

Original Review Article

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CONOTOXINS: AN EMERGING DRUG-BASED SOURCE AGAINST VARIOUS CANCERS AND LUNG DISEASES

Manisha Nag^{1,2}, Sweta Rani Chaurasia^{1,3}, Sana Tasneem^{1,4}, Pramod Kumar^{1,5}, Subhashini Singh Thakur^{1,5}, Sonia Choubey^{1,6}, Annie Jessica Toppo¹, Priyangulta Beck¹, Ganesh Chandr Baskey², Mukesh Nitin^{1*}

1. Department of Tech Biosciences, Digianalix, South Samaj Street Tharphakna Ranchi, -834001, Jharkhand, India.
2. Department of Zoology, Dr. Shyama Prasad Mukherjee University, Ranchi, -834008, Jharkhand, India.
3. Department of Biotechnology, M.S. Ramaiah University of Applied Sciences, Bengaluru - 560054, Karnataka, India.
4. Department of Zoology, Ranchi Women's College Ranchi, -834001, Jharkhand, India.
5. Department of Biotechnology, Marwari College Ranchi, -834008, Jharkhand, India.
6. Department of Botany (Biotechnology), Ranchi University, Ranchi, -834008, Jharkhand, India.

ABSTRACT: Conotoxins are proteinaceous compounds extracted from cone snail venom and have pharmacological effects against a variety of illnesses, including several types of cancer (including breast, lung, pancreatic cancer etc.) and numerous lung ailments (including asthma and tuberculosis). The property of conotoxins inhibit pro inflammatory conditions and tumor suppression thereby providing us a new idea of a potent drug target component for treating the diseases mentioned above. Based on the genomics literature review, selective proteins and biological compounds like conotoxins played very important role in curing variable diseases like cancer, asthma, tuberculosis etc, which was predicted through advanced computational biology and docking analysis.

Keywords: Conotoxin, Conus, Cancer, Lung diseases, Molecular docking.

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Corresponding Author: Dr. Mukesh Nitin* Ph.D.

Scientist and Head, Dept. of Tech Biosciences, Digianalix, INDIA

Genomic Scientific Consultant, Gentan, Izmir, TURKEY

Email- digianalix@gmail.com

1. INTRODUCTION

Cancer is a life-threatening and devastating disease that affects millions of people worldwide. In 2020, the world health organization (WHO) anticipated that there will be more than 13 million deaths and 21 million new cases of cancer by 2030. Despite significant advances in research and treatment, cancer still accounts for a major cause of morbidity and fatality. As a result, there is a critical need to develop new methods and approaches to cancer therapy that can improve outcomes for patients [1-4]. Cancer is defined as the uncontrolled growth and division of abnormal cells in the body due to some fault in the machinery of the cell cycle that controls cell proliferation [5,6]. Chemotherapy is one of the existing and effective methods to treat various types of cancerous cells. Although chemotherapy is quite effective in the treatment of cancer, it has many side effects and limited efficacy, especially in the advanced stages of the disease. Additionally, chemotherapy drugs often target both cancerous and healthy cells, leading to significant toxicity and adverse effects [7-9]. The potential utilization of natural products, particularly substances originating from the marine life, as a source of fresh cancer treatments has recently emerged as one of the most important fields of research. [10]. In particular, marine gastropods (conus snails) belonging to the Conidae family [11], contains a venom gland that secretes neurotoxin generally referred to as conopeptide or conotoxins, have emerged as a promising class of natural product for cancer therapy [12,13]. Conotoxins are classified into different families depending on the types of their molecular target and corresponding pharmacological activity [14,15]. Conotoxin has a wide range of structural and functional diversity and mainly targets membrane protein receptors, especially ion channel [15,16]. Membrane ion channels have a significant role in cell proliferation and play an important role in the development of cancer [17-25]. Ultimately, proliferation studies show that inhibition of K⁺ channel expression or channel blockade by any specific inhibitor reduces cell proliferation. Thus, selectively targeting and blocking ion channels would be a significant therapeutic approach for cancer therapy. Hence in this review, we will focus on the development of anticancer drugs using conotoxins for cancer therapy, along with the mechanism of action of conopeptide and the opportunities & obstacles involved in creating conotoxins-based cancer therapies. By analyzing and synthesizing the available data, we aim to provide a comprehensive and up-to-date perspective on the potential of conotoxins as cancer therapeutics and to identify key

areas for future research and development in this exciting and rapidly evolving field.

2. APPLICATIONS AND USAGE OF MARINE PRODUCTS

Marine habitats are an excellent source of bio-active natural compounds due to their unique and diverse chemical structures [26]. Biomedical exploration of marine natural compounds that exhibit various pharmacological activities, including anticancer, antimicrobial, antiviral, anti-inflammatory, and analgesic properties [27]. Applications of these compounds have shown potential for the treatment of various diseases, including cancer, infection, and inflammation [28,29]. For anti-cancer treatment, two marine natural products, ziconotide and brentuximab vedotin, have been commercialized over the past few years. Also, 3 out of 4 novel medications produced for the treatment of cancer have been derived from both marine and terrestrial natural products [30]. In 2011, another marine product synthesized as a drug (Adcetris) developed from sea hare for the treatment of cancer. Further, several other marine-derived products are under different phases of clinical trials in various countries for significant cancer treatment such as plitidepsin glembatumumab vedotin [31]. Toxins secreted from animals are peptides that interact with specific target molecules such as ion channels (e.g., sodium and/or potassium ion channels), thus affecting the neuromuscular, cardiovascular and immune systems [32,33].

3. CONE SNAILS (CONUS)

Cone snails are invertebrates belonging to the phylum Mollusca, class gastropods and genus conus, these marine gastropods consist of 700 species from the genus conus [34]. These are found in various marine habitats across the world including coral reefs and shallow sandy waters in Western Atlantic, Indian and Pacific oceans. However, they cannot survive in freshwater [35]. Predator cone snails have been a matter of growing interest due to their greatly evolved hunting approach that makes use of conotoxins to paralyze prey. Cone snails are usually slowly moving creatures surrounded by fast-moving prey, which presents a major existential challenge to these predators. However, they successfully deal with it and control it by developing a venomous apparatus, which is accountable for the synthesis, storage, and delivery of large amounts of the most sophisticated conotoxins peptides [36]. These toxins are effective in treating various diseases. Other factors include the fact that some toxins are extremely selective for particular receptors in the body and also marine snails are abundant & easily accessible source of venom toxins. They have a specialized radula tooth that is modified into a hollow, harpoon-like structure called a radula tooth. When the snail hunts for its prey, it extends this tooth and injects the venom into the prey. The venom of cone snails contains a complex mixture of toxins that can paralyze or kill their prey [37,38]. The venom gland is a modified salivary gland of marine cone snails that can secrete neurotoxic peptides in the large amount commonly known as conopeptide or conotoxins reached in disulphide bridges with pharmacological activities [39,40].

4. PROPERTIES OF CONOTOXIN

Conotoxins typically consist of 8- 35 amino acids and a high number of modifications such as being rich in disulphide bonds, hydroxylation and glycosylation. These modifications are responsible for the structural stability and specificity of conotoxins [41-44]. Cone snail venom conopeptide are encoded by gene superfamilies, and a single species consist of 100-400 venom peptides [45-47]. The nomenclature used for classifying conopeptides was proposed by Cruz et al. in 1985. It categorizes them into superfamilies based on sequence and framework homology, and then into pharmacological families based on the targets they interact with. Conopeptide with no disulfide comes under six groups, viz., the ntulakins (which target the neurotensin receptor), the conantokins (which target the N-methyl-D-aspartic acid receptor) [48], the conorfamides (thought to target the Rfamide receptor) [49], the conolysins (thought to target cellular membranes) [50], the conophans (target unknown) [51], and the conomarphins (target unknown) [52]. Peptides containing one disulfide bond are classified as either the contryphans (target undefined) [53] or the conopressins (vasopressin homologs) [54].

5. VARIANCE IN THE STRUCTURAL PROPERTIES

The disulfide-rich peptides are referred to as conotoxins and are classified into six different classes based on their structures. They are α conotoxins, ω conotoxins, δ conotoxins, μ conotoxins, κ conotoxins and contulakin[55].

(a) **α conotoxins**- These conotoxins have a compact globular structure stabilized by two disulfide bonds. They typically consist of 12- 19 amino acid residues [56,57]. α conotoxins are known to block nicotinic acetylcholine receptors in both neuronal and muscle types [58,59]. These peptides are used to treat various diseases, anxiety, Parkinson's disease, pain, hypertension, cancer, and also muscle relaxants [60].

(b) **ω conotoxins**- These conotoxins are typically 24-30 amino acid residues in length and contain three disulfide bonds. The first isolated ω conotoxins were GVIA from *Conus geographus* [61,62]. They block voltage-gated calcium channels in the nervous system [63,64]. ω conotoxins are the most selective inhibitor among the other conotoxins because of their therapeutic potential in the management of severe pain.

(c) **δ conotoxins**- These conotoxins are approximately 30 amino acid residues in length and contain three disulfide bonds [65]. They block voltage-gated sodium channels.

(d) **μ conotoxins**-These conotoxins have a compact globular structure stabilized by three disulfide bonds. They typically consist of 16-25 amino acid residues and are known to block voltage-gated sodium channels in the nervous system and caused paralyzing [65].

(e) **κ conotoxins**-The first κ conotoxin was PVIIA, isolated from the fish hunting cone *C. purpurascens*, it is a27 amino acid residue [66]. They block voltage-gated potassium channels.

6. MODULATION OF IMMUNE SYSTEM WITH RELATED TO CONOTOXIN

Conotoxins can highly regulate the immune system due to their therapeutic potential [67-71].

Conotoxins target ion channels that have vital roles in immune cells and immune-related diseases. One such ion channel immunomodulatory is Voltage-gated potassium channels that suppress immunomodulatory activation through blockade [72-76]. Voltage-gated potassium channels are involved in various immunological processes such as leukocyte, macrophage activation, lymphocytes, and proliferation [77,78]. K- conotoxins, PVIIA blocks the shaker potassium channels by binding to the specific binding site(triethanolamine) and this channel which is expressed in immuno-cells such as macrophages and T-lymphocytes, is a potent immunomodulatory target [79]. Other conotoxins are α -conotoxins, two PnIA and ImI α -conotoxins isolated from the venoms of *Conus pennaceus* and *Conus imperialis* respectively. PnIA block the molluscan neuronal acetylcholine receptors [80]. Later found that it was also an inhibitor of mammalian nAChRs. In the year 1994, another mammalian nAChR-specific conotoxin ImI was discovered [81]. ImI has IC₅₀ values of 220nM and 1800nM for the homomeric nAChRs consist two subunits $\alpha 7$ and $\alpha 9$ respectively [82]. $\alpha 7$ have a particular interest as a pharmacological target in the immune system. $\alpha 7$ nAChR involve in the production of antibody [83] and T-cell proliferation [84]. Literature suggests that anti-inflammatory effects were predominately produced by the activation of the $\alpha 7$ nAChR [85].

7. EFFECTS OF CONOTOXIN ON CANCER CELLS

The mechanism of action of conotoxins in cancer cells varies depending on the specific conotoxins and the type of cancer being targeted. Conotoxins in particular target the ion channel that communicates with cancer cells. One common mechanism by which conotoxins can affect cancer cells is by binding to and blocking ion channels that are overexpressed in cancer cells. The mechanism of conotoxins in cancer cells with nAChRs involves binding to a specific site on the receptor. The nicotinic acetylcholine receptors (nAChRs) are known to have a role in inflammation [86] and oncogenesis [87]. one of the frequently utilised techniques in the study of nAChRs are α conotoxins which specifically and efficiently inhibit different subtypes of nAChRs [88]. Several nAChR subunits, including $\alpha 3$, $\alpha 7$, $\alpha 9$ and $\beta 4$, are expressed in a variety of tumour cells and are involved in the control of cell proliferation, apoptosis, invasion, migration, and angiogenesis [89-92]. the α -conotoxin AuIB may prevent small-cell lung carcinomas (SCLC) cells from surviving by binding to nAChRs with the subunits $\alpha 3/\alpha 5/\beta 4$ [93]. In response to tumour microenvironments, $\alpha 9$ nAChRs play a crucial role in driving cancer cell proliferation, angiogenesis, cancer metastasis, and apoptosis suppression during carcinogenesis [94-96]. For examples, *Conus generalis* produced the α O-conotoxin GeXIVA, which effectively inhibits nAChRs $\alpha 9$ and $\alpha 10$ subunits [97]. In vivo, it also demonstrated significant neuropathic pain relief in one rat model [98,99]. One recent research also demonstrated that GeXIVA played a crucial role

in the prevention of the growth of cervical cancer cells [100].

8. INVESTIGATION

Few tumour causing genes were identified through literature review on different cancers conditions and a meta-analysis was carried out in three other lung disease conditions (TB, Asthama and Lung Cancer). The hub genes were identified from genomic data set of variable above mentioned lung diseases through NCBI GEO datasets-GSE54712, GSE43696, GSE20050. Further, effective significant differential expressed genes qualifying p -value < 0.05 were screened for hub genes studies through gene conservancy analysis. The selected and screened genes were further prepared as receptors, and then docked with the conotoxin variant taken (CONOTOXIN MII, CONOTOXIN GeXIVA, 3 alpha subunit CONOTOXIN). Analyzation of the effective ligand-receptor docked complex results given in (Table 1-2) and (Figure 1-2) shows that more effective docking score obtained is from conotoxin variant GeXIVA with receptor CSF1R is -287.13 and conotoxin variant GeXIVA with receptor NACHRs β 2 is -326.20. Moreover, MD simulations were carried out with best docked complex highlighting its eigenvalue of docked complex of conotoxin variant GeXIVA with receptor CSF1R is $1.715798e-05$ and conotoxin variant GeXIVA with receptor NACHRs β 2 is $1.231755e-05$ which shows better stability of docked complex as shown in (Figure 3-4).

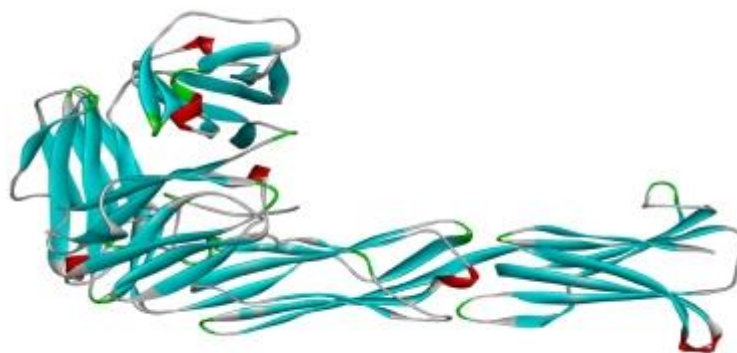


Figure 1: Docked complex Conotoxin GeXIVA with CSF1R receptor

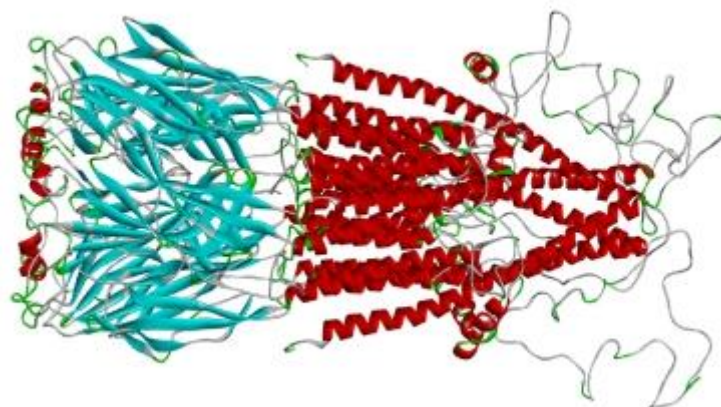


Figure 2: Docked complex Conotoxin GeXIVA with NACHRsβ2 receptor

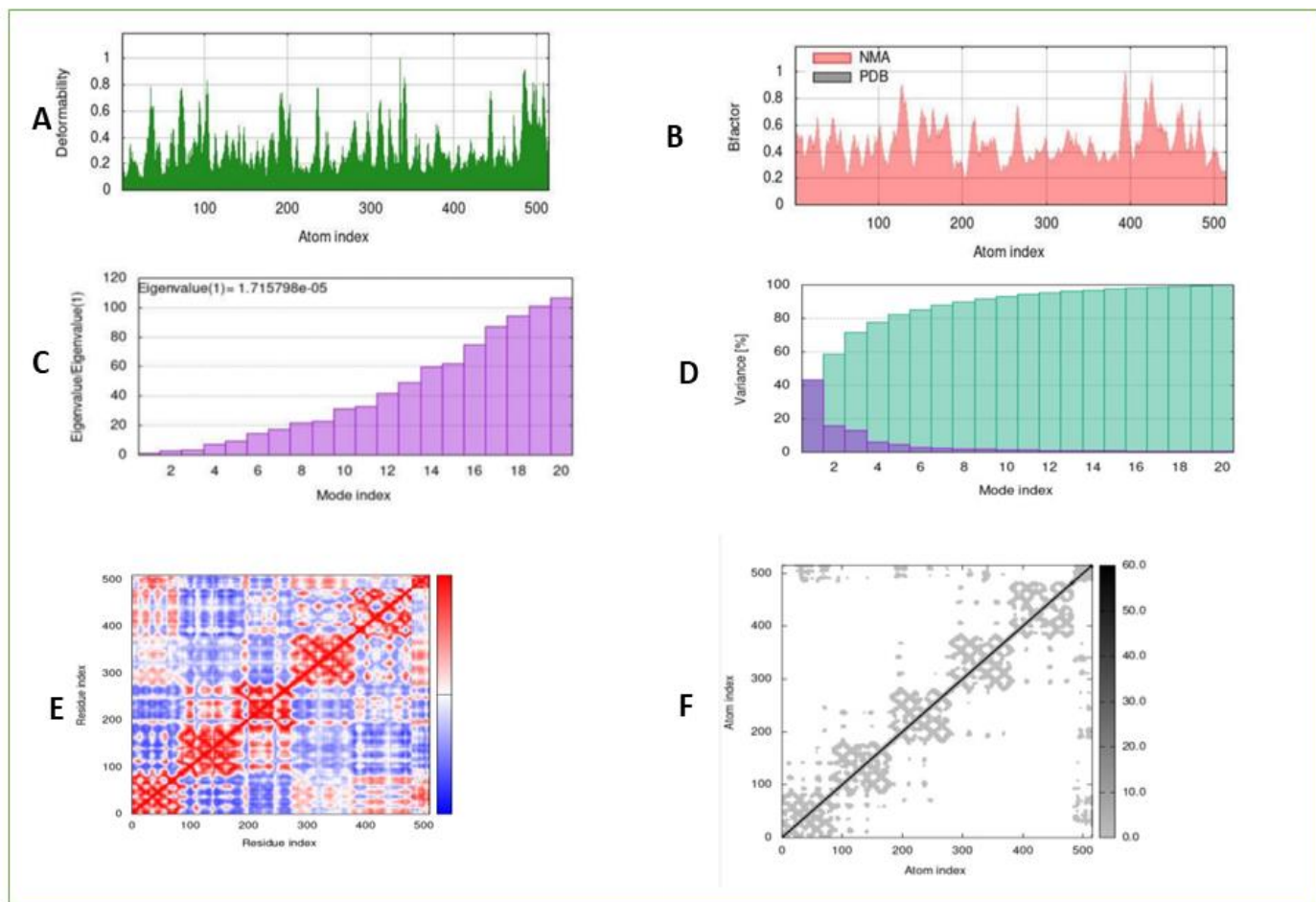


Figure 3. MD simulation result; A: deformability, B: b-factor, C: eigenvalue, D: variance, E: residue index, F: atom index of receptor CSF1R with ligand conotoxin GeXIVA

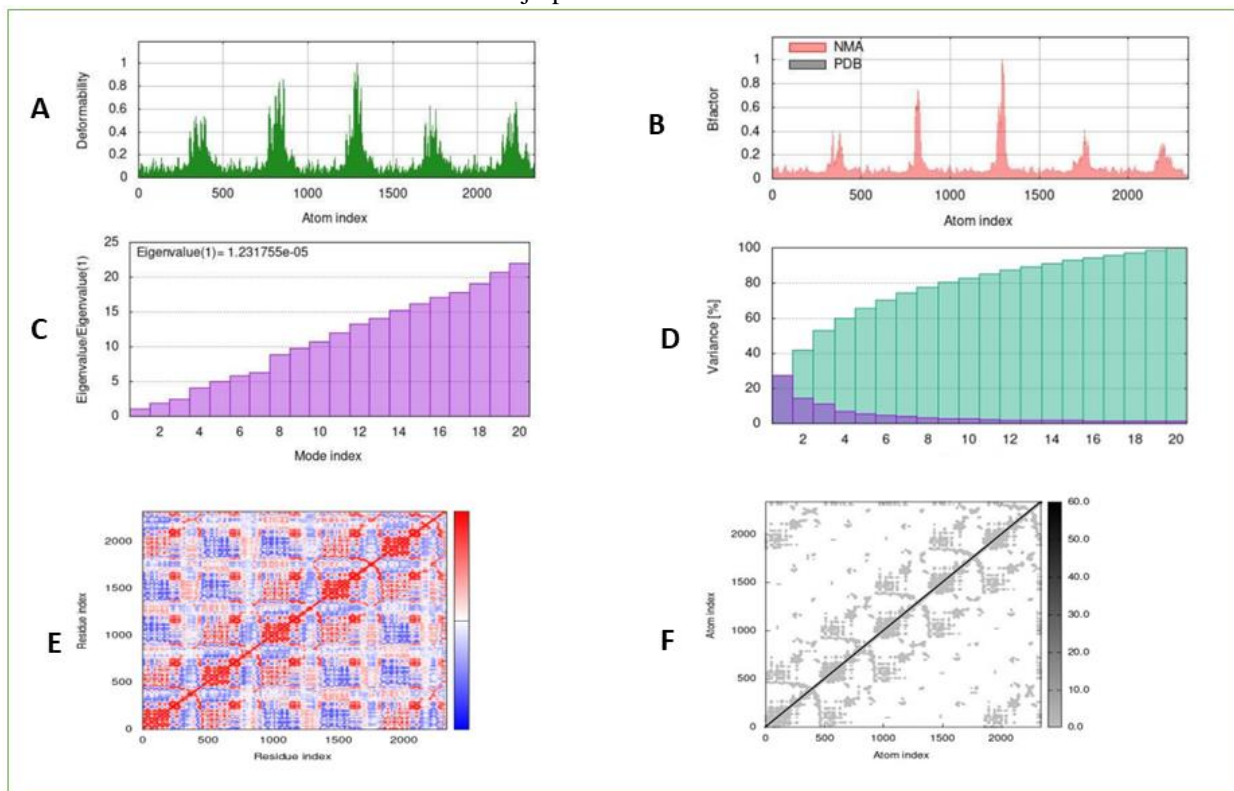


Figure 4. MD simulation result; A: deformability, B: b-factor, C: eigenvalue, D: variance, E: residue index, F: atom index of receptor NACHRs β 2 with ligand conotoxin GeXIVA

B-Factor: The experimental B-factor is taken from the corresponding PDB field and the calculated from NMA is obtained by multiplying the NMA mobility by $(8\pi^2)$. Be aware that many PDB files of averaged NMR models contain no B-factors (actually, the B-factor column gives an averaged RMS).

Eigenvalues: The eigenvalue associated to each normal mode represents the motion stiffness. Its value is directly related to the energy required to deform the structure. The lower the eigenvalue, the easier the deformation.

Variance: The variance associated to each normal mode is inversely related to the eigenvalue. Colored bars show the individual (red) and cumulative (green) variances.

Covariance map: Covariance matrix indicates coupling between pairs of residues, i.e. whether they experience correlated (red), uncorrelated (white) or anti-correlated (blue) motions.

Elastic network: The elastic network model defines which pairs of atoms are connected by springs. Each dot in the graph represents one spring between the corresponding pair of atoms. Dots are colored according to their stiffness, the darker grays indicate stiffer springs and vice versa.

9. DISCUSSION

The clinical investigation of these conotoxins has been hardly reported. So far, only a few studies have published the anticancer activity of conotoxins [101]. To date, several conotoxins have proceeded to the clinical research stage, including ω -MVIIA (chronic pain, marketed), ω -CVID (analgesia, phase IIa), contulakin-G (analgesia, phase I), conantokin-G (analgesia/anti-epileptic, phase Ib), χ MrIA (analgesia, phase IIa), α -Vc1.1 (analgesia, phase II, terminated), μ O-MrVIB (analgesia, phase II), etc. [102,103]. ω -conotoxin MVIIA (ziconotide is a synthetic version of the peptide) is a commercial conotoxin derived from the venom of *C. magus* to treat chronic pain in serious cancer and AIDS patients since its launch in 2004 approved by the U.S. Food and Drug Administration (FDA) [104,105]. Conotoxins are also rapidly undergoing development for the treatment of various health conditions like pain, Alzheimer's disease, Parkinson's disease, cardiac infarction, hypertension, and various other neurological diseases [106-108]. Conopeptide is a vast drug resource; cone snails can develop up to one million different bio-active compounds [109]. But less than 0.1% of these effective compounds have been structurally and functionally identified [110-112]. Hence, conotoxins can be a significant therapeutic approach for the development of drugs treating cancer disease.

10. CONCLUSION

In this study, we identified a special category of compound conotoxins, obtained from invertebrate sources like, molluscs which cover wide aspects of medicinal properties providing cure against various diseases with the use of computational biology and metanalysis. We successfully identified potential target receptors which plays vital role in various metabolic pathways for different kind of cancers, asthma and tuberculosis. Further, in our study we identified particular class of conotoxins (GeXIVA, MII, 3Alpha subunit) which can inhibit proliferation of cancerous cells and acts as tumor suppressors. Future studies on these variants can build a potential path for developing drugs against various diseases.

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ETHICS APPROVAL AND CONSENT TO PARTICIPATE

Not applicable.

HUMAN AND ANIMAL RIGHTS

No Animals/Humans were used for studies that are base of this research.

CONSENT FOR PUBLICATION

Not applicable.

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CONFLICT OF INTEREST

The authors have no conflict of interest.

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