A COMPARATIVE STUDY OF STEROCHEMICAL EFFECTS OF ANTI-PROSTATE AGENTS BY MOLECULAR DOCKING

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ABSTRACT

Objective: A comparative study of anti-prostate agents to investigate the stereochemical influences on binding affinity by molecular docking.

Methods: Structures of enantiomers (R and S stereoisomers) for known anti-prostate cancer (PCa) agents were drawn using ChemBioDraw 2D software. Thereafter, they were converted to 3D structures using the ChemBioDraw 3D software in which they were subjected to energy minimization using the MM2 method and then saved as PDB extension files which can be accessed using the ADT interface. AutoDock Vina (ADT) 1.5.6 software version was used for molecular docking study.

Results: A total of 12 different anti-Pca agents were selected and drawn including well-known drug R-bicalutamide. All molecules showed the binding affinity with respect to the nature of stereochmeistry. R-stereoisomers showed better interaction as well as binding affinity toward 1z95 (mutated androgen receptor protein involved in the progression of PCa) whereas their S-stereoisomers were found inferior in comparison.

Conclusion: This study showed that CB1-R and R-bicalutamide (with R-stereochemistry) were better in binding affinity comparative to their counterpart CB1-S and S-Bicalutamide (with S-stereochemistry). All the selected anti-Pca agents were showing the effect of stereocenreal center; therefore, we must choose the right kind of stereochmeistry while planning to develop the newer anti-Pcagents.

Keywords: Stereisomer, Molecular docking, Anti-prostate agent, AutoDock Vina, Androgen.

INTRODUCTION

Prostate cancer (PCa) has been described as a disease that affects the prostate region of the male genital organ due to the proliferation of androgen receptors and hormones present in the prostate organ [1]. Research shows in the preventive medicine reports that PCa continues to disproportionately affect African-American men in terms of mobility and mortality [2]. Furthermore, various examination studies, knowledge, beliefs, and behaviors on PCa screening [3] and recent researches also evaluate the effect of interventions that focus on enhancing the screening and about decision-making among African-American men [4,5].

As described and illustrated earlier, PCa is a leading cause of death in men. Globally, it is the most diagnosed/screened among men AutoDock Vina between the ages of 65 and 74 years [2]. The prediction or diagnosis differs greatly, but highly dependent on a number of factors such as stage of diagnosis, race, and age. Currently, the treatment of PCa includes androgen deprivation, surgery, radiation, endocrine therapy, radical prostatectomy, and chemotherapy [6]. Drugs like flutamide (Bulexin), hydroxyflutamide (Active form of Flutamide), bicalutamide (Casodex), nilutamide, and enzalutamide (formerly called MDV3100) (Xtandi) are all non-steroidal androgen receptor antagonists which were approved for the treatment of PCa [7].

Recent years, stereochemistry has gained great attention in pharmaceutical industries and regulatory authorities [8]. Stereochemistry is mostly dealt with compounds which possess chiral centers. Chiral compounds which are not superimposable in their mirror images are known as enantiomers and have opposite rotation of the direction of plane polarized light. R and S stereoisomers are the enantiomers represent the absolute stereochemistry of stereocenter in 3D structure. These enantiomers behave differently in interaction with chiral environments such as enzymes, receptors, and transporters. Generally, they differ in chemical and pharmacological properties such as bioavailability, the rate of metabolism, metabolites, excretion, potency, and selectivity for receptors, enzymes, and transporters [9]. 50% of marketed drugs are chiral and out of that 50% are available as racemate (an equal mixture of both enantiomers) mixtures [10]. The most important thing here is that these two enantiomers may have same pharmacological action, for example, ofloxacin (R-enantiomer) and levofloxacin (S-enantiomer) which are both antimicrobials of quinolones [11]. One enantiomer may be active and the other may be inactive, the examples are (S)-albuterol, a β2-adrenergic receptor agonist for treatment of asthma, and (S)-omeprazole, a proton pump inhibitor for treatment of gastroesophageal reflux has been shown to be superior to their racemate formulations in clinical trials cause their R-enantiomers are inactive [9]. One of them may be having some undesirable effects, like S-timolol, a β-blocker agent can cause severe bronchoconstriction, while R-isomer did not cause, due to this it is marketed as pure R-enantiomer [12]. Hence, to achieve better results in therapy and prevent harmful effects of unwanted enantiomer, the availability and isolation of a single enantiomer have become an important issue in pharmaceuticals, nowadays.

METHODS

Computer-aided drug design, especially molecular docking, is a very useful method to find out the potent molecule without investing extreme effort along with funds in research [13-17]. ADT 1.5.6 software [18] is
used by us to examine the activity in terms of binding affinity (Kcal/mol), and thereafter, the outcomes were compared in binding affinity score for best-docked conformation. The structures of various known anti-PCa agents were drawn with their stereochemistry with the help of ChemBioDraw ultra and further converted to the 3D structure using ChemBioDraw 3D. All the designed structures were optimized by energy minimization using MM2 method [19] and converted to readable format at the ADT interface. For a comparative study of stereoisomers of the anti-PCa agents, a protein 1z95 was selected and downloaded from protein data bank [20]. The outcomes of results were analyzed by AutoDock Vina (ADT) result which reveals close contact, hydrogen bond, hydrophilic, and hydrophobic interactions.

RESULTS AND DISCUSSION

The main purpose of this study is to find out the stereochemical effect of anti-PCa agents which are used the treatment of PCa or recently investigated. The molecules were selected from the available literature having a different class of compounds, that is, bicalutamide [21], propionamides [22], and thioxoimidazolidinones [23] as shown in Fig. 1.

All structures were drawn through ChemBioDraw 2D software (both R and S or RR and SS) to study the stereochemical effect and then converted into 3D structures using ChemBioDraw 3D software and later saved as pdb files so that it can be accessed by the ADT software. Androgen protein receptor was downloaded from the protein data bank site rcsb.org specifically protein with code 1z95 for this study.

All the selected molecules were analyzed using the ADT molecular modeling software. Initially, the protein validation has been done by extraction of ligand (Fig. 2). Preparations of ligands were done by the addition of polar hydrogens, detecting root, and converting it to pdbqt extension file, while 1z95 protein was prepared by removing water molecules, repairing missing atoms, adding polar hydrogens only, and subsequently adding the Kollman charges. Further, the grid box was generated keeping the ligand as a center (Fig. 3). From grid output file, the configuration file "conf.txt" was prepared and command prompt was used for ADT molecular docking by giving the command "program files\the scripps research institute\vina\vina.exe --config conf.txt --log log.txt." It generated the output file with the docking score or binding affinity (Kcal/mol). Similarly, all twelve molecules were studied and their binding affinities are shown in Table 1.

The study was based on twelve different molecules containing one or more stereocenter. Binding affinities represented in Table 1 revealed that stereochemistry played a critical role and considered for better interaction with the 1z95 protein. This was observed while comparison of the binding affinities of all R-stereoisomers with their S-stereoisomers. To know the effect of stereochemistry and various substituents, all the interactions of protein 1z95 were analyzed in detail. Different amino acid residues are found to be involved in the interactions at binding sites include ILE899, ASN705, LEU704, THR877, ARG752, MET745, MET895, GLN711, and LEU741. The aforementioned amino acids can be classified on the basis of their polarity, hydrophobicity, and charge such as ILE899, LEU704, MET745, MET895, GLN711, and LEU741. The aforementioned amino acids can be classified on the basis of their polarity, hydrophobicity, and charge such as ILE899, LEU704, MET745, MET895, and LEU741 are hydrophobic in nature, while ASN705, THR877, and GLN711 are polar amino acids present in the protein receptor 1z95, therefore, represent similar kind of interaction with the molecule of interest.

Both R or RR and S or SS were studied for their interactions with the 1z95 protein. These results suggested the essence of stereocenter on the binding affinity of the existing anti-PCa agents. In Table 1, all propanamide and thioxoimidazolidinone derivatives having R and RR stereocenter showed more binding affinity than the S and SS stereocenter ligands. Among them, molecules with higher binding affinity were discussed herein in detail. Molecule CA1-R (R-bicalutamide) and CA1-S (S-bicalutamide) were found to be with

-11.3 Kcal/mol and -10.3 Kcal/mol binding affinity, respectively, to 1z95 protein.
Fig 4: Interactions between ligand CA1-R (R-stereoisomer) and protein 1z95

Fig 5: Interactions between ligand CA1-S (S-stereoisomer) and protein 1z95

Fig 6: (a) Ligand CA1-R (R-stereoisomer) and (b) CA1-S (S-stereoisomer) with ribbon structure of protein 1z95
Table 1: Selected ligands with affinity

| S.No. | STRUCTURES (CODE) | Binding affinity (Kcal/mol) |
|-------|-------------------|----------------------------|
| 1     | R-Bicalutamide (CA1-R) | -11.3                     |
| 2     | S-Bicalutamide (CA1-S) | -10.3                     |
| 3     | (CB1-R)            | -10.2                     |
| 4     | (CB1-S)            | -9.8                      |
| 5     | (CB2-R)            | -9.6                      |
| 6     | (CB2-S)            | -9.6                      |
| 7     | (CC1-RR)           | -7.4                      |
| 8     | (CC1-SS)           | -9.1                      |
| 9     | (CC2-RR)           | -6.9                      |
| 10    | (CC2-SS)           | -8.2                      |
| 11    | (CC3-RR)           | -7.8                      |
| 12    | (CC3-SS)           |                           |

A detailed observation showed that CA1-R (R-bicalutamide) has favorable interaction with 1z95 androgen receptor (Fig. 4). The amine group of ARG752 of protein 1z95 has hydrogen bonding with the cyano group of CA1-R (R-bicalutamide), similar interaction showed by cyano group with MTH745 and GLU711. The carbonyl group of the ASN705 residue of the 1z95 is involved in a hydrogen bonding with the chiral hydroxyl group and sulphone group of CA1-R (R-bicalutamide). The interactions between the protein (1z95) and ligand CA1-S (S-bicalutamide) are less favorable (Figs. 5 and 6). Herein, also amine group of the ARG752 residue of the receptor protein is involved in hydrogen bonding with the cyano group of CA1-S (S-bicalutamide), while chiral hydroxyl group does not involve in hydrogen bonding resulting in low binding affinity. The overlay of CA1-S (S-bicalutamide) and CA1-R (R-bicalutamide) depicted in Fig. 7 showed the importance of chiral hydroxyl group which is opposite in case of CA1-S (S-bicalutamide), therefore, lacking hydrogen bonding and resulting low binding affinity.

CONCLUSION

The evolution of PCa occurs as a result of the overexpression of androgen receptor. There are several therapies available to treat PCa, but the most commonly used is chemotherapy include non-steroidal drugs. These drug molecules have a good involvement of hydroxyl group with particular stereochemistry. Thus, the present study is undertaken to investigate the role of the stereochemistry of bicalutamide, propanamide, and thioxoimidazolidinones as anti-PCa agents.

ADT molecular software was used for molecular docking of all stereoisomers on 1z95 (Androgen receptor) protein. Binding affinities revealed that molecule with score shows that R-stereochemistry is very important at the binding site resulting in higher binding affinity than its S-stereoisomer. Therefore, it could be suggested that while designing the novel anti-PCa agents, we must consider the stereochemistry at the stereocenter.

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CONFLICTS OF INTEREST

The author’s declared that they have no conflicts of interest.

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Fig. 7: Overlay of ligands CA1-R (R-stereoisomer) and CA1-S (S-stereoisomer) with their binding affinity
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