

# *In vitro* and *In silico* Antigout arthritic activities of Ethanolic and Aqueous stem extracts of *Cissus quadrangularis* – A TLR2 and TLR4 Receptor approach

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## ABSTRACT

The *in vitro* anti-gouty arthritic activity of ethanolic and aqueous extracts of *Cissus quadrangularis* stem was evaluated in terms of inhibition of xanthine oxidase, proteinase enzyme, protein denaturation and membrane stabilization. The Monosodium Urate (MSU) up-regulated expression patterns of toll-like receptors namely TLR-2 and TLR-4 were analyzed. The molecular docking was performed to select the antagonistic phyto ligand for receptors since their activation leads to the destructive immunological reactions occurring in gouty arthritis. The *in vitro* anti-gout arthritic activities were estimated using standard protocols. The MSU crystal-induced toll-like receptors gene expression was analyzed. Molecular docking and LIGPLOT analysis were performed with the GCMS-derived phytoconstituents in the ethanolic extract of *Cissus quadrangularis* with the receptors to find the interactions. Colchicine was used as a positive drug in the study. Among the two extracts, ethanolic extract revealed better *in-vitro* anti-arthritic activity in the present investigation. Out of the GCMS derived 13 phytoconstituents, based on docking score three main components namely Pentadecanoic acid, 14-methyl-,methyl ester, 10-Octadecenoic acid,methyl ester, 4-one,2-(3,4-dihydroxyphenyl)-2,3-dihydro-3,5,7-trihydroxy showed better antagonistic action in the LIGPLOT analysis. It may be concluded these three ligands may further develop as a potential anti-gouty arthritic drug.

## INTRODUCTION

Gout is a metabolic disorder with a worldwide distribution that is associated with an excess of circulating levels of uric acid (hyperuricemia). This results in the formation and deposition of monosodium urate crystals (MSU) in the joints causing inflammatory arthritis and severe pain. Hyperuricemia can be caused by impaired renal excretion or overproduction of uric acid due to overconsumption of purine-rich foods. Xanthine oxidase enzyme is responsible for the hyperuricemia and gouty catalyzing the oxidation of xanthine and hypoxanthine into uric acid (Unno *et al.*, 2004). The risk for supersaturation occurs and the urate crystallizes when the plasma concentration of monosodium urate (MSU) exceeds its solubility (around 7 mg/dl) concentration due to impaired glomerular filtration. Recognition of the naked

MSU crystal by Toll-like receptor TLR2 and TLR4, which are normally involved in triggering the innate host defense responses to infectious pathogens, was recently discovered to be a primary trigger of the inflammatory and degenerative tissue reactions associated with gouty arthritis (Liu-Bryan *et al.*, 2005; Liu-Bryan *et al.*, 2005). Nonsteroidal anti-inflammatory drugs (NSAIDs), as well as systemic and intra-articular corticosteroids, are used to treat acute gout (Getting *et al.*, 2002). In patients who cannot tolerate NSAIDs or systemic corticosteroids, oral colchicine is generally the next choice for primary therapy. But all these treatments have their own side effects such as gastrointestinal and liver toxicity.

It is a well-known fact that many Indian herbs are capable of a wide range of medicinal effects. From the time immemorial, mainly based on the practical experiences, these medicinal practices were developed and followed. *Cissus quadrangularis* is a medicinal plant which belongs to the Vitaceae family usually cultivated in India and Ceylon. In Ayurvedic system of medicine, this plant is used for the treatment of venereal diseases, gout, piles,

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leucorrhoea and syphilis (Yoganarsimhan, 2006). Siddha traditional medicine believes the use of this plant to heal broken bones, as an analgesic and a tonic (Mishra *et al.*, 2010). The pathophysiology of the gouty arthritis is characterized by multiple steps, where a potential drug should not only act to reduce the production of uric acid but should also reduce the immune response by inhibiting the activation of the Toll-like receptor and its downstream processes leading to the degenerative reactions involved in gouty arthritis. With the above scenario, the present investigation evaluated the and anti-gout activity of *Cissus quadrangularis* stem extracts through *in vitro* and *insilico* analysis.

## MATERIALS AND METHODS

### Chemicals

The chemicals used in the present study such as Ethanol, Bovine Serum Albumin, Acetyl Salicylic Acid were obtained from SD Fine Chemicals Ltd. Xanthine oxidase from bovine milk was purchased from Sigma ( $\times 4500$ ). Trypsin was purchased from SISCO Research Laboratories Pvt Ltd, Mumbai. The other chemicals used were of analytical grade.

### Collection of plant material

The stem of *Cissus quadrangularis* was purchased from koyambdu market, Chennai, India and were authenticated by Dr. Sankaranarayanan, Deputy Director, captain Srinivasamorthy Siddha medical institute Chennai, India. The voucher specimen is also available in herbarium file of the same Centre.

### Preparation of plant extract

The stem of *Cissus quadrangularis* (1 kg) was dried and ground into a coarse powder. The powder was extracted with 90% (v/v) ethanol at room temperature by cold maceration process for three days. The resultant extract was filtered and the excess solvent was removed by vacuum evaporator. The yield of the sample was calculated. Similarly, aqueous extract was prepared and the removal of water was carried out by freeze-drying and the yield of the sample is calculated. The prepared extracts were stored in the refrigerator until investigation.

### *In vitro* anti-arthritis activity of gouty arthritis

The anti-arthritis activity of gouty arthritis was analyzed using standard established protocols. The *in vitro* Xanthine oxidase inhibitory activity, protein denaturation inhibition assay, membrane stabilization study, and Proteinase enzyme inhibition study were studied to assess the anti-arthritis activity of the plant extracts EECQ and AECQ. The activity of xanthine oxidase was calculated using spectrophotometer method by Owens and Johns 1999. The protein denaturation inhibition process was evaluated using Mizushima and Kobayashi, 1968 method. The membrane stabilization of RBC was carried out according to the studies by Sakat *et al.* (2010). The proteinase enzyme inhibitory studies were performed by the method of Oyedapo *et al.* (1995).

### Determination of totalphenolic content

The total phenols present in the EECQ and AECQ extracts were calculated according to the methods of Slinkard and Singleton 1977. A quantity of 100 mg of the sample in 0.5 ml of

water is mixed with Folin–Ciocalteu’s reagent (diluted 1:10v/v) and 2 ml of  $\text{Na}_2\text{CO}_3$  (7.5%, v/v) solution. The whole set up was incubated for 90 min at 30°C, and the resultant color was measured at 765 nm with results expressed as gallic acid equivalents.

### Determination of total flavanoid content

The total flavonoid content present in the extracts were calculated by the method of Zhishen *et al.* (1999). 100 mg of test drugs were dissolved in 2 ml of distilled water and 0.15 ml of sodium nitrite solution. 0.15 ml of aluminium chloride solution was added after 6 minutes and incubated for 6 minutes. To the mixture 2 ml of 4%, NaOH was added followed by the addition of water to bring the volume to 5 ml and incubated for 15 min, color absorbance was determined at 510 nm and results were expressed as catechin equivalents.

### GC-MS analysis

For GC-MS analysis, the samples were injected into an HP-5 column (30 m  $\times$  0.25 mm i.d with 0.25  $\mu\text{m}$  film thickness), Agilent technologies 6890 N JEOL GC Mate II GC-MS model. The following chromatographic conditions were used: Helium as a carrier gas, the flow rate of 1 mL/min; and the injector was operated at 200°C and column oven temperature was programmed as 50-250°C at a rate of 10°C/min injection mode. The Following MS conditions were used: ionization voltage of 70 eV; ion source temperature of 250°C; the interface temperature of 250°C; mass range of 50-600 mass units.

### Monosodium urate crystals synthesis

MSU crystals were prepared according to Seegmiller *et al.*, 1962. 24 g of sodium hydroxide was solubilized in 200 ml of pyrogen-free double distilled water. 1 g of uric acid was added into the above solution and the pH was adjusted to 7.2 using 1N HCl. This solution was heated to 120°C for 6-hrs with gentle stirring. After incubation, the solution was left to cool at room temperature and kept at 4°C for overnight. The sedimented MSU crystals were collected by filtration and washed with absolute ethanol. Washed MSU crystals were dried under vacuum and sterilized by autoclaving at 121°C for 20 min. The sterilized crystals were prepared as 20 mg/ml stock in PBS and used for the assay.

### Cell treatment and quantitative Real-Time Polymerase Chain Reaction (RT-PCR) analysis of TLR4 and TLR2 expression

A quantity of  $1 \times 10^6$  cells/well of human PBMC were plated in 400  $\mu\text{l}$  volume on 24-well tissue culture plate and incubated for 30 min before compound treatment at 37°C under a humidified atmosphere of 95% air and 5%  $\text{CO}_2$ . Aliquotes of monosodium urate crystals were prepared in complete RPMI-1640 medium and transferred to the respective wells the plates were gently rocked for few seconds. 100  $\mu\text{l}$  of complete medium was added to the well served as untreated control. The plate was incubated at 37°C under a humidified atmosphere of 95% air and 5%  $\text{CO}_2$  for 20 hrs. After incubation, the cells were collected by centrifugation at 3000 rpm for 5 min at 4°C. The supernatants were discarded and the cell pellets were washed with Phosphate buffered saline before RNA isolation processes. Total RNA was isolated from treated human PBMC cells using the TRIzol Reagent (Sigma) according to the manufacturer’s instructions. The 2  $\mu\text{g}$

RNA was reverse transcribed by using the higher capacity cDNA reverse transcription kit (Applied Biosystems, CA, USA) and the mRNA expression was amplified by Quantitect SYBR1 PCR kit (QIAGEN, Valencia, CA, USA). Gene-specific primers for TLR4 and TLR2 expression were designed by referring to the NCBI/primer-BLAST tool software which is depicted in the Table 1. The RT-PCR data was collected and analyzed on MxPro Software from Agilent Technologies. The fold change in gene expression levels of target genes was calculated with normalization to  $\beta$ -actin values using the  $2^{-\Delta\Delta C_t}$  comparative cycle threshold method.

### Molecular docking analysis

*In silico* molecular docking studies were conducted with the help of patch dock to identify the best ligand for TLR2 and TLR4 proteins from the *Cissus quadrangularis* phyto compounds. The three-dimensional structures of toll-like receptor 4 (TLR-4) (PDB ID: 5NAO) and toll-like receptor 2 (TLR-2) (PDB ID: 2Z80) proteins were downloaded from protein data bank (<http://www.rcsb.org/pdb/>). The two-dimensional structures for GC-MS identified 13 phyto constituents from the *Cissus quadrangularis* were retrieved from PUB chemsite, and converted into three-dimensional structures using corina 3D converter. Using the patch dock server docking score regarding the ligand and protein interaction was identified. Using ligand showing appropriate docking score was selected and their further interactions with the proteins in terms of amino acid residues with the ligand and the active site were visualized and further confirmed using LIGPLOT. The docked results were saved as “pdb” file and binding affinity and molecular interaction between standard, test compounds and the receptor protein were visualized using PyMol Molecular Graphic System (Ver. 1.0) and Discovery Studio (Ver. 3.1) software, respectively. The standard drug colchicine is used as a reference drug.

### Statistical analysis

The values reported are Mean  $\pm$  SE. The statistical analysis was carried out using analysis of variance (ANOVA)

followed by Dunnett's 't' test. The p values < 0.05 were considered significant.

## RESULTS

### Xanthine oxidase inhibitory assay

The Xanthine oxidase inhibitory activity of EECQ and AECQ were depicted in Table 2. Between these two extracts the ethanolic extract of *Cissus quadrangularis* shows higher the maximum inhibitory activity to xanthine oxidase starting from 50  $\mu$ g concentration when compared to aqueous extract of *Cissus quadrangularis*. In the present study, the percentage of inhibition was almost similar to allopurinol the positive drug used.

### Inhibition of protein denaturation

The potential of inhibition of protein denaturation was studied (Table 3) and better inhibition was seen for EECQ than AECQ and at the concentration to 500  $\mu$ g and 1000  $\mu$ g of EECQ maximum inhibition potential was noted which was higher than the inhibition percentage of Acetylsalicylic acid the positive control used.

### Membrane stabilization study

The heat and hypotonic saline-induced damage to RBC membrane was stabilized by the plant extracts (Table 4) and were comparable with the positive drug Acetylsalicylic acid. The EECQ extract protected the RBC membrane than the AECQ.

### Proteinase inhibition study

The ability of the EECQ and AECQ to inhibit proteinase enzyme was shown in the Table 5. Though both the extracts showed proteinase inhibitory activity, the inhibition was found to higher for EECQ than AECQ extracts. In the present investigation the, 200  $\mu$ g to 1000  $\mu$ g of EECQ showed higher inhibitory activity than the Acetylsalicylic acid which was the positive drug used in this study.

**Table 1:** Details of Primers used for TLR-2, 4 and  $\beta$ -actin.

Gene	Primer	Primer sequence	Size of the amplicon (BP)
TLR-2 (NCBI Ref sequence NM_001318787.1)	Forward	GGCCAGCAAATTACCTGTGT	298
	Reverse	TTCTCCACCCAGTAGGCATC	
TLR-4 (NCBI Ref sequence NM_138557.2)	Forward	TGAGCAGTCGTGCTGGTATC	167
	Reverse	CAGGGCTTTTCTGAGTCGT	
$\beta$ -actin (NCBI Ref sequence NM_001101)	Forward	CATCGAGCACGGCATCGTCA	211
	Reverse	TAGCACAGCCTGGATAGCAAC	

**Table 2:** Effects of EECQ and AECQ on Inhibition of Xanthine Oxidase.

Conc of Extract (in $\mu$ g)	% Inhibition of EECQ	% Inhibition of AECQ	% Inhibition of Allopurinol
50	49.8 $\pm$ 1.41 a**	34.26 $\pm$ 0.9 b**	70.3 $\pm$ 0.7
100	56.8 $\pm$ 1.4 a**	48.9 $\pm$ 1.36 b**	72.0 $\pm$ 1.2
200	70.3 $\pm$ 1.05 a**	56.2 $\pm$ 0.75 b**	75.2 $\pm$ 0.76
500	83.5 $\pm$ 2.6 a ns	65.5 $\pm$ 1.45 b**	81.2 $\pm$ 1.15
1000	91.53 $\pm$ 1.35 a**	66.9 $\pm$ 1.2 b**	83.76 $\pm$ 1.49

Values are expressed in mean  $\pm$  SD (n = 3), statistical significant test for comparison was done by ANOVA followed by Dunnett's 't' test. Comparison between a: Allopurinol vs. EECQ b: Allopurinol vs. AECQ \*p < 0.05, \*\*p < 0.1 and ns: Non-Significant.

**Table 3:** Effects of EECQ and AECQ on Inhibition of Protein Denaturation.

Conc of Extract (in µg)	% Inhibition of EECQ	% Inhibition of AECQ	% Inhibition of Acetyl Salicylic Acid
50	49.8 ± 1.41 a**	34.26 ± 0.9 b**	70.3 ± 0.7
100	56.8 ± 1.4 a**	48.9 ± 1.36 b**	72.0 ± 1.2
200	70.3 ± 1.05 a**	56.2 ± 0.75 b**	75.2 ± 0.76
500	83.5 ± 2.6 a ns	65.5 ± 1.45 b**	81.2 ± 1.15
1000	91.53 ± 1.35 a**	66.9 ± 1.2 b**	83.76 ± 1.49

Values are expressed in mean ± SD (n = 3), statistical significant test for comparison was done by ANOVA followed by Dunnet's 't' test. Comparison between a: Acetylsalicylic acid vs. EECQ b: Acetylsalicylic acid vs. AECQ \*p < 0.05, \*\*p < 0.1 and ns: Non-Significant.

**Table 4:** Effects of EECQ and AECQ on membrane Stabilization.

Conc of Extract (in µg)	% stabilization of EECQ	% stabilization of AECQ	% stabilization of Acetyl Salicylic Acid
50	50.2 ± 0.95 a**	36.1 ± 1.06 b**	69.9 ± 1.3
100	61.9 ± 1.58 a**	51.2 ± 1.10 b**	74.2 ± 0.95
200	71.5 ± 1.25 a**	58.6 ± 1.41 b**	80.23 ± 1.0
500	78.33 ± 0.92 a**	63.9 ± 1.45 b**	82.8 ± 1.22
1000	82.6 ± 1.10 a ns	67.8 ± 2.4 b**	83.8 ± 1.5

Values are expressed in mean ± SD (n = 3), statistical significant test for comparison was done by ANOVA followed by Dunnet's 't' test. Comparison between a: Acetylsalicylic acid vs. EECQ b: Acetylsalicylic acid vs. AECQ \*p < 0.05, \*\*p < 0.1 and ns: Non-Significant.

**Table 5:** Effect of EECQ and AECQ on Proteinase Inhibition.

Conc. of Extract (in µg)	% Inhibition of EECQ	% Inhibition of AECQ	% Inhibition of Acetyl salicylic acid
50	13.49 ± 0.70 a**	10.57 ± 0.84 b**	23.27 ± 0.67
100	26.56 ± 0.60 aa ns	20.44 ± 0.75 b**	27.40 ± 1.29
200	40.20 ± 1.19 a*	35.72 ± 0.79 b ns	37.56 ± 0.86
500	80.34 ± 0.72 a ns	72.52 ± 0.75 b**	79.56 ± 0.86
1000	92.69 ± 0.74 a**	86.31 ± 0.78 b**	89.28 ± 0.68

Values are expressed in mean ± SD (n = 3), statistical significant test for comparison was done by ANOVA followed by Dunnet's 't' test. Comparison between a: Acetylsalicylic acid vs. EECQ b: Acetylsalicylic acid vs. AECQ \*p < 0.05, \*\*p < 0.1 and ns: Non-Significant.

### GCMS analysis

In the present investigation, the ethanolic extract of *Cissus quadrangularis* revealed better anti arthritic activity than its aqueous extract. So the EECQ was subjected to GC-MS analysis which revealed the presence of 13 chemical components (Figure 1).

### MSU effect on TLR-2 and TLR-4 gene expression

The effect of the MSU induced inflammatory response which is mediated by TLR-2 and TLR-4 receptors was studied using human peripheral blood mononuclear cells treated with monosodium urate crystals. The level of gene expression of these receptors with varying concentration of MSU crystals was studied using the RT-PCR method and depicted in Figure 2 and 3. In our present investigation, the expression pattern of TLR-2 was concentration-dependent that is at a lower concentration of MSU crystals such as 125 µg the gene expression was minimum and then gradually increases with increase in concentration and peaks at 1000 µg. Whereas TLR-4 gene expression was independent on the concentration of MSU and at lower concentration (125 µg)

itself it showed maximum gene expression.

### Molecular docking analysis

Molecular docking was performed for all the 13 compounds obtained from GCMS analysis with the TLR2 and TLR4 receptor using the patch dock sever and docking score was obtained (Table 6). Based on the docking score three compounds namely Pentadecanoic acid, 14-methyl-methyl ester, 10-octadecenoic acid, methyl ester, 4-one, 2-(3,4-dihydroxyphenyl)-2,3-dihydro-3,5,7-trihydroxy- were selected for the study of ligand-protein interaction using the LIGPLOT analysis. The number and nature of amino acid interaction through hydrogen bonding as well as hydrophobic interaction of the phyto compounds as well as the standard drug colchicine was explained in Table 7, Figure 3 and 4. From the previous studies it is noted that all these three compounds have the ability to bind with the amino acid arginine which was present in the active site of both TLR2 and TLR4 receptor by hydrogen bonding and thereby acting as antagonist to MSU crystals. Several amino acids in these two receptors are involved in the hydrophobic interactions with these phytochemicals but their interaction occurs in the binding site of the receptor rather than the active site.

**Table 6.** Docking score values of compounds with TLR-2 and TLR-4.

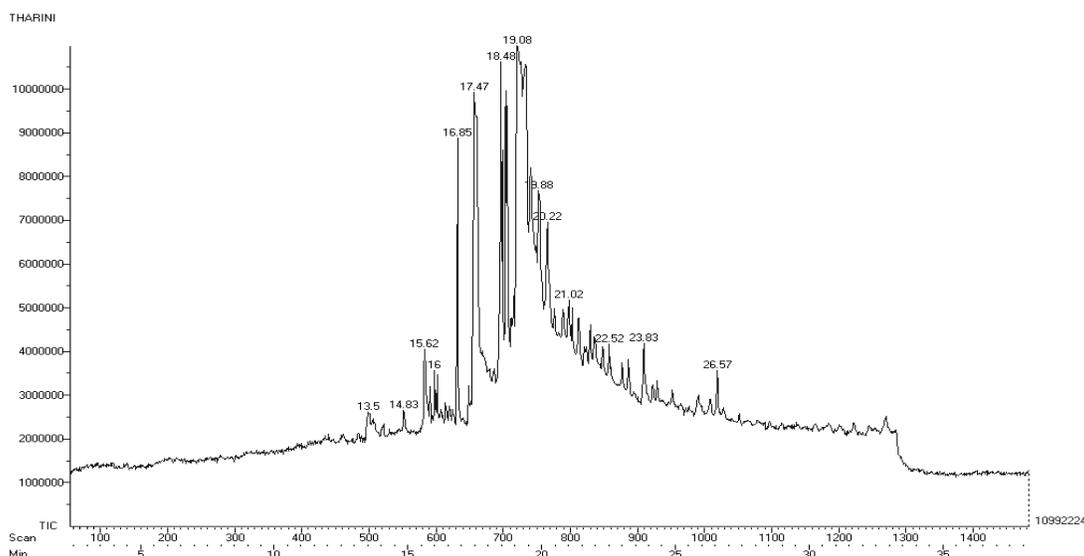
S.No	COMPOUND NAME	TLR-2	TLR-4
1	oxiraneoctanoic acid 3-octyl- methyl ester	-163.55	-4.23
2	1-Terpinen-4-ol	-87.12	-48.46
3	Flavone	-130.62	-51.91
4	Tetradecanoic acid	-86.86	-23.10
5	1-oxacyclopentadecan-2-one, 15-isopropenyl	-158.58	-6.31
6	Pentadecanoic acid, 14-methyl,-methyl ester	-131.38	-17.74
7	Hexadecanoic acide, ethyl ester	-156.67	-14.12
8	10-Octadecenoic acid, methyl ester	-140.91	-5.89
9	Oleic acid	-111.70	-37.48
10	Z,E-2-Methyl-3,13-octadecadien-1-ol	-131.39	-59.57
11	2-Oxo-5-benzoxy-6-methyl-4-phenyl-1,2,3,4-tetrahydropyrimidine	-212.66	-40.83
12	Isopropyl stearate	-145.05	-78.57
13	4-one,2-(3,4-dihydroxyphenyl)-2,3-dihydro-3,5,7-trihydroxy-	-97.48	-7.44
14	Colchicines	-145.54	-27.06

## DISCUSSION

In gout the uric acid in the form of monosodium urate crystals get deposited in the joints, synovial fluid and other tissues and causes an inflammatory and painful condition called as gouty arthritis (Albrecht *et al.*, 2014). The inflammatory response in gout is characterized by multiple steps and complex interactions between various cell types which ultimately leads to the ulceration of the joint cartilage, marginal osteophytosis, and chronic inflammation of synovial membrane (Dalbeth and Haskard, 2005; Carado *et al.*, 2006). The goals of gouty arthritis treatment are several fold. Initially, the drug should inhibit the overexpressed xanthine oxidase enzyme responsible for the production of uric acid. In the present investigation, the ethanolic extract of *Cissus quadrangularis* inhibited the xanthine

oxidase enzyme considerably. The presence of high content of phytosterol, calcium, ascorbic acid  $\beta$ -sitosterol,  $\delta$ -amyrin, carotene and  $\delta$ -amyron have the ability to modify the metabolic and physiological effects as reported by Metha *et al.* (2001); Shirwaikar *et al.* (2003). These may be responsible for the xanthine oxidase inhibitory activity observed in the present study.

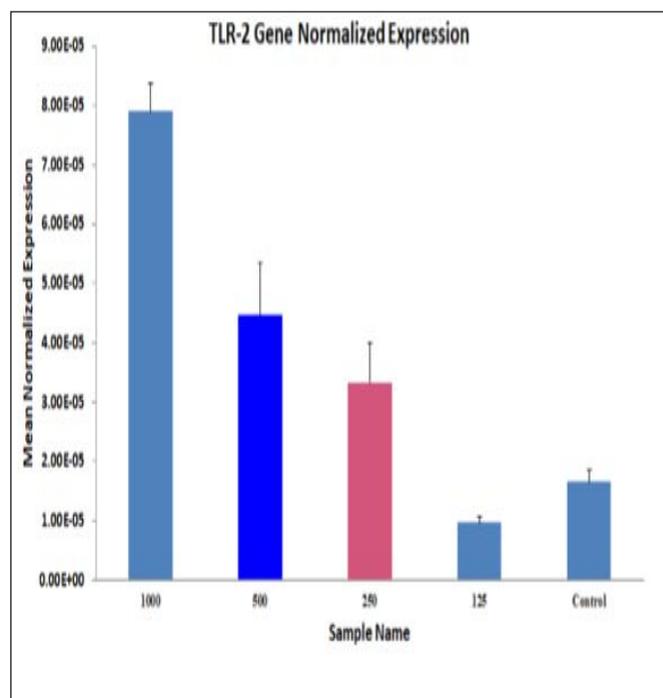
The binding of the MSU ligand as an agonist to TLR2 and TLR4 receptors in the neutrophils is the initial trigger to the immune response and inflammatory stimuli which orchestrate the multiple signaling pathways such as activating the NLRP3 inflammasome, resulting in processing and secretion of IL-1 $\beta$ , increased NF-kB activity, triggering the cytokines, such as interleukin-1 (IL-1), IL-6, IL-8, and tumor necrosis factor, matrix metalloproteinase (MMPs), and certain other mediators leading to the pathogenesis of gouty arthritis. (Haskard and Landis, 2002; Landis and Haskard, 2001). Similar to other forms of arthritis, there also the three basic interrelated processes like inflammation, synovial proliferation and joint tissue destruction that occurs in gouty arthritis. So the *in vitro* anti-arthritis activity in terms of inhibition of protein denaturation, membrane stabilization, and proteinase inhibition was studied for our plant extracts. The EECQ exhibited better anti gouty arthritic activity for membrane stabilization, inhibition of protease enzyme and protein denaturation. It is a well-known fact that the lysosomal membrane is similar to the RBC membrane and any constituent which protects the RBC membrane has the ability to protect the lysosomal membrane. This, in turn, inhibits the release of protein-degrading enzymes from the neutrophils at the inflammatory site and thereby tissue inflammation and damage (Chou CT, 1997). All types of inflammatory processes usually trigger directly or indirectly the different types of immunological mediators which subsequently induce the proteases and destruction of cells and tissues. Phytochemicals which specifically inhibit these inflammatory mediators have the protective effect as an anti-gout agent. According to Panthonget *et al.* (2007), the analgesic and anti-inflammatory activity of the plant extract is due to the presence of flavonoids especially luteolin and by  $\beta$ -sitosterol. Delapureta *et al.* (2005), have stated that the influx of neutrophils, as well as reduction of enzymes like MPO in the inflamed tissue, can be reduced by the  $\beta$ -sitosterol present in the *Cissus quadrangularis* extract.



**Fig. 1:** Shows the list of bioactive compounds present in the EECQ with their retention time.

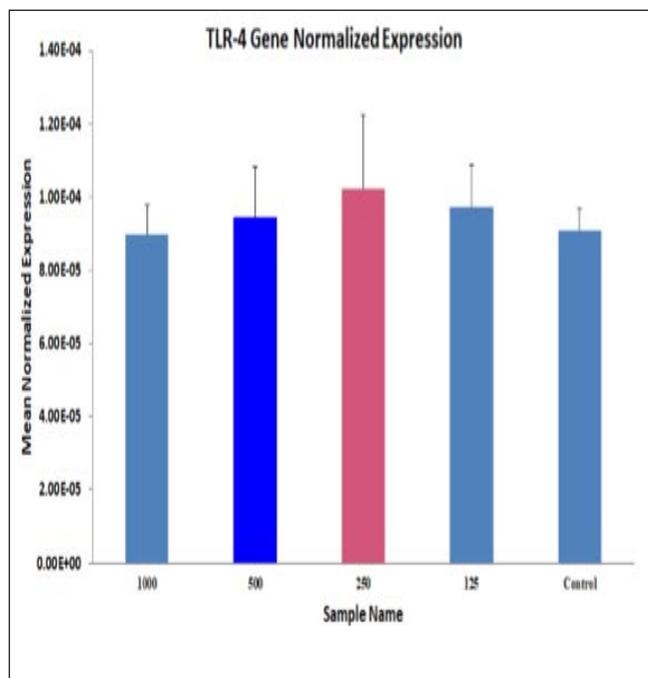
**Table 7:** Interaction with the TLR-2 and TLR-4 proteins with phytochemicals of EECQ.

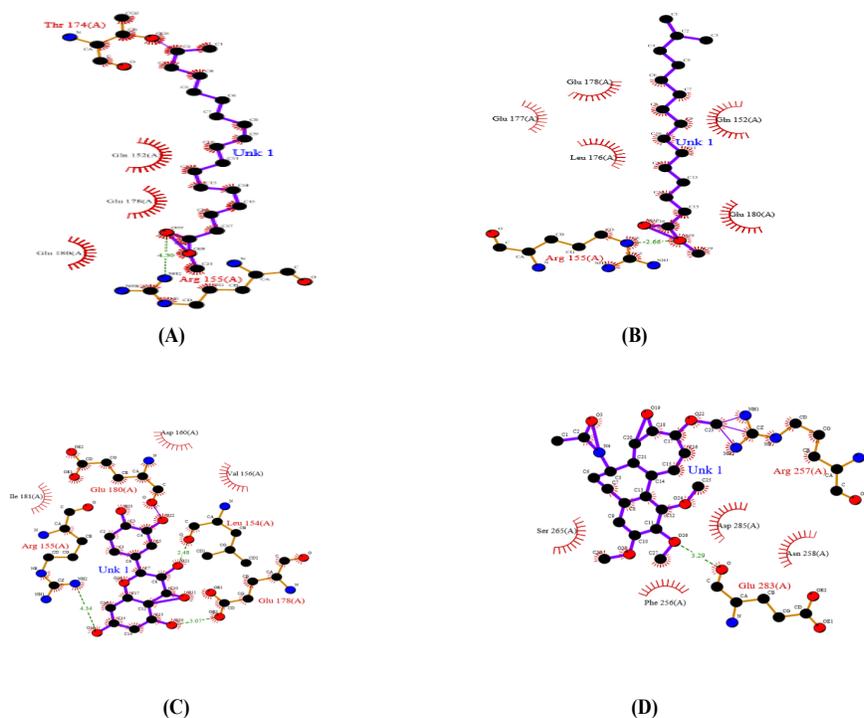
S. No	Compound	TLR-2 amino acid binding site		TLR-4 amino acid binding site	
		H bonding sites	Hydrophobic contact sites	H bonding sites	Hydrophobic contact sites
1	Pentadecanoic acid, 14-methyl-, methyl ester	1 Arg155(A)	87 Gln152(A), Leu176(A), Glu177(A), Glu178(A), Glu180(A).	1 Arg289(B)	63 Lys230(B), Arg234(B), Val259(B), Arg264(B), Ala271(B), Val316(B), Ser317(B)
2	10-Octadecenoic acid, methyl ester	1 Arg155(A)	91 Gln152(A), Thr174(A), Glu178(A), Glu180(A)	–	47 Arg234(B), Arg264(B), Arg289(B), Ala291(B), Tyr296(B), Ser317(B), Asn339(B), Lys341(B), Lys362(B)
3	4-one,2-(3,4-dihydroxyphenyl)-2,3-dihydro-3,5,7-trihydroxy-	3 Leu154(A), Arg155(A), Glu178(A).	75 Val156(A), Asp160(A), Glu180(A), Ile181(A).	–	62 Arg264(B), Leu293(B), Asp294(B), Tyr296(B), Thr319(B), Asn339(B), Lys341(B), Lys362(B).
4	Colchicines	2 Glu283(A), Asp286(A).	102 Phe256(A), Arg257(A), Asn258(A), Ser265(A), Asp285(A)	3 Asp294(B), Tyr296(B), Asn339(B).	95 Tyr292(B), Tyr295(B), Ser317(B).

**Fig. 2:** TLR-2 gene expression chart.

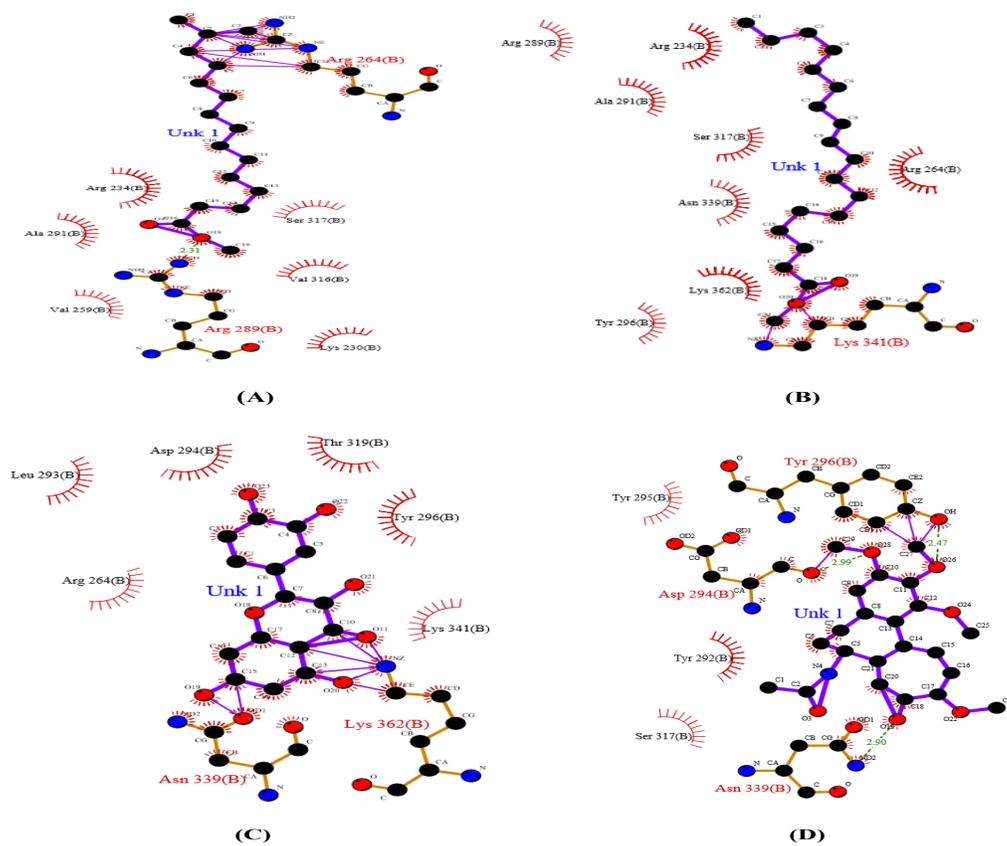
In the present study, we could able to conclude that there is an association between the MSU crystals and overexpression TLR 2 and TLR4 genes. The expression pattern was high for the TLR4 gene than the TLR 2 gene. So we also made an attempt to examine whether the phytochemicals present in the *Cissus quadrangularis* stem extract could act as an antagonist to these transmembrane protein receptors (TLR 2 and TLR4) and thereby inhibit the down streaming inflammatory processes. Based on the GC-MS analysis of the ethanolic extract of the *Cissus quadrangularis* stem 14

compounds were subjected to patch dock method of molecular docking analysis. Three compounds namely Pentadecanoic acid, 14-methyl-, methyl ester, 10-Octadecenoic acid, methyl ester, 4-one, 2-(3,4-dihydroxyphenyl)-2,3-dihydro-3,5,7-trihydroxy showed better activation energy and docking score. These three compounds were further subjected for the ligand-protein interactions for the possible antagonist action for the TLR 2 and TLR4 trans-membrane receptor. This blocking of the TLR 2 and TLR4 transmembrane receptor could hinder the further immunological reactions and protect the joints from destruction and pain.

**Fig. 3:** TLR-4 gene expression chart.



**Fig. 4:** Amino acid Interaction of TLR-2 protein with specific ligands. A) Pentadecanoic acid,14-methyl-,methyl ester. (B) 10-Octadecenoic acid, methyl ester. (C) 4-one,2-(3,4-dihydroxyphenyl)-2,3-dihydro-3,5,7-trihydroxy. (D) colchicines.



**Fig. 5:** Amino acid Interaction of TLR-4 protein with specific ligands. (A) Pentadecanoic acid,14-methyl-,methyl ester (B) 10-Octadecenoic acid, methyl ester. (C) 4-one,2-(3,4-dihydroxyphenyl)-2,3-dihydro-3,5,7-trihydroxy (D) colchicines.

## CONCLUSION

This knowledge will provide a new insight for the discovery of a single ligand which will be acting as an inhibitor for this TLR2 and TLR4 trans-membrane receptors thereby terminating the diverse immunochemical reactions leading to the pathogenesis of gouty arthritis. The single ligand may be used as a successful drug for the management of gouty arthritis which can replace the different types of drug which are at present in use for the treatment of gouty arthritis.

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