

**PENAMBATAN MOLEKULER FULVESTRANT PADA RESEPTOR
ESTROGEN DAN VARIAN POLIMORFISMENYA SECARA *IN SILICO***

SKRIPSI

diajukan sebagai salah satu syarat untuk memperoleh gelar Sarjana Sains
Program Studi Biologi



oleh
Dwi Fitriawati Fajrin
1900480

**PROGRAM STUDI BIOLOGI
FAKULTAS PENDIDIKAN MATEMATIKA DAN ILMU PENGETAHUAN ALAM
UNIVERSITAS PENDIDIKAN INDONESIA
BANDUNG
2023**

**PENAMBATAN MOLEKULER FULVESTRANT PADA RESEPTOR
ESTROGEN DAN VARIAN POLIMORFISMENYA SECARA *IN SILICO***

Oleh

Dwi Fitriawati Fajrin

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

© Dwi Fitriawati Fajrin 2023

Universitas Pendidikan Indonesia 2023

November 2023

Hak cipta dilindungi undang-undang

Skripsi ini tidak boleh diperbanyak seluruhnya atau sebagian dengan dicetak
ulang, difotokopi, atau cara lainnya tanpa izin dari penulis

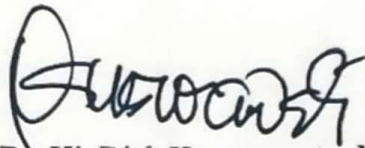
LEMBAR PENGESAHAN

(Dwi Fitriawati Fajrin)

**PENAMBATAN MOLEKULER FULVESTRANT PADA RESEPTOR
ESTROGEN DAN VARIAN POLIMORFISMENYA SECARA *IN SILICO***

Disetujui dan disahkan oleh:

Pembimbing I



Dr. Hj. Diah Kusumawaty, M.Si.

NIP: 197008112001122001

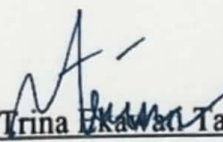
Pembimbing II



Dr. Any Aryani, M.Si.

NIP. 197105302001122001

Pembimbing III

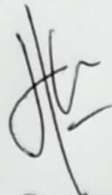


Prof. Dr. Trina Ekawati Tallei, M.Si.

NIP. 196605081995122001

Mengetahui,

Ketua Program Studi Biologi



Dr. H. Wahyu Surakusumah, M.T

NIP 197212301999031001

LEMBAR PERNYATAAN

Dengan ini saya menyatakan bahwa skripsi dengan judul “Penambatan Molekuler Fulvestrant pada Reseptor Estrogen dan Varian Polimorfismenya secara *In Silico*” 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 siap menanggung sanksi apabila di kemudian hari ditemukan adanya pelanggaran etika keilmuan atau ada klaim dari pihak lain terhadap keaslian karya saya ini.

Bandung, November 2023

Yang membuat pernyataan

Dwi Fitriawati Fajrin

NIM. 1900480

ABSTRAK

Penambatan Molekuler Fulvestrant pada Reseptor Estrogen dan Varian Polimorfismenya secara *In Silico*

Salah satu penyakit paling banyak menyebabkan kematian didunia adalah penyakit kanker. Kanker payudara adalah kasus paling banyak kanker yang diderita pada perempuan. ER atau *Estrogen Receptor* adalah reseptor yang berperan dalam perkembangan sel kanker payudara. Terdapat lebih dari 70% penderita kanker payudara memiliki status hormonal dengan kategori ER positif. ER alpha diketahui adalah target terapi dalam pengobatan kanker payudara ER positif. Diketahui pada ER alpha terdapat beberapa polimorfisme. Pada kasus polimorfisme ini diketahui dapat mempengaruhi fungsi protein salah satunya adalah resistensi terhadap terapi obat. Penelitian ini bertujuan untuk mengetahui interaksi Fulvestrant sebagai salah satu obat standar yang digunakan dalam pengobatan kanker payudara terhadap polimorfisme ER D538G, Y537S, Y537N, Y537C, dan L536R yang dilakukan secara *in silico*. Berdasarkan penelitian yang telah dilakukan menunjukkan bahwa Fulvestrant masih dapat direkomendasikan pada penderita kanker payudara dengan polimorfisme D538G, Y537S, Y537N, Y537C, dan L536R. Hasil temuan yang didapatkan diketahui polimorfisme pada asam amino 536, 537, dan 538 tidak memiliki peran yang signifikan dalam pengikatannya pada ligan. Hasil dari penelitian yang telah dilakukan ini dapat menjadi referensi pada penelitian yang akan datang terkait peran polimorfisme pada pengikatannya dengan ligan terutama pada reseptor ER.

Kata Kunci : Kanker Payudara, ER, Fulvestrant, Polimorfisme, Penambatan Molekuler

ABSTRACT

In Silico Approach for Molecular Docking of Fulvestrant with Estrogen Receptor and its Polymorphic Variants

One of the diseases that causes the most death in the world is cancer. The most frequent type of cancer among woman is breast cancer. ER or Estrogen Receptor is a receptor that has a role in the growth of breast cancer cells. More than 70% of breast cancer patient have the ER positive category hormonal status in. ER alpha is known to be a therapeutic target of the treatment in ER positive breast cancer. It is known that ER alpha contains several polymorphisms. In the case of this polymorphism, it is known that it can affect protein function, one of them is resistance to drug therapy. The aim of this research is to know the interaction of Fulvestrant as one the drugs that used in the breast cancer treatment with the ER polymorphisms D538G, Y537S, Y537N, Y537C, and L536R, by in silico method. Based on the research, it shows that Fulvestrant can still be recommended for breast cancer patient with the D538G, Y537S, Y537N, Y537C, and L536R polymorphisms. The findings obtained show that polymorphisms in amino acids 536, 537, and 538 do not have a significant role in binding to the ligand. The findings from this research can be used as a reference in future research regarding the role of polymorphism in binding to ligands, especially ER receptors.

Keywords: Breast Cancer, ER, Fulvestrant, Polymorphism, Molecular Docking

DAFTAR ISI

KATA PENGANTAR	v
ABSTRAK	vii
ABSTRACT	viii
DAFTAR ISI	ix
DAFTAR TABEL	xi
DAFTAR GAMBAR	xii
DAFTAR LAMPIRAN	xiv
BAB I PENDAHULUAN	1
1.1 Latar Belakang	1
1.2 Rumusan Masalah	4
1.3 Tujuan Penelitian	4
1.4 Manfaat Penelitian	4
1.5 Struktur Organisasi Skripsi	4
BAB II TINJAUAN PUSTAKA	6
2.1 Kanker	6
2.2 Kanker Payudara	8
2.3 <i>Estrogen Receptor</i> dan Varian Polimorfisme	11
2.4 Fulvestrant	16
2.5 Obat Presisi	18
2.6 Penambatan Molekuler	21
BAB III METODE PENELITIAN	25
3.1 Jenis Penelitian	25
3.2 Lokasi dan Waktu Penelitian	25
3.3 Alat dan Bahan	25
3.4 Prosedur Penelitian	27
3.4.1 Preparasi Sekuen <i>Estrogen Receptor</i>	27
3.4.2 <i>Multiple Sequence Alignment</i>	28
3.4.3 Pemodelan Protein	30
3.4.4 Validasi Pemodelan Protein	31
3.4.5 Visualisasi Variasi Polimorfisme <i>Estrogen Receptor</i>	32

3.4.6	<i>Superimpose Estrogen Receptor</i>	32
3.4.7	Preparasi Reseptor.....	33
3.4.8	Preparasi Ligan.....	34
3.4.9	Penambatan Molekuler.....	34
3.4.10	Visualisasi Hasil Penambatan Molekuler.....	35
3.4.11	Alur Kerja Penelitian.....	36
BAB IV HASIL DAN PEMBAHASAN		37
4.1	Struktur Reseptor dan Ligan.....	37
4.2	Penyiapan Struktur Reseptor	37
4.2.1	Sekuens Asam Amino <i>Estrogen Receptor</i>	37
4.2.2	<i>Multiple Sequence Alignment (MSA)</i>	39
4.2.3	Pemodelan Protein Reseptor	42
4.2.4	Validasi Pemodelan Protein Reseptor	44
4.2.5	Visualisasi Variasi Polimorfisme <i>Estrogen Receptor</i>	47
4.2.6	<i>Superimpose Estrogen Receptor</i>	49
4.2.7	Preparasi Reseptor.....	50
4.3	Preparasi Ligan Fulvestrant.....	50
4.3.1	Preparasi Ligan.....	51
4.4	Penambatan Molekuler dan Visualisasi.....	52
BAB V SIMPULAN, IMPLIKASI, DAN REKOMENDASI		58
5.1	Simpulan.....	58
5.2	Implikasi.....	58
5.3	Rekomendasi	58
DAFTAR PUSTAKA		59
LAMPIRAN.....		73

DAFTAR TABEL

Tabel 2.1 Subtipe Kanker Payudara.....	10
Tabel 2.2 Asam Amino	15
Tabel 3.1 Perangkat yang Digunakan	25
Tabel 3.2 Bahan yang Digunakan	26
Tabel 3.3 Laman Akses yang Digunakan	26
Tabel 3.4 Variasi Polimorfisme <i>Estrogen Receptor</i>	29
Tabel 4.1 Statistik Ramachandran Plot	45
Tabel 4.2 Struktur 3D Ligan	50
Tabel 4.3 Hasil Penambatan Molekuler Nilai Energi Ikatan Bebas Menggunakan Gnina	53

DAFTAR GAMBAR

Gambar 2.1 Sel kanker.....	7
Gambar 2.2 Struktur ER alpha dan ER beta	12
Gambar 2.3 Ilustrasi ER alpha dan ER beta (A/B)	12
Gambar 2.4 Mekanisme <i>Estrogen Receptor</i>	13
Gambar 2.5 Fulvestrant.....	16
Gambar 2.6 Ilustrasi Mekanisme Kerja Estradiol, Tamoxifen dan Fulvestrant....	17
Gambar 2.7 Elemen dalam <i>Molekuler Docking</i>	21
Gambar 2.8 Kuadran Plot Ramachandran.....	23
Gambar 3.1 Alur Kerja Penelitian.....	36
Gambar 4.1 Sekuens Asam Amino Lengkap <i>Estrogen Receptor Alpha</i>	38
Gambar 4.2 Sekuens Asam Amino <i>Estrogen Receptor Alpha Template</i>	39
Gambar 4.3 Penjajaran Sekuens Asam Amino <i>Estrogen Receptor Alpha</i>	40
Gambar 4.4 <i>Multiple Sequence Alignment</i> ER <i>Wild Type</i> , ER D538G, ER Y537S, ER Y537N, ER Y537C, ER L536 R.....	41
Gambar 4.5 Hasil Pemodelan Protein ER <i>Wild Type</i> (A), Polimorfisme D538G (B), Polimorfisme Y537S, (C) Polimorfisme Y537N, (D) Polimorfisme Y537N, (E) Polimorfisme Y537C, Dan (F) Polimorfisme L536R.....	43
Gambar 4.6 Plot Ramachandran Protein ER <i>Wild Type</i> (A), Polimorfisme D538G (B), Polimorfisme Y537S, (C) Polimorfisme Y537N, (D) Polimorfisme Y537N, (E) Polimorfisme Y537C, dan (F) Polimorfisme L536R.....	46
Gambar 4.7 ER <i>Wild Type</i> Asam Amino Posisi 536, 537 dan 538 (hijau) (A), Polimorfisme Varian ER L536R (abu-abu) (B), Polimorfisme Y537S (ungu) (C), Polimorfisme Y537N (kuning) (D), Polimorfisme Y537C (merah muda) (E), dan Polimorfisme D538G (biru) (F).....	48
Gambar 4.8 <i>Superimpose</i> Protein ER <i>Wild Type</i> (hijau) dengan Varian D538G (biru muda) (A), Polimorfisme Y537S (ungu) (B), Polimorfisme Y537N (kuning) (C), Polimorfisme Y537C (merah muda), dan (D)Polimorfisme L536R (abu-abu) (E).....	49

Gambar 4.9 Perbandingan Interaksi Fulvestrant dengan *Estrogen Receptor* WT
(A), Polimorfisme D538G(B), Y537S(C), Y537N(D), Y537C(E),
dan L536R (F)..... 55

DAFTAR LAMPIRAN

Lampiran 1. Preparasi Sekuen Reseptor <i>Estrogen Receptor</i>	73
Lampiran 2. <i>Multiple Sequence Alignment</i>	75
Lampiran 3. Persiapan Struktur 3D <i>Estrogen Receptor</i>	75
Lampiran 4. Validasi Struktur 3D <i>Estrogen Receptor</i>	76
Lampiran 5. Visualisasi Polimorfisme Menggunakan Perangkat Lunak Pymol ..	77
Lampiran 6. <i>Superimpose</i> Struktur 3D <i>Estrogen Receptor</i>	78
Lampiran 7. Preparasi Struktur 3D	78
Lampiran 8. Pengunduhan Struktur Ligan	79
Lampiran 9. Penambatan Molekuler Menggunakan Perangkat Lunak Gnina	80
Lampiran 10. Visualisai Penambatan Molekuler Menggunakan Perangkat Lunak Discovery Studio	80
Lampiran 11. Hasil Interaksi Ligan dan Reseptor	81

DAFTAR PUSTAKA

- Agus, N. A., Holik, H. A., & Achmad, A. (2022). Kemajuan Terkini dalam Radiofarmaka sebagai Agen Terapi Kanker Payudara. *Farmaka*, 20(2), 121–135. <http://jurnal.unpad.ac.id/farmaka/article/view/39356>
- American cancer society. (2022). *What is cancer?*. Diakses dari <https://www.cancer.org/cancer/understanding-cancer/what-is-cancer.html>
- Al-Mubarak, M., Sacher, A. G., Ocana, A., Vera-Badillo, F., Seruga, B., & Amir, E. (2013). Fulvestrant for advanced breast cancer: A meta-analysis. In *Cancer Treatment Reviews*. <https://doi.org/10.1016/j.ctrv.2013.03.004>
- Aliwikarta, K., Palupi, N. S., & Giriwono, P. E. (2016). Prevalensi Penyakit Kanker di Indonesia Berdasarkan Pola Konsumsi Pangan dan Gaya Hidup Cancer prevalence in Indonesia Based on Food Consumption Patterns and Lifestyle Newly Weds Foods Asia Pacific 2. *Jurnal Mutu Pangan*, 3(1), 71–78.
- Amelia, F. (2014). Studi Interaksi Ligan Peptidoid dan Peptida Dengan Enzim Protease NS3/NS2B Virus Dengue. *Jurnal Sainstek*, 6(1), 24–29.
- Ayaz, G., Yasar, P., Olgun, C. E., Karakaya, B., Kars, G., Razizadeh, N., Yavuz, K., Turan, G., & Muyan, M. (2019). Dynamic transcriptional events mediated by estrogen receptor alpha. *Frontiers in Bioscience - Landmark*, 24(2), 245–276. <https://doi.org/10.2741/4716>
- Baroroh, U., Muscifa, Z. S., Destiarani, W., Rohmatullah, F. G., & Yusuf, M. (2023). Molecular interaction analysis and visualization of protein-ligand docking using Biovia Discovery Studio Visualizer. *Indonesian Journal of Computational Biology (IJCB)*, 2(1), 22. <https://doi.org/10.24198/ijcb.v2i1.46322>
- Berman, H. M., Westbrook, J., Feng, Z., Gilliland, G., Bhat, T. N., Weissig, H., Shindyalov, I. N., & Bourne, P. E. (2000). The Protein Data Bank. *Nucleic Acids Research*, 28(1), 235–242. <https://doi.org/10.1038/s41577-020-00473-z>
- Boér, K. (2017). Fulvestrant in advanced breast cancer: evidence to date and place in therapy. *Therapeutic Advances in Vaccines*, 9(7), 465–479. <https://doi.org/10.1177/https>
- Bosco, K. H., & Brasseur, R. (2005). The Ramachandran plots of glycine and proline. *BMC Structural Biology*, 5(14), 1–11. <https://doi.org/10.1186/1472-6807-5-14>

- Breast Cancer Now.(2018). *Fulvestrant (Faslodex)*. Diakses dari <https://breastcancer.org/information-support/facing-breast-cancer/going-through-treatment-breast-cancer/hormone-therapy/fulvestrant-faslodex>
- Bross, P. F., Baird, A., Chen, G., Jee, J. M., Lostritto, R. T., Morse, D. E., Rosario, L. A., Williams, G. M., Yang, P., Rahman, A., Williams, G., & Pazdur, R. (2003). Fulvestrant in postmenopausal women with advanced breast cancer. *Clinical Cancer Research*, *9*(12), 4309–4317.
- Carugo, O., & Pongor, S. (2008). A normalized root-mean-square distance for comparing protein three-dimensional structures. *Protein Science*, *10*(7), 1470–1473. <https://doi.org/10.1110/ps.690101>
- Chen, P., Li, B., & Ou-Yang, L. (2022). Role of estrogen receptors in health and disease. *Frontiers in Endocrinology*, *13*, 1–25. <https://doi.org/10.3389/fendo.2022.839005>
- Chu, W. T., & Zheng, Q. C. (2013). Conformational changes of enzymes and DNA in molecular dynamics: Influenced by pH, temperature, and ligand. In *Advances in Protein Chemistry and Structural Biology* (1st ed., Vol. 92). Copyright © 2013 Elsevier Inc. All rights reserved. <https://doi.org/10.1016/B978-0-12-411636-8.00005-5>
- Coimbra, J. T. S., Feghali, R., Ribeiro, R. P., Ramos, M. J., & Fernandes, P. A. (2020). The importance of intramolecular hydrogen bonds on the translocation of the small drug piracetam through a lipid bilayer. *RSC Advances*, *11*(2), 899–908. <https://doi.org/10.1039/d0ra09995c>
- Cowley, S. M., Hoare, S., Mosselman, S., & Parker, M. G. (1997). Estrogen receptors α and β form heterodimers on DNA. *Journal of Biological Chemistry*, *272*(32), 19858–19862. <https://doi.org/10.1074/jbc.272.32.19858>
- Creighton, T. E. (1990). Protein folding. *Biochem*, *270*, 1–16. https://doi.org/10.1007/978-981-10-4968-2_2
- Csizmadia, P. (1999). MarvinSketch and MarvinView: Molecule Applets for the World Wide Web. *Third International Electronic Conference on Synthetic Organic Chemistry*. <https://doi.org/10.3390/ecsoc-3-01775>
- Dang, D. T. (2022). Molecular Approaches to Protein Dimerization: Opportunities for Supramolecular Chemistry. *Frontiers in Chemistry*, *10*, 1–12. <https://doi.org/10.3389/fchem.2022.829312>
- Daraz, F., Windarti, I., & Putu, R. A. S. (2023). Peran Obesitas dalam Metastasis Kanker Payudara. *Medula*, *13*(1), 172–178.

- EBE. (2015). *White Paper on Personalized Medicine*. Diakses dari: http://www.ebe-biopharma.eu/uploads/EBE-biopharma_WhitePaperOnPM_RGB.pdf
- Ekins, S., Mestres, J., & Testa, B. (2007). In silico pharmacology for drug discovery: Applications to targets and beyond. *British Journal of Pharmacology*, 152(1), 21–37. <https://doi.org/10.1038/sj.bjp.0707306>
- Elfita, L., Apriadi, A., Supandi, S., & Dianmurdedi, S. (2022). Studi Penambatan Molekuler dan Simulasi Dinamika Molekuler Senyawa Turunan Furanokumarin terhadap Reseptor Estrogen Alfa (ER- α) Sebagai Anti Kanker Payudara. *Jurnal Sains Farmasi & Klinis*, 9(3), 255. <https://doi.org/10.25077/jsfk.9.3.255-264.2022>
- Enmark, E., Peltö-Huikko, M., Grandien, K., Lagercrantz, S., Lagercrantz, J., Fried, G., Nordenskjöld, M., & Gustafsson, J. Å. (1997). Human estrogen receptor β -gene structure, chromosomal localization, and expression pattern. *Journal of Clinical Endocrinology and Metabolism*, 82(12), 4258–4265. <https://doi.org/10.1210/jcem.82.12.4470>
- Essoussi, N., Boujenfa, K., & Limam, M. (2008). A comparison of MSA tools. *Bioinformatics*, 2(9), 452–455. <https://doi.org/10.6026/97320630002452>
- Faisal, H. K. P., Zaini, J., & Yunus, F. (2020). Next-Generation Sequencing pada Kanker Paru Next-Generation Sequencing in Lung Cancer. *E Jurnal Kedokteran Indonesia*, 8(2). <https://doi.org/10.23886/ejki.8.11579>
- Ferguson, J. R., Harding, J. R., Killick, D. A., Lumbard, K. W., Scheinmann, F., & Stachulski, A. V. (2001). Putative metabolites of fulvestrant, an estrogen receptor downregulator. Improved glucuronidation using trichloroacetimidates. *Journal of the Chemical Society. Perkin Transactions 1*, 22, 3037–3041. <https://doi.org/10.1039/b105397n>
- Fidianingsih, I. (2013). 6721-11667-1-SM.pdf. In *Personalized Medicine Untuk Penderita Kanker* (Vol. 5, Issue 2).
- Fitri, D. E. W. (2023). *Penambatan Molekuler Pioglitazone Terhadap Varian Polimorfisme Reseptor PPAR γ Pada Diabetes Melitus Tipe 2*. (Skripsi). Fakultas Matematika dan Ilmu Pengetahuan Alam, Universitas Sam Ratulangi, Manado.
- Fuentes, N., & Silveyra, P. (2019). Estrogen receptor signaling mechanism. *Adv Protein Chem Struct Biol*, 116, 135–170. <https://doi.org/10.1016/bs.apcsb.2019.01.001.Estrogen>

- Gita, S. (2015). Peran Bioinformatika dalam Desain Kandidat Molekul Obat. *Biotrends*, *1*(1), 25–27. http://terbitan.biotek.lipi.go.id/index.php/biotrends/article/view/10%0Ahttps://www.researchgate.net/publication/317011679_Peran_Bioinformatika_dalam_Desain_Kandidat_Molekul_Obat
- GLOBOCAN/GCO(Global Cancer Observatory).(2021). *Cancer Today*. Diakses dari <https://gco.iarc.fr/>
- Green, S., Walter, P., Kumar, V., Krust, A., Bornert, J. M., Argos, P., & Chambon, P. (1986). Human oestrogen receptor cDNA: Sequence, expression and homology to v-erb-A. *Nature*, *320*(6058), 134–139. <https://doi.org/10.1038/320134a0>
- Gutnik, D., Evseev, P., Mirosnikov, K., & Shneider, M. (2023). Using AlphaFold Predictions in Viral Research. *Current Issues in Molecular Biology*. *45*(4), 3705-3732. <https://doi.org/10.3390/cimb45040240>
- Hakim, A., Widyanti, W., & Alfianto, U. (2018). HUBUNGAN ANTARA OBESITAS DENGAN RESEPTOR HORMONAL (Reseptor Estrogen dan Progesteron) DAN EKSPRESI HER-2/NEU PADA PASIEN KANKER PAYUDARA DI RS X SURAKARTA. *Biomedika*, *10*(1), 30–34. <https://doi.org/10.23917/biomedika.v10i1.5851>
- Haldosén, L. A., Zhao, C., & Dahlman-Wright, K. (2014). Estrogen receptor beta in breast cancer. *Molecular and Cellular Endocrinology*, *382*(1), 665–672. <https://doi.org/10.1016/j.mce.2013.08.005>
- Harbeck, N., Penault-Llorca, F., Cortes, J., Gnant, M., Houssami, N., Poortmans, P., Ruddy, K., Tsang, J., & Cardoso, F. (2019). Breast cancer. In *Nature Reviews Disease Primers* (Vol. 5, Issue 1). <https://doi.org/10.1038/s41572-019-0111-2>
- Harrod, A., Lai, C. F., Goldsbrough, I., Simmons, G. M., Oppermans, N., Santos, D. B., Györffy, B., Allsopp, R. C., Toghiani, B. J., Balachandran, K., Lawson, M., Morrow, C. J., Surakala, M., Carnevalli, L. S., Zhang, P., Guttery, D. S., Shaw, J. A., Coombes, R. C., Buluwela, L., & Ali, S. (2022). Genome engineering for estrogen receptor mutations reveals differential responses to anti-estrogens and new prognostic gene signatures for breast cancer. *Oncogene*, *41*(44), 4905–4915. <https://doi.org/10.1038/s41388-022-02483-8>
- Hendlich, M., Rippmann, F., & Barnickel, G. (1997). LIGSITE: Automatic and efficient detection of potential small molecule-binding sites in proteins. *Journal of Molecular Graphics and Modelling*, *15*(6), 359–363. [https://doi.org/10.1016/S1093-3263\(98\)00002-3](https://doi.org/10.1016/S1093-3263(98)00002-3)

- Hernando, C., Ortega-Morillo, B., Tapia, M., Moragón, S., Martínez, M. T., Eroles, P., Garrido-Cano, I., Adam-Artigues, A., Lluch, A., Bermejo, B., & Cejalvo, J. M. (2021). Oral selective estrogen receptor degraders (Serds) as a novel breast cancer therapy: Present and future from a clinical perspective. *International Journal of Molecular Sciences*, 22(15). <https://doi.org/10.3390/ijms22157812>
- Hernandez-Santoyo, A., Yair, A., Altuzar, V., Vivanco-Cid, H., & Mendoza-Barrer, C. (2013). Protein-Protein and Protein-Ligand Docking. *Protein Engineering - Technology and Application*, Figure 1. <https://doi.org/10.5772/56376>
- Hillisch, A., & Hilgenfeld, R. (2003). The role of protein 3D-structures in the drug discovery process. In *Modern Methods of Drug Discovery*. https://doi.org/10.1007/978-3-0348-7997-2_8
- Huang, T.S., & Ang, P.(2018).[Online]. TERAPI HORMONAL DALAM KANKER PAYUDARA LANJUT DAN CARA BARU UNTUK MENGATASI RESISTENSI. Diakses dari <https://oncocare.sg/id/blogs/terapi-hormonal-dalam-kanker-payudara-lanjut-dan-agen-baru-untuk-mengatasi-resisten/>
- Ibrahim, F., Huang, B., Xing, J. Z., Roa, W., & Gabos, S. (2009). Effects of Estrogen Contamination on Human Cells: Modeling and Prediction Based on Michaelis-Menten Kinetics. *Journal of Water Resource and Protection*, 01(05), 336–344. <https://doi.org/10.4236/jwarp.2009.15040>
- Jackson, A., & Powers, H. (2018). The Cancer Process. In *JAmerica Insitue for Cancer Research*. <https://doi.org/10.1001/jama.1930.02720200064036>
- Jenuth, J. P. (2000). *The NCBI. Publicly available tools and resources on the Web*. Methods in Molecular Biology (Clifton, N.J.). <https://doi.org/10.1385/1-59259-192-2:301>
- Jia, M., Dahlman-Wright, K., & Gustafsson, J. Å. (2015). Estrogen receptor alpha and beta in health and disease. *Best Practice and Research: Clinical Endocrinology and Metabolism*, 29(4), 557–568. <https://doi.org/10.1016/j.beem.2015.04.008>
- Jumper, J., Evans, R., Pritzel, A., Green, T., Figurnov, M., Ronneberger, O., Tunyasuvunakool, K., Bates, R., Židek, A., Potapenko, A., Bridgland, A., Meyer, C., Kohl, S. A. A., Ballard, A. J., Cowie, A., Romera-Paredes, B., Nikolov, S., Jain, R., Adler, J., ... Hassabis, D. (2021). Highly accurate protein structure prediction with AlphaFold. *Nature*, 596(7873), 583–589. <https://doi.org/10.1038/s41586-021-03819-2>

- Junaidin, Chaerani, S., & Fadla, N. H. (2019). *Stufi Homologi Enzim Tironase (Homo sapiens) dengan Mnegginakan Swis-Model. VI(1)*, 1–7.
- Kalontong, P. K., Safithri, M., & Tarman, K. (2022). Penambatan Molekul Senyawa Aktif Spirulina platensis sebagai Inhibitor TMPRSS2 untuk Mencegah Infeksi SARS-COV-2. *Jurnal Pengolahan Hasil Perikanan Indonesia*, 25(2), 253–267. <https://doi.org/10.17844/jphpi.v25i2.40645>
- Kamelia, T., & Putri, N. A. R. (2023). Perkembangan Precision Medicine di Indonesia. *Indonesian Journal Chest*, 10(1), 1–12.
- Karina, K., Rosadi, I., Rosliana, I., & Wahyuningsih, K. A. (2019). Studi in Silico: Prediksi Pengaruh Mutasi Titik Gen Vascular Endothelial Growth Factor Receptor-2 (Vegfr2) Terhadap Struktur Sekunder Protein. *Al-Kauniah: Jurnal Biologi*, 12(2), 220–228. <https://doi.org/10.15408/kauniah.v12i2.11804>
- Kemenkes RI.(2019). *Penyakit Kanker di Indonesia Berada Pada Urutan 8 di Asia Tenggara dan Urutan 23 di Asia*. Diakses dari: <http://p2p.kemkes.go.id/penyakit-kanker-di-indonesia-berada-pada-urutan-8-di-asia-tenggara-dan-urutan-23-di-asia/>
- Khan, A., Ashfaq-Ur-Rehman, Junaid, M., Li, C. D., Saleem, S., Humayun, F., Shamas, S., Ali, S. S., Babar, Z., & Wei, D. Q. (2020). Dynamics Insights Into the Gain of Flexibility by Helix-12 in ESR1 as a Mechanism of Resistance to Drugs in Breast Cancer Cell Lines. *Frontiers in Molecular Biosciences*, 6(January), 1–14. <https://doi.org/10.3389/fmolb.2019.00159>
- Klein, M. E., Parvez, M. M., & Shin, J. G. (2017). Clinical Implementation of Pharmacogenomics for Personalized Precision Medicine: Barriers and Solutions. *Journal of Pharmaceutical Sciences*, 106(9), 2368–2379. <https://doi.org/10.1016/j.xphs.2017.04.051>
- Kumar, P., & Arya, A. (2018). Ramachandran plot - A simplified approach. *Pathfinder Research and Training Foundation, India, January*, 6. https://www.researchgate.net/publication/330158666_Ramachandran_plot-_A_simplified_approach
- Kwakowsky, A., Milne, M. R., Waldvogel, H. J., & Faull, R. L. (2016). Effect of estradiol on neurotrophin receptors in basal forebrain cholinergic neurons: Relevance for Alzheimer’s disease. *International Journal of Molecular Sciences*, 17(12), 1–28. <https://doi.org/10.3390/ijms17122122>
- Labellapansa, A., Muhimmah, I., & Indrayanti. (2013). Klasifikasi Citra Imunohistokimia Sel Kanker Payudara HER2 Skore 1+ dan 3+. *Seminar Nasional Informatika Medis (SNIMed) IV*

- Laskowski, R. A. (2001). PDBsum: Summaries and analyses of PDB structures. *Nucleic Acids Research*, 29(1), 221–222. <https://doi.org/10.1093/nar/29.1.221>
- Laskowski, R. A., MacArthur, M. W., Moss, D. S., & Thornton, J. M. (1993). PROCHECK: a program to check the stereochemical quality of protein structures. *Journal of Applied Crystallography*, 26, 283–291. <https://doi.org/10.1107/s0021889892009944>
- Lewandowska, A. M., Rudzki, M., Rudzki, S., Lewandowski, T., & Laskowska, B. (2019). Environmental risk factors for cancer - review paper. *Annals of Agricultural and Environmental Medicine*, 26(1), 1–7. <https://doi.org/10.26444/aaem/94299>
- Liang, J., & Shang, Y. (2013). Estrogen and cancer. *Annual Review of Physiology*, 75(9), 9.1-9.16. <https://doi.org/10.1146/annurev-physiol-030212-183708>
- Liao, P. Y., & Lee, K. H. (2010). From SNPs to functional polymorphism: The insight into biotechnology applications. *Biochemical Engineering Journal*, 49(2), 149–158. <https://doi.org/10.1016/j.bej.2009.12.021>
- Lovell, S. C., Davis, I. W., Adrendall, W. B., de Bakker, P. I. W., Word, J. M., Prisant, M. G., Richardson, J. S., & Richardson, D. C. (2003). Structure validation by C alpha geomF. *Proteins: Structure, Function, and Genetics*, 437–450. <http://onlinelibrary.wiley.com/store/10.1002/prot.10286/asset/10286 ftp.pdf?v=1&t=gwhx9jy0&s=b3b0f129a5acf7f4513aea04d22aad7ee4f4a89d>
- Lopez, M.J., S. Shamim., Mohiuddin.(2013). Biochemistry, Essential Amino Acids. [Online]. Diakses dari <https://www.ncbi.nlm.nih.gov/books/NBK557845/>
- Lu, Y., & Liu, W. (2020). Selective Estrogen Receptor Degradors (SERDs): A Promising Strategy for Estrogen Receptor Positive Endocrine-Resistant Breast Cancer. *Journal of Medicinal Chemistry*, 63(24), 15094–15114. <https://doi.org/10.1021/acs.jmedchem.0c00913>
- Łukasiewicz, S., Czezelewski, M., Forma, A., Baj, J., Sitarz, R., & Stanisławek, A. (2021). Breast Cancer—Epidemiology, Risk Factors, Classification, Prognostic Markers, and Current Treatment Strategies— An Updated Review. *Cancers*, 13, 1–30.
- Luo, J., Wu, M., Gopukumar, D., & Zhao, Y. (2016). Big Data Application in Biomedical Research and Health Care: A literature Review. *Biomed Inform Insight*, 8, 1–10. <https://doi.org/10.1051/jphyscol:1981825>

- Mappa, A. I. D., Arisadewo, L., Anwar, K., & Hasugian, R. A. (2021). Pemanfaatan Big Data dalam Mengangani Data Peningkatan Pasien COVID-19 di Indonesia. *Jurnal Teknik Informatika Unika St. Thomas*, 6(1), 178–187.
- Mardianingrum, R., Bachtiar, K. R., Susanti, S., Aas Nuraisah, A. N., & Ruswanto, R. (2021). Studi In Silico Senyawa 1,4-Naphthalenedione-2-Ethyl-3-Hydroxy sebagai Antiinflamasi dan Antikanker Payudara. *ALCHEMY Jurnal Penelitian Kimia*, 17(1), 83. <https://doi.org/10.20961/alchemy.17.1.43979.83-95>
- Marshall, S. F., Clarke, C. A., Deapen, D., Henderson, K., Largent, J., Neuhausen, S. L., Reynolds, P., Ursin, G., Horn-Ross, P. L., Stram, D. O., Templeman, C., & Bernstein, L. (2010). Recent breast cancer incidence trends according to hormone therapy use: The California Teachers Study cohort. *Breast Cancer Research*, 12(1), 1–9. <https://doi.org/10.1186/bcr2467>
- Martaadisoebrata, D. (2019). Personalized Medicine pada Pelayanan Kesehatan Reproduksi Personalized Medicine in Reproductive Health Service. *Obstetrics & Gynecology Science*, 2(1), 1–13.
- Maryanto, B. (2017). Big Data dan Pemanfaatannya Dalam Berbagai Sektor. *Media Informatika*, 16(2), 14–19. https://jurnal.likmi.ac.id/Jurnal/7_2017/0717_02_BudiMaryanto.pdf
- Mathur, R., Rana, B. S., & Jha, A. K. (2018). SNP, single nucleotide polymorphism. *Encyclopedia of Animal Cognition and Behavior*. https://doi.org/https://doi.org/10.1007/978-3-319-47829-6_2049-1
- Mayrovitz, H. N. (2022). *Breast Cancer*. Excon Publication: Brisbane
- McNutt, A. T., Francoeur, P., Aggarwal, R., Masuda, T., Meli, R., Ragoza, M., Sunseri, J., & Koes, D. R. (2021). GNINA 1.0: molecular docking with deep learning. *Journal of Cheminformatics*, 13(1), 1–20. <https://doi.org/10.1186/s13321-021-00522-2>
- Menteri Kesehatan RI. (2022). Peraturan Menteri Kesehatan Republik Indonesia Nomor 13 Tahun 2022 Tentang Perubahan Atas Peraturan Menteri Kesehatan Nomor 21 Tahun 2020 Tentang Rencana Strategis Kementerian Kesehatan Tahun 2020-2024. [https://peraturan.bpk.go.id/Home/Download/212694/Permenkes Nomor 13 Tahun 2022.pdf](https://peraturan.bpk.go.id/Home/Download/212694/Permenkes%20Nomor%2013%20Tahun%202022.pdf)
- Morihito, R. V. S. ., Chungdinata, S. E., Nazareth, T. A., Pulukadang, M. I., Makalew, R. A. ., & Pinontoan, B. (2017). Identifikasi Perubahan Struktur Dna Terhadap Pembentukan Sel Kanker Menggunakan Dekomposisi Graf. *Jurnal Ilmiah Sains*, 17(2), 153. <https://doi.org/10.35799/jis.17.2.2017.17368>

- Movérare-Skrtic, S., Börjesson, A. E., Farman, H. H., Sjögren, K., Windahl, S. H., Lagerquist, M. K., Andersson, A., Stubelius, A., Carlsten, H., Gustafsson, J. Å., & Ohlsson, C. (2014). The estrogen receptor antagonist ICI 182,780 can act both as an agonist and an inverse agonist when estrogen receptor α AF-2 is modified. *Proceedings of the National Academy of Sciences of the United States of America*, *111*(3), 1180–1185. <https://doi.org/10.1073/pnas.1322910111>
- Muhammed, M. T., & Aki-Yalcin, E. (2022). Molecular Docking: Principles, Advances, and its Applications in Drug Discovery. *Letters in Drug Design & Discovery*, *19*, 1–16. <https://doi.org/10.2174/1570180819666220922103109>
- Naithani, N., Sinha, S., Misra, P., Vasudevan, B., & Sahu, R. (2021). Precision medicine: Concept and tools. *Medical Journal Armed Forces India*, *77*(3), 249–257. <https://doi.org/10.1016/j.mjafi.2021.06.021>
- Nathan, M. R., & Schmid, P. (2017). A Review of Fulvestrant in Breast Cancer. *Oncology and Therapy*. <https://doi.org/10.1007/s40487-017-0046-2>
- National Institutes of Health (US); Biological Sciences Curriculum Study. NIH Curriculum Supplement Series Bethesda (MD): National Institutes of Health (US). (2007). *Understanding Cancer*. Diakses dari <https://www.ncbi.nlm.nih.gov/books/NBK20362/>
- Nicoloso, M. S., Sun, H., Spizzo, R., Kim, H., Wickramasinghe, P., Shimizu, M., Wojcik, S. E., Ferdin, J., Kunej, T., Xiao, L., Manoukian, S., Secretò, G., Ravagnani, F., Wang, X., Radice, P., Croce, C. M., Davuluri, R. V., & Calin, G. A. (2010). Single-nucleotide polymorphisms inside microRNA target sites influence tumor susceptibility. *Cancer Research*, *70*(7), 2789–2798. <https://doi.org/10.1158/0008-5472.CAN-09-3541>
- Nurmayeni, & Windarti, I. (2022). Gambaran Subtipe Molekuler Kanker Payudara di Indonesia. *Agromedicine*, *9*(5), 145–148.
- Osborne, C. K., Wakeling, A., & Nicholson, R. I. (2004). Fulvestrant: An oestrogen receptor antagonist with a novel mechanism of action. *British Journal of Cancer*, *90*, S2–S6. <https://doi.org/10.1038/sj.bjc.6601629>
- Pagadala, N. S., Syed, K., & Tuszynski, J. (2017). Software for molecular docking: a review. *Biophysical Reviews*, *9*(2), 91–102. <https://doi.org/10.1007/s12551-016-0247-1>
- Park, S. H., Jung, I. K., & Kim, J. H. (2020). Effect of dietary changes from high-fat diet to normal diet on breast cancer growth and metastasis. *Journal of Nutrition and Health*, *53*(4), 369–380. <https://doi.org/10.4163/JNH.2020.53.4.369>

- Patel, A. (2020). Benign vs Malignant Tumors. *JAMA Oncology*, 6(9), 1488. <https://doi.org/10.1001/jamaoncol.2020.2592>
- Pham, T. H., Lecomte, S., Efstathiou, T., Ferriere, F., & Pakdel, F. (2019). An update on the effects of glyceollins on human health: Possible anticancer effects and underlying mechanisms. *Nutrients*, 11(1). <https://doi.org/10.3390/nu11010079>
- Pradana, M. S., Apsari, G. R., Adawiyah, R., Linatari, M. A., & Rahmayadi, D. (2019). Program Pengajaran Sequence Alignment Berbasis Bioinformatika di MA Matholi'ul Anwar Lamongan. *Jurnal Pengabdian Kepada Masyarakat*, 3(1), 101–106.
- Prieto-Martínez, F. D., Arciniega, M., & Medina-Franco, J. L. (2018). Acoplamiento Molecular: Avances Recientes y Retos. *TIP Revista Especializada En Ciencias Químico-Biológicas*, 21, 65–87. <https://doi.org/10.22201/fesz.23958723e.2018.0.143>
- Purba, E. F., & Simanjuntak, P. (2012). *Metode Penelitian*. Percetakan SADIA: Medan
- Puspawati, P. R., Kristina, S. A., & Wiedyaningsih, C. (2020). Dampak merokok terhadap kematian dini akibat kanker di Indonesia: estimasi years of life lost (YLL). *J Majalah Farmaseutik*, 16(1), 104. <https://doi.org/10.22146/farmaseutik.v16i1.49790>
- Rafeeq, M. M. (2022). Molecular docking analysis of phytochemicals with estrogen receptor alpha. *Bioinformation*, 18(8), 697–702. <https://doi.org/10.6026/97320630018697>
- Rahayu, Y. C., & Auerkari, E. (2004). Teknik Immuno Histokimia Sebagai Pendeteksi Penyakit Infeksi. *Jurnal Kedokteran Gigi Universitas Indonesia*, 11(2), 76–82.
- Rahmadhayanti, E., Hayati, L., & Saleh, M. (2014). Hubungan polimorfisme gen reseptor angiotensin II tipe 1 1166 A/C dengan kejadian preeklampsia. *Jurnal Kedokteran*, 46(1), 52–58.
- Reynaldi, M. A., & Setiawansyah, A. (2022). Potensi anti-kanker payudara tanaman songga (*Strychnos lucida* R.Br): Tinjauan interaksi molekuler terhadap reseptor estrogen- α in silico. *Sasambo Journal of Pharmacy*, 3(1), 30–35. <https://doi.org/10.29303/sjp.v3i1.149>
- Robertson, J. F. R., Come, S. E., Jones, S. E., Beex, L., Kaufmann, M., Makris, A., Nortier, J. W. R., Possinger, K., & Rutqvist, L. E. (2005). Endocrine treatment options for advanced breast cancer - The role of fulvestrant. *European Journal of Cancer*, 41(3), 346–356. <https://doi.org/10.1016/j.ejca.2004.07.035>

- Robertson, J. F. R., & Harrison, M. (2004). Fulvestrant: Pharmacokinetics and pharmacology. *British Journal of Cancer*, *90*, S7–S10. <https://doi.org/10.1038/sj.bjc.6601630>
- Robinson, M., Atmakusumah, T. D., Irawan, C., & Shatri, H. (2019). Immunohistochemistry Profile of Breast Cancer Patients that Get Anthracyclin – Based Chemotherapy in RSUD Kota Bogor. *Jurnal Penyakit Dalam Indonesia*, *6*(4), 173–177.
- Sadée, W., & Dai, Z. (2005). Pharmacogenetics/genomics and personalized medicine. *Human Molecular Genetics*, *14*(2), 207–214. <https://doi.org/10.1093/hmg/ddi261>
- Saini, A., Kumar, M., Bhatt, S., Saini, V., & Malik, A. (2020). Cancer Causes and Treatment. *Internation Journal of Pharmaceutical Sciences and Research*, *11*(7), 3109–3122. [https://doi.org/10.13040/IJPSR.0975-8232.11\(7\).3109-22](https://doi.org/10.13040/IJPSR.0975-8232.11(7).3109-22)
- Saragih, R. Y., Hidayah, N., & Roza, D. (2021). Docking Molekular Potensi Antikanker Payudara Protein3ert Dengan Senyawa Turunan Kuinin. *Prosiding Seminar Nasional Kimia & Pendidikan Kimia*, 376–381.
- Sari, S. E., Harahap, W. A., & Saputra, D. (2018). Pengaruh Faktor Risiko Terhadap Ekspresi Reseptor Estrogen Pada Penderita Kanker Payudara Di Kota Padang. *Jurnal Kesehatan Andalas*, *7*(4), 461. <https://doi.org/10.25077/jka.v7.i4.p461-468.2018>
- Saudale, F. Z. (2020). Pemodelan Homologi Komparatif Struktur 3d Protein dalam Desain dan Pengembangan Obat. *Al-Kimia*, *8*(1), 93–103. <https://doi.org/10.24252/al-kimia.v8i1.9463>
- Scabia, V., Ayyanan, A., De Martino, F., Agnoletto, A., Battista, L., Laszlo, C., Treboux, A., Zaman, K., Stravodimou, A., Jallut, D., Fiche, M., Bucher, P., Ambrosini, G., Sflomos, G., & Brisken, C. (2022). Estrogen receptor positive breast cancers have patient specific hormone sensitivities and rely on progesterone receptor. *Nature Communications*, *13*(1). <https://doi.org/10.1038/s41467-022-30898-0>
- Setiawan, H., & Irawan, M. I. (2017). Kajian Pendekatan Penempatan Ligan Pada Protein Menggunakan Algoritma Genetika. *Jurnal Sains Dan Seni ITS*, *6*(2), 2–6. <https://doi.org/10.12962/j23373520.v6i2.25468>
- Setiawan, I. M. A. (2023). Peran Pemeriksaan Imunohistokimia dalam Diagnosis dan Prognosis Kanker Payudara. *Cermin Dunia Kedokteran*. *50*(8), 443–446.
- Shah, R., Rosso, K., & David Nathanson, S. (2014). Pathogenesis, prevention, diagnosis and treatment of breast cancer. *World Journal of Clinical Oncology*, *5*(3), 283–298. <https://doi.org/10.5306/wjco.v5.i3.283>

- Shatoff, E., & Bundschuh, R. (2020). Single nucleotide polymorphisms affect RNA-protein interactions at a distance through modulation of RNA secondary structures. *PLoS Computational Biology*, *16*(5), 1–16. <https://doi.org/10.1371/journal.pcbi.1007852>
- Shen, L. X., Babilion, J. P., & Stanton, J. (1999). Single-nucleotide polymorphisms can cause different structural folds of mRNA. *Proceedings of the National Academy of Sciences of the United States of America*, *96*(14), 7871–7876. <https://doi.org/10.1073/pnas.96.14.7871>
- Shiovitz, S., & Korde, L. A. (2015). Genetics of breast cancer: a topic in evolution. *Annals of Oncology*, *26*(7), 1291–1299.
- Singh, V. ., Singh, A. ., Chand, R., & Kushwaha, C. (2011). Role of Bioinformatics in Agriculture and Sustainable Development. *International Journal of Bioinformatics Research*, *3*(2), 221–226. https://doi.org/10.1007/978-3-031-01949-4_44
- Sinha, T. (2018). Tumors: Benign and Malignant. *Cancer Therapy & Oncology International Journal*, *10*(3), 1–3. <https://doi.org/10.19080/ctoj.2018.10.555790>
- Soen, G. I. E., Marlina, & Renny. (2022). Implementasi Cloud Computing dengan Google Colaboratory Pada Aplikasi Pengolah Data Zoom Participants. *Journal Informatic Technology And Communication*, *6*(1), 24–30.
- Suhara.(2018). *Dasar-dasar Biokimia*. Prisma Press: Bandung
- Sulaiman, T. W., Fitriansyah, R. B., Alaudin, A. R., & Ratsanjani, M. H. (2023). Penerapan Big Data Dalam Kesehatan Masyarakat. *Jurnal Sains, Teknik, Dan Studi Kemasyarakatan*, *1*(3), 129–138.
- Sun, Z., Shi, Y., Shen, Y., Cao, L., Zhang, W., & Guan, X. (2015). Analysis of different HER-2 mutations in breast cancer progression and drug resistance. *Journal of Cellular and Molecular Medicine*, *19*(12), 2691–2701. <https://doi.org/10.1111/jcmm.12662>
- Sunarto, A. A. (2015). Perbandingan Program Sequence Alignment. *Jurnal Rekayasa Nusaputra*, *1*(1), 1–6
- Syahputra, G., Ambarsari, L., & Sumaryada, T. (2014). Simulasi Docking Kurkumin Enol, Bisdemetoksikurkumin dan Analognya sebagai Inhibitor Enzim12-Lipoksigenase. *Jurnal Biofisika*, *10*(1), 55–67.
- Teng, S., Madej, T., Panchenko, A., & Alexov, E. (2009). Modeling effects of human single nucleotide polymorphisms on protein-protein interactions. *Biophysical Journal*, *96*(6), 2178–2188. <https://doi.org/10.1016/j.bpj.2008.12.3904>

- Torres, P. H. M., Sodero, A. C. R., Jofily, P., & Silva-Jr, F. P. (2019). Key topics in molecular docking for drug design. *International Journal of Molecular Sciences*, 20(18), 1–29. <https://doi.org/10.3390/ijms20184574>
- Toy, W., Shen, Y., Won, H., Green, B., Sakr, R. A., Will, M., Li, Z., Gala, K., Fanning, S., King, T. A., Hudis, C., Chen, D., Taran, T., Hortobagyi, G., Greene, G., Berger, M., Baselga, J., & Chandarlapaty, S. (2013). ESR1 ligand-binding domain mutations in hormone-resistant breast cancer. *Nature Genetics*, 45(12), 1439–1445. <https://doi.org/10.1038/ng.2822>
- Triwani, T., & Saleh, I. (2015). Single Nucleotide Polymorphism Promoter -765g / C Gen Cox-2 Sebagai Faktor Risiko Terjadinya Karsinoma Kolorektal. *Biomedical Journal of Indonesia*, 1(1), 2–10.
- Trott, O., & Olson, A. J. (2010). Software News and Updates Gabbit — AutoDock Vina: Improving the Speed and Accuracy of Docking with a New Scoring Function, Efficient Optimization, and Multithreading. *Journal of Computational Chemistry*, 31(2), 455–461. <https://doi.org/10.1002/jcc>
- Tsunoda, D. F., Freitas, A. A., & Lopes, H. S. (2011). A genetic programming method for protein motif discovery and protein classification. *Soft Computing*, 15(10), 1897–1908. <https://doi.org/10.1007/s00500-010-0624-9>
- Tyagi, N., Sharma, G. N., Shrivastava, B., & Chaudhary, N. (2017). *An International Open access peer reviewed journal Cancer: An Overview Article Information* : 6(5), 2740–2747.
- Vinsentricia, A., Hami Seno, D. S., & Bintang, M. (2015). *In Silico* Analysis of *Curcuma longa* Against PCAF Histon Asetiltransferase. *Current Biochemistry*, 2(2), 52–62. <https://doi.org/10.29244/cb.2.2.52-62>
- Wadood, A., Ahmed, N., Shah, L., Ahmad, A., Hassan, H., & Shams, S. (2013). In-silico drug design: An approach which revolutionarised the drug discovery process. *OA Drug Design and Delivery*, 1(1), 1–4. <https://doi.org/10.13172/2054-4057-1-1-1119>
- Wibawa., I. D. N. (2017). Personalized medicine in Gastroentero-hepatology. *Denpasar Update in Gastroentero-Hepatology*, 4-9
- Widyastuti, M. D., Noviyanti, N. K. M., Sanjaya, I. K. N. S., & Susanti, N. M. P. (2020). Aktivitas Antihiperpigmentasi Likopen Secara in Silico. *Jurnal Kimia*, 14(2), 107. <https://doi.org/10.24843/jchem.2020.v14.i02.p01>
- Yaşar, P., Ayaz, G., User, S. D., Güpür, G., & Muyan, M. (2017). Molecular mechanism of estrogen–estrogen receptor signaling. *Reproductive Medicine and Biology*, 16(1), 4–20. <https://doi.org/10.1002/rmb2.12006>

- Yuan, S., Chan, H. C. S., & Hu, Z. (2017). Using PyMOL as a platform for computational drug design. *Wiley Interdisciplinary Reviews: Computational Molecular Science*, 7(2), 1–10. <https://doi.org/10.1002/wcms.1298>
- Zardecki, C., Dutta, S., Goodsell, D. S., Voigt, M., & Burley, S. K. (2016). RCSB Protein Data Bank: A Resource for Chemical, Biochemical, and Structural Explorations of Large and Small Biomolecules. *Journal of Chemical Education*, 93(3), 569–575. <https://doi.org/10.1021/acs.jchemed.5b00404>
- Zhang, C., Guo, S., Yang, L., Liu, J., Zheng, S., Zhong, Q., Zhang, Q., & Wang, G. (2017). Metabolism, pharmacokinetics, and bioavailability of ZB716, a Steroidal Selective Estrogen Receptor Downregulator (SERD). *Oncotarget*, 8(61), 103874–103889. <https://doi.org/10.18632/oncotarget.21808>