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**Original Article** 

## EVALUATION OF ANTI-CML ACTIVITY OF METHANOL AND AQUEOUS EXTRACTS OF BENKARA MALABARICA (LAM.) TIRVENG PLANT LEAVES

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## ABSTRACT

**Objective:** To investigate the phytoconstituents and *in vitro* cytotoxicity of methanol (MeOH) and aqueous (AQE) extracts of *Benkara malabarica* (Lam.) Triveng (*BM*) plant leaves.

**Methods:** Gas chromatography-mass spectrometry (GC MS) was carried out to disclose the principal phytoconstituents present in MeOH and AQE extracts of *BM. In vitro* cytotoxicity of *BM* extracts were determined by 3-(4,5-Dimethylthiazol-2-yl)-2,5-diphenyltetrazolium bromide (MTT) assay. Acridine orange (AO)/ethidium bromide (EB) and 4', 6-diamidino-2-phenylindole (DAPI) staining were performed to visualize morphological changes upon treatment of *BM* extracts. Fluorescence-activated cell sorting (FACS) was carried out to determine the apoptosis and cell cycle arrestability of *BM* extracts.

**Results:** GC MS analysis reported the presence of nine phytoconstituents in MeOH and AQE extracts of *BM*. The IC<sub>50</sub> of *BM* MeOH, AQE extracts treated K562 cells were 49.78±1.697, 15.47±1.19  $\mu$ g/ml for 48 h and found to be statistically significant (*p*<0.001). AO/EB and DAPI staining results anticipated the induction of apoptosis and DNA fragmentation upon treatment of *BM* extracts. FACS analysis revealed the SubG<sub>0</sub> cell populations increased in K562 cells treated by *BM* MeOH (18.15) and AQE (51.26) extracts.

**Conclusion:** The results of the present study uncovered that the *BM* AQE extract was more potent in inhibiting K562 cell proliferation through cell cycle arrest and apoptosis compared to the MeOH extract of *BM*.

Keywords: Phyto-constituents, Cytotoxicity, Benkara malabarica, GC MS analysis and K562 cells

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## INTRODUCTION

Medicinal plants are extensively used all over the world as folk medicine because they are cost-effective and inexpensive alternative sources of drugs due to their less side effects [1-3]. Plants are a rich source of secondary metabolites such as flavonoids, carotenoids, phenols, alkaloids, terpenoids, saponins and steroids. These secondary metabolites of plant origin have exhibited antimicrobial, antioxidant and anticancer activities [4, 5]. According to World Health Organization (WHO) reports, approximately 65-80 % of the world population use traditional medicine to treat various diseases. Interestingly, 50 % of all modern clinical drugs are of natural product origin [6-8].

Chronic myeloid leukemia (CML) is a hematoproliferative neoplasm that is marked by uncontrolled myeloid cell divisions in bone marrow. The hallmark of CML is the presence of shortened Philadelphia chromosome (Ph) which arises due to a reciprocal translocation between chromosome 9 and chromosome 22 [(9;22) (q34;q11)], leading to the creation of bcr-abl oncogene. This bcr-abl oncogene encodes a constitutively activated tyrosine kinase, BCR-ABL. The catalytically activated kinase, in turn, activates multiple cell proliferatory signalling pathways such as RAS, a small GTPase, mitogen-activated protein kinase (MAPK), signal transducers and activator of transcription (STAT) and phosphoinositide-3-kinase (PI3K) pathways [9-12]. Tyrosine kinase inhibitors (TKIs) like imatinib [13], nilotinib, dasatinib, bosutinib [14] and ponatinib [15] were approved by the US Food and Drug Administration (FDA). TKIs have changed the clinical course of CML; however, mutations in bcr-abl and multi-drug resistance (MDR) due to efflux of the drug because of overexpression of p-glycoprotein (p-gp) make TKIs less effective. Hence, there is a need for an alternative strategy to develop new BCR-ABL inhibitors. Clearly, natural products obtained from plants offer an alternate, effective and inexpensive source for CML therapy [16].

K562 is a BCR-ABL positive CML cell line that was established from a 53-year-old woman with CML. After 176 passages in 3.5 y, the K562

cell line still has active CML cells in blast crisis stage with Ph [17, 18]. In the present study, the *in vitro* cytotoxicity of *Benkara malabarica* (Lam.) *Tirveng* (*BM*) extracts were evaluated against K562 cells. Previous reports suggest that *BM* leaf and bark extracts reported anti-bacterial and anti-fungal activities [19]. *BM* methanol (MeOH) root extract had reported anti-convulsant activity against isoniazide induced convulsion in *in vivo* models [20].

## MATERIALS AND METHODS

#### **Chemicals and reagents**

All the chemicals and reagents used in this work were purchased from Hi-media Pvt. Ltd, Bombay, India.

#### **Cell culture**

K562 cell line was obtained from the National Centre for Cell Sciences (NCCS), Pune and were maintained in RPMI medium (pH 7.4) supplemented with 10 % fetal bovine serum (FBS) and antibiotics (100 U/ml penicillin and 100  $\mu$ g/ml streptomycin) at 37 °C in a humidified 5 % CO<sub>2</sub> atmosphere.

## Collection and identification of plant material

*BM* plant leaves were collected inside the Pondicherry University campus. The plant sample was identified and authenticated by Dr. N. Ayyappan, French Institute of Pondicherry (IFP), Puducherry. *BM* herbarium specimen was deposited at IFP with accession number HIFP27056.

#### Preparation of BM extracts

100 g of *BM* leaf powder was extracted with both 500 ml of MeOH and water (AQE) using soxhlet apparatus. Extract solution was then filtered through Whatman No.1 filter paper and concentrated using rota-vapour. The concentrated *BM* extracts were stored at 4 °C until further use.

#### Estimation of total phenolic content (TPC)

MeOH and AQE extracts of *BM* in the concentration (conc.) range of 40-200  $\mu$ g were taken and adjusted to 20  $\mu$ l with distilled water. Then, 50  $\mu$ l of Folin–Ciocalteu (FC) reagent (1/10 dilution) was added and incubated for 5 min at room temperature (RT). This was followed by the addition of 50  $\mu$ l of sodium carbonate solution (7.5%). The absorbance was measured at 760 nm. Gallic acid was used as a standard for the calibration curve. The TPC was expressed as gallic acid equivalents (GAE) [21].

#### Estimation of total flavonoid content (TFC)

TFC of the *BM* plant extracts was determined by the aluminium chloride method as described [22]. MeOH and AQE extracts of *BM* of varying conc. (40-200 µg) were taken and was made up to 10 µl with the solvent (MeOH and Distilled water). Then, 30 µl of sodium nitrite (0.03 %) was added and incubated for 5 min at RT followed by the addition of 30 µl of aluminium chloride (10 %) solution. Incubation was carried for 5 min at RT and 200 µl of 1 mmol sodium hydroxide was added. The total volume was then made up to 1000 µl with distilled water and the absorbance was measured at 510 nm. Various conc. of standard quercetin was used to make a standard calibration curve. TFC was expressed as quercetin equivalents (QE).

#### Gas chromatography-mass spectrometry (GC MS) analysis

GC MS analysis of MeOH and AQE extracts of *BM* was carried out using Clarus 500 Perkin–Elmer (auto system XL) of as previously described [23]. Using MS data library on a National Institute of Standards and Technology (NIST) Ver.2.1 and comparing the spectrum obtained through GC MS revealed compounds present in the *BM* extracts.

# 3-(4,5-dimethylthiazol-2-yl)-2,5-diphenyltetrazolium bromide (MTT) assay

K562 cells were plated in a 96-well tissue culture plates at a density of approximately 5,000 cells per well. The cells were then incubated with 100  $\mu$ l of MeOH and AQE extracts of *BM* in the conc. range (2, 4, 8, 16, 32 and 64  $\mu$ g) for 48 h. Untreated cells were used as a control and blank wells containing 100  $\mu$ l of medium only. After 48 h of incubation, 20  $\mu$ l of the MTT reagent (5 mg/ml) was added to each well and the plate was incubated for 4 h at 37 °C. Then the plate was centrifuged at 1500 rpm for 5 min. The supernatant was discarded and to the pellet 150  $\mu$ l of dimethyl sulphoxide (DMSO) was added to each well and absorbance was read at 595 nm using Aspinco biotech Elisa plate reader [24].

## Acridine orange (AO)/ethidium bromide (EB) staining

Approximately  $1\times10^5$ K562 cells were seeded in 6-well tissue culture plate. Cells were then treated with IC<sub>50</sub> conc. of *BM* MeOH and AQE extracts for 48 h. After 48 h of treatment, cells were harvested and washed once with phosphate buffer saline (PBS) (pH 7.4) and stained with AO/EB (1 µg/ml) for 10 min. The stained cells were then examined under a Nikon Eclipse Ti fluorescence microscope (Nikon Instruments Inc., NY, USA) using a UV filter (450-490 nm). Untreated cells were used as control [25].

## 4',6-diamidino-2-phenylindole (DAPI) staining

Approximately  $1 \times 10^5$  K562 cells were seeded into 6-well tissue culture plates. Then, the cells were treated with  $IC_{50}$  conc. of MeOH and AQE extracts of *BM* for 48 h following which cells were harvested and washed once with PBS (pH 7.4) and fixed with 4 % paraformaldehyde for 30 min. The fixed cells were stained with DAPI (1 µg/ml) for 10 min examined under a Nikon Eclipse Ti fluorescence microscope (Nikon Instruments Inc., NY, USA) using a UV filter (450-490 nm). Untreated cells were used as control [26].

#### Cell cycle analysis

Briefly, 1x10<sup>6</sup>K562 cells were seeded in 6 well plates and treated with the IC<sub>50</sub> conc. of *BM* extracts for 48 h at 37 °C. Subsequently, cells were harvested, washed with ice-cold PBS and fixed in 70 % ethanol at 4 °C for overnight. The fixed cells were then incubated with 0.5 ml of propidium iodide (PI) (50  $\mu$ g/ml) solution containing triton X-100 (0.1 %), sodium citrate (0.1 %) and RNase-A (25  $\mu$ g/ml) for 10 min under the dark. The percentage of cells in various phases of the cell cycle was assessed in a BD FACS CALIBUR instrument [27, 28].

#### Statistical analysis

All experiments were carried out in triplicates. The values are represented as a mean±standard deviation (SD). IC<sub>50</sub> values of *BM* extracts in MTT assay were determined using non-linear regression employing GraphPad prism software. MTT results were analysed using a one-way analysis of variance (ANOVA) followed by Tukey's multiple comparisons tests where *p* value<0.001 took as significant using SPSS software.

#### **RESULTS AND DISCUSSION**

#### **TPC and TFC**

Phenols and flavonoids are among the most widely occurring secondary metabolites present in plants. Phenols are aromatic compounds which contain one or several hydroxyl groups directly attached to the benzene ring. Phenolic compounds exhibit antioxidant, anti-mutagenic and anti-cancer activities [29, 30]. TPC in BM extracts was deduced from the standard curve of gallic acid (fig. 1A) (Y = 0.017x-0.004 and  $R^2 = 0.990$ ). The TPC content of MeOH and AOE extracts of *BM* had contained 238.5 and 312 ug/mg of GAE (fig. 1B). Flavonoids belong to polyphenolic compounds which are prevalent in plants. They contain two phenyl rings A, B and a heterocyclic ring C (commonly referred to as a C6-C3-C6 skeleton). Flavonoids exhibit antioxidant, anti-inflammatory, antibacterial, antiviral and anticancer activities [31, 32]. The TFC in BM extracts was derived from the standard curve of quercetin (Y = 0.002x-0.000 and R<sup>2</sup> = 0.989) (fig. 2A). TFC content of MeOH and AQE extracts of BM was found to be 798.5 and 333 µg/mg of QE (fig. 2B).



Fig. 1: TPC of *BM* extracts, A): Standard curve for gallic acid. B): TPC of MeOH and AQE extracts of *BM*. TPC was more in AQE compared to MeOH extract of *BM*. Values were represented as mean±SD (n=3)



Fig. 2: TFC of *BM* extracts, A): Standard curve for quercetin. B): TFC of MeOH and AQE extracts of *BM*. TFC was more in MeOH compared to AQE extract of *BM*, values were represented as mean±SD (n=3)



(B)

Fig. 3: GC MS chromatograph of *BM* extracts, A): GC MS chromatograph of *BM* MeOH extract showed various compounds at major peaks like 18.83-9-Octadecenoic acid [Z]-, methyl ester, 12.53-phenol,2,4-bis[1,1-dimethylethyl]-, 19.43-2-methyl-Z,Z-3,13-Octadecadienol and 21.6-benzene1,1-1<sup>1</sup>-[1-(2,2-dimethyl-3-butenyl)-1,3-propanediyl]bis-were present. B): GC MS chromatograph of *BM* AQE extract major peaks at 18.83, 17.15, 20.32 and 22.7 corresponds to 10-octadecenoic acid methyl ester, hexadecanoic acid methyl ester, Isopropyl stearate and 2-Cyclohexen-3,6-diol-1-one,2-tetradecanoyl

## GC MS analysis

GC MS analysis was based on the computer evaluation of mass spectra of samples through NIST based software, direct comparison

of peaks and retention time with those of standard compounds and computer matching with the NIST library. Both MeOH (fig. 3A) and AQE extract (fig. 3B) of *BM* showed nine important phytoconstituents.



Fig. 4: List of phyto-constituents identified in GC MS chromatograph of MeOH extract of *BM*, A): Phenol,2,4-bis[1,1-dimethylethyl], B): 2hexadecanol, C): 4H-1-Benzopyran-4-one,5-hydroxy-7-methoxy-2-phenyl, D): 11-Tricosene, E): 9-Octadecenoic acid [Z]-, methyl ester, F): 2-methyl-Z,Z-3,13-Octadecadienol, G): Ethanol,2-(9-octadecenyloxy) [Z]-, H): Benzene1,1-1<sup>1</sup>-[1-(2,2-dimethyl-3-butenyl)-1,3propanediyl]bis-and I): Quinoline-4-carboxylic acid, 2-(4-isopropylphenyl)

Various phytoconstituents present in MeOH and AQE extracts of BM were illustrated in fig. 4 and fig. 5.



Fig. 5: List of phyto-constituents identified in GC MS chromatograph of AQE extract of *BM*, A): Hexadecanoic acid methyl ester, B): 10-Octadecenoic acid methyl ester, C): 4H-1-Benzopyran-4-one,2-[3,4-dimethoxyphenyl]-7-hydroxy, D): 10,13-Eicosadienoic acid, methyl ester E): Isopropyl stearate, F): Tricosan-2-ol, G): 2-Cyclohexen-3,6-diol-1-one,2-tetradecanoyl, H): Pyrano[4,3-b]benzopyran-1,9dione,5a-methoxy-9a-methyl-3-[1-propenyl]perhydro-and I): Cyclohexan-1-one-3a,5a diacetic acid,2a(-5 hydroxy-2,4-dimethoxyphenyl)diethyl ester

## MTT assay

MTT assay was performed to determine the cytotoxicity of BM MeOH and AQE extracts (fig. 6) against K562 cells for 48 h. Both BM extracts decreased K562 cell proliferation in a dose-

dependent manner. The IC<sub>50</sub> of MeOH and AQE extracts of *BM* was found to be 49.78±1.697 and 15.47±1.19  $\mu$ g/ml respectively. The results of our assay revealed that AQE extract was more potent than the MeOH extract in reducing the K562 cell proliferation.



Fig. 6: MTT assay of *BM* extracts treated K562 cells for 48 h, values were represented as mean±SD (n=3). Statistical analysis of the experimental results was performed using ANOVA followed by Tukey's multiple comparisons to control in which \*\* represent *p*<0.001 took as significant

## AO/EB staining

Nuclear changes and apoptotic body formation are a characteristic feature of apoptosis, which can be visualized by AO/EB staining [33]. K562 untreated cells appear uniformly green, whereas dead cells appear orange in colour. AQE extract of *BM* (fig. 7C) was more potent in inducing cell death in K562 cells compared to the MeOH extract-treated cells (fig. 7B).

#### **DAPI** staining

Blue-fluorescent DAPI stain preferentially binds to the AT-rich regions of the minor groove of double-stranded (ds) DNA [34]. Binding of DAPI to dsDNA (resulting from DNA defragmentation) increase approximately 20-fold fluorescence enhancement, which was evident in the AQE treated K562 cells (fig. 7F) compared to MeOH extract (fig. 7E). Morphological changes observed in the treated cells displayed broken nuclei as discrete fragments.



Fig. 7: AO/EB and DAPI staining of K562 cells treated with IC<sub>50</sub> conc. of *BM* MeOH and AQE extracts for 48 h and images were captured under 20 X magnification. A): In AO/EB staining untreated K562 cells showed green fluorescence. B): *BM* MeOH and C): AQE extracts treated K562 cells undergoing apoptosis (represented by an arrow mark) appeared as an orange colour. D): In DAPI staining untreated K562 cells visualized as normal blue colour whereas E): *BM* MeOH and F): AQE extracts treated K562 cells undergoing DNA fragmentation (represented by an arrow mark) showed more intensity in blue colour

#### Cell cycle analysis

PI is used as a DNA stain in flow cytometry to evaluate cell viability or DNA content in cell cycle analysis. It is useful for differentiating necrotic (or) apoptotic and normal cells [35]. To determine whether the MeOH and AQE extracts of *BM* induced cell cycle arrest in the K562 cells, the DNA content was measured by

FACS using PI staining. There was a significant increase in SubG<sub>0</sub> population to 51.26 (fig. 8C) and 18.15 (fig. 8B) in K562 cells treated with AQE and MeOH extracts of *BM*. Concomitantly, a decrease in G<sub>0</sub>/G<sub>1</sub> (from 63.04 to 38.93 and 53.04) and S phases (from 24.73 to 6.20 and 13.67) for AQE and MeOH extracts of *BM* was observed. The results confirmed that K562 cells upon treatment of *BM* extracts induce apoptosis.



Fig. 8: Cell cycle analysis of *BM* extracts treated K562 cells. A): K562 untreated cells. B): induction of apoptosis (sub-G<sub>0</sub> phase) by *BM* MeOH, C): AQE extracts for 48 h revealed by PI staining

#### CONCLUSION

Plants are being investigated extensively for their pharmacological purpose as potent anti-cancer drugs. In the present study, *BM* MeOH and AQE crude extracts were being used to examine their cytotoxicity towards CML cell line. Secondary metabolites like phenols and flavonoids were estimated. TPC was more in AQE extract than the MeOH extract of *BM*. TFC was high in MeOH extract of *BM* compared to AQE extract of *BM*. GC MS analysis revealed nine important phytoconstituents in MeOH and AQE extracts of *BM*. In *vitro* cytotoxicity of these extracts was tested against K562 cells. AO/EB staining, DAPI staining and cell cycle analysis revealed that AQE extract is more potent than the MeOH extract of *BM* in inducing apoptosis in K562 cells. Further studies are required to provide more comprehensive data on the anti-CML activity of the AQE and MeOH extracts of *BM*.

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## **AUTHORS CONTRIBUTIONS**

All authors equally contributed to drafting the paper. All authors have read and approved the final manuscript.

#### ABBREVIATION

methanol–MeOH, aqueous–AQE, *Benkara malabarica* (Lam.) Tirveng–*BM*, gas chromatography mass spectrometry–GC MS, 3-(4,5-Dimethylthiazol-2-yl)-2,5-diphenyltetrazolium bromide–MTT, acridine orange–AO, ethidium bromide–EB, 4',6-diamidino-2phenylindole–DAPI, fluorescence-activated cell sorting–FACS, world health organization–WHO, chronic myeloid leukemia–CML, Philadelphia chromosome–Ph, mitogen activated protein kinase– MAPK, signal transducers and activator of transcription–STAT, phosphoinositide-3-kinase–PI3K, tyrosine kinase inhibitors–TKIs, food and drug administration–FDA, multi-drug resistance–MDR, pglycoprotein–p-gp, national centre for cell sciences–NCCS, fetal bovine serum–FBS, French institute of Pondicherry–IFP, Total phenolic content–TPC, concentration–conc, folin–ciocalteu–FC, room temperature–RT, gallic acid equivalents–GAE, total flavonoid content–TFC, quercetin equivalents–QE, national institute of standards and technology–NIST, dimethyl sulphoxide–DMSO, phosphate buffer saline–PBS, propidium iodide–PI, standard deviation–SD, analysis of variance–ANOVA, double-stranded–ds, department of science and technology–DST and sophisticated analytical instrument facility–SAIF.

## **CONFLICTS OF INTERESTS**

The authors declare that they have no conflict of interest

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