

DAFTAR PUSTAKA

- Allocati, N., Masulli, M., Alexeyev, M. F., dan Di Ilio, C. (2013). *Escherichia coli* in Europe: An overview. *Int J Environ Res Public Health*, 10(12), 6235–6254. <https://doi.org/10.3390/ijerph10126235>
- Bagattini, M., Crivaro, V., Popolo, A. Di, Gentile, F., Scarcella, A., Triassi, M., ... Sperimentale, M. (2006). Molecular epidemiology of extended-spectrum β -lactamase-producing *Klebsiella pneumoniae* in a neonatal intensive care unit. *J Antimicrob Chemother*, 979–982. <https://doi.org/10.1093/jac/dkl077>
- Barfield, W. D. (2016). Standard terminology for fetal , infant , and perinatal deaths. *J Pediatr*, 137(5). <https://doi.org/10.1542/peds.2016-0551>
- Belanger, L., Garenaux, A., Harel, J., Boulianne, M., Nadeau, E., dan Dozois, C. M. (2011). *Escherichia coli* from animal reservoirs as a potential source of human extraintestinal pathogenic *E. coli*. *FEMS Immunol Med Microbiol*, 62, 1–10. <https://doi.org/10.1111/j.1574-695X.2011.00797.x>
- Bettelheim, Breadon, A., Faiers, M., O'farrell, S., dan Shooter. (1974). The origin of O serotypes of *Escherichia coli* in babies after normal delivery. *J Hyg Cam*, 72(67), 67–70.
- Bingen, E., Denamur, E., Brahimi, N., dan Elion, J. (1996). Genotyping may provide rapid identification of *Escherichia coli* kl organisms that cause neonatal meningitis. 152–156.
- Birchenough, G. M. H., Johansson, M. E. V, Stabler, R. A., Dalgakiran, F., Hansson, G. C., dan Wren, B. W. (2013). Altered innate defenses in the neonatal gastrointestinal tract in response to colonization by neuropathogenic *Escherichia coli*. *Infect Immun*, 81(9), 3264–3275. <https://doi.org/10.1128/IAI.00268-13>
- Bramantono, Purwati, dan Hamidah. (2013). The prevalence of extended spectrum beta-lactamase (ESBL) in third generation cephalosporin usage among sepsis patients in the department of internal medicine RSUD Dr . Soetomo Surabaya. *Folia Medica Indonesia*, 49(4), 244–251.
- Brikun, I., Suziedelis, K., dan Berg, D. E. (1994). DNA sequence divergence among derivatives of *Escherichia coli* k-12 detected by arbitrary primer PCR (Random Amplified Polymorphic DNA) fingerprinting. *J Bacteriol*, 176(6), 1673–1682.
- Brook, I., Barrett, T. C., Brinkman, R. C., Martin, J. William, dan Finegold, M. S. (1979). Aerobic and anaerobic bacterial flora of the maternal cervix and

- newborn gastric fluid and conjunctiva : prospective study. *Pediatrics*, 63(3), 451–455.
- Brook's F Geo, Carroll KC, Morse SA, Mietzner T. (2013). Jawetz, Melnick, Adelberg's Medical microbiology 26th edition. Medical Microbiology. jawetz, melnick, dan adelberg. Jakarta: penerbit Kedokteran EGC
- Burdet, C., dan Clermont, O. (2014). *Escherichia coli* bacteremia in children age and portal of entry are the main predictors of severity. *Pediatr Infect Dis*, 33(2), 872. <https://doi.org/10.1097/INF.0000000000000309>
- Cengiz, M., Buyukcangaz, E., Arslan, E., Mat, B., Sahinturk, P., Sonal, S., ... Sen, A. (2012). Papers molecular characterisation of quinolone resistance in *Escherichia coli* from animals in Turkey. *BMJ*. <https://doi.org/10.1136/vr.100719>
- Chaudhuri, R. R., dan Henderson, I. R. (2012). The evolution of the *Escherichia coli* phylogeny. *Infection, Genetics and Evolution*, 12(2), 214–226. <https://doi.org/10.1016/j.meegid.2012.01.005>
- Clermont, O., Bonacorsi, S., dan Bingen, E. (2000). Rapid and simple determination of the *Escherichia coli* phylogenetic group. *Appl Environ Microbiol*, 66(10), 4555–4558.
- Clermont, O., Christenson, J. K., Denamur, E., dan Gordon, D. M. (2013). The Clermont *Escherichia coli* phylo-typing method revisited : improvement of specificity and detection of new phylo-groups. *Environ Microbiol Rep*, 5(1), 58–65. <https://doi.org/10.1111/1758-2229.12019>
- Clermont, O., Olier, M., Hoede, C., Diancourt, L., Brisse, S., Picard, B., ... Keroudean, M. (2011). Animal and human pathogenic *Escherichia coli* strains share common genetic backgrounds. *Infect Genet Evol*, 11, 654–662. <https://doi.org/10.1016/j.meegid.2011.02.005>
- Cole, B. K., Scott, E., Ilikj, M., Bard, D., Akins, D. R., dan Dyer, D. W. (2017). Route of infection alters virulence of neonatal septicemia *Escherichia coli* clinical isolates. *PLOS ONE*, 12(12), 1–22.
- Coque, T. M., Novais, Â., Carattoli, A., Poirel, L., Pitout, J., Peixe, L., ... Nordmann, P. (2008). Dissemination of clonally related *Escherichia coli* strains expressing. *Emerg Infect Dis*, 14(2), 195–200.
- Cunningham GF, et al. 2012. *William Obstetrics*. 23rd. Jakarta: EGC

- Darmstadt, G. L., Zaidi, A. K. M., dan Stoll, B. J. (2011). Neonatal infection : A global perspective. In *Oxford University Press* (Vol. 23). <https://doi.org/10.1093/heapol/czn001>
- Das, P., Singh, A. K., Mukherjee, S., Rajendran, K., Saha, D. R., Koley, H., dan Basu, S. (2013). Composition of *Escherichia coli* population in the neonatal gut : phylogroups and virulence determinants. *J Med Microbiol*, 62, 1680–1687. <https://doi.org/10.1099/jmm.0.052225-0>
- Dash, M., Padhi, S., Mohanty, I., Panda, P., dan Parida, B. (2013). Antimicrobial resistance in pathogens causing urinary tract infections in a rural community of Odisha , India. *J Family Community Med*, 20(1), 20–26. <https://doi.org/10.4103/2230-8229.108180>
- Dubois, V., Barbeyrac, B. De, Rogues, A., Arpin, C., Coulange, L., Andre, C., ... Quentin, C. (2010). CTX-M-producing *Escherichia coli* in a maternity ward: a likely community importation and evidence of mother-to-neonate transmission . *J Antimicrob Chemother*, 65(May), 1368–1371. <https://doi.org/10.1093/jac/dkq153>
- Dusé, A. G. (2015). Infection control in developing countries with particular emphasis on South Africa Infection control in developing countries with particular emphasis on South Africa. *S Afr J Inf Dis*, 8782(2), 37–41. <https://doi.org/10.1080/10158782.2005.11441230>
- Emacar, J. (2010). Antibiotic resistance patterns of *Escherichia coli* isolated from HIV-sero positive adults at Mbagathi District Hospital , Nairobi . *J Appl Biosci*, 27, 1705–1714.
- Escobar-páramo, P., Le, A., Gall, T. Le, Amorin, C., Gouriou, S., Picard, B., ... Denamur, E. (2006). Identification of forces shaping the commensal *Escherichia coli* genetic structure by comparing animal and human isolates. *Environ Microbiol*, 8(11), 1975–1984. <https://doi.org/10.1111/j.1462-2920.2006.01077.x>
- European CDC. (2014). Systematic review of the effectiveness of infection control measures to prevent the transmission of carbapenemase-producing Enterobacteriaceae through cross-border transfer of patients. In *Stockholm: ECDC*. <https://doi.org/doi 10.2900/418192>
- Fernández-Prada, M., Martínez-Ortega, C., Santos-Simarro, G., Morán-Álvarez, P., Fernández-Verdugo, A., dan Costa-Romero, M. (2019). Outbreak of extended-spectrum beta-lactamase-producing *Klebsiella pneumoniae* in a neonatal intensive care unit: Risk factors and key preventive measures for eradication in record time. *Anales de Pediatría (English Edition)*, 91(1), 13–20. <https://doi.org/10.1016/j.anpede.2018.11.002>

- Flokas, M. E., Karanika, S., Alevizakos, M., dan Mylonakis, E. (2017). Prevalence of ESBL-producing Enterobacteriaceae in pediatric bloodstream infections: a systematic review and meta-analysis. *PLOS ONE*, 1–13. <https://doi.org/10.1371/journal.pone.0171216>
- García, C., Horna, G., Linares, E., Ramírez, R., Tapia, E., Velásquez, J., ... Jacobs, J. (2012). Antimicrobial drug resistance in Peru. *Emerg Infect Dis*, 18(3), 18–19.
- Goering, R. V. (2010). Pulsed field gel electrophoresis : A review of application and interpretation in the molecular epidemiology of infectious disease. *Infect Genet Evol*, 10(7), 866–875. <https://doi.org/10.1016/j.meegid.2010.07.023>
- Gupta, A., Latta, P. Della, Todd, B., San, P., Haas, J., Wu, F., dan Rubenstein, D. (2014). Beta -lactamase – producing *Klebsiella pneumoniae* in a neonatal intensive care unit linked to artificial nails. *Infect Control Hosp Epidemiol*, 25(May), 210–215.
- Happy, T. A., Setyarini, W., & Ranuh, IGM Reza Gunadi Kuntaman, K. (2020). Prevalence ESBL Producing *Escherichia Coli* among Children in Indonesia. *Indian J of Public Health Research and Development*, 11(05), 804–809.
- Hasanpour, M., dan Najafi, A. (2017). Development of a multiple real-time PCR assay for phylogenetic analysis of *Escherichia coli* strains. *J Microbiol Methods*. <https://doi.org/10.1016/j.mimet.2017.03.009>
- Huang, Y., Zhuang, S., dan Du, M. (2007). Risk factors of nosocomial infection with bacteria in a neonatal intensive care unit in China. *J Clin Epidemiol*, 35(5), 339–345. <https://doi.org/10.1007/s15010-007-6356-9>
- Ikechukwu, O., Ifeanyichukwu, I., David, I. O., Agwu, E. U., dan Ogbonnaya, O. (2017). Antibiotic resistance and beta-lactamase genes detection among extended spectrum beta-lactamase (ESBL) -producing *Escherichia coli* and salmonella species isolated from cockroaches (*periplaneta americana*) in Abakaliki , South-east Nigeria. *World J Med Sci*, 14(4), 113–120. <https://doi.org/10.5829/idosi.wjms.2017.113.120>
- Iranpour, D., Hassanpour, M., Ansari, H., Tajbakhsh, S., Khamisipour, G., dan Najafi, A. (2015). Phylogenetic groups of *Escherichia coli* strains from patients with urinary tract infection in Iran based on the new clermont phylotyping method. *Biomed Res Int*, 2015, 5–12.
- Jacoby, G. A., dan Munoz-price, L. S. (2005). The new B -lactamases. *N Engl J Med*, 352(4), 380–391.

- Jauregui, F., Landraud, L., Passet, V., Diancourt, L., Frapy, E., Guigon, G., ... Brisse, S. (2008). Phylogenetic and genomic diversity of human bacteremic *Escherichia coli* strains. *BMC Med Genomics*, 14(9), 1–14. <https://doi.org/10.1186/1471-2164-9-560> 44
- Johnson, J. R. ., dan Russo, T. A. (2002). Extraintestinal patogenic *Escherichia coli*: “The other bad *E. coli*.” *J Lab Clin Med*, 139(3), 155–162. <https://doi.org/10.1067/mlc.2002.121550>
- Johnson, J. R., Oswald, E., Bryan, T. T. O., dan Kuskowski, M. A. (2002). Phylogenetic distribution of virulence-associated genes among *Escherichia coli* isolates associated with neonatal bacterial meningitis in the Netherlands. *J Infect Dis*, 185, 774–783.
- Kaper, J. B., Nataro, J. P., dan Mobley, H. L. T. (2004). Patogenic *Escherichia coli*. *Nat Rev Microbiol*, 2(February), 123–140. <https://doi.org/10.1038/nrmicro818>
- Kar, D., Bandyopadhyay, S., Bhattacharyya, D., Samanta, I., Mahanti, A., Nanda, P. K., ... Singh, R. K. (2015). Molecular and phylogenetic characterization of multidrug resistant extended spectrum beta-lactamase producing *Escherichia coli* isolated from poultry and cattle in Odisha , India. *Infect Genet Evol*, 29, 82–90. <https://doi.org/10.1016/j.meegid.2014.11.003>
- Karim, A., Poirel, L., Nagarajan, S., dan Nordmann, P. (2001). Plasmid-mediated extended-spectrum B-lactamase (CTX-M-3 like) from India and gene association with insertion sequence IS Ecp1. *FEMS Microbiol Lett*, 201, 237–241.
- Kazemnia, A., Ahmadi, M., dan Dilmaghani, M. (2014). Antibiotic resistance pattern of different *Escherichia coli* phylogenetic groups isolated from human urinary tract infection and avian colibacillosis. *Iran Biomed J*, 18(4), 219–224. <https://doi.org/10.6091/ibj.1394.2014>
- Kementarian, K. (2018). *Profil kesehatan Indonesia 2017* (R. Kurniawan, Yudianto, B. Hardhana, dan T. Siswanti, Eds.). Retrieved from <http://www.kemkes.go.id>
- Kilic, A., MUZ, A., Ertas, H. B., dan Özbey, G. (2009). Random Amplified Polymorphic DNA (RAPD) analysis of *Escherichia coli* isolated from chickens. *XXII World's Poultry Congres*, 23(1), 1–4.
- Kumar, S., Kumar, M., Raj, A., dan Prakash, J. (2017). Evaluation of genetic analysis of *Escherichia coli* isolated from two different environmental sources : sewage water verses soiled bedding materials of laboratory rodents. *Braz Arch Biol Technol*, 60(December), 1–10.

- Kuntaman, K., Santoso, S., Wahjono, H., dan Mertaniasih, N. M. (2011). The sensitivity pattern of extended spectrum beta lactamase-producing bacteria against six antibiotics that routinely used in clinical setting. *J Indon A 45 Assoc*, 61(12), 482–486.
- Luo, C., Walk, S., Gordon, D. M., Feldgarden, M., dan Tiedje, J. M. (2011). Genome sequencing of environmental *Escherichia coli* expands understanding of the ecology and speciation of the model bacterial species. *Proceeding of the National Academy of Sciences of the United State of America*, 108(17). <https://doi.org/10.1073/pnas.1015622108>
- Macgregor, R., dan Tunnessen, W. (2015). The incidence of patogenic organisms in the normal flora of the neonate ' s external ear and nasopharynx. *Clin Pediatr*, 12(12), 697–700.
- Mackie, R., Sghir, A., dan Gaskins, H. R. (1999). Developmental microbial ecology of the neonatal gastrointestinal. *Am J Clin Nutr*, 69, 1035S – 45S.
- Madico, G., Akopyants, N. S., dan Berg, D. E. (1995). Arbitrarily primed PCR DNA fingerprinting of *Escherichia coli* O157 : H7 strains by using templates from boiled cultures. *Clin Microbiol Infect*, 33(6), 1534–1536.
- Mammaia, C., Carlo, P. Di, Cipolla, D., Giuffre, M., Casuccio, A., Gaetano, V. Di, ... Giovanni Corsello. (2004). Surveillance of multidrug-resistant gram-negative bacilli in a neonatal intensive care unit : prominent role of cross transmission. *Am J Infect Control*, 35(4), 8–10. <https://doi.org/10.1016/j.ajic.2006.04.210>
- Marialouis, X. A., dan Santhanam, A. (2016). Antibiotic Resistance , RAPD- PCR typing of multiple drug resistant strains of *Escherichia coli* from urinary tract infection (UTI). *J Clin Diagn Res*, 10(3), 5–9. <https://doi.org/10.7860/JCDR/2016/16470.7389>
- Martinez, J. L., dan Rojo, F. (2011). Metabolic regulation of antibiotic resistance. *FEMS Microbiol Rev*, 35, 768–789. <https://doi.org/10.1111/j.1574-6976.2011.00282.x>
- Maurer, J. J., Lee, M. D., Lobsinger, C., Brown, T., Maier, M., Thayer, S. G., ... Thayer, S. G. (1998). American association of avian pathologists molecular typing of avian *Escherichia coli* isolates by random amplification of polymorphic DNA (Vol. 42).
- McMurry, L. M., dan Stuart B, L. (2011). The periplasmic protein mppA is not involved in regulation of marA in *Escherichia coli*. *Antimicrob Agents Chemother.*, 55(10), 6764. <https://doi.org/10.1128/AAC.05030-11>

- Moissenet, D., Clermont, O., Bingen, E., Arlet, G., Denamur, E., Me, A., ... Antoine, S. (2010). Meningitis caused by *Escherichia coli* producing TEM-52 extended-spectrum β -lactamase within an extensive outbreak in a neonatal ward: epidemiological investigation and characterization of the strain. *J Clin Microbiol*, 48(7), 2459–2463. <https://doi.org/10.1128/JCM.00529-10>
- Mruk, I., dan Kobayashi, I. (2014). To be or not to be : regulation of restriction – modification systems and other toxin – antitoxin systems. *Nucleic Acids Res*, 42(1), 70–86. <https://doi.org/10.1093/nar/gkt711>
- Mugnaioli, C., Luzzaro, F., De Luca, F., Brigante, G., Perilli, M., Amicosante, G., ... Rossolini, G. M. (2006). CTX-M-type extended-spectrum β -lactamases in Italy: Molecular epidemiology of an emerging countrywide problem. *Antimicrob Agents Chemother.*, 50(8), 2700–2706. <https://doi.org/10.1128/AAC.00068-06>
- Muinck, E. J. De, Øien, T., Storrø, O., Johnsen, R., Stenseth, N. C., Rønningen, K. S., dan Rudi, K. (2011). *Diversity , transmission and persistence of Escherichia coli in a cohort of mothers and their infants.* 3, 352–359. <https://doi.org/10.1111/j.1758-2229.2010.00231.x>
- Nicolas-Chanoine, M.-H., Blanco, J., Leflon-Guibout, V., Demarty, R., Alonso, M. P., Canic, M. M., ... Johnson, J. R. (2008). Intercontinental emergence of *Escherichia coli* clone O25 : H4-ST131 producing CTX-M-15. *J Antimicrob Chemother*, 61, 273–281. <https://doi.org/10.1093/jac/dkm464>
- Nielsen, K. L., Godfrey, P. A., Stegger, M., Andersen, P. S., Feldgarden, M., dan Fridomdt-Møller, N. (2015). Selection of unique *Escherichia coli* clones by Random Amplified Polymorphic DNA (RAPD): Evaluation by whole genome sequencing. *Microbiol Methode NIH Public Access*, 5, 101–103. <https://doi.org/10.1016/j.mimet.2014.05.018.Selection>
- Ntirenganya, C., Manzi, O., Muvunyi, C. M., dan Ogbuagu, O. (2015). High prevalence of antimicrobial resistance among common bacterial isolates in a tertiary healthcare facility in Rwanda. *J Trop Med Hyg*, 92(4), 865–870. <https://doi.org/10.4269/ajtmh.14-0607>
- Osawa, S. (1987). The guanine and cytosine content of genomic DNA and bacterial evolution. *Proceeding of the National Academy of Sciences of the United State of America*, 84(January), 166–169.
- Oteo, J., Cercenado, E., Fernández-romero, S., Saéz, D., Padilla, B., Zamora, E., dan Cuevas, O. (2012). Extended-spectrum-B- lactamase-producing *Escherichia coli* as a cause of pediatric infections : Report of a neonatal intensive care unit outbreak due to a CTX-M-14-producing strain.

Antimicrob Agents Chemother., 56(1), 54–58.
<https://doi.org/10.1128/AAC.05103-11>

Ouyang-latimer, J., Ajami, N. J., Jiang, Z., Okhuysen, P. C., Paredes, M., Flores, J., dan Dupont, H. L. (2010). Biochemical and genetic diversity of enterotoxigenic *Escherichia coli* associated with diarrhea in United States students in Cuernavaca and Guadalajara. *J Infect Dis*, 201(12), 1831–1838.
<https://doi.org/10.1086/652797>

Palma, N., Gomes, C., Riveros, M., García, W., Martínez-puchol, S., Ruiz-roldán, L., ... Ruiz, J. (2016). Virulence factors pro fi les and ESBL production in *Escherichia coli* causing bacteremia in Peruvian children. *Diagnostic Microbiology and Infectious Disease*, 86(1), 70–75.
<https://doi.org/10.1016/j.diagmicrobio.2016.05.017>

Picard, B., Garcia, S., Gouriou, P., Duriez, P., Brahim, N., Bingen, E., ... Debre, R. (1999). The Link between Phylogeny and Virulence in *Escherichia coli* Extraintestinal Infection †. *American Society for Microbiology*, 67(2), 546–553.

Pitout, J., Nordmann, P., Laupland, K. B., dan Poirel, L. (2005). Emergence of Enterobacteriaceae producing extended-spectrum β -lactamases (ESBLs) in the community. *J Antimicrob Chemother*, 56(May), 52–59.
<https://doi.org/10.1093/jac/dki166>

Rath, S., Dubey, D., Suhu, M. C., Debata, N. K., dan Pandhy, R. N. (2014). Surveillance of ESBL producing multidrug resistant *Escherichia coli* in a teaching hospital in India. *Asian Pac J Trop Dis*, 4(2), 140–149.
[https://doi.org/10.1016/S2222-1808\(14\)60331-5](https://doi.org/10.1016/S2222-1808(14)60331-5)

Rawat, D., dan Nair, D. (2010). Extended-spectrum β -lactamases in gram negative bacteria. *J Glob Infect Dis*, 2(3), 263–274. <https://doi.org/10.4103/0974-777X.68531>

Saiman, L., Cronquist, A., Wu, F., Rubenstein, D., Eisner, W., Kreiswirth, B. N., dan Latta, P. Della. (2014). An outbreak of methicillin-resistant *Staphylococcus aureus* in a neonatal intensive care unit. *Infect Control Hosp Epidemiol*, 24(5), 317–321.

Salamah, U., Kuntaman, K., Wasito, E. B., & Setyarini, W. (2019). *Perbandingan kelompok filogenetik bakteri Escherichia coli extended spectrum β -lactamase pada isolat komunitas dan patogen di RSUD Dr. Soetomo Surabaya*. Airlangga University.

- Salem, H. H., Huang, T.-H., Ali, B. A., dan Xie, Q. (2006). Differentiation of *Bacillus thuringiensis* and *Escherichia coli* by the Randomly Amplified Polymorphic DNA analysis. *J Appl Sci*, 6(7), 1540–1546.
- Schults, C., dan Suzanne, G. (2012). Plasmid-mediated resistance in Enterobacteriaceae changing landscape and implications for therapy. *Drugs*, 72(1), 1–16.
- Shakil, S., Akram, M., dan Khan, A. U. (2008). Tigecycline : A critical update. *J Chemother*, 20(4), 411–419.
- Shakil, S., Ali, A. S. Z., Akram, A. M., Ali, S. M., dan Khan, A. U. (2010). Risk factors for extended-spectrum β -lactamase producing *Escherichia coli* and *Klebsiella pneumoniae* acquisition in a neonatal intensive care unit. *J Trop Pediatr*, 56(2), 90–96. <https://doi.org/10.1093/tropej/fmp060>
- Shakil, S., Khan, R., Zarrilli, R., dan Khan, A. U. (2008). Aminoglycosides versus bacteria – a description of the action , resistance mechanism , and nosocomial battleground. *J Biomed Sci*, 15, 5–14. <https://doi.org/10.1007/s11373-007-9194-y>
- Song, K., Jeon, J. H., Park, W. B., Park, S., Kim, B., Oh, M., ... Kim, N. J. (2009). Clinical outcomes of spontaneous bacterial peritonitis due to extended-spectrum beta-lactamase-producing *Escherichia coli* and *Klebsiella* species: A retrospective matched case-control study. *BMC Infect Dis*, 6(9), 1–6. <https://doi.org/10.1186/1471-2334-9-41>
- Stoll, B. J., Hansen, N., Fanaroff, A. A., Wright, L. L., Carlo, W. A., Ehrenkranz, R. A., ... Poole, W. K. (2002). Late-onset sepsis in very low birth weight neonates: the experience of the NICHD neonatal research network. *Pediatrics*, 110(2), 285–291.
- Storberg, V. (2014). ESBL-producing Enterobacteriaceae in Africa a non-systematic literature review of research published 2008-2012. *Infect Ecol Epidemiol*, 1(4), 1–16.
- Tannock, G. W. (1995). Microecology of the gastrointestinal tract in relation to lactic acid bacteria. *Int Dairy J*, 5(95), 1059–1070.
- Tannock, G. W. (1999). The normal microflora : an introduction. In *Medical Importance of the Normal Microflora* (pp. 1–2). Kluwer Academic Publishers.
- Vila, J., Simon, K., Ruiz, J., Horcajada, J. P., Velasco, M., Barranco, M., dan Moreno, A. (1999). Are quinolone-resistant uropatogenic *Escherichia coli* less virulent ? *J Infect Dis*, 186, 1039–1042.

- Vogel, L., Oorschot, E. Van, Maas, H. M. E., Minderhoud, B., Dijkshoorn, L., dan Infect, C. M. (2000). Epidemiologic typing of *Escherichia coli* using RAPD analysis , ribotyping and serotyping. *Clin Microbiol Infect*, 6(2), 82–87
- Wang, G., Whittam, T. S., Berg, C. M., dan Berg, D. E. (1993). RAPD (arbitrary primer) PCR is more sensitive than multilocus enzyme electrophoresis for distinguishing related bacterial strains. *Nucleic Acids Res*, 21(25), 5930–5933.
- Weist, K., Wendt, C., Petersen, L. R., Versmold, H., dan Rüden, H. (2014). An outbreak of pyoderma among neonates caused by ultrasound gel contaminated with methicillin-susceptible *Staphylococcus aureus*. *Infect Control Hosp Epidemiol*, 21(12), 760–764.