

DAFTAR PUSTAKA

- Amack, S. C., & Antunes, M. S. (2020). CaMV35S Promoter – A Plant Biology and Biotechnology Workhorse in The Era of Synthetic Biology. *Current Plant Biology*, 24, 100179. <https://doi.org/10.1016/j.cpb.2020.100179>
- Aoyama, T., & Chua, N.-H. (1997). A Glucocorticoid-Mediated Transcriptional Induction System in Transgenic Plants. *The Plant Journal*, 11(3), 605–612.
- Apridamayanti, P., Pratiwi, R., Purwestri, Y. A., Tunjung, W. A. S., & Rumiya. (2017). Anthocyanin, Nutrient Contents, and Antioxidant Activity of Black Rice Bran of *Oryza sativa* L. 'Cempo Ireng' from Sleman, Yogyakarta, Indonesia. *Indonesian Journal of Biotechnology*, 22(1), 49–54.
- Chardin, C., Girin, T., Roudier, F., Meyer, C., & Krapp, A. (2014). The Plant RWP-RK Transcription Factors: Key Regulators of Nitrogen Responses and of Gametophyte Development. *Journal of Experimental Botany*, 65(19), 5577–5587. <https://doi.org/10.1093/jxb/eru261>
- Chatterjee, J., Dionora, J., Elmido-mabilangan, A., Wanchana, S., Thakur, V., Bandyopadhyay, A., Brar, D. S., & Quick, W. P. (2016). The Evolutionary Basis of Naturally Diverse Rice Leaves Anatomy. *PLoS ONE*, 11, 1–25. <https://doi.org/10.1371/journal.pone.0164532>
- Chongloi, G. L., Prakash, S., & Vijayraghavan, U. (2019). Regulation of Meristem Maintenance and Organ Identity During Rice Reproductive Development. *Journal of Experimental Botany*, 70(6), 1719–1736. <https://doi.org/10.1093/jxb/erz046>
- Counce, P. A., Keisling, T. C., & Mitchell, A. J. (2000). A Uniform, Objectives, and Adaptive System for Expressing Rice Development. *Crop Science*, 40(2), 436–443. <https://doi.org/10.2135/cropsci2000.402436x>
- Fan, M.-J., Wang, I.-C., Hsiao, Y., Lin, H., Tang, N.-Y., Hung, T.-C., Quan, C., Lien, J.-C., & Chung, J.-G. (2015). Anthocyanins from Black Rice (*Oryza sativa* L.) Demonstrate Antimetastatic Properties by Reducing MMPs and NF- κ B Expressions in Human Oral Cancer CAL 27 Cells. *Nutrition and Cancer*, 67(2), 327–338. <https://doi.org/10.1080/01635581.2015.990576>
- Fauzia, A. N. (2017). *Ekspresi Gen OsRDK4 Pada Padi Hitam Transgenik (Oryza sativa L. 'Cempo Ireng') Untuk Induksi Embriogenesis Somatik*. Tesis. Universitas Gadjah Mada.
- Fuchs, M., & Lohmann, J. U. (2020). Aiming for The Top: Non-Cell Autonomous Control of Shoot Stem Cells in *Arabidopsis*. *Journal of Plant Research*, 133, 297–309. <https://doi.org/10.1007/s10265-020-01174-3>
- Galli, M., & Gallavotti, A. (2016). Expanding the Regulatory Network for Meristem Size in Plants. *Trends in Genetics*, 32(6), 372–383. <https://doi.org/10.1016/j.tig.2016.04.001>
- Gao, H., Wang, W., Wang, Y., & Liang, Y. (2019). Molecular Mechanisms Underlying Plant Architecture and Its Environmental Plasticity in Rice. *Molecular Breeding*, 39(12). <https://doi.org/10.1007/s11032-019-1076-2>
- Gou, J., Fu, C., Liu, S., Tang, C., Debnath, S., Flanagan, A., Ge, Y., Tang, Y., Jiang, Q., Larson, P. R., Wen, J., & Wang, Z. Y. (2017). The miR156-SPL4 Module Predominantly Regulates Aerial Axillary Bud Formation and Controls Shoot

- Architecture. *New Phytologist*, 216(3), 829–840.
<https://doi.org/10.1111/nph.14758>
- He, W., & Zhang, X.-S. (2003). Responses of An Evergreen Shrub *Sabina vulgaris* to Soil Water and Nutrient Shortages in The Semi-Arid Mu Us Sandland in China. *Journal of Arid Environments*, 53(3), 307–316.
<https://doi.org/10.1006/jare.2002.1051>
- IRRI. (2013). *Standard Evaluation System for Rice*. International Rice Research Institute. Philippines. http://www.clrri.org/ver2/uploads/SES_5th_edition.pdf
- ITIS. (2022). *ITIS - Report: Oryza sativa*.
https://www.itis.gov/servlet/SingleRpt/SingleRpt?search_topic=TSN&search_value=41976#null
- Itoh, J.-I., Nonomura, K.-I., Ikeda, K., Yamaki, S., Inukai, Y., Yamagishi, H., Kitano, H., & Nagato, Y. (2005). Rice Plant Development: from Zygote to Spikelet. *Plant and Cell Physiology*, 46(1), 23–47.
<https://doi.org/10.1093/pcp/pci501>
- Jennings, P. R., Coffman, W. R., & Kauffman, H. E. (1979). *Rice Improvement*. International Rice Research Institute. Philippines.
https://doi.org/10.1007/978-1-4614-7903-1_16
- Jeong, S., Palmer, T. M., & Lukowitz, W. (2011). The RWP-RK Factor GROUNDED Promotes Embryonic Polarity by Facilitating YODA MAP Kinase Signaling. *Current Biology*, 21(15), 1268–1276.
<https://doi.org/10.1016/j.cub.2011.06.049>
- Jiajia, W., Jing, X., Qian, Q., & Guangheng, Z. (2020). Development of Rice Leaves: How Histocytes Modulate Leaf Polarity Establishment. *Rice Science*, 27(6), 468–479. <https://doi.org/10.1016/j.rsci.2020.09.004>
- Jiao, Y., Wang, Y., Xue, D., Wang, J., Yan, M., Liu, G., Dong, G., Zeng, D., Lu, Z., Zhu, X., Qian, Q., & Li, J. (2010). Regulation of OsSPL14 by OsMIR156 Defines Ideal Plant Architecture in Rice. *Nature Genetics*, 42(6), 541–544.
<https://doi.org/10.1038/ng.591>
- Kebrom, T. H., Burson, B. L., & Finlayson, S. A. (2006). Phytochrome B Represses Teosinte Branched1 Expression and Induces Sorghum Axillary Bud Outgrowth in Response to Light Signals. *Plant Physiology*, 140(3), 1109–1117. <https://doi.org/10.1104/pp.105.074856>
- Kebrom, T. H., Spielmeier, W., & Finnegan, E. J. (2013). Grasses Provide New Insights Into Regulation of Shoot Branching. *Trends in Plant Science*, 18(1), 41–48. <https://doi.org/10.1016/j.tplants.2012.07.001>
- Köszei, D., Johnston, A. J., Rutten, T., Czihal, A., Altschmied, L., Kumlehn, J., Wüst, S. E. J., Kirioukhova, O., Gheyselinck, J., Grossniklaus, U., & Bäuml, H. (2011). Members of The RKD Transcription Factor Family Induce An Egg Cell-Like Gene Expression Program. *Plant Journal*, 67(2), 280–291. <https://doi.org/10.1111/j.1365-313X.2011.04592.x>
- Kristantini, K., Sutarno, S., Wiranti, E. W., & Widyayanti, S. (2016). Kemajuan Genetik dan Heritabilitas Karakter Agronomi Padi Beras Hitam pada Populasi F2. *Jurnal Penelitian Pertanian Tanaman Pangan*, 35(2), 119.
<https://doi.org/10.21082/jpntp.v35n2.2016.p119-124>
- Kristantini, K., & Wiranti, E. W. (2017a). Clustering of 18 Local Black Rice Base

- on Total Anthocyanin. *Biology, Medicine, & Natural Product Chemistry*, 6(2), 47. <https://doi.org/10.14421/biomedich.2017.62.47-51>
- Kristamtini, K., & Wiranti, E. W. (2017b). Clustering of 18 Local Black Rice Base on Total Anthocyanin. *Biology, Medicine, & Natural Product Chemistry*, 6(2), 47–51. <https://doi.org/10.14421/biomedich.2017.62.47-51>
- Kristamtini, Taryono, Basunanda, P., Murti, R. H., & Supriyanta. (2012). Morphological of Genetic Relationships Among Black Rice Landraces From Yogyakarta and Surrounding Areas. *ARPN Journal of Agricultural and Biological Science*, 7(12), 982–989.
- Kumar, A., Batra, R., Gahlaut, V., Gautam, T., Kumar, S., Sharma, M., Tyagi, S., Singh, K. P., Balyan, H. S., Pandey, R., & Gupta, P. K. (2018). Genome-Wide Identification and Characterization of Gene Family for RWP-RK Transcription Factors in Wheat (*Triticum aestivum* L.). *PLoS ONE*, 13(12), 1–28. <https://doi.org/10.1371/journal.pone.0208409>
- Kushwaha, U. K. S. (2016). *Black Rice: Research, History and Development*. Springer International Publishing Switzerland. <https://doi.org/10.4159/9780674262539>
- Leegood, R. C. (2008). Roles of The Bundle Sheath Cells in Leaves of C3 Plants. *Journal of Experimental Botany*, 59(7), 1663–1673. <https://doi.org/10.1093/jxb/erm335>
- Li, G., Chen, X., Zhou, C., Yang, Z., Zhang, C., Huang, Z., Pan, W., & Xu, K. (2022). Vascular Bundle Characteristics of Different Rice Variety Treated with Nitrogen Fertilizers and Its Relation to Stem Assimilates Allocation and Grain Yield. *Agriculture*, 12(6). <https://doi.org/10.3390/agriculture12060779>
- Li, X., Qian, Q., Fu, Z., Wang, Y., Xiong, G., Zeng, D., Wang, X., Liu, X., Teng, S., Hiroshi, F., Yuan, M., Luo, D., Han, B., & Li, J. (2003). Control of Tillering in Rice. *Nature*, 422, 618–621. <https://doi.org/10.1038/nature01518>
- Lian, L., Xu, H., Zhang, H., He, W., Cai, Q., Lin, Y., Wei, L., Pan, L., Xie, X., Zheng, Y., Wei, Y., Zhu, Y., Xie, H., & Zhang, J. (2020). Overexpression of OsSPL14 Results in Transcriptome and Physiology Changes in Indica Rice 'MH86.' *Plant Growth Regulation*, 90, 265–278. <https://doi.org/10.1007/s10725-019-00569-0>
- Liang, W. hong, Shang, F., Lin, Q. ting, Lou, C., & Zhang, J. (2014). Tillering and Panicle Branching Genes in Rice. *Gene*, 537, 1–5. <https://doi.org/10.1016/j.gene.2013.11.058>
- Liao, Z., Yu, H., Duan, J., Yuan, K., Yu, C., Meng, X., Kou, L., Chen, M., Jing, Y., Liu, G., Smith, S. M., & Li, J. (2019). SLR1 Inhibits MOC1 Degradation to Coordinate Tiller Number and Plant Height in Rice. *Nature Communications*, 10(1), 1–9. <https://doi.org/10.1038/s41467-019-10667-2>
- Liu, X., Hu, Q., Yan, J., Sun, K., Liang, Y., Jia, M., Meng, X., Fang, S., Wang, Y., Jing, Y., Liu, G., Wu, D., Chu, C., Smith, S. M., Chu, J., Wang, Y., Li, J., & Wang, B. (2020). ζ -Carotene Isomerase Suppresses Tillering in Rice through the Coordinated Biosynthesis of Strigolactone and Absciscic Acid. *Molecular Plant*, 13(12), 1784–1801. <https://doi.org/10.1016/j.molp.2020.10.001>
- Liu, Y., Gu, D., Ding, Y., Wang, Q., Li, G., & Wang, S. (2011). The Relationship Between Nitrogen, Auxin and Cytokinin in The Growth Regulation of Rice

- (*Oryza sativa* L.) Tiller Buds. *Australian Journal of Crop Science*, 5(8), 1019–1026.
- Livak, K. J., & Schmittgen, T. D. (2001). Analysis of Relative Gene Expression Data Using Real-Time Quantitative PCR and The 2- $\Delta\Delta$ CT Method. *Methods*, 25(4), 402–408. <https://doi.org/10.1006/meth.2001.1262>
- Luo, L., Li, W., Miura, K., Ashikari, M., & Kyoizuka, J. (2012). Control of Tiller Growth of Rice by *OsSPL14* and Strigolactones, Which Work in Two Independent Pathways. *Plant and Cell Physiology*, 53(10), 1793–1801. <https://doi.org/10.1093/pcp/pcs122>
- Makino, A. (2011). Photosynthesis, Grain Yield P Nitrogen Utilization. *Plant Physiology*, 155(1), 125–129.
- Mardiah, Z., Septianingrum, E., Handoko, D. d., & Kusbiantoro, B. (2017). Improvement of Red Rice Eating Quality Through One-Time Polishing Process and Evaluation on Its Phenolic and Anthocyanin Content. *International Journal of Agriculture, Forestry and Plantation*, 5, 22–28.
- Mathan, J., Bhattacharya, J., & Ranjan, A. (2016). Enhancing Crop Yield by Optimizing Plant Developmental Features. *Development (Cambridge)*, 143(18), 3283–3294. <https://doi.org/10.1242/dev.134072>
- Mayer, K. F. X., Schoof, H., Haecker, A., Lenhard, M., Jürgens, G., & Laux, T. (1998). Role of *WUSCHEL* in Regulating Stem Cell Fate in The Arabidopsis Shoot Meristem. *Cell*, 95(6), 805–815. [https://doi.org/10.1016/S0092-8674\(00\)81703-1](https://doi.org/10.1016/S0092-8674(00)81703-1)
- Miura, K., Ikeda, M., Matsubara, A., Song, X.-J., Ito, M., Asano, K., Matsuoka, M., Kitano, H., & Ashikari, M. (2010). *OsSPL14* Promotes Panicle Branching and Higher Grain Productivity in Rice. *Nature Genetics*, 42(6), 545–549. <https://doi.org/10.1038/ng.592>
- Monobe, M. M. de S., & Silva, R. C. da. (2016). Gene Expression: An Overview of Methods and Applications for Cancer Research. *Veterinaria e Zootecnia*, 23(4), 532–546.
- Moon, S., Jung, K.-H., Lee, D.-E., Lee, D.-Y., Lee, J., An, K., Kang, H.-G., & An, G. (2006). The Rice *FON1* Gene Controls Vegetative and Reproductive Development by Regulating Shoot Apical Meristem Size. *Molecules and Cells*, 21(1), 147–152.
- Nafisa, A. R. (2021). *Analisis Faktor Transkripsi OsERF3, OsAP2, OsEREBP2 Tanaman Padi Hitam Transgenik (Oryza sativa L. “Cempo Ireng”) Overekspresi OsRKD3*. Universitas Gadjah Mada.
- Nagasawa, N., Miyoshi, M., Kitano, H., Satoh, H., & Nagato, Y. (1996). Mutations Associated with Floral Organ Number in Rice. *Planta*, 198, 627–633. <https://doi.org/10.1007/BF00262651>
- Nicolas, A., & Laufs, P. (2022). Meristem Initiation and de novo Stem Cell Formation. *Frontiers in Plant Science*, 13. <https://doi.org/10.3389/fpls.2022.891228>
- Nutan, K. K., Rathore, R. S., Tripathi, A. K., Mishra, M., Pareek, A., & Singla-Pareek, S. L. (2020). Integrating the Dynamics of Yield Traits in Rice in Response to Environmental Changes. *Journal of Experimental Botany*, 71(2), 490–506. <https://doi.org/10.1093/jxb/erz364>

- Oktaviani, D. F. (2020). *Ekspresi Gen OsRDK4 Pada Padi Hitam Transgenik (*Oryza sativa* L. 'Cempo Ireng') Generasi T2 Pembawa Konstruksi 35S::GAL4::OsRKD4::GR*. Universitas Gadjah Mada.
- Ouwerkerk, P. B., De Kam, R. J., Hoge, J. H. C., & Meijer, A. H. (2001). Glucocorticoid-Inducible Gene Expression in Rice. *Planta*, 213(3), 370–378. <https://doi.org/10.1007/s004250100583>
- Polato, N. (n.d.). *Oryza Illustrations*. Retrieved January 10, 2023, from https://archive.gramene.org/species/oryza/rice_illustrations.html
- Pratiwi, R., & Purwestri, Y. A. (2017). Black Rice as A Functional Food in Indonesia. *Functional Foods in Health and Disease*, 7(3), 182–194. <https://doi.org/10.31989/ffhd.v7i3.310>
- Purwestri, Y. A., Sari, R. D. K., Anggraeni, L. N., & Sasongko, A. B. (2015). *Agrobacterium tumefaciens* Mediated Transformation of rolC::Hd3a-GFP in Black Rice (*Oryza Sativa* L. cv. Cempo Ireng) to Promote Early Flowering. *Procedia Chemistry*, 14, 469–473. <https://doi.org/10.1016/j.proche.2015.03.063>
- Purwestri, Y. A., Susanto, F. A., & Fauzia, A. N. (2019). Flowering Gene Expression in Indonesian Long Harvest Black Rice (*Oryza sativa* L. 'Cempo Ireng'). *Australian Journal of Crop Science*, 13(6), 874–880. <https://doi.org/10.21475/ajcs.19.13.06.p1588>
- Rahim, M. A., Umar, M., Habib, A., Imran, M., Khalid, W., Lima, C. M. G., Shoukat, A., Itrat, N., Nazir, A., Ejaz, A., Zafar, A., Awuchi, C. G., Sharma, R., Santana, R. F., & Emran, T. Bin. (2022). Photochemistry, Functional Properties, Food Applications, and Health Prospective of Black Rice. *Journal of Chemistry*, 2022. <https://doi.org/10.1155/2022/2755084>
- Rebouillat, J., Dievart, A., Verdeil, J. L., Escoute, J., Giese, G., Breitler, J. C., Gantet, P., Espeout, S., Guiderdoni, E., & Périn, C. (2009). Molecular Genetics of Rice Root Development. *Rice*, 2, 15–34. <https://doi.org/10.1007/s12284-008-9016-5>
- Saichuk, J. (2009). *Louisiana Rice Production Handbook*. Louisiana State University Agricultural Center. USA. <http://www.lsuagcenter.com/~media/system/1/a/1/e/1a1e972d0509a6bb68d2637605c85038/pub2331riceproductionhandbook2014completebook.pdf>
- Sakamoto, T., & Matsuoka, M. (2004). Generating High-Yielding Varieties by Genetic Manipulation of Plant Architecture. *Current Opinion in Biotechnology*, 15, 144–147. <https://doi.org/10.1016/j.copbio.2004.02.003>
- Salgado, J. M., de Oliveira, A. G. C., Mansi, D. N., Donado-Pestana, C. M., & Bastos, Candido Ricardo Marcondes, F. K. (2010). The Role of Black Rice (*Oryza sativa* L.) in the Control of Hypercholesterolemia in Rats. *Journal of Medicinal Food*, 13(6), 1355–1362.
- Salsinha, Y. C. F., Maryani, Indradewa, D., Purwestri, Y. A., & Rachmawati, D. (2021). Leaf Physiological and Anatomical Characters Contribute to Drought Tolerance of Nusa Tenggara Timur Local Rice Cultivars. *Journal of Crop Science and Biotechnology*, 24(3), 337–348. <https://doi.org/10.1007/s12892-020-00082-1>
- Scofield, G. N., Hirose, T., Aoki, N., & Furbank, R. T. (2007). Involvement of The

- Sucrose Transporter, OsSUT1, in The Long-Distance Pathway for Assimilate Transport in Rice. *Journal of Experimental Botany*, 58(12), 3155–3169. <https://doi.org/10.1093/jxb/erm153>
- Shao, G., Lu, Z., Xiong, J., Wang, B., Jing, Y., Meng, X., Liu, G., Ma, H., Liang, Y., Chen, F., Wang, Y., Li, J., & Yu, H. (2019). Tiller Bud Formation Regulators MOC1 and MOC3 Cooperatively Promote Tiller Bud Outgrowth by Activating FON1 Expression in Rice. *Molecular Plant*, 12(8), 1090–1102. <https://doi.org/10.1016/j.molp.2019.04.008>
- Smith, C. W., & Dilday, R. H. (2003). *Rice: Origin, History, Technology, and Production*. John Wiley & Sons, Inc. United States of America.
- Song, X., Lu, Z., Yu, H., Shao, G., Xiong, J., Meng, X., Jing, Y., Liu, G., Xiong, G., Duan, J., Yao, X.-F., Liu, C.-M., Li, H., Wang, Y., & Li, J. (2017). IPA1 Functions As A Downstream Transcription Factor Repressed by D53 in Strigolactone Signaling in Rice. *Cell Research*, 27, 1128–1141. <https://doi.org/10.1038/cr.2017.102>
- Suzaki, T., Sato, M., Ashikari, M., Miyoshi, M., Nagato, Y., & Hirano, H.-Y. (2004). The Gene FLORAL ORGAN NUMBER1 Regulates Floral Meristem Size in Rice and Encodes A Leucine-Rich Repeat Receptor Kinase Orthologous to Arabidopsis CLAVATA1. *Development*, 131(22), 5649–5657. <https://doi.org/10.1242/dev.01441>
- Tanaka, W., Ohmori, Y., Ushijima, T., Matsusaka, H., Matsushita, T., Kumamaru, T., Kawano, S., & Hirano, H. (2015). Axillary Meristem Formation in Rice Requires the WUSCHEL Ortholog TILLERS ABSENT1. *Plant Cell*, 27(4), 1173–1184. <https://doi.org/10.1105/tpc.15.00074>
- Tripathi, K. K., Govila, O. P., Warriar, R., & Ahuja, V. (2011). *Biology of Oryza sativa L. (Rice)*. Department of Biotechnology, Ministry of Science and Technology, Ministry of Environment and Forests, Government of India. India.
- van den Elzen, P. J. M., Townsend, J., Lee, K. Y., & Bedbrook, J. R. (1985). A Chimaeric Hygromycin Resistance Gene as A Selectable Marker in Plant Cells. *Plant Molecular Biology*, 5(5), 299–302. <https://doi.org/10.1007/BF00020627>
- Waki, T., Hiki, T., Watanabe, R., Hashimoto, T., & Nakajima, K. (2011). The Arabidopsis RWP-RK Protein RKD4 Triggers Gene Expression and Pattern Formation in Early Embryogenesis. *Current Biology*, 21(15), 1277–1281. <https://doi.org/10.1016/j.cub.2011.07.001>
- Wang, B., Smith, S. M., & Li, J. (2018). Genetic Regulation of Shoot Architecture. *Annual Review of Plant Biology*, 69(1), 437–468. <https://doi.org/10.1146/annurev-arplant-042817-040422>
- Wang, L., Sun, S., Jin, J., Fu, D., Yang, X., Weng, X., Xu, C., Li, X., Xiao, J., & Zhang, Q. (2015). Coordinated Regulation of Vegetative and Reproductive Branching in Rice. *Proceedings of the National Academy of Sciences of the United States of America*, 112(50), 15504–15509. <https://doi.org/10.1073/pnas.1521949112>
- Wang, Y., & Li, J. (2008). Molecular Basis of Plant Architecture. *Annual Review of Plant Biology*, 59, 253–279.

<https://doi.org/10.1146/annurev.arplant.59.032607.092902>

- Wijayanti, P. (2018). *Karakter Fenotipik dan Molekuler Padi Hitam Transgenik (*Oryza sativa* L. 'Cempo Ireng') Pembawa Gen OsRKD4*. Universitas Gadjah Mada.
- Williams, M. L., Farrar, J. F., & Pollock, C. J. (1989). Cell Specialization Within The Parenchymatous Bundle Sheath of Barley. *Plant, Cell and Environment*, 12, 909–918.
- Wu, L.-L., Liu, Z.-L., Wang, J.-M., Zhou, C.-Y., & Chen, K.-M. (2011). Morphological, Anatomical, and Physiological Characteristics Involved in Development of The Large Culm Trait in Rice. *Australian Journal of Crop Science*, 5(11), 1356–1363.
- Xing, Y., & Zhang, Q. (2010). Genetic and Molecular Bases of Rice Yield. *Annual Review of Plant Biology*, 61, 421–442. <https://doi.org/10.1146/annurev-arplant-042809-112209>
- Xue, Z., Liu, L., & Zhang, C. (2020). Regulation of Shoot Apical Meristem and Axillary Meristem Development in Plants. *International Journal of Molecular Sciences*, 21(8). <https://doi.org/10.3390/ijms21082917>
- Yan, J. Q., Zhu, J., He, C. X., Benmoussa, M., & Wu, P. (1998). Quantitative Trait Loci Analysis for The Developmental Behavior of Tiller Number in Rice (*Oryza sativa* L.). *Theoretical and Applied Genetics*, 97, 267–274. <https://doi.org/10.1007/s001220050895>
- Yeung, E. C. T., Stasolla, C., Sumner, M. J., & Huang, B. Q. (2015). Plant Microtechniques and Protocols. In *Suparyanto dan Rosad (2015)* (Vol. 5, Issue 3). Springer International Publishing Switzerland.
- Yue, E., Li, C., Li, Y., Liu, Z., & Xu, J.-H. (2017). MiR529a modulates panicle Architecture Through Regulating SQUAMOSA PROMOTER BINDING-LIKE Genes in Rice (*Oryza sativa*). *Plant Molecular Biology*, 94, 469–480. <https://doi.org/10.1007/s11103-017-0618-4>
- Zawistowski, J., Kopec, A., & Kitts, D. D. (2009). Effects of A Black Rice Extract (*Oryza sativa* L. indica) on Cholesterol Levels and Plasma Lipid Parameters in Wistar Kyoto Rats. *Journal of Functional Foods*, 1(1), 50–56. <https://doi.org/10.1016/j.jff.2008.09.008>