

**BIOREMEDIASI LOGAM KROMIUM MENGGUNAKAN KOMBINASI  
FORMULA DARI KONSORSIUM BAKTERI *Brevibacillus borstelensis*,  
*Bacillus licheniformis*, DAN *Bacillus* sp.**

**SKRIPSI**

*Diajukan untuk memenuhi sebagian syarat memperoleh gelar Sarjana Sains  
Program Studi Biologi*



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BANDUNG  
2025**

## **LEMBAR HAK CIPTA**

### **BIOREMEDIASI LOGAM KROMIUM MENGGUNAKAN KOMBINASI FORMULA DARI KONSORSIUM BAKTERI *Brevibacillus borstelensis*, *Bacillus licheniformis*, DAN *Bacillus* sp.**

Oleh  
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Sebuah skripsi yang diajukan untuk memenuhi salah satu syarat memperoleh  
gelar Sarjana Sains pada Program Studi Biologi,  
Fakultas Pendidikan Matematika dan Ilmu Pengetahuan Alam

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Juli 2025

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**BIOREMEDIASI LOGAM KROMIUM MENGGUNAKAN KOMBINASI  
FORMULA DARI KONSORSIUM BAKTERI *Brevibacillus borstelensis*,  
*Bacillus licheniformis*, DAN *Bacillus* sp.**

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## **LEMBAR PERNYATAAN**

Dengan ini saya menyatakan bahwa skripsi dengan judul “Bioremediasi Logam Kromium Menggunakan Kombinasi Formula dari Konsorsium Bakteri *Brevibacillus borstelensis*, *Bacillus licheniformis*, dan *Bacillus* sp.” ini beserta seluruh isinya adalah benar-benar karya saya sendiri. Saya tidak melakukan penjiplakan atau pengutipan dengan cara-cara yang tidak sesuai dengan etika ilmu yang berlaku dalam masyarakat keilmuan. Atas pernyataan ini, saya bersedia menanggung risiko/sanksi apabila di kemudian hari ditemukan adanya pelanggaran etika keilmuan atau ada klaim dari pihak lain terhadap keaslian karya saya ini.

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Yang membuat pernyataan,

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Dalam proses penulisan skripsi ini, penulis menyadari bahwa masih terdapat ruang untuk perbaikan. Oleh karena itu, saran dan kritik yang membangun sangat diharapkan agar skripsi ini dapat menjadi referensi yang lebih baik di masa mendatang. Semoga hasil dari skripsi ini dapat memberikan kontribusi positif dalam pengembangan ilmu pengetahuan bagi penulis dan pembaca yang berkepentingan.

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## ABSTRAK

### Bioremediasi Logam Kromium Menggunakan Kombinasi Formula dari Konsorsium Bakteri *Brevibacillus borstelensis*, *Bacillus licheniformis*, dan *Bacillus* sp.

Logam kromium merupakan salah satu polutan prioritas karena menimbulkan berbagai resiko bagi organisme, mikroorganisme, dan lingkungan. Industri kerajinan kulit menjadi salah satu penghasil limbah yang mengandung logam kromium dari limbah penyamakan kulit. Upaya yang dapat dilakukan untuk mengurangi pencemaran logam kromium di lingkungan yaitu dengan metode bioremediasi. Bioremediasi dapat menggunakan mikroorganisme yaitu bakteri, penggunaan konsorsium bakteri dianggap lebih efektif dibandingkan dengan bakteri tunggal. Penelitian ini bertujuan untuk menganalisis kemampuan formula konsorsium bakteri yang berpotensi sebagai agen bioremediasi logam kromium secara *in vitro*. Bakteri diisolasi dari tanah area *rhizosfer* tumbuhan yang berada di sekitar bak penampungan limbah di wilayah Sukaregang, Kabupaten Garut. Bakteri diseleksi dengan memilih bakteri yang memiliki resistensi tertinggi lalu diidentifikasi berdasarkan aktivitas biokimia dan gen 16S rRNA. Kemampuan bioremediasi logam kromium ditunjukkan berdasarkan interaksi antar bakteri dan persentase pengurangan Cr(VI) oleh konsorsium bakteri. Identifikasi bakteri berdasarkan aktivitas biokimia yang dilengkapi dengan analisis gen 16S rRNA menunjukkan ketiga isolat bakteri teridentifikasi sebagai *Brevibacillus borstelensis*, *Bacillus licheniformis*, dan *Bacillus* sp.. Hasil uji interaksi antar bakteri menunjukkan bahwa bakteri penyusun konsorsium saling bersinergi. Seluruh konsorsium bakteri mampu mengurangi Cr(VI) namun memiliki kemampuan yang berbeda. Konsorsium dengan kombinasi tiga bakteri yaitu *Brevibacillus borstelensis*, *Bacillus licheniformis*, dan *Bacillus* sp. memiliki persentase pengurangan Cr(VI) tertinggi yaitu sebesar 39,38%.

**Kata Kunci:** Bioremediasi, Gen 16S rRNA, Konsorsium, Kromium, Penyamakan Kulit

## ***ABSTRACT***

### ***Bioremediation of Chromium Metal Using Formula Combination of Bacterial Consortium *Brevibacillus borstelensis*, *Bacillus licheniformis*, and *Bacillus* sp.***

*Chromium metal is one of the priority pollutants because it poses various risks to organisms, microorganisms, and the environment. The leather craft industry is one of the producers of waste containing chromium metal from tanning waste. Efforts that can be made to reduce chromium metal pollution in the environment are by bioremediation methods. Bioremediation can use microorganisms, namely bacteria, the use of bacterial consortia is considered more effective than single bacteria. This study aims to analyze the ability of bacterial consortium formulas that have the potential as chromium metal bioremediation agents in vitro. Bacteria were isolated from the soil of the rhizosphere area of plants around the waste storage tank in the Sukaregang area, Garut Regency. Bacteria were selected by selecting bacteria that had the highest resistance and then identified based on biochemical activity and the 16S rRNA gene. The ability of chromium metal bioremediation was shown based on the interaction between bacteria and the percentage of Cr(VI) reduction by the bacterial consortium. Identification of bacteria based on biochemical activity supported by 16S rRNA gene analysis showed that the three bacterial isolates were identified as *Brevibacillus borstelensis*, *Bacillus licheniformis*, and *Bacillus* sp.. The results of the interaction test between bacteria showed that the bacteria that make up the consortium synergize with each other. All bacterial consortia are able to reduce Cr(VI) but have different abilities. The consortium with a combination of three bacteria, namely *Brevibacillus borstelensis*, *Bacillus licheniformis*, and *Bacillus* sp. has the highest Cr(VI) reduction percentage of 39.38%.*

***Keywords:*** Bioremediation, 16S rRNA Gene, Consortium, Chromium, Leather Tanning

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## DAFTAR PUSTAKA

- Abdullah, K. (2018). *Berbagai Metodologi dalam Penelitian Pendidikan dan Manajemen*. Gowa: Gunadarma Ilmu.
- Ablaye, N., Samba, M. M., Madiop, G., Aziz, C. A., & Kandioura, N. (2019). Practical identification key of *Digitaria ciliaris* (Retz.) Koeler and *Digitaria horizontalis* Willd.(Poaceae): two harmful weeds of crops in Senegal. *GSC Biological and Pharmaceutical Sciences*, 8(2), 101–105. <https://doi.org/10.30574/gscbps.2019.8.2.0144>
- Ahsan, S., Mohona, E. F., Shaon, M. A. R., Turzo, S. M. S. M., & Rahman, S. (2024). Identification of Bacteria Capable of Chromium Bioremediation from the Environment. *Plant Tissue Culture and Biotechnology*, 34(1), 71–81. <https://doi.org/10.3329/ptcb.v34i1.74356>
- Aké, A. H. J., Rochdi, N., Jemo, M., Hafidi, M., Ouhdouch, Y., & El Fels, L. (2024). Cr (VI) removal performance from wastewater by microflora isolated from tannery effluents in a semi-arid environment: A SEM, EDX, FTIR and zeta potential study. *Frontiers in Microbiology*, 15, 1423741. <https://doi.org/10.3389/fmicb.2024.1423741>
- Akib, M. A. (2014). *Prosedur Rancangan Percobaan: Aplication Of The Model In Different Environmental Conditions*. Sengkang: Lampena Intimedia.
- Allan, R. N., Lebbe, L., Heyrman, J., De Vos, P., Buchanan, C. J., & Logan, N. A. (2005). *Brevibacillus levickii* sp. nov. and *Aneurinibacillus terranovensis* sp. nov., Two Novel Thermoacidophiles Isolated from Geothermal Soils of Northern Victoria Land, Antarctica. *International Journal of Systematic and Evolutionary Microbiology*, 55(3), 1039–1050. <https://doi.org/10.1099/ijss.0.63397-0>
- Al-Shuhaib, M. B. S., & Hashim, H. O. (2023). Mastering DNA Chromatogram Analysis in Sanger Sequencing for Reliable Clinical Analysis. *Journal of Genetic Engineering and Biotechnology*, 21(1). <https://doi.org/10.1186/s43141-023-00587-6>
- Alyousif, N. A. (2022). Distribution, Occurrence and Molecular Characterization of *Bacillus* Related Species Isolated from Different Soil in Basrah Province, Iraq. *Biodiversitas*, 23(2), 679–686. <https://doi.org/10.13057/biodiv/d230209>
- Amann, R. I., Ludwig, W., & Schleifer, K. H. (1995). Phylogenetic Identification and In Situ Detection of Individual Microbial Cells Without Cultivation. *Microbiological Reviews*, 59(1), 143–169. <https://doi.org/10.1128/mmbr.59.1.143-169.1995>
- Amin, A., Naveed, M., Sarwar, A., Rasheed, S., Saleem, H. G. M., Latif, Z., & Bechthold, A. (2022). In Vitro and In Silico Studies Reveal *Bacillus cereus* AA-18 as a Potential Candidate for Bioremediation of Mercury-Contaminated Wastewater. *Frontiers in Microbiology*, 13, 1–13. <https://doi.org/10.3389/fmicb.2022.847806>
- Applied Biosystems. (2010). BigDye® Terminator v3.1 Cycle Sequencing Kit Protocol.

- Apriliani, N. I. (2024). *Bioremediasi Kromium Menggunakan Kombinasi Dua Rhizobakteri Indigenous Secara In Vitro*. (Skripsi). Universitas Pendidikan Indonesia, Bandung.
- Arif, Ade Alfarez, D., & Rizky Ramadhan, M. (2023). Anova dan Tukey HSD Perbandingan Produksi Padi Antara Tiga Kabupaten di Provinsi Jambi. *Multi Proximity: Jurnal Statistika Universitas Jambi*, 2(1), 23–31. <https://doi.org/10.22437/multiproximity.v2i1.25908>
- Arsyadi, A., & Khusnuryani, A. (2016). “Isolasi Bakteri Indigenous Pereduksi Krom (VI) dari Limbah Cair Laboratorium Biologi UIN Sunan Kalijaga Yogyakarta”. Dalam Setyawan, A. D., Ridwan, M., Lestari, D. F., Pamungkas, D. W., Kharismamurti, K., Romadhon, M. A., & Liza, N (Penyunting). *Prosiding Seminar Nasional Biodiversitas* (hlm. 100–103). Surakarta: Kelompok Studi Biodiversitas Program Studi Biologi FMIPA UNS.
- Asif, S., Khan, M., Waqar Arshad, M., & Shabbir, M. I. (2021). PCR Optimization for Beginners: A Step by Step Guide. *Research in Molecular Medicine*, 9(2), 81–102. <https://doi.org/10.32598/rmm.9.2.1189.1>
- Asri, A. C., & Zulaika, E. (2016). Sinergisme Antar Isolat Azotobacter yang Dikonsorsiumkan. *Jurnal Sains dan Seni ITS*, 5(2), 57–59.
- Astutik, L. W. (2015). *Resistensi Dan Potensi Azotobacter Sebagai Bioremoval Timbal (Pb)*. (Skripsi). Institut Teknologi Sepuluh Nopember, Surabaya.
- Ayele, A., & Godeto, Y. G. (2021). Bioremediation of Chromium by Microorganisms and Its Mechanisms Related to Functional Groups. *Hindawi : Journal of Chemistry*. 2021. 1–21. <https://doi.org/10.1155/2021/7694157>
- Aziema, N. (2024). *Uji Kemampuan Formula Konsorsium Bakteri Rhizosfer dalam Bioremediasi Logam Kromium Secara In Vitro*. (Skripsi). Universitas Pendidikan Indonesia, Bandung.
- Azubuike, C. C., Chikere, C. B., & Okpokwasili, G. C. (2016). Bioremediation Techniques—Classification Based on Site of Application: Principles, Advantages, Limitations and Prospects. *World Journal of Microbiology and Biotechnology*, 32(11). 1–21. <https://doi.org/10.1007/s11274-016-2137-x>
- Bahiru, D. B., Teju, E., Kebede, T., & Demissie, N. (2019). Levels of Some Toxic Heavy Metals (Cr, Cd and Pb) in Selected Vegetables and Soil Around Eastern Industry Zone, Central Ethiopia. *African Journal of Agricultural Research*, 14(2), 92–101. <https://doi.org/10.5897/ajar2018.13324>
- Bakshi, A., & Panigrahi, A. K. (2022). Chromium Contamination in Soil and Its Bioremediation: An Overview. Dalam J. A. Malik (Penyunting), *Advances in Bioremediation and Phytoremediation for Sustainable Soil Management: Principles, Monitoring and Remediation* (hlm. 229–248). Cham: Springer International Publishing. [https://doi.org/10.1007/978-3-030-89984-4\\_15](https://doi.org/10.1007/978-3-030-89984-4_15)
- Barco, R. A., Garrity, G. M., Scott, J. J., Amend, J. P., Nealson, K. H., & Emerson, D. (2020). A Genus Definition for Bacteria and Archaea Based on a Standard Genome Relatedness Index. *Ecological and Evolutionary Science*, 11(1), 1–20. <https://doi.org/10.1128/mBio.02475-19>

- Barghouthi, S. A. (2011). A Universal Method for the Identification of Bacteria Based on General PCR Primers. *Indian Journal of Microbiology*, 51(4), 430–444. <https://doi.org/10.1007/s12088-011-0122-5>
- Barillot, C. D. C., Sarde, C. O., Bert, V., Tarnaud, E., & Cochet, N. (2013). A Standardized Method for The Sampling of Rhizosphere and Rhizoplan Soil Bacteria Associated to A Herbaceous Root System. *Annals of Microbiology*, 63(2), 471–476. <https://doi.org/10.1007/s13213-012-0491-y>
- Basiludin, B., & Kurniawan, M. H. S. (2023). Analisis Dampak Pandemi Covid-19 terhadap City Pair Penerbangan yang Terdampak Menggunakan Uji Friedman dan Analisis Regresi Linear. *Emerging Statistics and Data Science Journal*, 1(2), 215–226.
- Batool, R., Yrjälä, K., & Hasnain, S. (2014). Impact of Environmental Stress on Biochemical Parameters of Bacteria Reducing Chromium. *Brazilian Journal of Microbiology*, 45(2), 573–583. [www.sbmicrobiologia.org.br](http://www.sbmicrobiologia.org.br)
- Bini, C., Buffa, G., Fontana, S., & Wahsha, M. (2013). Are Plants Growing at Abandoned Mine Sites Suitable for Phytoremediation of Contaminated Soils?. *Geophysical Research Abstracts*, 15.
- Bredow, C., Azevedo, J. L., Pamphile, J. A., Mangolin, C. A., & Rhoden, S. A. (2015). In Silico Analysis of The 16S rRNA Gene of Endophytic Bacteria, Isolated From The Aerial Parts and Seeds of Important Agricultural Crops. *Genetics and Molecular Research*, 14(3), 9703–9721. <https://doi.org/10.4238/2015.August.19.3>
- Bruslind, L. (2019). *General Microbiology*. Corvallis: Oregon State University.
- Bustin, S. A., Benes, V., Garson, J. A., Hellemans, J., Huggett, J., Kubista, M., Mueller, R., Nolan, T., Pfaffl, M. W., Shipley, G. L., Vandesompele, J., & Wittwer, C. T. (2009). The MIQE Guidelines: Minimum Information for Publication of Quantitative Real-time PCR Experiments. *Clinical Chemistry*, 55(4), 611–622. <https://doi.org/10.1373/clinchem.2008.112797>
- Cao, Z., Yan, W., Ding, M., & Yuan, Y. (2022). Construction of Microbial Consortia for Microbial Degradation of Complex Compounds. *Frontiers in Bioengineering and Biotechnology*, 10, 1–14. <https://doi.org/10.3389/fbioe.2022.1051233>
- Cappuccino, J. G., & Sherman, N. (2014). *Microbiology A Laboratory Manual Tenth Edition*. Boston: Pearson Education.
- Cervantes, C., Campos-García, J., Devars, S., Gutiérrez-Corona, F., Loza-Tavera, H., Torres-Guzmán, J. C., & Moreno-Sánchez, R. (2001). Interactions of chromium with microorganisms and plants. *FEMS Microbiology Reviews*, 25(3), 335–347. <https://doi.org/10.1111/j.1574-6976.2001.tb00581.x>
- Chen, W., Li, W., Wang, T., Wen, Y., Shi, W., Zhang, W., Guo, B., & Yang, Y. (2022). Isolation of Functional Bacterial Strains From Chromium-Contaminated Site and Bioremediation Potentials. *Journal of Environmental Management*, 307. <https://doi.org/10.1016/j.jenvman.2022.114557>
- Christofan, V. B., (2025). *Bioremediasi Logam Kromium Menggunakan Kombinasi Formula Dari Konsorsium Bakteri Bacillus anthracis, Bacillus licheniformis, dan Bacillus sp. (Skripsi)*. Universitas Pendidikan Indonesia, Bandung.

- Clarridge, J. E. (2004). Impact of 16S rRNA Gene Sequence Analysis for Identification of Bacteria on Clinical Microbiology and Infectious Diseases. *Clinical Microbiology Reviews*, 17(4), 840–862. <https://doi.org/10.1128/CMR.17.4.840-862.2004>
- Coreño-Alonso, A., Cruz-Jiménez, G., López-Martínez, L., Reyna-López, G. E., & Acevedo-Aguilar, F. J. (2017). A Rapid, Eco-Friendly, and Reliable Microplate Method for Determination of Cr(VI). *Turkish Journal of Chemistry*, 41(3), 420–425. <https://doi.org/10.3906/kim-1609-53>
- Costa-Santos, M., Mariz-Ponte, N., Dias, M. C., Moura, L., Marques, G., & Santos, C. (2021). Effect of *Bacillus* spp. And *Brevibacillus* sp. on The Photosynthesis and Redox Status of *Solanum Lycopersicum*. *Horticulturae*, 7(2), 1–14. <https://doi.org/10.3390/horticulturae7020024>
- Crossley, B. M., Bai, J., Glaser, A., Maes, R., Porter, E., Killian, M. L., Clement, T., & Toohey-Kurth, K. (2020). Guidelines for Sanger Sequencing and Molecular Assay Monitoring. *Journal of Veterinary Diagnostic Investigation*, 32(6), 767–775. <https://doi.org/10.1177/1040638720905833>
- Dave, S., & Bhatt, N. (2018). Biotransformation of Cr (VI) by Newly Invented Bacterial Consortium SN6. *Journal of Pure and Applied Microbiology*, 12(3), 1375–1384. <https://doi.org/10.22207/JPAM.12.3.40>
- de Vos, P., Garrity, G., Jones, D., Krieg, N. R., Ludwig, W., Rainey, F. A., Schleifer, K.-H., & Whitman, W. B. (2009). *Bergey's Manual of Systematic Bacteriology: Volume 3: The Firmicutes, Second Edition*. Dordrecht: Springer.
- Deb, J. S., & Roy, A. (2021). Isolation of Chromium Resistance Bacteria From Loam Soil Sample of River Ganga. *World Journal of Pharmaceutical and Life Science*, 7(9), 145–157.
- Denaya, S., Yulianti, R., Pambudi, A., & Effendi, Y. (2021). “Novel Microbial Consortium Formulation as Plant Growth Promoting Bacteria (PGPB) Agent”. Dalam *IOP Conference Series: Earth and Environmental Science*, 637(1). <https://doi.org/10.1088/1755-1315/637/1/012030>
- Dinoto, A., Handayani, R., Setianingrum, N., & Julistiono, H. (2020). Culturable Gut Bacteria of Ikan Batak (*Neolissochilus sumatrana* Weber & De Beaufort, 1916) Collected in Toba Samosir, Indonesia. *Biodiversitas*, 21(10), 4483–4488. <https://doi.org/10.13057/biodiv/d211003>
- Divyasree, P., & Subramanian, J. J. (2014). Comparative Studies on The Bioremediation of Hexavalent and Trivalent Chromium using *Citrobacter freundii*: Part I-Effect of parameters controlling Biosorption. *Int. J. Environ. Res*, 8(4), 1127–1134.
- Dixit, R., Wasiullah, Malaviya, D., Pandiyan, K., Singh, U. B., Sahu, A., Shukla, R., Singh, B. P., Rai, J. P., Sharma, P. K., Lade, H., & Paul, D. (2015). Bioremediation of Heavy Metals From Soil and Aquatic Environment: An Overview of Principles and Criteria of Fundamental Processes. *Sustainability (Switzerland)*, 7(2), 2189–2212. <https://doi.org/10.3390/su7022189>

- Elahi, A., & Rehman, A. (2019). Comparative behavior of two gram positive Cr<sup>6+</sup>-resistant bacterial strains *Bacillus aerius* S1 and *Brevibacterium iodinum* S2 under hexavalent chromium stress. *Biotechnology Reports*, <https://doi.org/10.1016/j.btre.2019.e00307>.
- Elshafey, N., Elkashef, A., Hamedo, H., & Mansour, M. (2022). Molecular Identification of Halophilic Microorganisms Producing Exo-Halozymes by A Culture-Dependent Approach From The North Sinai Solar saltern. *The Egyptian Journal of Experimental Biology*, 18(2), 187–195. <https://doi.org/10.5455/egyjebb.20220726105555>
- Eng, A. Y., Narayanan, A., Alster, C. J., & DeAngelis, K. M. (2023). Thermal Adaptation of Soil Microbial Growth Traits in Response to Chronic Warming. *Applied and Environmental Microbiology*, 89(11). <https://doi.org/10.1128/aem.00825-23>
- Errington, J. (2003). Regulation of endospore formation in *Bacillus subtilis*. *Nature Reviews Microbiology*, 1(2), 117–126. <https://doi.org/10.1038/nrmicro750>
- Fathima, A., Manikandamathan, V. M., Jonnalagadda, R. R., & Unni Nair, B. (2020). Chromium-catechin Complex, Synthesis and Toxicity Check Using Bacterial Models. *Heliyon*, 6(8). <https://doi.org/10.1016/j.heliyon.2020.e04563>
- Fatwa, B. E., Yoswaty, D., & Effendi, I. (2021). Identification of Indigenous Bacteria from Dumai Sea Waters Using 16S rRNA Method. *Journal of Coastal and Ocean Sciences*. 2(3).
- Gao, K., Mu, C. L., Farzi, A., & Zhu, W. Y. (2020). Tryptophan Metabolism: A Link Between the Gut Microbiota and Brain. *Advances in Nutrition*, 11(3), 709–723. <https://doi.org/10.1093/advances/nmz127>
- Garrity, G. M., Bell, J. A., & Lilburn, T. G. (2004). *Taxonomic Outline of The Prokaryotes Bergey's Manual ® of Systematic Bacteriology* (2nd Edition). New York: Springer. <https://doi.org/10.1007/bergeysoutline200405>
- Gupta, A. K., Ganjewala, D., Goel, N., & Khurana, N. (2013). Bioremediation of Tannery Chromium: A Microbial Approach. *Research Journal of Pharmacy and Technology*. 7(1). 118–122.
- Hadi, A. E., Khalisha, A., Pambudi, A., & Effendi, Y. (2021). “Potential of Bacteria Consortium as Growth Controller of Pathogenic Fungi *Fusarium oxysporum* F. sp. *cubense* (Foc)”. Dalam *IOP Conference Series: Earth and Environmental Science*, 637(1). <https://doi.org/10.1088/1755-1315/637/1/012029>
- Hamed, M. T., Elwakil, B. H., Hagar, M., Ghareeb, D. A., & Olama, Z. A. (2025). Chromium bioremediation mechanistic action assessment using bacterial consortium isolated from Egyptian Petroleum Refining Company. *Scientific African*, 28, e02642. <https://doi.org/10.1016/j.sciaf.2025.e02642>
- Han, S., Li, Y., & Gao, H. (2022). Generation and Physiology of Hydrogen Sulfide and Reactive Sulfur Species in Bacteria. *Antioxidants*, 11(12). <https://doi.org/10.3390/antiox11122487>

- Hasan, M. I., Bag, S., Halder, D., Bhowmik, S., Chakraborty, A., & Ghosh, A. (2024). Simultaneous Removal of Malachite Green and Lead From Water by Consortium Dry-biomasses of *Bacillus licheniformis* AG3 and *Bacillus cereus* M116. *Scientific Reports*, 14(1). <https://doi.org/10.1038/s41598-024-70658-2>
- Hinkelmann, K. (2015). History and Overview of Design and Analysis of Experiments. Dalam A. Dean, M. Morris, J. Stufken, & D. Bingham (Ed.), *Handbook of Design and Analysis of Experiments* (hlm. 3–62). Boston: Taylor & Francis Group.
- Hitsmi, M., Firdaus, Ml., & Nurhamidah. (2020). Pengembangan Metode Citra Digital Berbasis Aplikasi Android Untuk Analisis Ion Logam Cr(VI). *Jurnal Pendidikan dan Ilmu Kimia*. 4(2), 117–124.
- Hossan, S., Hossain, S., Islam, M. R., Kabir, M. H., Ali, S., Islam, M. S., Imran, K. M., Moniruzzaman, M., Mou, T. J., Parvez, A. K., & Mahmud, Z. H. (2020). Bioremediation of Hexavalent Chromium by Chromium Resistant Bacteria Reduces Phytotoxicity. *International journal of environmental research and public health*, 17(17), 6013. <https://doi.org/10.3390/ijerph17176013>
- Huang, Y., Tang, J., Zhang, B., Long, Z. E., Ni, H., Fu, X., & Zou, L. (2023). Influencing Factors and Mechanism of Cr(VI) Reduction by Facultative Anaerobic *Exiguobacterium* sp. PY14. *Frontiers in Microbiology*, 14. <https://doi.org/10.3389/fmicb.2023.1242410>
- Ibrahim, K. M., & Peterson, P. M. (2014). *Grasses of Washington*, D.C. Washington D.C: Smithsonian Institution Scholarly Press.
- Iskandar, A. U., Ethica, S. N., Sukeksi, A., Mukaromah, A. H., Sulistyaningtyas, A. R., & Darmawati, S. (2021). “Molecular systematic and phylogenetic analysis of indigenous bacterial isolates with potential as bioremediation agent based on 16S rRNA gene analysis”. Dalam *IOP Conference Series: Earth and Environmental Science*, 743(1). <https://doi.org/10.1088/1755-1315/743/1/012010>
- Jacob, F., & Monod, J. (1961). Genetic egulatory Mechanisms in The Synthesis of Proteins. *Journal of Molecular Biology*, 3(3), 318–356. [https://doi.org/10.1016/S0022-2836\(61\)80072-7](https://doi.org/10.1016/S0022-2836(61)80072-7)
- Jiang, H., Dong, H., Zhang, G., Yu, B., Chapman, L. R., & Fields, M. W. (2006). Microbial Diversity in Water and Sediment of Lake Chaka, an Athalassohaline Lake in Northwestern China. *Applied and Environmental Microbiology*, 72(6), 3832–3845. <https://doi.org/10.1128/AEM.02869-05>
- Johnson, J. S., Spakowicz, D. J., Hong, B. Y., Petersen, L. M., Demkowicz, P., Chen, L., Leopold, S. R., Hanson, B. M., Agresta, H. O., Gerstein, M., Sodergren, E., & Weinstock, G. M. (2019). Evaluation of 16S rRNA Gene Sequencing for Species and Strain-level Microbiome Analysis. *Nature Communications*, 10(1). <https://doi.org/10.1038/s41467-019-13036-1>
- Kadhum, H. A., & Hasan, T. H. (2019). The Study of *Bacillus subtilis* Antimicrobial Activity on Some of The Pathological Isolates. *International Journal of Drug Delivery Technology*, 9(2), 193–196. <https://doi.org/10.25258/ijddt.9.2.12>

- Kalsoom, Batool, A., Din, G., Din, S. U., Jamil, J., Hasan, F., Khan, S., Badshah, M., & Shah, A. A. (2021). Isolation and screening of chromium resistant bacteria from industrial waste for bioremediation purposes. *Brazilian journal of biology*, 83, e242536. <https://doi.org/10.1590/1519-6984.242536>
- Khalil, A. B., Sivakumar, N., Arslan, M., Saleem, H., & Qarawi, S. (2018). Insights into *Brevibacillus borstelensis* AK1 through Whole Genome Sequencing: a Thermophilic Bacterium Isolated from a Hot Spring in Saudi Arabia. *BioMed Research International*, 2018(1). <https://doi.org/10.1155/2018/5862437>
- Khanam, R., Al Ashik, S. A., Suriea, U., & Mahmud, S. (2024). Isolation of Chromium Resistant Bacteria from annery Waste and Assessment of Their Chromium Reducing Capabilities – A Bioremediation Approach. *Heliyon*, 10(6). <https://doi.org/10.1016/j.heliyon.2024.e27821>
- Khuzaimah, I., & Astuti, D. (2022). "Literature Study of the Effectiveness of Bioremediation Methods on Chromium Reduction in Tanning Wastewater". Dalam *The 16<sup>th</sup> Research Colloquium* (hlm. 349–364).
- Kim, M., Oh, H. S., Park, S. C., & Chun, J. (2014). Towards a Taxonomic Coherence Between Average Nucleotide Identity and 16S rRNA Gene Sequence Similarity for Species Demarcation of Prokaryotes. *International Journal of Systematic and Evolutionary Microbiology*, 64, 346–351. <https://doi.org/10.1099/ijss.0.059774-0>
- Kredich, N. M. (2008). Biosynthesis of Cysteine. *EcoSal Plus*, 3(1). <https://doi.org/10.1128/ecosalplus.3.6.1.11>
- Kuncoro, Y. M., & Soedjono, E. S. (2022). Studi Pustaka: Teknologi Pengolahan Air Limbah pada Industri Penyamakan Kulit. *Jurnal Teknik ITS*, 11(3), 142–149.
- Kuzyakov, Y., & Razavi, B. S. (2019). Rhizosphere Size and Shape: Temporal Dynamics and Spatial Stationarity. *Soil Biology and Biochemistry*, 135, 343–360. <https://doi.org/10.1016/j.soilbio.2019.05.011>
- Lalucat, J., Mulet, M., Gomila, M., & García-Valdés, E. (2020). Genomics in Bacterial Taxonomy: Impact on the Genus *Pseudomonas*. *Genes*, 11(2), 139. <https://doi.org/10.3390/genes11020139>
- Langley, S., & Beveridge, T. J. (1999). Effect of O-side-chain-lipopolysaccharide chemistry on metal binding. *Applied and environmental microbiology*, 65(2), 489–498. <https://doi.org/10.1128/AEM.65.2.489-498.1999>
- Lewaru, S., Riyantini, I., Mulyani, Y., Fakultas, A., Dan Ilmu, P., Unpad, K., Dosen, S., Perikanan, F., & Ilmu, D. (2012). Identifikasi Bakteri Indigenous Pereduksi Logam Berat Cr (VI) Dengan Metode Molekuler Di Sungai Cikijing Rancaekek, Jawa Barat. *Jurnal Perikanan dan Kelautan*, 3(4), 81–92.
- Li, J., Gu, T., Li, L., Wu, X., Shen, L., Yu, R., ... & Zeng, W. (2020). Complete Genome Sequencing and Comparative Genomic Analyses of *Bacillus* sp. S3, a Novel Hyper Sb (III)-oxidizing Bacterium. *BMC microbiology*, 20(106), 1–19. <https://doi.org/10.1186/s12866-020-01737-3>

- Li, M. H., Gao, X. Y., Li, C., Yang, C. L., Fu, C. A., Liu, J., Wang, R., Chen, L. X., Lin, J. Q., Liu, X. M., Lin, J. Q., & Pang, X. (2020). Isolation and Identification of Chromium Reducing *Bacillus cereus* Species from Chromium-contaminated Soil for he Biological Detoxification of Chromium. *International Journal of Environmental Research and Public Health*, 17(6). <https://doi.org/10.3390/ijerph17062118>
- Liu, H., Bergman, N. H., Thomason, B., Shallom, S., Hazen, A., Crossno, J., Rasko, D. A., Ravel, J., Read, T. D., Peterson, S. N., Yates, J., & Hanna, P. C. (2004). Formation and Composition of the *Bacillus anthracis* Endospore. *Journal of Bacteriology*, 186(1), 164–178. <https://doi.org/10.1128/JB.186.1.164-178.2004>
- Liu, L., Li, W., Song, W., & Guo, M. (2018). Remediation Techniques for Heavy Metal-contaminated Soils: Principles and Applicability. *Science of The Total Environment*, 633, 206–219.
- Liu, W., Li, L., Khan, Md. , A., & Zhu, F. (2012). Popular Molecular Markers in Bacteria. *Molecular Genetics, Microbiology, and Virology*, 27(3), 103–107.
- Long, D., Tang, X., Cai, K., Chen, G., Shen, C., Shi, J., Chen, L., & Chen, Y. (2013). Cr(VI) Resistance and removal by Indigenous Bacteria Isolated from Chromium-contaminated Soil. *Journal of Microbiology and Biotechnology*, 23(8), 1123–1132. <https://doi.org/10.4014/jmb.1301.01004>
- Madigan, M. T., Martinko, J. M., Bender, K. S., Buckley, D. H., & Stahl, D. A. (2015). *Brock Biology Of Microorganisms: Vol. 14th Edition*. Boston: Pearson Education.
- Mala, J. G. S. M., Takeuchi, S., Dhanasingh Sujatha, & Mani, U. (2020). Microbial Chromate Reductases: Novel and Potent Mediators in Chromium Bioremediation-A Review. *Applied Microbiology: Theory & Technology*, 32–44. <https://doi.org/10.37256/amtt.112020222>
- Matzinger, K. (2004). The Importance of 16S rRNA Bacterial Spore Identification. *Spore News*, 1(4). Lakewood: Mesa Labs
- Meitiniarti, V. I., Putri, E. K., Runutu, A. E., Nugroho, R. A., & Kasmiyati, S. (2022). Isolation and identification of Cr-resistant bacteria from the rhizosphere of *Tagetes* sp. and their ability to reduce Cr (VI). *Biodiversitas Journal of Biological Diversity*, 23(8).
- Mitri, S., & Richard Foster, K. (2013). The Genotypic View of Social Interactions in Microbial Communities. *Annual Review of Genetics*, 47, 247–273. <https://doi.org/10.1146/annurev-genet-111212-133307>
- Mohapatra, R. K., Pandey, S., Thatoi, H., & Panda, C. R. (2017). Reduction of chromium (VI) by marine bacterium *Brevibacillus laterosporus* under varying saline and pH conditions. *Environmental Engineering Science*, 34(9), 617–626. <https://doi.org/10.1089/ees.2016.0627>
- Mohseni, M., Khosravi, F., & Chaichi, M. J. (2014). Bioremediation Activity of Pb (II) Resistance *Citrobacter* sp. MKH2 Isolated from Heavy Metal Contaminated Sites in Iran. *Journal of Sciences, Islamic Republic of Iran*. 25(2), 105–110. <https://www.researchgate.net/publication/285132288>

- Monga, A., Fulke, A. B., & Dasgupta, D. (2022). Recent Developments in Essentiality of Trivalent Chromium and Toxicity of Hexavalent Chromium: Implications on Human Health and Remediation Strategies. *Journal of Hazardous Materials Advances*, 7. <https://doi.org/10.1016/j.hazadv.2022.100113>
- Mueller-Dombois, D., & Ellenberg, H. (1974). *Aims and Methods of Vegetation Ecology*. New York: John Wiley & Sons. <https://doi.org/10.2307/213332>
- Munawwaroh, I., Saliaputri, L., Herdiyani, S. M., Puspasari, T. T., Winarni, S., Studi, P., Statistika, S., Mipa, F., Padjadjaran, U., & Mei, D. (2023). Implementasi Analisis Variansi Pada Desain Bujur Sangkar Youden Untuk Eksperimen. *Equator: Journal of Mathematical and Statistical Sciences (EJMSS)*, 2(1), 10–16.
- Murray, P. R., Rosenthal, K. S., & Pfaller, M. A. (2016). *Medical Microbiology* (8th ed.). Philadelphia: Elsevier Inc.
- Murthy, M. K., Khandayataray, P., Padhiary, S., & Samal, D. (2023). A review on chromium health hazards and molecular mechanism of chromium bioremediation. Dalam *Reviews on Environmental Health*, Vol. 38, Nomor 3, (hlm. 461–478). <https://doi.org/10.1515/reveh-2021-0139>
- Mythili, K., & Karthikeyan, B. (2011). Bioremediation of Cr (VI) from Tannery Effluent Using *Bacillus* spp and *I* spp. *International Multidisciplinary Research Journal*, 6, 38–41. <http://irjs.info/>
- Nagai, T., Iwamoto, E., Sakai, T., Arima, T., Tensha, K., Lida, Y., Lida, T., & Nakai, T. (2008). Characterization of *Edwardsiella ictaluri* Isolated from Wild ayu plecoglossus altivelis in Japan. *Fish Pathology*, 43(4). <https://doi.org/10.3147/jsfp.43.158>
- National Center for Biotechnology Information. (2024). *PubChem Element Summary for AtomicNumber 24, Chromium*. Diakses pada 4 November 2024 dari <https://pubchem.ncbi.nlm.nih.gov/element/Chromium>
- Nelson, D. L., & Cox, M. M. (2013). *Lehninger Principles of Biochemistry* (6th ed.). New York: W. H. Freeman and Company.
- Ningtyas, M. D., Isdiantoni, Ekawati, I., Koentjoro, M. P., & Prasetyo, E. N. (2020). “Microbial Consortium Synergism for Promising Freshwater Culture Probiotic”. Dalam H. Habiddin, S. Majid, S. Ibnu, N. Farida, & W. Dasna (Penyunting), *AIP Conference Proceedings Volume 2215*. <https://doi.org/10.1063/5.0000813>
- Niu, D., Zuo, Z., Shi, G. Y., & Wang, Z. X. (2009). High Yield Recombinant Thermostable  $\alpha$ -amylase Production Using an Improved *Bacillus licheniformis* system. *Microbial Cell Factories*, 8(58), 1-7. <https://doi.org/10.1186/1475-2859-8-58>
- Nugraha, C. A. (2019). *Isolasi dan Identifikasi Molekuler Bakteri Selulotik Pencernaan Rayap (Cryptotermes sp.) Menggunakan Gen 16S rRNA*. (Skripsi). Universitas Pendidikan Indonesia, Bandung.
- Oliveira, H. (2012). Chromium as an Environmental Pollutant: Insights on Induced Plant Toxicity. *Journal of Botany*, 2012, 1–8. <https://doi.org/10.1155/2012/375843>

- Osadebe, A. U., & Imasuen, K. (2024). Tolerance of *Xanthomonas campestris* for Heavy Metals Typically Associated with Brownfield Land. *Archives of Ecotoxicology*, 6(3), 65–71. <https://doi.org/10.36547/ae.2024.6.3.65-71>
- Ozimede, C. O., Obute, G. C., & Nyananyo, B. L. (2019). Morphological and Anatomical Diversity Study on three Species of *Amaranthus* namely; *A. hybridus* L., *A. viridis* L. and *A. spinosus* L. from Rivers State, Nigeria. *Journal of Applied Sciences and Environmental Management*, 23(10), 1875–1880. <https://dx.doi.org/10.4314/jasem.v23i10.17>
- Panda, S. K., & Choudhury, S. (2005). Chromium Stress in Plants. *Braz. J. Plant Physiol.*, 17(1).
- Panneerselvam, P., Choppala, G., Kunhikrishnan, A., & Bolan, N. (2013). Potential of Novel Bacterial Consortium for The Remediation of Chromium Contamination. *Water, Air, and Soil Pollution*, 224(12). <https://doi.org/10.1007/s11270-013-1716-9>
- Panneerselvam, P., Sharma, L., Nayak, A. K., Senapati, A., Tshering, J., Prabhukarthikeyan, B. S. R., Kumar, U., Kumar, A., Prasanna, G., Debasis, P., Anandan, M. A., & Nayak, P. K. (2020). Development of Microbial Consortium for Plant Growth Promotion, Nutrient and Disease Management in Rice-horticulture Based Cropping System in Sikkim. *NRRI Reasearch Bulletin*, 28.
- Paredes-Sabja, D., Shen, A., & Sorg, J. A. (2014). *Clostridium difficile* Spore Biology: Sporulation, Germination, and Spore Structural Proteins. *Trends in Microbiology*, 22(7), 406–416. <https://doi.org/10.1016/j.tim.2014.04.003>
- Patel, K. D., Bhanshali, F. C., Chaudhary, A. V., & Ingle, S. S. (2013). A new enrichment method for isolation of *Bacillus thuringiensis* from diverse sample types. *Applied biochemistry and biotechnology*, 170(1), 58–66. <https://doi.org/10.1007/s12010-013-0145-y>
- Pérez-Bustamante, I. S., Cruz-Flores, R., López-Carvallo, J. A., & Sánchez-Serrano, S. (2024). Effect of the 16S rRNA Gene Hypervariable Region on the Microbiome Taxonomic Profile and Diversity in the Endangered Fish *Totoaba macdonaldi*. *Microorganisms*, 12(11). <https://doi.org/10.3390/microorganisms12112119>
- Pérez-Cobas, A. E., Gomez-Valero, L., & Buchrieser, C. (2020). Metagenomic Approaches in Microbial Ecology: An Update on Whole-genome and Marker Gene Sequencing Analyses. *Microbial Genomics*, 6(8), 1–22. <https://doi.org/10.1099/mgen.0.000409>
- Pietikäinen, J., Pettersson, M., & Bååth, E. (2005). Comparison of Temperature Effects on Soil Respiration and Bacterial and Fungal Growth Rates. *FEMS Microbiology Ecology*, 52(1), 49–58. <https://doi.org/10.1016/j.femsec.2004.10.002>
- Processors & Growers Research Organisation. (2021). *Guide when Sampling Soil for Microbe Analysis*. Diakses pada 27 Januari 2025 dari <https://www.pgro.org/>

- Rahayu, D. R., & Mangkoedihardjo, S. (2022). Kajian Bioaugmentasi untuk Menurunkan Konsentrasi Logam Berat di Wilayah Perairan Menggunakan Bakteri (Studi Kasus: Pencemaran Merkuri di Sungai Krueng Sabee, Aceh Jaya). *Jurnal Teknik ITS*, 11(1), 15–22.
- Ramli, N. N., Othman, A. R., Kurniawan, S. B., Abdullah, S. R. S., & Hasan, H. A. (2023). Metabolic pathway of Cr(VI) reduction by bacteria: A review. *Microbiological Research*, 268.
- Ravisankar, V., Joshua, R. E., & Rajkumar, D. (2024). Synergistic effects of microbes and plant for remediation of chromium contaminated soil. *Global NEST Journal*, 26(1).
- Reddy, G. K. K., Kavibharathi, K., Singh, A., & Nancharaiah, Y. V. (2023). Growth-dependent Cr(VI) Reduction by *Alteromonas* sp. Under Haloalkaline Conditions: Toxicity, Removal Mechanism and Effect of Heavy Metals. *Research Square*. <https://doi.org/10.21203/rs.3.rs-3590297/v1>
- Rinanda, T. (2011). Analisis Sekuensi 16S rRNA di Bidang Mikrobiologi. *Jurnal Kedokteran Syiah Kuala*, 3, 172–177. <http://rdp.cme.msu.edu/html/>
- Ripanda, A., Luanda, A., Sule, K. S., Mtabazi, G. S., & Makangara, J. J. (2023). *Galinsoga parviflora* (Cav.): A comprehensive review on ethnomedicinal, phytochemical and pharmacological studies. *Heliyon*, 9(2023). <https://doi.org/10.1016/j.heliyon.2023.e13517>
- Robles-Rodríguez, C., Cardoso-Carmona, D., González-Dávalos, L., Lozano-Flores, C., Páez-Trejo, A., Shimada, A., & Mora, O. (2023). Characterization of an Acidogenic Bacterial Consortium as Probiotic and Its Effect on Rumen Fermentation In Vitro and In Vivo. *Ruminants*, 3(4), 324–346. <https://doi.org/10.3390/ruminants3040028>
- Rosmania, & Yanti, F. (2020). Perhitungan Jumlah Bakteri di Laboratorium Mikrobiologi Menggunakan Pengembangan Metode Spektrofotometri. *Jurnal Penelitian Sains*, 22(2), 76–86. <http://ejurnal.mipa.unsri.ac.id/index.php/jps/index>
- Rousk, J., Bååth, E., Brookes, P. C., Lauber, C. L., Lozupone, C., Caporaso, J. G., Knight, R., & Fierer, N. (2010). Soil Bacterial and Fungal Communities Across a pH Gradient in an Arable Soil. *ISME Journal*, 4(10), 1340–1351. <https://doi.org/10.1038/ismej.2010.58>
- Roychowdhury, R., Zaman, S., Roy, M., & Mitra, A. (2022). Isolation and Characterization of Arsenic and Cadmium Resistant *Brevibacillus* sp. from Fly Ash Sample of Mejia Thermal Power Station (MTPS), West Bengal, India. *Techno International Journal of Health, Engineering, Management and Science*, 2, 35–39.
- Ryan, K. J., & Ray, C. G. (2004). *Sherris Medical Microbiology* (4th ed.). New York: McGraw-Hill.
- Sanchez-Hachair, A., & Hofmann, A. (2018). Hexavalent Chromium Quantification in Solution: Comparing Direct UV-visible Spectrometry with 1,5-diphenylcarbazide Colorimetry. *Comptes Rendus Chimie*, 21(9), 890–896. <https://doi.org/10.1016/j.crci.2018.05.002>

- Sanjay, M. S., Sudarsanam, D., Raj, G. A., & Baskar, K. (2020). Isolation and identification of chromium reducing bacteria from tannery effluent. *Journal of King Saud University - Science*, 32(1), 265–271. <https://doi.org/10.1016/j.jksus.2018.05.001>
- Sarankumar, R. K., Arulprakash, A., Devanesan, S., Selvi, A., AlSalhi, M. S., Rajasekar, A., & Ahamed, A. (2020). Bioreduction of Hexavalent Chromium by Chromium Resistant Alkalophilic Bacteria Isolated from Tannery Effluent. *Journal of King Saud University - Science*, 32(3), 1969–1977. <https://doi.org/10.1016/j.jksus.2020.02.010>
- Sari, A. P. N. (2015). *Resistensi dan Potensi Bacillus Sebagai Bioremoval Logam Kromium (Cr)*. (Skripsi). Institut Teknologi Sepuluh Nopember, Surabaya.
- Sari, D. P. (2022). *Potensi Kultur Isolat Bakteri dan Fungi Sebagai Agen Bioremediasi Logam Krom Secara In Vitro*. (Skripsi). Universitas Pendidikan Indonesia, Bandung.
- Shan, B., Hao, R., Xu, H., Zhang, J., Li, J., Li, Y., & Ye, Y. (2022). Hexavalent Chromium Reduction and Bioremediation Potential of *Fusarium proliferatum* S4 Isolated from Chromium-contaminated soil. *Environmental Science and Pollution Research*, 29(52). <https://doi.org/10.1007/s11356-022-21323-6>
- Shanker, A. K., Cervantes, C., Loza-Tavera, H., & Avudainayagam, S. (2005). Chromium Toxicity in Plants. *Environment International*, 31(5), 739–753. <https://doi.org/10.1016/j.envint.2005.02.003>
- Songnaka, N., Nisoa, M., Atipairin, A., Wanganuttara, T., & Chinnawong, T. (2022). Enhanced Antibacterial Activity of *Brevibacillus* sp. SPR19 by Atmospheric and Room Temperature Plasma Mutagenesis (ARTP). *Scientia Pharmaceutica*, 90(2). <https://doi.org/10.3390/scipharm90020023>
- Sugiyono. (2013). *Metode Penelitian Kuantitatif Kualitatif dan R&D*. Bandung: Alfabeta.
- Sulistiani, W. S., Widowati, H., Sari, K., & Sutanto, A. (2021). Effect of chromium metal accumulation on the magnesium absorption and chlorophyll content in vegetables. *Makara Journal of Science*, 25(1), 5.
- Sunandi, E., Nugroho, S., & Rizal, J. (2009). Rancangan Acak Lengkap Dengan Subsample. *E-jurnal Statistika*. 80-101.
- Suparman, N. F. N., & Retnaningrum, E. (2023). Isolasi, Identifikasi, dan Uji Potensi Bakteri Laut sebagai Agen Bioremediasi Logam Berat Kromium (Cr). *Jurnal Sumberdaya Alam dan Lingkungan*, 10(3), 114–125. <https://doi.org/10.21776/ub.jsal.2023.010.03.3>
- Tariq, M., Zahoor, M., Yasmeen, T., Naqqash, T., Rashid, M. A. R., Abdullah, M., Rafiq, A. R., Zafar, M., Irfan, I., & Rasul, I. (2025). Biocontrol efficacy of *Bacillus licheniformis* and *Bacillus amyloliquefaciens* against rice pathogens. *PeerJ*, 13. <https://doi.org/10.7717/peerj.18920>
- Tjitosoepomo, G. (2018). *Morfologi tumbuhan*. Yogyakarta: Gadjah Mada University Press.
- Upadhyay, N., Vishwakarma, K., Singh, J., Mishra, M., Kumar, V., Rani, R., Mishra, R. K., Chauhan, D. K., Tripathi, D. K., & Sharma, S. (2017). Tolerance and Reduction of Chromium(VI) by *Bacillus* sp. MNU16 Isolated

- from Contaminated Coal Mining Soil. *Frontiers in Plant Science*, 8. <https://doi.org/10.3389/fpls.2017.00778>
- Usmadi. (2020). Pengujian Persyaratan Analisis. *Inovasi Pendidikan*, 7(1), 50–62.
- Vijayaraj, A. S., Mohandass, C., Joshi, D., & Rajput, N. (2018). Effective bioremediation and toxicity assessment of tannery wastewaters treated with indigenous bacteria. *3 Biotech*, 8(10). <https://doi.org/10.1007/s13205-018-1444-3>
- Vos, M., Quince, C., Pijl, A. S., de Hollander, M., & Kowalchuk, G. A. (2012). A Comparison of rpoB and 16S rRNA as Markers in Pyrosequencing Studies of Bacterial Diversity. *PLoS ONE*, 7(2). <https://doi.org/10.1371/journal.pone.0030600>
- Wafi, O. M. A. A., (2025). *Bioremediasi Logam Kromium Menggunakan Kombinasi Formula Dari Konsorsium Bakteri Bacillus parllicheniformis, Bacillus licheniformis, dan Brevibacillus borstelensis*. (Skripsi). Universitas Pendidikan Indonesia, Bandung.
- Warmiati, W. (2024). Reaction Stability Test of Hexavalent Chromium Complex with 1,5-Diphenylcarbazide in Analysis using UV Visible Spectrophotometer. *Indonesian Journal of Chemical Research*, 11(3), 218–222. <https://doi.org/10.30598/ijcr>
- Wilbur, S., Abadin, H., Fay, M., Yu, D., & Tencza, B. (2012). *Toxicological Profile For Chromium*. Atlanta: Agency for Toxic Substances and Disease Registry. <https://stacks.cdc.gov/view/cdc/12385>
- Willey, J. M. ., Sandman, K. M. ., & Wood, D. H. . (2020). *Prescott's microbiology (11th ed.)*. New York: McGraw-Hill Education.
- Woese, C. R. (1987). Bacterial Evolution. *Microbiological Reviews*, 51(2), 221–271.
- Wuana, R. A., & Okieimen, F. E. (2011). Heavy Metals in Contaminated Soils: A Review of Sources, Chemistry, Risks and Best Available Strategies for Remediation. *ISRN Ecology*, 2011, 1–20. <https://doi.org/10.5402/2011/402647>
- Yan, G., Gao, Y., Xue, K., Qi, Y., Fan, Y., Tian, X., Wang, J., Zhao, R., Zhang, P., Liu, Y., & Liu, J. (2023). Toxicity Mechanisms and Remediation Strategies for Chromium Exposure in the Environment. *Frontiers in Environmental Science*. 11. <https://doi.org/10.3389/fenvs.2023.1131204>
- Yu, J., Kim, P. D., Jang, Y., Kim, S. K., Han, J., & Min, J. (2022). Comparison of polylactic acid biodegradation ability of *Brevibacillus brevis* and *Bacillus amyloliquefaciens* and promotion of PLA biodegradation by soytone. *Biodegradation*, 33(5), 477–487. <https://doi.org/10.1007/s10532-022-09993-y>
- Zhalnina, K., Dias, R., de Quadros, P. D., Davis-Richardson, A., Camargo, F. A. O., Clark, I. M., McGrath, S. P., Hirsch, P. R., & Triplett, E. W. (2015). Soil pH Determines Microbial Diversity and Composition in the Park Grass Experiment. *Microbial Ecology*, 69(2), 395–406. <https://doi.org/10.1007/s00248-014-0530-2>