Original Article

Brazilian green propolis suppresses acetaminophen-induced hepatocellular necrosis by modulating inflammation-related factors in rats

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Abstract: Propolis is a resin-like material produced by honey bees from bud exudates and sap of plants and their own secretions. An ethanol extract of Brazilian green propolis (EEBGP) contains prenylated phenylpropanoids and flavonoids and has antioxidative and anti-inflammatory effects. Acetaminophen (*N*-acetyl-*p*-aminophenol; APAP) is a typical hepatotoxic drug, and APAP-treated rats are widely used as a model of drug-induced liver injury. Oxidative stress and inflammatory reactions cause APAP-induced hepatocellular necrosis and are also related to expansion of the lesion. In the present study, we investigated the preventive effects of EEBGP on APAP-induced hepatocellular necrosis in rats and the protective mechanism including the expression of antioxidative enzyme genes and inflammation-related genes. A histological analysis revealed that administration 0.3% EEBGP in the diet for seven days reduced centrilobular hepatocellular necrosis. EEBGP administration did not significantly change the mRNA expression levels of antioxidant enzyme genes in the liver of APAP-treated rats but decreased the mRNA expression of cytokines including *II10* and *II1b*, with a significant difference in *II10* expression. In addition, the decrease in the mRNA levels of the *II1b* and *II10* genes significantly correlated with the decrease in the percentage of hepatocellular necrosis. These findings suggest that EEBGP could suppress APAP-induced hepatocellular necrosis by modulating cytokine expression. (DOI: 10.1293/tox.2018-0027; J Toxicol Pathol 2018; 31: 275–282)

Key words: acetaminophen, antioxidative enzyme, Brazilian green propolis, chemokine, cytokine, hepatocellular necrosis

Introduction

Propolis is a resin-like material produced by honey bees by mixing their secretions with bud exudates and sap of plants. The constituents of propolis vary depending on the vegetation in the region where it is produced, which could alter its biological activities¹. Brazilian green propolis (BGP) mainly consists of *Baccharis dracunculifolia* D.C. (Asteraceae) exudates, and an ethanol extract of BGP (EE-BGP) contains prenylated phenylpropanoids such as drupanin, baccharin, and artepillin C and flavonoids such as kaempferol and kaempferide^{2, 3}. BGP is used in traditional drugs or supplements because it has a wide range of beneficial biological activities including antibiotic, antivirus, anti-

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oxidative, and anti-inflammatory effects^{2, 3}. Previous studies have shown that EEBGP and its major ingredient, artepillin C, directly eliminate reactive oxygen species (ROS) as radical scavengers and induce the activity of antioxidative enzymes *in vitro*^{4–6}. Oral administration of EEBGP prevented hepatic damage induced by α -naphthyl-isothiocyanate⁷ or water immersion restraint stress in rats⁸. Because EEBGP increased antioxidative enzyme activity in the liver and reduced lipid peroxide levels in the blood, the hepatoprotective effect was thought attributable to the suppression of oxidative stress^{7, 8}.

Acetaminophen (*N*-acetyl-*p*-aminophenol; APAP) is a commonly used analgesic and antipyretic agent. It can cause hepatotoxicity, the severity of which can worsen along with a dose increase, in humans and animals. APAP-treated rats and mice are widely used as models of drug-induced liver injury⁹. Most of the APAP absorbed through the small intestine is excreted in urine after conjugation with glucuronic acid or sulfate in the liver. A part of it is metabolized to its active metabolite, *N*-acetyl-*p*-benzoquinoneimine (NAPQI), by cytochrome P450 2E1 (CYP2E1), after which NAPQI is detoxified by conjugation with glutathione and excreted in bile^{10–12}. When a high dose of APAP is taken, a

large amount of NAPQI is formed, causing mitochondrial dysfunction and ROS production, which impairs cellular macromolecules, ultimately resulting in hepatocellular necrosis^{13–15}. Hepatocellular necrosis worsens with a decrease in the activity of antioxidative enzymes such as Mn-superoxide dismutase or glutathione peroxidase 1 in rodents^{16, 17}. In contrast, substances having antioxidative capacity reduce APAP-induced liver injury. In animal experiments, food-derived substances such as extracts of tomato¹⁸, Zingiberaceae species19, 20, grape seed21, and tea polyphenols22 reduced oxidative stress by increasing the activity of antioxidative enzymes and suppressing hepatocellular necrosis induced by APAP or other chemicals. The inflammatory reaction is also related to an increase in hepatocellular necrosis after APAP administration. Kupffer cells, the resident liver macrophages, respond to damage-associated molecular patterns released from necrotic cells and attract inflammatory cells such as monocytes and neutrophils by producing cytokines and chemokines. However, massive activation of an inflammatory response impairs normal hepatocytes, resulting in expansion of the necrosis area23-25. Increased levels of cytokines such as interleukin (IL)-1 β , IL-6, and IL-10 and chemokines such as CC-motif chemokine ligand 2 (CCL2) and CXC-motif chemokine ligand 2 (CXCL2) were observed in the liver after APAP administration²⁶⁻³¹. APAP-induced hepatocellular necrosis in mice decreased when IL-1^β production or signaling^{26–28} or CCL2 signaling²⁹ was disrupted. EEBGP has been shown to suppress the increase in the levels of inflammatory cytokines and chemokines in the blood of lipopolysaccharide-treated mice³². Therefore, EEBGP could be an attractive candidate agent to suppress APAPinduced inflammatory responses and tissue damage. However, to date, the hepatoprotective effect of EEBGP has not been investigated in models of APAP-induced liver injury.

In the present study, first, we investigated whether EEBGP prevents APAP-induced hepatocellular necrosis. Furthermore, to elucidate the underlying protective mechanism, we examined the gene expression level of antioxidative enzymes and inflammatory-related genes in the rat liver after APAP administration and further investigated the correlation between the change in gene expression and the magnitude of the necrosis area.

Materials and Methods

Samples

EEBGP was obtained by extracting a BGP block with 95% ethanol (API Co., Ltd, Gifu, Japan). The extract was dried and mixed with dextrin powder (Matsutani Chemical Industry Co., Ltd., Hyogo, Japan) at a ratio of 2:3, and then the mixed powder was blended into a normal diet (CE-2, CLEA Japan, Inc., Tokyo, Japan) at the rate of 0.3%.

Animals and experimental design

Male Wistar/ST rats (5 weeks old) were purchased from Japan SLC (Shizuoka, Japan) and were maintained under standard conditions at $23 \pm 1^{\circ}$ C in $55 \pm 10\%$ humidity

on a controlled light/dark schedule (light on from 08:00 to 20:00). The animals were adapted to the rearing environment for approximately one week and then divided into control and EEBGP groups. The animals in the control group were given the normal diet, while those on in the EEBGP group were given a 0.3% EEBGP-containing diet, whereby the amount of daily intake of EEBGP was 291 mg/kg, for consecutive 7 days. All animals had free access to tap water. APAP (Nacalai Tesque, Kyoto, Japan) was dissolved in 0.5% (w/v) methylcellulose (Shin-Etsu Chemical, Tokyo, Japan) solution. All animals were administered APAP by oral gavage at 800 mg/kg at 16:00 on the 8th day and fasted until necropsy. The APAP dose was determined according to a previous report in which hepatotoxicity was induced by a single administration of 8 g/day of APAP in humans³³, which is two-fold the daily maximum dose. Thus, the APAP dose in the present study was equivalent to a toxic dose in humans. We previously observed that hepatocellular necrosis developed in rats that were administered 800 mg/kg APAP at 16:00 but not in most rats that were administered the same dose of APAP at 09:00 (unpublished data); this finding is consistent with a report that that shows the daily oscillation of APAP-induced liver injury in mice³⁴. In the present study, animals with centrilobular necrosis were evaluated (control group, n=4; EEBGP group, n=5). Twenty-four hours after APAP administration, the animals were euthanized by exsanguination under isoflurane (Wako Pure Chemical Industries, Osaka, Japan) anesthesia, and blood samples were obtained by drawing blood from an abdominal vein into tubes containing a serum-separating agent (Terumo, Tokyo, Japan). Liver samples were immediately collected thereafter, and part of the left lateral lobe was cut into small pieces and frozen in liquid nitrogen. The samples were then stored at -80°C for gene expression analysis. The rest of the left lateral lobe and the right median lobe were fixed in 10% neutral-buffered formalin (Wako Pure Chemical Industries) and then embedded in paraffin for a histological analysis.

The study was performed in accordance with the laboratory animal welfare regulations of our company based on the Standards Relating to the Care and Management of Laboratory Animals and Relief of Pain (Notice No. 88 of the Ministry of Environment dated April 28, 2006). The protocol of the animal study was approved by the Animal Ethics Committee of API Co., Ltd. (protocol #43-006).

Histopathological analysis

The paraffin-embedded tissue specimens were stained with hematoxylin and eosin (H&E, Wako Pure Chemical Industries) and observed under light microscopy. For quantitatively evaluating APAP-induced liver necrosis, photomicrographs were randomly taken at 50× magnification (8 images for each animal), and the percentage of the necrotic area was determined using the Image J software (Rasband W.S., U.S.NIH, Bethesda, MA, USA).

Biochemical analysis

Blood samples were centrifuged at $2,000 \times g$ for 15 min at 4°C and serum was collected to determine alanine aminotransferase (ALT) and aspartate aminotransferase (AST) levels by using a FUJI DRI-CHEM 7000i (Fujifilm, Tokyo, Japan).

Determination of the expression of antioxidative enzyme genes and inflammation-related genes

The mRNA expression levels of seven antioxidative enzymes genes including superoxide dismutase 1 (Sod1), superoxide dismutase 2 (Sod2), catalase (Cat), glutathione peroxidase 1 (Gpx1), glutathione peroxidase 4 (Gpx4), peroxiredoxin 1 (Prdx1), and glutathione reductase (Gsr) and eight inflammation-related genes including cytokines IL-1 β (II1b), IL-6 (II6), IL-10 (II10), and tumor necrosis factor- α (Tnf); chemokines CCL2 (Ccl2), CCL3 (Ccl3), and CXCL2 (Cxcl2); and nitric oxide synthase 2 (Nos2) were determined. Liver tissues (approximately 50 mg) were homogenized in 1 mL of ISOGEN® II (Nippon Gene, Tokyo, Japan) by using a Polytron® PT1300D homogenizer (Kinematica, Lucerne, Switzerland); then sterilized ultrapure water was added to the homogenates. The homogenates were centrifuged at 12,000 \times g for 15 min at 4°C to collect the supernatants for further analysis. Total RNA was extracted and purified using 4-bromoanisole (Wako Pure Chemical Industries) and 8 mM lithium chloride solution (Nacalai Tesque). cDNA was synthesized from the total extracted RNA with SuperScript[™] III First-Strand Synthesis SuperMix for qRT-PCR (Thermo Fisher Scientific, Waltham, MA, USA) and was used for quantitative polymerase chain reaction (PCR) analyses performed using an Applied Biosystems 7300 Real-Time PCR System (Thermo Fisher Scientific). The PCR conditions were 40 cycles at 95°C for 15 s and 60°C for 1 min. Primers for the genes and glyceraldehyde-3-phosphate dehydrogenase (Gapdh) gene were designed using Primer-BLAST (Table 1; https://www.ncbi.nlm.nih.gov/tools/primer-blast/) based on the sequences obtained from the NCBI

Table 1. PCR Primers

Gene name	F or R	Primer sequence	Accession No.
Reference gene			
Gapdh	F	AAGGTGAAGGTCGGAGTCAAC	NM 002046.5
	R	GGGGTCATTGATGGCAACAATA	—
Antioxidative enz	zyme gene	s	
Sod1	F	CAGGATTAACTGAAGGCGAGCAT	NM 017050.1
	R	GCTGGACCGCCATGTTTCTT	
Sod2	F	GCTGGCTTGGCTTCAATAAGG	NM_017051.2
	R	GTGCTCCCACACATCAATCC	
Cat	F	AGCCAGAAGAGAAACCCACAAA	NM_012520.2
	R	CCATTCATGTGCCGATGTCC	
Gpx1	F	CGGGACTACACCGAAATGAATG	NM_030826.4
	R	TCACCTCGCACTTCTCAAACAA	
Gpx4	F	GTGCATCGTCACCAACGTG	NM_017165.3
	R	CGGCTGCAAACTCCTTGATT	
Prdx1	F	TCAGATCCCAAGCGCACCAT	NM_057114.1
	R	AGCGGCCAACAGGAAGATCA	
Gsr	F	GTGACCACGAGGAAGACGAAA	NM_053906.2
	R	CATCTCATCGCAGCCAATCC	
Inflammation-rel	ated genes		
Cytokines			
Il1b	F	TGAGGACCCAAGCACCTTCT	NM_031512.2
	R	TGGGAACATCACACACTAGCA	
Il6	F	GTCAACTCCATCTGCCCTTCA	NM_012589.2
	R	TGTCAACAACATCAGTCCCAAGA	
<i>Il10</i>	F	AGTGGAGCAGGTGAAGAATGA	NM_012854.2
	R	CAGTAGATGCCGGGTGGTT	
Tnf	F	GGTTCCGTCCCTCTCATACAC	NM 012675.3
	R	TCCACATCTCGGATCATGCTTT	
Chemokines			
Ccl2	F	CTCTCTTCCTCCACCACTATGC	NM_031530.1
	R	GTAGTTCTCCAGCCGACTCATT	_
Ccl3	F	ACAAGCGCACCCTCTGTT	NM 013025.2
	R	GAATTTGCCGTCCATAGGAGAA	_
Cxcl2	F	GCGCCCAGACAGAAGTCATA	NM 053647.1
	R	CGAGGCACATCAGGTACGAT	—
Production of inflammatory mediators			
Nos2	F	TCAGGCTTGGGTCTTGTTAGC	NM 012611.3
	R	ATGTCTGTGACTTTGTGCTTCTG	—

F, forward primer; R, reverse primer.

Reference Sequence Database (https://www.ncbi.nlm.nih. gov/refseq/). We normalized the mRNA level of each gene to that of *Gapdh*.

Statistical analysis

Statistical analyses were performed using Ekuseru-Toukei 2012 (Social Survey Research Information, Tokyo, Japan). After examining the homogeneity of variance between the control group and EEBGP group with the F-test, we used Student's *t*-test and Welch's *t*-tests for parametric and nonparametric analyses, respectively. Differences between the two groups were considered significant when the *p*-value was less than 0.05 (p<0.05). Correlations between the necrosis area and the mRNA levels of inflammationrelated genes were determined by calculating Spearman's rank correlation coefficient.

Results

Effects of EEBGP administration on APAP-induced liver injury

Abnormal clinical symptoms and the effect on body weight were not observed in animals administered EEBGP (data not shown). APAP administration induced hepatocellular necrosis with inflammatory cell infiltration extending from the centrilobular to the intermediate zone of the liver in the control group (Fig. 1Aa). The animals in EEBGP group also developed hepatocellular necrosis with inflammatory cell infiltration, but the lesions were limited to a smaller region around the central veins (Fig. 1Ab). The area of necrosis, $28.2 \pm 5.0\%$, significantly decreased in the EEBGP group compared with that in the control group, which was $45.9 \pm 5.3\%$ (p<0.05, Fig. 1B). Consistent with the reduction in the histological damage, serum ALT and AST levels in the EEBGP group were much lower than those in the control group, without statistical significance (Fig. 1C).



Fig. 1. Effect of EEBGP administration on hepatocellular necrosis. A: Representative photographs of the liver after 24 h of APAP 800 mg/kg administration. A-(a): Control group. Centrilobular hepatocellular necrosis with inflammatory cell infiltration is observed. A-(b): EEBGP group. Hepatocellular necrosis with inflammatory cell infiltration is limited to the region around the central veins. CV, central vein; PT, portal triad. Tissue specimens were stained with hematoxylin and eosin (100×). B: Percentage of necrosis area in the liver of the control and EEBGP groups. The vertical bars represent the mean values and standard deviation for each group. **p*<0.05 versus the control group, assessed by Student's *t*-test. C: The serum ALT and AST levels in the EEBGP and control groups. The vertical bars represent the mean values and standard deviation for each group. **p*-aminophenol; ALT, alanine aminotransferase; AST, aspartate aminotransferase

Effects of EEBGP administration on the mRNA expression of antioxidative enzyme genes and inflammation-related genes in the liver

There were no significant differences in the mRNA levels of antioxidant enzyme genes, including Sod1, Sod2, Cat, Gpx1, Gpx4, and Gsr, between the control and EEBGP groups (Fig. 2). On the other hand, EEBGP administration suppressed expression of the inflammation-related genes. The mRNA level of the *Ill0* gene in the EEBGP group significantly decreased compared with that in the control group (p<0.05, Fig. 3). The mRNA level of the *Illb* gene in the EE-BGP group also decreased, but the decrease was not significant (p=0.06, Fig. 3). The mRNA levels of the *Il6* and *Cxcl2* genes in the EEBGP group decreased to about half of those in the control group, but the intergroup differences were not significant (Fig. 3). The mRNA levels of the Tnf, Ccl2, Ccl3, and Nos2 genes were not markedly different between the control and EEBGP groups (Fig. 3). An analysis of the correlation between the percentage of hepatocellular necrosis area and changes in the expression of inflammation-related



Fig. 2. mRNA levels of the antioxidative enzymes in the liver of the control and EEBGP groups. The mRNA levels are normalized to that of *Gapdh* expression. The vertical bars represent the mean values and standard deviation for each group. EEBGP, ethanol extract of Brazilian green propolis

genes revealed that the decreases in the mRNA levels of the *Illb* and *Ill0* genes were significantly related to the decrease in hepatocellular necrosis (p<0.05 and p<0.01, respectively, Fig. 4), and the percentage of hepatocellular necrosis tended to decrease along with the decreases in the mRNA levels of the *Il6* and *Cxcl2* genes (Fig. 4).

Discussion

In APAP-induced liver injury, the APAP metabolite NAPQI causes mitochondrial dysfunction and produces ROS. ROS impair phospholipid membranes and proteins and result in hepatocellular necrosis^{13–15}. In addition, APAPinduced hepatocellular necrosis is aggravated by massive activation of inflammatory reactions, which is caused by cytokine and chemokine production by Kupffer cells in response to hepatocellular necrosis^{23–25}. In the present study, EEBGP administration for 7 days in diet before APAP administration decreased the area of hepatocellular necrosis. It is possible that EEBGP decreased hepatocellular necrosis via the following mechanism: (i) elimination of ROS, (ii) moderation of the inflammatory reaction, and (iii) regulation of the metabolism and excretion of APAP and NAPQI.

A previous study showed elevation of the mRNA and protein levels of antioxidative enzymes and the decrease of the ROS level in RAW294.6 cells following exposure to EEBGP⁶. The mRNA levels of antioxidative enzymes were also upregulated in a mouse model of amyloid A (AA) amyloidosis by a dietary intake of EEBGP for 17 days before and after the induction of amyloidosis³⁵. On the other hand, we could not detect any changes in the mRNA levels of antioxidative enzymes in the present study. In the present study, we focused on a preventive effect of EEBGP, and therefore, the rats were administered EEBGP before APAP administration. In contrast, the mice were continuously given EE-BGP before and after the induction of amyloidosis in the AA amyloidosis model³⁵. The differences in the timing of administration could account for the discrepancy in the effects



Fig. 3. mRNA levels of the inflammation-related genes in the liver of the control and EEBGP groups. A: Cytokines. B: chemokines and nitric oxide synthase 2. The mRNA levels are normalized to that of *Gapdh* expression. The vertical bars represent the mean values and standard deviation for each group. *p<0.05 versus the control group, assessed by Student's *t*-test. EEBGP, ethanol extract of Brazilian green propolis



Fig. 4. Correlation between the percentage of necrosis area and mRNA level of inflammation-related genes in the liver of the EEBGP and control groups. A: *111b*, B: *116*, C: *1110*, and D: *Cxc12* gene. The mRNA levels are normalized to that of *Gapdh* expression. The vertical axis represents the individual mRNA levels, and the horizontal axis represents the individual necrosis area. The R-value (correlation coefficient) and the p-value were assessed by Spearman's rank correlation coefficient. EEBGP, ethanol extract of Brazilian green propolis

of EEBGP on the expression of antioxidative enzymes. In the present study, a decrease in the mRNA levels of pro-inflammatory cytokine and chemokine genes such as II1b, Il6, and Cxcl2 was observed in the liver samples obtained from the EEBGP group. Notably, the decrease in *Illb* expression significantly correlated with the decrease in the percentage of hepatocellular necrosis. It has been clearly demonstrated that the activation of IL-1 α and IL-1 β has an important role in the aggravation of APAP-induced hepatocellular necrosis using anti-IL-1 antibody-treated mice or IL-1 receptor-deficit mice, in which APAP-induced hepatotoxicity was markedly suppressed^{27, 28}. It has also been reported that the suppression of IL-1 β reduced the production of other cytokines and tissue damage caused by inflammation in mice with liver injury³⁶. On the other hand, when inflammation was caused by hepatic damage, anti-inflammatory cytokines as well as pro-inflammatory cytokines were induced^{37, 38}. An increase in the mRNA expression of the Ill0 gene, which is a well-known anti-inflammatory cytokine, after APAP administration was reported in mice^{30, 31}. Kanno et al. showed that mRNA levels of the Ill0 gene in the blood were higher in mice with more severe APAP-induced toxicity³¹; therefore, the decreased *1110* level in the EEBGP group might be largely attributable to the decreased necrosis rather than

a direct effect of EEBGP on *1110* expression in the present study. The significant correlation between the mRNA levels of the *1110* gene and the percentage of the hepatocellular necrosis might support this notion.

In the present study, we detected the gene expression changes in the cytokines and chemokines but not the antioxidative enzymes, which might be attributable to the sampling time after APAP administration. It was previously reported that, after APAP administration, liver glutathione was depleted within 1 h in mice^{39, 40}and that this was rapidly followed by production of reactive oxygen spices⁴⁰⁻⁴². Therefore, it is possible that the antioxidative enzymes were induced earlier than 24 h after APAP administration. On the other hand, it has been shown that the inflammatory reaction continued until 24 h after APAP administration in mice⁴².

A previous report showed that oral administration of an ethanol extract of poplar-derived propolis for seven consecutive days decreased CYP2E1 activity and increased the activity of sulfotransferase and glutathione *S*-transferase, resulting in a decrease in APAP-induced liver injury⁴³. It was also reported that oral administration of caffeic acid phenethyl ester (CAPE), a unique constituent of poplar-derived propolis, for three consecutive days decreased the level of CYP2E1 protein and its enzymatic activity⁴⁴. Although the constituents of propolis vary depending on the vegetation in the region where it is produced and EEBGP does not contain CAPE⁵, we cannot deny the possibility of an effect of EEBGP on drug-metabolic enzymes such as CYP2E1.

In conclusion, dietary administration of EEBGP decreased the area of APAP-induced hepatocellular necrosis in the present study. EEBGP administration did not change the mRNA levels of antioxidative enzyme genes in the liver but decreased the mRNA levels of inflammation-related genes including cytokine and chemokine genes, some of which correlated with the decrease in the necrosis area. In the liver, not only APAP but also ischemia-induced liver injury causes an acute inflammatory reaction, and viral infection, alcohol consumption, or fatty liver cause chronic inflammation, which ultimately progresses to serious liver diseases such as cirrhosis and cancer^{45, 46}. The results of the present study suggest the possibility of a modulating effect of EE-BGP on hepatic inflammation caused by tissue damage, and EEBGP could be a good candidate to suppress acute inflammation and progression of chronic inflammation.

Disclosure of Potential Conflicts of Interest: There are no conflicts of interest to declare.

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