Kuijiean Suppress Inflammation in Ulcerative Colitis Rat Models by Phosphorylation Level of HuR

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Abstract

In this study used a TNBS/ethanol method and combined complex factors established the rat model of ulcerative colitis and treatment with the Xipayi Kuijiean (KJA) to conduct whole process. We detected the expression level of hnRNA and mRNA in the five inflammation factor such as IL-1α, IL-1β, IL-10, IL-17, TNF-α and regulatory protein NF-xB and HuR in UC with complex factor group (UC group), normal group, UC treated with KJA group (KJA group) and negative control group. In UC group, we assessed the colonic damage and colon length by naked eye decision scores, KJA significantly reduced the severity of colitis inflammation. The immune dysfunction tends to be basically restored in colon tissue of the KJA group. Furthermore, in UC group hnRNA and mRNA level are increased, but mRNA level significantly higher than hnRNA. The post-transcriptional level of inflammatory related factors was mainly followed by the reduction of mRNA level after treated with KJA. These results demonstrate that post-regulation level of HuR has an important role in the development of UC.

Keywords: TNBS/ethanol method and combined complex factors induce rat model; Ulcerative colitis; qRT-PCR; Xipayi Kuijiean; HuR

Introduction

Inflammatory bowel disease (IBD) is a chronic inflammatory disease of the gastrointestinal tract, which clinically contains Crohn’s disease (CD), ulcerative colitis (UC), and other conditions [1,2]. In recent years, the UC disease obviously increasing both at home and abroad, and closely associated with the onset of colorectal cancer, the main risk of colorectal cancer is found to be high in UC patients include longer duration, greater anatomical extent of colonic involvement [3].

The etiology of UC includes immunity, heredity, environment, intestinal flora and mental factors, but the pathogenesis is not clear [4]. Among them, immunological mechanism is concerned by more and more researchers, especially the intestinal mucosal immune system plays a key role in the occurrence and development of UC [5,6].

The imbalance between pro-inflammatory and anti-inflammatory cytokines that occurs in UC results in disease progression and tissue damage and limits the resolution of inflammation. Cytokines play an important role in the immune pathogenesis of ulcerative colitis, where they drive and regulate multiple aspects of intestinal inflammation [7,8].

There are pro-inflammatory cytokines which include IL-1, IL-2, IL-6, IL-8, IL-12, TNF-a, IFN-gamma etc, and anti-inflammatory cytokines with transforming growth factor beta (TGF-β), IL-1 receptor antagonist (IL-1Ra), IL-4, IL-10, IL-13, etc. These cytokines in intestinal immune regulation of UC patients, play an important role in the inflammatory response. The functions instead of balance between the two kinds of cytokines were damaged and balance disorders or various cytokines not normally present in a cell, which can lead to the occurrence of UC [9].

Xipayi Kuijiean (KJA) is a prescription for treating UC [10]. Here, we provide data supporting such studies, indicating that KJA administered in the TNBS/ethanol and combined with complex factors established rat model of UC. In previous research KJA can suppress ulcerative colitis in rat model because of the strong anti-inflammatory properties of KJA [11].

This study mainly aims at observing the curative effectiveness of KJA on pro-inflammatory cytokines (IL-1α, IL-1β and TNF-α) as well as on anti-inflammatory cytokine (IL-10) in UC rats to confirm and explore the mechanism in treating UC. The expression level of hnRNA and mRNA of IL-1α, IL-1β, IL-10, TNF-α, mRNA of IL-17 increased in colon tissue of UC group.

The expression of cytokines was regulated by NF-xB transcriptional and HuR post-transcriptional levels. NF-xB performs a pivotal function in the expression of many genes involved in immune and inflammatory responses, including ones that contribute to UC.

In normal intestinal epithelial cells, inactive NF-xB present in the cytoplasm by binding to their inhibitory subunit, IκB. In response to stimuli, IκB proteins are phosphorylated, ubiquitinated, and degraded. The ultimate result is the translocation of NF-xB into the nucleus, where it can bind specific DNA binding sites and regulate the transcription of target genes, such as IL-1β and TNF-α [12-15].

In our previous study, the phosphorylation level of NF-xB is increased in UC model group, that indicate changes of transcriptional levels of various genes in UC are likely to be related to the regulation of NF-xB [16].

In UC, activation of NF-xB and the release of inflammatory mediators, leads to the damage of the colonic mucosa. NF-xB often
needs of specific sequences with some other regulatory factor to form a complex, the role of NF-kB promote or inhibit depends on the combination of other regulatory factors, however which regulatory factors affect to the NF-kB activity need to further study. RNA binding proteins associated with RNA binding protein HuR, Tristetraproline (TPP) and other proteins in cell [17].

HuR is one of the important post-transcription factors to regulate gene expression and has very important biological function in the body, it is not only the important regulatory factors of life activity, or important regulatory factors in the process of activated immune cells and its dysfunction related to inflammation, tumor and other diseases [18].

Therefore, the study on the expression and regulation of HuR and its inflammatory factors is of great significance in the development of UC and the research on drug intervention. In this study using Western blot detected the phosphorylation levels of NF-kB in nucleus and HuR in cytoplasm. The phosphorylation level of NF-kB and HuR are raised in negative control group but they were decreased in the KJA group [16].

That indicated the NF-kB and HuR are activated in UC group, treated with KJA could inhibit the activation or phosphorylation of NF-kB and HuR, and they regulate transcription of cytokines and stability of the mRNA respectively, could affect the expression level of cytokines.

qRT-PCR used to detected the expression level of hnRNA and mRNA of IL-1α, IL-1β, IL-10, TNF-α, mRNA of IL-17 from different group rats colon tissue, analyze the regulation mechanism of the expression level of candidate genes expression in the whole process of UC.

Material and Methods

Materials

**Chinese traditional medicine:** Xipaiy Kuijiean (KJA) Enema (Preparation Standard of Medical Institutions Formulated by the Food and Drug Administration of Xinjiang Uygur Autonomous Region ZJZ-0001-2013) is composed of gallic (liquid herbal extract, 50 mL/bag) and saline were provided by genomics technology service headquarters; 2 x SYBR Green - Mix (QIAGEN, Germany), 2 x Taq PCR MasterMix purchased from Beijing TIANGEN; primers using Primer 5.0 software design.

**Chemical reagent:** TNBS(sigmaUSA, number#SLB 6263V)Chloral hydrate, anhydrous ethanol, liquid paraffin, ether and physiological saline were provided by genomics technology service headquarters; Tritol from Invitrogen used for RNA extraction; 2 x SYBR Green - Mix (QIAGEN, Germany); 2 x Taq PCR MasterMix purchased from Beijing TIANGEN; primers using Primer 5.0 software design.

**Instruments and equipment:** FLI- 2999 HT artificial climate box; a gradient PCR (BIO- RAD in Cycler TM Thermal Cycler, USA); Real-time fluorescent quantitative PCR (ABI 7500 Real-time PCR, United States); High-speed centrifuge at low temperature ( Eppendorf centrifuge 5417r, Germany); coagulation glue imager (BIO-RAD, USA); Ultramicro spectrophotometer (Thermo Scientific Nanodrop 2000/USA); –80°C ultra-low temperature freezer (Thermo Forma series, The United States).

**Primer design:** For mRNA and hnRNA primer sequences were identified from the genomic database (NCBI, GenBank), using Primer 5 design software or blast Primer.

**Antibody and reagent for Western blot:** Phospho-NF-kB p65 antibody from Cell signaling technology. HuR antibody and beta actin antibody from Santa Cruz Biotechnology. SDS, TEMED, Tris-base and BCA protein assay reagent kit was provided by Shanghai sangon biological engineering co. LTD.

Method

The grouping experimental animals: SPF male Wistar rats (180 ± 30 g) aged 6 to 8 weeks provided by Animal Experimental Center of Xinjiang Medical University. The rats were randomly divided into normal group (10 rats), TNBS induced ulcerative colitis model group (15 rats), complex factor induced ulcerative colitis and treated with KJA (15 rats) and negative control group (20 rats) each 5 rats per cage. They were adaptively fed for seven days before building models.

**Establish experiment rat models:** Control group were housed in a controlled environment (temperature 24-25°C, humidity 70%-75%) and were fed on a normal laboratory diet. All experiments were conducted according to the guidelines of the local ethics committee at Xinjiang Medical University.

**Establish the complex factors induced UC rat model group and KJA group:** According to the literature [19] using artificial climate box to create an environment (relative temperature is 26-28°C, relative humidity of 36-40%, and light), put animals to 10 h a day [21-23], feed 300 g per cage for daily and 200 ml water [24,25], according to body weight, rats feed with 2 ml/100 g dose gastric medicine per daily.

A chronic stimulation is given that include: intermittent plantar stimulation (output voltage is 20-40 V, the interval time of voltage is 2-5 V, 30 min per day) [26] clip tail (clamp the tip of the rat tail within 2-3 cm, each to the bite, Fighting, treatment once a day for 5 min) [27]; give noise to rats with 100 db with fast rhythm of playback music, continues to play 2 hours per day for 10-13 days [28].

TNBS enema was used to induce UC model, the rats fasted for 1 day but were allowed to drink water freely. After anesthetized by 10% chloralhydrate solution (According to the weight of 0.3 ml/100 g, an intraperitoneal injection), a catheter 3 mm in diameter was slowly inserted about 8 cm into the anus, then 50% ethanol solution with 70 mg/kg TNBS was slowly injected.

Drugs were slowly administrated to make sure that the liquid did not flow out, and the rats were kept upside down for 1 min after the injection, in order to facilitate the distribution of TNBS in the gut [29].

With the above method to establish the complex factor induced UC model. At the same time every day morning and night per day enema to UC rats 1.35 ml/100 g dose of KJA total of 10 days. After established UC models, fasting 3 days, during these days a dose of 2 ml/100 g dose lavage of 20% glucose is given twice a day in the morning and evening to fill the stomach. Every cage supply 200 ml of 5% glucose water. For negative control groups, using distilled water instead of KJA.

**Collection of tissue samples:** The rat was anesthetized and the abdomen opened and the blood was taken from the abdominal aorta. Take colon tissue from the anus 5 cm-8 cm, cut open along the mesenteric margin, and gently wash the intestine with saline solution (feces, secretions, and blood Liquid), rinse clean and unfold the colon, surface of the mucosa, to observe the colonic mucosal injury [30].

For histopathological observation, the lesion area was taken and immediately placed in the 40 g/L formaldehyde solution fixed 24 h, the
paraffin bag is buried, stain with HE dye. The colon tissues placed and kept in -80°C.

**Total RNA extraction:** The total RNA was extracted from the colon tissues by the Trizol (Invitrogen, Gaithersburg, MD, USA) one-step method. After enrichment of RNA by isopropyl alcohol precipitation, total RNA was column purified by using a NucleoSpin® RNA clean-up kit (MACHEREY-NAGEL, Germany). The concentration and purity of total RNA were determined by a spectrophotometer, and the quality assessment was conducted by the integrity of 28S and 18S rRNA.

**Primer design:** IL-1α, IL-1β, IL-10, IL-17, TNF-α primers designed 2-3 pairs of primers are designed for mRNA and hnRNA candidates for candidate genes (Tables 1 and 2). Applied the method of quantitative RT-PCR to detect the expression level of hnRNA and mRNA in the five inflammation factor such as IL-1α, IL-1β, IL-10, IL-17, TNF-α in normal group, UC model group, KJA group and negative control group. The condition of qPCR reaction: degeneration at 95°C for 10 min, 95°C for 15 s, each primer annealed temperature is 1 min.

| Gene   | Primer sequences          | Length(bp) | Anneling T(°C) |
|--------|---------------------------|------------|----------------|
| IL-1α  | Forward: ACATGTATGCCTACTCATCAGG<br>Reverse: TCCCGAAATCTCCTTCAGCCAC  | 127        | 57             |
| IL-1β  | Forward: AGGAGAGACGAAGCAAACGACAA<br>Reverse: GTTTGAGTCACCTCCTCCA  | 122        | 57             |
| IL-10  | Forward: GCTCAGACACTGCTATGTTG<br>Reverse: GTTGTCAGCTGTTGCTTTC  | 183        | 67             |
| IL-17  | Forward: TCCATGTTGCTGATGCTT<br>Reverse: AGGTTGAAGTGAACCGGTG  | 208        | 67             |
| TNF-α  | Forward: CACCAACGCTCTCTCTGTCA<br>Reverse: GGCGTTGCTACCGAAGGGT  | 143        | 59             |
| β-actin| Forward: AGCCATGTGCTAGCTGGCAG<br>Reverse: ACCCTCATAGATGGGCACAG  | 115        | 57             |

Table 1: The primer list of qRT-PCR (for mRNA).

**Statistical analysis**

Data analysis was performed using SPSS software v13.0. Values were reported as mean ± SD. ANOVA followed by Student's t-test was used for multiple comparisons of the data. Statistical significance was set at an alpha value of p<0.05.

| Gene   | Primer sequences          | Length(bp) | Anneling T(°C) |
|--------|---------------------------|------------|----------------|
| IL-1α  | Forward: ACGGCTAAGTTTCAATCA<br>Reverse: CACCGAAGACCTTTACAT  | 270        | 58             |
| IL-1β  | Forward: ACTGTTCCAGACCCATAC<br>Reverse: CACTCTGAAAGTGGGAGG  | 288        | 58             |
| IL-10  | Forward: TCCCTTTTCTCTACTGAGG<br>Reverse: ACACCTTTGTCTTGGAGG  | 232        | 60             |
| TNF-α  | Forward: GTGCTGCTGCTCTCTGCTC<br>Reverse: CACGCTTGTGCTTGCTTCC  | 298        | 62             |
| IL-17  | Forward: AGACTACCTCACCCTCTCCAC<br>Reverse: TACAGGCTTGTGGGAGTAGAGG  | 219        | 63             |

Table 2: The primer list of qRT-PCR (for hnRNA).

**Results**

Compared with the normal group, in UC group general state has undergone obvious changes, these characterize are indicated in Table 3A and 3B. Compare to negative control group, in KJA group, the rat hair becoming lustrous, mental state is better, have more activity, eat regular, the food quantity is normal, formed defecate.

The mortality was obviously lower than negative control group (Figures 1A and 1B). Those symptoms in negative control group are appearing seriously.
The KJA on the ulcerative colitis is very effectiveness (Figures 2A-2D). Observed each group's colon tissue can found the therapeutic effect of KJA. To investigate mucosal inflammation, we performed H&E staining and demonstrated the KJA group and negative control group pathological results.

Colon tissue in KJA group (Figures 3A and 3C), all the colonic mucosal epithelial part defect, some glandular arrangement rules, the structure is clear, inflammatory cells infiltration of sub mucosa, and part of the mucous membrane hyperplasia of thickening, a small amount of mucosal epithelium was mild hyperplasia, ulcer area is narrow.

In the negative control group (Figures 3B and 3D), mucosal epithelial cell necrosis, inflammatory cells infiltrating into the sub mucosa and muscularis, there are a large number of neutrophil infiltration, and accompanied by obvious hyperaemia edema.

### Table 3A: The inflammation scores of colon tissue from KJA and negative control group.

| GROUP          | N  | Level of degree | X2 | P   |
|----------------|----|-----------------|----|-----|
| KJA group      | 9  | 0, 1, 2, 3, 4   | 15 | 0.02 |
| Negative control group | 6  | 0, 0, 0, 0, 6   |    |     |

**Histological parameters and description**

| Score | Observe the colon tissue by naked eye | Epithelial cell | Inflammatory cell |
|-------|--------------------------------------|-----------------|-------------------|
| 0     | no damage                            | normal          | not infiltration  |
| 1     | mildly congested, edema              | loss of goblet cells | infiltrate to bottom of the mucosal base, fibrosis |
| 2     | intestine swollen, mucosa is coarse  | loss large part of goblet cells | a large number of chronic inflammatory |
|       | and granular                         |                 |                   |
| 3     | intestine swollen with necrosis and  | loss of part of goblet, columnar cell | the mucosa and has a large number of neutrophils infiltrating |
|       | ulcer                                |                 |                   |
| 4     | hyperemic, mucosal necrosis and ulcer| loss of part of goblet, columnar cell | mucosal necrosis, colon wall necrosis, ulceration is greater than 1 cm |

**Table 3B:** The inflammation score and its description.

Using western blot analysis, detect phosphorylation levels of each groups of the nucleus NF-kB and cytoplasm HuR proteins. Compared with the control group, phosphorylation levels of NF-kB and HuR proteins are obviously increased in UC group and negative control group. But both of NF-kB and HuR proteins are decreased in KJA group. The evaluation results of their expression levels in each experiment groups showed in Figure 4A and 4B.

hnRNA expression and mRNA expression of several cytokines such as IL-1α, IL-1β, IL-10, IL-17 and TNF-α in all experimental model groups (Tables 4 and 5). Also the expression levels of each genes hnRNA in all experimental groups in Figure 5. Pro-inflammatory cytokines are implicated in ulcerative colitis, with anti-inflammatory cytokine therapy are potential targets for remission in ulcerative colitis [19].

Compare the expression levels of hnRNA and mRNA of IL-1α, IL-1β, IL-10, IL-17 and TNF-α which is resulted in both of hnRNA and mRNA expression levels are increased, among them the mRNA expression level are significantly higher than hnRNA expression level. The role of post-transcriptional level is very important for the further investigation of treatment of inflammation diseases.
Figure 2: Colon tissue morphology of different group. Colon tissue specimens the mucosal hyperemia edema and ulcer formation, the colon tissue and surrounding tissue adhesion, whole intestinal wall necrosis (B); KJA effects on the ulcerative colitis colon tissue, does not appear in the erosion, the range of ulcer significantly reduced, ulcer healing, mucosal congestion of blood edema decreased (C); In the negative control group (D) the colon thickens, wall of the intestine is highly congested with blood, mucosal necrosis and ulcer formation, erosion, the range of ulcer is larger.

Figure 3: HE staining of colon tissue. Colonic mucosa epithelial part defect, the structure is clear, inflammatory cells infiltration of sub mucosa, some mucosa epithelial hyperplasia and granulation tissue formation, and part of the mucous membrane hyperplasia of thickening, a small amount of mucosal epithelium was mild atypical hyperplasia, ulcer area is narrow (A, C). Mucosal epithelial cell necrosis, glands fossae disappear, inflammatory cells infiltrating into the sub mucosa and muscularis, hyperaemia edema (A, D).

Figure 4A: protein phosphorylation levels of in Nuclear NF-κB and cytoplasmic HuR. lane 1, 2 for Normal group; lane3, 4 for UC group; lane 56 for UC with KJA treated group lane; 7,8 for UC negative group. The phosphorylation levels of nucleus NF-kB and cytoplasm HuR protein are increased in UC group colon tissue, and in KJA group there phosphorylation levels are decreased.

Figure 4B: Semi-quantitative grayscale analysis of NF-κB and HuR protein phosphorylation levels. A: normal B: UC; C: with KJA; D: UC negative.

Figure 5: The differences in gray analysis were expressed in the differences of the genes IL-1α, IL-1β, IL-10, IL-17, and TNF-α. 1: Normal; 2: UC; 3: UC with KJA; 4: UC negative.

Discussion

Ulcerative colitis is a chronic inflammatory bowel disease that affects millions of people worldwide, and is significantly associated colorectal cancer risk [31-33]. At the present, the explanation of pathogenesis of UC is not yet complete, and the therapeutic effect is not ideal.

Recently the immunomodulatory agent originating from herbal medicine represents a promising approach for UC therapy, as shown by the variety of clinical trials and experimental studies currently
underway [34]. Traditional Chinese medicinal enemas can effectively inhibit regional mucosal inflammatory factors and improve disorders associated with immunity [35].

In our previous study investigated and indicated treatment of UC rats with KJA significantly reduced the NF-κB and IL-1β in the intestinal mucosa as compared with UC rats treated with water. In previous research, KJA can significantly effect to the morphology and pathological changes of colon tissue in UC rats [20].

In UC group, rats significantly decreased activity, loss of weight, like crouching, the hair color to dim and lackluster, mental depression, appear the loose, mucous, purulent and bloody stool. In previous study, UC model rats colon inflammatory cells infiltration, the decrease of microvilli in columnar cells, increase of mucous of granula in goblet-cells and mitochondrial vacoules in colon tissue on histomorphology [36].

Colon tissue specimens in UC model, the mucosal hyperemia edema and ulcer formation and the shape of linear or focal, whole intestinal wall necrosis, mucosa thickening, and intestinal stenosis. In the negative control group, rats treated with water and the symptoms clearly same as UC group rats. The colon thickens, wall of the intestine is highly congested with blood, mucosal necrosis and ulcer formation, erosion, appeared big lump.

These pathological symptoms are in KJA group, the range of ulcer significantly reduced, did not see colon thickened, color returns to normal. That shows the KJA could affect to improve the colon tissue and cure the inflammation of ulcer.

The pathological alterations in colon of KJA group were manifested in Figures 3A and 3C. Compared to the negative control group, KJA group significantly showed the structure is clear, ulcer is narrowed, part of the mucous membrane hyperplasia. As we analyzed above, the KJA group exhibited a greater inhibitor effect of inflammation score and symptoms.

In the present study, KJA significantly ameliorated inflammatory cell infiltration in KJA group that compared with complex factors induced UC group. The Xipayi Kuijiean (KJA), which was made from gallic, was used for enema to rats model groups, can treat many kinds of ulcers and the usual medicines for inflammation [37]. After gallic tannis bind to proteins, except for the convergence, there are anti-inflammatory, analgesia, antivirus, anti-immunity, decreasing blood lipid concentration and blood sugar and other functions [38].

The phosphorylation levels of nucleus NF-kB and cytoplasm HuR protein are increased in UC group colon tissue, and in KJA group they are decreased. This result indicate rats were treated with KJA can be suppress the nucleus NF-kB and cytoplasm HuR protein activation or phosphorylation, than might regulate the transcription activity of cytokine and the stability of mRNA, consequently be affected the gene expression of cytokines. This analysis showed that KJA markedly suppressed phosphorylation of NF-kB p65 and also inhibited NF-kB p65 translocation to the nucleus in the inflamed colon tissue [39].

Compared the normal group and UC group, hnRNA expression level increased in cytokines of IL-1α, IL-1β, IL-10 and TNF-α, and these differences were statistically significant (P<0.05). Compared with KJA group and the negative control group, hnRNA expression level of IL-10 and TNF-α are decreased in KJA group, the difference was statistically significant (P<0.05), and hnRNA expression level of IL-1α, IL-β, IL-17 is no statistical significance (P>0.05).

The mRNA expression was significantly increased of IL-1α, IL-1β, IL-10, IL-17 and TNF-α in UC group (P<0.05). In KJA group, the mRNA expression level of IL-1α, IL-1β, IL-17, TNF-α was significantly decreased (P<0.05). But the mRNA expression level of IL-10 was not significantly changed (P>0.05).

IL-1 is an important medium that is closely related to immune regulation, inflammation and tissue injury. The important role of IL-1 in immunity and inflammation induced with many effects in the process of protein expression, for example: chemokines, cytokines, nitric oxide synthase (iNOS), matrix metalloproteinases (MMPS) and an enzyme called cyclooxygenase 2 (cox-2), etc [40].

IL-10 reduce the Major histocompatibility complex (MHC), increase the soluble TNF-α, IL-1Ra and other several anti-inflammatory protein expression, inhibition of IL-1β, IL-4, IL-5, IL-6, IL-2, IL-8, TNF-α, IFN-γ, and the secretion of iNOS and COX-2 and adjust the differentiation and proliferation of immune cells, thereby limiting and termination of the inflammatory response [42-56].

### Table 4: The hnRNA differential expression of IL-1α, IL-1β, IL-10, IL-17, TNF-α in colon tissues from different group. NOTE *compare with the normal groups P<0.05, compare with the negative groups P<0.05.

| Genes       | IL-1α | IL-1β | IL-10 | IL-17 | TNF-α |
|-------------|-------|-------|-------|-------|-------|
| Normal      | 0.55 ± 0.71 | 0.99 ± 0.71 | 0.90 ± 0.22, 0.92 ± 0.53 | 1.01 ± 0.62 |       |
| U C group   | 14.79 ± 7.08* | 28.03 ± 11.85* | 6.63 ± 4.98*, 0.36 ± 0.17 | 2.27 ± 0.96* |       |
| UC treated with KJA | 0.46 ± 0.38 | 0.61 ± 0.52 | 0.90 ± 0.38*, 0.53 ± 0.52 | 0.75 ± 0.31* |       |
| UC negative group | 1.34 ± 1.23 | 0.93 ± 0.90 | 1.91 ± 0.62, 0.81 ± 0.77 | 1.92 ± 0.51 |       |

### Table 5: The mRNA differential expression of IL-1α, IL-1β, IL-10, IL-17, TNF-α in colon tissues from different group NOTE *compare with the normal groups P<0.05, compare with the negative groups P<0.05.

| Genes       | IL-1α | IL-1β | IL-10 | IL-17 | TNF-α |
|-------------|-------|-------|-------|-------|-------|
| Normal      | 0.83 ± 0.19 | 1.05 ± 0.09 | 1.16 ± 0.25 | 1.09 ± 0.44 | 0.69 ± 0.35 |
| U C group   | 114.45 ± 50.59* | 306.06 ± 72.25* | 11.97 ± 6.10* | 5.62 ± 4.27* | 4.93 ± 2.99* |
| UC treated with KJA | 0.48 ± 0.40* | 0.57 ± 0.41* | 1.15 ± 0.83 | 0.30 ± 0.29* | 0.549 ± 0.36* |
| UC negative group | 1.45 ± 0.94 | 2.04 ± 0.89 | 1.24 ± 0.56 | 1.19 ± 0.62 | 1.37 ± 0.56 |
IL-17 is an important cytokines induced by T cells and promotes inflammation. The mRNA expression level significantly increased of IL-17 in the UC group, but hnRNA expression level of was not statistically significant (P>0.05). That might be the stability of mRNA expression of IL-17 was improved by raising the level of HuR phosphorylation in the UC group. It can also be synergistic with TNF-α to amplify its inflammatory effect [46-47].

Our observations showed KJA inhibits the expression of pro-inflammatory cytokines, that indicate the regulation of post-transcriptional level and HuR are important to understanding the mechanism of UC. Thereby we conclude that KJA exerts its therapeutic effects by inhibit the HuR phosphorylation level in the UC.

Compare the hnRNA and mRNA expression levels between the UC group and KJA group shown by Table 6. The mRNA expression levels were significantly higher than hnRNA levels in IL-1α, IL-1β, IL-17 (P<0.05). This study reveals that the mRNA expression levels of IL-1α, IL-1β and IL-10 in UC group were all increased. In particular, the increase of mRNA expression of IL-1α, IL-1β in UC group higher than that of normal group, and these two genes expression significantly higher than IL-10.

This result indicates the increased phosphorylation level of HuR causes colon tissue inflammation and lead to increase the mRNA amount in UC group. Treated with KJA, the phosphorylation level of HuR and mRNA decreased, resulted in no any difference on expression level between the mRNA and hnRNA, shown as Table 6. In this study, the expression level of mRNA higher than hnRNA of IL-1α, IL-1β which might indicate HuR play an important regulatory role on mechanism of UC.

Table 6: Compare the hnRNA and mRNA expression levels of IL-1α, IL-1β, IL-10, TNF-α in colon tissues from UC group treated with KJA.

| Genes  | mRNA | hnRNA | P   | UC treated with KJA | p   |
|--------|------|-------|-----|---------------------|-----|
| IL-1α  | 114.45 ± 50.49 | ± 14.79 ± 7.08 | 0.03 | 0.48 ± 0.40 | ± 0.46 ± 0.38 | ± 0.93 |
| IL-1β  | 306.05 ± 72.25 | ± 28.03 ± 11.85 | 0 | 0.61 ± 0.52 | ± 0.93 ± 0.90 | ± 0.48 |
| IL-10  | 11.97 ± 6.10 | 6.63 ± 4.96 | 0.1 | 1.15 ± 0.83 | ± 0.90 ± 0.39 | ± 0.46 |
| TNF-α  | 3.90 ± 1.80 | 2.27 ± 1.96 | 0.14 | 0.55 ± 0.36 | ± 0.75 ± 0.31 | ± 0.27 |
| IL-17  | 5.61 ± 4.27 | 0.36 ± 0.16 | 0.04 | 0.30 ± 0.29 | ± 0.53 ± 0.52 | ± 0.4 |

In conclusion, gene expression of IL-1α, IL-1β and IL-17 and mRNA stability might be regulated by HuR; The TNF-α expression activity mainly based on NF-kB regulation. Since in the further study, the role of HuR post-transcriptional regulating mRNA is very important than transcription regulation of NF-kB. HuR can lead to another new way to relieve inflammation in UC, and is suitable method for curing UC.

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