

DAFTAR PUSTAKA

- Abbas, A. K., Lichtman, A. H., & Pillai, S. (2015). *Cellular and Molecular Immunology – Edisi 8* dari [https://books.google.co.id/books?hl=en&lr=&id=RWYWBAQAQBAJ&oi=fnd&pg=PP1&dq=\(Abbas+AK,+Lichtman+A H,+Pillai+S,+2015\)&ots=fv17e3AAH0&sig=wrL8I7GsW6MUccYECeZvzCuFSHk&redir_esc=y#v=onepage&q&f=false](https://books.google.co.id/books?hl=en&lr=&id=RWYWBAQAQBAJ&oi=fnd&pg=PP1&dq=(Abbas+AK,+Lichtman+A H,+Pillai+S,+2015)&ots=fv17e3AAH0&sig=wrL8I7GsW6MUccYECeZvzCuFSHk&redir_esc=y#v=onepage&q&f=false), diakses pada 4 Maret 2024.
- Anonim. “Pedoman Pencegahan dan Pengendalian MPOX (Monkeypox) 2023.” Diakses pada 10 Juni. 2023. dari <https://infeksiemerging.kemkes.go.id/document/pedoman-pencegahan-dan-pengendalian-MPOX-Monkeypox-2023/view>, diakses pada 10 Juni 2023
- Armanto Makmun, S. F. H. (2020). TINJAUAN TERKAIT PENGEMBANGAN VAKSIN COVID – 19. *Molucca Medica*, 13.
- Boltjes, A., Movita, D., Boonstra, A., & Woltman, A. M. (2014). “The role of Kupffer cells in hepatitis B and hepatitis C virus infections”. *Journal of hepatology*, 61(3), Halaman 660–671. doi; 10.1016/j.jhep.2014.04.026, diakses pada 4 Maret 2024.
- Eslami, S., Glassy, M. C., & Ghafouri-Fard, S. (2022). “A comprehensive overview of identified mutations in SARS CoV-2 spike glycoprotein among Iranian patients.” *Gene*, 813, 146113. doi: 10.1016/j.gene.2021.146113 diakses pada 30 November 2023.
- Garg, A., & Gupta, D. (2008). “VirulentPred: a SVM based prediction method for virulent proteins in bacterial pathogens”. *BMC bioinformatics*, Halaman 9, 62. doi; 10.1186/1471-2105-9-62, diakses pada 5 Maret 2024.
- Grote, A., Hiller, K., Scheer, M., Münch, R., et al. (2005). “JCat: a novel tool to adapt codon usage of a target gene to its potential expression host.” *Nucleic acids research*, 33(Web Server issue), Halaman 526–531. doi: 10.1093/nar/gki376, diakses pada 3 Maret 2024.

- Mubarak, A., Alturaiki, W., & Hemida, M. G. (2019). "Middle East Respiratory Syndrome Coronavirus (MERS-CoV): Infection, Immunological Response, and Vaccine Development". *Journal of immunology research*, 2019, 6491738. <https://doi.org/10.1155/2019/6491738>, diakses pada 15 Maret 2024.
- Babkin, I. v., Babkina, I. N., & Tikunova, N. v. (2022). "An Update of Orthopoxvirus Molecular Evolution". *Viruses 2022, Vol. 14*, Halaman 388, 14(2), 388. <https://doi.org/10.3390/V14020388>, diakses pada 5 Maret 2024.
- Behmard, E., Soleymani, B., Najafi, A., & Barzegari, E. (2020). "Immunoinformatic design of a COVID-19 subunit vaccine using entire structural immunogenic epitopes of SARS-CoV-2". *Scientific Reports*, 10(1). doi: 10.1038/S41598-020-77547-4, diakses pada 7 Maret 2024.
- Latimer, B., Toporovski, R., Yan, J., Pankhong, P., Morrow, et al. (2014). "Strong HCV NS3/4a, NS4b, NS5a, NS5b-specific cellular immune responses induced in Rhesus macaques by a novel HCV genotype 1a/1b consensus DNA vaccine." *Human vaccines & immunotherapeutics*, 10(8), Halaman 2357–2365. doi: 10.4161/hv.29590, diakses pada 8 Maret 2024.
- Levast, B., Awate, S., Babiuk, L., Mutwiri, G. et al (2014). "Vaccine Potentiation by Combination Adjuvants". *Vaccines*, 2(2), Halaman 297–322. doi: 10.3390/vaccines2020297, diakses pada 24 Maret 2024
- Yu, C. S., Cheng, C. W., Su, W. C., Chang, K. et al (2014). "CELLO2GO: a web server for protein subCELLular LOcalization prediction with functional gene ontology annotation." *PloS one*, 9(6), e99368. doi: 10.1371/journal.pone.0099368, diakses pada 4 Maret 2024
- CDC. (2022, December 16). *2022 MPOX Outbreak Global Map | MPOX | Poxvirus | CDC*. <https://www.cdc.gov/poxvirus/Monkeypox/response/2022/world-map.html>, diakses pada 4 April 2024

- Chaplin D. D. (2010). "Overview of the immune response." *The Journal of allergy and clinical immunology*, 125(2 Suppl 2), Halaman 3–23. doi: 10.1016/j.jaci.2009.12.980, diakses pada 10 Maret 2024
- Jones D. T. (1999). "Protein secondary structure prediction based on position-specific scoring matrices." *Journal of molecular biology*, 292(2), Halaman 195–202. doi: 10.1006/jmbi.1999.3091, diakses pada 1 Maret 2024.
- Kozakov, D., Beglov, D., Bohnuud, T., Mottarella, S. E et al (2013). "How good is automated protein docking?". *Proteins*, 81(12), Halaman 2159–2166. doi: 10.1002/prot.24403, diakses pada 25 Maret 2024.
- Kozakov, D., Hall, D. R., Xia, B., Porter, K. A., et al (2017). "The ClusPro web server for protein-protein docking." *Nature protocols*, 12(2), Halaman 255–278. doi: 10.1038/nprot.2016.169, diakses pada 10 April 2024.
- Gasteiger, E., Hoogland, C., Gattiker, A., Duvaud, S., et al (2005). "Protein Identification and Analysis Tools on the ExPASy Server". *The Proteomics Protocols Handbook*. Halaman 571–607. doi: 10.1385/1-59259-890-0:571, diakses pada 19 Juli 2024.
- Farzan, M., Farzan, M., Mirzaei, Y., Aliman, S., et al (2023). "Immunoinformatics-based multi-epitope vaccine design for the re-emerging Monkeypox virus." *International immunopharmacology*, 123, 110725. doi: 10.1016/j.intimp.2023.110725, diakses pada 12 Januari 2024.
- Fleri, W., Paul, S., Dhanda, S. K., Mahajan, S., Xu, X., Peters, B., & Sette, A. (2017). "The Immune Epitope Database and Analysis Resource in Epitope Discovery and Synthetic Vaccine Design." *Frontiers in immunology*, 8, Halaman 278. doi: 10.3389/fimmu.2017.00278, diakses pada 30 November 2023
- Castiglione, F., Mantile, F., De Berardinis, P., & Prisco, A. (2012). "How the interval between prime and boost injection affects the immune response in a computational model of the immune system." *Computational and*

- mathematical methods in medicine*, 2012, 842329. doi: 10.1155/2012/842329, diakses pada 5 April 2024.
- Vogel F. R. (2000). "Improving vaccine performance with adjuvants." *Clinical infectious diseases : an official publication of the Infectious Diseases Society of America*, 30 Suppl 3, Halaman 266–270. doi: 10.1086/313883, diakses pada 7 April 2024.
- Flower, D. R. (2007). "Immunoinformatics and the in silico prediction of immunogenicity". An introduction. *Methods in Molecular Biology (Clifton, N.J.)*, 409, Halaman 1–15. doi: 10.1007/978-1-60327-118-9_1/COVER, diakses pada 7 April 2024.
- Gamage, D. G., Gunaratne, A., Periyannan, G. R., & Russell, T. G. (2019). "Applicability of Instability Index for In vitro Protein Stability Prediction." *Protein and peptide letters*, 26(5), Halaman 339–347. doi: 10.2174/0929866526666190228144219, diakses pada 12 Maret 2024.
- Ghafouri, F., Conan, R. A., Noorbakhsh, F., Samimi, H., & Haghpanah, V. (2020). "An in-silico approach to develop of a multi-epitope vaccine candidate against SARS-CoV-2 envelope (E) protein." Research square, rs.3.rs-30374. doi: 10.21203/rs.3.rs-30374/v1, diakses pada 30 November 2023
- Nagpal, G., Usmani, S. S., Dhanda, S. K., Kaur, H., et al (2017). "Computer-aided designing of immunosuppressive peptides based on IL-10 inducing potential". *Scientific Report*, 7(42851). Halaman 3-6. doi: 10.1038/srep42851, diakses pada 19 Juli 2024.
- Harris E. (2022). "What to Know About Monkeypox." *JAMA*, 327(23), Halaman 2278–2279. doi: 10.1001/jama.2022.9499, diakses pada 19 Juli 2024.
- Hidayat, S., & Syahputra, A. A. (2020). "Sistem Imun Tubuh Pada Manusia". *Visual Heritage: Jurnal Kreasi Seni dan Budaya*, vol 2(3), Halaman.144-149. file:///C:/Users/user/Downloads/898-3487-1-PB%20(2).pdf, diakses pada 18 Maret 2024.

- Doytchinova, I. A., & Flower, D. R. (2007). "VaxiJen: a server for prediction of protective antigens, tumour antigens and subunit vaccines." *BMC bioinformatics*, 8, Halaman 4. doi: 10.1186/1471-2105-8-4, diakses pada 18 Maret 2024.
- Ladnyj, I. D., Ziegler, P., & Kima, E. (1972). "A human infection caused by monkeypox virus in Basankusu Territory, Democratic Republic of the Congo." *Bulletin of the World Health Organization*, 46(5), Halaman 593–59, <https://www.ncbi.nlm.nih.gov/pmc/articles/PMC2480792/>, diakses pada 14 April 2024.
- Dimitrov, I., Flower, D. R., & Doytchinova, I. (2013). "AllerTOP--a server for in silico prediction of allergens." *BMC bioinformatics*, 14 Suppl 6(Suppl 6), Halaman 4. doi: 10.1186/1471-2105-14-S6-S4, diakses pada 14 April 2024.
- Dimitrov, I., Naneva, L., Doytchinova, I., & Bangov, I. (2014). "AllergenFP: allergenicity prediction by descriptor fingerprints." *Bioinformatics (Oxford, England)*, 30(6), Halaman 846–851. doi: 10.1093/bioinformatics/btt619, diakses pada 15 April 2024.
- Yang, J. X., Tseng, J. C., Yu, G. Y., Luo, Y., et al (2022). "Recent Advances in the Development of Toll-like Receptor Agonist-Based Vaccine Adjuvants for Infectious Diseases." *Pharmaceutics*, 14(2), Halaman 423. doi: 10.3390/pharmaceutics14020423, diakses pada 16 April 2024.
- Thakar, J., Pilione, M., Kirimanjeswara, G., Harvill, E. T., & Albert, R. (2007). "Modeling systems-level regulation of host immune responses." *PLoS computational biology*, 3(6), e109. doi: 10.1371/journal.pcbi.0030109, diakses pada 16 April 2024.
- Kemkes. (2023). *Kasus Monkeypox Mulai Terdeteksi di Indonesia*. <https://upk.kemkes.go.id/new/kasus-Monkeypox-mulai-terdeteksi-di-indonesia>, diakses pada 13 Januari 2024.

- Kemkes. (2022). *Penyakit Cacar Monyet (Monkeypox) dan yang Perlu Kita Tahu Tentangnya*. <http://www.b2p2vrp.litbang.kemkes.go.id/mobile/berita/baca/419/Penyakit-Cacar-Monyet-Monkeypox-dan-yang-Perlu-Kita-Tahu-Tentangnya>, diakses pada 12 Januari 2024.
- Kmiec, D., & Kirchhoff, F. (2022). "Monkeypox: A New Threat?". *International journal of molecular sciences*, 23(14), 7866. doi: 10.3390/ijms23147866, diakses pada 8 Maret 2024
- Krisnasari, A. S. (2018). *Imunoserologi: Teknologi Laboratorium Medik*. Jakarta: Penerbit Buku Kedokteran EGC
- Lo, Y. T., Pai, T. W., Wu, W. K., & Chang, H. T. (2013). "Prediction of conformational epitopes with the use of a knowledge-based energy function and geometrically related neighboring residue characteristics. *BMC Bioinformatics*, 14(SUPPL4), Halaman 1–10. doi: 10.1186/1471-2105-14-S4-S3/TABLES/3, diakses pada 8 Mei 2024.
- Zhang L. (2018). "Multi-epitope vaccines: a promising strategy against tumors and viral infections." *Cellular & molecular immunology*, 15(2), Halaman 182–184. doi: 10.1038/cmi.2017.92, diakses pada 8 Mei 2024.
- McGuffin, L. J., Bryson, K., & Jones, D. T. (2000). "The PSIPRED protein structure prediction server." *Bioinformatics (Oxford, England)*, 16(4), Halaman 404–405. doi: /10.1093/bioinformatics/16.4.404, diakses pada 6 Maret 2024.
- Magnus, P. von, Andersen, E. K., Petersen, K. B., & Birch-Andersen, A. (1959). "A POX-LIKE DISEASE IN CYNOMOLGUS MONKEYS." *Acta Pathologica Microbiologica Scandinavica*, 46(2), Halaman 156–176. doi: 10.1111/J.1699-0463.1959.TB00328, diakses pada 6 Maret 2024.
- Malonis, R. J., Lai, J. R., & Vergnolle, O. (2020). "Peptide-Based Vaccines: Current Progress and Future Challenges." *Chemical reviews*, 120(6),

- Halaman 3210–3229. <https://doi.org/10.1021/acs.chemrev.9b00472>, diakses pada 6 Maret 2024.
- Mansur, D. S., Smith, G. L., & Ferguson, B. J. (2014). “Intracellular sensing of viral DNA by the innate immune system.” *Microbes and Infection*, 16(12), Halaman 1002–1012. doi: 10.1016/J.MICINF.2014.09.010, diakses pada 8 April 2024.
- Marco, M. del M. F. de, Alejo, A., Hudson, P., Damon, I. K., & Alcami, A. (2010). “The highly virulent variola and Monkeypox viruses express secreted inhibitors of type I interferon.” *The FASEB Journal*, 24(5), Halaman 1479–1488. doi: 10.1096/FJ.09-144733, diakses pada 10 April 2024.
- Murphy, Kenneth dan Casey Weaver. (2017) *Immunobiology*. Washington Garland Science, Taylor & Francis Group, LLC.
- Netea, M. G., Van der Meer, J. W., Sutmoller, R. P., Adema, G. J., & Kullberg, B. J. (2005). “From the Th1/Th2 paradigm towards a Toll-like receptor/T-helper bias.” *Antimicrobial agents and chemotherapy*, 49(10), Halaman 3991–3996. doi: 10.1128/AAC.49.10.3991-3996.2005, diakses pada 10 April 2024.
- Saadi, M., Karkhah, A., & Nouri, H. R. (2017). “Development of a multi-epitope peptide vaccine inducing robust T cell responses against brucellosis using immunoinformatics based approaches.” *Infection, genetics and evolution : journal of molecular epidemiology and evolutionary genetics in infectious diseases*, 51, Halaman 227–234. doi: 10.1016/j.meegid.2017.04.009, diakses pada 8 Juli 2024.
- Kavoosi, M., Creagh, A. L., Kilburn, D. G., & Haynes, C. A. (2007). “Strategy for selecting and characterizing linker peptides for CBM9-tagged fusion proteins expressed in *Escherichia coli*.” *Biotechnology and bioengineering*, 98(3), Halaman 599–610. doi: 10.1002/bit.21396, diakses pada 23 April 2024.

- Shah, M., Anwar, M. A., Kim, J. H., & Choi, S. (2016). "Advances in Antiviral Therapies Targeting Toll-like Receptors." *Expert opinion on investigational drugs*, 25(4), Halaman 437–453. doi: 10.1517/13543784.2016.1154040, diakses pada 19 April 2024.
- Van Regenmortel MHV (1996). "Mapping Epitope Structure and Activity: From One-Dimensional Prediction to Four-Dimensional Description of Antigenic Specificity." *Methods (San Diego, Calif.)*, 9(3), Halaman 465–472. doi: 10.1006/meth.1996.0054, diakses pada 19 April 2024.
- Soltan, M. A., Eldeen, M. A., Elbassiouny, N., Kamel, H. L., et al (2021). "In Silico Designing of a Multitope Vaccine against *Rhizopus microsporus* with Potential Activity against Other Mucormycosis Causing Fungi". *Cells*, 10(11), Halaman 3014. doi: 10.3390/cells10113014. diakses pada 22 April 2024.
- Akhtar, N., Kaushik, V., Grewal, R. K., Wani, A. K., et al (2022). "Immunoinformatics-Aided Design of a Peptide Based Multiepitope Vaccine Targeting Glycoproteins and Membrane Proteins against Monkeypox Virus." *Viruses*, 14(11), Halaman 2374. doi: 10.3390/v14112374 Diakses pada 22 April 2024.
- Rapin, N., Lund, O., Bernaschi, M., & Castiglione, F. (2010). "Computational immunology meets bioinformatics: the use of prediction tools for molecular binding in the simulation of the immune system." *PloS one*, 5(4), Edisi 9862. doi: 10.1371/journal.pone.0009862, diakses pada 1 April 2024.
- Raghav, P. K., Kumar, R., Kumar, V., & Raghava, G. P. S. (2019). "Docking-based approach for identification of mutations that disrupt binding between Bcl-2 and Bax proteins: Inducing apoptosis in cancer cells." *Molecular genetics & genomic medicine*, 7(11), Edisi 910. doi: 10.1002/mgg3.910, diakses pada 1 April 2024
- Raghav, P. K., Raghav, A., Lathwal, A., Saxena, A., (2023). "Experimental and clinical data analysis for identification of COVID-19 resistant ACE2

- mutations.” *Scientific reports*, 13(1), Halaman 2351. doi: 10.1038/s41598-022-20773-9, diakses pada 2 April 2024
- Radji, M. (2009). Vaksin DNA: Vaksin Generasi Keempat. *Majalah Ilmu Kefarmasian*, 6(1), Halaman 4. doi: 10.7454/psr.v6i1.3433, diakses pada 1 April 2024
- Rao, A. K., Petersen, B. W., Whitehill, F., Razeq, J. H., (2022). “Use of JYNNEOS (Smallpox and Monkeypox Vaccine, Live, Nonreplicating) for Preexposure Vaccination of Persons at Risk for Occupational Exposure to Orthopoxviruses. Recommendations of the Advisory Committee on Immunization Practices — United States, 2022.” *MMWR. Morbidity and Mortality Weekly Report*, 71(22), Halaman 734–742. doi:10.15585/MMWR.MM7122E1, diakses pada 1 April 2024
- Reed, K. D., Melski, J. W., Beth Graham, M., Regnery, R. L., (2004). “The Detection of Monkeypox in Humans in the Western Hemisphere.” <https://doi.org/10.1056/NEJMoa032299>, 350(4) Halaman 342–350. doi: 10.1056/NEJMoa032299, diakses pada 18 April 2024
- Rokom. (2023). “Kasus Monkeypox Bertambah di Indonesia Akibat Sex Berisiko”. dari <https://sehatnegeriku.kemkes.go.id/baca/umum/20231023/1544066/kasus-monkeypox-bertambah-di-indonesia-akibat-sex-berisiko/>., diakses pada 1 November 2023.
- Rokom. (2022). “Monkeypox Menular Kalau Sudah Bergejala”. dari <https://sehatnegeriku.kemkes.go.id/baca/rilismedia/20220727/2840760/monkeypox-menular-kalau-sudah-bergejala/>., diakses pada 1 November 2023.
- Arai, R., Ueda, H., Kitayama, A., Kamiya, N., & Nagamune, T. (2001). “Design of the linkers which effectively separate domains of a bifunctional fusion protein.” *Protein engineering*, 14(8), Halaman 529–532. doi: 10.1093/protein/14.8.529 Diakses pada 19 April 2024

- Shantier, S. W., Mustafa, M. I., Abdelmoneim, A. H., Fadl, H. A., Elbager, S. G., & Makhawi, A. M. (2022). "Novel multi epitope-based vaccine against monkeypox virus: vaccinomic approach." *Scientific reports*, 12(1), Halaman 15983. doi: 10.1038/s41598-022-20397-z Diakses pada 19 April 2024
- Sharun, K., Tiwari, R., Patel, S. K., Karthik, K., et al (2020). Coronavirus disease 2019 (COVID-19) in domestic animals and wildlife: advances and prospects in the development of animal models for vaccine and therapeutic research. *Human Vaccines & Immunotherapeutics*, 16(12), Halaman 3043. doi: 10.1080/21645515.2020.1807802 Diakses pada 1 April 2024
- Shchelkunov, S. N., Totmenin, A. v., Safronov, P. F., Mikheev, M. v., et al (2002). "Analysis of the Monkeypox virus genome." *Virology*, 297(2), Halaman 172–194. doi: 10.1006/viro.2002.1446 Diakses pada 2 April 2024
- Stratmann, T. (2015). "Cholera Toxin Subunit B as Adjuvant—An Accelerator in Protective Immunity and a Break in Autoimmunity." *Vaccines*, 3(3), Halaman 579–596. doi: 10.3390/vaccines3030579 Diakses pada 8 April 2024
- Aiman, S., Alhamhoom, Y., Ali, F., Rahman, N., et al (2022). "Multi-epitope chimeric vaccine design against emerging Monkeypox virus via reverse vaccinology techniques- a bioinformatics and immunoinformatics approach." *Frontiers in immunology*, 13, Halaman 985450. doi: 10.3389/fimmu.2022.985450 Diakses pada 11 Mei 2024
- Abdi, S. A. H., Ali, A., Sayed, S. F., Abutahir, Ali, A., & Alam, P. (2022). "Multi-Epitope-Based Vaccine Candidate for Monkeypox: An In Silico Approach." *Vaccines*, 10(9), Halaman 1564. doi: 10.3390/vaccines10091564, diakses pada 18 April 2024.

- Islam, S. I., Mou, M. J., Sanjida, S., Tariq, M (2022). "Designing a novel mRNA vaccine against *Vibrio harveyi* infection in fish: an immunoinformatics approach." *Genomics & informatics*, 20(1), Edisi 11. doi: 10.5808/gi.21065, diakses pada 20 April 2024.
- Dhanda, S. K., Vir, P., & Raghava, G. P. (2013). "Designing of interferon-gamma inducing MHC class-II binders." *Biology direct*, 8, Halaman 30. doi: 10.1186/1745-6150-8-30, diakses pada 1 Juni 2024.
- Dhanda, S. K., Gupta, S., Vir, P., & Raghava, G. P. (2013). "Prediction of IL4 inducing peptides." *Clinical & developmental immunology*, 2013, 263952. doi: 10.1155/2013/263952, diakses pada 1 Juni 2024.
- Syafa'ah, I., & Yudhawati, R. (2019). "Peran Imunitas Mukosa terhadap Infeksi *Mycobacterium Tuberculosis*." *Jurnal Respirasi*, 2(2), Halaman 61. doi: 10.20473/JR.V2-I.2.2016.61-68, diakses pada 1 Juni 2024.
- Taupiqurrohman, O., Yusuf, M., Nuswantara, S., & Subroto, T. (2016). "PERANCANGAN VAKSIN VIRUS PAPILLOMA MANUSIA TIPE-16 BERBASIS EPIPOPE DENGAN BERBANTUKAN IMUNOINFORMATIKA." <https://doi.org/10.24114/jpkim.v8i3.4535>, diakses pada 2 Juni 2024.
- Tan, C., Zhu, F., Pan, P., Wu, A., & Li, C. (2023). "Development of multi-*epitope* vaccines against the *Monkeypox* virus based on envelope proteins using immunoinformatics approaches." *Frontiers in immunology*, 14, 1112816. <https://doi.org/10.3389/fimmu.2023.1112816>, diakses pada 2 Juni 2024.
- Kar, T., Narsaria, U., Basak, S., Deb, D., et al (2020). "A candidate multi-*epitope* vaccine against SARS-CoV-2." *Scientific reports*, 10(1), Halaman 10895. <https://doi.org/10.1038/s41598-020-67749-1>, diakses pada 1 Juni 2024.
- WHO. (2022a, May). *MPOX (Monkeypox) outbreak 2022 - Global*. dari <https://www.who.int/emergencies/situations/Monkeypox-oubreak-2022>, diakses pada 1 Juni 2024.

WHO. (2022b, June 27). *Multi-country Monkeypox outbreak: situation update*.
dari <https://www.who.int/emergencies/disease-outbreak-news/item/2022-DON396>, diakses pada 1 Juni 2024.

Hu, W., Yen, Y. T., Singh, S., Kao, C. L., et al (2012). "SARS-CoV regulates immune function-related gene expression in human monocytic cells." *Viral immunology*, 25(4), Halaman 277–288. doi: 10.1089/vim.2011.0099, diakses pada 4 Maret 2024.

McCollum, A. M., & Damon, I. K. (2014). "Human monkeypox." *Clinical infectious diseases : an official publication of the Infectious Diseases Society of America*, 58(2), Halaman 260–267. doi: 10.1093/cid/cit703, diakses pada 11 Juni 2024.



