

A Monte Carlo simulation study of vinblastine and vincristine as clinical drugs

Shiva Joohari¹ and Majid Monajjemi^{2*}

¹Ph.D. Student, Department of Chemistry, Tehran Science and Research Branch, Islamic Azad University, Tehran, Iran

²Department of Chemistry, Tehran Science and Research Branch, Islamic Azad University, Tehran, Iran

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ABSTRACT

In this study, Monte Carlo statistical mechanical simulations for vinblastine and vincristine were carried out in standard manner using the Metropolis sampling technique in canonical (T, V, N) ensemble. Geometrical optimizations of vinblastine and vincristine were carried out with the HF method coupled to 6-31G(d) basis sets for all atoms. Simulation was done by four force fields of MM⁺, BIO⁺, AMBER and OPLS. Some important energy parameters such as Potential Energy and Total Energy in ten different simulating temperatures (300, 302, 304, 306, 308, 310, 312, 314, 316 and 318 Kelvin) were used for computation.

Keywords: Vinblastine; Vincristine; Monte Carlo

INTRODUCTION

The vinca alkaloids are a subset of drugs derived from the Madagascar periwinkle plant. They were discovered in the 1950's by Canadian scientists, Robert Noble and Charles Beer [1, 2]. Vinca alkaloids have been found to control diabetes, high blood pressure, and are the drugs that have even been used as disinfectants. There are four major vinca alkaloids (vinblastine, vinorelbine, vincristine, and vindesine) that are applied in clinical usages [3].

The vinca alkaloids are cytotoxics so that they halt the division of cells and cause cell death. During cell division, vinca alkaloid molecules bind to the

building blocks of a protein called tubulin and prevent from its formation. Tubulin protein normally works in cells to create "spindle fibers," (also called microtubules). These microtubules provide cells with both the structure and flexibility they need to divide and replicate. Without microtubules, cells cannot divide. The vinca alkaloid's mechanism in a nutshell: by occupying tubulin's building block structure, vinca alkaloids inhibit cancer cells division. The antitumor activity of vinblastine is ascribed to primarily inhibition of mitosis at metaphase through its interaction with tubulin. Vinblastine

*Corresponding author: m_monajjemi@yahoo.com

binds to the microtubular proteins of the mitotic spindle causes microtubule crystallization so that mitotic arrest or cell death [4, 5].

Vincristine's inhibition of microtubule formation is especially powerful. Dynamic character of tubulin protein is the reason for this fact. Its long chain of building blocks is always growing in some places and breaking in others. The less contiguous parts of a tubulin molecule have pieces only two building blocks long, called dimers. Vincristine has a high affinity for tubulin dimers, and the reaction between vincristine and the dimers is rapidly reversible. That means a vincristine molecule will attach to a dimer at one site, break off, and then reattach at another site. This keeps two sites per dimer "poisoned" and unable to reassemble into the protein. So vincristine's ability to destabilize tubulin is especially good [6, 7].

The aim of this work is to understand molecular mechanic of vinblastine and vincristine drugs, which will be useful for designing anticancer drugs.

THEORETICAL

Background and Computational Methods

The Monte Carlo method was invented by scientists working on the atomic bomb in the 1940s, who named it for the city in Monaco famed for its casinos and games of chance. Its core idea is to use random samples of parameters or inputs to explore the behavior of a complex system or process [8]. Molecular mechanic simulation method are especially useful in studying systems with a large number of coupled degrees of freedom, such as liquids disordered materials, strongly coupled solids and cellular structures.

Simulation refers to methods aimed at generating a representative sampling of a system at a finite temperature [9]. Monte

Carlo (MC) methods are a class of computational algorithms that rely on repeated random sampling to compute their results.

The systems can be studied in the coarse-grained or *ab initio* frameworks depending on the desired accuracy. Computer simulations allow us to monitor the local environment of a particular molecule to see if some chemical reaction is happening for instance. We can also conduct thought experiments when the physical experiments are not feasible, for instance breaking bonds, introducing impurities at specific sites, changing the local/global structure, or introducing external fields. Biological systems such as proteins [10] membranes [11], images of cancer [12], are being studied by means of computer simulations.

Thermodynamic averages of molecular properties can be determined from MC methods, as can minimum energy structures. MC simulations require only the ability to evaluate the energy of the system, which may be advantageous if calculating the first derivative is difficult or time-consuming. Furthermore, since only a single particle is moved in each step, only the energy changes associated with this move must be calculated, not the total energy for the whole system. A disadvantage of MC methods is the lack of the time dimension and atomic velocities, and they are not suitable for studying time-dependent phenomena or properties depending on momentum.

Geometrical optimizations of vinblastine and vincristine were carried out with the HF method coupled to 6-31G(d) basis sets for all atoms. Simulation was done in MM⁺, BIO⁺, AMBER and OPLS force fields. Some important energy parameters such as Potential Energy and Total Energy in ten different simulating temperatures (300, 302, 304, 306, 308,

310, 312, 314, 316 and 318 Kelvin) were used for computation.

RESULTS AND DISCUSSION

Monte Carlo statistical mechanical simulations were carried out in standard manner using the Metropolis sampling technique [13] in canonical (T, V, N) ensemble.

The two compounds are very similar in the structure and molecular structure of vinblastine and vincristine has been presented in scheme 1.

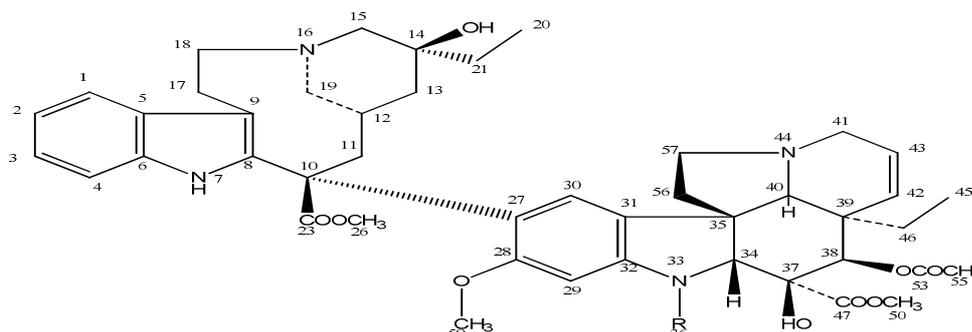
The main purpose is to find for the lowest energy, in which the molecule is in its most stable state. In this study AMBER, MM⁺, BIO⁺ and OPLS force fields were chosen. The total Potential Energy is the sum of mentioned contribution interactions based on the force fields. Therefore, force fields are a series of functional energy

parameters that evaluate performance and calculate the Potential Energy of molecule in various positions of its constituent atoms and bonds [14, 15].

In this work, we have determined all possible potential and total energy of vinblastine and in vincristine by Monte Carlo method at 300, 302, 304, 306, 308, 310, 312, 314, 316, and 318 K. The obtained valuable data for thermodynamic parameters (E potential, E total), analyzed under the different simulation procedure, various temperatures values every 10 ps span are listed in tables. Calculations of potential energy and total energy were performed by four force fields (AMBER, BIO⁺, MM⁺ and OPLS). For example, the results obtained by AMBER and MM⁺ for vinblastine and vincristine have been tabulated as tables 1-4.

Table 1. Calculated potential energy for vinblastine, belong to AMBER force field at ten different temperature

Method	Potential Energy (Kcal/mol)									
	AMBER/ Monte Carlo									
Time (PS)	300	302	304	306	308	310	312	314	316	318
10	50449000	53512342	53512342	53512342	53128563	52690522	52806650	52806650	52806651	52806650
20	1314905.8	1626318.6	1626318.6	1785355.3	1204928.9	1787593.5	2115727.6	2115727.6	2094548.3	2127119.6
30	271681.5	189228.57	189522.42	105404.19	163599.14	165276.17	116559.33	116262.21	217476.76	270463.7
40	52361.15	29296.18	37555.35	21771.44	28030.01	35234.38	18926.18	23716.78	43225.02	62895.78
50	13874.383	8497.899	10712.909	8287.879	7107.155	8711.508	8189.063	7894.82	11596.633	16761.794
60	6035.871	4427.751	5305.803	4684.934	4082.47	4404.972	4590.423	4416.86	4773.3	5101.984
70	2658.939	3458.772	3628.05	3561.057	3091.596	3166.71	3482.746	3371.445	3144.896	3418.002
80	2018.184	2980.936	2950.486	2871.28	2612.269	2720.612	2756.059	2844.769	2472.099	2707.019
90	1604.06	2608.213	2570.094	2488.824	2243.976	2475.265	2199.5	2568.873	1946.18	2265.44
100	1288.7445	2293.1791	2301.0309	2252.82	1903.6555	2263.2345	1756.3964	2332.8191	1582.1818	1831.39818



Scheme 1. Molecular structure of vinblastine and vincristine (Vincristine: R= CHO; Vinblastine: R= CH₃).

Table 2. Calculated potential energy for vinblastine, belong to MM+ force field at ten different temperature

Method	Potential Energy (Kcal/mol)									
	MM+ / Monte Carlo									
Time (PS)	300	302	304	306	308	310	312	314	316	318
10	4534.575	4534.575	4518.281	4537.923	4557.658	4406.06	4393.835	4393.835	4394.035	4382.339
20	3141.478	3118.733	3138.051	3011.141	2919.735	2892.965	2962.516	2962.516	2891.541	2913.151
30	2338.082	2294.103	2298.376	2156.649	2122.811	2072.862	2164.998	2164.998	2103.993	2177.182
40	1753.396	1795.26	1737.617	1644.85	1766.982	1628.7	1729.651	1729.651	1703.937	1678.862
50	1378.158	1369.994	1345.728	1240.39	1483.564	1346.885	1381.282	1379.448	1443.559	1357.129
60	1118.115	1057.000	1142.777	975.0752	1306.345	1145.577	1092.2192	1109.6975	1146.613	1085.6503
70	899.4439	839.9989	979.0861	809.4138	1120.114	971.6776	880.545	874.4816	981.1758	931.5397
80	736.228	700.2869	741.6991	688.4232	899.2035	852.353	763.7964	747.9081	838.3322	790.2792
90	633.1893	626.3875	628.5347	612.8694	721.9906	719.4925	665.6212	654.7784	742.1789	667.0155
100	550.19736	558.6757	573.32891	562.37427	624.42518	666.90045	580.88282	564.29009	647.041545	585.594727

Comparisons of potential energy levels in different temperatures are displayed in Figs. 1a- 1d for vinblastine and 2a- 2d for vincristine. The energy plots versus time steps for four force fields have been illustrated in figure 2a-2d and 3a-3d. The plots show the same behavior for four methods at all 10 temperature intervals.

According to results observed in tables 1 and 2 and figs 1 for vinblastine potential energy in different time steps and various force fields shows that maximum quantity observed in 310 K in OPLS force field and minimum quantity observed in 300 K, 550.19 Kcal/mol in MM+ force field.

According to results observed in tables 3 and 4 and figs 2 for vincristine potential energy in different time steps and various force fields shows that maximum quantity observed in 310 K in AMBER force field and minimum quantity observed in 310 K, 278.51 Kcal/mol in MM+ force field.

According to results observed in tables 5 and fig. 3 for vinblastine total energy in different temperature and various force fields shows that maximum quantity observed in 308 K, 2526.39 Kcal/mol in OPLS method.

According to results observed in tables 6 and fig. 4 for vincristine total energy in different temperature and various force

fields shows that maximum quantity observed in 306 K, 570.447 Kcal/mol in BIO+ method.

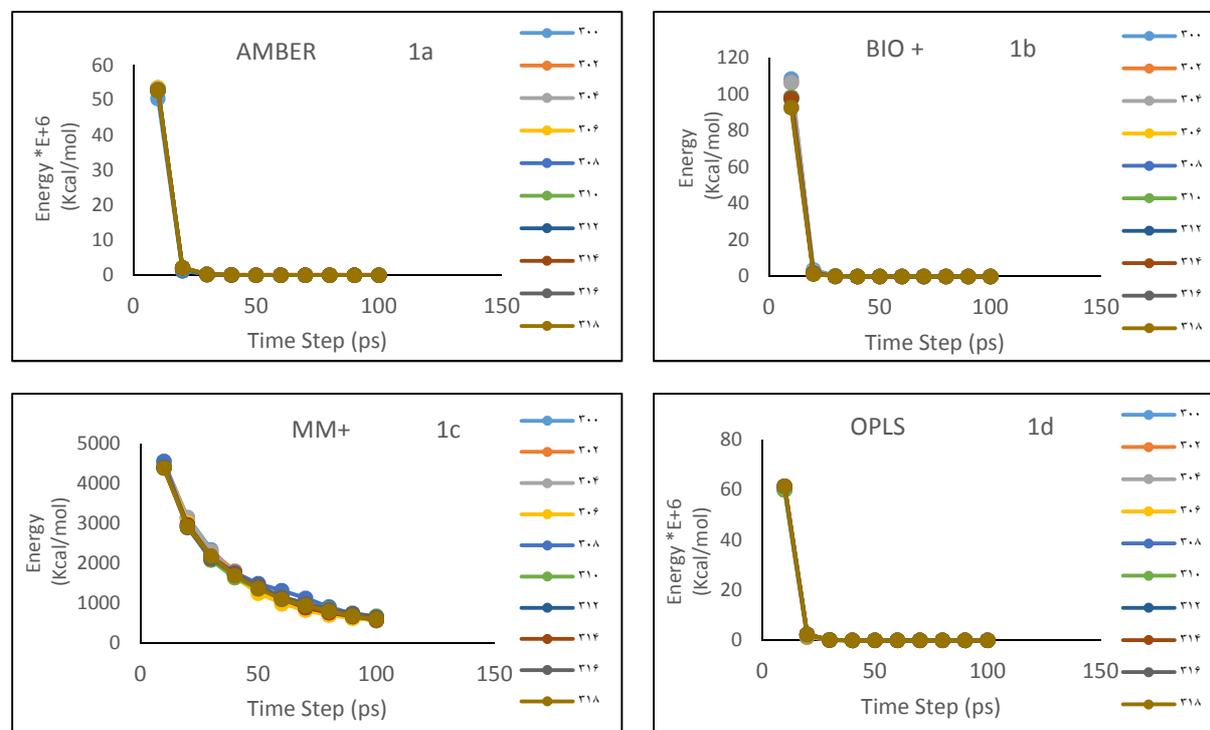


Fig. 1. The graphs of vincristine potential energy belong to a) AMBER b) BIO⁺, c) MM⁺, d) OPLS force fields.

Table 3. Calculated potential energy for vincristine, belong to AMBER force field at ten different temperature

Method	Potential Energy (Kcal/mol)									
	AMBER/ Monte Carlo									
Time (PS)	300	302	304	306	308	310	312	314	316	318
10	66669750	66671240	66671240	66671240	66671240	81036240	70086826	70075328	70075328	58988803
20	3200478.3	4887313.4	4887313.8	4887313.9	3814583.8	5311674.9	969475.5	924610.3	924609.5	177970.3
30	70518.05	58130.19	331198.4	70424.57	60196.7	65283.15	66477.72	39605.25	39520.64	7816.392
40	5442.965	3206.117	158067.4	6133.808	5069.302	6810.503	8880.389	4711.518	3892.231	1888.933
50	1933.352	1047.783	718566	1465.579	1354.736	1779.538	1911.726	1801.541	1556.908	981.6114
60	1159.876	669.871	488821.1	926.0091	724.1822	919.1727	1109.991	1048.303	956.0189	705.5823
70	860.7192	523.628	499640.3	616.8188	498.0776	666.7751	695.2647	727.495	683.5751	543.4913
80	619.9293	455.3749	375344.2	488.0953	430.2659	542.5169	504.1217	576.4321	543.4305	460.7368
90	470.1542	403.445	396553.9	418.8983	404.8856	471.2364	428.6849	487.3585	474.307	430.5735
100	404.9508	383.0372	368279.55	381.96991	382.40409	424.43782	400.20609	427.94445	443.43745	400.167

Table 4. calculated potential energy for vincristine, belong to MM⁺ force field at ten different temperature

Method	Potential Energy (Kcal/mol)									
	MM ⁺ / Monte Carlo									
Time (PS)	300	302	304	306	308	310	312	314	316	318
10	1811.181	1825.863	1774.751	1774.751	1774.751	1781.948	1781.948	1791.754	1791.754	1786.094
20	1172.071	1196.137	1068.278	1069.0069	1079.8519	1144.8081	1088.374	1128.662	1128.6629	1094.697
30	774.9873	805.4633	688.3277	691.7222	725.0924	699.1269	687.7324	739.8002	752.2391	701.8544
40	558.8123	545.9983	514.4215	518.433	554.1168	510.242	498.7968	551.805	555.8914	529.5489
50	464.0144	427.0653	426.5527	441.7188	469.7542	416.2247	430.5498	459.9416	446.453	447.3894
60	401.0245	375.6622	378.397	388.0903	420.5896	361.8982	382.0928	396.2465	395.9465	398.4633
70	372.1776	335.9098	349.9951	362.4731	382.0106	329.2889	349.4232	349.6155	352.1546	371.3351
80	345.6443	322.6189	332.1923	339.363	355.0489	309.4171	328.8694	312.5646	342.5879	342.5787
90	316.2172	308.121	309.9506	329.2566	327.4962	288.6556	321.3545	303.098	323.4118	322.2277
100	305.6673	294.8295	293.7925	306.93409	307.82491	278.51809	306.16309	287.0697	310.96718	318.11172

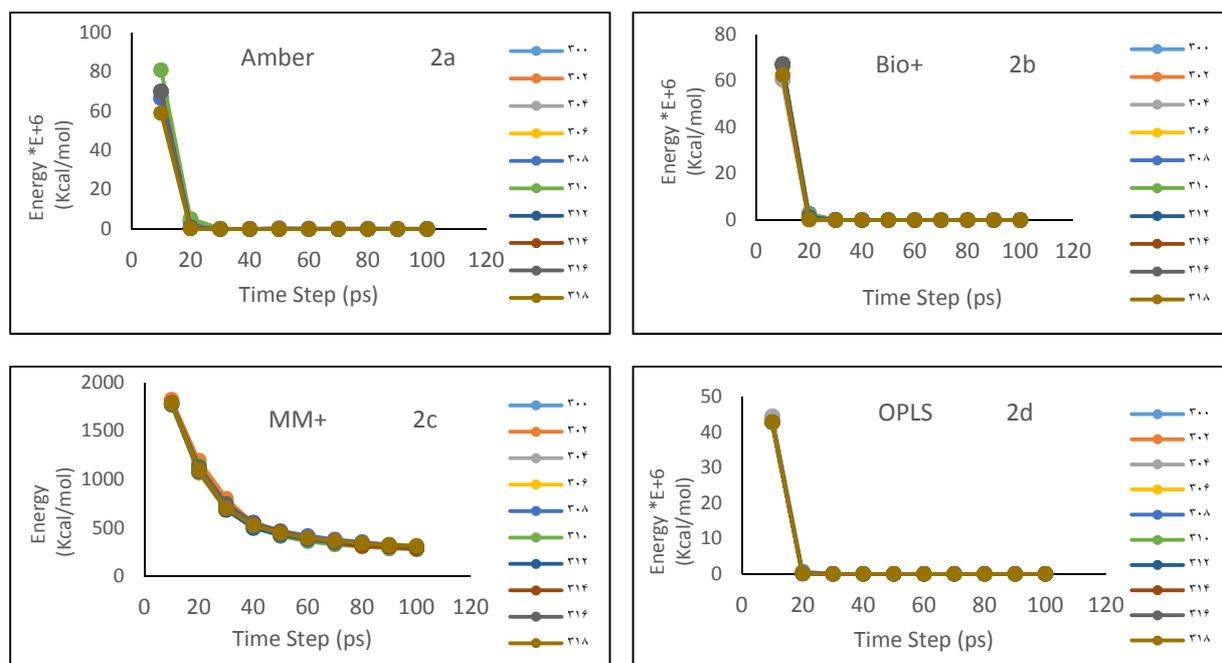

Fig. 2. The graphs of vincristine potential energy belong to a) AMBER b) BIO⁺, c) MM⁺, d) OPLS force fields.

Table 5. Computed total energy (kcal/ mol) for vinblastine, belong to AMBER, MM⁺, BIO⁺ and OPLS force fields under ten different temperature

Method	Total Energy (Kcal/mol)									
	300	302	304	306	308	310	312	314	316	318
AMBER	1283.59	2271.41	2307.07	2228.64	1817.02	2288.61	1718.96	2322.42	1527.61	1770.11
OPLS	2164.13	2136.51	1893.53	2272.42	2526.39	1877.34	2032.83	2142.88	2135.64	2111.96
MM ⁺	630.38	628.254	650.87	644.359	702.702	756.339	655.522	648.057	707.722	667.406
BIO ⁺	1792.13	2495.95	1744.98	1844.41	1259.92	1430.14	1967.26	1445.2	1483.22	1613.85

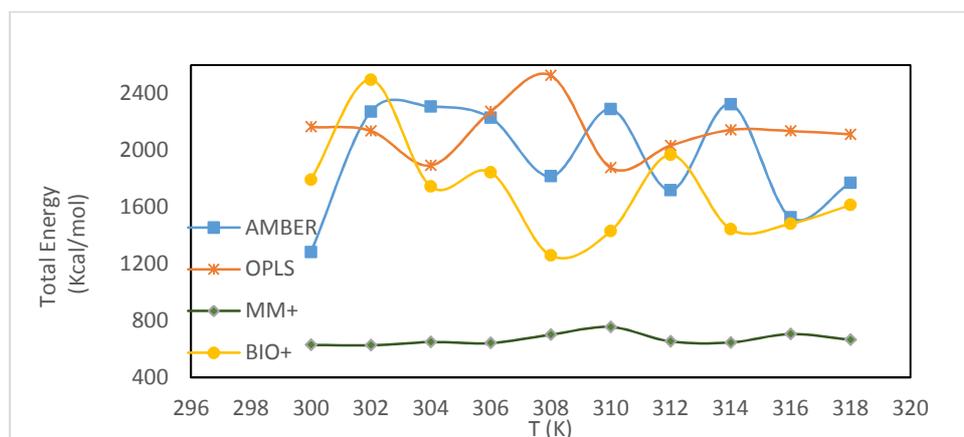


Fig. 3. The graphs of vinblastine Total Energy.

Table 6. Computed total energy (kcal/ mol) for vincristine, belong to AMBER, MM⁺, BIO⁺ and OPLS force fields under ten different temperature

Method	Total Energy (Kcal/mol)									
	300	302	304	306	308	310	312	314	316	318
AMBER	489.792	483.865	504.532	482.932	483.375	534.712	502.087	525.551	537.601	505.981
OPLS	452.847	471.442	445.115	469.3	430.017	436.682	445.061	460.554	454.705	435.433
MM ⁺	416.127	404.437	394.893	401.57	414.904	381.269	411.173	387.991	420.581	425.974
BIO ⁺	545.059	547.836	550.403	570.447	552.624	555.436	517.037	572.967	562.428	570.926



Fig. 4. The graphs of vincristine Total Energy.

CONCLUSIONS

Performance of the Molecular Mechanic investigation of vincristine and vinblastine gave the potential energy and total energy by Monte Carlo method in different temperature. The study showed that the system has the different level of energy and the different stability which is caused

by the forces from inside the system. The best spatial conformity which means the highest stability level or the lowest level of energy was found. Also, it is obvious from the above diagrams that maximum amount of potential energy is related to 312K, OPLS method for vinblastine and 310K,

AMBER method for vincristine. The highest level of total energy observed in OPLS method for vinblastine and BIO⁺ for vincristine. So with considering high amount of total energy, there will be minimum stability in this method.

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