

## Daftar Pustaka

- [1] Setyorini, Kuspriyanto, D. H. Widyanoro, and A. Pancoro, “The implementation of bit-parallelism for DNA sequence alignment,” *J. Phys. Conf. Ser.*, vol. 835, no. 1, 2017, doi: 10.1088/1742-6596/835/1/012004.
- [2] J. Gregor and M. G. Thomason, “Dynamic Programming Alignment of Sequences Representing Cyclic Patterns,” *IEEE Trans. Pattern Anal. Mach. Intell.*, vol. 15, no. 2, pp. 129–135, 1993, doi: 10.1109/34.192484.
- [3] G. Myers, “A fast bit-vector algorithm for approximate string matching based on dynamic programming,” *J. ACM*, vol. 46, no. 3, pp. 395–415, 1999, doi: 10.1145/316542.316550.
- [4] J. Xiong, *Essential bioinformatics*, vol. 148. 2006.
- [5] D. Huson, “3 Bit-Vector-Based Approximate String Matching Ukkonens algorithm,” pp. 12–19, 2004.
- [6] G. Attardi, T. Flagella, and P. Iglieo, “A customisable memory management framework for C++,” *Softw. - Pract. Exp.*, vol. 28, no. 11, pp. 1143–1184, 1998, doi: 10.1002/(sici)1097-024x(199809)28:11<1143::aid-spe194>3.0.co;2-7.