

**SIMULASI DINAMIKA MOLEKULER ANDROGRAFOLID
DAN 2 TURUNANNYA TERHADAP
ACETYLCHOLINESTERASE DAN *BETA-SECRETASE 1*
SEBAGAI KANDIDAT ANTI ALZHEIMER**

SKRIPSI

**ALMIRA PINKAN CARISSA
A221089**



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YAYASAN HAZANAH
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Sebagai salah satu syarat untuk memperoleh gelar Sarjana Farmasi

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Kutipan atau saduran baik sebagian ataupun seluruh naskah harus menyebutkan nama pengarang dan sumber aslinya, yaitu Sekolah Tinggi Farmasi Indonesia

Skripsi ini saya persembahkan untuk Mama tercinta dan keluarga tercinta.
Terima kasih telah menjadi alasan saya untuk terus berjuang, menjadi rumah
tempat saya pulang, dan menjadi sumber kekuatan terbesar dalam
perjalanan akademik ini.

ABSTRAK

Penyakit Alzheimer merupakan penyakit neurodegeneratif progresif yang ditandai oleh penurunan fungsi kognitif dan memori, dengan *acetylcholinesterase* (AChE) dan β -secretase 1 (BACE1) sebagai dua target terapeutik utama. Penelitian ini bertujuan mengevaluasi potensi andrografolid, asam andrografik, dan 14-deoksi-11,12-didehidroandrografolid sebagai kandidat inhibitor multitarget AChE dan BACE1 melalui simulasi dinamika molekuler selama 10 ns, dengan donepezil dan verubecestat sebagai kontrol positif serta E20 dan M7D sebagai ligan alami. Parameter yang dianalisis meliputi *Root Mean Square Deviation*, *Root Mean Square Fluctuation*, okupansi ikatan hidrogen, dan energi bebas ikatan dengan metode MM-GBSA. Hasil simulasi menunjukkan bahwa seluruh kompleks uji mampu mempertahankan kestabilan konformasi, dengan 14-deoksi-11,12-didehidroandrografolid menunjukkan profil RMSD ligan paling stabil terhadap kedua reseptor, sementara fluktuasi residu terutama terjadi pada daerah loop dan terminal yang secara alami bersifat fleksibel. Analisis ikatan hidrogen menunjukkan interaksi persisten dengan residu-residu kunci pada sisi aktif kedua enzim. Berdasarkan perhitungan MM-GBSA, 14-deoksi-11,12-didehidroandrografolid menunjukkan afinitas pengikatan paling kompetitif terhadap AChE dengan nilai energi bebas total sebesar -42,9 kkal/mol, sedangkan andrografolid menunjukkan afinitas terbaik terhadap BACE1 dengan nilai -34,7 kkal/mol, mendekati ligan alami M7D. Secara keseluruhan, andrografolid dan turunannya berpotensi dikembangkan sebagai kandidat inhibitor multitarget AChE dan BACE1 untuk pengembangan agen terapeutik Alzheimer, meskipun validasi in vitro dan in vivo tetap diperlukan.

Kata kunci: Alzheimer, andrografolid, AChE, BACE1, *molecular docking*, dinamika molekuler, MM-GBSA.

ABSTRACT

Alzheimer's disease is a progressive neurodegenerative disease characterized by a decline in cognitive and memory functions, with acetylcholinesterase (AChE) and β -secretase 1 (BACE1) as two major therapeutic targets. This study aimed to evaluate the potential of andrographolide, andrographic acid, and 14-deoxy-11,12-didehydroandrographolide as multitarget inhibitors of AChE and BACE1 through 10 ns molecular dynamics simulations, with donepezil and verubecestat as positive controls and E20 and M7D as native ligands. The analyzed parameters included root mean square deviation, root mean square fluctuation, hydrogen bond occupancy, and binding free energy calculated using the MM-GBSA method. The simulation results showed that all test complexes were able to maintain conformational stability, with 14-deoxy-11,12-didehydroandrographolide exhibiting the most stable ligand RMSD profile toward both receptors, while residue fluctuations mainly occurred in loop and terminal regions that are naturally flexible. Hydrogen bond analysis revealed persistent interactions with key residues in the active sites of both enzymes. Based on the MM-GBSA calculations, 14-deoxy-11,12-didehydroandrographolide showed the most competitive binding affinity toward AChE, with a total free energy value of -42.9 kcal/mol, whereas andrographolide showed the best affinity toward BACE1, with a value of -34.7 kcal/mol, approaching that of the native ligand M7D. Overall, andrographolide and its derivatives have the potential to be developed as multitarget inhibitors of AChE and BACE1 for the development of therapeutic agents for Alzheimer's disease, although in vitro and in vivo validation is still required.

Keywords: Alzheimer's disease, andrographolide, AChE, BACE1, molecular docking, molecular dynamics, MM-GBSA.

KATA PENGANTAR

Bismillahirrahmanirrahim,

Puji dan syukur penulis panjatkan ke hadirat Allah SWT atas segala berkah rahmat dan ridha-Nya penulis dapat menyelesaikan penelitian dan penulisan skripsi yang berjudul **“Simulasi Dinamika Molekuler Andrografolid Dan 2 Turunannya Terhadap Acetylcholinesterase Dan Beta-secretase 1 Sebagai Kandidat Anti Alzheimer”**.

Penelitian dan penulisan skripsi ini dilakukan untuk memenuhi salah satu syarat untuk mendapatkan gelar sarjana pada jurusan Farmasi Sekolah Tinggi Farmasi Indonesia

Penulis mengucapkan terima kasih kepada dosen pembimbing bapak apt. Seno Aulia Ardiansyah, M.Si dan ibu apt. Maria Ulfah, M.Si atas bimbingan, nasihat, dukungan, serta pengorbanan yang diberikan. Pada kesempatan ini, tidak lupa penulis mengucapkan terima kasih yang sebesar-besarnya kepada :

1. Dr. apt. Adang Firmansyah, M.Si., selaku Ketua Sekolah Tinggi Farmasi Indonesia,
2. Dr. apt. Diki Payugo, M.Si., selaku Wakil Ketua I Bidang Akademik,
3. Dr. apt. Hesti Riasari, M.Si., selaku Ketua Program Studi Sarjana Farmasi,
4. Dr. apt. Diki Prayugo, M.Si, selaku Wali Dosen yang telah banyak memberikan bimbingan dan arahan kepada penulis,
5. Seluruh staf dosen, staf administrasi, serta karyawan Sekolah Tinggi Farmasi Indonesia,
6. Mama dan keluarga yang senantiasa memberikan doa, kasih sayang, dukungan moril maupun materiil, serta semangat yang tiada henti kepada penulis. Tanpa pengorbanan, kesabaran, dan cinta kasih Mama, penulis tidak akan mampu sampai pada tahap ini.
7. Seluruh Angkatan 2022, serta teman-teman terdekat saya yang selalu saling menyemangati dan mendukung satu sama lain.

Penulis menyadari bahwa skripsi ini masih memiliki keterbatasan. Oleh sebab itu, penulis sangat mengharapkan kritik, saran, dan masukan yang konstruktif dari berbagai pihak sebagai bahan evaluasi untuk penyempurnaan karya ilmiah di masa mendatang. Besar harapan penulis agar skripsi ini dapat memberikan kontribusi positif, menambah wawasan bagi pembaca, serta menjadi salah satu referensi yang bermanfaat dalam pengembangan ilmu pengetahuan, khususnya di bidang yang berkaitan dengan penelitian ini.

Bandung, Juli 2026
Penulis

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