Melatonin attenuates acute pancreatitis-associated lung injury in rats by modulating interleukin 22

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Abstract

AIM: To investigate whether therapeutic treatment with melatonin could protect rats against acute pancreatitis and its associated lung injury.

METHODS: Seventy-two male Sprague-Dawley rats were randomly divided into three groups: the sham operation (SO), severe acute pancreatitis (SAP), and melatonin treatment (MT) groups. Acute pancreatitis was induced by infusion of 1 mL/kg of sodium taurocholate (4% solution) into the biliopancreatic duct. Melatonin (50 mg/kg) was administered 30 min before pancreatitis was induced, and the severity of pancreatic and pulmonary injuries was evaluated 1, 4 and 8 h after induction. Serum samples were collected to measure amylase activities, and lung tissues were removed to measure levels of mRNAs encoding interleukin 22 (IL-22) and T helper cell 22 (Th22), as well as levels of IL-22.

RESULTS: At each time point, levels of mRNAs encoding IL-22 and Th22 were significantly higher ($P < 0.001$) in the MT group than in the SAP group (0.526 ± 0.143 vs. 0.156 ± 0.027, respectively, here and throughout, after 1 h; 0.489 ± 0.150 vs. 0.113 ± 0.014 after 4 h; 0.524 ± 0.168 vs. 0.069 ± 0.013 after 8 h; 0.378 ± 0.134 vs. 0.122 ± 0.015 after 1 h; 0.205 ± 0.041 vs. 0.076 ± 0.019 after 4 h; 0.302 ± 0.108 vs. 0.045 ± 0.013 after 8 h, respectively) and significantly lower ($P < 0.001$) in the SAP group than in the SO group (0.156 ± 0.027 vs. 1.000 ± 0.010 after 1 h; 0.113 ± 0.014 vs. 1.041 ± 0.235 after 4 h; 0.069 ± 0.013 vs. 1.110 ± 0.213 after 8 h; 0.122 ± 0.015 vs. 1.000 ± 0.188 after 1 h; 0.076 ± 0.019 vs. 0.899 ± 0.125 after 4 h; 0.045 ± 0.013 vs. 0.991 ± 0.222 after 8 h, respectively). The mean pathological scores for pancreatic tissues in the MT group were significantly higher ($P < 0.01$) than those for samples in the SO group (1.088 ± 0.187 vs. 0.488 ± 0.183 after 1 h; 2.450 ± 0.212 vs. 0.469 ± 0.242 after 4 h; 4.994 ± 0.184 vs. 0.513 ± 0.210 after 8 h), but were significantly lower ($P < 0.01$) than those for samples in the SAP group at each time point (1.088 ± 0.187 vs. 1.969 ± 0.290 after 1 h; 2.450 ± 0.212 vs. 3.344 ± 0.386 after 4 h; 4.994 ± 0.184 vs. 6.981 ± 0.301 after 8 h). The severity of SAP increased significantly ($P < 0.01$) over time in the SAP group (1.088 ± 0.187 vs. 2.450 ± 0.212 between 1 h and 4 h after inducing pancreatitis; and 2.450 ± 0.212 vs. 4.994 ± 0.184 between 4 and 8 h after inducing pancreatitis).

CONCLUSION: Melatonin protects rats against acute pancreatitis-associated lung injury, probably through the upregulation of IL-22 and Th22, which increases the innate immunity of tissue cells and enhances their regeneration.

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INTRODUCTION

Severe acute pancreatitis (SAP) is an inflammatory disease characterized by tissue edema, acinar necrosis, hemorrhage, and fat necrosis in the pancreas. Subsequent entry of proteolytic enzymes and vasoactive mediators into the systemic circulation always leads to multi-organ complications[1,2]. Pulmonary dysfunction is the most prominent manifestation of extra-abdominal organ dysfunction in pancreatitis, and its severity ranges from mild oxygenation abnormalities to severe acute respiratory distress syndrome (ARDS)[3]. Respiratory failure is associated with approximately 60% of the deaths that occur within the first week of pancreatitis[3-5]. Thus, the success of efforts to identify new therapeutic strategies for SAP will likely depend largely on understanding the molecular pathogenesis of damage to both the pancreas and the lungs.

Interleukin 22 (IL-22) is a member of the IL10 family of cytokines. It is an important effector of activated IL-22-producing T helper cells 22 (Th22), Th1 cells, and Th17 cells, as well as cytotoxic T-cell subsets, γδT cells, natural killer (NK), and NKT cells[5-7]. Activated T cells and NK cells are major sources of IL-22, and the highest levels of IL-22 expression have been detected in CD4+ memory cells. In contrast, IL-22 is not produced by either resting or activated monocytes, or B cells[8]. Like all other members of the IL10 family, IL-22 exerts its biological effects via heterodimeric transmembrane receptor complexes that each comprise of a type-1 and a type-2 receptor chain. These receptor chains belong to the cytokine receptor family class 2[9].

Since its discovery in 2000, several research groups have studied the unique biological activities of IL-22 and its significance in diseases. IL-22 protects tissues from damage, enhances tissue repair, maintains tissue integrity, increases innate immunity, and promotes anti-microbial defenses[8,9]. Therapy involving IL-22 improves liver regeneration after 70% hepatectomy[11] and also ameliorates high-fat diet-induced liver lipogenesis and hepatic steatosis[2,23]. Similarly, therapy with IL-22 counteracts the destructive nature of inflammatory bowel disease and ulcerative colitis[12,13] and prevents lung inflammation and fibrosis[14,15].

The recently discovered Th22 subset of T cells, which differs from Th17 and other known T-cell subsets based on its unique function[15], produces cytokines such as IL-22, IL-26, and IL-13. Th22 cells express several fibroblast growth factors, are associated with epidermal repair responses, and synergize with tumor necrosis factor α to induce a characteristic Th22 signature in keratinocytes[16]. Moreover, Th22 cells may regulate epidermal responses in inflammatory skin diseases and could be involved in unknown pathways that control tissue immunity and remodeling[17].

Although most melatonin is produced by the pineal gland, it can also be secreted by extrapineal organs and tissues such as the gastrointestinal tract, the retina and lens, skin, immune and hematopoietic cells, and some reproductive organs[16,17]. Several studies have demonstrated the beneficial effects of melatonin on SAP. As early as 1999, melatonin was reported to attenuate pancreatic edema and lipid peroxidation in caerulein-induced SAP[18]. Subsequent studies confirmed that pretreatment with melatonin significantly decreases all investigated inflammatory parameters associated with SAP[19,20]. Moreover, melatonin exerts an anti-inflammatory effect by inhibiting nuclear factor kappa B, a transcription factor with a central role in the development of inflammatory diseases. The present study was undertaken to examine the role of IL-22 in the pathogenesis of AP-induced lung damage, as well as the capacity of melatonin to protect against AP-associated lung injury.

MATERIALS AND METHODS

Animals

A total of 72 clean-grade healthy adult male Sprague-Dawley (SD) rats, each weighing 200-250 g, were obtained from the Experimental Animal Center of Wenzhou Medical College, Wenzhou, China. All animals were fed standard rat chow and had unlimited access to water under conditions that provided a constant room temperature of 25 °C and a 12-h/12-h day/night cycle. All animals were acclimated for at least a week before experimentation. Animals were deprived of rat chow for 12 h before experimentation but were allowed unlimited access to water throughout the experimental period. All procedures were performed in accordance with the guidelines for animal experiments of Wenzhou Medical College, Wenzhou, China.

Animal groups and procedures

SD rats were anesthetized by intraperitoneal (i.p.) injection of 10% chloralhydrate (2 mL/kg; Solarbio, Beijing, China) and were randomly assigned to a sham operation group (SO group; n = 24), a SAP group (n = 24), or a melatonin treatment group (MT group; n = 24). For each animal in the SAP group, a laparotomy was performed through a midline incision, and 1 mL/kg of 4% sodium taurocholate (Sigma, St. Louis, United States) was retrogradely injected into the biliopancreatic duct through the papilla using a segmental epidural catheter via a microinjection pump at a rate of 0.2 mL/min. A microclip was placed in the hepatic portion of the biliopancreatic duct to avoid reflux before injection of sodium taurocholate.
Rats in the SO group were subjected to the same surgical procedure but without infusion of 4% sodium taurocholate. In the MT group, melatonin (50 mg/kg body weight; Sigma) was administered i.p. 30 min before the injection of taurocholate. After each operation, the abdomen of each rat was closed in two layers. All surgical procedures were performed using sterile techniques.

The rats were killed by exsanguination at defined time points (1, 4 and 8 h; n = 8 for each time point) after the surgical procedure. Postcava puncture was used to obtain 3 mL blood from each animal to assay levels of serum amylase. The concentration of serum amylase (U/L) was assayed using a fully automatic biochemical analyzer (Hitachi, Tokyo, Japan). Lung tissues were harvested immediately before death, and tissue samples were fixed in 4% paraformaldehyde for immunohistochemical analysis. Other tissues were removed and stored in liquid nitrogen for the determination of IL-22 and Th22 mRNA levels using real-time reverse-transcription polymerase chain reaction and IL-22 protein levels using enzyme-linked immunosorbent assay (ELISA).

Histological analysis and pathological scores of pancreatic tissues

Samples of lung and pancreatic tissues were fixed in 40 g/L formaldehyde, embedded with paraffin wax, sectioned (4 μm thick), stained with hematoxylin and eosin (H and E), and observed using light microscopy. Tissue alteration was assessed (20 fields/section) by an experienced histologist who was blinded to the experimental protocol. Ten randomly selected visual fields in each pathological section were observed (H and E staining; × 400) using a high-powered microscope (CX31, Tokyo, Japan) and scored by two experienced pathologists using established standards[19], respectively. The maximum possible score for each visual field was 12, and the pathological score for each section was determined by calculating the mean of the scores from the two pathologists for the 10 visual fields.

Real-time, fluorescence-based quantitative polymerase chain reaction

Total RNA was extracted from lung tissues using TRIzol reagent (Invitrogen, Carlsbad, United States), and cDNA was synthesized using the first strand cDNA synthesis kit (MBI Fermentas, Burlington, Canada). mRNAs encoding IL-22 and Th22 were detected using the PCR mix kit (Daweike, Shanghai, China), and real-time quantitative PCR was performed using the ABI 7500 Sequence Detection System (Applied Biosystems Inc, Carlsbad, United States). For PCR, we used the following primers for IL-22 (forward: 5’-GCCAGCTGCTGCTTCTCGT-3’, reverse: 5’-CTGGCCTCCTTGGCCAGCAT-3’), Th22 (forward: 5’-GGGTCTCAGGGTCTGGCCGC-3’, reverse: 5’-CCTCAGTTTACCGAGAACCCCA-3’), and β-actin (forward: 5’-CGGTTCACCCGGCAGTAA-3’, reverse: 5’-CGACGGAGCAGCCAGCGAT-3’, Generay, Shanghai, China). The cDNA was de-natured at 95 °C for 5 min and amplified over the course of 40 cycles, each involving 95 °C (15 s), 60 °C (45 s), and 72 °C (60 s), with a final extension at 72 °C (5 min). The samples were tested in triplicate, and the expression of IL-22 and Th22 mRNA was calculated using the $2^{-\Delta\Delta CT}$ method.

**Sandwich enzyme-linked immunosorbent assays**

Concentrations of IL-22 were measured using commercially available sandwich ELISA kits (all from R and D Systems, Rapidbio, United States). IL-22 levels were assayed using microtiter wells coated with purified rat IL-22 and a solid-phase IL-22-specific antibody. The horseradish peroxidase (HRP)-labeled secondary antibody was detected with the tetramethylbenzidine substrate solution, which creates a blue reaction product. The HRP enzyme-catalyzed reaction was terminated by the addition of a sulfuric acid solution, and the color change was measured spectrophotometrically at a wavelength of 450 nm. The concentration of IL-22 in the samples was then determined by comparing the $A_{450\ nm}$ of the samples to a standard curve.

**Statistical analysis**

Statistical analysis were carried out using the SPSS software (ver. 15.0; SPSS Inc., Chicago, IL, United States). All data are expressed as the mean ± SD. One-way analysis of variance were used to investigate differences among the SO, SAP, and MT groups at each time point, and post hoc comparisons were performed between samples in the SO and SAP groups to check for statistical significance. Differences were considered significant at P < 0.05.

**RESULTS**

**Histopathological examination of lung tissues and pathological scores of pancreatic tissues after SAP induction**

The appearance of lung tissue samples collected 1 h (Figure 1A), 4 h (Figure 1B), and 8 h (Figure 1C) after the induction of SAP was much more severely affected than the appearance of tissue from animals in the SO group (Figure 1D), which tended to remain morphologically normal even 8 h after surgery. Microscopic examination of lung tissue from animals in the SAP group provided extremely clear evidence of edema, hemorrhage complicated by microthrombosis, cell degeneration, and inflammation (Figure 1A-C). The extent of damage observed in the lungs of SAP animals was largely alleviated in lung tissues of animals in the MT group (Figure 1E-G).

Pancreatic tissue samples were also collected and analyzed from each group of animals. At each time point, the mean pathological score for pancreatic tissues in the MT group was significantly higher (P < 0.01) than for samples in the SO group, but significantly lower (P < 0.01) than for samples in the SAP group at each time point. The severity of SAP increased significantly over time in
the SAP group \((P < 0.01, \text{Figure } 1H)\).

**Serum amylase levels after SAP induction**

Levels of serum amylase at each time point were markedly elevated in the SAP group relative to the SO \((P < 0.01)\) and MT \((P < 0.05)\) groups (Figure 2). In addition, the levels of amylase in the MT group increased compared with those in the SO group. In the SAP group, amylase levels increased over time, indicating the development of SAP. There was no significant difference between the levels of serum amylase at the three time points in the SO group (Figure 2).

**Expression of IL-22 and Th22 mRNAs after SAP induction**

Levels of the IL-22 and Th22 mRNAs in lung tissues were significantly decreased within 1 h after the induction of SAP \((P < 0.01; \text{Figure } 3A \text{ and } B)\), and levels of both mRNAs remained low during the remaining time points. Importantly, this decrease in both mRNAs in the SAP group relative to the SO group was partly rescued in the MT group. Differences between levels in the SAP group and the MT group were significant \((P < 0.01)\), and expression differed significantly between the MT group and the SO group \((P < 0.01, \text{Figure } 3A \text{ and } B)\).

**Levels of IL-22 in lung tissue after SAP induction**

The level of IL-22 in lung tissue in the MT group was significantly lower than that in the SO group and significantly higher than that in the SAP group at each time point \((P < 0.01; \text{Figure } 4)\). The level of IL-22 in the lung tissue samples decreased significantly over time in the SAP group \((P < 0.01)\), although IL-22 levels did not differ significantly between each pair of time points in the SO group.

**DISCUSSION**

Patients with SAP and the associated multiple-organ failure in which it frequently results\(^{[20]}\) require admission to a high-dependency or intensive-care unit. Despite recent progress in revealing the mechanisms that underlie SAP, which include inflammation, cell injury, and cell death, the exact pathogenesis of SAP is not fully understood. The uncontrolled generation of reactive oxygen species (ROS) during SAP causes oxidative damage, both in the pancreas and distant organs. The lung is especially vulnerable to damage by ROS because it possesses the largest endothelial surface area of any organ in the body\(^{[20]}\).

IL-22 is a member of the IL-10 cytokine family, which additionally comprises IL-10, IL-19, IL-20, IL-24,
In this study, we measured the levels of mRNAs encoding IL-22 and Th22 in rat pulmonary tissues from rats in the sham operation, severe acute pancreatitis, and melatonin treatment groups. A: Expression of IL-22 mRNA 1, 4 and 8 h after induction of SAP. B: Expression of Th22 mRNA 1, 4 and 8 h after induction of SAP. Levels of mRNAs encoding IL-22 and Th22 decreased with continued progression of SAP (P < 0.01). Compared with the SO group, the levels of IL-22 and Th22 mRNAs in pulmonary tissues from the MT group were 0.3-4 times lower (P < 0.01). Figure 4 Levels of interleukin 22 in lung tissue samples from rats in the sham operation, severe acute pancreatitis, and melatonin treatment groups. Levels of IL-22 at 1, 4 and 8 h after induction of SAP. Levels of IL-22 decreased with continued progression of SAP (P < 0.01). Compared with the SO group, the levels of IL-22 in the MT group were 0.3-4 times lower (P < 0.01). Figure 5 Levels of interleukin 22 in lung tissue samples from rats in the sham operation, severe acute pancreatitis, and melatonin treatment groups. Levels of IL-22 at 1, 4 and 8 h after induction of SAP. Levels of IL-22 decreased with continued progression of SAP (P < 0.01). Compared with the SO group, the levels of IL-22 in the MT group were 0.3-4 times lower (P < 0.01). Figure 6 Levels of interleukin 22 in lung tissue samples from rats in the sham operation, severe acute pancreatitis, and melatonin treatment groups. Levels of IL-22 at 1, 4 and 8 h after induction of SAP. Levels of IL-22 decreased with continued progression of SAP (P < 0.01). Compared with the SO group, the levels of IL-22 in the MT group were 0.3-4 times lower (P < 0.01).
suggests that the alleviation of acute pancreatitis-associated lung injury by melatonin is mediated, at least in part, by upregulation of IL-22 and Th22 activities.

In summary, this study indicates, for the first time, that the downregulation of pulmonary IL-22 and Th22 may be associated with the pathogenesis of acute pancreatitis-associated lung injury. Melatonin, functioning as a protectant and antioxidant, may reduce lung injury by increasing IL-22 and Th22 activity.

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