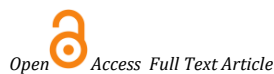
Available online on 15.11.2023 at <http://jddtonline.info>

Journal of Drug Delivery and Therapeutics

Open Access to Pharmaceutical and Medical Research

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Research Article

In-Vitro and *In-Silico* Approach Distinguish ER- α and HER-2 Antagonistic Properties of Indian Herbal Formulation on Breast Cancer

Munir Khan¹, Tapas K. Sur², Avijit Saha², Chandreyi Ghosh³, Tuhin K. Biswas⁴, Sirshendu Chatterjee³, Srikanta Pandit^{4*}

¹Health Reactive, Mumbai, India

²R.G. Kar Medical College & Hospital, Kolkata, India

³Techno India University, Kolkata, India

⁴J.B. Roy State Ayurvedic Medical College & Hospital, Kolkata, India

Article Info:



Article History:

Received 23 Aug 2023
Reviewed 02 Oct 2023
Accepted 23 Oct 2023
Published 15 Nov 2023

Cite this article as:

Khan M, Sur TK, Saha A, Ghosh C, Biswas TK, Chatterjee S, Pandit S, *In-Vitro* and *In-Silico* Approach Distinguish ER- α and HER-2 Antagonistic Properties of Indian Herbal Formulation on Breast Cancer, *Journal of Drug Delivery and Therapeutics*. 2023; 13(11):6-12

DOI: <http://dx.doi.org/10.22270/jddt.v13i11.6274>

*Address for Correspondence:

Dr. Srikanta Pandit, Associate Professor, J.B. Roy State Ayurvedic Medical College & Hospital, Kolkata, 170-172 Raja Dinendra Street, Kolkata, India, Pin-700004

Abstract

Objectives: The anticancer effect of an Indian herbal preparation was studied under a cancer cell line, as well as the *in silico* computational methods that explain the probability of protein ligands binding to ER- α and HER-2 receptors.

Method: The *in vitro* anticancer activity of Body Revival® suspension (BR) was determined using cytotoxicity tests, cell invasion and migration assays, and metastatic protein expression assays using MCF-7 breast cancer cells. The computational predictive biological method was applied to find out the pharmacodynamic and pharmacokinetic interactions between the active molecules present in the BR and ER- α /and HER-2 of breast cancer.

Results: BR showed significant and dose dependent cytotoxic effects on MCF-7 cells. The 50% effective cytotoxic dose of BR was 34.27 μ l/ml. It restricted invasion (26%) and migration (28%) of cancer cells than BSA control. MMP-9 and IL-6 concentration were reduced significantly ($p < 0.001$) after treatment. Cucurbitacin B had maximum *in silico* binding energy score (-7.8) with ER- α , while symconoside B had with HER-2 (-8.4); but, among the other interactions between the two ligands and receptors, withaferin A had the highest affinity (-15.3). Additionally, withaferin A, symconoside A, and symconoside B curcubitacin A demonstrated bioavailability and fulfilled safety standards.

Conclusion: Body Revival® showed as a powerful multi-target inhibitor of ER- α and HER-2 that has prospective anticancer action without side effects, and may be useful in the therapy management following a successful trial in breast cancer patients.

Keywords: Breast cancer, Cytotoxicity, MCF-7, ER- α , HER-2, Herbs, *In Silico*

INTRODUCTION

Breast cancer is the most common cancer diagnosed worldwide, with an estimated 2.3 million new cases in 2020 alone. Due to the long-standing predominance of risk factors related to reproduction, hormones, and lifestyle in North America, Canada, and Western Europe, incidence rates in these countries have been higher¹⁻². However, Asian nations like China, Japan, and India have also seen an increase in the prevalence of breast cancer³. Recent data suggest, 1 in 9 Indian female has a lifelong risk of developing breast cancer⁴.

Breast cancer is a genetically and clinically heterogeneous disease. Due to a lack of diagnostic markers, it has been difficult to measure the progression of this disease⁵. Breast cancers were categorized using conventional tumor classification such as fibroepithelial, myoepithelial, and mesenchymal neoplasms⁶⁻⁷. Human epidermal growth factor receptor-2 (HER-2) and hormone receptors (ER and PR) are the most relevant clinical markers that are now frequently employed in stratifying the disease⁸. HER-2 is positive in 1 in 5

individuals with breast cancer, while ER- α is positive in 1 in 3 cases⁹. On that molecular classes, this disease has now recognized as estrogen receptor positive (ER⁺), HER-2-positive (HER2⁺), triple-negative (TN), and unclassified¹⁰. Chemotherapeutics are used for its treatment, which either acts on one or more of these up- or down-regulating receptor signaling pathways to confront the deadly disease, but they are accompanied by serious side effects, including emesis, anorexia, diarrhea, skin rashes, hot flashes, headaches, fever, exhaustion, and hair loss¹¹⁻¹³. Numerous herbal medications are used in conjunction with chemotherapy/or radiation therapy to overcome these challenges and increase the effectiveness of cancer treatment while minimizing side effects and consequences¹⁴⁻¹⁶.

Body Revival® (BR), a polyherbal suspension has been developed (M/s Health Reactive, Mumbai) to treat cancers. Each 5 ml BR contained dry water extract of *Aegle marmelos* fruit pulp (150 mg), *Acorus calamus* rhizome (175 mg), *Rubia cordifolia* root (200 mg), *Symplocos racemosa* stem bark (95

mg), *Withania somnifera* root (325 mg), *Blumea lacera* fruit (115 mg), *Rumex vesicarius* whole plant (240 mg), *Cucumis melo* seed (200 mg), and honey (qs). These substances, together with their active constituents, have anticancer characteristics, including β -asarone, cucurbitacin B, methylglyoxal, quercetin, symconoside A, symconoside B and withaferin¹⁷. In addition, it has a substantial quantity of polyphenols such gallic acid, p-coumaric acid, quercetin, and apigenin¹⁸. BR was reported to prevent myocardial infarction and decrease vascular platelet aggregation in animal models¹⁹. Furthermore, it enhanced cellularity in bone marrow and leukocyte, granulocyte, and lymphocyte counts in peripheral blood of immunosuppressive animals¹⁸. Recent study suggested BR improved quality of life, especially in the psychological and physical spheres of daily living of breast cancer patients²⁰. However, at this time, its mode of action in cancer environments is a matter of debate. Hence, extensive research was carried out using *in vitro* and *in silico* computational methodologies to ascertain the potential effect of BR on breast cancer cells and the underlying protein-ligands binding interaction probabilities with ER- and HER-2 receptors.

MATERIAL AND METHODS

In vitro anticancer activities

The following verified standard procedures for screening cancer drugs were employed.

Cytotoxicity Test

MCF-7 cells were maintained in Dulbecco's modified eagle medium (DMEM) containing 10% FBS and incubated at 37°C in CO₂ incubator (Thermo Fisher, USA). Streptomycin and penicillin (100 µg/ml) was used to avoid any contamination. Approximately, 1×10⁴ cells were grown separately in 96 well plates and treated with varying concentrations of BR (*i.e.*, 0, 6.25, 12.5, 25, 50, 100 µl/ml) at 37°C for 24h. In the following day, post-treated cells were washed with PBS, and incubated with MTT (1 mg/ml stock) at 37°C for 4h. The absorbance was measured at 570 nm using a micro-plate reader (Sinothinker, China). All tests were done in triplicate. The 50% cytotoxic concentration of the test compound was identified for treated cell line²¹.

Cell Invasion and Migration Assay

1×10⁴ MCF-7 cells were cultured in 24 well plates at 37°C in CO₂ incubator for 24h. The post-treated cells were then further incubated for another 24h after constructing scratches on the monolayer cells, and the migration was observed using an inverted microscope (Zeiss, Germany). In invasion assay, 1×10⁴ MCF-7 cells were seeded in Transwell chamber (ECM 555, Sigma, USA) for overnight at 37°C in a CO₂ incubator. Non-invaded cells were stained with Trypan blue dye (Sigma, USA) and washed sequentially to remove death cells. The invasive cells in the matrix were observed under an inverted microscope and the corresponding intensities were measured using a micro-plate reader at 450 nm²².

Metastatic Protein Expression Assay

Briefly, 1×10³ MCF-7 cells were incubated with BSA vehicle or BR for 24 h at 37°C. Thereafter, relative protein expression analysis was carried out using matrix metalloproteinase-9 (MMP-9) and inflammatory cytokine, interleukin-6 (IL-6) specific ELISA kits (RayBio, USA) following the supplier's protocol. The micro-plate was read at 450 nm.

In Silico Molecular Docking

The computational predictive biological method was applied to find out the pharmacodynamic and pharmacokinetic

interactions between the molecules present in BR and breast cancer receptors.

(i) Receptors preparation: The crystal structures of ER- α and HER-2 were downloaded from the Protein data bank (<https://www.rcsb.org/>). The structure of the protein was validated using SAVES 6.0 server. Energy minimization was done using SPDBV software. The structural quality of the target protein was determined using PROCHECK server.

(ii) Determination of active sites: The presence of amino acids in the active site was determined by the CASTp web server²³.

(iii) Ligand preparation: The structure data format of the selected 10 bioactive compounds of BR (apigenin, cucurbitacin B, gallic acid, methylglyoxal, p-coumaric acid, quercetin, symconoside A, symconoside B, withaferin and β -asarone) was retrieved from the PubChem database (www.pubchem.ncbi.nlm.nih.gov). Gasteiger charges (polar hydrogen charges) were drafted and non-polar hydrogen molecules were combined with carbons. The protein and ligand were converted to PDBQT format using Autodock 4.2 tools.

(iv) Molecular docking: The docking of all 10 compounds was done into a 3D X-ray structure by Autodock 4.2 and AutodockVina. This is a fruitful automated method to investigate the binding of macromolecule and ligands. With the help of Autodock tools, Gasteiger charges and hydrogen atoms were added to the protein and for simulation AutodockVina was used. The algorithm that AutodockVina uses is the Broyden-Fletcher-Goldfarb-Shanno algorithm that improves the accuracy of docking and prediction of the binding mode. Finally, the binding complexes were visualized by Bovia Discovery Studio Visualizer²⁴.

Pharmacokinetic, toxicity and safety studies:

Absorption, Distribution, Metabolism, and Excretion (ADME)

ADME were measured at the SwissADME website (<https://www.swissadme.ch>). The following parameters such as aqueous solubility (LogS), skin permeation (Log kp), bioavailability Score, human intestinal absorption, blood-brain barrier and CYP2C9 inhibitors were measured and compared.

Toxicity and Safety Prediction

The tolerance capacity of animal models as well as human before application and ingestion are important. To predict the toxicity level, an online server named pkCSM (<http://biosig.unimelb.edu.au/pkcsml/>) was used where the structure can be drawn otherwise input the SMILES which were downloaded from Drugbank, Pubchem or Zinc15 database. The pkCSM allows the study of toxicological effects by analyzing AMES toxicity, oral rat chronic and acute toxicity and maximum tolerated dose for human.

Statistical Analysis

The research results were input in the electronic data-sheet for statistical analysis using SPSS version 20 (IBM, Chicago, USA). Categorical variables were presented as percentages. All data were presented as mean and standard deviation (SD) and statistically analyzed by t-test. P-value ≤ 0.05 was considered significant.

RESULTS

The results of cytotoxic effects of different doses of BR in MCF-7 cells were presented in Fig. 1A. The 50% cytotoxic dose of BR was noted 34.27 µl/ml ($y = -0.964x + 83.04$; $r^2 = 0.825$). In transwell cell membrane, BR exhibited 34% invasion (34 ± 8.2)

by low dose (10 μ l/ml) and 26% by high dose (10 μ l/ml) than BSA control (Fig. 1B). Moreover, BR showed 42% (42 \pm 9.44) migration at low dose and 28% (28 \pm 10.23) at high dose, compared to 70% (70 \pm 8.25) migration in the BSA control (Fig. 1C).

MMP-9 or metastatic matrix protein expression of MCF-7 cells showed dose dependent inhibition after BR treatment (Fig.

1D). BR at the dose of 10 μ l/ml inhibited 36% (56 \pm 5.28 pg/ml) and 52% (42 \pm 9.14) of MMP-9 respectively than BSA control (88 \pm 3.24). BR demonstrated dose dependent down regulation of IL-6 concentrations in the MCF-7 cells (Fig. 1E). At the dose of 10 μ l/ml BR reduced 32.9% (63 \pm 8.3) IL-6 and at the dose of 20 μ l/ml (48 \pm 7.6) diminished 48.9% compared to BSA control (94 \pm 5.8).

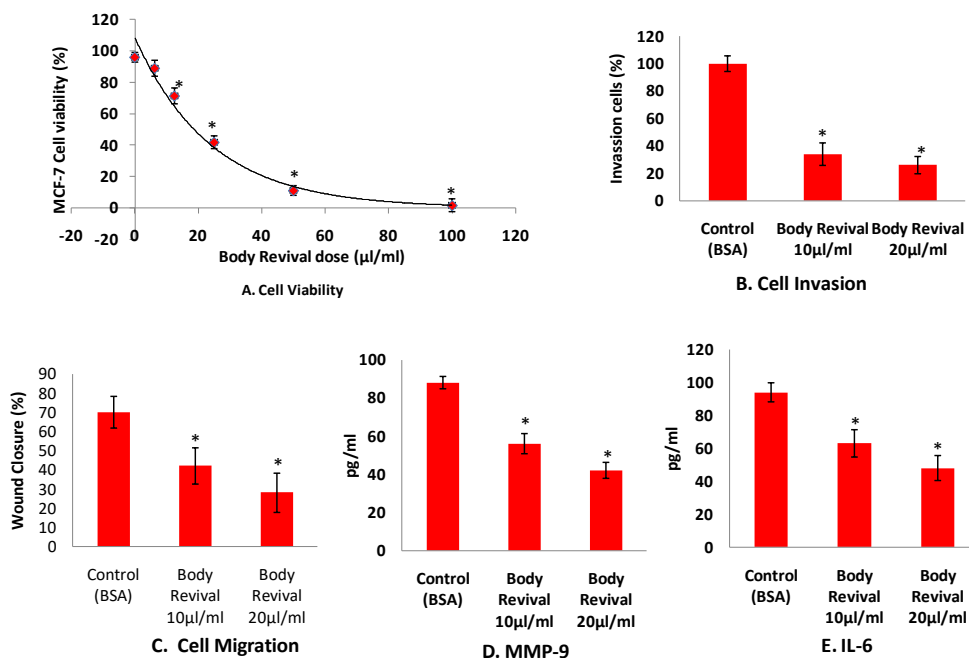


Figure 1: *In vitro* anti-cancer effect of BR on MCF-7 cells

The result of *in silico* molecular docking analysis of protein-ligands binding was described in Table 1. The binding energy score of cucurbitacin B (-7.8) was the maximum, followed by withaferin (-7.7), symconoside A (-7.2), symconoside B (-6.8) and quercetin (-6.3) with ER- α indicating their strong therapeutic inhibitory properties with the protein ligands binding. Furthermore, the binding energy score of

symconoside B (-8.4) was the highest, followed by symconoside A (-7.9), withaferin A (-7.6), quercetin (-7.5), cucurbitacin B (-6.9), apigenin (-6.9) and gallic acid (-6.4) with HER-2 also. With regard to interactions between ligands and receptors (ER- α and HER-2), withaferin A had the highest affinity (-15.3).

Table 1: *In silico* protein-ligands binding energy of active components of BR

Active Components	Protein - Ligands binding energy (kcal/mol)		
	ER- α (5J1H)	HER2 (5T92)	Total
Apigenin	-5.5	-6.9	-12.4
Cucurbitacin B	-7.8	-6.9	-14.7
Gallic acid	-5.2	-6.4	-11.6
methylglyoxal	-2.8	-3.3	-6.1
p-Coumaric acid	-5.2	-6.1	-11.3
Quercetin	-6.3	-7.5	-13.8
Symconoside A	-7.2	-7.9	-15.1
Symconoside B	-6.8	-8.4	-15.2
withaferin	-7.7	-7.6	-15.3
β -asarone	-4.5	-4.9	-9.4

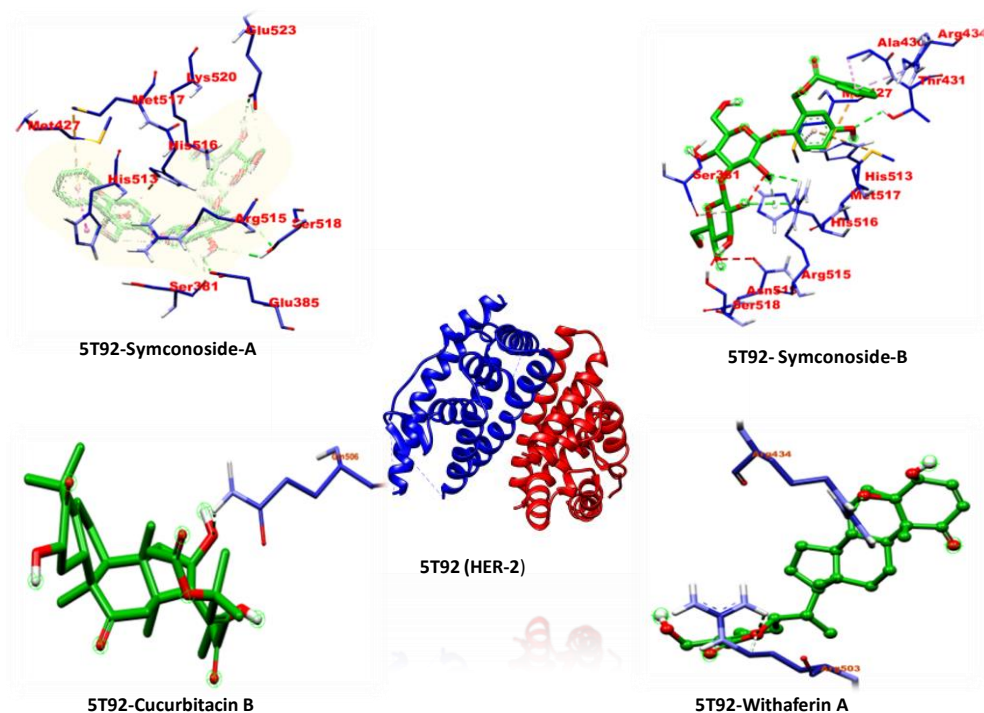


Figure 2: *In silico* HER-2 receptor binding sites of 4 most active components of BR

The four most active ingredients present in BR, symconoside A, symconoside B, curcubitacin, and withaferin A with HER-2, were shown in Fig. 2 as protein-ligand binding sites with amino acid consequences. Figures showed symconoside A binds with 10 amino acids: Ser_381, Glu_385, Met_427, His_513, Arg_515, His_516, Met_517, Ser_518, Lys_520 and

Glu_525 of HER-2; symconoside B binds with 11 amino acids: Ser_381, Met_427, Ala_430, Thr_431, Ala_434, His_513, Arg_515, His_516, Met_517, Ser_518 and Arg_519 of HER-2; curcubitacin B binds with 1 amino acid: Gln_506 of HER-2; and withaferin A binds with 2 amino acids: Arg_434 and Arg_503.

Table 2. *In silico* pharmacokinetic, ADME and toxicological properties of active components of BR

Properties	Symconoside A	Symconoside B	Curcubitacin B	Withaferin A
Molecular formula	C ₂₆ H ₃₂ O ₁₄	C ₂₆ H ₃₂ O ₁₄	C ₃₂ H ₄₆ O ₈	C ₂₈ H ₃₈ O ₆
Molecular weight (g/mol)	568.52	568.52	558.7	470.6
H-bond donor	14	8	3	6
H-bond acceptor	8	14	8	2
Lipinski violation	3	3	1	0
Skin permeation (LogKp)	-2.73	-2.73	-3.50	-3.02
Aqueous solubility (LogS)	-2.606	-2.79	-4.28	-4.46
Bioavailability score	0.17	0.17	0.55	0.55
Human intestinal absorption	32.58 (Low)	30.24 (Low)	79.86 (High)	86.31 (High)
Caco-2 permeability	-0.651	-0.261	0.582	-0.651
Blood Brain Barrier	-1.45	-1.52	-1.17	-0.03
CYP2C9 inhibitor	No	No	No	No
AMES Toxicity	No	No	No	No
Rat Oral Chronic Toxicity (log mg/kg bw/day)	5.42	5.53	1.66	0.95
Rat Oral Acute Toxicity (LD ₅₀) log mg/kg bw	2.50	2.71	3.82	2.78
Maximum tolerable dose (human) log mg/kg/day	-0.008	-0.194	-0.77	-0.41

Table 2 shown the *in silico* anticipated ADME and toxicity evaluations of the four major interacting molecules found in BR, including symconoside A, symconoside B cucurbitacin B, and withaferin A. The strongest H-bond donor was symconoside A, although symconoside B also has the ability to act as an H-bond acceptor. The Lipinski rule was only not broken by withaferin A. Withaferin A and cucurbitacin B both had a 0.55 bioavailability score. Withaferin A showed the highest permeability in Caco-2 cells (0.885 cm/sec). Withaferin A had an intestine absorption rate of 86.32, followed by cucurbitacin B at 79 and symconosides at 30 to 32. The blood-brain barrier (BBB) can be more easily crossed by withaferin A than by the other three substances. No active ingredients have been found to be harmful to AMES or to inhibit CYP2C9. Symconoside B, cucurbitacin B, and withaferin A all had human tolerated doses (log mg/kg/day) of -0.194, -0.77, and -0.412, respectively, demonstrating their non-toxic nature.

DISCUSSION

Alkaloids, flavonoids, terpenoids and polyphenols compounds are a few examples of natural compounds that have been extensively used in preclinical research of breast cancer over the past 20 years due to their abundance in the natural world, low toxicity, and high efficacy. Recently published studies revealed active components of BR possess anticancer effects. These components might improve the immune system's capacity to defend DNA against damage, reduce oxidative damage, or initiate the apoptotic process¹⁷⁻¹⁸. However, its role in cancer has not been explored as a composite. Cancer immunotherapy has gained increasing attention over the past few decades and has grown into an excellent option for cancer treatment²⁵. Clinical adjunct therapies for the treatment of cancer have a long history in traditional Indian medicine.

MTT cell proliferation assay is one of the most widely used for evaluating anticancer activity of both synthetic derivatives and natural products. The viable cells contain NAD(P)H-dependent oxido-reductase enzymes, which reduce the MTT to formazan. In the present study, BR dose dependently and significantly enhanced the cytotoxicity and cell viability was seen to drop in MCF-7 breast cancer cells. This single study unequivocally demonstrated BR has anticancer effect. In a previous study, withaferin A, one of the active ingredients in BR, demonstrated cytotoxicity against four human cancer cell lines: DU-145 for the prostate, HCT-15 for the colon, A-549 for the lung, and IMR-32 for the neuroblastoma²⁶. Similarly, cucurbitacin B has been found to have anticancer properties in human leukemia cells²⁷. Moreover, symconosides showed cytotoxic action on Hep3B hepatocellular carcinoma cells²⁸.

In the transwell cell migration assay, the ability of cells to chemotactically move in the direction of a chemo-attractant is quantified. Cell migration studies tally the quantity of cells that pass through a porous membrane, whereas cell invasion procedures quantify cell movement across extracellular matrix, a critical step in angiogenesis. The topology of the extracellular environment, adhesion, confinement, and stiffness are the main physical factors affecting cell movement²⁹⁻³⁰. Additionally, it might evaluate distinct migratory capacities brought on by the over-expression of a receptor²¹. Present study demonstrated BR has the ability to prevent cancer cells from spreading over normal cells into the surrounding tissues. Fruit pulp extract of *Aegle marmelos*, one of the important components of BR exhibited anti-proliferative activity through suppressing the breast tumor growth rate³¹.

Matrix metalloproteinases (MMPs) are a family of zinc-dependent endopeptidases. MMP-9 plays vital roles in cancer cell invasion and tumor metastasis. It supports angiogenesis by weakening matrix barriers and has the ability to reduce

tumor neovascularization. It also plays a part in the breakdown of the basement membrane³². Breast cancer tissues have high levels of MMP-9, which is directly linked to lymph node metastases and tumor stage³³. Triple-negative and HER-2-positive breast tumors clearly exhibit over expression of MMP-9³⁴. It can serve as a guide for determining the prognosis and course of treatment for breast cancer. Hence, development of MMP-9 inhibitors is an important area for breast cancer research³⁵. In the present study, metastatic matrix protein expression, particularly MMP-9 showed dose dependent inhibition after BR treatment in MCF-7. Withaferin A showed a significant correlation with a decrease of MMP-9 mRNA expression levels in metastatic Caski cell line³⁶.

Furthermore, many cancers, including breast cancer, have been shown to over express the cytokine interleukin-6 (IL-6), which is found in the tumor microenvironment. In the tumor microenvironment, fibroblasts associated with the tumor and tumor cells are the main producers of IL-6³⁷⁻³⁸. The immunopathogenic role of IL-6 and its signaling in breast cancer tumor development, metastasis, and treatment resistance has been shown in numerous investigations³⁹. It is clear that the presence of high levels of IL-6 in breast cancer tissues encouraged the production of Jagged-1, which in turn helped the cancer cells proliferate and maintain their aggressive nature⁴⁰. Therefore, it would seem that IL-6 targeting and/or its receptor in combination with other effective anticancer medicines could be a potent therapeutic approach for breast cancer therapy⁴¹. In this study, BR treatment showed significant inhibition of IL-6 concentrations in the MCF-7 metabolites. Other studies confirmed that withaferin A blocked IL-6 and TNF- α -induced cancer cell invasion and thereby eliminated the interactions between STAT3, STAT1, and NF- κ B and suppressed STAT3 phosphorylation⁴².

Breast stem cell proliferation, differentiation, and cell death are regulated by ER- α and HER-2 signaling pathways, and breast cancer is mostly caused by the over expression of these signaling pathways. Hence, ER- α and HER-2 antagonists have received a lot of interest as possible anti-cancer drugs⁴³. The target of the present grid based *in silico* docking study was to screen out the potential antagonists of ER- α and HER-2 from the most effective compounds or ligands present in BR, like apigenin, cucurbitacin B, gallic acid, methylglyoxal, p-coumaric acid, quercetin, symconoside A, symconoside B, withaferin A and β -asarone¹⁷. The present study revealed cucurbitacin B has the highest binding score with ER- α , followed by withaferin A, symconoside A, and B, indicating their strong antagonistic properties in breast cancer. In addition, symconoside B has the greatest binding score to HER-2, followed by symconoside A, withaferin A, quercetin, cucurbitacin B, and apigenin. Withaferin A also displayed the most encouraging potentiating qualities when taking into account the overall protein-ligands binding affinity for ER- α and HER-2. Withaferin A has previously been shown to treat down-regulation of ER- α protein expression, which correlates with a decline in nuclear level, suppression of mRNA level, and inhibition of E2-dependent activation of ERE2e1b-luciferase reporter gene⁴⁴. Thus, four of the ten active molecules—symconosides A and B, cucurbitacin B, and withaferin A of BR—exhibited synergistic therapeutic potentials for breast cancer. In order to predict preclinical toxicological endpoints, clinical side effects, and ADME characteristics of these substances, *in silico* approaches were further explored. This study offers a powerful systems pharmacology approach for identification of promising and safe molecules from BR for development of breast cancer therapy.

Withaferin A interacts with the positively charged residual amino acids of HER-2 at Arg_434 and Arg_503 and possesses 6

hydrogen donor bonds and 2 hydrogen acceptor bonds; whereas symconoside A interacts with 10 residual amino acids, symconoside B with 11, and cucurbitacin B with a single (Gln₅₀₆). Since withaferin A adheres to all five of Lipinski's principles, it may be termed orally bioavailable than other three components. Withaferin A has the greatest aqueous solubility, whereas symconoside is poorly soluble. The Caco-2 human colon cancer cell line is an example of an experimental screen used in drug discovery to measure membrane permeability and estimate human oral absorption. The most rapid rate of oral absorption and Caco-2 permeability was found in withaferin A. The blood-brain barrier (BBB) has been described as a dynamic interface that regulates the passage of substances between the blood and the brain to maintain the best possible circumstances for neuronal and glial activity⁴⁵. BBB prevents the entry of harmful substances into the brain. All four substances were expected to pass the BBB.

The Ames test is typically used in predicted toxicity models to assess potential carcinogenic/mutagenic effects of substances. In the current investigation, the Ames test revealed that none of the bioactive components were carcinogenic. Moreover, Cytochrome P450 2C9 (CYP2C9) enzyme in liver is involved in drug metabolism and excretion. CYP2C9 inhibition may lead to toxic drug accumulation and hazardous drug-drug interactions in the body. Present *in silico* study highlighted that the components present in BR did not have any role in CYP2C9 inhibition. Additionally, they failed to exhibit oral acute and chronic toxicity in animal models, and they were categorized as class V according to the poisonous class of the Globally Harmonized System of classification of chemical labels.

Therefore, taking into account *in vitro* and *in silico* studies, it may be concluded that the bioactive ingredients present in Body Revival®, exhibit as potent multi-target inhibitors of ER- α and HER-2 with potential anticancer activity without side effects and may be helpful in the treatment management after successful trial in breast cancer patients.

Acknowledgement

We thank CEO of Health Reactive, Mumbai for providing the test samples.

Conflict of Interest

There is no conflict of interest.

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