



**COMPARISON OF GENOTOXICITY PRODUCED BY HYDRO ALCOHOLIC EXTRACT OF
CURCUMA AROMATICA SALISB, CURCUMA ZEDOARIA WITH CURCUMIN BY
AMES TEST, COMET ASSAY AND MICRONUCLEUS TEST**

Srividya A.R^{*1}, Dhanbal S.P², Sathish Kumar M.N³ and Vishnuvarthan V.J.³

¹Department of Pharmaceutical biotechnology, JSS College of Pharmacy Udhagamandalam, Tamilnadu, India

²Department of Pharmacognosy and Phytomedicine, JSS College of Pharmacy, Udhagamandalam, Tamilnadu, India

³Department of Pharmacology, JSS College of Pharmacy, Udhagamandalam, Tamilnadu, India

*Corresponding Author Email: pharmarsrividya@yahoo.com

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ABSTRACT

This paper deals with the comparison of genotoxicity produced by the hydro alcoholic extract of *Curcuma aromatica* salisb and *Curcuma zedoaria* (christm) rosecoe with curcumin by Ames test, comet assay and Micronucleus test. *Curcuma aromatica* and *Curcuma Zedoaria* belongs to the family Zingiberaceae. In the presence of metabolic activation factor (S9), the revertant colonies were found to be more in the case of TA98 and TA 100 strains. When compared to positive control the numbers of revertant colonies were found to be more or less equivalent to negative control which confirmed the absence of genotoxic effect of test compounds. In the comet assay, among the three tested compounds, curcumin was found to be more genotoxic in nature when compared by *Curcuma aromatica*, *Curcuma zedoaria*. The damage of the cells was found to be more in the presence of metabolic activation factor than the absence of metabolic activation factor. This confirmed that the phytoconstituents are not genotoxic in nature but their metabolites that are formed in the presence of metabolic activation factor is mainly responsible for their genotoxic effect. In the Micronucleus test, when compared to the positive control (Methyl methane sulfonate at the concentration of 50 µg/ml), all the test compounds such as hydro alcoholic extract of *Curcuma aromatica*, *Curcuma zedoaria*, and Curcumin at the concentration of 50µg/ml showed less toxicity to HEp-2 cells. These results might either due to the presence of low concentration of curcumin in the extracts or due to the presence of other phytoconstituents that are present in the extract.

Keywords: *Curcuma aromatica*, *Curcuma zedoaria*, Ames test, comet assay, micronucleus test, Curcumin.

INTRODUCTION

The potential consequence of genetic damage that a cell incur is called genotoxicity. In all the cells and organism the damage response mechanism is available. Genotoxins are the substances which causes the genotoxicity in a cell¹. In India majority of the population uses traditional natural preparations from the plant material for the treatment of various diseases and hence it is necessary to assess clastogenic potential of the traditional plant extract when associated with other substances. Along with the increase in genotoxicity; risk of developing cancer increases². In everyday life, usage of antimutagens and anticarcinogens are suggested and most effective procedure for preventing human cancer and genetic diseases. In medicinal plants, a bioactive component acts as a strategy to block or reverse carcinogens at the early stages. In cancer treatment medicinal plants were considered effective and inexpensive. Herbal plants were proved antimutagenic agents and helps in inhibiting carcinogenic effect of some chemical mutagens, herbal plants were proved as an antimutagenic agent. Currently there is enormous interest in developing new plant based pharmaceutical products³. *Curcuma* species are perennial herbs belonging to the family Zingiberaceae. From that species *Curcuma aromatica* and *Curcuma zedoaria* has been selected for this study. Traditionally *Curcuma aromatica* rhizome is used as tonic, carminatives, astringent, bitters, in sprain and in snake bite and also for treating skin eruption due to infection. α - Curcumene and β Curcumene, d and p-methoxy cinnamic acid are present in this rhizome. Curcumin is the coloring matter in these rhizomes⁴. Traditional uses of *Curcuma zedoaria* rhizome are Carminative, stomachic, gastrointestinal stimulant, diuretic, expectorant demulcent, rubifacient. It is also used in flatulence and dyspepsia. For checking leucorrhoea discharge fresh roots are used and it is

also used as blood purifier. This rhizome is used in the treatment of Goiter according to the information that is present in the Ayurvedic pharmacopoeia of India. Number of terpenoids including Curcumin, Curcumenone, Curdione, Curcumenol, Curzerenone, Furanogermenone, Germacrone, Germacrone epoxide a volatile oil (1.0-1.5) resembling ginger oil and starch (50%) are present in the dried rhizome. It is also used as an antitumor, antiallergenic and antimicrobial agent^{5,6}. Curcumin which is an active component is present in the range of 2-5%. It is an orange crystalline powder which is insoluble in water. For centuries, Curcumin has been consumed as a dietary spice at dose up to 100mg/day. The biological activities that are reported for Curcumin are blood cholesterol reduction, LDL peroxidation prevention, platelet aggregation inhibition, thrombosis suppression, Myocardial infarction curing, type –II diabetes associated symptoms suppression, rheumatoid arthritis, Multiple Sclerosis, Alzheimer's disease, human immune deficiency virus (HIV) replication inhibition, wound healing enhancement, liver protection from injury, bile secretion increase, cataract formation prevention, Protects from Pulmonary toxicity and fibrosis prevention. It also possesses anti-leshmaniasis anti atherosclerotic and anticancer activities^{2,4}. It proved to be a powerful antioxidant⁷. We have performed the genotoxicity studies of the same plants by various methods such as chromosomal aberration studies², DNA sugar damage test, SOS Chromotest, plasmid nicking assay and sequential analysis⁴. Due to the diversity of the end points it is clear that the potential genotoxicity or mutagenicity of a compound cannot be assessed by a single assay system⁸. By any single genotoxic test procedure the mutagenic potential of an agent can't be reliably determined because currently available mutagenicity assay have their own strength and weakness⁹. In this paper we dealt with the

genotoxic activities of hydro alcoholic extract of *Curcuma aromatica*, *Curcuma zedoaria* and Curcumin by Ames reversion assay, Micronucleus test and Comet assay.

MATERIALS AND METHODS

The rhizomes of *Curcuma aromatica* salisb and *Curcuma zedoaria* (Christm) Roscoe are purchased from PSS Herbs Pvt. Ltd, Kerala, India and authenticated by Dr. S. Rajan, Botanist, Udthagamandalam, Tamilnadu, India. Curcumin was sponsored by Sami Labs Pvt. Ltd, Bangalore, India. The rhizomes of *Curcuma aromatica* salisb, *Curcuma zedoaria* (Christm) Roscoe are shade dried; milled and coarse powder is separated. 50% hydro alcoholic extracts are prepared for *Curcuma aromatica* salisb, *Curcuma zedoaria* (Christm) Roscoe rhizomes and these prepared extracts are used for antioxidant and genotoxicity studies.

Ames test

It is the method for evaluating the mutagenic properties of extracts by bacterial reverse mutation assay using various tester strains. This test was carried with certain modification according to Meshram *et al* 1992. TA 98, TA 100 of *Salmonella typhimurium* are the histidine deficient (His⁻) mutant tester strain. The bacterial reverse mutation test uses amino acid requiring strains of *Salmonella typhimurium* to detect the point mutation which involves substitution, addition or deletion of one or few DNA base pairs. The principle of this bacterial reverse mutation test is that it detects mutation which revert mutation present in the test strains and restore the functional capability of the bacteria to synthesize an essential amino acid. The revertant bacteria are detected by their ability to grow in the absence of essential amino acids required by the parent tester strains. TA-98 strain is used to identify the frame shift mutation. TA-100 strain is used to identify the base pair substitution. These mutant strains are obtained from Amala Cancer Institute, Thirussur, India. On receipt these strains are labeled in in-house and kept under refrigerated conditions. One loopful of microorganism is removed and inoculated into a conical flask containing nutrient broth. The inoculated flask is incubated at 37^o C for about 14-18 hours to achieve a density of 1-2x10⁹CFU/ml. 100µl of overnight culture (1-2x10⁹CFU/ml) is used for the bacterial reverse mutation assay. Extracts is made to soluble in reverse osmosis water and 50µg/ml concentrations of the extracts are prepared. Vehicle control is plated for tester strains in the presence and absence of S9 mixture. In the absence of S9 mix, 2- nitrofluorene is used as positive control for the strain TA98. In the presence of S9 mix benzo [a] pyrene is used as positive control. In the absence of S9 mix, sodium azide is used as positive control and 2- aminoanthracene is used as positive control in the presence of S9 mix for the strain TA-100. The tester strain does not possess enzymes systems which are present in mammals to metabolize pro- mutagens to active electrophilic metabolite capable of reacting with DNA. Sometimes these pro-mutagens interact with mammalian enzyme system and yield mutagenic metabolite. Hence it is necessary to add external metabolic activation system that is post mitochondrial fraction (S9). Presence / absence of genotoxicity is detectable, based up on the number of revertant colonies per plate^{10,11}.

Procedure

The revertant colonies are counted, their mean and standard deviation is calculated. Positive result is defined as a dose

dependent increase or a reproducible increase at one or more concentrations in the number of revertant colonies per plate in at least one strain with or without metabolic activation system. Bacterial culture is inoculated from the frozen cultures to 10ml of nutrient broth and incubated at 37^oC for 12hours to obtain approximately 1x10⁹cells/ml. To 2ml of top agar medium containing 0.6% agar, 0.6 % NaCl and traces of histidine (21µg/ml) and biotin (24.4µg/ml) and 0.1ml of cell suspension, 0.5ml of extract solution (50µl/ml concentration) are added. Solvent control, positive controls, S9 mix (0.5/ml plate, 10 %v/v) are also added on the top of the nutrient agar medium. Experiments are carried out in triplicate for each tested compounds. Then the plates are incubated upside down for 2days at 37^oC and the resulting histidine revertant are counted.

Comet assay

Comet assay was being carried out with the human blood. 4ml of heparinized whole blood is mixed with 3ml of RPMI medium. The medium is centrifuged at 4^oC for 15minutes at 1600rpm. Buffy coat containing the lymphocytes is washed in RPMI 1640 medium. Then the cells are resuspended in RPMI 1640 medium and the concentration of the cells is adjusted in such a way that 1ml should contain 10⁶cells approximately. By Trypan blue staining method the cell viability is counted and checked. The cells that exceeded 90% viability are used in genotoxicity assay. Under alkaline method the comet assay was carried out according to the method of Singh *et al*. 1988 with slight modification. 0.9ml of the cell suspension is mixed with 0.1ml of the *Curcuma aromatica*, *Curcuma zedoaria* and Curcumin. And then it is incubated at 37^oC for 30minutes. By centrifugation, the cells are collected and suspended in low melting point agarose in the phosphate buffer saline with the pH7.4. The cells are immediately pipetted onto a glass slide which was already coated with a layer of normal melting point agarose. Then the glass slide is kept at 4^oC for 5minutes to get solidify. Over this agarose layer melting agarose is added as the third layer and the slide was kept at 4^oC for 5minutes. For about an hour the slides are immersed in lysing solution (2.5M NaCl, 10mM Na₂EDTA, 10mM Tris pH 10, 1% Sodium sarcocinate, 1% Triton X-100 and 10% dimethyl sulfoxide) at 4^oC. The slides are placed in a horizontal gel electrophoresis. Using 25V and 300mA electrophoresis is performed for 20minutes. With the neutralizing buffer (400mM, Tris Buffer pH7.5) the slides are washed and then stained with ethidium bromide. Using a fluorescence microscope cells are visualized. Using the computerized image analysis system the data are analyzed. By the ratio of tail to head length the extend of DNA damages is ascertained and scored as follows:- <5.0 % -No damage, 5.1-20.0% - Low damage, 20.1-40.0%-medium damage, 40.1-95.0%- high damage, >95.1%-complete damage.¹²⁻¹⁴

Micronucleus assay

Micronucleus assay was carried out by using Human larynx epidemoid carcinoma cells (HEp-2) cells which was are obtained from the National cell culture, Pune, India. In 25cm² flasks containing 5ml of complete medium, HEp-2 cells are seeded and maintained as monolayer growing at 37^oC. 10% fetal calf serum is supplemented to the culture medium along with 2.38mg/ml of HEPES and 0.01mg/ml of Streptomycin and 0.005mg/ml of Penicillin. After thawing 10⁶cells are seeded in several flasks in complete medium for 48hours till the completion of two cycles. The cells are washed with PBS

and treated with the test compounds such as hydro alcoholic extract of *Curcuma aromatica*, *Curcuma zedoaria* and Curcumin at 50µg/ml concentration in serum free medium over a period of 2 hours. The cells that were treated with MMS (Methyl methane sulfonate) at the concentration of 100µg/ml are used as positive control. The cells that did not receive any drug are used as negative control. To evaluate the mutagenicity, the cells are washed twice with 5.0ml of PBS (pH 7.4). To block cytokinesis and to yield binucleated cells, to a fresh complete culture medium Cytochalasin- B is added at the concentration of 3µg/ml and kept for 28 hours. The cells are rinsed twice with 5ml of PBS. After trypsinisation the cells are centrifuged for 5 minutes at 900rpm. In ice-cold hypotonic solution (1% sodium citrate + one drop of 10% formaldehyde) the pellets are resuspended and carefully homogenized with Pasteur pipette. Homogenized cell suspension is once again centrifuged for 5 minutes at 900rpm. The pellets are resuspended in methanol and acetic acid in the ratio 3:1 and the cells were homogenized carefully with the Pasteur pipette. On a previously cleaned slide, the fixed cells are then dropped and covered with a film of ice cold distilled water for 5 minutes. The cells are stained in 3% giemsa dissolved in phosphate buffer (Na₂HPO₄- 0.06M and KH₂PO₄- 0.06M- pH 6.8). Then the cells are washed with water, dried and kept at 4°C until it is scored. Using a microscope at 400x magnification, 2000 cells per slide is analyzed. The criteria that is followed for the identification of binucleated cells and micronuclei as follows:^{15,16}

1. Shape should be round for both the nuclei and micronuclei.
2. When compared to the main nuclei, the micronuclei should be 1/3 small.
3. The main nuclei should not touch the micronucleus.
4. The color and intensity of the micronuclei must be same as the main nuclei.

RESULTS

Ames Reverse mutation assay

In this assay, two strains were used. They are TA 98 and TA 100. TA 98 is used to determine frame shift mutation that is caused by the test compounds. TA100 is used to determine base pair substitution. Both the strains were obtained from Amala Cancer Research Institute, Thirussur, Kerala, India. To determine the frame shift mutation, in the absence of S9 factor, 2- nitrofluorene and in the presence of S9 factor Benzo[a]pyrene was used as the positive control which produced the number of revertant colonies such as 682±86 and 510±44 respectively at the concentration of 50µg/ml. In the negative control plate the revertant colonies produced were 12±1, 16±2 in the absence and presence of S9 respectively. In the presence of S9 factor revertant colonies found to decrease in the case of positive control where as for the test compounds, revertant colonies found to increase in the presence of S9 factor.

In the absence and presence of S9 factor

Revertant colonies were found to be more at the concentration of 50µg/ml for *Curcuma aromatica* in the absence of S9 factor 14±2 and in the presence of S9 factor the number of revertant colonies were 18±3. For *Curcuma zedoaria*, in the absence of S9 factor 10±1 and in the presence of S9 factor the number of revertant colonies was 16±1. The number of revertant colonies produced by curcumin was found to be 23±1 in the absence of S9 factor

and 26±3 in the presence of metabolic activation factor. The results are tabulated in Table 1.

TA 100 Control

To detect the base pair substitution type of mutation, in the absence of S9 factor sodium azide and in the presence of S9 factor 2- aminoanthracene were used as positive control which gave the number of revertant colonies such as 1501±132 and 308±20 at the concentration of 50µg/ml. In the negative control plates the number of revertant colonies was 14±2 and 78±18 in the absence and presence of S9 factor respectively.

In the Absence and presence of S9 factor

In the absence of S9 factor, the number of revertant colonies was found to be 16±3 for *Curcuma aromatica* at the concentration of 50µg/ml. At the concentration of 50µg/ml the numbers of revertant colonies were 18±1 for *Curcuma zedoaria* and 25±2 for Curcumin. In the presence of S9 factor the revertant colonies were 64±10 for *Curcuma aromatica* at the concentration of 50µg/ml respectively. At the concentration of 50µg/ml, the numbers of revertant colonies were 80±8 for *Curcuma zedoaria* and 86±7 for curcumin. When compared to positive control the numbers of revertant colonies were found to be more or less equivalent to negative control which confirms the absence of genotoxic effect of test compounds. In the presence of S9 factor Benzo [a] pyrene was used as the positive control, 2- nitrofluorene was used as the positive control in the absence of S9 factor. The results are tabulated in Table 2.

Comet assay

In control samples 93±3 cells were found to have no damage in the presence of metabolic activation factor where as in the absence of metabolic activation factor 97±3 cells were found to have no damage in the control sample. At the concentration of 50µg/ml; In the absence of metabolic activation factor *Curcuma aromatica* salisb, No damage was found in 70±3 cells, low damage was found to be 22±5 and medium damage was found in 8±4 cells. In cells there was no high damage and complete damage occurred in the cells. In the cells treated with *Curcuma zedoaria* (Christm) Roscoe the normal cells were found to be 79±3 cells. In 15±2 cells low damage was observed and in 6±3 cells medium damage was observed. In the cells treated with *Curcumin*, normal cells were found to be 62±4, 28±3 cells showed low damage and medium damage had occurred in 10±4 cells.

At the concentration of 50 µg/ml

In the presence of metabolic activation factor *Curcuma aromatica* salisb caused high damage in 6±5 cells, medium damage was found in 12±3 cells, low damage was found in 24±5 cells and normal cells were found to be 49±2 cells. *Curcuma zedoaria* (Christm) Roscoe caused high damage in 5±4 cells, medium damage in 11±5 cells and low damage in 30 ±1 cells and normal cells were found to be 54±3 cells. Curcumin caused complete damage in 4±3, high damage in 8±3 cells. It caused medium damage in 10±2 cells and low damage in 37±2 cells and normal cells were found to be 41±3 cells. Among the three tested compounds, curcumin was found to be more genotoxic in nature when compared by *Curcuma aromatica*, *Curcuma zedoaria*. These results might either due to the presence of low concentration of curcumin in the extracts or due to the presence of other phytoconstituents that are present in the extract. The damage

of the cells was found to be more in the presence of metabolic activation factor than the absence of metabolic activation factor. This confirms that the phytoconstituents are not genotoxic in nature but their metabolites that are formed in the presence of metabolic activation factor is mainly responsible for their genotoxic effect. These results are tabulated in the Table 3.

Micronucleus test

Curcuma aromatica at the concentration of 50µg/ml produced the micronuclei at the rate of 12.333±2.223, 15.666±1.778, in the absence and presence of metabolic activation factor respectively. *Curcuma zedoaria* at the concentration of 50µg/ml produced the micronuclei at the

rate of 9.666±1.999, 14.666±2.666 in the absence and presence of metabolic activation factor respectively. Curcumin at the concentration of 50µg/ml produced the micronuclei at the rate of 14.666±1.555, 19.333±1.111 in the absence and presence of metabolic activation factor respectively. Positive control (Methyl methane sulfonate) produced the micronuclei at the rate of 25.666±1.555, 35.66±2.886 in the presence and absence of metabolic activation factor respectively. When compared to the positive control (Methyl methane sulfonate at the concentration of 50µg/ml), all the test compounds such as hydro alcoholic extract of *Curcuma aromatica*, *Curcuma zedoaria* and Curcumin at the concentration of 50µg/ml showed less toxicity to HEp-2 cells. The results are tabulated in Table 4.

Table 1: Ames test result for the Strain TA98

Test compounds	Mean ±SEM (-S9)	Mean ±SEM (+S9)
	Concentrations	Concentrations
	50µg/ml	50µg/ml
Positive control	682±86	510±44
Negative control	12±1	16±2
<i>Curcuma aromatica</i> salisb	14±2	18±3
<i>Curcuma zedoaria</i> (Christm.) Roscoe	10±1	16±1
Curcumin	23±5	26±3

Table 2: Ames test result for TA 100

Test compounds	Mean ±SEM (-S9)	Mean ±SEM (+S9)
	Concentrations	Concentrations
	50µg/ml	50µg/ml
Positive control	1501±132	308±20
Negative control	14±2	78±18
<i>Curcuma aromatica</i> salisb	16±3	64±10
<i>Curcuma zedoaria</i> (Christm.) Roscoe	18±1	80±8
Curcumin	25±2	86±7

Table 3: Genotoxicity of *Curcuma aromatica*, *Curcuma zedoaria* and Curcumin in human lymphocytes in the comet assay

Sample	Cell number				
	No damage	Low damage	Medium damage	High damage	Complete damage
In the absence of metabolic activation factor					
Control	93±3	3±3	0	0	0
<i>Curcuma aromatica</i>	70±3	22±5	8±4	0	0
<i>Curcuma zedoaria</i>	79±3	15±2	6±3	0	0
Curcumin	62±4	28±3	10±4	0	0
In the presence of metabolic activation factor					
Control	97±3	5±2	-	-	-
<i>Curcuma aromatica</i>	49±2	24±5	12±3	6±5	0
<i>Curcuma zedoaria</i>	54±3	30±1	11±5	5±4	0
Curcumin	41±3	37±3	10±2	8±3	4±3

Table 4: Mean Frequencies of Micronucleus observed in HEp-2 cells

Treatment	Repetition			Mean ±standard deviation
	1	2	3	
Negative control	6	3	4	4.33±1.086
Positive control- in the absence of S9 factor	40	32	35	35.66±2.886
Positive control- in the presence of S9 factor	25	28	24	25.666±1.555
<i>Curcuma aromatica</i> salisb				
In the absence of S9 factor	13	15	09	12.333±2.223
In the presence of S9 factor	18	16	13	15.666±1.778
<i>Curcuma zedoaria</i> (Christm.) Roscoe				
In the absence of S9 factor	8	9	12	9.666±1.999
In the presence of S9 factor	12	17	15	14.666±2.666
Curcumin				
In the absence of S9 factor	14	17	13	14.666±1.555
In the presence of S9 factor	21	18	19	19.333±1.111

Several substances in plants express cytotoxic and genotoxic activities and show correlation with the incidence of tumors. Therefore, understanding the health benefits and or potential toxicity of the plants is important¹². Although plant extracts have been used in the treatment of diseases according to knowledge accumulated over centuries, it is also known that many plants synthesize toxic substances, which in nature acts as defense against infections, insects and herbivores. Some substances present in some medicinal plants are potentially toxic and carcinogenic and it has also been reported that some traditional medicines may have genotoxic potentials. Assessment of the potential genotoxicity of traditional medicine is indeed an important issue as damage to the genotoxic materials may lead to critical mutation and therefore also to an increased risk of cancer and other disease. Major bioactive phytochemicals that have been associated with many plants are different types of saponins and flavonoid. In over all safety evaluation of the botanicals, a modest trend towards increasing the inclusion of information, a genotoxicity appeared in peak in the last four years because of significant awareness of the impact of genotoxicity¹⁷. In *in-vitro* assay, recent investigations have revealed that many plants used as food or in traditional medicine have mutagenic hazard¹⁸. The isolated compounds from the plants such as Quercetin, furoquinoline, alkaloids and isothiocyanates were considered to be mutagens. It is very difficult to speculate the compounds that are responsible for mutagenic response detected with plant extracts because they are complex mixtures of organic compounds. Short term *in-vitro* and *in vivo* studies as well as long term carcinogenicity studies, with chemically treated animals confirmed that phytochemicals could also possess antimutagenicity and anticarcinogenic effect. Epidemiological studies also supported that the chemo preventive effect in which phytochemicals exhibit genotoxic/mutagenic effect by themselves or potential the effect of other xenobiotics¹⁹. Many literatures stressed the importance of carrying out the genotoxicity studies for the medicinal plants. Botanicals contain multiple chemical constituents which may be pharmacologically active with significant proportions of chemically undefined constituents, the genotoxic information obtained from studies using a whole herbal or multicomponent herb product is relatively lacking. The presence of phytoconstituents including tannins, catechins, flavonones, isoflavones are responsible for the possible genotoxic effects of plant extracts. Flavonoids inhibit topoisomerase I and II enzyme which will interfere with the replication and transcription process, inhibiting the relegation of DNA-double strand breaks and enhancing the formation of cleavable DNA- enzyme complexes. Phenolic rich extracts could lead to accumulated DNA breaks and mutation, thus contributing significantly to genotoxicity. Because of these above statement, a systematic phytoconstituents analysis has been carried out for *Curcuma aromatica* Salisb and *Curcuma zedoaria* (Christm) Roscoe. 50% hydro alcoholic extract of *Curcuma aromatica* salisb found to contain alkaloids, tannin and flavonoids where as *Curcuma zedoaria* (Christm) Roscoe contain alkaloids and flavonoids. Total phenol and total flavonol content was found to be more in *Curcuma aromatica* than *Curcuma zedoaria*. The percentage of curcumin was found to be $6.14 \pm 2.13\%$ and $3.84 \pm 1.76\%$ for *Curcuma aromatica* and *Curcuma zedoaria* respectively. By any single genotoxic test procedure the mutagenic potential of an agent can't be reliably determined because currently available mutagenicity assays have their own strength and weakness⁹. Ames test is an *in vitro*

mutagenicity assay which is useful for prescreening potential carcinogens, as approximately 80% of carcinogens are proved to be mutagenic⁹. In Ames test where the target organism is unable to oxidize chemicals, so necessary co factors must be supplied for an exogenous activation. The major group of chemical carcinogen is activated by S9, generally comprises the hepatic post-mitochondrial fraction from rats pre-treated with Aroclor 1254, a mixture of polychlorinated bi phenyl which serve as a potent inducer of cytochrome P₄₅₀ families in particular P₄₅₀ I and P₄₅₀ II. A clear mutagenic response is seen if the source of the liver preparation is obtained from Arclor 1254- treated animals. In the evaluation of genotoxicity of new chemicals, established mutagens are utilized as positive control to ensure not only the responsiveness of the bacterial strains but also the efficiency of the activating system during the routine employment of the Ames test. To mutate all *Salmonella typhimurium* strain 2- amino anthracene is most widely used mutagen which appears to be activated by hepatic system derived from all animal's species including man. N-hydroxylation catalyzes the activation of 2-amino anthracene since Aroclor 1254 is an established inducer of P₄₅₀- I family. The mutagenicity of 2-aminoanthracene was markedly decreased when the animals are treated with Arcoclor 1254. Microsome from the untreated animals could bioactivate 2-amino anthracene to mutagens²⁰. TA 98, TA 100 of *Salmonella typhimurium* are the histidine deficient (His⁻) mutant tester strain. In many of the tester strain TA 98, TA 100 of *Salmonella typhimurium* does not have intrinsic potential to induce mutation by increasing the number of His⁺ colonies. Exogenous metabolic activation is required for most of the chemicals to form an ultimate mutagenic species to induce mutation in *in- vitro* assays including Ames Salmonella test. Some electrophilic chemicals are mutagenic and directly act with DNA. For metabolic activation studies, in general 4-5% Aroclor 1254- inducer S9 fraction in the S9 mixture is used. In this study 1% S9 factor was used. To induce the mutagenic response, few chemicals require an elevated level of S9 fraction 10-30% in the S9 mixture¹⁰. Biological samples may cause problems in assays for mutagenicity using Ames test/ Salmonella test because of the presence of autotrophic growth factors. In each of the colonies growing on the minimal agar plates, histidine added by biological samples to the test system may extend the autotrophic growth phase of plated bacteria thereby increase the probability of spontaneous reversion to prototrophy. Histidine- related growth factors added in the plate incorporation test may give a false positive due to the consequence of extended autotrophic growth²¹. Some 20years ago, Ostling and Johnson developed the comet assay, which has its origin in the micro gel electrophoresis technique which can be performed in different ways and has been used for many different purposes during the last few years. By using the alkaline version of the assay it is possible to detect not only DNA single strand breaks and alkali labile sites but also DNA/DNA and DNA/Protein cross-links²². The assay which is capable of detecting the DNA damage in individual cells is the comet assay. At the time of lysis, increased DNA migration results from the induction of DNA- single strand breaks, alkali labile sites and incomplete excision repair sites cell death arising from a non- DNA mediated process or apoptosis is associated with increased DNA migration accompanied with DNA fragmentation. In extreme cases like the apoptotic cells, the head and tail are well separated and with an increasing number of breaks, DNA pieces migrate

freely into the tail of the comet. Information about the number of strand breaks is provided by the intensity of fluorescence in the tail relative to the head. Over the wide range of damage, tail length, percentage of total DNA in the tail and tail moment all reflect DNA damage, though the percentage tail DNA generally seems to be most useful. A measure of both the smallest detectable size of migrating DNA which is reflected in the comet tail length and the number of relaxed/broken pieces which is represented by the intensity of DNA in the tail is incorporated in the tail moment. Inter individual differences such as the age of blood donors and their physical activities, smoking habit and cell cycle status are important and may reflect differences in the repair of an induced DNA damage likely to add complexity to the problem. During comet formation in both alkaline and neutral assay system, the chromatin structure which is fundamental to the replication and transcription activity affects the role of DNA²³. Among the three tested compounds Curcumin was found to be more genotoxic in nature when compared by *Curcuma aromatica*, *Curcuma zedoaria*. These results might either due to the presence of low concentration of Curcumin in the extracts or due to the presence of other phytoconstituents that are present in the extract. The damage of the cells was found to be less in the absence of metabolic activation factor than the presence of metabolic activation factor. This confirms the original parent molecule is not genotoxic in nature but its metabolites are producing the genotoxicity effect in various models.

Micronucleus test

A micronucleus test is a test used in toxicological screening for potential genotoxic compounds. The assay is now recognized as one of the most successful and reliable assay for genotoxic carcinogens, i.e carcinogens that act by causing genetic damage and is the OECD guidelines for the testing of chemicals. There are two versions of this test; one is *in vivo* and other *in vitro*. A micronucleus is the erratic (third) nucleus that is formed during the anaphase of mitosis or meiosis. Micronuclei are cytoplasmic bodies having a portion of acentric chromosome or whole chromosome which was not carried to the opposite poles during the anaphase. Their formation result in the daughter cell lacking a part or all of a chromosome. These chromosome fragments or whole chromosomes normally develop nuclear membrane and forms as micronuclei as a third nucleus. After cytokinesis, one daughter cell ends up with one nucleus and the other ends up with one large and one small nucleus, i.e., micronuclei. There is a chance of more than one micronucleus forming when more genetic damage has happened. The micronucleus test is used as a tool for genotoxicity assessment of various chemicals. For the safety evaluation of new drugs or industrial chemicals in Japan as well as in other countries, the micronucleus test has been widely used as a sensitive *in vivo* genotoxicity test and has been recommended for use in regulatory guidelines. From guidelines to guidelines, the experimental protocol of the test varies slightly in details such as number of animals, sex, dosing and sampling time. Before the start of full scale experimentation, a pilot preliminary experiment was to be carried out for each test substances because the optimum response may vary according to the chemical concerned²⁴.

CONCLUSION

Among the three tested compounds curcumin produced the higher genotoxic effect in all the tested models. Among the

plant extracts, *Curcuma aromatica* showed more genotoxic effect than *Curcuma zedoaria* but when compared to the positive control the genotoxic effect was found to be very very less and it was almost similar to that of negative control. This confirms that Curcumin alone in the presence of metabolic activation factor produced genotoxicity in somewhat higher amount than its absence.

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