iron rice. *Front. Plant Sci*. 4(1):1-15. <https://doi.org/10.3389/fpls.2013.00156>.
- Wang, Q., M. Chen, Q. Hao, H. Zeng, He. 2021. Research and Progress on the Mechanism of Iron Transfer and Accumulation in Rice Grains. *Plants*. 10:1-18. <https://doi.org/10.3390/plants10122610>.
- Wen-yue, C., C. Hai-rui, B. Jin-song , Z. Xiang-sheng, and S. Qing-yao. 2006. A Simplified Rice DNA Extraction Protocol for PCR Analysis. *Rice Science*. 13(1): 67-70. <http://www.ricescience.org>.
- Wu, D., G. He, W. Tian, M. Saleem, D. Li, Y. Huang, L. Meng, Y. He, Y. Liu, and T. He. 2021. OPT gene family analysis of potato (*Solanum tuberosum*) responding to heavy metal stress: Comparative omics and co-expression networks revealed the underlying core templates and specific response patterns. *International Journal of Biological Macromolecules*. 188. 892–903. <https://doi.org/10.1016/j.ijbiomac.2021.07.183>.
- Yen, M. R., Y. H. Tseng, and M. H. Saier Jr. 2001. *Maize Yellow Stripe1*, an iron-phytosiderophore uptake transporter, is a member of the *oligopeptide transporter (OPT)* family. *Microbiology (Reading, England)*. 147(11): 2881–2883. <https://doi.org/10.1099/00221287-147-11-2881>.
- Yusuf, Z.K. 2010. Polymerase Chain Reaction (PCR). *Saintek*. 5 (6): 1-6.
- Zhang, X., D. Zhang, W. Sun, and T. Wang. 2019. The adaptive mechanism of plants to iron deficiency via iron uptake, transport, and homeostasis.

International Journal of Molecular Sciences. 20(10): 1–14.
<https://doi.org/10.3390/ijms20102424>.

Zhang C, K.I. Shinwari, L. Luo, and L. Zheng. 2018a. *OsYSL13* Is Involved in Iron Distribution in Rice. *International Journal of Molecular Sciences*. 19(11):3537. <https://doi.org/10.3390/ijms19113537>.

Zhang, C., W. Lu, Y. Yang, Z. Shen, J.F. Ma, and L. Zheng. 2018b. *OsYSL16* is Required for Preferential Cu Distribution to Floral Organs in Rice. *Plant and Cell Physiology*. 59 (10):2039–2051. <https://doi.org/10.1093/pcp/pcy124>.

Zheng, L., M. Fujii, N. Yamaji, A. Sasaki, M. Yamane, I. Sakurai, K. Sato, and J.F. Ma. 2011. Isolation and Characterization of a Barley Yellow Stripe-Like Gene, *HvYSL5*. *Plant and Cell Physiology*. 52(5): 765–774. <https://doi.org/10.1093/pcp/pcr009>.